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A dissertation submitted for the degree of
Doctor of Philosophy

Identification Of Yeast Genes Enabling Efficient Oenological Fermentation Under Nitrogen-Limited Conditions

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Declaration

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Abstract

Nitrogen deficiency can often lead to slow or sluggish fermentation, resulting in wine out of specification and at risk of oxidation and microbial contamination. Problems due to nitrogen deficiency can be rectified by optimising grape chemistry (through vineyard fertilization), or more commonly supplementing the fermentation with ammonium salts. An alternative is to use wine yeast that can utilize nitrogen efficiently and complete fermentation more reliably. However, to develop ‘nitrogen efficient’ yeast, it is important to understand how such yeast can utilize nitrogen effectively by identifying genes that influence fermentation performance over a range of nitrogen concentrations. Past research related to the identification of genes influencing nitrogen efficiency under fermentative conditions is largely confined to laboratory yeast.

Investigation of the ~5,000 non-essential genes in yeast is possible through research tools such as deletion libraries (collections of strains, each with a single gene deletion). Several genome-wide studies have successfully used deletion libraries in the auxotrophic background of laboratory yeast to investigate phenotypes in response to exposure to single stress factors associated with fermentation. However, the need to supplement with amino acids to overcome auxotrophies makes quantitative physiological studies in nitrogen limiting conditions impractical. Therefore, in this study, we have used a prototrophic deletion collection in both laboratory and wine yeast backgrounds to identify genes influencing fermentation performance.

Screening (micro-fermentation; 600 µL) of the prototrophic laboratory yeast deletion library (BY4741; 5,372 deletants) and the partial wine yeast library (AWRI1631; 1,844 deletants) for growth and consumption of sugar and nitrogen under limiting (75 mg FAN L⁻¹) and non-limiting nitrogen (450 mg FAN L⁻¹) conditions identified deletants with improved fermentation. To better understand the role of individual genes in fermentation, candidate gene sets from each screen were compared to each other and to other published data sets from genome wide transcriptomic analyses related to fermentation.

Wine yeast deletants that enabled shortened micro-fermentation duration in low nitrogen conditions were further investigated, since the experiment best represented nitrogen deficient grape must associated with problematic fermentation. Fifteen deletants completed fermentation quicker than the wildtype (c.a. a 15-59% time reduction) when tested in larger (100 mL) fermentations. This group of genes were annotated to biological processes including protein modification, transport, metabolism and ubiquitination (*UBC13*, *MMS2*, *UBP7*, *UBI4*, *BRO1*, *TPK2*, *EAR1*, *MRP17*, *MFA2* and *MVB12*), signalling (*MFA2*) and amino acid metabolism (*AAT2*). Among

the genes identified, *MFA2* (mating a-factor), which conferred a 34% decrease in fermentation duration, was further investigated. We were interested to understand how deletion of this mating type gene affected fermentation since a link between these metabolic pathways would be novel.

The 15 strains identified in this study, which were fermentation proficient in a 'wine-like', limited nitrogen condition, provide a basis to better understand how yeast adapt to nitrogen limitation during fermentation. Furthermore, the corresponding genes can be targeted in second generation strain improvement programs, using tools such as CRISPR (yet to be approved by relevant regulatory bodies) to generate nitrogen efficient yeast to reduce the need to supplement low nitrogen fermentations.

List of abbreviations

Abbreviation	Full term
Δ	Gene deletion
$^{\circ}\text{C}$	Degree celsius
3'	Three prime of the nucleic acid sequence
5'	Five prime of the nucleic acid sequence
AAPs	Amino Acid Permeases
APO	Ascomycete Phenotype Ontology
ATP	Adenosine Triphosphate
AUC	Area Under the Curve
Bp	Base pair
CDGJM	Chemically Defined Grape Juice Medium
cDNA	Complementary deoxyribonuclease acid
CO₂	Carbon dioxide
CRISPR	Clustered Regularly Interspaced Short Palindromic Repeats
DAP	Diammonium Phosphate
DNA	Deoxyribonucleic Acid
E	Environment
FAN	Free Amino Nitrogen
FD	Fermentation Duration
G	Genotype
GE	Genotype and environment
GM	Genetically Modified
GO	Gene Ontology
GS	Glutamate Synthetase
HMR	Hidden MAT right
HN	High Nitrogen
HOG	High Osmolarity Glycerol
KEGG	Kyoto Encyclopaedia of Genes and Genomes
LN	Low Nitrogen
LYDL	Laboratory Yeast Deletion Library
MAPK	Mitogen Activated Protein Kinase
N	Nitrogen
NCR	Nitrogen Catabolite Repression
NOPA	o-phthalodialdehyde/N-acetyl-L-cysteine
NREL	Normalized Relative Expression Level
OD	Optical Density
ORF	Open Reading Frame
PCR	Polymerase Chain Reaction
PM	Plasma Membrane

qPCR	Quantitative Polymerase Chain Reaction
QTL	Quantitative Trait Loci
RNA	Ribonucleic Acid
RNA-Seq	RNA sequencing
SD	Standard Deviation
SGD	<i>Saccharomyces</i> Genome Database
SNPs	Single Nucleotide Polymorphisms
SPS	Ssy1-Ptr3-Ssy5
TCA	Tricarboxylic Acid Cycle
WYDL	Wine Yeast Deletion Library
YAN	Yeast Assimilable Nitrogen
YPD	Yeast Extract Peptone Dextrose

CHAPTER 1

Literature review and summary of research aims

1. Introduction

This literature review was written during the first six months of Ph.D. candidature (August 2013), as the Core Component Structured Program (CCSP). The purpose of this literature review is to provide the background material to the project and identify a research topic to address the project aims. For more updated literature, please refer to the introduction sections for the individual thesis chapters.

1.1 Yeast and winemaking

Winemaking dates back to 6000 BC, with the earliest evidence found in Asia Minor (Mesopotamia and the Caucasus). Historically, winemaking is associated with the spread of civilization, initially through the Mediterranean and later in the 15th and 16th century to the New World - the Americas and South Africa (Pretorius, 2000). However, wine microbiology is only a historically recent practice, with Louis Pasteur in 1863 showing that yeast, namely *Saccharomyces cerevisiae*, are responsible for alcoholic fermentation. This knowledge that yeast can convert grape sugars (mainly glucose and fructose) into ethanol, carbon dioxide, and other by-products has contributed significantly to the control of the winemaking process (Pretorius 2000). Today, *S. cerevisiae* is most commonly used in wine fermentations because of its reliable fermentative properties and desirable sensory contribution to wine (Pretorius 2000; Astorga et al. 2007). Yeast affects the aromatic profile through the synthesis of flavour compounds as well as the modification and liberation of those from grape-derived (non-volatile) precursors (Bartowsky et al. 2007).

While grape musts are a complex mixture of nutrients, they are harsh environments for yeast growth and metabolic activity. Grape musts generally contain high concentrations of sugars and are low in pH (Backhus et al. 2001). During the progression of fermentation, yeast cells can be subjected to extreme temperatures, progressive depletion of nutrients and increasing ethanol content, which affects their growth and viability (Pretorius 2000; Mendes-Ferreira et al. 2007a). However, wine yeast have distinct physiological properties that help them adapt to harsh oenological conditions (Rossignol et al. 2003). Although wine yeast are usually able to complete fermentation with nutrients naturally present in grape juice, sometimes this concentration is not sufficient (Ugliano et al. 2009; Pretorius 2000; Barbosa et al. 2009). Complete fermentation is defined as when the residual fermentable sugar in wine reaches <2-4 g L⁻¹ and it is one of the primary objectives for the production of dry table wines (Alexandre and Charpentier 1998). To complete fermentation of the sugars, typically yielding 12 to 15% alcohol, yeast requires high concentrations of nutrients such as nitrogen (Vilanova et al. 2007). However, the concentration of nitrogen in grape juice can also vary widely and sufficient nitrogen is very important for completing

fermentation (Bell and Henschke 2005). *S. cerevisiae* can use a variety of nitrogen sources, however, some are not assimilable (Beltran et al. 2004; Crépin et al. 2012). Nitrogen compounds readily consumed by yeast cells are termed as Yeast Assimilable Nitrogen (YAN) and in grape juice typically comprise ammonium salts, amino acids and peptides (Bell and Henschke 2005, Ugliano et al. 2007).

Insufficient YAN in the grape juice often results in problematic slow or sluggish fermentation if not rectified (Alexandre and Charpentier 1998; Bisson 1999; Taillandier et al. 2007). Grape must nitrogenous content depends on the grape variety, time of harvest, root stock, nitrogen fertilization, berry maturation, vine water status, soil type and fungal infection (Alexandre and Charpentier 1998; Salmon and Barre 1998; Pretorius 2000). Nitrogen present in the grape must range from 60-2400 mg L⁻¹ with the minimum requirement of YAN for a standard fermentation being 120-140 mg L⁻¹ (Alexandre and Charpentier 1998; Conradie 2001; Mendes-Ferreira et al. 2004). Ammonium and amino acids are the primary YAN compounds present in the grape juice, making up to 40% and 51-92%, respectively of the total assimilable nitrogen (Beltran et al. 2004; Crepin et al. 2012). These are readily taken up by yeast, with 50% to 100% consumed during the first 20 hours, depending on the initial amount present (Jiranek et al. 1995a; Salmon and Barre 1998; Carrasco et al. 2003; Vilanova et al. 2007).

Historically, fermentation was performed using indigenous microflora present on grapes, but today commercial starter cultures predominate. Pure culture inoculation usually results in rapid and reliable fermentations that are relatively predictable in flavour and thus quality of wine (Pretorius 2000). Although there are over 200 commercially available starter cultures (Bartowsky et al. 2007), there is still a focus on improving wine yeast strains to enhance the fermentation process and produce a wine of improved quality (Pretorius 2000; Donalies et al. 2008). This is being driven by the need for more reliable strains that are able to tolerate more extreme fermentation conditions (which are becoming common) as well as strains that result in wines with unique flavour and aroma profiles (Pretorius 2000).

Efforts involving development of certain traits of wine yeast, often involve using laboratory strains initially as they are readily genetically modified (Donalies et al. 2008). Laboratory strains are simple models for genetic studies as they are heterothallic, that is they carry mutations that prevent them from switching mating type, are generally haploid or diploid (rather than polyploidy or aneuploidy), and are engineered to have auxotrophies (Astorga et al. 2007; Donalies et al. 2008). Auxotrophy is a condition where the organism is unable to synthesize certain enzymes and requires specific growth factors supplemented in the medium to compensate the nutritional

deficiency (Pronk 2002; Mülleider et al. 2012). Auxotrophic markers are important selection tools used to select against revertants after genetic modifications (Pronk 2002). While laboratory strains are readily manipulable, they lack the phenotypic characters needed to produce quality wine (Astorga et al. 2007; Donalies et al. 2008). It is preferable to use wine yeast strains in such studies, however they are homothallic, are typically diploid, if not polyploid, aneuploid or allopolyploid, and tend to sporulate poorly. Being prototrophic, they require dominant selectable markers such as antibiotic resistance, to be suitable for genetic studies.

Gene deletions are important tools in determining gene function, and in providing an insight into further strain improvement (Birrell et al. 2001; Giaever et al. 2002). Gene functionality can be partially determined by analysing the phenotype of a yeast strain (deletant) carrying a gene deletion (Winzeler et al. 1999; Ando et al. 2006). With recent developments in genetic engineering and the complete sequencing of *S. cerevisiae*, it is possible to easily delete or introduce a gene into the yeast genome (Donalies et al. 2008). To this end, a consortium was set up to construct a complete collection (library) of gene deletions (i.e. the gene knock-out collection) of *S. cerevisiae* by systematically inactivating every single gene (Winzeler et al. 1999; Birrell et al. 2001). This approach has enabled research aimed at understanding the biochemical and physiological functions of an entire biological system.

This Ph.D. project involves the use of yeast deletion libraries to identify genes influencing fermentation performance in different nitrogen conditions, to better understand the process of fermentation, enabling a guided approach to strain optimization of 'industry ready' wine yeast.

1.2 Importance of nitrogen during alcoholic fermentation

During fermentation, YAN compounds present in grape juice are metabolised to supply an intracellular pool of polyamines, amino acids, nucleotide bases and their derivatives. These products are needed for cell biomass formation, protein synthesis, and growth, and the excess (e.g. arginine) is stored in the vacuoles until required (Sablayrolles et al. 1996; Hofman-Bang 1999; Carrasco et al. 2003; Vilanova et al. 2007). However, yeast is unable to assimilate all forms of nitrogen present in grape juice and can differentiate between preferred and non-preferred nitrogen sources. Examples of preferred nitrogen sources are ammonium, asparagine, glutamate, and glutamine. Proline, allantoin and urea are examples of non-preferred nitrogen sources (Hofman-Bang 1999; Crépin et al. 2012).

As a consequence nitrogen availability during fermentation promotes biomass formation and fermentation rate and shortens fermentation duration (Alexandre and Charpentier 1998;

Beltran et al. 2005). When nitrogen is supplemented during fermentation, the timing of addition influences its affect. For instance, nitrogen addition during the cell growth phase enhances biomass formation, whereas addition during early stationary phase stimulates fermentation resulting in shortened fermentation duration, with little increase in biomass (Beltran et al. 2005). Nitrogen additions made after half of the sugar has been utilised can have a reduced effect (Bell and Henschke 2005). In addition to biomass formation and fermentation rate, the concentration and type of nitrogen present during fermentation also influences the production of many volatile and non-volatile compounds. These compounds are responsible for desirable aromas in wine; however, an excessive concentration of these can be detrimental to wine quality (Bely et al. 2003; Vilanova et al. 2007).

1.3 Adverse effects of limited nitrogen

Nitrogen limitation during fermentation can negatively affect yeast growth and metabolism, resulting in reduced sugar catabolism, sluggish or stuck fermentation resulting in wine that may have residual sugars, is prone to spoilage, out of specification and therefore decreased quality (Alexandre and Charpentier 1998). Typically fermentation takes approximately 7-10 days to complete, while slow and sluggish fermentation can exceed a month or more. Excessive fermentation time can be costly since tank space is often in short supply and the extra labor is required

Additionally, nitrogen limitation can lead to the formation of H_2S as part of the sulfate assimilation pathway. During this process H_2S is liberated as an intermediate product formed by the reduction of sulfate or sulphite (Bell and Henschke 2005; Jiranek et al. 1995b). However, in the presence of suitable and sufficient nitrogen sources H_2S is sequestered by O-acetyl homoserine, which is derived from nitrogen metabolism (Ono et al. 1999). On the other hand insufficient nitrogen leads to the accumulation of H_2S inside the cell that later diffuses into the fermenting must producing objectionable 'rotten-egg' odour (Bell and Henschke 2005; Jiranek et al. 1995b). However, in Australia, grape juices have typically low concentrations of nitrogen and winemakers commonly supplement these juices with nitrogen salts often curatively (Alexandre and Charpentier 1998).

1.4 Nitrogen supplementation and side effects

Problems due to nitrogen deficiency can be rectified by optimizing vineyard fertilization or more commonly, supplementing the fermentation medium with ammonium salts (Jiranek et al. 1995a), usually diammonium phosphate (DAP) (Gutiérrez et al. 2013b). These supplementations

are typically carried out empirically without determining the nitrogen requirement of the organism or the nitrogen content of the grape juice (Jiranek et al. 1995a). High nitrogen supplementation contravenes the wine industry's desire to reduce the amount of additives used in the production of wines and the level of DAP supplementation is commonly regulated throughout international markets. Therefore there is a need to adhere to the demands of both regulatory authorities and consumers (Pretorius 2000). Moreover, poorly considered supplementation during fermentation can result in a wine with a high residual nitrogen content after fermentation, thereby encouraging microbial instability and production of undesirable compounds (such as volatile phenols, ethyl carbamate and N-heterocyclic compounds; Bell and Henschke 2005) that reduce the quality of wine (Jiranek et al. 1995a; Taillandier et al. 2007; Gutiérrez et al. 2013b). Therefore it is important to have a sound knowledge of the nitrogen content of the grape juice and the nitrogen requirement of the yeast strain used in fermentation to avoid excessive additions (Gutiérrez et al. 2013b). Knowledge of nitrogen content will render supplementation as an option rather than a routine practice and reduced use of supplements is in line with the industry's aim in producing quality wine (Jiranek et al. 1995a).

1.5 Yeast nitrogen transport and metabolism

Through general and specific transport systems or permeases (membrane proteins) yeast transports from the grape juice into the cell. The high capacity general permeases include Gap1p, Can1p (arginine carriers) and the ammonium transporters (Mep1p, Mep2p, Mep3p). Among the MEP proteins, Mep2p shows the highest affinity for ammonium followed by Mep1p and Mep3p (with lower affinity). Apart from these, there are specific transporters which carry one or a few amino acids, listed in Table 1 (Beltran et al. 2004; Crépin et al. 2012; Ljungdahl and Daignan-Fornier 2012).

Table 1. Plasma membrane-localized transporters of nitrogen substrates

Systematic name	Gene name	Functional description	Substrate specificity
Ammonium permease family			
YGR121C	<i>MEP1</i>	Medium-affinity, high-capacity ammonium permease	Ammonium
YNL142W	<i>MEP2</i>	High-affinity, low-capacity ammonium permease	Ammonium
YPR138C	<i>MEP3</i>	Low-affinity, high-capacity ammonium permease	Ammonium
Amino acid permease family			
YKR039W	<i>GAP1</i>	General, high-capacity, amino acid permease	All L-amino acids, D-amino acids, GABA, peptides, polyamines
YGR055W	<i>MUP1</i>	High-affinity methionine permease	Cys
YBR069C	<i>TAT1</i>	Tyrosine and tryptophan permeases	Val, Thr, His (low-affinity), Leu
YOL020W	<i>TAT2</i>	High-affinity tryptophan permease	Phe, Tyr, Trp, Gly, Ala
YDR508C	<i>GNP1</i>	High-affinity glutamine permeases	Thr, Gln, Ser, Cys, Leu, Met, Asn
YCL025C	<i>AGP1</i>	Broad substrate range, medium capacity permease	Val, Ile, Phe, Met, Ser, Leu, Trp, Met, Cys, Ala
YBR068C	<i>BAP2</i>	Branched-chain amino acid permease	Val, Ile, Phe, Met, Leu, Trp, Met, Cys, Ala
YDR046C	<i>BAP3</i>	Branched-chain amino acid permease	Val, Ile, Phe, Tyr, Leu, Trp, Met, Cys, Ala, Thr
YNL268W	<i>LYP1</i>	Lysine permease	Lys, Met
YGR191W	<i>HIP1</i>	Histidine permease	His
YOR348C	<i>PUT4</i>	High-affinity proline permease	Pro
YNL270C	<i>ALP1</i>	Arginine permease	Arg
YEL063C	<i>CAN1</i>	Arginine permease	Arg
YNL268W	<i>LYP1</i>	Lysine permease	Lys, Met
Other transporters/ permeases			
YLL055W	<i>YCT1</i>	Cysteine transporter	Cys
YHL061C	<i>DUR3</i>	High-affinity polyamine	Urea

Adapted from Ljungdahl and Daignan-Fornier (2012)

These specific and general transport permeases are regulated by different molecular regulatory mechanisms such as Nitrogen Catabolite Repression (NCR) (Beltran et al. 2004; Gutiérrez et al. 2013b) and the SPS sensor (Forsberg and Ljungdahl 2001; Ljungdahl 2009). NCR controls the general (Gap1p, Agp1p) and specific amino acid carriers (Put4p), and the MEP permeases. SPS regulates the synthesis of branched-chain amino acid permeases encoded by *BAP2* and *BAP3*, the high-affinity glutamine transporter Gnp1p, the tyrosine and tryptophan carriers Tat1p and Tat2p, the dicarboxylic amino acid permease Dip5p, and the high-affinity methionine permease Mup1p. To a lesser extent and with lower efficiency, these permeases can also catalyze the importation of several other amino acids (Crépin et al. 2012).

1.5.1 Nitrogen Catabolite Repression (NCR)

Different mechanisms regulate the process of nitrogen transport into the yeast cells. One of the primary mechanism is NCR, which represses the transcription of genes that encode for permeases needed in the uptake and catabolism of non-preferred or poorly used nitrogen sources (Pretorius 2000; Beltran et al. 2004; Jiménez-Martí and Del Olmo 2008). Genes such as *URE2* and the four GATA-binding transcription factors, which are *GLN3*, *GAT1* (positive regulators) and *DAL80*, *DEH1/GZF3* (negative regulators; Salmon and Barre 1998; Hofman-Bang 1999). In the presence of preferred nitrogen sources, Gln3p and Gat1p (the positive activators of NCR) remain in the cytoplasm conjugated with Ure3p, which does not enable the NCR-sensitive gene to use the non-preferred nitrogen sources.

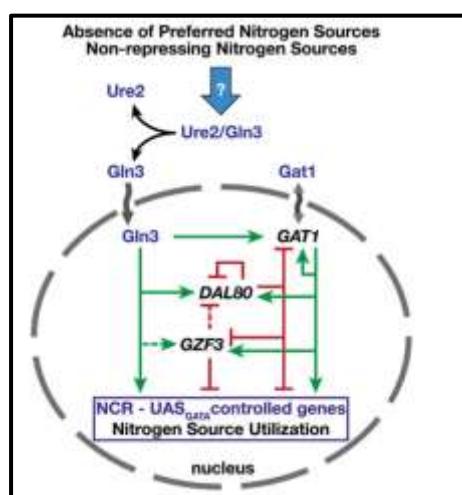


Figure 1. The nitrogen catabolite repression (NCR) circuit. The *URE3* and the four GATA factors: *GLN3* and *GAT1* (positive regulator) and *DAL80* and *GZF3* (negative regulators). All cross regulate each other's expression based on the presence of preferred and non-preferred nitrogen sources, corresponding to their positive and negative role in allowing the expression of NCR-sensitive genes for the uptake of nitrogen sources. Figure and text adapted from Ljundahl and Daignan-Fronier (2012).

On the other hand, depletion of preferred nitrogen sources liberates Gln3p from the URe3p/Gln3p complex and transports Gln3p and Gat1p to the nucleus, derepressing the expression of NCR-sensitive genes allowing the cells to uptake non-preferred sources (Jiménez-Martí and Del Olmo 2008; Crépin et al. 2012). Dal80p and Gzf3p act as repressors of NCR-sensitive genes. All the four GATA factors have zinc-finger DNA binding motifs that enable binding to the GATAAG sequence present in the promoter of the NCR-sensitive genes (Georis et al. 2009; Georis et al. 2011). Also, mutation of *URE3* leads to localization of Gln3p to the nucleus and lifts derepression in spite of the availability of preferred sources (Salmon and Barre 1998). However, Gln3p plays a significant role in NCR, as cells lacking *GLN3* were unable to derepress NCR-sensitive genes (Mitchell and Magasanik 1984; Minehart and Magasanik 1991). Therefore all the four GATA factors control each other's regulation (cross-regulation), contributing either positive or negative regulation based on their role (reviewed by Ljungdahl and Daignan-Fornier (2012); Fig.1).

1.5.2 Plasma membrane Ssy1-Ptr3-Ssy5 (SPS) sensor

The second mechanism by which nitrogen transport is regulated is the SPS sensor. The plasma membrane of the yeast cell contains three core components, Ssy1, Ptr3 and Ssy5, which together are known as the SPS sensor. SPS senses extracellular amino acids, inducing the transcription of genes encoding for particular amino acid permeases (AAPs) to import the amino acids into the cell. Ssy1p senses and activates Stp1p and Stp2p (latent cytoplasmic proteins). These proteins enter the nucleus, bind to the permease genes, transcribing and translating into AAPs, which are then targeted to the plasma membrane for the uptake of the particular amino acids (Ljungdahl 2009; Crépin et al. 2012; Fig. 2).

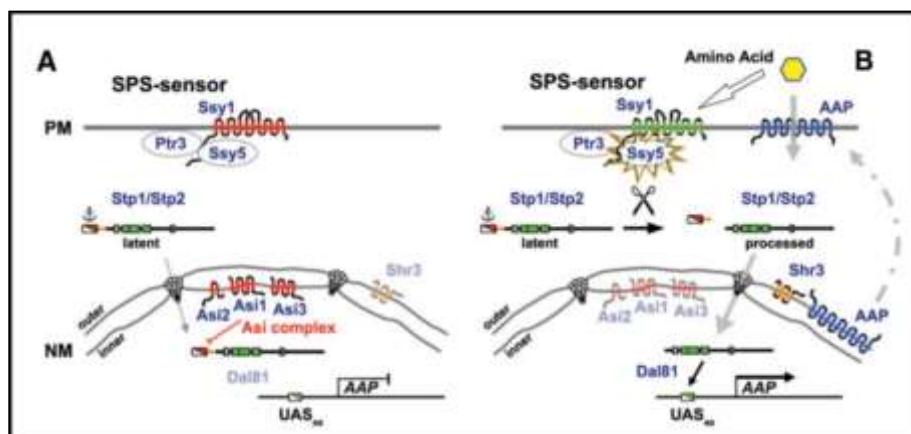


Figure 2. Schematic diagram of the major regulatory steps of the SPS-sensing pathway. (A) The non-induced dormant state in the absence of inducing amino acids: low levels of amino acid permeases (AAPs) are present in the plasma membrane (PM). (B) The induced state in the presence of extracellular amino acids: de-repressed AAP gene expression leads to increased levels of AAPs in the PM and enhanced rates of amino acid uptake (Ljungdahl 2009).

1.5.3 Nitrogen metabolism

Once YAN sources are transported into the cell, the nitrogen is either directly used for protein synthesis or degraded to glutamate and glutamine through the biosynthetic pathways in the presence of ammonium (Hofman-Bang 1999). Glutamate and glutamine are amino acids that provide 85% and 15%, respectively, of total cellular nitrogen (Cooper 1982) and ammonium serves as an amino group donor to produce these compounds. The ammonium needed for the synthesis of glutamate and glutamine is either directly transported into the cell by the MEP transporters or is generated by the degradation of some less preferred nitrogen sources (ter Schure et al. 2000; Magasanik and Kaiser 2002; Fig. 3).

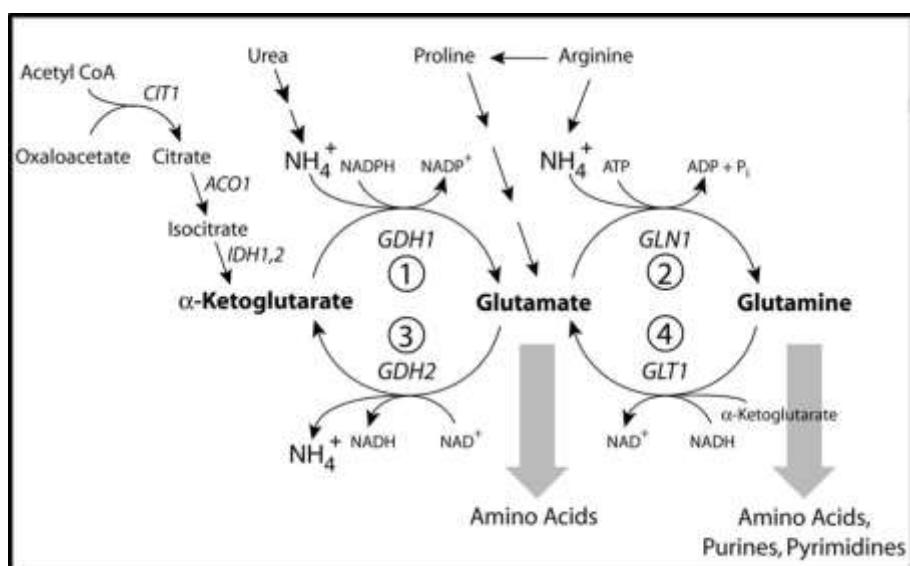


Figure 3. Central pathway for nitrogen metabolism. All cellular nitrogen compounds are synthesized from either glutamate or glutamine. (1) Glutamate is synthesized by ammonia combining with α -ketoglutarate, which is synthesized from acetyl CoA and oxaloacetate in the early steps of the TCA cycle. (2) Glutamine is synthesized by the combination of ammonia with glutamate. (4) Glutamine reacts with α -ketoglutarate, catalysed by glutamate synthetase encoded by *GLT1* resulting in the formation of glutamate (3) Glutamate reacts with NAD-dependent glutamate dehydrogenase (NAD-GDH) encoded by *GDH2* to produce ammonium. The pathways for utilization of other nitrogen sources like urea, proline and arginine, are also shown. The *S. cerevisiae* gene for each of the enzymatic steps is designated in italics (Magasanik and Kaiser 2002).

Yeast converts nitrogen sources into glutamate and glutamine using the Tricarboxylic Acid Cycle (TCA). During fermentation, metabolism of carbon sources takes place through the glycolytic pathway resulting in Acetyl CoA, which enters the TCA cycle to synthesize α -ketoglutarate. This α -ketoglutarate serves as the precursor for the production of glutamate and glutamine in the presence of ammonia. When ammonia is the only source of nitrogen, α -ketoglutarate reacts with ammonia in the presence of the NADP-linked enzyme, glutamate dehydrogenase (NADP-GDH) encoded by the genes *GDH1* and *GDH2* to produce glutamate (Reaction 1). The glutamate combines with ammonia in the presence of the enzyme glutamine synthetase (GS) encoded by

GLN1 to produce glutamine (Reaction 2) (Hofman-Bang 1999; ter Schure et al. 2000; Magasanik and Kaiser 2002). When glutamine is the only source of nitrogen in the medium, Reactions 3 and 4 take place to produce ammonium, where glutamine reacts with α -ketoglutarate catalysed by glutamate synthetase encoded by *GLT1* resulting in the formation of glutamate. This glutamate then reacts with NAD-dependent glutamate dehydrogenase (NAD-GDH) encoded by *GDH2* to produce ammonium (Hofman-Bang 1999; ter Schure et al. 2000; Magasanik and Kaiser 2002).

1.6 Improvement of industrial wine yeast strains

The ability of yeast to assimilate and metabolize compounds such as YAN dramatically influences the capabilities of *S. cerevisiae* to convert grape juice into wine. For this reason, when formulating new strategies to improve wine technology and quality, efficient assimilation by yeast of nutrients such as nitrogen is an important aspect to be considered. Techniques previously used to produce wine of better quality and taste involved the isolation and screening of new yeast from grape and wine samples. This technique has helped in identifying several strains with desirable winemaking characteristics that are used commercially today. However, to improve specific traits of wine yeast, alternative strategies have to be formulated.

Methods such as sexual (mating, sporulation) and parasexual (protoplasmic fusion) breeding, random mutagenesis and genetic engineering are examples of strategies currently used for strain improvement (Cebollero et al. 2007; Donalies et al. 2008). These strain improvement methods have several advantages and disadvantages when compared with one another. The limitation with breeding includes a lack of selection markers and genetic instability of industrial yeast (Cebollero et al. 2007; Donalies et al. 2008). Random mutagenesis is simple, rapid and does not require deeper background knowledge on the gene and its mechanism. However, presence of multiple gene copies in the wine yeast often makes the selection of recessive mutations complicated. Other limitations of random mutagenesis include chances of missing genes contributing to a phenotype and difficulty in studying the entire genome (Winzeler et al. 1999; Giaever et al. 2002). Also, usage of correct dosage of mutagen to produce the desired phenotype and unknown mutations leading to the loss of desirable traits (Pretorius 2000; Steensels et al. 2014) creates further complexities. Conversely, while genetic engineering, allows for controlled modification of genes, it requires a deeper background knowledge of the gene and mechanism that has to be improved (Cebollero et al. 2007). To date, *Saccharomyces cerevisiae* strain ML01 is the first registered recombinant strain, for wine making in the USA. Whereas most wine yeast cannot readily use malic acid, ML01, can complete malolactic fermentation, having the *Oenococcus oeni* gene for malolactic enzyme and the malate permease gene from

Schizosaccharomyces pombe (Husnik et al. 2006). Whilst the use of genetically engineered strains in food production have received only minimal acceptance worldwide, genetic engineering has led to the successful construction of gene deletion libraries. These mutant collections serve as a platform to study gene functionality, which has potential application in understanding which processes to target, regarding the optimisation strategies for future strain improvement (Winzeler et al. 1999; Astorga et al. 2007).

1.7 Using a gene deletion library as a tool to study gene function

Gene deletion libraries can represent a powerful tool for identifying gene function (Winzeler et al. 1999) and thus future target strategies for wine yeast improvement (Donalies et al. 2008). Deletion libraries are usually created by the precise deletion of every single gene or Open Reading Frame (ORF), between its start and stop codon of a given genome. Each gene can be deleted using gene replacement technologies incorporating Polymerase Chain Reaction (PCR) based strategies to generate a cassette targeted at a specific gene and often containing a dominant selectable marker (*KanMX*; Fig. 4). The *KanMX* deletion cassette contains an antibiotic resistance marker coupled on either side with molecular barcodes or tags, made of a 20 base pair oligonucleotide sequences, a “barcode” which allows identification of each deletion mutant (Winzeler et al. 1999; Birrell et al. 2001; Giaever et al. 2002). Based on this strategy, Winzeler and co-workers (1999) successfully constructed a collection of 6925 deletants in a laboratory yeast background. However, as laboratory yeast do not often have desirable winemaking traits, for the specific purpose of physiological studies under winemaking-like conditions, the Australian Wine Research Institute (AWRI) has constructed a partial deletion library in the wine yeast AWRI1631, a *MATa* haplotype of N96 (Borneman et al. 2008; Varela et al. 2012). Unfortunately due to technical difficulties this currently still remains as only a partially complete library.

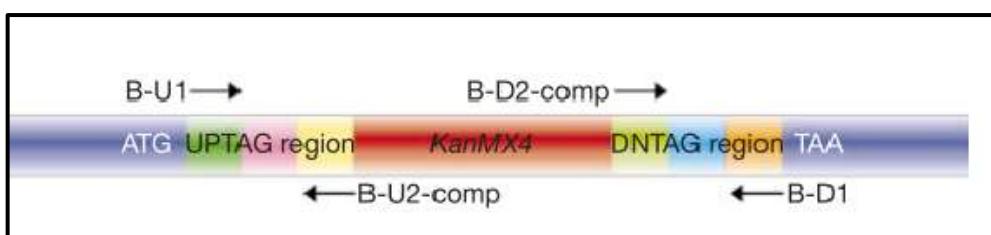


Figure 4. The *KanMX* deletion cassette, containing an antibiotic marker coupled with UPTAG and DNTAG barcode sequences or tags on either side. The B-U1, B-U2-comp, B-D1 and B-D2-comp are the biotin-labelled deletion-specific primers used to amplify the unique UPTAG and DOWNTAG sequence (Giaever et al. 2002).

1.7.1 Previous studies on deletion libraries under fermentation conditions

Several phenotypic studies have investigated gene function during fermentation with the use of deletion mutant collections in a laboratory yeast background. Some examples include the

identification of genes essential for tolerance to ethanol and alcohols (Fujita et al. 2006; van Voorst et al. 2006; Teixeira et al. 2009), high pressure and low temperature (Abe and Minegishi 2008), acetic acid (Mira et al. 2010), oxidative and chemical stress (Tucker and Fields 2004), high glucose (Teixeira et al. 2010), butanol (González-Ramos et al. 2013) and growth under anaerobic conditions (Reiner et al. 2006). A previous study from this laboratory did identify several mutations allowing nitrogen efficient fermentation, yet this was far from comprehensive as a transposon library was utilised and the identity of only a handful of mutants determined (Gardner et al. 2005). To date, there is no published comprehensive study identifying genes related to nitrogen efficiency under fermentative conditions by individually screening every mutant of a deletion library.

1.8 Limitations of current laboratory yeast deletion libraries

Screening for fermentation performance and other physiological studies have successfully used the available laboratory yeast deletion libraries constructed on BY4741, BY4742 and BY4743 backgrounds. However, these possess auxotrophic markers, or mutated genes that encode enzymes involved in the synthesis of amino acids and/or nucleotides (Pronk 2002; Mülleider et al. 2012). These auxotrophic strains are routinely grown either in complex medium (e.g. containing yeast extract and peptone) or defined synthetic media supplemented with sufficient amounts of the appropriate growth factor (Pronk 2002). Several studies showed that such supplementation often produces misleading results that limit data interpretation. Supplementation can block some biological pathways or can even create a combinatorial influence on metabolic networks, which make quantitative comparisons of growth difficult. In addition to interference in metabolic and physiological responses, supplementation can also reduce the life span of auxotrophic cells (Pronk 2002; Canelas et al. 2010; Mülleider et al. 2012).

1.8.1 Requirement for prototrophy and the available prototrophic libraries

To reduce the limitation caused by auxotrophic markers in physiological studies, restoring prototrophy is essential in auxotrophic strains or to generate prototrophic strains carrying a selectable marker to counter select against contamination (Pronk 2002; Canelas et al. 2010; Mülleider et al. 2012). Such strains would have the functional genes required to synthesize enzymes necessary for growth, and therefore should have similar growth kinetics to the wild type strain (Pronk 2002; Mülleider et al. 2012). This would allow gene deletions, which affect processes related to cell growth to be easily identified during a library screen. Recently, Mülleider and co-workers (2012) created a complete collection of gene knock out mutants (deletants) where prototrophy was restored using the pHLUM vector (Fig. 5) in the gene deletion library of BY4741.

The functional genes expressed on the plasmid, complement the mutated genes (auxotrophic markers) on the genome.

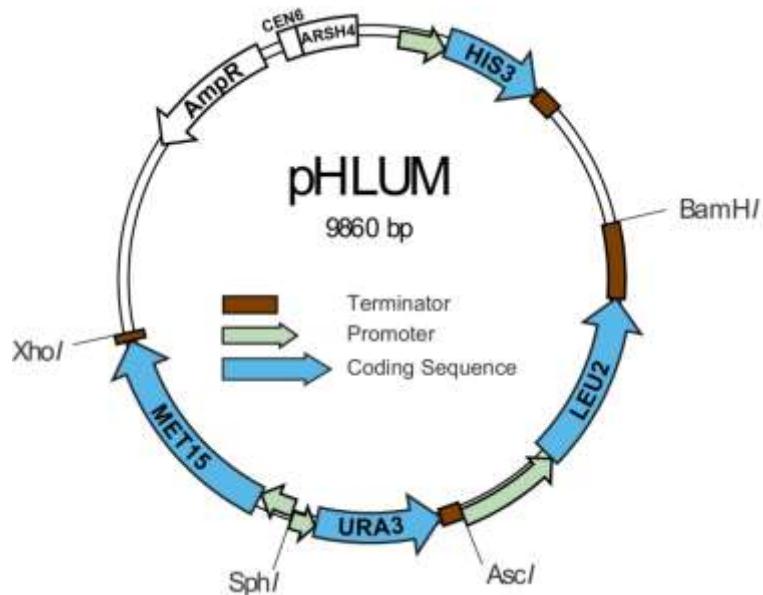


Figure 5. pHLUM vector used to restore prototrophy in the gene deletion library collection of BY4741 (Mulleder et al. 2012)

The construction of a prototrophic gene knock out collection in the haploid wine strain AWRI1631 using similar deletion cassettes to that of the laboratory deletion libraries (Varela et al. 2012) represents a further advancement. This partial library constructed by the AWRI has no known auxotrophic markers and represents 30% of the non-essential genes in the yeast genome. Both these libraries are suitable to study gene function under winemaking-like conditions, allowing for the investigation of fermentation performance under varying nutrient conditions. This Ph.D. project aims to use both these prototrophic mutant collections as screening tools, to identify genes related to nitrogen assimilation during alcoholic fermentation in a synthetic grape juice

1.9 Nitrogen efficient genes and strains of *Saccharomyces cerevisiae*

Past research on yeast transcriptomics aids better understanding on how yeast alter their gene expression pattern based on nitrogen conditions during fermentation. Transcriptomic studies have identified genes that change expression pattern during fermentation under conditions of nitrogen limitation and starvation (Rossignol et al. 2003; Mendes-Ferreira et al. 2007a; Mendes-Ferreira et al. 2007b) and addition of DAP or amino acid to nitrogen-depleted medium (Marks et al. 2003; Mendes-Ferreira et al. 2007b; Jiménez-Martí and Del Olmo 2008). For example, Rossignol et al. 2003 identified significant transcriptional reprogramming of genes (especially the genes associated with stress response) at the beginning of stationary phase as nitrogen became depleted. Likewise, Backhus et al. (2001) used a microarray DNA analysis and compared gene

expression in different nitrogen conditions during fermentation. In high nitrogen, they noticed upregulation of genes involved in the biosynthesis of macromolecular precursors and low nitrogen genes involved in translation and carbon metabolism were upregulated. Mendes and co-worker (2007b) identified 36 genes that were highly expressed in a low nitrogen condition that could be used to monitor nitrogen to avoid stuck/sluggish fermentation arising due to nitrogen depletion. On the other hand, transcriptomic studies undertaken to understand the effect of ammonium or amino acid addition to nitrogen-depleted medium identified that ammonium addition led to an upregulation of genes involved in amino acid synthesis whereas amino acid addition initiated protein synthesis (Jiménez-Martí and Del Olmo 2008). Similarly, Marks et al. (2003) reported a higher expression of genes involved in amino acid biosynthesis on DAP addition.

In an attempt to study yeast gene expression in limited nitrogen followed by nitrogen supplementation, Mendes et al. (2007a) identified variations in the expression of many genes associated with cellular processes such as carbohydrate metabolism, oxidative stress and protein synthesis during fermentation in limited nitrogen. However, refeeding of nitrogen affected the glycolysis and metabolic pathway which allowed yeast to overcome stress and restart fermentation (Mendes et al. 2007a).

Alternatively, in a study comparing gene expression between strains with similar phenotypic and genotypic characters but differing nitrogen requiring strains, Contreras et al. (2012) identified genes related to nitrogen uptake and metabolism. Genes such as *URE2*, *MET31* and *ATO2* and *URA2*, *HXT12*, *STR3* and *IIV5* were identified between strains that differed in ammonium and amino acid consumption, respectively. Therefore past transcriptomic studies provide knowledge on the expression pattern of yeast genes based on the nitrogen conditions during fermentation. Whereas, investigating individual deletants of the non-essential genes will enable identification of genes that are proficient in conditions such as limited nitrogen during fermentation. However, there is only limited research directly related to the identification of nitrogen efficient mutants. Gardner and co-workers (2005) used transposon mutagenesis to construct a library of ~5000 yeast mutants, disrupting individual genes in W303 (*MATa, ura3Δ*). The authors screened for nitrogen efficient mutants in a nitrogen-limited: carbon-excess (75 mg FAN L⁻¹, 200 g glucose L⁻¹) medium, two genes were identified (*NGR1* and *GID7*) that when deleted resulted in a shortening of fermentation duration.

1.10 Bioinformatics as tools in determining gene function

Large data sets, for instance those generated during genome scale library screening require specific analytical tools to deliver objective results. Tools such as Gene Ontology (GO;

<http://www.geneontology.org>) enable identified genes from a phenotypic study to be classified in relation to gene function. GO provides functional and descriptive information on the molecular function, biological process and cellular component using domain-specific ontologies. Domain specific ontologies are vocabularies that are similar in all the organisms. GO are constructed and supported by researchers from many model organism databases and bioinformatics groups through several consortia (Blake and Harris 2002; Blake et al. 2013). An example of a database group that provides information on budding yeast genome is the *Saccharomyces* Genome Database (SGD, <http://www.yeastgenome.org>). SGD maintains information on yeast mutant phenotypes using vocabularies such as the Ascomycete Phenotype Ontology (APO), yeast biochemical pathways, systematic nomenclature and also the reference genome sequence. SGD is constructed by collecting and organizing experimental results from the peer-reviewed literature. Hence, SGD is considered to be an encyclopaedia of the yeast genome that can facilitate experimental design and results (Cherry et al. 2012). This project involves the use of tools such as GO and SGD to identify the function of genes linked to nitrogen efficiency.

1.11 Aims and objectives of the project

Based on this literature review outlining the importance of nitrogen during oenological fermentation and its assimilation in the context of generating nitrogen efficient wine yeast, the following gaps were identified:

1. Past transcriptomic studies provide knowledge on the expression pattern of yeast genes in different nitrogen conditions during fermentation. However, identification of individual nitrogen efficient mutants has been limited to those identified with the use of a transposon mutagenesis system not covering the whole genome and identifying only a few involved genes. Analysis of the fermentation performance of deletion mutants of each non-essential gene is required.
2. Past studies have used tools such as the yeast gene deletion libraries, with which screening of individual deletants of non-essential genes has identified genes required for fermentation related processes. Examples include the identification of genes essential for tolerance to ethanol and other alcohols, high pressure and low temperature, acetic acid, oxidative and chemical stress, high glucose and growth under anaerobic conditions. However, there is no study using the deletion libraries to identify deletants influencing fermentation in sufficient and limiting nitrogen conditions.
3. The laboratory yeast deletion libraries utilized in past studies contained auxotrophic markers, which required medium supplementation to overcome auxotrophy. Recent investigations report that auxotrophic markers can influence biological parameters like growth and aging,

which in turn leads to a bias in physiological and metabolic studies. Also, the need for supplementation does not allow for analysis under limited nitrogen conditions. Prototrophic yeast deletion libraries should be screened for fermentation performance under limited in comparison to sufficient nitrogen conditions. To highlight those genes specifically influencing timely completion of fermentation when nitrogen is scarce.

The main objective of this research is to:

1. Screen the prototrophic laboratory yeast deletion libraries (in the laboratory yeast BY4741 and the wine yeast AWRI1631) to identify genes influencing fermentation duration in a chemically defined grape juice medium containing either sufficient or limiting nitrogen (Chapter 1).
2. Compare and further investigate (using Gene Ontology tools and past literature) identified genes from screening experiments between the two libraries, in different nitrogen conditions and to published related data sets to understand how the gene deletants confer fermentation efficiency (Chapter 2 and 3).
3. Further characterize genes found here to be associated with fermentation efficiency in limited nitrogen conditions to understand how deletion enables fermentation proficiency in low nitrogen wine-like juice (Chapter 3 and 4).

CHAPTER 2

**Identification of yeast genes related to fermentation efficiency in limited
and sufficient nitrogen**

2.1 Introduction

Yeast requires sufficient macro-nutrients such as nitrogen to complete fermentation. According to previous reports, the minimum requirement ranges from 120-140 mg L⁻¹ (Bely et al. 1990; Cramer et al. 2002; Martínez-Moreno et al. 2012) and 200-267 mg L⁻¹ (Mendes-Ferreira et al. 2004). However, nitrogen in grape juice is often limited and requires supplementation to prevent problems such as sluggish fermentation and, in extreme cases, fermentation arrest (Alexandre and Charpentier 1998; Bell and Henschke 2005). Commercial strains differ in their nitrogen requirement (Barbosa et al. 2009; Gutierrez et al. 2012; Martínez-Moreno et al. 2012; Brice et al. 2013; Gutiérrez et al. 2013a), thus yeast with lower nitrogen requirements are more likely to withstand nitrogen deficiency or starvation, thereby utilising the available nitrogen for growth and metabolism, needed to complete fermentation (Gardner et al. 2002; Brice et al. 2013). Such strains are therefore useful in enabling fermentation completion in nitrogen deficient grape must.

The differences in nitrogen requirement may be due to strains possessing high genetic diversity due to a large number of allelic variants (Salinas et al. 2016). Previous QTL analyses have identified some of the genetic basis of these variations related to nitrogen consumption during fermentation. Brice et al. (2014) identified polymorphism in genes (*GCN1*, *MDS3*, *ARG81* and *BIO3*) involved in nitrogen metabolism and signalling using bi-parental crosses. Similarly, Jara et al. (2014) identified variations in *GLT1*, *ASI1* and *GDH2* genes (involved in nitrogen signalling) in strains from a different lineage (i.e. wine and sake). Although classical QTL studies involving bi-parental crosses have mapped traits to loci with large phenotypic effects, these have failed to identify loci contributing to minor phenotypic changes (Bloom et al. 2015; Cubillos et al. 2017). Fulfilling this gap, a recent QTL study used multi-parental cross mapping (Cubillos et al. 2013) together with RNA-Seq to identify variations in genes involved in the major metabolic pathways (i.e. *ARO1*, biosynthesis of aromatic amino acid precursors; *PDC1*, pyruvate decarboxylase; *CPS1*, permease involved in degradation of protein into amino acids; *ASI2*, SPS signalling; *LYP1*, lysine permease and *ALP1*, arginine transporter) that are the basis for strains that are better adapted to different nitrogen conditions during fermentation (Cubillos et al. 2017). While commercially available strains are robust, the ability to complete fermentation in nitrogen deficient juices remains a concern to winemakers. To date, research has focused on improving utilisation of non-preferred nitrogen such as proline (Poole et al. 2009; Long 2014) and arginine (Martin et al. 2003) or increasing nitrogen efficiency, measured by shortened fermentation duration when nitrogen is limiting (Gardner et al. 2002; Zhang 2014). Although some genes related to nitrogen

utilisation and efficiency have been identified (Gardner et al. 2002; Zhang 2014), the knowledge underlying these processes remains limited, with other genes yet to be identified.

Deletion libraries provide the means to determine other genes related to nitrogen efficiency through high-throughput screening of fermentation performance under limited nitrogen conditions. Whilst commercially available gene deletion libraries in haploid or diploid laboratory yeast represent most of the yeast genome, a drawback regarding fermentation studies is that they require supplementation because of the presence of auxotrophic markers (Winzeler et al. 1999). Regardless, they have been successfully employed to study growth sensitivity to single fermentation-related stresses. Examples include tolerance to ethanol (Fujita et al. 2006; van Voorst et al. 2006; Teixeira et al. 2009), acetic acid (Mira et al. 2010), high glucose (Teixeira et al. 2010), growth under anaerobic conditions (Reiner et al. 2006) and adaptation to low temperature (Salvadó et al. 2016). Similarly, work by Walker et al. (2014) identified genes essential to successful completion of fermentation when supplied with high sugar (200 g L⁻¹) and sufficient nitrogen (450 mg L⁻¹) in juice-like conditions which is valuable. However, these auxotrophic strains do not allow fermentation studies under suboptimal nitrogen concentrations. Given that grape must nitrogen content varies widely (50-350 mg L⁻¹; Ugliano et al., 2007) and the risk of stuck and sluggish fermentation increases at <150 mg L⁻¹ (Alexandre and Charpentier 1998; Bell and Henschke 2005), it is necessary to investigate the genes affecting fermentation under both high and low nitrogen conditions.

Comparative transcriptomic analysis has allowed a better understanding of how yeast differs in their physiological response to nitrogen composition/content during fermentation. Several studies using different nitrogen requiring yeast (both laboratory and wine strains) grown under Low Nitrogen (LN) and High Nitrogen (HN) conditions have alluded to the genetic basis of their adaptation to these circumstances. For example, studies include: in HN, Barbosa et al. (2015) (670 mg L⁻¹) and Contreras et al. (2012) (518 mg L⁻¹) and in LN, Barbosa et al. (2015) (67 mg L⁻¹), Mendes et al. (2007b) (66 mg L⁻¹) and Brice et al. (2013) (100 mg L⁻¹). Additionally, Rossignol et al. (2003) used grape must with 300 mg L⁻¹ (considered sufficient nitrogen) to examine gene expression of the entire genome during alcoholic fermentation. Also, exploration of global gene expression in response to various nitrogen sources has been examined (Scherens et al. 2006; Godard et al. 2007), along with the effect of DAP addition during fermentation (Marks et al. 2003) and ammonium and amino acid addition to nitrogen-depleted medium (Jiménez-Martí and Del Olmo 2008) has revealed modifications in gene expression in order to adapt to various nitrogen conditions. Whilst transcriptomic studies have involved prototrophic yeast (mentioned above),

deletion libraries analysed to date have been auxotrophic, which require supplementation. Prototrophic strains are preferable to auxotrophic strains for physiological studies (Canelas et al. 2010), because experimental protocols can be designed to avoid potential metabolic effects associated with supplementation (Mülleder et al. 2012; Pronk 2002), since none is required, a necessity in low nitrogen studies. Hence, this study reports the two prototrophic deletion collections, one in a laboratory background and the other in a wine yeast background in HN (450 mg L⁻¹) and LN (75 mg FAN L⁻¹) to identify genes influencing fermentation performance.

2.2 Materials and methods

Strains and media used in this study

Prototrophic deletion libraries

A Laboratory Yeast Deletion Library (LYDL) containing 5372 deletants in BY4741 (*MATa*; *his3Δ1*; *leu2Δ0*; *met15Δ0*; *ura3Δ0*), where prototrophy was restored by complementation using a CEN6/ARSH4 low copy plasmid, pHLM (Mülleder et al. 2012) was used. In addition a partial Wine Yeast Deletion Library (WYDL) containing 1844 deletants in AWRI1631 (a haplotype of wine strain N96; Borneman et al. 2008) with gene deletions constructed as described in Varela et al. 2012 was used.

Replication of the libraries

The libraries were inoculated into 200 µL YPD medium (1% yeast extract, 2% bacto peptone, 2% D-glucose; Guthrie and Fink 2002) in 96-well microtiter plates, using a 96-pinned replicator, and incubated for 2 days at 28 °C. These cultures were used to inoculate plates for the screening experiment, before addition of glycerol (to 15%) for storage at -80 °C. Liquid handling was done using an 8-channel pipetting CAS-3800 robot (Corbett Robotics, Australia).

A schematic of the subsequent experimental work flow is shown in Fig. 1.

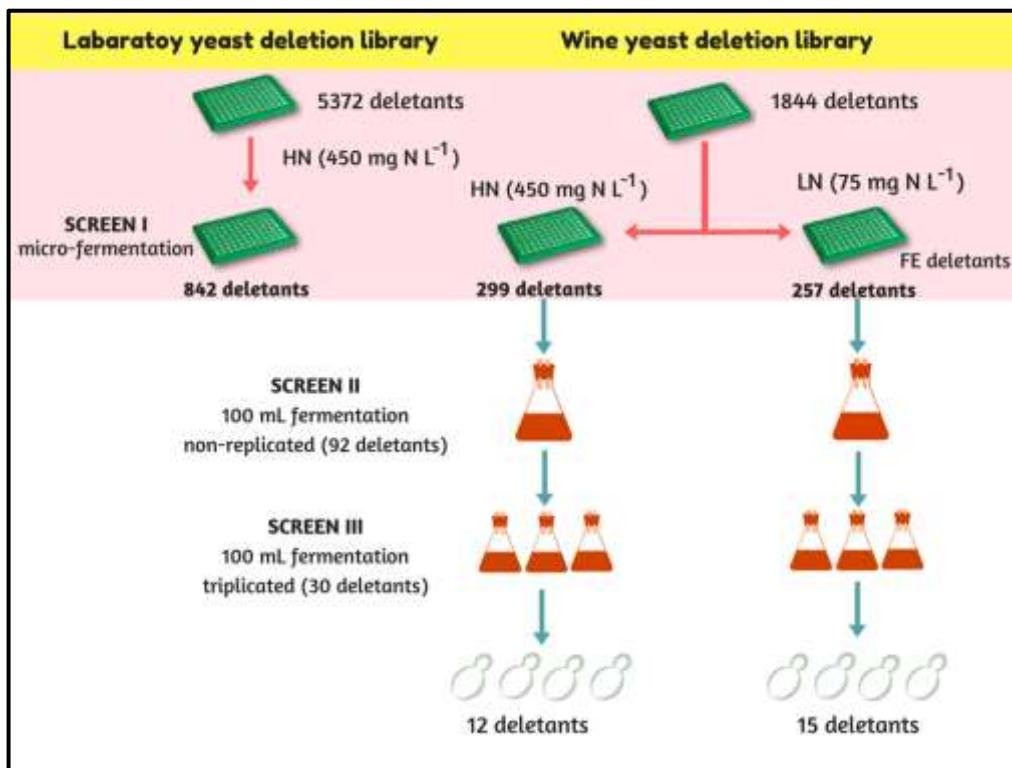


Figure 1. Outline of fermentation screens. Micro-fermentations (600 μL) and laboratory-scale fermentations (100 mL) and the number of gene deletants shortlisted at each stage as fermentation efficient (FE). Evaluation of FE deletants from the WYDL_LN and WYDL_HN screens in 100 mL fermentations identified 15 and 12 FE mutants.

Micro-fermentation (Screen I)

Micro-fermentations were conducted in 600 μL of Chemically Defined Grape Juice Medium (CDGJM; Jiranek et al. 1995a; Walker et al. 2014) in deep 96-well plates. The CDGJM contained 3 g L^{-1} of grape polyphenol extract to mimic red wine fermentations (Walker et al. 2014). The CDGJM was supplemented with different amounts of nitrogen to simulate LN (75 mg L^{-1}) and HN (450 mg L^{-1}) conditions. Four replicates microtiter plate (library) were inoculated with 1% (6 μL) of the overnight YPD culture. A replicate was removed at intervals during fermentation for determination of residual sugar and nitrogen concentration. This sacrificial approach to collect samples over time was used to prevent cross-contamination and experimental artefacts arising from multiple sampling of the same plate during fermentation.

Laboratory-scale fermentations (Screen II and III)

Fermentations (100 mL) involved an initial non-replicated screen of 92 candidates with a 1% (culture v/v) inoculation rate. Thirty strains were chosen from these based on their sugar utilisation and re-evaluated in a triplicated screen using a standardized inoculation rate of 1×10^7 cells mL^{-1} . All the laboratory-scale fermentations were conducted on an automated 96-flask

fermentation platform (Peter et al. 2017). Briefly, the platform was built on a Freedom EVO® robotic work desk (Tecan) and consisted of four 24-position water-jacketed carriers to maintain temperature, and stirrers to continuously agitate the fermentations. Each flask comprised of a 100 mL Schott bottle with an inverted internal fermentation lock and silicon sealed sampling port in the lid. The liquid handling arm of the platform collected ~1 mL of samples at programmed intervals into 96-well plates. Samples were stored at -20 °C for subsequent analysis of residual sugar and nitrogen.

Determination of residual sugar and nitrogen concentration

Residual glucose and fructose were measured enzymatically as described by Boehringer-Mannheim (1989) with modifications to a 96-well format described by Liccioli et al. (2011b). Residual free amino acid was determined by the NOPA (o-phthaldialdehyde/N-acetyl-L-cysteine) procedure (Dukes and Butzke 1998).

Calculation of area under the curve to determine fermentation performance

The residual sugar values measured at each sampling point from the 100 mL fermentations were used to plot fermentation curves for every deletant. The Area Under the Curve (AUC) of each profile was then calculated using the composite trapezoid rule using GraphPad Prism 6 (GraphPad software INC., La Jolla, C.A., USA) as described by Liccioli et al. (2011a). Deletants were considered to be fermentation efficient when the AUC values of the deletants were significantly less than that of the wildtype using a two way ANOVA (Dunnett's multiple comparison test).

Interpretation of screening results and selection of efficient candidates

Deletants were chosen from the initial micro-fermentations (LYDL_HN, WYDL_HN and WYDL_LN) based on their ability to catabolise at least twice the sugar of the wildtype at one or two sequential time points. These 'Fermentation Efficient' (FE) strains were compared between each screen and also with other previously reported gene sets (Marks et al. 2003; Scherens et al. 2006; Mendes-Ferreira et al. 2007b; Contreras et al. 2012; Barbosa et al. 2015). In the WYDL screen, FE deletants in the HN screen, had a sugar content < 16.5 g L⁻¹ for the deletants vs ~33 g L⁻¹ for the wildtype. In the LN screen, the residual sugar was < 24.5 g L⁻¹ (FE deletants) vs ~48 g L⁻¹ (wildtype). However, in the LYDL_HN screen, a more stringent cut-off was set since the library contained more deletants (5372) than the WYDL (1844); selection was based on two time-points. Candidates with <34 g L⁻¹ sugar compared to the wildtype (69 g L⁻¹) at the first time-point, and <11 g L⁻¹ sugar vs wildtype (23 g L⁻¹) at the second time-point, were shortlisted.

Besides these LYDL candidates, 92 best performing strains were chosen from the WYDL based on residual sugar and, in the case of HN fermentation, nitrogen content for laboratory-scale trials. From the WYDL_LN screen, candidates with residual sugar < 48 and 26 g L⁻¹ at time-point 1 and time-point 2 (when the wildtype was ~88 and 48 g L⁻¹) and from the WYDL_HN, candidates with residual sugar < 90 and 40 g L⁻¹ (while the wildtype was ~99 and 33 g L⁻¹) and nitrogen > 200 mg L⁻¹ were chosen for further evaluation.

Identification of gene function and enrichment of the identified gene data sets using computational software tools

The function of candidate genes identified from the micro-fermentation screen was sought using the *Saccharomyces* Genome database (<http://www.yeastgenome.org/>; Cherry et al. 2012). Gene sets were analysed using the Gene Ontology (GO) software GoToolbox and GO-Stats (http://genome.crg.es/GO_ToolBox/; Martin et al. 2004) to determine enrichment of genes based on their GO terms. This computational tool uses a hypergeometric distribution with hypothesis (Bonferroni) correction to calculate P-values (Martin et al. 2004) and the cut-off for significance was set as P < 0.01. The gene set were also analysed using KEGG (Kyoto Encyclopaedia of Genes and Genomes) pathway database (http://www.genome.jp/kegg/tool/map_pathway) (Kanehisa and Goto 2000; Kanehisa et al. 2004; 2016; 2017) to classify genes based on their involvement in distinct cellular pathways.

2.3 Results

Genes whose deletion resulted in shorter fermentation.

Prototrophic yeast deletion libraries in a laboratory yeast (BY4741) and wine yeast (AWRI1631) were used to identify genes influencing fermentation duration in the HN and LN conditions. The wine yeast deletion library (WYDL) containing 1844 deletants was analysed in micro-fermentation in LN (WYDL_LN; 75 mg L⁻¹; Appendix 3, Pg 163) and HN (WYDL_HN; 450 mg L⁻¹; Appendix 2, Pg 149). The Laboratory Yeast Deletion Library (LYDL) containing 5372 deletants was tested only in HN (LYDL_HN; Appendix 1, Pg 108; Fig. 1). The instability of the plasmid pHLUM in some of the BY4741 strains of the LYDL deletants (100 mL fermentations; data not shown), precluded the screen under LN conditions.

Deletants were selected based on their sugar utilisation profile in micro-fermentations in comparison to their wildtype. Those classified as FE (see Materials and Methods) were shortlisted for *in silico* analysis. Micro-fermentation of the LYDL_HN, WYDL_HN and WYDL_LN identified 842, 299 and 257 FE deletants, respectively (Appendix 4-6, Pg 177-187).

Evaluation of selected yeast deletants in laboratory-scale fermentations

Ninety two of the best performing FE candidates from each of the WYDL micro-fermentation (WYDL_HN and WYDL_LN) were re-evaluated in 100 mL fermentations, as the larger scale provided better control of inoculum, anaerobiosis and a larger sample volume. Since one of the aims was to further characterise a set of the most FE deletants, fermentation progress for individual strains was plotted as residual sugar (g L^{-1}) versus time (Fig. 2A and 2B), and the AUC was determined to rank the strains based on their overall performance during fermentation. Thirty-seven deletants were identified from the WYDL_HN screen, having AUC values 85-99% of the wildtype (5975 - 6990 compared to 7006 for the wildtype; Fig. 3A) and 51 deletants from the WYDL_LN had AUC values ranging from 7185 - 9959 (or 72-99%) compared to 9999 (100%) for wildtype (Fig. 3B).

To further define the fermentation performance of the shortlisted candidates, the 30 best performing deletants the WYDL_LN screen (having the smallest AUC values of the wildtype, indicative of FE), were evaluated in triplicate 100 mL fermentations (Fig. 1). In the WYDL_HN, 12 deletants were observed to have an AUC of up to 9% less than the wildtype. The AUC values for the deletants were 4824 – 5284 (91-99%) compared to 5314 for the wildtype (Fig. 3A). The small reduction in fermentation duration can be attributed to the high amount of nitrogen (450 mg L^{-1}), which resulted in fermentation being completed in 68 h. However, in WYDL_LN (75 mg L^{-1}), 15 strains finished fermentation in a significantly shorter time (121 to 249 h compared to 293 for the wildtype; Fig. 3B). The deleted genes of these deletants were *TDA7*, *UBP7*, *AIM26*, *MFA2*, *UBC13*, *UBI4*, *MMS2*, *MRP17*, *MKT1*, *MVB12*, *AAT2*, *BRO1*, *INA1*, *TPK2* and *EAR1* (Peter et al. 2017). Four genes (*AAT2*, *YPR1*, *INA1*, and *MMS2*) of the 15 were also present within the 12 strains highlighted in the WYDL_HN fermentations.

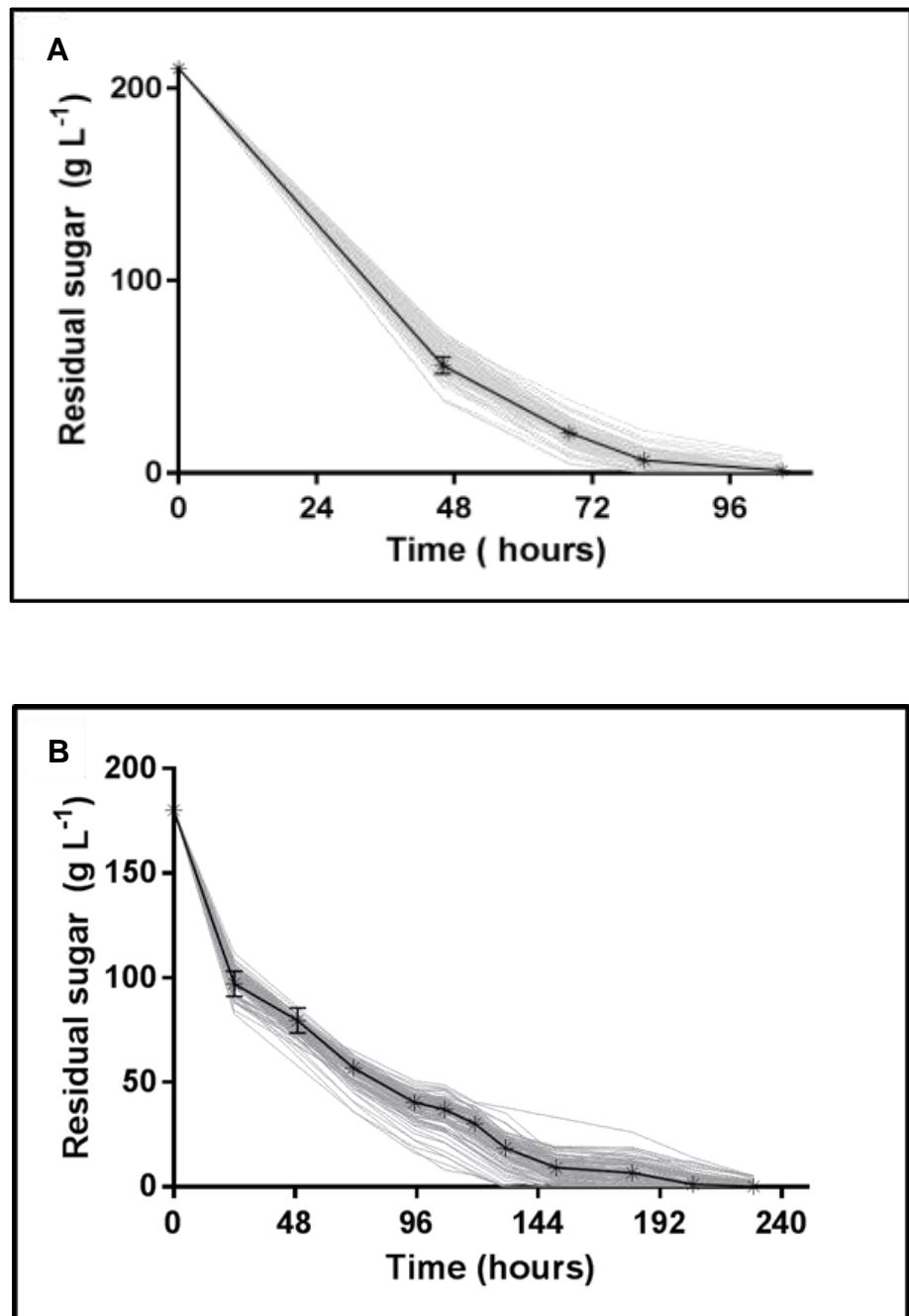


Figure 2. Sugar utilisation of the wildtype, AWRI1631 and 92 shortlisted deletants. Fermentations (100 mL) were conducted at 28 °C in CDGJM with (A) 450 mg L⁻¹ (HN) and (B) 75 mg L⁻¹ (LN). AWRI1631 (black line) 4 replicates and 92 deletants (grey lines) single cultures. The standard deviation of the mean is represented as error bars.

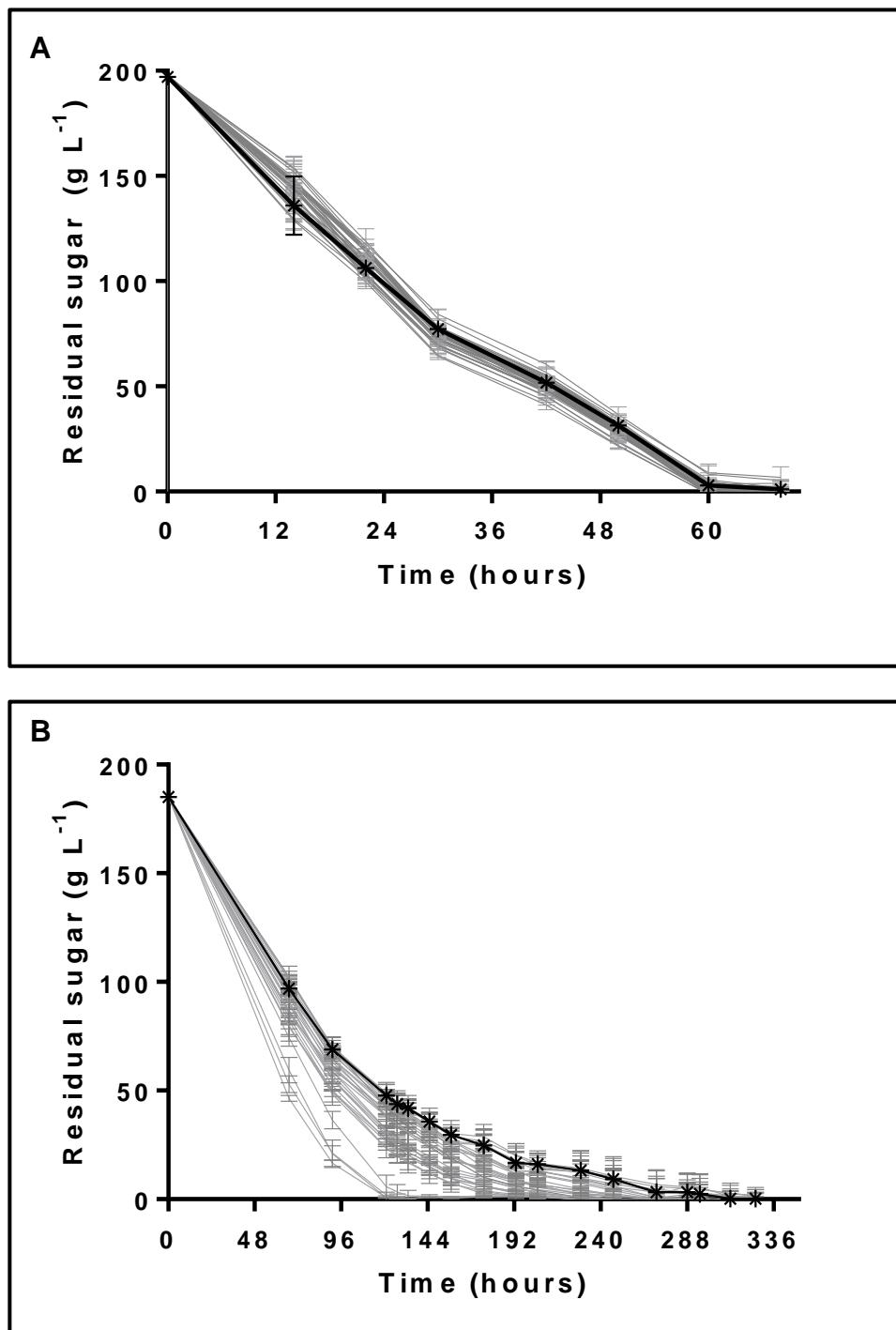


Figure 3. Sugar utilisation profile of the wildtype, AWRI1631 and 30 shortlisted deletants. Fermentations (100 mL) were conducted at 28 °C in CDGJM with (A) 450 mg L^{-1} (HN) and (B) 75 mg L^{-1} (LN). AWRI1631 (black line) and 30 deletants (grey lines). Standard deviation of the mean (triplicates) is represented as error bars.

***In silico* analysis of the FE gene set reveals genes common and unique to each micro-screen**

Comparison of the FE deletants identified in all micro-fermentation screens (Screen I, Fig. 1) - WYDL (HN and LN) and LYDL (HN), revealed genes common and unique to each experimental condition (Fig. 4). Twenty four genes (as deletions) were common to all three data sets (Table 1). Furthermore, 27 genes were presented only in LYDL_HN and WYDL_HN, whilst 87 genes were present in the WYDL_HN and WYDL_LN data sets. Nineteen genes were also represented in both LYDL_HN and WYDL_LN data sets. A large number of genes identified as deletions capable of shortening fermentation, were exclusive to the individual screens: LYDL_HN (748), WYDL_HN (161) and WYDL_LN (127). The gene deletant found in two or all data sets originating from the micro-fermentation screen are presented in Table 1.

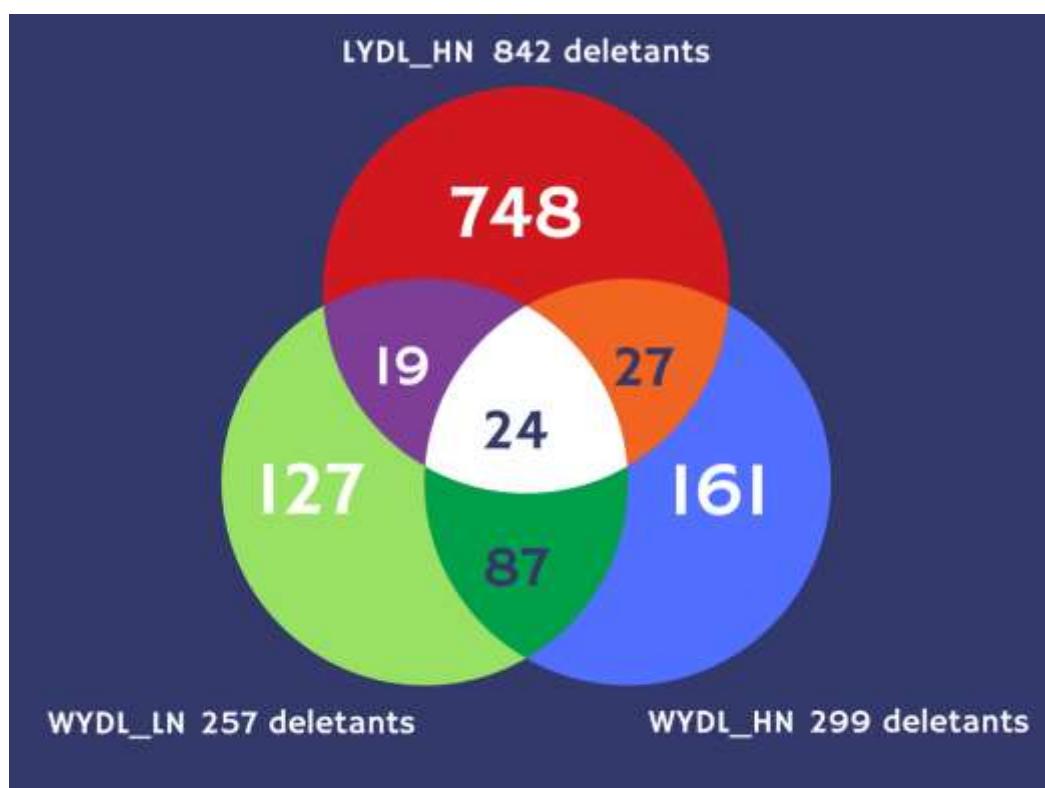


Figure 4. Comparison of the FE gene sets identified in the study. The Venn diagram shows common and unique FE deletants capable of fermenting twice more sugars than the wildtype during micro-fermentation of WYDL (HN and LN), and LYDL (HN).

Table 1. Genes of FE deletants in two or all data sets from the micro-fermentation screening of WYDL (HN and LN) and LYDL (HN)

Library Screen(s)	Identified Genes
WYDL_HN and WYDL_LN (87 genes)	AAT2, AGE2, AIM25, AZF1, BEM2, BOL1, CCP1, CWH41, CYS4, DOA1, DOT5, EAR1, EMC4, AU1, FLC3, FLO11, GGA1, GRX4, HAP5, HCS1, IGO1, INA1, IST1, JID1, KAP122, KSS1, MAL32, MFA2, MIT1, MKT1, MPC1, MRH4, MRP17, NAS2, NIS1, OAC1, OSM1, PBP1, PEP8, PEX31, PGM1, PHB1, PHB2, POM34, PTC2, RMD5, ROM1, SAK1, SAS5, SAY1, SHO1, SIP2, SMA1, SPP1, SPR2, SPT3, SWA2, SWP82, TDA7, TPC1, TPK2, TVP38, TWF1, UBC8, UBP7, UBX3, VMA21, VMA3, VPS68, XPT1, YMC1, YPR1, YPT10, YRF1-3, YER137C, YIL055C, YKL070W, YKL071W, YKR051W, YGR015C, YGR168C, YMR254C, YOR225W, YPR063C, YPR071W, YKL018C-A, YMR105W-A
LYDL_HN and WYDL_HN (27 genes)	AFT2, DLD1, DOG2, EMC5, FMP48, GET1, HSP150, HUT1, IDS2, KEL3, MEI5, MET10, MSH5, OPI3, OYE3, PPT2, RME1, SKG3, SLI1, STB5, TIM21, YDL114W, YGR045C, YGR051C, YOR062C, YPL191C, YPL247C
common to LYDL_HN, WYDL_HN and WYDL_LN (24 genes)	RAD16, TOG1, YLF2, CKA1, MSN4, VPS60, MMS2, PDE1, BBC1, VPS55, ENT4, IOC2, FAR8, RSF1, GIS2, HXT14, SDD3, DBP1, MLH3, RPL7B, YHL008C, YGL082W, YGR066C, YGR237C

Micro-fermentations (600 µL) were conducted in CDGJM as described in Materials and Methods. Deletants resulting in improved fermentation performance (i.e. fermenting twice more sugars than the wildtype) were shortlisted: WYDL_HN (299 genes), WYDL_LN (257 genes) and LYDL_HN (842 genes). Genes represented in 2 or 3 data sets are represented (Fig. 4). Genes denoted in bold represent the genes identified among the 15 genes, whose deletion shortened fermentation duration in CDGJM (LN; 75 mg L⁻¹; 100 mL fermentation; Screen III).

Gene Ontology and KEGG pathway analysis of the FE genes

The gene deletants identified in micro-screens as FE were classified using GO toolbox GO-Stats into specific Gene Ontology categories based on their role in biological processes. GO analysis of the 24 FE genes common to all the three micro-screens and the 27 FE genes represented in LYDL_HN and WYDL_HN highlighted enrichment for 7 different processes (Table 2 and 3). However, GO analysis of the 87 FE genes common to WYDL_HN and WYDL_LN, identified enrichment of genes that were for 34 different processes (Table 4). Kegg pathway analysis of this 87 FE genes highlighted 6 genes (*SHO1, BEM2, KSS1, ROM1, FLO11* and *MFA2*) that are targets of the MAPK (Mitogen Activated Protein Kinase) signalling pathway. Additionally, GO analysis of the 161 and 127 genes unique to WYDL_HN and WYDL_LN, respectively, identified enrichment of genes for process mainly involved in catabolic process and regulation in the HN (Fig. 5A) condition and in the LN conditions the data set was enriched for process such as ubiquination and protein turnover (Fig. 5B)

Table 2. GO enrichment analysis of the genes common to LYDL_HN, WYDL_HN and WYDL_LN.

GO_ID	TERM	P-VALUE	GENES
GO:0009214	cyclic nucleotide catabolic process	3.46E-03	<i>PDE1</i>
GO:0006198	cAMP catabolic process	3.46E-03	<i>PDE1</i>
GO:0019932	second-messenger-mediated signalling	4.56E-03	<i>PDE1, RSF1</i>
GO:0045324	late endosome to vacuole transport	6.48E-03	<i>VPS60, VPS55</i>
GO:0006821	chloride transport	6.90E-03	<i>YHL008C</i>
GO:0051599	response to hydrostatic pressure	6.90E-03	<i>MSN4</i>
GO:0050826	response to freezing	6.90E-03	<i>MSN4</i>

Table 3. GO enrichment analysis of the genes common to WYDL_HN and LYDL_HN.

GO_ID	TERM	P-VALUE	GENES
GO:0015785	UDP-galactose transport	3.78E-03	<i>HUT1</i>
GO:0050896	response to stimulus	4.60E-03	<i>AFT2, SLI1, DOG2, HSP150, STB5, GET1, OYE3, MEI5, MSH5, MET10</i>
GO:0007126	meiosis	4.71E-03	<i>MEI5, IDS2, MSH5, RME1</i>
GO:0051327	M phase of meiotic cell cycle	4.71E-03	<i>MEI5, IDS2, MSH5, RME1</i>
GO:0051321	meiotic cell cycle	4.79E-03	<i>MEI5, IDS2, MSH5, RME1</i>
GO:0042221	response to chemical stimulus	5.04E-03	<i>AFT2, SLI1, STB5, GET1, OYE3, MET10</i>
GO:0042493	response to drug	9.40E-03	<i>SLI1, STB5, MET10</i>

Table 4. GO enrichment analysis of the genes common to WYDL_HN and WYDL_LN.

GO_ID	TERM	P-VALUE	GENES
GO:0032511	late endosome to vacuole transport via multivesicular body sorting pathway	3.00E-05	<i>EAR1, IST1, GGA1, VPS68</i>
GO:0032509	endosome transport via multivesicular body sorting pathway	3.88E-05	<i>EAR1, IST1, GGA1, VPS68</i>
GO:0045861	negative regulation of proteolysis	4.19E-04	<i>PHB2, PHB1</i>
GO:0045324	late endosome to vacuole transport	7.73E-04	<i>EAR1, IST1, GGA1, VPS68</i>
GO:0016197	endosome transport	7.83E-04	<i>EAR1, IST1, PEP8, GGA1, VPS68</i>
GO:0042177	negative regulation of protein catabolic process	8.28E-04	<i>PHB2, PHB1</i>
GO:0009895	negative regulation of catabolic process	8.28E-04	<i>PHB2, PHB1</i>
GO:0006511	ubiquitin-dependent protein catabolic process	1.17E-03	<i>RMD5, DOA1, UBP7, EAR1, SWP82, NAS2, UBC8, GGA1</i>
GO:0043328	protein targeting to vacuole during ubiquitin-dependent protein catabolic process via the MVB pathway	1.36E-03	<i>EAR1, GGA1</i>
GO:0065007	biological regulation	1.46E-03	<i>PTC2, SAK1, NIS1, AZF, RMD5, MKT1, YRF1-3, SIP2, MFA2, SHO1, TPK2, SPP1, KAP122, ROM1, SAS5, TWF1, BEM2, DOT5, SWP82, PHB2, HAP5, PBP1, AGE2, KSS1, VMA21, SPT3, PEP8, AIM25, PHB1, UBC8, GRX4, PGM1, YPT10</i>
GO:0019941	modification-dependent protein catabolic process	1.50E-03	<i>RMD5, DOA1, UBP7, EAR1, SWP82, NAS2, UBX3, UBC8, GGA1</i>
GO:0043632	modification-dependent macromolecule catabolic process	1.81E-03	<i>RMD5, DOA1, UBP7, EAR1, SWP82, NAS2, UBX3, UBC8, GGA1</i>
GO:0050789	regulation of biological process	1.90E-03	<i>PTC2, SAK1, NIS1, AZF1, RMD, MKT1, SIP2, MFA2, SHO1, TPK2, SPP1, ROM1, SAS5, TWF1, BEM2, DOT5, SWP82, PHB2, HAP5, PBP1, AGE2, KSS1, VMA21, SPT3, AIM25, PHB1, UBC8, GRX4, PGM1, YPT10</i>
GO:0051603	proteolysis involved in cellular protein catabolic process	1.91E-03	<i>RMD5, DOA1, UBP7, EAR1, SWP82, NAS2, UBX3, UBC8, GGA1</i>
GO:0040007	growth	1.91E-03	<i>SIP2, SHO1, TPK2, ROM1, BEM2, KSS1, SPT3</i>

GO:0042221	response to chemical stimulus	1.98E-03	<i>PTC2, AZF1, SAY1, MFA2, YKL070W, CCP1, KAP122, CYS4, DOT5, YPR1, KSS1, YKL071W, GRX4</i>
GO:0032269	negative regulation of cellular protein metabolic process	2.03E-03	<i>TWF1, PHB2, PHB1</i>
GO:0050794	regulation of cellular process	2.25E-03	<i>PTC2, SAK1, NIS1, AZF1, RMD5, MKT1, SIP2, MFA2, SHO1, TPK2, SPP1, ROM1, SAS5, TWF1, BEM2, DOT5, SWP82, PHB2, HAP5, PBP1, AGE2, KSS1, VMA21, SPT3, PHB1, UBC8, GRX4, PGM1, YPT10</i>
GO:0007165	signal transduction	2.34E-03	<i>PTC2, SIP2, MFA2, SHO1, TPK2, ROM1, BEM2, AGE2, KSS1, YPT10</i>
GO:0001403	invasive growth in response to glucose limitation	2.40E-03	<i>SIP2, TPK2, KSS1, SPT3</i>
GO:0030162	regulation of proteolysis	2.80E-03	<i>PHB2, PHB1</i>
GO:0051248	negative regulation of protein metabolic process	2.96E-03	<i>TWF1, PHB2, AIM25, PHB1</i>
GO:0007034	vacuolar transport	3.44E-03	<i>EAR1, TVP38, IST1, PEP8, GGA1, VPS68</i>
GO:0051246	regulation of protein metabolic process	4.14E-03	<i>PTC2, SAK1, MKT1, MFA2, TWF1, BEM2, PHB2, PBP1, VMA21, AIM25, PHB1, UBC8, PGM1</i>
GO:0007166	cell surface receptor linked signal transduction	4.32E-03	<i>PTC2, MFA2, SHO1, KSS1</i>
GO:0045721	negative regulation of gluconeogenesis	4.68E-03	<i>RMD5, UBC8</i>
GO:0006508	proteolysis	6.54E-03	<i>RMD5, DOA1, UBP7, EAR1, SWP82, PHB2, NAS2, UBX3, PHB1, UBC8, GGA1</i>
GO:0051181	cofactor transport	6.99E-03	<i>FLC3, TPC1</i>
GO:0006109	regulation of carbohydrate metabolic process	7.43E-03	<i>RMD5, HAP5, UBC8</i>
GO:0030447	filamentous growth	7.44E-03	<i>SIP2, SHO1, TPK2, KSS1, SPT3</i>
GO:0045912	negative regulation of carbohydrate metabolic process	8.29E-03	<i>RMD5, UBC8</i>
GO:0010677	negative regulation of cellular carbohydrate metabolic process	8.29E-03	<i>RMD5, UBC8</i>
GO:0080090	regulation of primary metabolic process	8.51E-03	<i>PTC2, SAK1, AZF1, RMD5, MKT1, MFA2, SPP1, SAS5, TWF1, BEM2, DOT5, SWP82, PHB2, HAP5, PBP1, KSS1, VMA21, SPT3, AIM25, PHB1, UBC8, PGM1</i>
GO:0007154	cell communication	8.57E-03	<i>PTC2, SIP2, MFA2, SHO1, TPK2, ROM1, BEM2, AGE2, KSS1, YPT10</i>

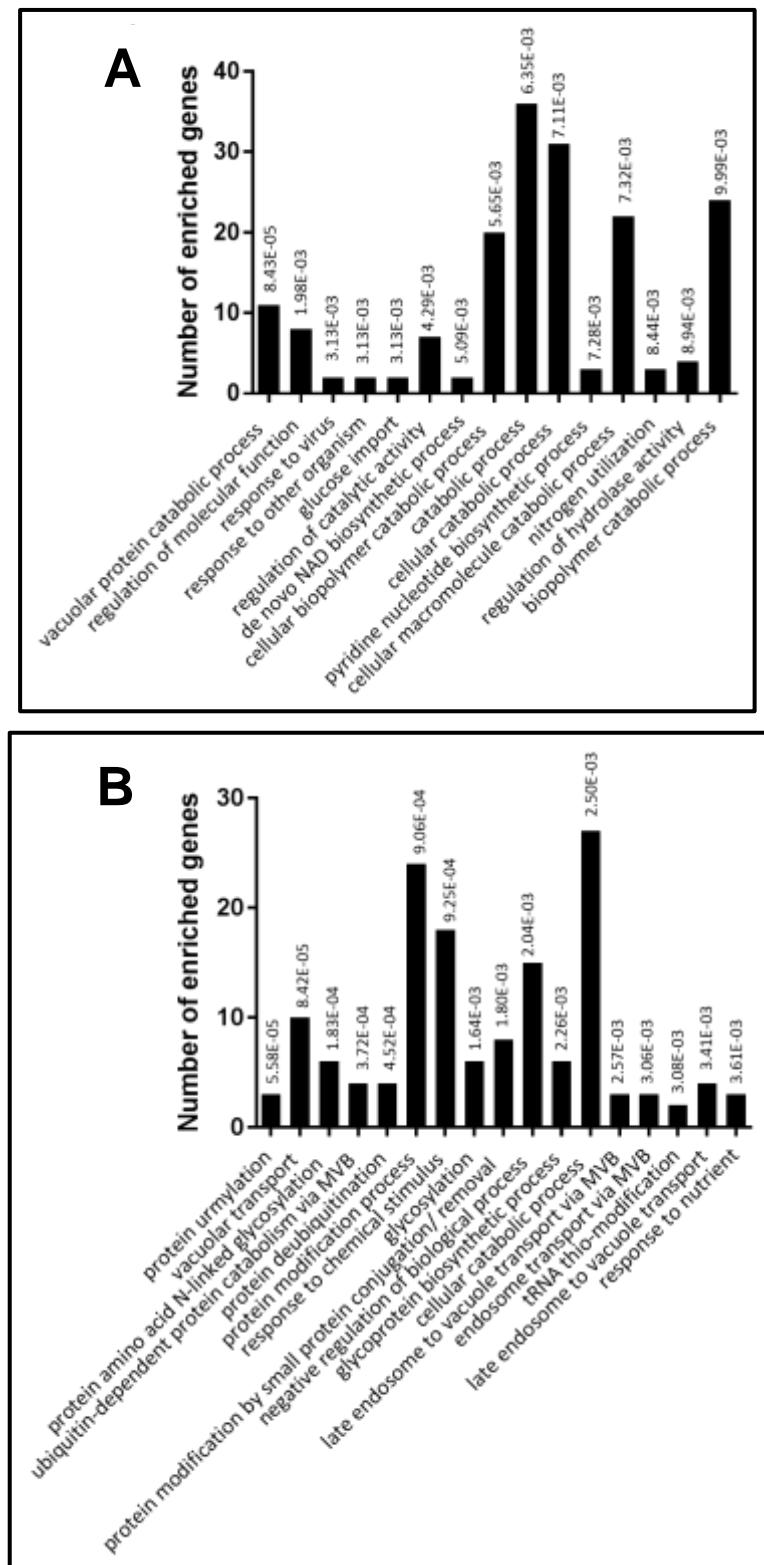


Figure 5. Enrichment of FE genes annotated to specific GO terms from SGD using GO ToolBox GO-Stats. Number of genes found in the indicated GO term which shows significant enrichment under the two growth conditions. **(A)** Genes specific to WYDL_HN. **(B)** Genes specific to WYDL_LN. P-value marked above each bar indicates significance of enrichment.

Genes influencing fermentation efficiency that are unique to each experimental screen and influenced by genotype, environment or both

Large number of FE deletants identified in the micro-fermentation were unique to each screen (Fig. 4): 748 (LYDL_HN), 161 (WYDL_HN) and 127 (WYDL_LN).

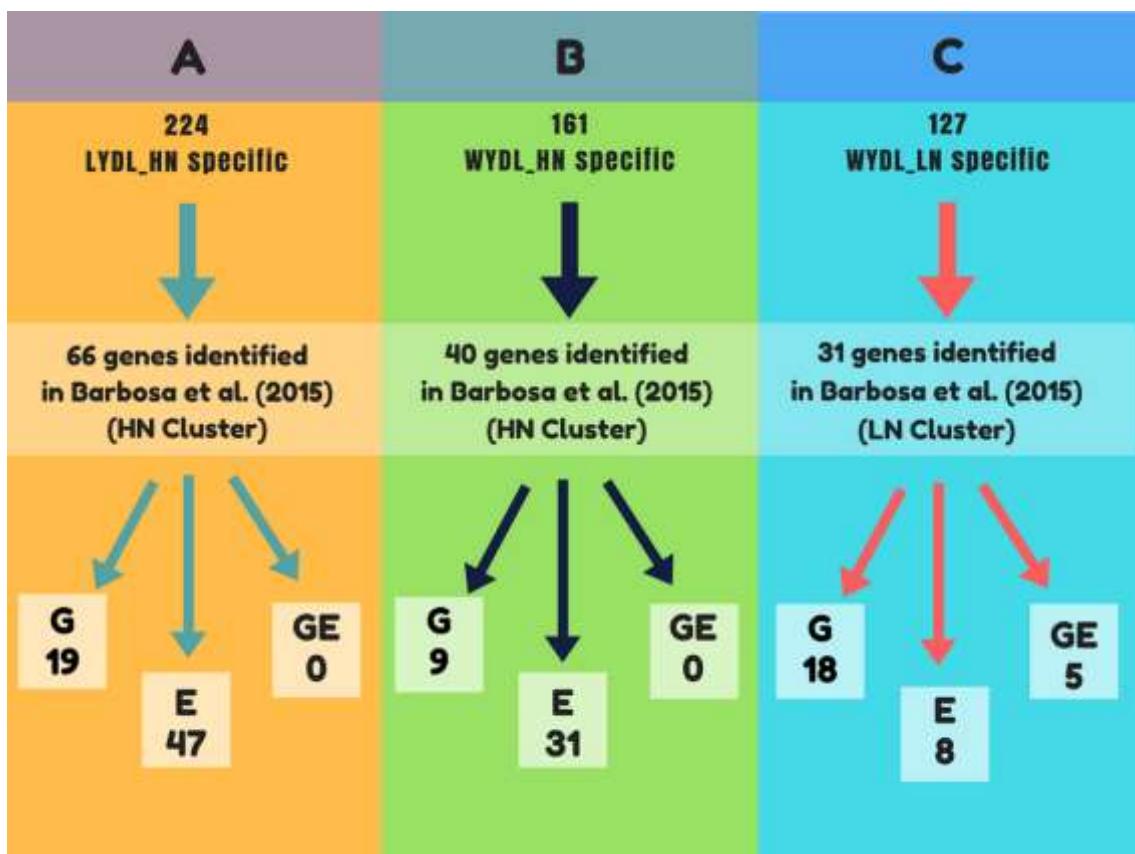


Figure 6. Comparison of FE genes (Screen I) to genes affected by Genotype (G), Environment (E) or both (GE) under HN and LN conditions, as reported by Barbosa et al. (2015). **(A)** LYDL_HN genes identified in the HN condition. **(B)** WYDL_HN genes identified in the HN condition. **(C)** WYDL_LN genes identified in the LN condition.

The identified 161 and 127 FE deletants in the WYDL are likely to be affected by Environment (E; high and low nitrogen). Comparison of the LYDL_HN data set to the WYDL, which represents only 30% of the LYDL, revealed that 73% (548 deletants of the 748 identified as FE and unique to this screen) were not present in the WYDL_HN and the remaining 224 did not exhibit improved fermentation in the WYDL. Therefore the 224 genes are likely to be dependent on Genotype (G). Comparative transcriptomic analysis of wine yeast strains CEG, VL1 and QA23 by Barbosa et al. (2015) identified genes whose expression was influenced by Genotype (G), Environment (E; high and low nitrogen availability) and both Genotype and Environment (GE). Comparison of genes unique to each screen identified in our study, which are likely to be dependent on G and E to clusters reported by Barbosa et al. (2015) as influenced by G, E and GE,

in HN and LN, identified in HN, 19 and 47 (from LYDL) and 9 and 31 (from WYDL) genes to be affected by G and E, respectively, and in LN, 18, 8 and 5 (from WYDL) genes to be affected by G, E and GE, respectively (Fig. 6, Table 5).

Table 5. Fermentation efficient genes influenced by Genotype (G), Environment (E) and interaction between both (GE)

Genes dependency	Genes
Genotype in HN: 19 genes	YBR214W (SDS24), YCL032W (STE50), YEL003W (GIM4), YHL021C (AIM17), YHL022C (SPO11), YHR154W (RTT107), YHR156C (LIN1), YKL010C (UFD4), YJR109C (CPA2), YJR110W (YMR1), YJR111C (PXP2), YLR267W (BOP2), YOL110W (SHR5), YOR197W (MCA1), YOR221C (MCT1), YOR323C (PRO2), YPR040W (TIP41), YPL107W, YPL162C
Environment in HN: 31 genes	YGR061C (ADE6), YDL156W (CMR1), YGL191W (COX13), YIR032C (DAL3), YDL178W (DLD2), YPR023C (EAF3), YDL234C (GYP7), YFR024C-A (LSB3), YGL086W (MAD1), YPR167C (MET16), YFR011C (MIC19), YLR265C (NEJ1), YGL013C (PDR1), YDL093W (PMT5), YGL037C (PNC1), YGR123C (PPT1), YER185W (PUG1), YKL038W (RGT1), YPL208W (RKM1), YER161C (SPT2), YGR129W (SYF2), YPR133W-A (TOM5), YFR007W (YFH7), YHR132C (ECM14), YDL175C (AIR2), YDL243C (AAD4), YFR024C-A (LSB3), YDL144C, YLL053C, YGL242C, YPR148C
Environment in LN: 8 genes	YDR370C (DXO1), YDR379W (RGA2), YGR162W (TIF4631), YGR232W (NAS6), YJR124C (MGR2), YLR052W (IES3), YPR069C (SPE3), YJR124C
GE interaction in LN: 5 genes	YIL015W (BAR1), YIL041W (GVP36), YGR112W (SHY1), YLL039C (UBI4), YPL084W (BRO1)

***In silico* analysis of 257 genes whose deletion confers improved fermentation under limited nitrogen in a wine yeast**

The 257 FE genes (Screen I: WYDL_LN micro-screen), were compared to genes down-regulated under different nitrogen conditions, reported in several transcriptomic studies (Marks et al. 2003; Scherens et al. 2006; Mendes-Ferreira et al. 2007b; Contreras et al. 2012; Barbosa et al. 2015). A total of 39 genes were down-regulated in response to either nitrogen content or composition, or yeast nitrogen requirement (Table 6). Among the 39, 14 and 10 genes were identified to be down-regulated in LN (66 - 67 mg L⁻¹) in the studies conducted by Barbosa et al. (2015) and Mendes-Ferreira et al. (2007b), respectively. Further, 13 genes were down-regulated upon DAP addition following nitrogen starvation (Marks et al. 2003), 6 genes in different nitrogen sources (Scherens et al. 2006) and 3 genes down-regulated in strains of differing nitrogen requirement (Contreras et al. 2012).

Table 6. Thirty nine genes identified in this study that are down-regulated in response to nitrogen content, composition, or yeast requirement

Transcriptomic Study	Down-regulated genes
Low Nitrogen 67 mg L ⁻¹ : 14 genes, Barbosa et al. (2015)	YLR413W (INA1) , YIL085C (KTR7), YLR150W (STM1), YMR265C, YER156C, YLL039C (UBI4) , YNL255C (GIS2), YBL024W (NCL1), YPR063C, YPL198W (RPL7B), YFR032C-A (RPL29), YOR213C (SAS5), YGR283C, YBL028C
Differing nitrogen sources i.e. glutamine, proline, rapamycin: 6 genes, Scherens et al. (2006)	YLR027C (AAT2) , YLR413W (INA1) , YGR155W (CYS4), YDR385W (EFT2), YGR155W, YDR385W
Different nitrogen requirement HN and LN strain: 3 genes, Contreras et al. (2012)	YNL190W, YNL229C (URE2), YJR102C (VPS25)
Nitrogen-starvation/low nitrogen 66 mg L ⁻¹ : 10 genes, Mendes-Ferreira et al. (2007b)	YIL035C (CKA1), YKL120W (OAC1), YJR102C (VPS25), YGR105W (VMA21), YBL024W (NCL1), YLR292C (SEC72), YGR085C (RPL11B), YJR124C, YLR413W (INA1) , YNL084C (END3)
DAP addition 20% w/v DAP addition after 30% sugars were consumed: 13 genes, Marks et al. (2003)	YEL012W (UBC8), YER167W (BCK2), YIL055C, YDR320C (SWA1), YDR386W (MUS81), YGL087C (MMS2) , YGR105W (VMA21), YGR206W (MVB12) , YPL203W (TPK2) , YDL091C (UBX3), YDR392W (SPT3), YGR066C, YPR061C (JID1)

257 FE genes identified in the WDYL_LN micro-fermentation (Fig. 1) were compared with genes highlighted in transcriptomic studies, specifically those down-regulated in response to different nitrogen content, composition or yeast nitrogen requirement. Reported are the genes from previous studies that were also identified in this study as were down-regulated. Genes in bold were shown to have improved fermentation efficiency under LN conditions in 100 mL fermentations (Fig. 3B)

2.4 Discussion

Micro-fermentation of prototrophic deletion libraries to identify genes that confer fermentation efficiency under different nitrogen conditions

Two prototrophic yeast deletion libraries were used to identify genes, whose deletion conferred fermentation efficiency under high (450 mg L⁻¹) and low (75 mg L⁻¹) nitrogen conditions (Fig. 1). The LYDL (5372 deletants), represented most of the non-essential genes in the genome, with prototrophy being restored by plasmid complementation (pHLUM; Mülleder et al. 2012). The WYDL (1844) constructed in a prototrophic wine yeast, AWRI1631 (Borneman et al. 2008; Varela et al. 2012), represented 30% of the LYDL. A preliminary screen was undertaken (Screen I: LYDL_HN, Appendix 1, Pg 108), but because of observed pHLUM plasmid loss from the relatively rich CDGJM, further screen in this medium (i.e. LYDL_LN) was abandoned. Instead, screening

was undertaken using the WDYL under both conditions (Appendix 2 and 3, Pg 149 and 163), with the LYDL_HN data used to verify potential gene candidates, including those absent in the WYDL.

Laboratory-scale fermentation of the shortlisted candidates

Ninety-two of the best performing strains from each of the WYDL micro-screens (HN and LN) were evaluated unreplicated as 100 mL fermentations (Screen II; Fig. 2A and 2B). In the HN condition, fermentation duration was expected to be comparable between deletants and wildtype, as nitrogen was non-limiting. Therefore, FE deletants were instead defined as those that completed fermentation with a smaller AUC and nitrogen consumption. In the LN condition, deletants that were able to complete fermentation in a shorter duration were considered FE. In this way, 30 of the best performing candidates were chosen from each of the conditions (HN and LN) and evaluated in triplicated 100 mL fermentations incorporating a standardised inoculation rate (Screen III; Fig. 3A and 3B). Of the 30, we identified 15 deletants in AWRI1631 (i.e. 16% of those tested) that completed fermentation in a significantly shorter time under LN conditions; ranging from 41 to 78% of the wildtype. These candidates, warranted further investigation and are potential targets for strain improvement (Fig. 3B; Peter et al. 2017).

In screen III of the 30 best-performing wine yeast deletants under HN conditions, fermentation was rapid (67 h) due to the ready availability of nitrogen, therefore, only minor differences existed between fermentations completed prior to the wildtype (Fig. 3A). However, lowering of the fermentation temperature from 28 °C to 22 °C as well as reducing the nitrogen concentration to levels similar to industrial fermentations (i.e. 300-350 mg L⁻¹), was intended to provide better differentiation in fermentation performance between strains.

In silico analysis of the FE gene sets from the micro-fermentations

In micro-fermentation (Screen I), a cut-off based on deletants that were able to ferment twice the sugar of the wildtype at the same time point(s), were used to identify yeast gene deletants, which conferred fermentation efficiency. Deletants with this phenotype totalled 842 (LYDL_HN; Appendix 4, Pg 177), 299 (WYDL_HN; Appendix 5, Pg 184) and 257 (WYDL_LN; Appendix 6, Pg 187). *In silico* comparison of these gene sets identified common and unique genes (Fig. 4). Interestingly, 24 genes were common to all three groups (Table 1), suggesting that these genes are capable of reducing fermentation duration independent of background and nitrogen conditions. The presence of 27 genes in the WYDL_HN and LYDL_HN (Table 1) indicates that these genes influence FE independent of strain background. These genes may be viewed as environment (condition) dependent as they result in FE in both the LYDL_HN and WYDL_HN but

are not present in the WYDL_LN data set (discussed below). However, this idea cannot be confirmed, as the LYDL_LN screen was not undertaken. Eighty seven genes were identified as conferring fermentation efficiency regardless of nitrogen content (WYDL HN and WYDL LN; Table 1). As such these genes seem unlikely to be directly related to nitrogen metabolism but perhaps play a role in tolerance to high sugar conditions. These gene deletions are potential targets for yeast strain improvement specifically for high sugar fermentations, an increasingly common problem of oenological fermentation. Finally, nine genes of this set (Table 1: 87 genes; *AAT2*, *EAR1*, *INA1*, *MFA2*, *MKT1*, *MRP17*, *TDA7*, *UBP7* and *TPK2*) and *MMS2* (Table 1: 24 genes) were amongst the 15 deletants capable of significantly reducing fermentation duration in 100 mL trials (Screen III: WYDL_LN; Peter et al. 2017).

Classification of genes into specific Gene Ontology categories identifies enrichment in the 87 genes common to WYDL_HN and WYDL_LN

GO Toolbox GO-Stats calculates the statistically relevant over-representation (or enrichment) of GO terms associated with a gene set, relative to the genomic distribution of GO terms. An enrichment of genes in a set annotated to a particular GO term implies the importance of that GO cluster to fermentation completion. Classification of genes using this tool, based on their role in a biological process (Martin et al. 2004) revealed an enrichment for genes that were for a process. GO analysis of the 87 FE genes common to the WYDL_HN and WYDL_LN conditions highlighted enrichment for 34 different processes (Table 4). Of the observed enrichment, we identified a set of genes involved in cell communication (*PTC2*, *SIP2*, *MFA2*, *SHO1*, *TPK2*, *ROM1*, *BEM2*, *AGE2*, *KSS1* and *YPT10*; Table 4). Further Kegg pathway analysis of the 87 FE genes highlighted 6 genes (*SHO1*, *BEM2*, *KSS1*, *ROM1*, *FLO11* and *MFA2*), which were associated with cell communication and also target of the MAPK (Mitogen Activated Protein Kinase) signalling pathway (Fig. 7). The MAPK pathway is an intracellular signalling pathway that helps yeast sense and adapt to a changing environment. The MAPK pathway is a cascade of three sequentially activating kinases; a MAPK kinase kinase (MAPKKK), which activates the MAPK kinase (MAPKK), which activates MAPK. *Saccharomyces cerevisiae* has at least five signalling pathways that contain the MAP kinase cascade (Levin and Errede 1995; O'Rourke and Herskowitz 1998). The 6 genes highlighted in the Kegg pathway analysis are involved in the following MAPK pathways; Pheromone response pathway (*MFA2*, mating a-factor and *FUS3*), Cell wall integrity pathway (*BEM2*), High Osmolarity Glycerol (HOG) pathway (*SHO1*) and Filamentous growth pathway (*KSS1* and *FLO11*) (Saito and Tatebayashi 2004; Saito 2010). Further investigation of these genes

and others in the pathway is required to understand how their deletion confers fermentation efficiency.

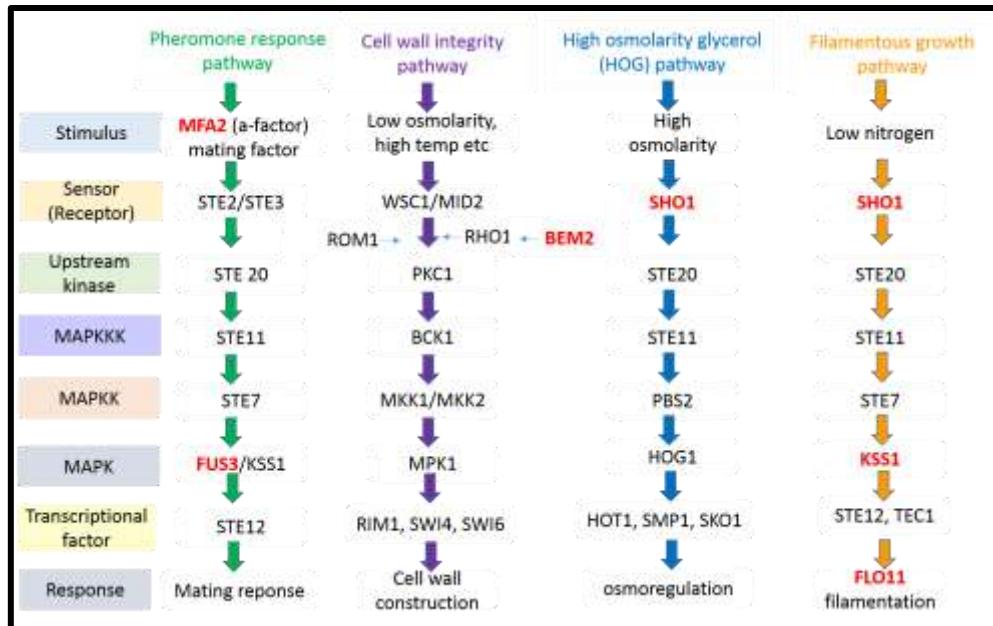


Figure 7. Yeast signalling pathways involving MAPK. Four of the MAPK signalling pathways (Pheromone response pathway, green; Cell wall integrity pathway, purple; High Osmolarity Glycerol (HOG) pathway, blue and Filamentous growth pathway, orange) are targets in the pathway from external stimulation to response by the yeast cells. Genes identified as FE from the WYDL_HN and WYDL_LN are shown in red (adapted from Saito and Tatebayashi 2004).

Genes that confer fermentation efficiency, only in one condition.

GO analysis of the genes specific to WYDL_HN (161) and WYDL_LN (127) identified enrichment of genes for processes associated with cellular catabolic process and regulation in HN and ubiquitination and protein turnover in the LN condition (Fig. 5A and 5B). These findings suggest that under LN conditions, removal of key proteins within the ubiquitin protein degradation pathway may slow turnover of transporters and other proteins critical to maintaining cellular metabolism thereby enabling rapid fermentation. Nitrogen depletion occurs approximately halfway through exponential growth, irrespective of initial nitrogen (Varela et al. 2004). During the late stationary phase, yeast requires energy mainly for cell maintenance in response to diffusion of ions into the cytoplasm due to increasing ethanol content, and for protein turnover due to nitrogen depletion (Varela et al. 2004). In another study, Canelas et al. (2010) proposed that as protein synthesis is an energy consuming process requiring more ATP, strains increase glycolytic flux by driving more ATP and co-factors into protein synthesis. Nevertheless, deletion of genes associated with protein degradation could result in energy saving by reducing protein turnover and extending

the presence of transporter(s)/ion pump(s) in the plasma membrane thereby allowing continued sugar uptake and glycerol efflux. An increased sugar uptake could increase ATP production to support proton and ion homeostasis during ethanol stress (Alexandre et al. 2001; Viana et al. 2012; Fig. 8A and 8B).

Although the start of nitrogen depletion triggers enormous physiological changes such as autophagy, nitrogen recycling, decrease of protein synthesis and the storage of glycogen and trehalose (Rossignol et al. 2003; Brice et al. 2013), the mechanisms involved in sensing external nitrogen and initiating a cellular response to combat nitrogen starvation remain unclear. Interestingly, there are distinct genetic mechanisms by which yeast with differing nitrogen requirement respond to nitrogen limitation/starvation (Brice et al. 2013). In yeast requiring HN, genes related to protein degradation, nitrogen recycling and the stress response are upregulated, whilst in LN requiring yeast, it is those related to synthetic processes such as protein synthesis and RNA processing that are upregulated (Brice et al. 2014). Although the nitrogen requirement profile of AWRI1631 is unknown, it would appear that deletion of genes related to catabolism has a positive but minor effect when yeast are grown with HN, whereas in the LN conditions, dysfunctional ubiquitin-mediated protein turnover is more effective.

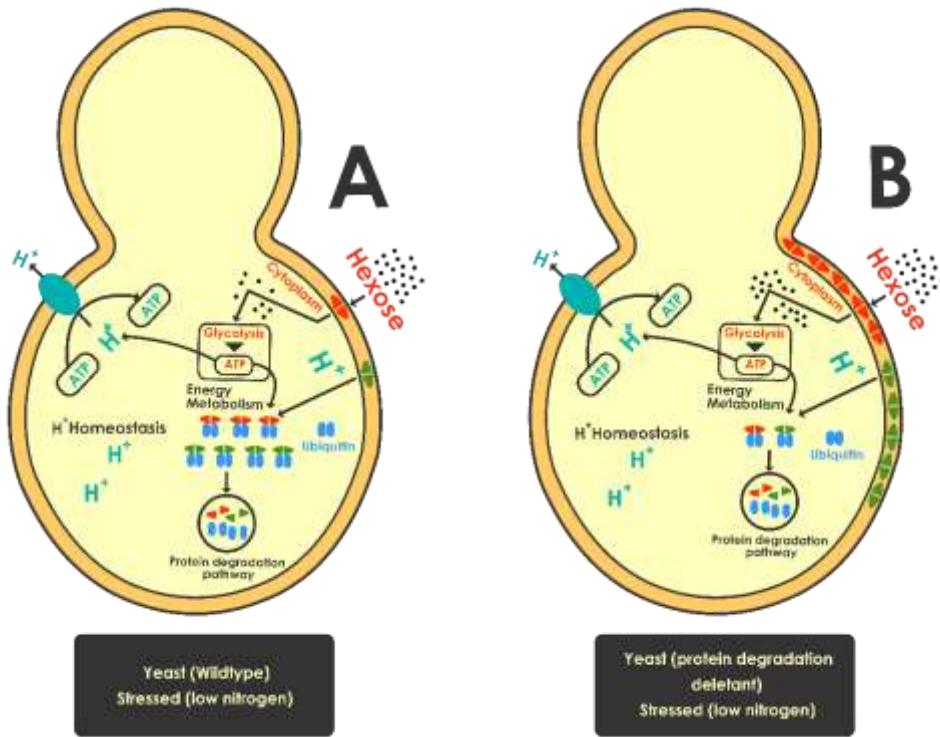


Figure 8. Yeast metabolism (glycolysis, protein degradation and H^+ homeostasis) in LN fermentations. (A) In the wildtype; transporters on the plasma membrane mediate nutrient transfer (hexose transporters , amino acid and nitrogen transporters) across the cell membrane. Transported hexoses are metabolised through the glycolytic pathway producing energy (ATP). When limited nitrogen is available these transporters are down-regulated and degraded, thus fewer are available. The protein degradation pathway aids protein turnover during nitrogen depletion and the H^+ ATPase pumps H^+ ions out of the cell during ethanol stress, which requires energy. (B) It is proposed in deletants of protein degradation pathway genes, protein degradation is reduced enabling more transporters to be present on the plasma membrane to actively transport more sugars resulting in more energy to be used for other activities such as H^+ homeostasis, cell maintenance and fermentation performance.

Barbosa et al. (2015), in a comparative transcriptomic analysis of 3 wine yeast (CEG, VL1 and QA23) in HN and LN fermentation conditions, classified genes into different clusters dependent on whether their pattern of expression was affected by Genotype (G; strain-specific), Environment (E; mechanism-specific) or both Genotype and Environment (GE). The genes specific to each screen (161, WYDL_HN; 127, WYDL_LN and 224, LYDL_HN) were compared to the HN and LN clusters reported by Barbosa et al. (2015) (Fig. 6). Nineteen of the 224 FE genes (LYDL_HN), shown to be FE specifically in the laboratory yeast, can be considered as genotype specific as their deletion did not result in FE in the WYDL_LN (Table 6, Fig. 6). Barbosa et al. (2015) demonstrated differential expression of these 19 genes depending on wine strain. Of the 19 genes, 6 genes (*STE50*, *SDS24*, *LIN1*, *MCA1*, *RTT107* and *AIM17*) represented significant enrichment for processes associated with stress response. The genes from the WYDL_HN (161) and WYDL_LN (127) may be environment (in this case, nitrogen) dependent given the differences

in nitrogen availability in these media. In the two respective screens, 31 and 8 genes were also identified as affected by the environment by Barbosa et al. (2015) (Table 5). Enriched GO terms of these 31 genes were mainly related to negative regulation of cellular processes such as transcription, metabolism, gene expression and biosynthesis (*PDR1*, *SPT2*, *EAF3*, *MAD1*, *PNC1*, and *RGT1*), regulation of glucose transport (*RGT1*) and sulfate reduction (*MET16*). The identified 8 genes highlighted enrichment for processes such as spermidine metabolism (*SPE3*), proteasome regulatory (*NAS6*) and stress granule assembly (*TIF4631*). Similar to Barbosa et al. (2015), in this study the number of genes identified to be affected by environment in HN (161 and 31 genes) was comparatively more than in the LN (127 and 8 genes) condition. In LN fermentations, early depletion of nitrogen allows yeast cells to adjust growth rate to limited nitrogen and maintain homeostasis. In this case, transcriptional levels are more stable in LN when compared to HN fermentations (Barbosa et al. 2015), therefore deleting a gene that is highly active during HN fermentation could possibly affect the sugar uptake or catabolic pathway enabling fermentation proficiency. However, these genes require further investigation, to elucidate the underlying mechanisms behind fermentation efficiency and nitrogen availability.

Interestingly, 5 genes identified as FE (WYDL_LN) were reported by Barbosa and co-workers (2015) as influenced by GE (Table 6). The authors suggest that those genes affected by GE interaction, respond differently to the situation depending on the strain. These genes are referred to as being genetically variable for transcriptional plasticity (Landry et al. 2006; Barbosa et al. 2015). In this study, this gene set was significantly enriched for the processes of ubiquitination, protein turnover (*UBI4* and *BRO1*) and peptide catabolism (*BAR1*).

Comparison of the WYDL_LN genes that result in reduced fermentation duration when deleted with published transcriptomic data sets

The 257 genes identified in the WYDL_LN micro-screen as FE upon deletion, were compared to down-regulated genes from transcriptional studies conducted in various nitrogen conditions (Marks et al. 2003; Scherens et al. 2006; Mendes-Ferreira et al. 2007b; Contreras et al. 2012; Barbosa et al. 2015). 39 genes were matched to the collated transcriptomic data (Table 6). These gene deletions enabling fermentation efficiency in this study have also been shown to be down-regulated in circumstances of different nitrogen conditions, suggesting that the expression of these genes are down-regulated in adverse conditions such as limiting nitrogen. Further investigation of these genes is needed to explore the mechanism that assists in adaptation to LN conditions and improved fermentation performance. Further, these strains could be targets for yeast strain improvement specific for LN conditions.

2.5 Conclusion

Screening of the prototrophic deletion libraries in this study identified interesting genes that confer fermentation efficiency upon deletion in high and low nitrogen conditions. 161 genes (WYDL_HN) were specifically over represented for cellular catabolic process and regulation and 127 genes (WYDL_LN) that were related to ubiquitination and protein turnover. In addition, 87 genes common to the WYDL_HN and WYDL_LN conditions, several genes were enriched for protein degradation. In general, these findings suggest that deletion of genes associated with protein degradation pathways such as transport (via endosome/vacuoles) and catabolism confer improved fermentation phenotype. Deletion of genes such as *BRO1* (Nikko et al. 2003), *MVB1* (Oestreich et al. 2007) and *EAR1* (Léon et al. 2008) have been previously shown to interfere with the degradation of amino acid permeases. This occurs either by hindering movement of these permeases (Gap1p) to the vacuolar lumen (*ear1Δ*, Leon et al. 2008) or recycling Gap1p to the cell membrane without ubiquitination in the presence of ammonium (Nikko et al. 2003). However, it is not known if these and other gene deletions identified in this study affect the degradation of other amino acid permeases or amino acid catabolic enzymes. Both could play a role in the observed enhanced fermentation performance. Furthermore, of the genes whose deletion was identified to confer FE in WYDL_LN, *INA1*, *UBI4*, *MMS2*, *MVB12* and *TPK2* have been shown to be down-regulated in prototrophic wine yeast (Marks et al. 2003; Mendes-Ferreira et al. 2007a; Barbosa et al. 2015) and *INA1* and *AAT2* have been shown to be down-regulated in BY4741 (laboratory strain; Scherens et al. 2006) during fermentation under different N conditions. Likewise, genes such as *BRO1* and *UBI4* are referred to as being genetically variable for transcriptional plasticity (Landry et al. 2006; Barbosa et al. 2015). For instance, these genes respond differently to environment based on the genotype of the strain. Further investigations of these genes are required to better understand how their deletion confers fermentation efficiency. These questions are worth further investigations to extend to the knowledge of fermentation efficiency.

CHAPTER 3

**Use of a wine yeast deletion collection reveals genes that influence
fermentation performance under low nitrogen conditions**

(Submitted for publication)

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Contribution to the Paper	Designed and performed experiments, analysed data and interpreted data and drafted/constructed manuscript	
Overall percentage (%)	70	
Certification:	This paper reports on original research I conducted during the period of my Higher Degree by Research candidature and is not subject to any obligations or contractual agreements with a third party that would constrain its inclusion in this thesis. I am the primary author of this paper.	
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Co-Author Contributions

By signing the Statement of Authorship, each author certifies that:

- I. the candidate's stated contribution to the publication is accurate (as detailed above);
- II. permission is granted for the candidate to include the publication in the thesis; and
- III. the sum of all co-author contributions is equal to 100% less the candidate's stated contribution.

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Contribution to the Paper	Oversaw the experimental design, supervised the research and revised the manuscript.
Signature	Date 10/6/17

Name of Co-Author	Tom A. Lang
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Signature	Date 14/08/17

Name of Co-Author	Vladimir Jiranek
Contribution to the Paper	Oversaw the experimental design, supervised the research, revised the manuscript and submitted the manuscript as corresponding author.
Signature	Date 9.8.17

Use of wine yeast deletion collection reveals genes that influence fermentation performance under low nitrogen conditions

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3.1 Abstract

A deficiency of nitrogenous nutrients in grape juice can cause stuck and sluggish alcoholic fermentation, which has long been a problem in winemaking. Nitrogen requirements vary between wine yeast strains, and the ability of yeast to assimilate nitrogen depends on the nature and concentration of nitrogen present in the medium. In this study a wine yeast gene deletion collection (1844 deletants in the haploid AWRI1631 background) was screened to identify genes whose deletion resulted in a reduction in the time taken to utilise all sugars when grown in a chemically defined grape juice medium supplemented with limited nitrogen (75 mg L⁻¹ as a free amino acid mixture). Through micro-scale and laboratory-scale fermentations 15 deletants were identified that completed fermentation in a shorter time than the wildtype (c.a. 15-59% time reduction). This group of genes was annotated to biological processes including protein modification, transport, metabolism and ubiquitination (*UBC13*, *MMS2*, *UBP7*, *UBI4*, *BRO1*, *TPK2*, *EAR1*, *MRP17*, *MFA2* and *MVB12*), signalling (*MFA2*) and amino acid metabolism (*AAT2*). Deletion of *MFA2*, encoding mating factor-a, resulted in a 55% decrease in fermentation duration. *Mfa2Δ* was chosen for further investigation to understand how this gene deletion conferred fermentation efficiency in limited nitrogen conditions.

3.2 Introduction

To complete fermentation successfully (i.e. utilisation of all fermentable sugars), yeasts require sufficient Yeast Assimilable Nitrogen (YAN). The presence of sufficient YAN supports appropriate cell growth and metabolism (Bell and Henschke 2005; Ugliano et al. 2007). The rate at which a cell grows and maintains its metabolic activity can influence the regulation of other pathways (e.g. sugar metabolism), which in turn affects fermentation kinetics (Alexandre and Charpentier 1998; Bely et al. 1990; Sablayrolles et al. 1996). YAN metabolism also leads to the production of various aroma and flavour compounds, which contribute to the organoleptic properties of the wine (Vilanova et al. 2007). Therefore YAN availability is important in fermentation by significantly influencing cell growth, rate of sugar catabolism and wine flavour (Bell et al. 1979; Henschke and Jiranek 1993; Ugliano et al. 2007).

Although YAN is one of the main nutrient groups needed during fermentation, it is often not present in sufficient amounts in grape must to enable yeast to utilise all sugar. The amount and composition of YAN present in the juice can vary depending on factors such as the grape variety, viticultural management practices, soil, climate and degree of ripeness (Alexandre and Charpentier 1998; Bell and Henschke 2005; Henschke and Jiranek 1993). For example the YAN content in Australian grape juices varies between 50-350 mg L⁻¹ (Ugliano et al. 2007). YAN concentrations below 150 mg L⁻¹ increase the risk of stuck and sluggish fermentation as a consequence of yeast metabolism being affected (Jiranek et al. 1995a; Alexandre and Charpentier 1998; Bell and Henschke 2005). Stuck and sluggish fermentations result in wine left with residual sugar, which is not desirable in certain wine styles (Bisson 1999). Moreover, unfermented sugars can support spoilage organisms (McClellan et al. 1989) producing undesirable aromas and flavours, as well as volatile acidity (Alexandre and Charpentier 1998). The depletion of YAN in juices can also lead to the liberation of excessive H₂S in wine (Jiranek et al. 1995b; Ugliano et al. 2009).

In order to minimise problematic fermentations due to YAN insufficiency, it is common practice for winemakers to supplement fermentations with inorganic ammonium salts. These supplementations are often carried out empirically without determining the nitrogen requirement of the inoculated strain or the nitrogen content of the grape juice (Jiranek et al. 1995a; Torrea et al. 2011). Poorly considered supplementation could result in wine with increased levels of residual nitrogen that encourage microbial instability in the bottled wine, and thus reduce quality (Bell and Henschke 2005; Jiranek et al. 1995a). Also, YAN supplementation plays an important role in modulating the production of both primary metabolites and volatile compounds, which can affect

wine aroma and flavour (Torrea et al. 2011) positively or negatively if not balanced or over threshold (Ugliano et al. 2007).

An alternative paradigm to supplementation is to use yeast strains that are able to efficiently utilise the YAN available in juice in order to complete fermentation. Commercial wine yeasts have different requirements or demands for YAN (Jiranek et al. 1995a; Brice et al. 2014; Gutiérrez et al. 2012; Gutierrez et al. 2013). Yeasts with lower YAN demands are less likely to undergo nitrogen (N) starvation, enabling successful completion of fermentation in YAN-deficient juices (Gardner et al. 2002; Manginot et al. 1998). Therefore identification of genetic determinants conferring N-efficient behaviour has great merit. Previous work by this group utilising a transposon mutant library in a wine yeast background identified genes impacting N-efficiency (Gardner et al. 2005). This study builds on that work by screening a library of ORF deletants constructed in a wine yeast background.

Accordingly, the fermentation capabilities of a partial wine yeast deletion collection were analysed in high-throughput micro-fermentations under limited-N conditions and mutants grouped according to their performance. Those with the shortest fermentation were tested in laboratory-scale cultures to confirm their ability to complete N-limited fermentation quickly. The deleted genes of these mutants are of interest as they represent potential targets for strain improvement. Among the genes highlighted was *MFA2*, encoding mating pheromone a-factor. This is the first report describing the dual function of *MFA2* in mating as well as fermentation.

3.3 Material and methods

Strains and media

The parental prototrophic haploid strain AWRI1631 (*MATa*) was derived from a widely used commercial strain (N96; Borneman et al. 2008). A wine yeast deletion collection containing 1844 mutants was prepared in AWRI1631 by replacement of non-essential genes with the *KanMX* cassette (Wach et al. 1994) as described for a subset of these deletants (Varela et al. 2012). Strains were routinely grown on YPD medium (1% yeast extract, 2% bactopeptone, 2% D-glucose; Guthrie and Fink 2002) with 200 mg L⁻¹ of G418 sulfate for deletant strains and with 2% agar when solid media were required for plating.

Chemically Defined Grape Juice Medium (CDGJM; McBryde et al. 2006) containing 200 g L⁻¹ sugar as equimolar amounts of glucose and fructose with 75 mg L⁻¹ of a mixture of amino acids and ammonium chloride (Henschke and Jiranek 1993) was used for micro-scale and laboratory-scale fermentations. To mimic red grape juice, polyphenol extract powder (Cat: Tppr,

OenoProd, Sarl) dissolved in 1.67 mL g⁻¹ of 100% ethanol, was added to sterile CDGJM at the rate of 3 g L⁻¹ (Walker et al. 2014).

Micro-fermentation of the wine yeast deletion collection in CDGJM

Measurement of cell growth

The wine yeast deletion collection and the wildtype strain, AWRI1631, were inoculated from -80 °C glycerol stocks (2 µL) into 96-well microtiter plates (Corning Costar®; Cat No. CLS3596) containing 200 µL of YPD and incubated statically for 48 h at 28 °C. Cells were resuspended by agitation and 2 µL of the culture were inoculated into 200 µL of CDGJM in 96-well microtiter plates and covered with clear breathable sealing film (Axygen; AXY BF-400) to allow measurement of growth as optical density at 600 nm using a spectrophotometric plate reader (Tecan M200 Infinite). All liquid handling was performed using automatic platforms fitted with 8-channel pipetting head (CAS-3800, Corbett Robotics, Australia).

Measurement of fermentation progress

Fermentation progress was measured using a sacrificial sampling approach as described in Liccioli et al. (2011b). In brief, four copies of the same plate were prepared by transferring 6 µL of overnight YPD culture into 600 µL of CDGJM in deep 96-well plates (Axygen; Cat No. P-DW-20-C) and covered with a cloth membrane (Diversified Biotech; Cat No. BEM-1). One plate of the 4 copies was collected and stored at -20 °C at each sampling point during fermentation to enable subsequent determination of residual sugar content. This approach was used to eliminate the potential influence of sampling from the small volume fermentations. Residual glucose and fructose were quantified enzymatically as described in Boehringer-Mannheim (1989) with final volumes adjusted to 200 µL for analysis in 96-well microtiter plates and reagents being purchased from Megazyme. Liquid handling during plate preparation and enzymatic analysis was performed robotically.

Evaluation of strains in laboratory scale fermentations

Flask fermentations (100 mL) in this study were conducted in CDGJM using either an automated 96-flask fermentation platform or 250 mL Erlenmeyer flasks fitted with an airlock (Walker et al. 2003). The automated fermentation platform built on a Tecan Freedom EVO consisted of four 24-position water jacketed carriers, to maintain temperature with magnetic stirrers to continuously agitate the fermentations. Each flask was comprised of a customized 100 mL Schott bottle with a fermentation lock and septum-sealed sampling port. The liquid handling arm of the Freedom EVO

collected samples (~1 mL) at regular intervals for off-line analysis. 200 µL from each sample was used to measure growth (see above) and the remaining samples were stored at -20 °C prior to the enzymatic determination of residual sugar.

Calculation of area under the curve and fermentation duration to determine fermentation performance

Fermentations were considered complete and total fermentation duration determined when residual sugar content was less than 2.0 g L⁻¹. The residual sugar values measured at each sampling point were used to plot fermentation curves for each mutant from the 100 mL fermentations. The Area Under the Curve (AUC) of each profile was then calculated using the composite trapezoid rule using GraphPad Prism 6 (GraphPad software INC., La Jolla, C.A., USA) as described by Liccioli et al. (2011a). Where the AUC values of the deletants were less than that of the wildtype, the deletants were considered to have an improved fermentation efficiency.

Statistical analysis

A two way ANOVA (Dunnett's multiple comparison test) was performed to determine the statistical significance between the fermentation performance of the individual mutants and the wildtype during fermentation. A scatter plot representing correlation between residual sugar and growth was performed using Pearson r correlation. Observed differences in the expression level of gene(s) in the wildtype and deletants were statistically tested using a two way ANOVA (Bonferroni's multiple comparisons test). All of these statistical analyses was performed using GraphPad Prism 6 (GraphPad software INC., La Jolla, C.A., USA).

Primers used in this study

The primers and corresponding sequences used in this study are shown in Table 1. Primers were sourced from Sigma Aldrich (Australia).

Confirmation of identity of individual deletants

The identity of individual deletants was authenticated by amplification of the deletion cassette using gene-specific primers (Table 1) and comparison of the DNA sequence to the reference genome of S288c using WU-BLAST2 (<http://www.yeastgenome.org/blast-sgd>). For deletants incorrectly labelled, the barcode region of the *KanMX* deletion cassette was amplified using primers U1 and D1 (Table 1). The tool FASTA barcodes (<http://www.ttuhscl.edu/som/cbb/FASTAbarcodes/>) was used to identify the query sequence to the up and down tags of the deletion library (McMahon et al. 2011).

Table 1. Primers used in this study.

Gene name	Forward primer sequence (5'-3')	Reverse primer sequence (5'-3')
TDA7 ^a	TAATGCGATTATTCGCTTGTAAAT	GTATGTTAAAATGCCTCCAGATTG
UBP7 ^a	GCATCCGGTAATACTAACAGAGAA	AATAATAATGGTGGTAATGGCAATG
AIM26 ^a	GCTGGTATTGACAGGTAACTATGCT	GAATCCAGTTCGTTCTCAAACCTAG
MFA2 ^a	TGATAGTTCCCTTCCGTTAAGTG	TGGCTCAAAACTTTTCACTTTAC
UBI4 ^a	AATAATCCTGGATAAACCAATTG	CTTTCCCTTTGGTAGTCACAATA
MKT1 ^a	ATTAAGAACAGACGAAAAATGGT	AACTTCAATCTATCCAGTAACGGT
MMS2 ^a	CACCACTATTGCTCATTTGTACTG	TATTATTATTGGCTGGACTGGAG
MRP17 ^a	ATACCCACTCAAACAAACTCATTG	CTTAGTCAGGATTGTCGTTAGCAT
MVB12 ^a	TGATTCATGTATTTGGTCAGA	GAAATTGACGATGGAGAAGAAGTAA
AAT2 ^a	ATACACAATTACTCCAGTAGCTGCC	TATCCTTATGTTCATGGGAGTTGT
BRO1 ^a	GCCTCCCTTTCTTGATGTATATT	ACAGTCTAGCACGAAAAAGAAGAGA
TPK2 ^a	TACAATTCTGGCCTTACCTAAA	TAATTTGCACTGAGATCATGAGA
INA1 ^a	CTCAGTACTCGTTAGCTTGAATCC	TGGGCACTTACTCATACATCAAATA
EAR1 ^a	TACAGGCTTAAGAACCTTGCTCTA	TCATACGTTCTTGTGACACATT
MFA1 ^a	GCCCACCTTATTCTTGTCTT	AAGAAACAATGAAGCAAGATAAGGA
MFA1 ^c	ATGCAACCCTACCGCTAC	CTAAGCAATAACACATGCTGGG
MFA2 ^c	GATCACCACTGCTCCACAC	TTAAGCGATAACACAGGCAG
TAF10 ^d	ATATTCCAGGATCAGGTCTCCGTAGC	GTAGTCTCTCATTCTGTTGATGTTGTTGTTG
ALG9 ^d	CACGGATAGTGGCTTGGTGAACAATTAC	TATGATTATCTGGCAGCAGGAAAGAACTTGGG
U1 ^b	GATGTCCACGAGGTCTCT	
D1 ^b		CGGTGTCGGTCTCGTAG
KanB ^b	CTGCAGCGAGGAGCCGTAAT	
KanC ^b		TGATTTGATGACGAGCGTAAT

^a Deletion cassette amplification.^b Identification of clones that were incorrectly labelled in the wine yeast library.^{a, b} Primer sequences were from *Saccharomyces* Genome Deletion Project

(http://wwwsequence.stanford.edu/group/yeast_deletion_project/Deletion_primers_PCR_sizes.txt).

^c Primers for qPCR target genes were designed using Primer3 software (http://simgene.com/Primer3).^d Primers for qPCR reference genes were adapted from Teste et al. (2009).

Construction of an *MFA1* knock-out in AWRI1631

An *MFA1* deletion strain was constructed in the haploid wine strain AWRI1631 using a PCR based deletion strategy (Baudin et al. 1993; Wach et al. 1994). The *mfa1Δ::KanMX* deletion cassette was sourced from the *MFA1* deletion clone from the prototrophic BY4741 collection (Mulleder et al. 2012). The cassette was PCR amplified using VELOCITY DNA polymerase (Bioline 21098) with gene specific primers A and D (Table 1) according to the manufacturer's instructions. The identity of the PCR product was confirmed by sequencing and transformed into AWRI1631 using the lithium acetate method (Gietz and Schiestl 2007). Transformants were selected on YPD plates containing G418 sulfate (200 mg L⁻¹) and the deletions were confirmed by PCR amplification and sequencing.

Quantitative PCR for gene expression studies

To study gene expression, a qPCR experiment was conducted following the MIQE guidelines (Bustin et al. 2010; Taylor and Mrkusich 2014). Samples of the culture harvested during fermentation at similar residual sugar concentrations (approximately 128, 83, 35 and 10 g L⁻¹) were used to compare transcript abundance between strains. Cells were collected by centrifugation, and resuspended in TRIzol® Reagent (Invitrogen; 15596026) followed by snap freezing in liquid N and storage at -80 °C. Total RNA was extracted using Zymo Research kits (Direct-zol RNA MiniPrep R2050) as per the manufacturer's instructions. RNA purity and integrity were estimated from OD₂₆₀/OD₂₈₀ (Tecan M200 Infinite) and gel electrophoresis. cDNA was synthesized from the total RNA (BioRad iScript kit; 170-8897) and any contaminating DNA removed with an Ambion TURBO DNasefree™ kit (AM1907). Relative quantification of the transcripts was performed on a Biorad CFX-96 Real Time PCR system with Ssofast™ Evagreen Supermix (BioRad; 172-5203) with primers as listed in Table 1. The amplification cycle included an initial denaturation at 95 °C for 30 sec followed by 40 cycles at 95 °C for 5 sec and 60 °C for 5 sec. A sample maximisation strategy was used with all six biological replicates being amplified in duplicate. At the end of each run a melting curve was prepared for all samples from 65 to 95 °C. Transcript abundance of the target genes was normalized with abundance of reference genes (*TAF10*, *ALG9*; Teste et al. 2009) in the same sample. The stability of the housekeeping genes was determined using qbase+ software (Vandesompele et al. 2002) and primer efficiency was determined by running a standard curve using a serially diluted cDNA sample. The mean fold-change of expression for given genes per sample for different time points was determined using a relative quantification method (Schmittgen and Livak 2008) using the qbase+ software (Vandesompele et al. 2002), Excel (Microsoft Office 2013) and GraphPad Prism (GraphPad Prism 6 – GraphPad software INC., La Jolla, C.A., USA).

3.4 Results

Construction of a wine yeast deletion library

To ensure the most industrially relevant genetic background was used in this study, a wine yeast deletion library was constructed in the haploid wine yeast derivative AWRI1631 (Borneman et al. 2008). Using the approach reported previously (Varela et al. 2012) a total of 1844 mutants were constructed (Appendix 3, Pg 163). All the deletants were checked by PCR amplification and sequencing of the deletion cassette.

High throughput screening (micro-fermentation) of the wine yeast deletion collection

In order to identify genes that influence fermentation duration when only supplied with minimal N (75 mg L^{-1}), the sugar utilisation profile of clones from this wine yeast deletion collection was compared to that of the wildtype strain, AWRI1631, in a high throughput micro-fermentation screening in CDGJM (Fig. 1).

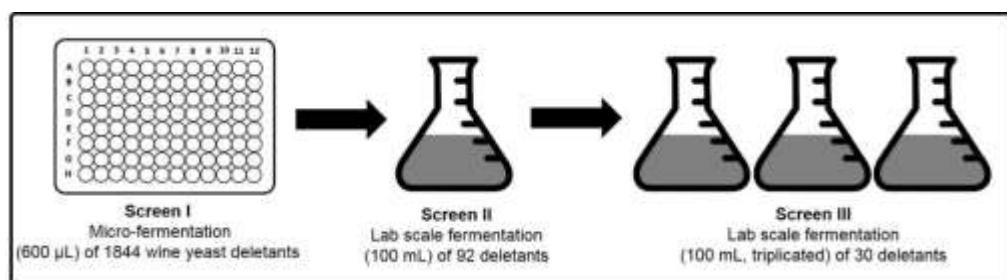


Figure 1. Schematic representation of the screening of the wine yeast deletion library for fermentation performance in a chemically defined grape juice medium with limited nitrogen (75 mg N L^{-1}).

To ensure supplied nitrogen would protract fermentation preliminary experiments were also undertaken with the addition of sufficient N (450 mg L^{-1}), the wildtype strain was able to utilise all available sugar in ~ 60 h in comparison to ~ 200 h when only minimal N is available (data not shown). Additional fermentation wells containing the wildtype were sampled at a higher frequency to accurately determine the most appropriate sampling points (A-D) for the deletants (Fig. 2A). Initially the wildtype consumed sugars slowly (43 g L^{-1} by 96 h), however fermentation became more vigorous after 96 h. Using this information, residual sugars were measured for the 1844 deletants at time points B (121 h) and C (146 h). The extent of sugar catabolism by the wine yeast deletants was highly variable (Fig. 2B), when the concentration of residual sugars for the wildtype was $42 \pm 8 \text{ g L}^{-1}$ (time point C). Given this range the deletants were classified into three groups: 1040 with residual sugar around that of the wildtype ($30 - 60 \text{ g L}^{-1}$), 282 with more residual sugar

($> 60 \text{ g L}^{-1}$) than the wildtype and 522 with markedly less residual sugar ($< 30 \text{ g L}^{-1}$) than the wildtype group (Appendix 7, Pg 189). Apart from the sugar utilisation profile, growth was monitored for the wildtype and the deletants during the early stages of micro-fermentation (Fig. 2A and Appendix 7, Pg 189). The wildtype reached stationary phase at 113 h with an optical density of 1.77. Therefore, using optical density at 113 h and residual sugars at 146 h (time point C) a correlation was plotted to identify whether the fermentation abilities of the deletants were dependent on growth (Fig. 3). A weak negative correlation was observed, emphasizing a small effect of growth on fermentation progression (i.e. increase in growth of the deletants leading to a decrease in residual sugar). Further analysis of the 522 deletants that completed fermentation in a shorter time than the wildtype, identified that the majority (408 i.e. 78%) of the deletants had higher optical density than the wildtype.

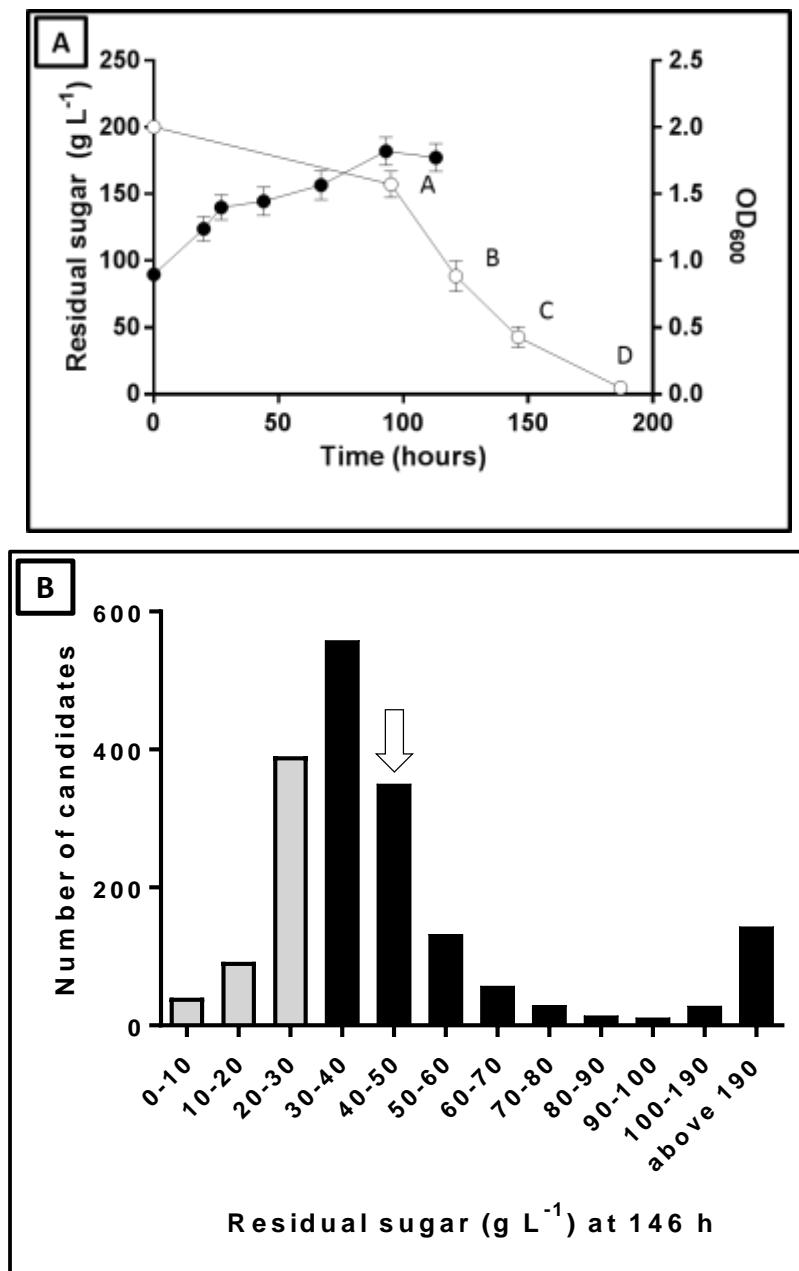


Figure 2. (A) Fermentation performance of AWRI1631 during micro-fermentation. Sugar utilisation (open symbols) at four time points: A (95 h), B (121 h), C (146 h) and D (187 h). Growth as estimated by optical density at 600 nm (solid symbols) Sixteen replicates were analysed. **(B)** Histogram representing the distribution of deletion mutants (number) based on the residual sugar content at time point C (146 h). The arrow indicates the residual sugar concentration range of the wildtype, AWRI1631 ($42 \pm 8 \text{ g L}^{-1}$) at time point C. Candidates for further analysis were chosen from the groups with grey shaded bars.

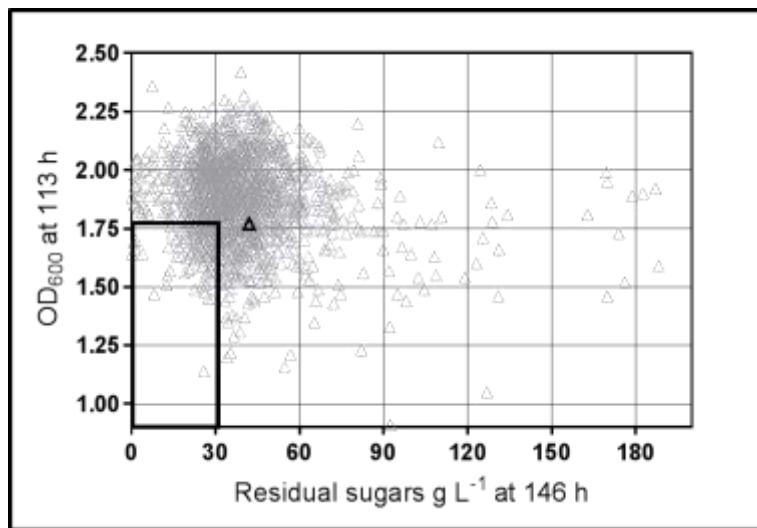


Figure 3. Scatter plot of the residual sugar (at time point C, 146 h) and growth (OD_{600} at 113 h) of the deletion mutants (grey) and the wildtype (black) during micro-fermentation. Deletants (114) exhibiting shorter fermentation duration and decreased growth in comparison to the wildtype are highlighted in a bold box. Correlation coefficient (r) = -0.1616.

Based on the results of this screening exercise, 92 candidates showing the fastest fermentation and no growth defects were chosen for further evaluation (Appendix 8, Pg 190). These candidates were selected as they were observed to utilise significantly more sugar than the wildtype, having less than 48 and 26 $g\ L^{-1}$ of residual sugar at time points B and C, respectively – i.e. approximately half of that of the wildtype (88 and 42 $g\ L^{-1}$ of residual sugar at time point B and C, respectively). Of the possible 522 candidates, an arbitrary total of 92 of the fastest deletants was chosen since this number (plus controls) could be accommodated in the 96-position fermentation platform.

Confirmation of deletant fermentation performance in laboratory scale fermentations

The fermentation performance of the 92 shortlisted deletants was confirmed in laboratory-scale (100 mL) fermentations (screen II) since this format allowed greater control of experimental conditions and better representation of an anaerobic, wine-like fermentation (Walker et al. 2014). Results of this experiment highlighted 48 deletants that completed fermentation in less time than the wildtype (Fig. 4A), with 30 utilising all sugar within 131 to 205 h compared to 229 h for the wildtype. As well as total fermentation time, the Area Under the Curve (AUC) of fermentation curves of individual deletants was also used for a more holistic view of the pattern of sugar utilisation by which deletants were ranked (Liccioli et al. 2011a). AUC values calculated for the 30 strains were 7185 to 9516 (i.e. 72-95 % of the wildtype at 9999; Appendix 9, Pg 191). These 30 candidates were further assessed in a third screen conducted in triplicate (Fig. 4B). Fifteen of these deletants

utilised all sugars in significantly less time than the wildtype (Table 2). *Tda7Δ* and *aim26Δ* were the most rapid deletants, completing fermentation in 121 h, which was 172 h ahead of the wildtype (293 h). Closely following were *ubp7Δ* and then *mfa2Δ* (Table 2). The remaining eleven deletants exhibited fermentation durations ranging from 63 to 85% of the wildtype. The AUC for individual deletants representing sugar utilisation over time was calculated as being 54–85% of AWRI1631, with *tda7Δ* (AUC = 8911) being the most efficient and *ear1Δ* (AUC = 14073) being the least efficient compared to the wildtype (AUC = 16399; Table 2).

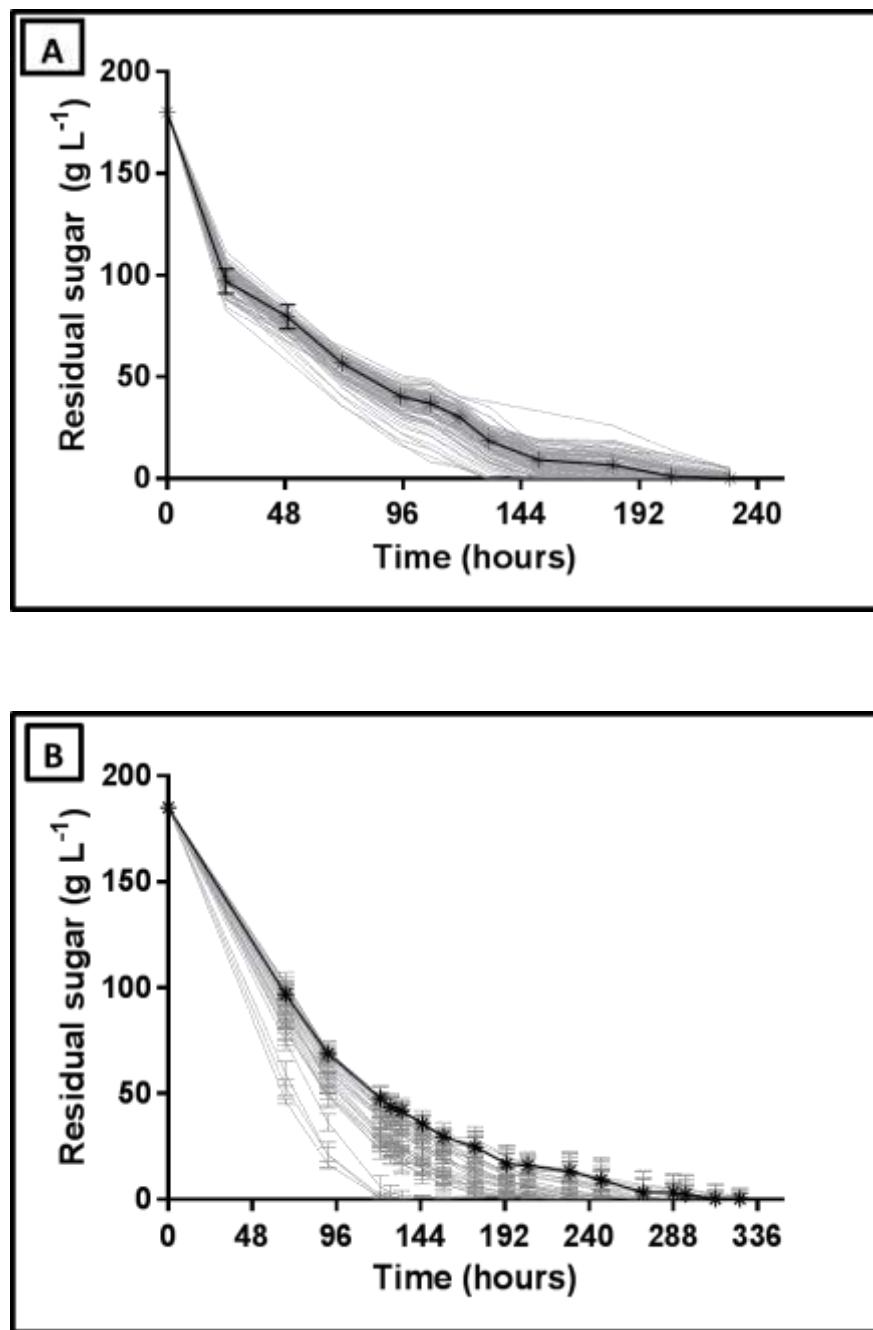


Figure 4. Sugar utilisation of the wildtype (AWRI1631, black line) compared to the shortlisted 92 deletants (grey lines) over time. Fermentation was conducted in 100 mL of CDGJM with 75 mg N L^{-1} at 28 °C on a robotic sampling platform. **(A)** Screen II of the shortlisted 92 yeast deletants (single cultures) compared with the wildtype (4 replicates). **(B)** Screen III of the selected 30 candidates in triplicate, along with the wildtype strain (6 replicates). Standard deviation of the mean is represented as error bars

Table 2. Fermentation performance of fermentation efficient deletants.

Deleted gene	Gene function	FD (h) (FD ± SD)	FD % of wildtype	AUC (AUC ± SD)	AUC % of wildtype (AUC % ± SD)
<i>TDA7^a</i>	Topoisomerase I Damage Affected	121 ± 0	41 ± 0	8911 ± 114	54 ± 0
<i>UBP7^a</i>	Ubiquitin-specific Protease	123 ± 3	42 ± 1	9332 ± 275	57 ± 1
<i>AIM26^a</i>	Altered Inheritance rate of Mitochondria	121 ± 0	41 ± 0	9602 ± 482	58 ± 2
<i>MFA2^a</i>	Mating Factor A	131 ± 12	45 ± 4	10825 ± 431	66 ± 2
<i>UBC13^a</i>	Ubiquitin-conjugating	185 ± 17	63 ± 6	12205 ± 645	74 ± 3
<i>UBI4^a</i>	Ubiquitin	193 ± 0	66 ± 0	12225 ± 306	74 ± 1
<i>MMS2^b</i>	Methyl MethaneSulfonate sensitivity	209 ± 18	71 ± 6	12475 ± 720	76 ± 4
<i>MRP17^a</i>	Mitochondrial Ribosomal Protein	201 ± 7	69 ± 2	12643 ± 85	77 ± 1
<i>MKT1^a</i>	Maintenance of K2 Killer Toxin	191 ± 15	65 ± 5	12712 ± 360	77 ± 2
<i>MVB12^a</i>	MultiVesicular Body sorting factor of 12 kilodaltons	193 ± 0	66 ± 0	13239 ± 211	81 ± 1
<i>AAT2^b</i>	Aspartate AminoTransferase	215 ± 28	73 ± 10	13273 ± 986	81 ± 6
<i>BRO1^b</i>	BCK1-like Resistance to Osmotic shock	227 ± 21	78 ± 7	13387 ± 699	81 ± 4
<i>INA1^c</i>	Indicator of Abscission	225 ± 48	77 ± 16	13567 ± 1375	82 ± 8
<i>TPK2^a</i>	Takashi's Protein Kinase	201 ± 7	69 ± 2	13593 ± 184	83 ± 1
<i>EAR1^d</i>	Endosomal Adaptor of Rsp5p	249 ± 38	85 ± 13	14073 ± 269	85 ± 6
<i>Wildtype</i>		293 ± 21	100	16399 ± 752	100

Fermentation performance was calculated as AUC (Area Under the Curve) and FD (Fermentation Duration) of the 15 shortlisted gene deletants (from Screen III) and percentage difference in AUC and FD of the deletants in comparison to the wildtype, AWRI1631 (i.e. AUC % = (mutant AUC/parent AUC)*100 and FD % = (deletant FD/wildtype FD)*100). Fermentation duration corresponds to the time (h) by which residual sugar was < 2 g L⁻¹. Values represent the average of FD/AUC ± SD. Deletants significantly different in fermentation completion when compared to the wildtype are represented by ^a, p <0.0001; ^b, p<0.001; ^c, p <0.01, ^d, p <0.1 (two-way ANOVA).

Characterisation of the deletants *mfa1Δ* and *mfa2Δ*

To date, *MFA2* has only been annotated to mating and cell-type regulation (Galgoczy et al. 2004; Johnson 1995; Michaelis and Herskowitz 1988). Our results of improved fermentation performance under N-limited conditions for *mfa2Δ* alludes to a dual function of this gene. In order to explore the link between efficient fermentation and the yeast mating factor, the function of the two genes encoding mating a-factor, *MFA2* and *MFA1*, were further studied. The sugar utilisation and growth profile of *mfa1Δ* was similar to the wildtype (AWRI1631), whilst growth and fermentation duration of *mfa2Δ* were reduced (Fig. 5).

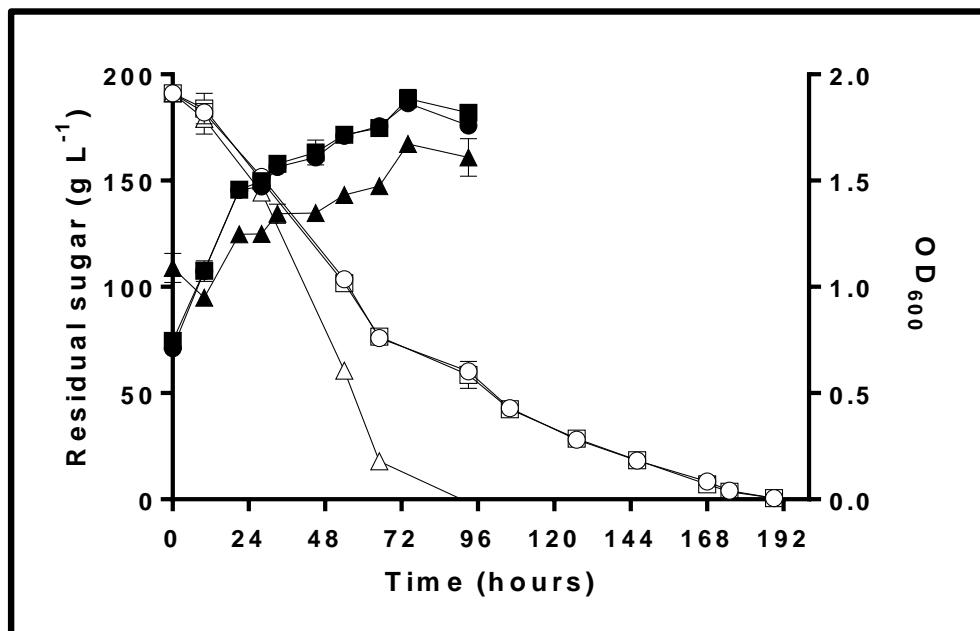


Figure 5. Fermentation performance of *mfa1Δ*, *mfa2Δ* and the wildtype AWRI1631. Fermentations (100 mL) were conducted in triplicate in CDGJM with 75 mg N L⁻¹. Data points represent the average value \pm standard deviation. Sugar utilisation over time of the individuals (open symbols). Growth as estimated by optical density at 600 nm of the individuals (solid symbols). Circle; AWRI1631, square; AWRI1631 *mfa1Δ* and triangle; AWRI1631 *mfa2Δ*.

The expression of *MFA1* and *MFA2* during fermentation was also examined. RNA was extracted from cultures of mutant (AWRI1631 *mfa2Δ*) and parental (AWRI1631) strains when the sugar content was approximately 128, 83, 35 and 10 g L⁻¹ in each culture. Differential expression of *MFA1* and *MFA2* was observed in the wildtype, with *MFA2* being expressed to a greater extent (~1.93 fold NREL) compared to *MFA1* (~0.17 fold NREL) at the first time point (Fig. 6A and B). The expression of both genes markedly declined at 83 and 35 g L⁻¹ of residual sugar (corresponding to 58.5% and 82.5% sugar consumption, respectively). In the case of *MFA2*, expression recovered in the wildtype in the final sample (95% of sugar consumed; Fig. 6B). The

absence of *MFA2* transcript in the AWRI1631 *mfa2Δ* strain confirmed the deletion of the *MFA2* gene (Fig. 6B). Interestingly, expression of *MFA1* was upregulated by ~6- to 7-fold in the *MFA2* deletant at the initial sampling point (128 g L⁻¹). Although *MFA1* expression declined subsequently, it remained ~4-fold higher than in the wildtype (Fig. 6A).

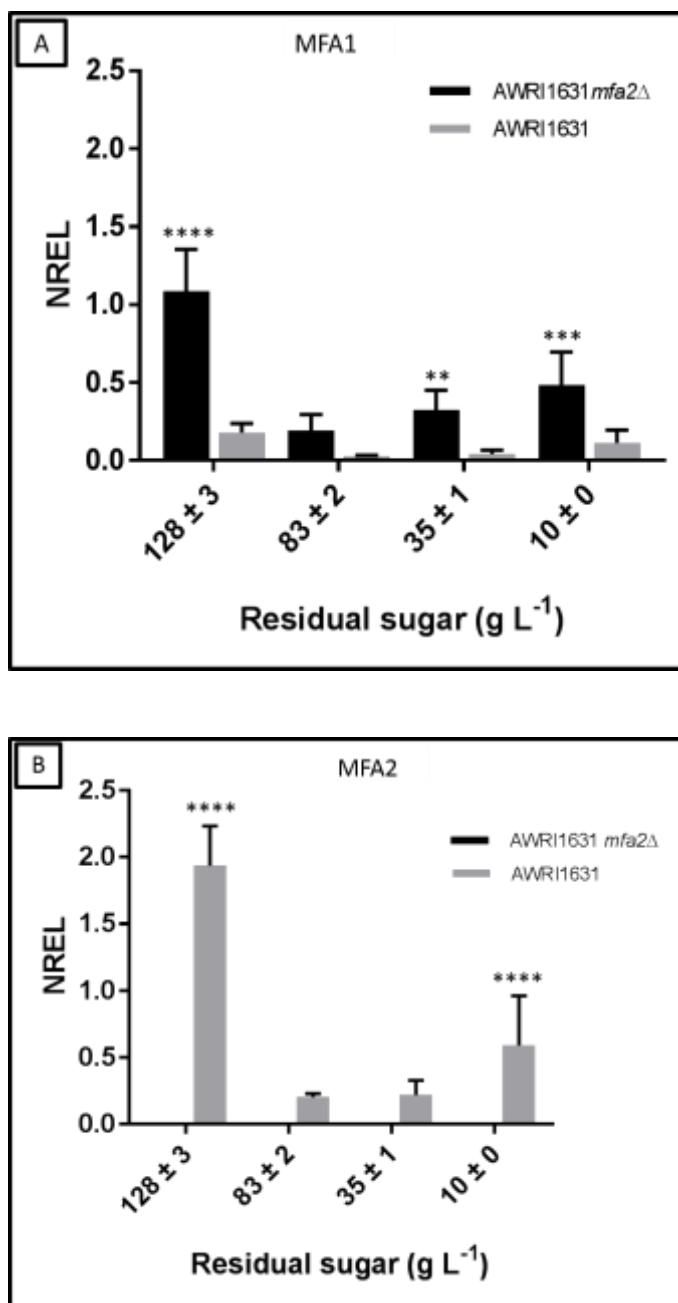


Figure 6. Expression of *MFA1* (A) and *MFA2* (B) in AWRI1631 and AWRI1631 *mfa2Δ* strains. Normalized relative expression level (NREL) is shown at four selected residual sugar levels (g L⁻¹). Two-way ANOVA was used to determine the significant differences in expression levels between *mfa2Δ* and the parent. Significant difference is indicated as **, p < 0.01; ***, p < 0.001; ****, p < 0.0001.

3.5 Discussion

Yeasts differ in fermentative capabilities in N deficient must, which is dependent upon their demand for nitrogen (N) to maintain glycolytic flux – the genetic basis of which is the focus of recent research (Brice et al. 2014; Brice et al. 2013; Tesnière et al. 2015). In order to generate N-efficient strains it is important to identify genes and processes that allow the yeast to perform well in low N juices. Gene deletions provide a paradigm to better understand the mechanisms behind the improved fermentation phenotype exhibited by these strains. Although there are several strategies that are now available for strain improvement, whole genome investigations using collections of yeast mutants with single defined gene deletions (Giaever et al. 2002; Winzeler et al. 1999) provide opportunities to readily study gene functionality, and so identify targets for strain improvement (Donalies et al. 2008). Some studies have successfully used deletion collections in auxotrophic laboratory yeast backgrounds to investigate growth phenotypes in response to exposure to single stress factors associated with fermentation. Examples include the identification of genes essential for tolerance to ethanol and other alcohols (Fujita et al. 2006; Teixeira et al. 2009; van Voorst et al. 2006), high pressure and low temperature (Abe and Minegishi 2008), acetic acid (Mira et al. 2010), oxidative and chemical stress (Tucker and Fields 2004), high glucose (Teixeira et al. 2010), anaerobic growth (Reiner et al. 2006) and low temperature (Salvadó et al. 2016). Most recently this approach has been used by this group to define the Fermentome, genes essential to allow the successful completion of fermentation under the multi-stress conditions of a high sugar, juice-like medium (Walker et al. 2014). Alternatively, genes important to the response of yeast to stress have been identified through transcriptional analyses following exposure to stresses such as N limitation (Backhus et al. 2001; Brice et al. 2013; Contreras et al. 2012; Mendes-Ferreira et al. 2007a; Mendes-Ferreira et al. 2007b; Rossignol et al. 2003). But research in a wine yeast background to identify genes related to N-efficiency in wine-like fermentations has been limited to the screening of a random collection of transposon mutants of non-essential genes (Gardner et al. 2005) or a few deletion strains (Zhang 2014). The present study sought to fill this knowledge gap.

The need to supply sufficient N (both in amount and specific amino acid composition) to meet the auxotrophic requirements of laboratory strains used in previous studies, has meant that investigations related to N efficiency were limited to high N content (Walker et al. 2014). Several studies have shown that supplementation can affect metabolism through the blockage of pathways, or by combinatorially influencing metabolic networks, thereby making quantitative comparisons of growth and physiological studies difficult (Mulleder et al. 2012; Pronk 2002). The recent availability of prototrophic libraries in laboratory yeast (Mulleder et al. 2012) and wine yeast

(Varela et al. 2012) has made it possible to conduct studies under limited-N conditions, overcoming previous shortcomings.

In this study, a prototrophic wine yeast deletion library (Varela et al. 2012) was used to identify genes influencing fermentation in a wine-like limited N medium. Based on the sugar utilisation profile of the wildtype, preliminary screening via micro-fermentations (Appendix 3, Pg 163) of the wine yeast deletion collection allowed mutants to be segregated into groups based on the extent of sugar utilisation. Fermentation by 282 of the mutants (15%, Fig. 2B, (Appendix 7, Pg 189) was protracted indicating that these genes were essential for the timely completion of fermentation. Walker et al. (2014) used the deletion library in the auxotrophic laboratory strain BY4743 to identify 93 genes (i.e. the fermentome) essential for fermentation in CDGJM under high N (450 mg N L^{-1}) and high sugar (200 g L^{-1}) conditions. Interestingly, despite differences in the experimental conditions of these studies (i.e. N availability and strain background) our list of 282 protracted deletants included 15 of the fermentome genes (Table 3) identified by Walker et al. (2014), underscoring the universal importance of these. A further 14 deletants of fermentome genes, however, proved non-essential in the present study (i.e. in the AWRI1631 background), since they were not protracted compared to the wildtype. Further work to explore the influence of strain background and growth conditions will help resolve such apparent differences. More importantly, the wine yeast deletion library needs to be expanded to a complete collection from the current 1844 deletants or at least the remaining 64 fermentation essential genes identified by Walker et al. (2014) in order to assess their essentiality in the conditions of the present study.

Table 3. List of genes essential in timely completion of fermentation.

ORF	Gene name	Gene function description
<i>YEL051W</i>	<i>VMA8</i>	Subunit D of the V1 peripheral membrane domain of V-ATPase
<i>YHL020C</i>	<i>OPI1</i>	Transcriptional regulator of a variety of genes
<i>YKR007W</i>	<i>MEH1</i>	Component of the EGO and GSE complexes
<i>YFR053C</i>	<i>HXK1</i>	Hexokinase isoenzyme 1
<i>YDR247W</i>	<i>VHS1</i>	Cytoplasmic serine/threonine protein kinase
<i>YGR063C</i>	<i>SPT4</i>	Spt4p/5p (DSIF) transcription elongation factor complex subunit
<i>YJR033C</i>	<i>RAV1</i>	Subunit of RAVE complex (Rav1p, Rav2p, Skp1p)
<i>YLL007C</i>	<i>LMO1</i>	Homolog of mammalian ELMO (Engulfment and cell MOtility)
<i>YMR263W</i>	<i>SAP30</i>	Component of Rpd3L histone deacetylase complex
<i>YNL076W</i>	<i>MKS1</i>	Pleiotropic negative transcriptional regulator
<i>YOR209C</i>	<i>NPT1</i>	Nicotinate phosphoribosyltransferase
<i>YOR221C</i>	<i>MCT1</i>	Predicted malonyl-CoA:ACP transferase
<i>YOR265W</i>	<i>RBL2</i>	Protein involved in microtubule morphogenesis
<i>YPR036W</i>	<i>VMA13</i>	Vacuolar Membrane Atpase
<i>YPR074C</i>	<i>TKL1</i>	TransKetoLase

ORF; Open Reading Frame.

Twenty eight percent of the wine yeast deletion library mutants (522 deletants) completed fermentation in a shorter time, alluding to the functional genes negatively influencing fermentation progress. Whether these genes are directly related to nitrogen metabolism or instead are general regulators of fermentation is worthy of future study. Of these, 114 deletants were capable of enabling shorter fermentation time in spite of exhibiting decreased growth. In general fermentation rate has been reported to correlate with biomass (Varela et al. 2004), further evaluation of this group of genes may reveal exceptions. However these 522 genes are potential targets for guided strain improvement and thus were assessed in greater detail. A shortlist of the best performing 92 strains (< 48 and < 26 g L⁻¹ of sugar at time point B and C, respectively – i.e. approx. 50% of that of the wildtype) were chosen for further evaluation in 100 mL fermentations. Fifteen strains completed fermentation in between 41 and 69% of the time taken by the parent (Table 2). Gene Ontology (GO) analysis (Appendix 10, Pg 192) revealed that 11 out of these 15 genes were over-

represented for the GO terms related to different stages of ubiquitination and protein modification/turnover or metabolic processes: protein polyubiquitination (*UBI4*, *UBC13* and *MMS2*), protein deubiquitination (*UBP7*, *UBI4* and *BRO1*), protein modification by small protein conjugation (*UBP7*, *UBI4*, *BRO1*, *MMS2* and *UBC13*), post-translational protein modification (*UBP7*, *TPK2*, *UBI4*, *BRO1*, *MMS2* and *UBC13*) protein metabolic process (*UBP7*, *TPK2*, *UBI4*, *BRO1*, *MFA2*, *MMS2*, *MKT1*, *EAR1*, *UBC13* and *MRP17*) and endosomal transport (*MVB12*, *EAR1* and *BRO1*). The remaining four genes include *AAT2* (involved in aspartate aminotransferase in nitrogen metabolism) and three with unknown function, *INA1*, *TDA7* and *AIM26*.

Ubiquitination is a protein degradation process that is required in many cellular processes such as the cell cycle, endocytosis and stress response (Hershko 1997; Staub and Rotin 2006). In a study of the stress response of yeast under sake brewing conditions, 235 genes related to stress response or ethanol fermentation were deleted and investigated for their impact on ethanol production (Wu et al. 2009). Amongst the 235 genes were *UBI4*, *UBP7*, *UBC13* and *TPK2*, which have also been identified in our study. Wu and co-workers (2009) demonstrated that deletion of ubiquitin related genes such as *UBI4* and *UBC13*, increased fermentation abilities producing significantly more ethanol than the parental strain. Furthermore, enhanced fermentation abilities of *ubi4Δ* were confirmed in a number of laboratory yeast strain backgrounds (Wu et al. 2009).

An important group of proteins regulated by ubiquitination are plasma membrane proteins, in particular, amino acid permeases. Ubiquitination of these initiates endocytosis and the eventual degradation of the ubiquitin bound protein in the vacuole/lysosome via the multivesicular body (MVB) pathway (Hicke 2001). Genes such as *BRO1* and *EAR1* (in combination with *SSH4*) play a role in the regulation of Gap1p (general amino acid permease) degradation by ubiquitination in the presence of ammonium (Leon et al. 2008; Nikko et al. 2003). However, deletion of *BRO1* (Nikko et al. 2003), *MVB12* (Oestreich et al. 2007) or *EAR1* (in combination with *ssh4Δ*; Leon et al. 2008) have been shown to disrupt this pathway thereby interfering with the degradation process. Interestingly, *ear1Δ* (in combination with *ssh4Δ*) results in the localisation of Gap1p to the vacuolar membrane rather than the vacuolar lumen, where it would otherwise be degraded (Leon et al. 2008) and *bro1Δ* results in the recycling of Gap1p to the cell membrane without ubiquitination in the presence of ammonium (Nikko et al. 2003), essentially desensitising *GAP1* to nitrogen catabolite repression at the protein level. It is possible to consider that the lack of regulation of amino acid permeases by the MVB pathway may play a role in successful low N fermentation. However, it is not known how the gene deletions in this study affect the degradation of other amino

acid permeases or amino acid catabolic enzymes, which could also play a role in the observed enhanced fermentation performance.

MFA2 was further investigated to understand how deletion of this mating gene positively affected fermentation. To date *MFA2* has only been annotated to mating and cell-type regulation (Galgoczy et al. 2004; Johnson 1995; Michaelis and Herskowitz 1988), *MFA2*, together with *MFA1*, considered functionally redundant, are responsible for the production of a signalling molecule (a-factor) in a *MATa* cell (Michaelis and Herskowitz 1988). Both *MFA1* and *MFA2* produce a precursor (36 and 38 amino acids, respectively), which undergoes several modification steps involving different genes before the mature a-factor molecule is transported outside the cell through the ABC transporter *STE6* (review by Michaelis and Barrowman (2012)). We hypothesized that mating pheromone production and processing is an energy consuming process. Therefore in a haploid heterothallic strain, where no mating or switching of mating types occurs, the ability to down-regulate synthesis/processing of mating factor-a, may lead to a metabolic efficiency that benefits fermentation. This finding is in keeping with the finding that deletants of *MFA2* are haploproficient in either carbon- or nitrogen-limited media (Delnieri et al. 2007). Given that deletion of *MFA2* resulted in a significant increase (34%) in sugar catabolism compared to the parent, we chose to also examine *MFA1*, a functionally redundant gene encoding a-factor. As the *mfa1Δ* deletant was absent in the deletion collection used in this work, the *MFA1* deletant (*mfa1Δ::KanMX*) was constructed in the same background as *mfa2Δ*, AWRI1631, to determine whether an *mfa1Δ* exhibited a similar fermentation phenotype. In fact, *mfa1Δ* deletant performed no differently to the wildtype in terms of fermentation and growth (Fig. 5), suggesting potentially different roles for the *MFA1* and *MFA2* gene products.

In the wildtype background, transcription of *MFA2* was dominant to that of *MFA1* (Fig. 6A and B). However, when *MFA2* was deleted, *MFA1* produced a pattern of expression through fermentation that implied a compensation for the *MFA2* gene product in *mfa2Δ*. Contrasting with these results, others have shown *MFA1* and *MFA2* to be transcribed at similar levels and to be functionally equivalent. Thus deletion of one of the mating factor genes did not prevent the strain being mating proficient (Chen et al. 1997; Michaelis and Herskowitz 1988). However, here we show that these genes are not functionally equivalent, at least in terms of their impact on fermentation performance. Further work is needed to determine the basis for the link between these genes and fermentation, the extent to which they are functionally equivalent in mating type functions and interactions with related genes, and the basis for their differential expression even in the wildtype.

Novel interactions between genes have been identified when unexpected phenotypes result from mutations in two or more genes (Costanzo et al. 2010; Dixon et al. 2009). Nevertheless, gene interaction networks (adapted from Costanzo et al. 2016 and the *Saccharomyces Genome Database*), already indicate that *MFA1* and *MFA2* genes interact with different sets of genes, including genes involved in either carbohydrate metabolism, transport or storage (Table 4). These genes will be examined for their potential role in fermentation speed in a *mfa2Δ* strain.

Table 4. List of genes interacting with *MFA2* involved in carbohydrate metabolism/ transport or storage.

Type of interaction	Gene	Gene function
Negative	<i>GAL10</i>	GALactose metabolism
Negative	<i>GID7</i>	Glucose Induced Degradation deficient
Negative	<i>HXT5</i>	HeXose Transporter
Negative	<i>STL1</i>	Sugar Transporter-Like protein
Negative	<i>PCL6</i>	Pho85 CycLin

Adapted from Costanzo et al. (2016).

3.6 Conclusion

Nitrogen efficient strains offer a means for avoiding or reducing the problems associated with the fermentation of grape juices low in N, but the basis for these efficiency differences is not known. Motivated by this and recognising the fact that wine and laboratory yeast strains can behave differently under wine-like conditions, we generated and exploited a partial wine yeast deletion collection to identify genes whose deletion improved fermentation performance in low N media. Fifteen deletants were identified through a robust 3-stage screening process. The unexpected link between one of these gene deletions, *mfa2Δ*, and fermentation remains to be explored at a more fundamental level. Further candidates would likely be found with completion of the library with all non-essential genes, with all showing promise as the basis for construction of optimised strains for industry use, perhaps by genome editing (Bao et al. 2014). Their use would avoid the need for YAN supplementation or at least reduce the incidence of problem fermentations in unsupplemented juices.

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CHAPTER 4

Investigation of the effect of an *MFA2* deletion on fermentation duration

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Overall percentage (%)	80		
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By signing the Statement of Authorship, each author certifies that:

- I. the candidate's stated contribution to the publication is accurate (as detailed above);
- II. permission is granted for the candidate to include the publication in the thesis; and
- III. the sum of all co-author contributions is equal to 100% less the candidate's stated contribution.

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Investigation of the effect of an *MFA2* deletion on fermentation duration

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4.1 Abstract

Nitrogen insufficiency during oenological fermentation can lead to problems such as stuck and sluggish fermentation. Although supplementation can alleviate these problems, an alternative option is to conduct fermentation with yeast strains that utilize nitrogen efficiently thereby ensuing fermentation completion. We previously reported that deletion of *MFA2* (mating a-factor) in AWRI1631, a derivative of the wine related strain, N96, resulted in a 34% reduction of fermentation duration. Since *MFA2* has not previously been reported to impact fermentation, we sought to investigate *MFA2* further in this context, and to determine whether the effect of *MFA2* gene deletion was dependent on the mating type, genotype, and gene dosage. We found that the shortened fermentation by the *MFA2* deletion was not only specific to the N96 strain background but also specific to *MATa* cells. Shorter fermentation was also observed in diploids of N96 when *mfa2Δ* was sourced from the AWRI1631 *mfa2Δ* (*MATa*) strain. Further, these *MFA2* deletants also showed reduced growth, which could be due to increased protein turnover and the strain increasing glycolytic flux by driving more energy into protein synthesis rather than biomass formation.

4.2 Introduction

Traditional winemaking relies on the microflora present on grapes and winery equipment for the fermentation of grape juice to wine, whereas modern winemaking often uses commercially cultured strains (Pretorius 2000). Commercial strains with well-defined technological properties are a valuable tool to help winemakers manage wine styles, limit fermentation problems and contribute to quality (Pretorius 2000; Schuller and casal 2005). One of the expected qualities of such strains is an ability to ferment juices to "dryness" promptly. For this reason, commercially available starter cultures are usually robust fermenters able to withstand harsh winemaking conditions such as high alcohol concentrations and depletion of nutrients (Bradbury et al. 2006).

Yeast assimilable sources of nitrogen are often low in grape juices, which can be a cause of protracted fermentations (Bisson 1999) and the production of off aromas (Jiranek et al. 1995b). It is therefore commonplace to supplement juice/must with inorganic sources of nitrogen (commonly diammonium phosphate; DAP) (Jiranek et al. 1995a). However as nitrogen concentration and composition in grapes is known to vary significantly depending on environmental factors, vineyard practices and grape variety (Alexandre and Charpentier 1998, Bell and Henschke 2005), supplementation may prove to be insufficient or excessive. Overuse of DAP can lead to overproduction of some compounds (e.g. acetate esters, ethyl acetate), which negatively influence wine aroma and flavour (Ugliano et al. 2007). In some cases, residual nitrogen can encourage microbial instability and reduce the quality of the wine (Bell and Henschke 2005). Thus the nitrogen concentration in grape juice and must influences both the kinetics of the fermentation as well as aroma and flavor compound formation (Bell and Henschke 2005; Jiranek et al. 1991).

There is a wide variability between wine yeast strains in their nitrogen requirement (Jiranek et al. 1995a; Brice et al. 2014; Crepin et al. 2012; Gutiérrez et al. 2013a; Julien et al. 2000). For instance, some strains require more nitrogen to maintain high viability and metabolic activity (Brice et al. 2013). Therefore, the use of yeast 'nitrogen efficient' strains represents an innovative way to reduce the incidence of fermentation problems associated with nitrogen deficient juices (Jiranek et al. 1995a; Gardner et al. 2002), without resorting to supplementation, which can have the limitations detailed above. Although research in oenological microbiology has identified a few genes influencing nitrogen efficiency (Gardner et al. 2005; Zhang 2014), there are likely other genes and mechanisms to uncover.

Previously we identified genes influencing fermentation performance in low nitrogen conditions (Peter et al. 2017). Upon deletion, these genes give rise to improved fermentation performance, thereby making them possible targets for strain improvement (Astorga et al. 2007,

Donalies et al. 2008), potentially using genome editing techniques, should strains produced in this way be acceptable for use in the wine industry (Bao et al. 2014). Among the shortlisted mutants, *mfa2Δ* catabolised all sugar in about 2/3rd of the time taken by the wildtype. *MFA2* encodes the mating a-factor pheromone precursor and has not previously been linked to fermentation kinetics. We therefore sought to further characterise the effect of *mfa2Δ* on fermentation duration.

Most wine strains are diploid (Pretorius 2000). Since the deletant was originally highlighted in the haploid AWRI1631 background, the role of ploidy and gene dosage was therefore investigated using diploid heterozygous and homozygous *MFA2* deletion strains. Comparisons were made for the fermentation performance of haploid strains derived from N96 (AWRI1633 *MATα*, AWRI1631 *MATα*) and L2056 (C911A *MATα*, C9 11D *MATα*).

4.3 Materials and methods

Strains, media and culture conditions

The strains used in this study are listed in Table 1. Strains C911A and C911D are genetically related but non-identical haploid derivatives (Walker et al. 2003) of commercial strain L2056 (Lallemand, Australia). Strains AWRI1631 and AWRI1633 are haploid derivatives of commercial strain N96 (Anchor Yeast, South Africa). AWRI1633 was constructed using AWRI1631 (*MATα*; lacking HO gene) by introducing a plasmid borne copy of the HO gene, driven by the GAL4 promoter, which induced switching of mating type from *MATα* to *MATα*. The plasmid in the resulting AWRI1633 was cured on 5-FOA plates. All strains were maintained on Yeast Peptone Dextrose (YPD) medium (1% yeast extract, 2% bactopeptone, 2% D-glucose). G418 sulfate (200 mg L⁻¹) was added for antibiotic selection. Fermentations were conducted in Chemically Defined Grape Juice Medium (CDGJM, McBryde et al. 2006) containing 200 g L⁻¹ sugar with low nitrogen concentration (75 mg N L⁻¹ as a mix of amino acids and ammonium chloride; Henschke and Jiranek 1993) with 3 g L⁻¹ of polyphenol extract to mimic red grape juice (Walker et al. 2014).

Table 1. Strains used in this study.

Strains	Genotype	Source
L2056	<i>MATα/α</i>	Lallemand Pty Ltd, Australia
L2056 <i>MFA2 mfa2Δ</i>	<i>MATα/α, MFA2/mfa2Δ::KANMX4</i>	This study
L2056 <i>mfa2Δ mfa2Δ</i>	<i>MATα/α, mfa2Δ::KANMX4/mfa2Δ::KANMX4</i>	This study
C911A	L2056 <i>hoΔ MATα</i>	Walker et al. (2003)
C911D	L2056 <i>hoΔ MATα</i>	Walker et al. (2003)
C911A <i>mfa2Δ</i>	<i>MATα, mfa2Δ::KANMX4</i>	This study
C911D <i>mfa2Δ</i>	<i>MATα, mfa2Δ::KANMX4</i>	This study
C911A/C911D	<i>MATα/α, MFA2/MFA2</i>	This study
C911A <i>mfa2Δ/C911D</i>	<i>MATα/α, C911A mfa2Δ::KANMX4/ C911D MFA2</i>	This study
C911A/C911D <i>mfa2Δ</i>	<i>MATα/α, C911A MFA2/ C911D mfa2Δ::KANMX4</i>	This study
C911A <i>mfa2Δ/C911D mfa2Δ</i>	<i>MATα/α, C911A mfa2Δ::KANMX4/C911D mfa2Δ::KANMX4</i>	This study
AWRI1631	<i>N96 hoΔ MATα</i>	Borneman et al. (2008)
AWRI1631 <i>mfa2Δ</i>	<i>MATα, mfa2Δ::KANMX4</i>	Varela et al. (2012)
AWRI1633	<i>N96 hoΔ MATα</i>	AWRI, Australia
AWRI1633 <i>mfa2Δ</i>	<i>MATα, mfa2Δ::KANMX4</i>	This study
AWRI1631/AWRI1633	<i>MATα/α, MFA2/MFA2</i>	This study
AWRI1631 <i>mfa2Δ/1633</i>	<i>MATα/α, 31 mfa2Δ::KANMX4/33 MFA2</i>	This study
AWRI1631/1633 <i>mfa2Δ</i>	<i>MATα/α, 31 MFA2/33 mfa2Δ::KANMX4</i>	This study
AWRI1631 <i>mfa2Δ/1633 mfa2Δ</i>	<i>MATα/α, 31 mfa2Δ::KANMX4/33 mfa2Δ::KANMX4</i>	This study

Primers used in this study

Primers were sourced from Sigma Aldrich. The *MFA2* deletion cassette was amplified using primers *MFA2 Fwd* (TGATAGTTCCCTTTCCGTTAAGTG) and *MFA2 Rvs* (TGGCTCAAAACTTTTCACTTTAC), whilst the mating type of a strain was identified with three primers (Huxley et al. 1990): *MAT* (AGTCACATCAAGATCGTTATGG); *MATα* (ACTCCACTTCAAGTAAGAGTTG) and *MATα* (GCACGGAATATGGGACTACTTCG).

Construction of *MFA2* deletants

To construct *MFA2* deletion mutants in haploid strains AWRI1633, C911A and C911D and diploid strain L2056, the *mfa2Δ::KanMX* deletion cassette was PCR amplified from genomic DNA of AWRI1631 *mfa2Δ* (Peter et al. 2017) and used to replace the endogenous *MFA2* gene by

homologous recombination (Gietz and Woods 2002). Transformants were selected on YPD agar containing 200 mg L⁻¹ G418 sulfate (Ameresco) and confirmed by sequencing of the PCR amplicon (ORF::KanMX) encompassing the deletion cassette (Primers A and D; Table 2). The mating type was confirmed by PCR amplification across the HMR region (Huxley et al. 1990). Heterozygous diploid strain L2056 *MFA2/mfa2Δ* was sporulated in liquid medium (Codón et al. 1995) and individual tetrads dissected by micro manipulation (Guthrie and Fink 1991). L2056 *mfa2Δ/mfa2Δ* was then identified by PCR amplification of the *mfa2Δ::KanMX* deletion cassette for both alleles.

The following diploid wine yeast was constructed through mating the corresponding haploid *MATa* and *MATα* strains: C911A/C911D, C911A *mfa2Δ/C911D*, C911A/C911D *mfa2Δ*, C911A *mfa2Δ/C911D mfa2Δ*, AWRI1631/1633, AWRI1631 *mfa2Δ/1633*, AWRI1631/1633 *mfa2Δ* and AWRI1631 *mfa2Δ/1633 mfa2Δ*. Fresh cultures from YPD plates were mixed thoroughly on thin YE PD plates and incubated for 4 hours at 30 °C. Actively mating cells were isolated using a micromanipulator and their identity confirmed by sequence analysis of PCR amplicons encompassing the deletion cassettes.

Fermentation conditions

Tripletate 100 ml fermentations in CDGJM (75 mg N L⁻¹) were carried out at 28 °C using 250 ml flasks fitted with airlocks (Walker et al. 2003), with agitation. Fermentations were monitored by measuring residual sugar enzymatically (Walker et al. 2014) and considered finished when the residual sugars were <2 g L⁻¹. Growth was estimated at regular intervals by measurement of optical density (OD₆₀₀) at 600 nm. Yeast cultures at the end of fermentation were stained with propidium iodide and analyzed by flow cytometry for total cell numbers using a Guava easyCyte™ 12 HT FACS analyzer. Where haploid cultures tended to flocculate, they were sonicated for 30 seconds before staining. Dry cell weight was also determined in duplicate from terminal samples (Salmon 1997).

Statistical analysis

Fermentation duration, total cell count and biomass yield are presented as a mean of triplicates along with their standard deviation (SD). Student's t-test (two tailed) was used to identify statistically significant differences between two samples with a p-value <0.05.

4.4 Results

Effect of deletion of *MFA2* on fermentation performance of wine yeast derivatives

Previously we have reported that deletion of *MFA2* in the haploid wine yeast strain AWRI1631 resulted in a shortening of a nitrogen-limited fermentation by ~34% (Peter et al. 2017). To determine the dependence of this phenotype on strain background and ploidy, we first evaluated the fermentation performance of *MFA2* gene deletions in both haploid mating types of derivatives of two wine strains, N96 and L2056. Then the influence of gene dosage was assessed by comparing fermentation performance of the diploid versions, both homozygous and heterozygous for *MFA2*.

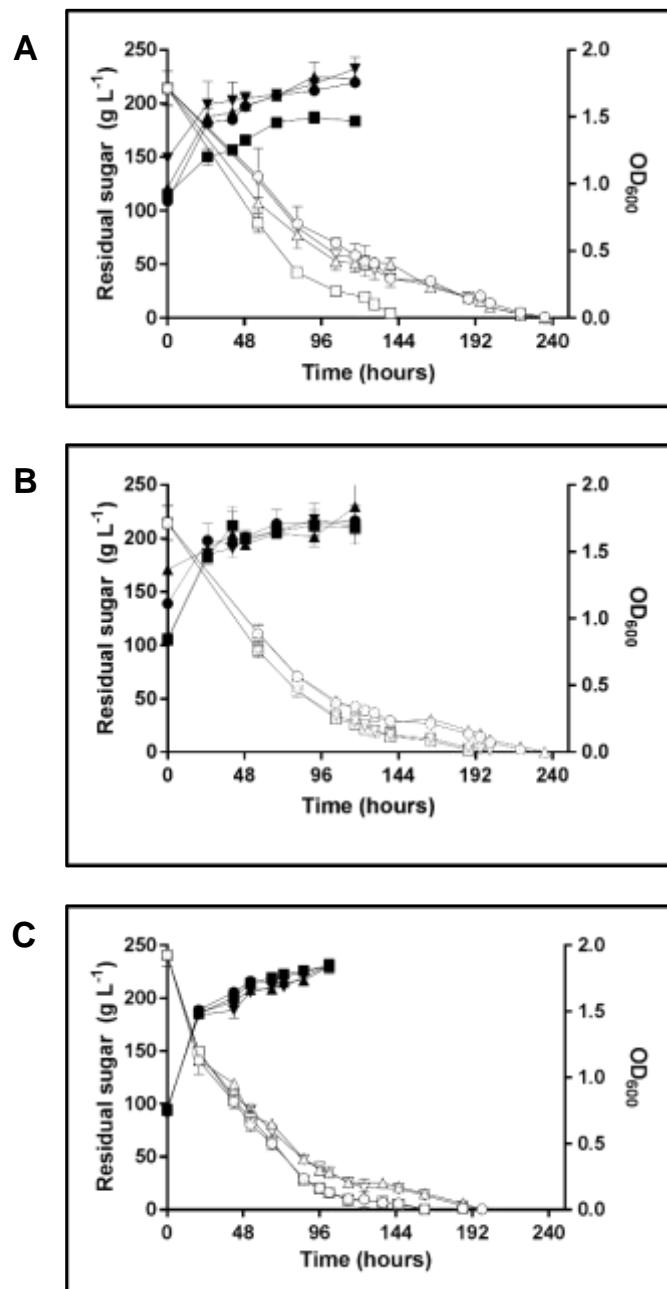


Figure 1. Fermentations of *MFA2* deletants. Fermentations (100 mL) were conducted in triplicate in CDGJM with 75 mg N L⁻¹. Sugar consumption was monitored enzymatically and reported as total sugars (open symbols). Growth was monitored as optical density at 600 nm (solid symbols). Data points are mean values of triplicate data with standard deviation as error bars. **(A)** AWRI1631 and AWRI1633 (haploids); Circle, AWRI1631; square, AWRI1631 *mfa2*; upright triangle, AWRI1631; inverted triangle, AWRI1633 *mfa2Δ*. **(B)** AWRI1631/AWRI1633 (diploids); Circle, AWRI1631/AWRI1633; square, AWRI1631 *mfa2Δ*/1633; upright triangle, AWRI1631/1633 *mfa2Δ*; inverted triangle, AWRI1631 *mfa2Δ*/AWRI1633 *mfa2Δ*. **(C)** C911A and C911D (haploids); Circle, C911A; square, C911A *mfa2Δ*; upright triangle, C911D; inverted triangle, C911D *mfa2Δ*.

Table 2. Fermentation duration, total cell count and biomass yield of AWRI1631, AWRI1633 and their *MFA2* deletants.

Strains	MT	FD (h)	FD% of WT	TCC (x 10 ⁷)	TBY (g L ⁻¹)	B / cell (pg)	B% of WT
AWRI1631	MATa	235 ± 0	100 ± 0	12.3 ± 0.2	4.1 ± 0.2	33.3	100
AWRI1631 <i>mfa2Δ</i>	MATa	146 ± 0*	62 ± 0	8.19 ± 0.2*	2.0 ± 0.4*	25.2	76
AWRI1633	MATa	230 ± 9	100 ± 0	14.0 ± 0.8	4.4 ± 0.0	31.7	100
AWRI1633 <i>mfa2Δ</i>	MATa	230 ± 9	100 ± 6	13.6 ± 0.7	4.5 ± 0.3	33.6	106
AWRI 1631/AWRI1633	MATa/α	230 ± 9	100 ± 0	9.60 ± 0.5	4.2 ± 0.1	44.5	100
AWRI1631 <i>mfa2Δ/1633</i>	MATa/α	190 ± 5*	82 ± 2	8.52 ± 1.2	2.9 ± 0.1*	34.0	77
AWRI1631/1633 <i>mfa2Δ</i>	MATa/α	235 ± 0	104 ± 0	9.44 ± 0.1	4.4 ± 0.2	46.6	105
AWRI1631 <i>mfa2Δ/1633 mfa2Δ</i>	MATa/α	201 ± 0*	87 ± 0	8.65 ± 0.9	3.2 ± 0.2*	37.0	83

WT, Wildtype; MT, Mating type; FD, Fermentation duration; TCC, Total cell count (cells mL⁻¹ ± SD); TBY, Total biomass yield; B, Biomass; Wildtype strains are written in bold text and *MFA2* deletants are written in plain text.

* Values that are significantly different to the wildtype (p< 0.05, two-tailed Student's t-test).

In the AWRI1631 and AWRI1633 backgrounds, only AWRI1631 *mfa2Δ* (MATa) was capable of reducing fermentation duration (by 38%), which is similar to our previous finding (Peter et al. 2017). For instance, deletion of *MFA2* from AWRI1633 (MATa) did not result in reduced fermentation as occurred for AWRI1631 *mfa2Δ* (c.a. 230 vs. 146 hrs, Fig. 1A and Table 2). To evaluate the influence of ploidy, diploid heterozygous and homozygous strains were made by mating wildtype or deletant forms of AWRI1631 and AWRI1633. Investigation of these diploids, identified strains with shortened fermentation duration when the *MFA2* deletion was in the AWRI1631 background. For example, AWRI1631/1633 *mfa2Δ* completed fermentation in a similar time to the wildtype (235 vs. 230 h, Fig. 1B and Table 2). Interestingly the heterozygous diploid (AWRI1631 *mfa2Δ/1633*) was slightly more efficient than the homozygous diploid (AWRI1631 *mfa2Δ/1633 mfa2Δ*), completing fermentation in 82% of the time of the wildtype (compared to 87%, Fig. 1B and Table 2). Conversely, deletion of *MFA2* from L2056 derivatives (C911A, C911D) did not shorten fermentation, either in the haploid (Fig. 1C and Table 3) or diploid homozygous or heterozygous deletions (see Appendix 11 and 12, Pg 194 and 195). Notably, strains C911A and

C911D, derived from the same background had a difference of 36 hours in fermentation duration (c.a. 162 vs. 198 hours, Fig. 1C and Table 3).

In addition to fermentation performance, deletion of *MFA2* in AWRI1631 affected growth, as estimated by a 25% reduction in optical density for AWRI1631 *mfa2Δ* compared to the wildtype (Fig. 1A). No effect was observed in AWRI1633 *mfa2Δ*. Total cell number, dry cell biomass and subsequently biomass/cell was calculated from terminal fermentation samples (Table 2 and 4). AWRI1631 *mfa2Δ* (*MATa*) exhibited a reduced total cell number and biomass yield and therefore the biomass per cell was less than the wildtype (c.a.76%). In AWRI1633 *mfa2Δ* (*MATa*) and AWRI1631/AWRI1633 *mfa2Δ* (heterozygous diploid) deletion of *MFA2* did not result in a reduction in the cell number or biomass yield. In fact, it resulted in a slight increase in biomass yield (~6%). However, deletion of *MFA2* in heterozygous AWRI1631 *mfa2Δ*/AWRI1633 and homozygous AWRI1631 *mfa2Δ*/AWRI1633 *mfa2Δ* diploids resulted in a reduction of total cell number and biomass yield, with biomass per cell calculated to be 77 and 83% of the wildtype AWRI1631/1633, respectively (Table 2).

In L2056, heterozygous and homozygous *MFA2* deletants resulted in a reduced total cell number, biomass yield and biomass per cell (c.a.77 and 85%). C911D *mfa2Δ* (*MATa*) and C911A/C911D *mfa2Δ* had similar total cell numbers to their wildtype, however, a significant difference in biomass yield, with biomass per cell reduced to 59 and 70% of their respective wildtypes. Compared to C911A *mfa2Δ* (*MATa*), C911A *mfa2Δ*/C911D and C911A *mfa2Δ*/C911D *mfa2Δ*, deletion of *MFA2* did not result in a noticeable difference in cell number or biomass yield (Table 3).

Table 3. Fermentation duration, residual sugar, total cell count and biomass yield of L2056, C911A, C911D and their *MFA2* deletants.

Strains	MT	FD (h)	FD % of WT	TCC (x 10 ⁷)	TBY (g L ⁻¹)	B / cell (pg)	B% of WT
L2056	<i>MATa/a</i>	142 ± 6	100 ± 0	9.6 ± 1.0	3.5 ± 0.2	36.4	100
L2056 <i>MFA2 mfa2Δ</i>	<i>MATa/a</i>	145 ± 0	102 ± 0	13.2 ± 1.2*	3.7 ± 0	28.1	77
L2056 <i>mfa2Δ mfa2Δ</i>	<i>MATa/a</i>	151 ± 1	102 ± 3	14.2 ± 0.2*	4.3 ± 0.3*	30.9	85
C911A	<i>MATa</i>	162 ± 0	100 ± 0	19.3 ± 0.9	5.9 ± 1.4	30.6	100
C911A <i>mfa2Δ</i>	<i>MATa</i>	162 ± 0	100 ± 0	21.1 ± 1.1	7.0 ± 1.4	33.2	108
C911D	<i>MATa</i>	198 ± 0	100 ± 0	20.6 ± 1.4	6.7 ± 1.3	32.6	100
C911D <i>mfa2Δ</i>	<i>MATa</i>	198 ± 0	100 ± 0	20.4 ± 1.3	3.9 ± 0.1*	19.1	59
C911A/C911D	<i>MATa/a</i>	186 ± 0	100 ± 0	13.4 ± 1.4	4.0 ± 0.1	29.5	100
C911A <i>mfa2Δ/C911D</i>	<i>MATa/a</i>	186 ± 0	100 ± 0	12.8 ± 1.2	3.7 ± 0.1*	29.0	98
C911A <i>/C911D mfa2Δ</i>	<i>MATa/a</i>	186 ± 0	100 ± 0	12.5 ± 1.3	2.6 ± 0.7*	20.5	70
C911A <i>mfa2Δ/ C911D mfa2Δ</i>	<i>MATa/a</i>	186 ± 0	100 ± 0	12.9 ± 0.8	3.7 ± 0*	28.7	97

WT, Wildtype; MT, Mating type; FD, Fermentation duration; TCC, Total cell count (cells mL⁻¹ ± SD); TBY, Total biomass yield; B, Biomass; Wildtype strains are written in bold text and *MFA2* deletants are written in plain text.

* Values that are significantly different to the wildtype ($p < 0.05$, two-tailed Student's t-test)

4.5 Discussion

Role of *MFA2* deletion in fermentation is specific to AWRI1631 (*MATa* haploid of N96)

In this study, we report that deletion of *MFA2* in AWRI1631 (*MATa* haploid of N96) shortened fermentation duration by 38% in CDGJM supplied with limited (75 mg N L⁻¹) nitrogen. To determine whether mating type influenced fermentation phenotype, *MFA2* was deleted in AWRI1633 (*MATa* haploid of N96). Replacement of the *MFA2* gene in AWRI1633 by the *mfa2::KanMX* deletion cassette (amplified from AWRI1631 *mfa2Δ*), did not affect fermentation, with the mutant behaving similarly to the wildtype. Gene dosage was also investigated to assess whether deletion of one or both copies of *MFA2* in the diploid could influence fermentation duration and growth. Interestingly, in the diploid (AWRI1631/1633) strains capable of reduced fermentation only when *mfa2Δ* originated from AWRI1631. Minor differences were observed between the

heterozygous (AWRI1631 *mfa2Δ*/AWRI1633) and homozygous (AWRI1631 *mfa2Δ*/AWRI1633 *mfa2Δ*) diploids, which completed fermentation in 18 and 13% less time taken than the wildtype, respectively. These findings confirm that the shortened fermentation of the *MFA2* deletion in N96 derivatives was specific to being carried on genetic material derived from a *MATa* cell. The finding that homologous replacement of the *MFA2* gene by the *mfa2Δ::KanMX* cassette (from AWRI1631 *mfa2Δ*) in AWRI1633 (*MATa*) did not result in altered fermentation alludes to a functional and interactional difference of *MFA2* in the two cell types. *MFA2*, together with *MFA1*, is responsible for the production of mating a-factor pheromone solely in *MATa* cells (Michaelis and Herskowitz 1988) whereas, *MFα1* and *MFα2* produce mating α-factor pheromone merely in *MATα* cells. However, deletion of *MFA1* from AWRI1631 (*MATa*) did not result in a similar reduction in fermentation duration (Peter et al. 2017), and further investigation of *MFα1* and *MFα2* is required. Previous studies have reported that the expression of ‘a-specific’ genes is repressed in *MATa* and diploid (*MATa/MATα*) cells in cell-type regulation (Galgoczy et al. 2004; Johnson 1995; Johnson and Herskowitz 1985).

These findings are novel as this is the first report of *MFA2* having dual effects, on mating and fermentation, specifically in *MATa* cells. Interestingly, *MATa2* is known to exhibit pleiotropy, with reports of the gene being involved in cross talk between cell-type regulation and the HOG pathway for pheromone-responsive genes in *MATα* cells during mating (Baltanás et al. 2013; Galgoczy et al. 2004; O'Rourke and Herskowitz 1998). Another gene, *MDF1* (*YCL058C*), formally known as *FYV5*, is also involved in the regulation of mating. *Mdf1p* is thought to promote vegetative growth in *MATα* cells in a rich medium, by suppressing mating efficiency, through binding to *MATa2* (Li et al. 2010). Furthermore, Li et al. (2014) propose that *Mdf1p* shortens the lag phase, by interacting with *Snf1p*, central to non-fermentable carbon utilisation, enabling the rapid consumption of glucose during early exponential growth in rich media. It is unclear whether deletion of these genes, as well as others such as *MCM1*, involved in cell-type regulation, affected fermentation in other strains. Also, how deletion of *MFA2* shorten fermentation duration or provided yeast with an adaptive advantage in grape juice to enable improved fermentation performance is yet to be determined.

AWRI1631 *mfa2Δ* have improved fermentation capacity for a given cell population

Deletion of *MFA2* in AWRI1631 resulted in shortened fermentation under limited nitrogen conditions. However, growth was significantly affected resulting in decreased cell number and biomass, with biomass per cell reduced by 24%. Our findings are contrary to the consensus that

fermentation rate is correlated with biomass, which is particularly important in nitrogen deficient conditions, common to problematic 'sluggish' fermentations. Thus in comparison, Varela et al. (2004) demonstrated that reduced metabolic activity (sugar catabolism) under limited nitrogen could be ameliorated by concentration of the biomass. In general, maximum fermentation rate (as CO₂ evolution) is related to maximum cell population for a given medium (Albertin et al. 2011), which further supports the hypothesis of a 'critical' biomass being required for efficient fermentation. This relationship holds true for yeast with high nitrogen demand (requirements), whereby nitrogen supplementation leads to higher fermentation rates through increased biomass and overall population numbers. However, in the case of yeast with lower nitrogen demand, higher fermentation rates are achieved with fewer cells, which are more metabolically active through increased glycolytic flux (Gutierrez et al. 2012). Canelas and co-workers (2010) made similar observations for chemostat cultures of CEN.PK113-7D having increased glucose uptake and reduced biomass yield compared to YSBN2. Further, expression of genes associated with biosynthesis and degradation of amino acid and protein significantly increased in CEN.PK113-7D. The authors proposed that as protein synthesis is an energy consuming process requiring more ATP, the strain increases glycolytic flux by driving more ATP and co-factors into protein (enzyme) synthesis rather than biomass formation (Canelas et al. 2010).

Similarly, deletion of *MFA2* in AWRI1631, which results in greater fermentation and reduced biomass formation may be due to increased protein turnover. Given that *MFA2* does interact with *UBI4*, encoding ubiquitin, which is conjugated to proteins targeted for degradation via the ubiquitin-26S proteasome (Drygin website; (Costanzo et al. 2010; Koh et al. 2010), it would be worth to investigate whether ubiquitin and protein turnover are affected in AWRI *mfa2Δ*.

Fermentation improvement in yeast by *MFA2* deletion is dependent upon genotype

To examine whether the effect of *MFA2* deletion in reducing fermentation duration was universal, we constructed and evaluated the fermentation performance of an *MFA2* deletion in C911A (*MATα*) and C911D (*MATα*); haploids of wine yeast Lalvin L2056. The failure of both *mfa2Δ* strains to shorten fermentation confirmed that the fermentation phenotype was not only reliant on *MATα* cell types but dependent on yeast genotype. This phenotypic variation between the two strains may reflect the degree of heterozygosity between L2056 and N96 (Borneman et al. 2016). Whole genome sequencing has identified a high level of inter-strain variation, as a result of evolutionary processes including single nucleotide polymorphisms (SNP's), Ty elements (number and arrangement), gene transfer, as well as insertion and deletion events. These variations are

sometimes specific to a strain, and the high variability in genes may be a reason for phenotypic diversification (Borneman et al. 2011; Borneman et al. 2008; Novo et al. 2009).

Although deletion of *MFA2* in C911D (*MAT α*) did not result in the same ability to ferment sugars as the original *mfa2 Δ* deletion in AWRI1631 (*MAT α* haploid of N96), similarities between the strains were noted for growth and biomass formation. In both yeast genotypes, deletion of *MFA2* in the *MAT α* strain resulted in reduced biomass formation. Conversely, deletion of *MFA2* in the corresponding *MAT α* haploids, AWRI1633 *mfa2 Δ* and C911A *mfa2 Δ* , resulted in a slight increase in biomass per cell (6-8% wildtype) whilst fermentation performance was comparable to the wildtypes.

In the diploid N96 and L2056 derivatives, *mfa2 Δ* on the *MAT α* allele resulted in a reduction in the biomass yield. Also, we observed that heterozygous deletions were more affected than the homozygous deletions, which may be due to a ‘rescue effect’ rendered by the *MFA2* deletion originating from the *MAT α* strain, which results in increased biomass formation. For example, the calculated biomass per cell of the heterozygous diploids were 77, 70 and 77 % of the wildtype for AWRI1631 *mfa2 Δ /1633*, C911A/C911D *mfa2 Δ* and L2056 *mfa2 Δ* , respectively. Whereas the homozygous diploids resulted in 83, 85 and 97% of biomass per cell of the wildtype for AWRI1631 *mfa2 Δ /AWRI1633 mfa2 Δ* , C911A *mfa2 Δ /C911D mfa2 Δ* and L2056 *mfa2 Δ /mfa2 Δ* , respectively. Although L2056 and C911A/C911D strains are expected to be similar, they appear to differ in behaviour. Such a difference could be due to genetic differences arising through segregation of chromatids and genetic recombination during several rounds of mating and sporulation (meiosis) performed to construct these strains. Also with the heterozygous L2056 deletion, we are unsure of the mating type of allele bearing the *MFA2* deletion.

Gene dosage affecting fitness and fermentation profile in diploids

Past studies have investigated gene dosage to assess fitness profile, by deleting a copy of a gene in a diploid, which could result in growth advantage (haploproficiency) or disadvantage (haploinsufficiency) (Delneri et al. 2008; Piggott et al. 2011; Pir et al. 2012). Competitive fitness studies of the diploid heterozygous deletion collection grown in different conditions of limiting and non-limiting nutrients (Table 5) have shown the *MFA2/mfa2 Δ* strain to be haploproficient (Delneri et al. 2008; Piggott et al. 2011).

Table 5. Fitness proficiency of heterozygous *MFA2* deletants in different culture conditions reported from previous studies.

Deletants (heterozygous)	Fitness proficiency	Culture condition	Reference
<i>MFA2/mfa2Δ</i>	HP	Synthetic grape juice	(Piggott et al. 2011)
<i>MFA2/mfa2Δ</i>	HP	C- and N- limitation	(Delneri et al. 2008)
<i>MFA2/mfa2Δ</i>	HP	C-, N- and P- limitation (dilution rate in chemostat 0.1,0.2 h ⁻¹)	(Pir et al. 2012)

HP denotes haploproficiency, C-, N-, P- denotes carbon, nitrogen and phosphorous respectively.

Contrary to these findings, specifically of Piggott et al. (2011) where research was in a synthetic grape juice, we have demonstrated that *MFA2/mfa2Δ* had reduced fitness in terms of growth (reduced cell number and biomass) although it was still capable of fermenting sugars quicker than the wildtype (conditional to the *mfa2Δ* originating from the *MATα* wildtype). The exact construction method of the heterozygous deletion strains (BY4743) used by Piggott et al. (2011) are not stated, thus it is possible that the differences observed may be due to the *mfa2Δ* carrying the *MATα* allele or more likely; genetic differences between the strains across the two studies or differences in cultural conditions and experimental design. In addition this study tested individual strains rather than looking at fitness in a pooled collection of the strains. When strains are evaluated individually for fermentation performance, growth does not necessarily relate to increased fermentation capacity (Canelas et al. 2010; Gutierrez et al. 2012). Here we report that deletion of *MFA2* in AWRI1631 (*MATα*) reduced fermentation duration under limited nitrogen conditions; this is a desirable attribute given that the yeast is more likely to succeed in nitrogen deficient juices and less liable to undergo starvation, leading to 'sluggish' or even 'stuck' fermentation.

In summary, strains which have a lower nitrogen demand, are considered beneficial, given that they do not require the same degree of supplementation as those of higher nitrogen demand and are likely to complete a nitrogen-limited fermentation with fewer difficulties. Thus seeking the genetic basis for such differences in nitrogen efficiency involving the *MFA2* gene was highlighted. We have demonstrated that deletion of *MFA2* gene shortens fermentation duration. However, the ability to do so was dependent upon genotype and mating type (i.e. only in AWRI1631, *MATα*). These findings pose several questions as to why the improved phenotype associated with this

particular gene deletion is limited to this strain. RNAseq analysis, whilst beyond the scope of this study, would be worthwhile in understanding the transcriptional differences between these strains during fermentation. Before concluding that *mfa2Δ* does not improve fermentation efficiency in other wine strains, it would be important to investigate the effect of this deletion in other wine yeast backgrounds. Also, further investigation into the expression of the genes associated with processes like biosynthesis and degradation of amino acid and proteins (as reported in Canelas et al. (2010)) might aid in uncovering the genetic basis of fermentation efficiency of AWRI1631 *mfa2Δ*.

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CHAPTER 5

General discussion, conclusions and future directions

5.1 General discussion and conclusions

Lack of sufficient yeast assimilable nitrogen is often the root cause of stuck and sluggish fermentations. The development of wine yeast that can efficiently utilise available nitrogen in deficient musts to maximise sugar catabolism would be of great benefit. These yeasts together with better management practices, such as appropriate timing and choice of nitrogen supplementation, will significantly reduce the occurrence of problematic fermentations and the associated reduction in wine quality. The availability of nitrogen efficient strains would also ideally reduce the wine maker's dependence on supplementation, and enable increased efficiency in vinification and tank throughput. Economic savings would be made due to a reduction in labour and processing costs (re-inoculation and supplementation) and a need to blend away poor quality wines, and would lead to overall increase in wine quality and value.

The generation of nitrogen efficient strains requires a greater understanding of nitrogen metabolism in relation to wine making fermentations. This project used the approach of screening prototrophic deletion libraries in laboratory and wine yeast backgrounds, to identify genes influencing fermentation efficiency under different nitrogen conditions. This effort resulted in revelation of cellular processes that influence fermentation efficiency and which are specific to a nitrogen insufficient environment.

5.1.1 Functional genomic tools identify key genes influencing fermentation in sufficient and limiting nitrogen conditions

Deletants with reduced fermentation duration were identified from both wine and laboratory yeast libraries with sufficient and limited nitrogen availability.

As a result of the screening experiments, we determined the following:

1. High throughput micro-fermentation identified gene deletions that allowed twice the sugar of the wildtype to be fermented (LYDL_HN, 748 deletants; WYDL_HN, 299 deletants and WYDL_LN, 257 deletants). Examination of the performance of deletants in limiting nitrogen revealed a subset of 127 genes that only exhibited improved fermentation in low nitrogen condition (WYDL). Deletants were also identified, which worsen fermentation performance (50% less sugar fermented vs the wildtype): (LYDL_HN, 501 deletants; WYDL_HN; 205 deletants and WYDL_LN; 187 deletants). Several of the fermentation essential genes identified as part of the laboratory yeast Fermentome (Walker et al. 2014), have also been demonstrated to be required by wine yeast in limited nitrogen. By comparison, the set of genes that

reduced fermentation time upon deletion are potential candidates for yeast strain improvement, whether as natural gene variants (having beneficial SNP/indels), CRISPR genome editing, mutagenesis or breeding, etc.

2. Comparison of the genes (those capable of catabolising more sugar within a specified time frame) from the micro-screens, identified genes that were common across strain backgrounds and/or were independent of initial nitrogen concentration. This data set was also compared to published data sets revealing genes shown to be regulated at the transcriptional level under similar conditions or affected by strain background. A Gene Ontology analysis of each data set revealed that genes common to WYDL_HN and WYDL_LN were enriched for genes involved in protein transport, catabolism, ubiquitination and cell communication. Of those associated with cell communication, 6 genes (*SHO1*, *BEM2*, *KSS1*, *ROM1*, *FLO11* and *MFA2*) were highlighted to be targets of the MAPK pathway. Further GO analysis of the 161 and 127 genes unique to WYDL_HN and WYDL_LN conditions, identified enrichment of genes for processes mainly involved in catabolic process and regulation in the high nitrogen condition. In the low nitrogen conditions, the gene set was enriched for processes such as ubiquitination and protein turnover.

5.1.2. Screening of the wine yeast deletion library for fermentation efficient mutants in limiting nitrogen

Physiological studies have traditionally been conducted in laboratory yeast, which are easily genetically manipulated, and have enabled identification of a myriad of pathways related to yeast metabolism. This research is now being undertaken in industrial yeast to better understand the adaptive mechanisms behind fermentation (brewing, bread-making, sake and wine production).

In this study, the primary focus was the investigation of how wine yeast withstand fermentation stresses associated with nitrogen deficiency – a common cause of sluggish and stuck fermentations. A wine yeast deletion library was screened when insufficient nitrogen was available, to identify genes that confer fermentation efficiency when under nitrogen limitation. From the micro-fermentation screen (WYDL_LN), we shortlisted 96 candidates that were observed to utilise significantly more sugar than the wildtype, having approximately half the residual sugar of that of the wildtype, these were evaluated in laboratory-scale fermentation trials (100 mL). The main findings were the:

1. Identification of 15 strains that were fermentation efficient, reducing fermentation duration by 15-59%.
2. Gene Ontology analysis revealing enrichment for genes in processes related to ubiquitination and protein modification/turnover or metabolic processes: protein polyubiquitination (*UBI4*, *UBC13* and *MMS2*), protein deubiquitination (*UBP7*, *UBI4* and *BRO1*), protein modification by small protein conjugation (*UBP7*, *UBI4*, *BRO1*, *MMS2* and *UBC13*), post-translational protein modification (*UBP7*, *TPK2*, *UBI4*, *BRO1*, *MMS2* and *UBC13*), protein metabolic process (*UBP7*, *TPK2*, *UBI4*, *BRO1*, *MFA2*, *MMS2*, *MKT1*, *EAR1*, *UBC13* and *MRP17*) and endosomal transport (*MVB12*, *EAR1* and *BRO1*).

5.1.3 Investigation of an *MFA2* deletion in AWRI1631

Of the 15 wine deletants identified as fermentation efficient in limiting nitrogen, a deletion of *MFA2* (mating a-factor; involved in mating and cell-type regulation), resulted in a 55% decrease in fermentation duration. *MFA2* was further investigated to better understand how its deletion conferred fermentation efficiency, since this gene and the process of yeast mating have not previously been associated with fermentation. We also investigated the fermentation performance of the functionally redundant *MFA1*, encoding a-factor. Fermentation dynamics of *mfa1Δ* and *mfa2Δ* were studied in detail as well as the expression profile of *MFA1* and *MFA2*. The following observations were noted:

1. Unlike *MFA2*, deletion of *MFA1* did not confer fermentation efficiency.
2. Gene expression studies revealed upregulation of *MFA2* in AWRI1631 when grown in CDGJM with limiting nitrogen. Deletion of *MFA2* from AWRI1631 resulted in an increased expression of *MFA1*.
3. *MFA1* was upregulated by ~6 to 7-fold in the *MFA2* deletant at the initial sampling point (128 g L⁻¹ residual sugar). Although *MFA1* expression declined subsequently, it remained ~4-fold higher than the wildtype

5.1.4 Fermentation performance of *mfa2Δ* is dependent on strain background, mating type and gene dosage

Further to understanding the role of *MFA2* in fermentation, studies were conducted to determine whether the effect of *mfa2Δ* was dependent on mating type, genotype, and/or gene dosage. Homozygous and heterozygous diploid *MFA2* deletants were constructed in genetic backgrounds derived from N96 (AWRI1633 MAT α ; AWRI1631 MAT α) and L2056 (C911A MAT α ;

C9 11D MAT_a). These strains were compared to the equivalent *MFA2* wildtype strains in 100 mL fermentations. The following observations were made:

1. The fermentation efficient phenotype associated with *mfa2Δ* was dependent upon genotype and mating type, with the phenotype confined to AWRI1631 MAT_a.
2. The diploid AWRI1631/1633 strains enabled quicker fermentation when the *mfa2Δ* allele originated from the 'a' mating type strain (AWRI1631 *mfa2Δ MATa*).

5.2 Future directions

1. The involvement of *MFA2* and potentially the mating pheromone pathway in fermentation is novel. We hypothesize that mating pheromone production and processing is an energy consuming process. Therefore, down-regulating the synthesis/processing of mating a-factor, could lead to a saving in energy. This energy might be redirected to other processes such as sugar uptake and catabolism. Deletion of *MFA2* resulted in significantly increased transcription of *MFA1* (functionally redundant to *MFA2*), but deletion of *MFA1* did not result in fermentation improvement. Over-expression of *MFA1* when wildtype *MFA2* is present remains to be examined. The results suggest that deletion of *MFA2* could also affect other interacting genes. Of these *GAL10*, *GID7*, *HXT5*, *STL1* and *PCL6* are known to be involved in carbohydrate metabolism/transport and storage (Costanzo et al. 2016). Further investigation of these genes, and specifically their interaction with *MFA2* during fermentation is required to determine the mechanism(s) by which *MFA2* influences sugar uptake and metabolism.
2. Deletion of *MFA2* from the wine yeast strain L2056 was found not to improve fermentation efficiency. It is also important to generate this deletion in several other strains (as MAT_a haplotypes) and evaluate their performance before determining that this fermentation phenotype is specific to only AWRI1631 *mfa2Δ*. Transcriptomic analysis (RNA-Seq) would help to decipher the transcriptional differences between AWRI1631 *mfa2Δ* and wildtype during fermentation, enabling a better understanding of the dual function of this gene in cell-type recognition during mating and fermentation.
3. High-through screening exercises are powerful tools for identification of gene function through phenotypic studies, as they allow simultaneous evaluation of thousands of strains. This study encompassed identification of genes through genome-wide

screening of fermentation performance in juice-like conditions with different nitrogen content. These genes (and processes) add to our current understanding of the role of nitrogen in fermentation. Not only were genes identified as critical to fermentation (leading to protraction when deleted), several were shown to result in fermentation proficiency, and represent likely targets using non-GM techniques in the generation of nitrogen efficient strains.

4. Screening of the laboratory yeast deletion library when sufficient nitrogen was available led to identification of 548 genes that enabled a ~50% reduction in fermentation duration. These genes were absent in the wine yeast deletion collection. Construction of these in a wine yeast background and subsequent evaluation of fermentation performance is worthwhile to determine whether they are potential candidates for wine yeast improvement. Also, growth of the laboratory yeast deletion library in a nutrient rich media without selective pressure, resulted in plasmid instability, possibly due to the rapid multiplication of yeast cells, failing to replicate the plasmid along with the chromosomal DNA (Kleinman et al. 1986; Caunt et al. 1988; Impoolsup et al. 1989). Contrary to Mülleder et al. (2012), this study suggests that the library is better grown in minimal media (ammonium sulfate only) to ensure retention of plasmid-based prototrophy.
5. In addition to the deletants identified as able to reduce fermentation duration, deletants extending fermentation were also identified in each micro-screen (LYDL_HN, 501 deletants; WYDL_HN, 205 deletants and WYDL_LN, 187 deletants), which were not tested further. These genes are worth further investigating to help increase understanding of the underlying processes and pathways involved.
6. Laboratory-scale (100 mL) testing of the deletants shortlisted from micro-fermentations identified candidates in the limiting nitrogen (75 mg L^{-1}) conditions. By comparison in sufficient nitrogen (450 mg L^{-1}), fermentation of the deletants and the wildtype was so rapid reducing the resolution between the strains. Therefore it is suggested that the available nitrogen be reduced from 450 mg L^{-1} to ~ 300 mg L^{-1} (industry standard), for similar investigations to provide greater separation of the strains.
7. In the micro-fermentations, 257 (WYDL_LN) and 299 (WYDL_HN) deletants were identified as potentially fermentation efficient, however only 92 were tested from each as this arbitrary number conveniently fit in an automated 96-flask fermentation platform (Peter et al. 2017) with the controls. The number of identified and confirmed

- candidates has therefore likely not been exhausted and additional genes may be found through further evaluation.
8. As is the case for deletion libraries, over expression libraries (Gelperin et al. 2005; Jones et al. 2008) serve as a powerful tool to study gene function. The function of the identified genes could also be further determined by using an over-expression library to examine impact on a fermentation efficient phenotype, including whether it is restored to the wildtype.
 9. The 15 wine yeast gene deletions identified to confer fermentation efficiency in limited nitrogen are worthy targets for industrial strain improvement. Should genome editing became acceptable for industrial strain construction and use in the wine industry (Bao et al. 2014), deletants constructed by this approach may be able to be used directly for the management of problematic fermentations associated with nitrogen deficient juice. Alternatively these genes may serve as targets for non-recombinant approaches. The 15 genes were mainly associated with protein turnover and ubiquitination. Of the genes involved in the degradation pathway, *BRO1*, *MVB12* and *EAR1* have been extensively studied. Reports have shown that these genes regulate the degradation of amino acid permeases through ubiquination. Deletion of some of these genes disrupts this pathway reducing the rate of permease turnover (Nikko et al. 2003; Oestreich et al. 2007; Léon et al. 2008). In the conditions used in this study this may lead to increased retention of permeases on the cell membrane, thus increasing nutrient (i.e. nitrogen and sugar) uptake. The present study could be extended by investigating the ubiquitination process in these deletants during fermentation by using antibodies to study cellular localisation of plasma membrane transporters which might reveal the connection between ubiquitination and fermentation efficiency.
 10. Gene deletion may affect other interacting genes, therefore investigating the key genes interacting with those highlighted genes may address how these deletants elicit their effect on fermentation efficiency. Traditional gene expression profiling by methods such as qPCR is effective in defining the expression pattern of genes during fermentation. A comparison of changes in the expression levels of interacting genes (mainly involved in adaptation to low nitrogen, nitrogen metabolism, stress regulation, sugar uptake and metabolism) in the wildtype and deletants would help define the role of the deletions in the network. Alternatively, gene array techniques such as micro or macro arrays or RNA sequencing that permit high throughput genome wide transcript

profiling could be used to study the expression of all annotated genes at the same time. Past comparative transcriptomic analysis has enhanced understanding of how yeasts differ in their physiological response to nitrogen composition/content during fermentation (Rossignol et al. 2003; Mendes-Ferreira et al. 2007a; Barbosa et al. 2015). Rossignol et al. (2003) reported over a thousand genes that changed expression at different stages of fermentation to adapt to the changing environment. Therefore genome wide expression profiling of the identified deletants during fermentation would help identify the key gene sets and processes contributing to fermentation proficiency in low nitrogen conditions.

11. The genetic basis of a specific phenotype can be identified through QTL mapping. This technique has been successfully used to identify the genetic basis for physiological traits in nitrogen consumption (Brice et al. 2014; Jara et al. 2014; Cubillos et al. 2017). Similar to these studies, to understand the effect of genes influenced by high nitrogen and low nitrogen during fermentation, bi-parental/ multi-parental crosses between the high nitrogen and low nitrogen users can be used to identify the genetic basis of variation in the F2 progeny, which will help relate the nitrogen efficiency phenotype to genotype.

Overall, this study has identified several genes influencing fermentation in various nitrogen conditions and strains backgrounds. We have identified gene sets that are needed for successful fermentation in high and low nitrogen conditions. Deletions that resulted in protracted fermentation, highlight the importance of those genes in stressful conditions such as low nitrogen. The set of gene deletions that positively influence fermentation are potential candidates for yeast strain improvement and the knowledge generated here could be used to formulate strategies to produce improved wine yeast for the future.

APPENDICES

Appendix 1. Micro-fermentation data of the LYDL_HN (all candidates).

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	102 h	
YAL002W	VPS8	0.95	1.19	1.35	1.60	1.70	1.90	4.29	3.37	49.28	15.27	53.57	16.64	310.36		
YAL004W	YAL004w	0.90	1.20	1.41	1.73	1.88	2.10	3.61	0.94	25.85	-0.03	29.46	0.91	196.73		
YAL005C	SSAT	0.90	1.17	1.37	1.70	1.83	2.04	11.99	3.34	50.16	29.13	62.14	32.47	296.70		
YAL007C	ERP2	0.95	1.18	1.43	1.79	1.95	2.16	3.16	1.56	28.96	8.67	32.12	10.23	220.25		
YAL009W	FOD14	0.91	1.20	1.42	1.73	1.87	2.07	7.96	1.62	36.53	7.66	44.49	9.28	231.12		
YAL010C	MDM10	0.92	1.08	1.24	1.58	1.73	1.95	7.98	1.68	42.70	12.51	50.69	14.20	268.66		
YAL011W	SWC3	0.90	1.04	1.19	1.52	1.67	1.90	31.88	16.39	78.25	59.17	110.13	75.56	217.41		
YAL012W	CYS3	0.86	1.10	1.08	1.06	1.30	1.48	21.46	11.08	65.79	47.89	87.25	58.97	301.29		
YAL013W	DEP1	0.88	1.12	1.30	1.59	1.72	1.91	20.82	10.47	65.98	59.69	86.80	70.16	197.36		
YAL014C	SYNB	0.94	1.12	1.31	1.71	1.87	2.09	3.77	1.31	28.30	2.83	32.06	4.13	210.20		
YAL015C	NTG1	0.89	1.10	1.24	1.55	1.67	1.88	7.16	1.66	39.40	10.57	46.56	9.01	245.97		
YAL016C_B	YAL016c_b	0.85	1.08	1.16	1.63	1.80	1.94	16.37	5.00	69.13	32.31	85.49	37.32	329.74		
YAL016W	TP03	0.95	1.01	1.05	0.95	0.96	0.93	123.63	122.41	149.02	114.63	272.64	237.04	413.99		
YAL017W	PSK1	0.90	1.15	1.35	1.68	1.81	2.01	16.14	1.87	62.23	23.90	78.38	17.26	213.62		
YAL018C	YAL018c	0.92	1.17	1.33	1.65	1.79	2.00	6.34	1.41	38.41	8.11	44.76	9.52	235.37		
YAL019W	FUN30	0.89	1.10	1.28	1.64	1.78	2.00	8.46	1.94	40.97	17.39	49.43	19.32	199.32		
YAL020C	ATS1	0.89	1.15	1.36	1.71	1.85	2.05	13.67	1.87	65.48	19.64	79.14	21.51	217.19		
YAL021C	CCR4	0.89	1.09	1.12	1.22	1.38	1.49	81.62	77.21	104.20	124.61	185.82	201.83	386.01		
YAL022C	FUN26	0.89	1.09	1.25	1.53	1.70	1.94	2.21	1.20	16.63	-0.42	18.84	0.78	164.88		
YAL023C	PMT2	0.88	1.00	1.09	1.29	1.39	1.57	5.22	1.26	34.84	5.87	40.06	7.14	186.79		
YAL024C	LTE1	0.84	1.07	1.09	1.27	1.54	1.84	23.39	9.97	76.96	52.90	100.35	62.87	314.74		
YAL026C	DRS2	0.88	1.08	1.23	1.52	1.63	1.82	16.25	2.86	53.59	21.91	69.84	24.77	271.22		
YAL027W	SAW1	0.91	1.11	1.33	1.57	1.73	1.92	11.70	1.29	48.92	9.87	60.62	11.16	218.15		
YAL028W	FRT2	0.89	1.15	1.38	1.72	1.86	2.07	10.37	1.47	45.32	7.95	55.69	9.42	234.02		
YAL029C	MYO4	0.94	1.12	1.29	1.58	1.71	1.94	9.36	1.47	44.26	8.57	53.62	10.04	193.06		
YAL030W	SNC1	0.91	1.14	1.33	1.71	1.85	2.08	3.57	1.41	23.15	2.46	26.72	3.86	236.39		
YAL031C	GIP4	0.89	1.12	1.30	1.61	1.76	1.98	3.24	1.26	29.52	0.69	32.76	1.35	242.63		
YAL034C	FUN19	0.90	1.15	1.33	1.65	1.79	2.01	10.77	1.53	46.84	13.43	57.61	14.95	243.27		
YAL035W	FUN12	0.93	1.18	1.17	1.24	1.57	1.87	47.93	29.86	106.84	92.72	154.77	122.58	385.40		
YAL036W	FUN12	0.96	1.11	1.13	1.16	1.43	1.75	39.20	17.47	88.54	68.72	127.75	86.18	375.21		
YAL036C	RBG1	0.88	1.13	1.30	1.67	1.84	2.08	1.75	1.19	9.74	-1.21	11.48	-0.02	230.07		
YAL037C_A	YAL037c_a	0.90	1.18	1.31	1.70	1.90	2.03	13.70	5.43	65.62	35.73	79.32	41.16	324.86		
YAL037W	YAL037w	0.93	1.16	1.41	1.76	1.92	2.14	7.65	1.48	36.43	12.07	43.48	13.49	217.77		
YAL038C	CTC3	0.89	1.03	1.05	0.93	0.98	0.98	16.48	2.05	126.35	119.25	140.38	246.60	266.37	569.46	
YAL040C	CLN2	0.88	1.09	1.25	1.61	1.78	2.00	14.20	1.58	52.50	18.22	69.29	19.43	299.31		
YAL042W	YAL042w	0.88	1.14	1.22	1.70	1.84	1.98	5.41	1.28	34.43	9.21	39.41	7.47	222.02		
YAL043C	Unknown	0.90	1.14	1.35	1.69	1.84	2.04	3.75	1.19	26.55	1.74	30.30	2.83	200.02		
YAL044C	GCV3	0.88	1.07	1.22	1.54	1.68	1.90	6.39	1.27	39.46	11.29	47.85	12.56	297.21		
YAL044W_A	YAL044w_a	0.89	1.13	1.31	1.73	2.00	2.17	17.99	4.82	69.27	33.77	87.26	38.59	327.23		
YAL045C	YAL045c	0.94	1.17	1.36	1.74	1.88	2.07	6.70	1.38	36.39	7.66	43.09	8.38	279.26		
YAL046C	AIM1	0.88	1.09	1.21	1.47	1.60	1.82	5.67	1.32	35.56	9.21	41.22	10.53	243.42		
YAL047C	SPC72	0.91	1.17	1.17	1.16	1.31	1.45	65.07	52.24	112.69	105.74	177.76	157.98	331.35		
YAL048C	GEM1	0.85	0.98	1.00	0.86	0.85	0.82	126.38	127.94	124.25	141.24	250.63	269.18	510.03		
YAL048C	GEM1	0.92	1.11	1.11	1.61	1.74	1.82	57.17	36.83	105.89	87.43	163.08	124.89	339.27		
YAL049C	AIM2	0.94	1.23	1.43	1.82	1.98	2.19	1.79	1.28	13.10	-0.32	14.89	0.96	262.16		
YAL049C	AIM2	1.00	1.39	1.52	1.97	2.10	2.17	12.08	0.76	59.94	6.79	72.02	7.55	303.81		
YAL051W	OAF1	0.91	1.17	1.33	1.72	1.89	2.12	1.44	1.45	8.55	-1.06	9.99	0.39	224.42		
YAL053W	FLC2	0.93	1.22	1.37	1.78	1.81	1.95	1.54	1.20	5.62	-1.35	7.16	-0.15	244.72		
YAL054C	ACS1	0.95	1.30	1.48	1.85	2.05	2.17	17.35	7.83	59.10	39.29	76.45	47.12	341.08		
YAL055W	PEX22	0.97	1.18	1.42	1.81	1.97	2.18	19.06	3.29	59.79	28.55	78.84	31.84	248.31		
YAL056W	GPB2	0.89	1.14	1.36	1.72	1.90	2.13	1.39	0.68	13.27	0.53	14.66	1.22	7.67		
YAL058C_A	Unknown	0.97	1.41	1.60	1.96	2.15	2.28	10.31	6.02	44.50	30.65	54.81	36.67	307.85		
YAL059W	CNE1	0.89	1.13	1.35	1.69	1.84	2.05	10.66	1.52	43.11	11.99	53.77	13.50	237.51		
YAL059W	ECM1	0.90	1.12	1.32	1.71	1.87	2.08	10.26	1.68	46.10	7.47	56.36	8.55	235.12		
YAL060W	BHD1	0.89	1.15	1.38	1.73	1.88	2.08	19.22	4.17	66.02	30.59	85.24	34.76	298.12		
YAL061W	BHD2	0.94	1.25	1.49	1.88	2.02	2.21	2.68	1.22	21.40	0.69	24.09	1.91	160.85		
YAL062W	GDH3	0.95	1.18	1.38	1.77	1.92	2.11	10.18	1.87	47.13	14.45	57.31	16.33	194.70		
YAL064C_A	YAL064c_a	0.98	1.46	1.67	1.93	2.05	2.29	3.77	3.59	40.51	38.80	44.28	42.39	229.37		
YAL065C	YAL065c	0.90	1.14	1.32	1.66	1.83	2.03	10.9	1.19	6.11	-1.19	7.19	0.00	58.78		
YAL066W	YAL066w	0.92	1.23	1.46	1.82	1.97	2.17	7.28	1.23	36.43	6.82	43.71	7.51	239.22		
YAL067C	SEOT	0.90	1.17	1.36	1.71	1.85	2.04	5.81	1.78	36.35	1.26	38.93	1.16	218.17		
YAL067W_A	YAL067w_a	0.99	1.34	1.51	1.91	2.04	2.18	12.54	5.44	63.57	37.28	76.11	42.72	315.35		
YAL068C	PAU8	0.94	1.22	1.41	1.75	1.93	2.14	6.28	1.54	45.15	15.37	50.45	16.27	288.38		
YAL069C_A	ERH1	0.95	1.08	1.28	1.72	1.90	2.10	3.60	1.01	43.85	7.35	50.00	8.61	177.71		
YAL070W	YAL070w	0.92	1.15	1.34	1.67	1.87	2.01	11.35	1.68	43.86	7.35	50.00	8.61	251.66		
YAL071C	SWH1	0.92	1.22	1.41	1.74	1.87	2.08	11.69	1.52	49.07	11.89	60.16	13.41	251.66		
YAL071C	BUD14	0.93	1.19	1.37	1.77	1.97	2.24	1.81	1.09	5.35	-1.50	7.16	-0.41	342.79		
YAL071W	ADE1	0.93	1.17	1.21	1.60	1.78	1.97	10.19	1.27	107.19	122.68	118.22	229.88	220.13	495.70	
YAL071C	KH2	0.98	1.2													

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YBL042C	FU11	0.96	1.32	1.51	1.92	2.07	2.21	12.49	7.29	51.23	40.15	63.72	47.44	339.41		
YBL043W	ECM13	0.94	1.34	1.50	2.22	2.35	2.40	10.95	6.06	49.93	34.75	60.88	40.81	327.60		
YBL044W		0.95	1.13	1.11	1.16	1.26	1.43	78.69	59.18	114.07	104.90	192.76	164.08	458.53		
YBL044W	YBL044w	0.91	1.31	1.25	1.00	1.02	0.97	136.13	126.45	147.88	135.38	284.01	261.83	527.07		
YBL044W	YBL044w	0.87	1.04	1.01	0.81	0.87	0.82	124.70	132.69	130.49	128.14	235.19	260.83	521.15		
YBL045C	CORT	0.89	1.27	1.30	1.54	1.79	1.94	9.04	3.28	58.69	34.36	67.74	37.85	260.00		
YBL045C	CORT	0.89	1.08	1.03	0.68	0.91	0.63	126.55	124.42	129.16	121.12	235.70	245.54	524.53		
YBL045W	PSV4	0.88	1.14	1.26	1.60	1.76	1.80	9.09	3.22	46.94	8.11	56.02	11.39	318.96		
YBL045W	PSV4	0.88	1.18	1.20	1.55	1.61	1.87	9.04	3.23	46.94	8.17	56.02	11.39	318.96		
YBL049W	YBL049w	1.00	1.36	1.57	1.95	2.12	2.23	5.98	3.22	36.40	12.40	42.29	15.62	302.23		
YBL049W	MCH1	0.92	1.23	1.40	1.76	1.92	2.04	3.98	3.09	24.17	7.95	28.15	11.84	232.35		
YBL051C	PIN4	0.89	1.06	1.02	1.25	1.45	1.67	12.50	6.87	58.38	43.90	70.88	50.77	331.39		
YBL052C	SAS3	0.92	1.27	1.41	1.82	1.98	2.11	4.96	3.47	30.72	17.24	35.57	20.72	300.38		
YBL053W	YBL053w	0.86	1.17	1.26	1.64	1.81	1.97	12.50	6.87	50.95	32.48	63.45	37.95	337.78		
YBL054W	TOD6	0.83	1.10	1.17	1.51	1.67	1.84	12.72	5.94	53.81	35.98	66.53	41.92	340.02		
YBL055C	YBL055c	0.92	1.23	1.37	1.77	1.94	2.10	10.72	5.47	50.18	34.05	60.90	39.52	323.62		
YBL056W	PTC1	0.88	1.14	1.22	1.58	1.71	1.88	11.83	5.30	53.29	31.36	65.12	36.66	323.97		
YBL057C	PTH2	0.86	1.12	1.20	1.53	1.70	1.85	8.26	5.33	43.99	32.80	52.25	38.12	320.34		
YBL058W	SHP1	0.83	1.09	1.07	1.00	1.12	1.35	61.95	36.08	107.64	82.09	169.59	118.17	347.39		
YBL059W	YBL059w	0.88	1.22	1.40	1.78	1.94	2.08	19.35	3.23	67.46	6.71	86.81	9.94	312.28		
YBL060W	YEL1	0.83	1.05	1.16	1.51	1.70	1.86	8.16	3.24	41.74	10.15	49.90	13.39	304.43		
YBL061C	SKT5	0.95	1.21	1.36	1.77	1.95	2.11	3.75	3.17	22.26	8.93	26.01	12.10	279.96		
YBL062W	YBL062w	0.91	1.17	1.29	1.68	1.87	2.02	4.83	3.25	31.52	13.43	36.35	16.68	287.33		
YBL063W	KIP1	0.89	1.04	0.98	1.01	1.11	1.16	30.94	28.76	75.90	74.61	106.84	103.37	335.37		
YBL064C	PRX1	0.93	1.20	1.26	1.69	1.68	2.04	22.11	8.46	67.38	45.36	89.50	53.82	336.93		
YBL065W	YBL065w	0.89	1.20	1.28	1.62	1.79	1.97	11.32	6.01	50.43	37.31	61.76	43.32	324.97		
YBL066C	SEF1	0.88	1.17	1.26	1.63	1.82	2.00	12.78	3.94	53.62	24.71	66.40	28.65	327.93		
YBL067C	UBP13	0.90	1.15	1.26	1.63	1.78	1.94	18.88	9.41	61.62	43.85	80.60	53.26	343.42		
YBL068W	PRS4	0.89	1.15	1.22	1.56	1.74	1.93	9.18	7.64	49.27	41.36	58.45	49.00	314.05		
YBL069W	AST1	0.92	1.24	1.34	1.72	1.89	2.06	6.71	3.63	39.73	19.87	46.44	23.50	303.34		
YBL070C	YBL070c	0.85	1.08	1.14	1.43	1.61	1.79	6.49	4.53	37.83	26.15	44.32	30.68	308.90		
YBL071C	YBL071c	0.90	1.20	1.41	1.83	1.99	2.13	6.99	4.57	41.08	18.14	48.08	21.66	301.45		
YBL071C-B	YBL071c-b	0.90	1.18	1.29	1.70	1.92	2.11	13.95	6.84	67.95	38.96	81.89	45.79	317.64		
YBL071W-A	K1111	0.89	1.11	1.16	1.48	1.73	1.93	24.10	16.99	55.70	103.92	72.70	309.22			
YBL072C	RPS6a	0.90	1.32	1.43	1.77	2.05	2.22	2.25	1.41	30.07	11.00	32.32	12.41	281.07		
YBL072C	RPS6a	0.83	1.16	1.21	1.63	1.82	2.00	12.78	3.94	53.62	24.71	66.40	28.65	327.93		
YBL073C	SSA3	0.92	1.19	1.35	1.44	1.91	2.08	4.69	3.17	29.54	9.66	34.22	12.17	289.10		
YBL078C	ATG6	0.92	1.15	1.24	1.67	1.84	2.03	13.06	7.48	55.17	48.21	50.27	28.65	326.44		
YBL079W	NUP170	0.84	1.10	1.08	1.39	1.57	1.83	18.44	20.75	65.91	69.93	84.45	86.38	331.44		
YBL079W	NUP170	0.89	1.24	1.29	1.56	1.84	2.08	19.69	7.22	75.49	52.90	95.00	69.13	309.78		
YBL080C	YBL080c	0.94	1.32	1.48	1.91	2.06	2.16	8.35	6.07	46.58	37.93	54.84	44.00	293.80		
YBL080C	PET111	0.89	1.23	1.28	1.93	1.91	2.08	13.29	12.53	143.88	142.73	269.17	216.31	534.44		
YBL080C	PET111	0.86	1.05	1.01	0.86	0.89	0.81	125.21	123.78	126.77	120.07	251.99	242.85	539.43		
YBL081W	YBL081w	0.94	1.29	1.42	1.79	1.96	2.11	7.98	4.53	44.67	31.85	52.55	36.78	309.92		
YBL082C	ALG3	0.94	1.24	1.33	1.60	1.74	1.90	11.65	4.60	46.13	28.61	57.78	33.21	319.25		
YBL083C	YBL083c	0.94	1.20	1.27	1.51	1.64	1.80	8.40	5.82	43.70	32.48	52.10	38.30	308.58		
YBL085W	BOH1	0.97	1.33	1.47	1.81	1.98	2.12	9.76	6.78	48.45	39.15	58.21	45.93	321.78		
YBL086C	YBL086c	0.96	1.32	1.50	1.89	2.05	2.18	9.94	5.75	48.17	34.07	58.11	39.82	319.28		
YBL087C	RPL23a	0.89	1.12	1.19	1.52	1.71	1.91	5.15	3.28	32.61	13.25	37.76	16.52	288.31		
YBL088C	TEL1	0.96	1.28	1.44	1.79	1.95	2.10	8.09	4.86	43.92	31.09	52.02	35.96	305.13		
YBL089W	AVT5	0.87	1.14	1.28	1.62	1.77	1.93	5.17	3.18	33.23	8.96	38.40	12.13	290.05		
YBL090W	MRP21	0.89	1.19	1.16	1.01	1.00	1.01	129.97	127.87	147.05	140.58	277.02	268.45	531.40		
YBL090W	MRP21	0.86	1.08	1.03	0.85	0.88	0.83	121.69	121.22	126.62	121.45	248.31	242.67	539.93		
YBL091C	MAP2	0.96	1.29	1.45	1.83	1.99	2.14	12.57	6.63	53.94	38.50	66.51	45.13	294.80		
YBL091C-A	SCS22	0.95	1.40	1.58	1.99	2.16	2.24	3.59	1.49	39.92	22.14	43.51	23.63	250.91		
YBL091C-A	SCS22	0.93	1.25	1.41	1.86	2.03	2.15	10.06	1.76	61.29	21.75	71.35	28.91	198.19		
YBL093C	ROX3	0.92	1.16	1.26	1.56	1.79	1.99	1.22	0.85	8.73	0.76	9.96	1.61	254.69		
YBL094C	YBL094c	0.90	1.22	1.33	1.64	1.84	2.02	1.13	0.81	8.63	0.73	8.64	1.11	207.00		
YBL094C	SCD2	0.93	1.26	1.45	1.86	2.04	2.14	18.97	6.95	45.74	27.56	65.51	24.49	307.00		
YBL095C	FLR1	0.89	1.16	1.26	1.60	1.78	1.97	8.21	1.59	49.67	16.28	57.88	17.78	298.36		
YBL096C	HHF1	0.84	1.08	1.14	1.41	1.69	1.91	4.85	1.66	36.86	7.22	41.71	14.38	290.17		
YBL097C	HHT1	0.92	1.33	1.48	1.88	2.08	2.21	15.07	8.88	62.58	47.98	77.66	56.85	316.43		
YBR001C	NTH2	0.94	1.34	1.59	2.00	2.17	2.29	25.81	12.14	83.65	62.29	108.66	74.43	357.43		
YBR003W	COO1	0.89	1.19	1.38	1.79	1.98	2.10	14.15	1.38	67.35	24.75	81.50	26.12	324.11		
YBR005W	RCR1	1.00	1.26	1.46	1.65	1.88	2.13	13.06	7.14	65.73	48.50	78.79	55.63	331.81		
YBR006W	UGA2	0.86	1.18	1.31	1.67	1.87	2.05	9.62	4.92	54.32	36.90	63.94	41.81	302.44		
YBR007C	DSF2	0.93	1.30	1.48	1.86	2.04	2.14	14.97	6.55	67.54	47.56	86.51	54.52	134.82		
YBR025C	OLA1	0.87	1.13	1.25	1.65	1.87	2.06	19.22	4.39	65.82	39.07	85.04	43.46	127.56		
YBR026C	ETR1	0														

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE			FRUCTOSE			TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YBR072c-A	YBR072c-a	0.94	1.25	1.41	2.01	2.00	2.11	14.58	8.23	65.98	39.76	80.57	47.99	325.47	325.47	325.47	325.47	
YBR072W	HSP26	0.86	1.14	1.30	1.57	1.77	1.92	5.81	2.63	44.28	29.66	50.09	32.29	247.24	247.24	247.24	247.24	
YBR073W	RDH54	0.93	1.29	1.50	1.85	2.01	2.15	10.55	2.53	57.21	31.74	67.76	34.27	283.50	283.50	283.50	283.50	
YBR074W	YBR074w	0.85	1.19	1.39	1.78	1.97	2.09	10.34	2.36	56.00	30.17	66.34	32.53	274.49	274.49	274.49	274.49	
YBR074W	YBR074w	0.93	1.27	1.45	1.82	1.98	2.13	8.29	2.00	51.51	20.25	59.80	22.25	298.42	298.42	298.42	298.42	
YBR075W	YBR075w	0.92	1.26	1.41	1.75	1.94	2.08	6.92	2.10	48.27	27.48	55.19	29.57	296.99	296.99	296.99	296.99	
YBR076W	ECM20	0.86	1.25	1.44	1.78	2.07	2.09	10.16	2.68	54.07	27.59	64.63	29.59	306.16	306.16	306.16	306.16	
YBR076W	ECM4	0.86	1.26	1.43	1.85	2.03	2.09	9.46	2.57	53.93	27.59	64.63	29.59	306.16	306.16	306.16	306.16	
YBR076W	ECM23	0.86	1.16	1.38	1.80	2.01	2.18	1.11	0.74	9.19	1.62	10.30	2.36	239.21	239.21	239.21	239.21	
YBR081C	SP77	0.94	1.01	1.04	1.07	1.24	1.34	75.34	61.81	113.12	104.40	188.46	161.21	422.00	422.00	422.00	422.00	
YBR081C	SP77	0.87	1.05	1.06	1.02	1.16	1.24	58.80	43.03	112.52	94.88	171.73	137.92	400.38	400.38	400.38	400.38	
YBR082C	UBC4	0.92	1.05	1.16	1.46	1.73	1.94	25.69	9.83	79.87	57.50	105.56	67.33	308.36	308.36	308.36	308.36	
YBR083W	TEC1	0.90	1.12	1.32	1.75	1.93	2.13	0.77	0.66	9.49	2.32	10.26	2.98	280.48	280.48	280.48	280.48	
YBR084C-A	RPL19a	0.93	1.10	1.26	1.76	2.03	2.27	1.42	0.63	25.51	6.54	26.93	7.17	211.82	211.82	211.82	211.82	
YBR084W	MIS1	0.93	1.20	1.43	1.88	2.05	2.19	1.15	0.66	20.97	5.17	22.12	5.83	281.70	281.70	281.70	281.70	
YBR085C-A	YBR085c-a	0.94	1.24	1.40	1.75	1.96	2.07	2.07	0.85	24.42	7.17	26.49	8.02	314.59	314.59	314.59	314.59	
YBR085W	AAC3	0.92	1.10	1.35	2.01	2.01	2.05	4.64	0.85	38.97	12.18	43.61	13.03	289.10	289.10	289.10	289.10	
YBR090C	YBR090c	0.94	1.17	1.38	1.92	2.11	2.26	1.93	0.65	28.45	8.37	30.38	9.03	294.44	294.44	294.44	294.44	
YBR090C-A	YBR090c-A	0.97	1.21	1.45	1.89	2.08	2.22	2.93	0.62	35.35	4.96	38.28	5.58	300.77	300.77	300.77	300.77	
YBR092C	PHO3	0.96	1.22	1.49	1.94	2.11	2.25	3.90	0.64	39.69	7.73	43.59	8.36	304.21	304.21	304.21	304.21	
YBR093C	PHO5	0.92	1.10	1.31	1.82	2.02	2.18	10.57	1.81	56.76	26.39	67.33	28.20	313.54	313.54	313.54	313.54	
YBR094W	PBY1	0.86	0.99	1.16	1.57	1.77	1.94	8.26	1.37	53.64	24.60	61.91	25.97	327.40	327.40	327.40	327.40	
YBR095C	RXT2	0.91	1.22	1.47	1.84	2.00	2.12	7.84	1.86	51.11	35.83	64.96	37.68	301.19	301.19	301.19	301.19	
YBR097W	VPS15	0.93	1.04	1.09	1.13	1.32	1.47	43.48	38.41	118.82	84.91	162.30	123.31	375.18	375.18	375.18	375.18	
YBR098W	MM54	0.97	1.20	1.42	1.86	2.06	2.20	6.25	1.90	47.03	28.69	53.28	30.58	316.61	316.61	316.61	316.61	
YBR098W	MM54	0.89	1.24	1.37	1.77	1.95	2.11	13.89	2.26	66.13	23.79	80.02	26.06	318.83	318.83	318.83	318.83	
YBR099C	YBR099c	0.97	1.24	1.49	1.92	2.12	2.26	1.34	0.78	25.09	8.61	26.43	9.39	291.31	291.31	291.31	291.31	
YBR100W	YBR100W	0.96	1.25	1.51	1.95	2.16	2.31	0.74	0.73	9.69	2.11	10.43	2.84	293.95	293.95	293.95	293.95	
YBR100W	YBR100W	0.90	1.04	1.21	1.71	1.93	2.14	1.39	0.77	25.37	7.70	26.76	12.47	301.30	301.30	301.30	301.30	
YBR101C	FEST	0.85	1.08	1.08	1.14	1.32	1.51	47.34	34.85	94.28	83.27	141.62	118.12	339.36	339.36	339.36	339.36	
YBR103W	SIF2	1.01	1.20	1.38	1.81	2.04	2.22	12.09	2.47	56.51	29.72	68.59	32.19	300.63	300.63	300.63	300.63	
YBR104W	YMC2	0.88	1.08	1.27	1.66	1.91	2.09	3.48	0.79	37.59	9.69	41.07	9.88	307.04	307.04	307.04	307.04	
YBR105C	VID24	0.94	1.16	1.38	1.80	1.96	2.10	4.77	1.27	42.60	21.61	47.37	22.88	303.48	303.48	303.48	303.48	
YBR106W	PHOB8	0.97	1.23	1.48	1.89	2.08	2.16	32.47	23.52	80.26	71.83	112.73	95.84	378.03	378.03	378.03	378.03	
YBR107W	IM3	1.03	1.23	1.43	1.84	2.13	2.27	17.14	1.70	70.18	37.40	87.32	30.77	325.13	325.13	325.13	325.13	
YBR108W	ANL2	0.86	1.18	1.34	1.79	2.03	2.17	3.32	1.14	37.15	9.81	40.47	10.05	297.44	297.44	297.44	297.44	
YBR111C	VS41	0.88	1.26	1.48	1.97	2.16	2.28	1.35	0.68	22.03	4.92	23.58	5.68	284.49	284.49	284.49	284.49	
YBR111W-A	SUS1	0.85	1.20	1.31	1.64	1.89	2.05	21.86	8.02	78.08	40.57	89.94	48.59	329.17	329.17	329.17	329.17	
YBR112C	VCY9	0.88	1.10	1.05	0.96	1.03	1.13	65.57	46.73	120.64	84.23	186.21	136.96	404.70	404.70	404.70	404.70	
YBR113W	YBR113c	0.98	1.28	1.54	1.96	2.13	2.24	3.06	0.85	38.14	13.28	38.20	14.15	298.62	298.62	298.62	298.62	
YBR114W	RAD16	0.88	1.33	1.60	2.05	2.23	2.36	0.75	0.71	3.08	1.50	3.83	2.21	265.39	265.39	265.39	265.39	
YBR115C	LYS2	0.94	1.13	1.34	1.77	2.00	2.14	7.55	2.33	48.40	29.59	55.95	31.93	319.14	319.14	319.14	319.14	
YBR116C	YBR116c	0.82	1.31	1.57	1.97	2.17	2.29	5.53	1.46	46.55	25.19	52.08	26.65	304.45	304.45	304.45	304.45	
YBR119W	MUD1	1.00	1.25	1.50	1.93	2.11	2.24	2.41	0.95	32.48	8.80	34.89	9.76	293.74	293.74	293.74	293.74	
YBR120C	CBP6	0.92	1.07	1.24	1.55	1.78	1.99	0.74	0.77	3.62	1.17	4.36	1.94	284.42	284.42	284.42	284.42	
YBR121C	GRS1	0.91	1.11	1.30	1.76	1.97	2.13	12.41	6.13	59.51	45.45	71.92	51.57	305.73	305.73	305.73	305.73	
YBR122C	MRP136	0.93	1.14	1.22	1.91	2.01	2.05	11.19	1.16	52.76	25.67	64.16	25.67	306.22	306.22	306.22	306.22	
YBR123C	AGP2	0.90	1.07	1.11	1.45	1.77	1.93	19.40	3.85	60.53	9.51	91.81	7.47	283.13	283.13	283.13	283.13	
YBR133C	HSL7	0.92	1.20	1.39	1.82	2.03	2.15	4.54	2.06	41.17	25.27	45.71	27.33	262.12	262.12	262.12	262.12	
YBR133C	HSL7	0.87	1.13	1.20	1.40	1.68	1.80	18.45	4.02	67.92	33.91	86.37	37.93	315.04	315.04	315.04	315.04	
YBR134W	YBR134w	0.91	1.14	1.32	1.77	1.99	2.13	8.70	5.04	58.38	42.10	67.08	47.14	292.59	292.59	292.59	292.59	
YBR137W	YBR137w	0.89	1.16	1.35	1.80	1.97	2.14	6.35	1.31	47.56	20.64	53.91	7.85	296.15	296.15	296.15	296.15	
YBR138C	YBR138c	0.88	1.05	1.21	1.65	1.84	2.04	5.34	1.32	49.03	34.63	54.37	37.95	296.32	296.32	296.32	296.32	
YBR161W	CSH1	0.90	1.11	1.30	1.79	1.98	2.07	1.94	2.05	2.05	29.63	28.22	31.58	30.27	282.36	282.36	282.36	282.36
YBR162C	TOS1	0.91	1.20	1.36	1.63	1.85	2.05	10.12	5.20	59.48	43.83	69.60	49.02	312.74	312.74	312.74	312.74	
YBR162W-A	YSY6	0.89	1.05	1.24	1.67	1.90	2.09	3.39	1.09	37.17	15.88	40.56	16.97	295.97	295.97	295.97	295.97	
YBR162W-A	YSY6	0.90	1.18	1.30	1.70	1.89	1.98	7.57	1.50	49.63	24.19	57.20	26.09	279.90	279.90	279.90	279.90	
YBR163W	EX05	0.92	1.10															

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE			FRUCTOSE			TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YBR217W	ATG12	0.98	1.24	1.45	1.95	2.12	2.23	27.27	11.64	86.74	63.49	114.01	75.12	319.86	319.86	319.86	319.86	
YBR218C	PVC2	0.92	1.19	1.31	1.75	1.97	2.15	6.89	1.59	54.92	29.35	61.81	30.95	311.62	311.62	311.62	311.62	
YBR219C	YBR219c	0.95	1.19	1.33	1.86	2.09	2.26	6.47	0.67	48.28	21.08	54.75	21.75	310.35	310.35	310.35	310.35	
YBR220C	YBR220c	0.91	1.15	1.31	1.82	2.04	2.22	0.59	0.11	20.43	5.66	21.02	5.77	293.33	293.33	293.33	293.33	
YBR221W	PDB1	0.92	1.12	1.29	1.83	2.00	2.18	0.26	0.04	11.36	2.16	11.62	2.20	287.30	287.30	287.30	287.30	
YBR221W-A	YBR221w-a	0.90	1.37	1.56	1.95	2.13	2.24	9.22	2.42	53.66	22.57	62.88	24.98	311.60	311.60	311.60	311.60	
YBR222C	PCP20	0.98	1.03	1.10	1.26	1.35	2.03	0.36	0.04	22.24	3.48	22.60	3.52	302.39	302.39	302.39	302.39	
YBR223C	TDP1	0.87	1.08	1.19	1.75	1.88	2.11	4.10	1.68	47.09	16.96	51.19	18.04	311.35	311.35	311.35	311.35	
YBR224W	YBR224w	0.98	1.14	1.24	1.83	2.03	2.29	9.20	2.45	53.66	22.57	62.88	24.98	311.60	311.60	311.60	311.60	
YBR225W	YBR225w	0.98	1.23	1.44	2.11	2.13	2.22	19.71	15.70	71.61	68.82	91.31	83.75	337.65	337.65	337.65	337.65	
YBR226C	YBR226c	0.94	1.19	1.33	1.78	1.99	2.14	0.05	-0.02	3.80	0.85	3.85	0.84	286.78	286.78	286.78	286.78	
YBR227C	MCX1	0.93	1.23	1.40	1.87	2.07	2.21	0.34	0.05	16.56	4.11	16.90	4.16	280.42	280.42	280.42	280.42	
YBR228W	SLX1	0.91	1.16	1.36	1.86	2.05	2.17	1.88	0.33	30.25	11.71	32.13	12.05	280.44	280.44	280.44	280.44	
YBR229C	ROT2	0.93	1.15	1.33	1.83	2.01	2.15	8.56	2.78	49.02	19.13	57.58	19.91	311.62	311.62	311.62	311.62	
YBR230C	OM14	0.92	1.19	1.38	1.91	2.08	2.23	4.13	0.36	44.76	15.29	48.89	15.64	298.80	298.80	298.80	298.80	
YBR230W-A	YBR230w-a	0.90	1.29	1.45	1.79	1.97	2.14	3.46	0.87	39.49	9.63	42.96	10.50	279.89	279.89	279.89	279.89	
YBR231C	SWC3	1.03	1.21	1.21	1.51	1.85	2.08	63.42	54.28	108.57	108.04	171.93	162.31	399.58	399.58	399.58	399.58	
YBR232C	YBR232c	0.93	1.27	1.48	1.88	2.04	2.16	14.29	3.14	54.14	15.48	68.43	18.61	331.96	331.96	331.96	331.96	
YBR233W	PBP2	0.91	1.13	1.31	1.86	2.05	2.19	8.86	0.74	52.79	21.64	61.65	22.38	315.23	315.23	315.23	315.23	
YBR235W	YBR235w	0.92	1.17	1.32	1.90	2.07	2.22	4.53	0.13	48.47	13.19	53.01	13.31	293.43	293.43	293.43	293.43	
YBR238C	YBR238c	0.93	1.17	1.37	1.86	2.06	2.18	19.77	9.91	71.33	60.73	91.10	70.65	341.77	341.77	341.77	341.77	
YBR239C	ERT1	0.90	1.11	1.28	1.83	2.05	2.18	0.66	0.16	21.19	6.34	21.84	6.51	293.88	293.88	293.88	293.88	
YBR240C	THI2	0.95	1.16	1.30	1.80	2.02	2.16	8.05	1.42	56.87	28.21	64.92	29.63	312.00	312.00	312.00	312.00	
YBR241C	YBR241c	1.01	1.25	1.44	1.98	2.21	2.36	0.29	0.14	16.43	2.63	16.72	2.77	300.90	300.90	300.90	300.90	
YBR242W	YBR242w	0.88	1.10	1.24	1.77	1.99	2.14	2.29	0.07	5.78	0.26	6.06	0.33	292.48	292.48	292.48	292.48	
YBR244W	GPK2	0.98	1.26	1.48	1.98	2.15	2.26	8.18	1.00	56.09	25.13	64.27	26.12	314.21	314.21	314.21	314.21	
YBR245C	ISW1	0.92	1.10	1.26	1.76	2.01	2.21	0.02	0.01	4.11	-0.03	4.12	-0.03	279.32	279.32	279.32	279.32	
YBR246W	RRT2	0.94	1.11	1.21	1.67	1.89	2.11	4.67	0.47	50.45	17.58	55.12	18.06	309.04	309.04	309.04	309.04	
YBR248C	HIS7	0.89	1.10	1.09	0.92	0.91	126.49	122.92	125.01	122.94	251.50	245.86	540.08	540.08	540.08	540.08		
YBR249C	ARO4	0.99	1.13	1.17	1.62	1.93	2.13	30.66	15.35	84.93	66.27	115.59	81.62	353.06	353.06	353.06	353.06	
YBR250W	SPD2	0.92	1.10	1.25	1.76	1.97	2.11	7.76	0.63	52.50	21.48	60.27	22.10	306.67	306.67	306.67	306.67	
YBR251W	MGPSS	0.94	0.91	0.89	0.92	0.94	0.93	118.94	130.46	125.24	136.93	244.17	267.39	533.36	533.36	533.36	533.36	
YBR255W	MT4	0.92	1.15	1.35	1.88	2.07	2.17	5.42	2.33	43.67	28.94	49.09	31.27	324.19	324.19	324.19	324.19	
YBR258C	SHG1	0.94	1.16	1.34	1.87	2.01	2.23	0.06	0.06	8.52	1.17	8.58	1.23	294.22	294.22	294.22	294.22	
YBR259W	YBR259w	0.94	1.17	1.38	1.89	2.07	2.19	11.00	3.82	59.39	40.34	70.39	44.15	316.37	316.37	316.37	316.37	
YBR260C	RGD1	0.92	1.15	1.27	1.83	2.03	2.21	10.99	3.56	54.62	42.31	65.80	45.87	319.24	319.24	319.24	319.24	
YBR261C	TAE1	0.91	1.12	1.27	1.81	2.02	2.26	3.06	0.24	43.90	17.83	46.95	18.05	307.39	307.39	307.39	307.39	
YBR262C	ANL4	0.93	1.13	1.31	1.83	2.03	2.26	0.95	0.24	63.75	9.85	64.70	10.00	304.78	304.78	304.78	304.78	
YBR263C	YBR263c	0.93	1.13	1.30	1.82	2.01	2.21	11.50	1.13	52.26	25.17	69.54	26.27	315.35	315.35	315.35	315.35	
YBR264C	YPT10	0.94	1.14	1.33	1.87	2.07	2.23	1.95	0.07	34.19	8.06	36.14	8.13	222.22	222.22	222.22	222.22	
YBR266C	YBR266c	0.95	1.07	1.10	1.59	1.87	2.09	40.58	26.43	160.19	83.70	140.77	110.13	375.58	375.58	375.58	375.58	
YBR267W	REH1	0.99	1.07	1.08	1.23	1.61	1.81	13.09	2.74	68.72	10.99	82.63	12.73	255.44	255.44	255.44	255.44	
YBR268W	MRLP137	0.93	0.98	0.97	0.99	1.01	1.21	121.49	144.26	124.45	136.71	245.92	280.97	536.15	536.15	536.15	536.15	
YBR269C	FMP21	0.92	1.31	1.45	1.84	2.02	2.21	6.15	0.15	48.57	7.54	54.71	8.49	271.62	271.62	271.62	271.62	
YBR270C	BIT2	0.90	1.06	1.22	1.62	1.81	2.00	2.88	0.66	33.75	8.12	36.64	8.78	290.97	290.97	290.97	290.97	
YBR271W	YBR271w	0.93	1.41	1.58	1.96	2.12	2.18	23.69	1.15	75.48	16.42	99.09	17.56	282.29	282.29	282.29	282.29	
YBR272C	HSM3	0.92	1.02	1.14	1.51	1.72	1.92	11.50	5.34	58.30	20.33	69.80	20.62	310.73	310.73	310.73	310.73	
YBR273C	UBX7	0.89	1.33	1.51	1.90	2.06	2.12	18.42	1.09	77.36	14.58	95.78	15.67	223.23	223.23	223.23	223.23	
YBR274W	CHK1	0.95	1.37	1.52	1.90	2.09	2.21	12.57	1.11	56.49	15.83	63.66	16.20	268.85	268.85	268.85	268.85	
YBR275C	RIF1	0.92	1.02	1.13	1.62	1.83	2.00	4.36	1.10	34.98	12.06	39.34	13.15	288.52	288.52	288.52	288.52	
YBR276C	PPS1	0.95	1.12	1.30	1.78	1.98	2.16	5.94	0.85	44.28	14.00	50.22	14.85	294.72	294.72	294.72	294.72	
YBR278W	DPB3	0.91	1.25	1.36	1.77	1.96	2.08	21.76	6.66	73.47	48.50	95.23	55.15	251.80	251.80	251.80	251.80	
YBR279W	PAF1	0.91	1.06	1.24	1.61	1.83	2.01	2.72	1.02	33.19	12.49	35.91	13.51	264.83	264.83	264.83	264.83	
YBR280C	SAF1	0.97	1.10	1.26	1.76	1.98	2.15	16.06	2.64	65.20	21.50	76.09	21.62	336.42	336.42	336.42	336.42	
YBR281C	DUG2	0.86	1.22	1.38	1.80	1.99	2.08	13.85	2.60	63.57	20.87	74.47	21.72	200.03	200.03	200.03	200.03	
YBR282W	MBP27	0.90	1.13	1.36	1.75	1.96	2.19	15.48	2.70	72.94	8.20	88.43	8.91	307.45	307.45	307.45	307.45	
YBR292C	VBL4	0.90	1.35	1.55	1.92	2.11	2.29	15.23	1.67	68.59	15.98	82.23	17.05	226.46	226.46	226.46	226.46	
YBR293W	YCL041	0.98	1.23	1														

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)		
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h		
YCL057C-A	YCL057c-a	0.91	1.18	1.34	1.74	1.97	2.08	11.48	3.95	59.05	27.79	70.53	31.73	306.32			
YCL057W	PRD1	0.94	1.24	1.42	1.93	2.10	2.20	11.55	1.21	62.99	12.59	74.54	13.80	320.68			
YCL058C	FVVS	0.96	1.22	1.24	1.34	1.63	1.94	35.69	18.79	76.43	61.83	112.12	80.62	348.85			
YCL060C	YCL060c	0.97	1.17	1.22	1.68	1.96	2.17	17.77	2.60	60.99	33.86	98.77	36.46	335.23			
YCL061C	MRC1	0.98	1.28	1.41	1.84	2.07	2.27	0.85	0.79	3.81	0.76	4.67	1.54	258.38			
YCL062W	YCL062w	0.91	1.10	1.12	1.52	1.80	1.99	16.22	9.15	69.40	60.11	85.62	69.26	312.34			
YCL063W	VAC07	0.93	1.20	1.31	1.77	1.97	2.12	1.41	1.48	20.66	18.15	22.08	19.54	307.12			
YCL064C	YAC14	0.94	1.14	1.19	1.56	1.76	1.99	3.03	2.63	32.63	24.44	35.96	22.71	270.84			
YCL065W	VBA3	0.93	1.13	1.35	1.54	2.05	2.22	1.34	1.49	1.60	18.28	20.36	19.76	21.96	270.84		
YCL074W	YCL074w	0.96	1.34	1.52	1.88	2.03	2.15	19.45	14.20	58.45	47.83	77.90	62.03	344.63			
YCL075W	YCL075w	0.95	1.34	1.54	1.89	2.04	2.15	20.07	8.49	59.89	43.11	79.96	52.61	345.35			
YCL076W	YCL076w	0.99	1.34	1.51	1.92	2.10	2.26	3.97	3.30	25.35	15.70	28.32	19.00	300.74			
YCR0001W	YCR001w	0.98	1.31	1.54	2.05	2.22	2.34	1.09	1.25	5.81	1.20	6.90	1.05	267.33			
YCR0002C	CDC10	0.94	1.11	1.22	1.72	1.95	2.05	21.94	20.98	71.79	71.97	93.72	92.95	290.05			
YCR0003W	MSP1_32	0.94	1.16	1.30	1.63	1.89	2.07	1.66	1.49	13.93	7.72	15.58	9.21	252.40			
YCR0004C	CP24	0.89	1.07	1.16	1.72	1.96	2.09	12.61	13.88	65.88	62.59	78.49	76.48	320.87			
YCR0005C	CIT2	0.98	1.28	1.44	1.89	2.11	2.22	1.99	1.12	29.19	3.47	31.18	4.59	305.66			
YCR0006C	YCR006c	0.94	1.21	1.37	1.81	2.02	2.11	7.56	5.24	60.25	6.66	313.90					
YCR0007C	YCR007c	1.02	1.35	1.53	1.98	2.16	2.23	9.15	1.20	62.29	10.00	71.44	11.21	308.10			
YCR0008W	SAT4	0.95	1.20	1.35	1.80	2.03	2.12	3.38	1.52	41.53	18.54	44.92	20.06	312.69			
YCR0009C	RVT161	0.93	1.13	1.23	1.61	1.84	1.93	1.26	1.17	3.75	1.97	5.02	3.14	263.26			
YCR010C	ADY2	1.00	1.30	1.51	2.00	2.18	2.28	1.28	1.80	27.43	20.00	29.81	21.80	288.32			
YCR011C	ADP1	0.91	1.18	1.37	1.88	2.08	2.18	2.46	3.12	31.88	31.82	34.34	34.94	287.81			
YCR014C	POL4	0.91	1.18	1.40	1.95	2.16	2.26	1.23	1.07	8.94	3.05	10.17	4.12	277.39			
YCR015C	YCR015c	0.94	1.16	1.33	1.79	2.00	2.11	8.11	3.44	55.29	37.05	63.40	40.49	300.61			
YCR016W	YCR016w	0.95	1.22	1.34	1.73	1.87	2.02	0.93	0.90	3.88	0.97	4.81	1.87	267.63			
YCR017C	CWH43	0.95	1.12	1.25	1.70	1.90	1.99	8.28	8.00	57.69	49.40	65.97	57.40	295.27			
YCR019W	MAK32	0.90	1.14	1.30	1.78	2.00	2.14	1.25	1.14	3.77	2.22	5.01	3.36	274.96			
YCR020C	PET18	1.04	1.38	1.58	2.06	2.22	2.31	11.60	1.37	59.47	13.16	71.07	14.53	321.78			
YCR020C-A	MAK31	1.01	1.46	1.69	2.17	2.32	2.42	1.18	1.11	13.41	0.32	14.59	1.43	290.53			
YCR020B-W	HTL1	0.93	1.31	1.48	1.79	1.94	2.03	11.84	2.72	62.52	37.70	74.36	40.42	285.33			
YCR021C	HSP30	0.99	1.38	1.58	2.02	2.17	2.25	1.06	1.68	9.72	1.99	10.78	3.07	286.77			
YCR022C	YCR022c	0.92	1.26	1.42	1.88	2.03	2.12	1.20	1.14	6.64	8.82	7.78		275.82			
YCR023C	YCR023c	1.00	1.42	1.68	2.12	2.27	2.33	1.97	1.11	25.16	10.99	27.13	12.10	280.88			
YCR024C	SLMS	0.89	1.14	1.17	1.43	1.66	1.84	6.43	1.56	52.33	23.28	58.76	24.84	248.94			
YCR024C-A	MPM1	0.90	1.31	1.47	1.81	2.03	2.15	16.73	1.15	70.50	16.50	87.23	17.85	323.36			
YCR025C	YCR025c	0.93	1.20	1.34	1.74	1.94	2.11	2.44	1.75	6.78	30.23	2.41	32.54	2.49	307.49		
YCR026C	MPBP1	0.81	1.14	1.40	1.69	1.86	1.93	1.27	1.63	1.75	25.32	14.42	26.95	2.17	228.76		
YCR027C	YCR027c	0.91	1.24	1.40	1.80	2.01	2.11	1.72	1.75	1.88	21.00	15.88	21.00	14.25	342.45		
YCR028C	RRB1	0.91	1.26	1.35	1.88	2.06	2.13	1.98	1.86	15.86	1.33	70.43	16.88	86.29	22.00	340.01	
YCR029C	FEN2	0.92	1.23	1.33	1.72	1.97	1.98	2.27	2.24	32.68	12.74	91.40	12.07	81.01	12.07	239.75	
YCR030C	RRB1	0.94	1.34	1.52	1.95	2.11	2.21	4.01	1.37	41.84	21.75	45.84	23.12	185.63			
YCR031C	PHOB7	0.92	1.32	1.54	1.88	2.05	2.19	1.34	1.20	23.07	12.42	29.05	12.00	239.75			
YCR032C	PHOB7	0.89	1.18	1.22	1.42	1.58	1.68	30.47	27.48	89.19	88.14	119.67	115.61	235.11			
YCR033C	YCR043c	0.93	1.32	1.42	1.77	1.94	2.06	5.41	4.49	51.99	42.91	57.40	47.40	291.56			
YCR044C	PER1	0.90	1.02	1.11	1.41	1.68	1.88	2.24	0.63	26.26	2.07	28.49	2.70	273.89			
YCR045C	RRT12	0.91	1.28	1.40	1.73	1.91	2.06	0.88	0.83	16.85	2.07	17.73	2.69	272.03			
YCR046C	IMG1	0.94	0.98	0.98	0.94	0.94	0.92	142.06	130.00	142.27	131.91	284.33	261.91	192.52			
YCR047C	YIH1	0.90	1.28	1.45	1.82	2.06	2.12	20.85	1.10	76.39	17.81	97.23	18.91	322.60			
YCR047C	TUD1	0.87	1.03	1.19	1.77	1.99	2.15	6.78	1.49	51.92	26.40	58.71	27.89	290.90			
YCR048W	TAH1	0.84	1.05	1.02	1.01	1.21	1.58	42.14	23.08	91.19	72.05	133.33	95.13	347.34			
YCR048W	ARE1	0.90	1.18	1.31	1.66	1.87	2.07	1.57	1.30	15.43	9.00	17.00	10.30	265.83			
YCR048W	ARE1	0.91	1.13	1.34	1.90	2.07	2.22	6.32	0.94	47.01	13.04	53.33	13.98	295.45			
YCR049C	YCR049c	0.89	1.36	1.47	1.81	1.98	2.06	10.46	0.84	61.06	12.11	71.53	12.95	314.80			
YCR050C	YCR050c	0.86	1.19	1.29	1.64	1.85	2.03	1.28	0.74	23.23	2.76	24.51	3.50	303.13			
YCR051W	YCR051w	0.90	1.27	1.39	1.76	1.93	2.06	11.76	0.88	63.86	11.70	75.62	12.58	312.88			
YCR053W	THR4	0.94	1.17	1.20	1.44	1.77	2.01	14.68	4.16	70.08	42.82	84.77	46.98	284.86			
YCR059C	YIH1	0.90	1.28	1.45	1.82	2.06	2.12	20.85	1.10	76.39	17.81	97.23	18.91	322.60			
YCR060W	TAH1	0.87	1.03	1.19	1.77	1.99	2.15	6.78	1.49	51.92	26.40	58.71	27.89	290.90			
YCR061W	YCR061w	0.91	1.29	1.43	1.80	2.02	2.12	5.80	0.71	46.99	11.38	52.78	12.09	302.73			
YCR061W	SRB8	0.90	1.26	1.38	1.74	1.94	2.04	9.17	2.01	69.53	22.64	78.70	23.82	308.76			
YCR062W	YCR062w	0.93	1.09	1.21	1.70	1.93	2.12	11.70	6.74	62.49	47.48	74.19	54.22	310.70			
YCR063W	BUD1	0.93	1.02	1.08	1.20	1.49	1.81	69.94	56.11	16.55	84.91	16.49	14.01	336.86			
YCR063W	YCR063w	0.95	1.25	1.36	1.77	1.97	2.02	1.57	1.24	32.86	3.26	63.58	3.68	237.41			
YCR069W	RAD18	0.88	1.24	1.40	1.75	1.94	2.06	9.72	0.84	60.15	27.97	69.87	22.81	311.66			
YCR076C	SE04	0.92	1.07	1.23	1.67	1.91	2.08	7.61	0.57	44.43	40.43	62.04	44.79	394.45			
YCR086W	ATG15	0.88	1.22	1.35	1.73	1.91	2.02	23.82	1.63	80.55	6.76	104.37	81.49	316.58			
YCR089W	CPR4	0.89	1.08	1.28	1.77	1.98	2.15	0.77	0.77	11.05	6.27	11.82	7.30	287.72			
YCR071C	IMG2	1.0															

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YDL039C	YDL039c	0.88	1.02	1.19	1.68	1.90	2.06	6.44	6.22	51.58	48.17	58.03	54.39	297.46		
YDL039C	PRM7	0.85	1.08	1.27	1.75	1.97	2.15	0.88	1.02	12.47	13.55	13.35	14.57	267.39		
YDL040C	NAT1	0.85	0.94	0.95	0.84	0.84	0.80	126.50	121.44	146.38	111.37	272.88	232.81	324.81		
YDL041W	YDL041w	0.91	1.17	1.45	1.94	2.11	2.26	2.43	1.08	30.11	14.99	32.54	16.07	287.37		
YDL044C	MFT2	0.92	1.14	1.12	1.06	1.00	0.97	120.42	124.65	133.91	129.12	254.34	253.78	536.65		
YDL045W-A	MGP70	0.89	1.26	1.43	1.27	2.59	2.24	2.43	1.37	33.25	18.83	35.67	20.20	288.80		
YDL046W	NPC2	0.88	1.13	1.34	1.91	2.03	2.02	2.43	1.08	32.59	25.81	34.53	27.67	291.51		
YDL047W	YDL047w	0.94	1.14	1.16	1.22	1.49	1.54	0.95	1.04	12.34	12.50	12.57	12.43	293.45		
YDL048C	STP4	0.92	1.19	1.44	1.87	2.04	2.17	1.82	2.11	1.82	1.86	17.12	11.61	271.31		
YDL049C	KHH1	0.92	1.08	1.07	0.93	0.93	0.90	122.38	128.62	126.27	128.82	249.65	247.44	536.46		
YDL050C	YDL050c	0.89	1.07	1.26	1.72	1.84	2.04	2.09	3.32	1.83	37.25	22.42	49.57	24.05	288.07	
YDL051W	LHP1	0.91	1.09	1.30	1.78	2.00	2.15	2.93	2.75	34.46	31.54	37.39	34.29	292.19		
YDL052C	SLC1	0.88	1.02	1.16	1.64	1.87	2.03	7.44	4.15	50.94	36.88	58.38	41.03	316.44		
YDL053C	PBP4	0.87	1.04	1.22	1.69	1.91	2.06	10.61	4.35	54.39	42.01	65.00	46.36	306.93		
YDL054C	MCH1	0.86	1.04	1.18	1.60	1.82	1.93	8.69	2.92	58.68	35.87	67.37	38.79	313.33		
YDL055W	MBP1	0.89	1.06	1.20	1.56	1.75	1.85	25.65	19.87	73.83	75.51	99.48	95.38	350.90		
YDL057W	YDL057w	0.90	1.07	1.07	0.92	0.92	0.89	124.19	123.54	132.81	136.26	256.99	259.80	537.90		
YDL059C	RAD59	0.83	0.98	1.16	1.59	1.83	1.99	5.31	3.70	46.69	38.72	52.00	42.42	305.97		
YDL061C	RPS29b	0.95	1.07	1.17	1.56	1.89	2.12	5.09	4.90	43.06	44.12	48.15	49.02	264.62		
YDL062W	YDL062w	0.88	1.07	1.06	0.93	0.95	0.96	95.05	88.27	125.55	118.15	220.60	206.42	417.98		
YDL063C	YDL063c	0.86	1.06	1.05	0.92	1.01	1.26	73.13	57.05	114.02	102.18	187.15	159.23	369.98		
YDL065C	PEX19	0.93	1.12	1.31	1.79	1.98	2.14	8.46	3.91	52.23	39.72	60.69	43.64	293.02		
YDL066W	IDP1	0.94	1.18	1.39	1.80	1.99	2.10	5.97	2.42	48.68	30.47	54.65	32.88	303.70		
YDL067C	Cox9	0.99	1.08	1.09	1.01	1.02	1.00	123.50	118.50	121.56	115.56	245.06	234.06	276.01		
YDL068W	YDL068w	0.88	1.08	1.08	1.00	1.25	1.54	34.86	20.57	89.28	73.33	124.14	93.90	300.24		
YDL069C	CBS1	0.94	1.15	1.13	1.00	1.00	0.97	123.87	130.70	127.52	100.76	251.39	231.46	510.03		
YDL069C	CBS1	0.92	1.11	1.10	0.96	0.97	0.94	122.48	120.90	135.97	132.67	258.45	253.57	537.92		
YDL070W	BDF2	0.89	1.03	1.19	1.70	1.95	2.12	4.39	5.39	42.26	43.42	46.65	48.80	294.61		
YDL071C	YDL071c	0.89	1.08	1.23	1.67	1.93	2.09	15.03	6.70	64.17	46.45	79.20	53.15	328.48		
YDL073W	YDL073w	0.84	1.06	1.28	1.73	1.94	2.05	15.35	5.98	61.85	43.52	77.21	49.49	322.62		
YDL074C	BRE1	0.95	1.32	1.53	1.93	2.09	2.24	22.16	3.04	67.75	31.55	89.92	34.59	329.07		
YDL075W	RPL37a	1.04	1.23	1.35	1.88	2.14	2.32	8.40	0.93	50.71	6.77	59.11	7.69	295.85		
YDL076C	RXT3	0.87	1.09	1.33	1.71	1.83	2.07	13.65	11.15	67.61	60.80	81.26	71.95	328.06		
YDL077C	VAM6	0.93	1.12	1.33	1.78	2.00	2.15	7.88	5.77	54.89	45.80	62.87	51.57	297.02		
YDL078C	VAM6	0.94	1.25	1.35	1.56	1.69	1.79	45.76	23.12	100.19	64.70	145.95	87.82	404.38		
YDL079C	MHD3	0.88	1.02	1.23	1.61	1.86	1.96	1.82	1.72	27.29	25.81	29.11	27.53	283.24		
YDL079C	MHK1	0.88	1.02	1.35	1.64	2.04	2.08	2.98	3.65	36.06	37.25	39.42	41.06	285.36		
YDL080C	TRIM3	0.91	1.11	1.27	1.64	1.84	1.94	3.33	3.31	39.48	38.11	43.41	41.62	226.11		
YDL081C	RPP1a	0.87	1.01	1.21	1.28	1.54	1.81	5.21	5.11	55.17	47.73	65.00	52.23	359.05		
YDL082W	YDL082w	0.95	1.05	1.16	1.62	1.88	2.05	2.84	4.45	36.09	42.95	38.83	47.40	271.81		
YDL082W	RPL12a	0.92	1.29	1.38	1.78	2.03	2.21	9.36	3.95	51.43	28.12	69.79	32.98	297.42		
YDL083C	RPS16b	0.88	0.99	1.07	1.46	1.79	2.02	10.63	11.46	60.03	62.18	70.67	73.64	306.34		
YDL085C-A	YDL085c-a	0.93	1.20	1.34	1.75	1.95	2.08	14.47	3.84	67.07	31.20	81.55	35.94	318.84		
YDL085W	NDE2	0.89	1.06	1.24	1.73	1.96	2.11	12.12	4.71	63.46	42.12	75.58	46.83	317.34		
YDL086W	YDL086w	0.89	1.09	1.27	1.75	1.98	2.08	21.22	6.06	73.83	48.92	95.05	54.97	339.59		
YDL088C	ASMA	0.95	1.42	1.62	1.99	2.19	2.22	16.81	1.56	63.79	23.93	80.40	25.50	326.75		
YDL089W	NUR1	0.88	1.35	1.52	1.88	2.11	2.16	14.54	9.79	62.42	50.39	76.96	60.18	321.97		
YDL090C	RAM1	0.95	1.29	1.40	1.69	1.95	2.06	39.75	34.10	87.59	87.60	127.34	121.70	366.10		
YDL091C	UBX3	0.94	1.36	1.55	1.92	2.13	2.17	11.57	4.14	55.41	35.35	66.98	39.49	319.63		
YDL093W	PM75	0.90	1.33	1.56	1.93	2.13	2.19	10.01	1.17	52.83	16.63	62.83	17.79	312.99		
YDL094C	YDL094c	0.87	1.21	1.40	1.88	2.20	2.21	7.21	1.47	55.23	18.18	62.45	19.64	315.62		
YDL095W	PM71	1.01	1.36	1.55	1.92	2.10	2.18	14.95	1.66	61.67	23.19	76.62	24.84	318.48		
YDL096C	YDL096c	1.01	1.42	1.57	1.93	2.10	2.19	12.99	6.63	60.09	45.67	73.08	52.30	325.50		
YDL099W	BUG1	0.95	1.30	1.47	1.88	2.05	2.20	16.25	4.64	63.35	48.04	79.60	43.09	344.78		
YDL100C	GET3	0.94	1.31	1.43	1.81	2.04	2.21	3.23	3.61	37.75	34.51	40.98	38.13	312.78		
YDL101C	DUN1	0.92	1.22	1.26	1.64	1.93	2.15	2.79	4.31	35.14	38.68	37.92	42.98	295.59		
YDL104C	QRT7	0.95	1.23	1.22	1.71	1.97	2.17	12.49	2.01	77.63	52.61	100.08	60.51	262.69		
YDL106C	YDL106w	0.95	1.36	1.53	1.89	2.10	2.19	1.51	1.61	4.75	4.26	5.26	5.26	226.44		
YDL106C	RPL18a	0.95	1.21	1.33	1.74	1.95	2.14	23.52	9.67	74.64	54.87	92.15	44.74	363.93		
YDL106C	RPL18a	0.92	1.25	1.37	1.73	1.92	2.10	17.13	14.44	68.18	60.87	86.31	75.61	348.19		
YDL121C	YDL121c	0.92	1.43	1.65	2.01	2.16	2.27	2.37	6.31	30.55	44.18	32.92	50.49	300.63		
YDL122W	YDL122w	0.98	1.34	1.48	1.91	2.28	2.30	12.23	6.12	61.74	41.75	73.88	47.87	316.14		
YDL123W	SN44	1.00	1.45	1.64	2.02	2.18	2.28	27.07	14.65	80.50	62.52	107.57	77.16	366.38		
YDL124W	YDL124w	0.99	1.30	1.45	1.96	2.06	2.26	11.61	4.99	57.48	33.77	69.09	37.77	357.39		
YDL125C	HWT1	0.95	1.35	1.49	1.89	2.17	2.25	21.78	9.56	74.38	52.01	96.17	61.57	316.19		
YDL127W	PCL2	0.94	1.30	1.42	1.80	2.01	2.17	14.49	14.43	64.63	62.74	81.33	77.18	329.26		
YDL128W	VCX1	0.90	1.29	1.44	1.81	2.00	2.19	1.86	1.32	22.43	2.71	24.29	4.03	276.69		
YDL129W	YDL129w	0.95	1.33	1.48	1.94	2.00	2.10	1.24	1.19	4.28	1.88	5.32	2.27	241.31</		

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE			FRUCTOSE			TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YDL197C	ASF2	1.00	1.40	1.66	2.10	2.26	2.35	21.70	15.41	70.81	64.15	92.51	79.56	346.63	346.63	346.63	346.63	
YDL198C	GGC1	0.92	1.24	1.34	1.76	1.96	2.07	6.71	1.60	54.61	24.30	61.32	25.90	209.79	209.79	209.79	209.79	
YDL199C	YDL199c	0.97	1.37	1.55	1.97	2.14	2.24	23.56	10.47	78.01	60.49	101.57	70.95	341.81	341.81	341.81	341.81	
YDL200C	MGT1	0.98	1.41	1.58	2.00	2.15	2.26	26.34	10.80	79.11	58.48	105.45	69.29	352.30	352.30	352.30	352.30	
YDL201W	TRM6	0.93	1.31	1.45	1.91	2.10	2.22	29.90	18.66	83.58	68.72	113.48	87.38	352.00	352.00	352.00	352.00	
YDL202W	MPL11	0.88	0.96	0.98	0.88	0.89	0.85	121.75	122.80	144.55	110.34	206.30	233.14	429.91	429.91	429.91	429.91	
YDL203C	ACK1	0.98	1.26	1.34	1.63	2.06	2.21	28.59	16.73	82.75	66.09	111.34	82.82	330.42	330.42	330.42	330.42	
YDL204W	RTN2	0.86	1.25	1.33	1.82	2.08	2.20	11.49	12.16	61.16	60.98	72.65	73.14	315.06	315.06	315.06	315.06	
YDL205C	YDL205c	1.06	1.37	1.44	1.99	2.03	2.04	4.60	1.11	40.13	14.16	48.16	14.16	224.14	224.14	224.14	224.14	
YDL206W	US44	1.00	1.37	1.62	2.00	2.15	2.25	23.55	7.63	76.24	51.25	98.89	58.88	281.44	281.44	281.44	281.44	
YDL211C	YDL211c	1.02	1.41	1.61	1.98	2.15	2.24	21.46	7.58	75.24	48.38	86.70	55.97	300.82	300.82	300.82	300.82	
YDL213C	NCR8	1.02	1.43	1.63	2.01	2.18	2.30	21.33	9.40	74.89	55.44	96.22	64.84	333.35	333.35	333.35	333.35	
YDL214C	PBR2	0.98	1.35	1.57	1.94	2.11	2.22	18.16	4.68	71.69	43.87	89.85	48.55	329.41	329.41	329.41	329.41	
YDL215C	GDH2	0.98	1.34	1.51	1.93	2.11	2.25	21.49	9.40	74.86	58.30	96.36	67.70	312.93	312.93	312.93	312.93	
YDL216C	RBF1	0.93	1.27	1.39	1.67	1.86	2.02	24.26	14.31	78.92	70.81	103.19	85.12	319.51	319.51	319.51	319.51	
YDL218W	YDL218w	0.98	1.37	1.52	1.96	2.14	2.27	22.29	4.58	74.49	43.05	96.79	48.03	309.76	309.76	309.76	309.76	
YDL219W	DID1	0.93	1.27	1.37	1.78	1.97	2.11	25.80	9.35	76.23	56.81	102.13	66.56	294.47	294.47	294.47	294.47	
YDL222C	FMP45	0.90	1.28	1.39	1.91	2.09	2.20	23.05	5.31	76.02	45.35	99.07	50.86	232.88	232.88	232.88	232.88	
YDL223C	HBT1	0.89	1.20	1.32	1.78	1.98	2.13	26.94	10.61	79.00	56.35	105.94	66.96	236.65	236.65	236.65	236.65	
YDL224C	WH4	0.96	1.24	1.30	1.77	2.01	2.19	5.57	7.80	47.76	50.31	53.33	58.12	241.33	241.33	241.33	241.33	
YDL225W	SHS1	0.94	1.20	1.33	1.68	2.00	2.07	36.53	19.34	89.97	72.17	126.50	91.51	319.14	319.14	319.14	319.14	
YDL226C	GCS1	0.97	1.33	1.55	1.92	2.09	2.20	19.74	3.17	74.75	34.88	94.49	38.05	271.60	271.60	271.60	271.60	
YDL227C	H6	1.05	1.38	1.54	1.88	2.04	2.18	22.15	14.12	77.91	65.35	100.05	79.48	317.12	317.12	317.12	317.12	
YDL229W	SSB1	1.02	1.39	1.57	1.93	2.12	2.25	15.57	2.88	65.58	32.48	81.15	35.37	332.98	332.98	332.98	332.98	
YDL230W	PTP1	0.97	1.33	1.54	1.93	2.10	2.20	19.09	6.22	77.22	50.50	96.31	56.72	342.46	342.46	342.46	342.46	
YDL231C	BRE4	1.03	1.40	1.63	2.07	2.24	2.34	13.77	5.62	65.79	45.65	79.57	51.27	321.31	321.31	321.31	321.31	
YDL232W	OST4	0.98	1.38	1.58	2.00	2.17	2.28	21.88	11.97	77.53	63.05	99.41	75.02	339.49	339.49	339.49	339.49	
YDL233W	YDL233w	0.95	1.31	1.45	1.81	2.00	2.14	27.67	7.02	83.90	50.70	111.58	57.72	353.17	353.17	353.17	353.17	
YDL234C	GYP7	0.94	1.31	1.43	1.86	2.06	2.21	25.47	4.78	79.38	41.79	104.85	46.57	335.96	335.96	335.96	335.96	
YDL236W	PHO13	0.96	1.36	1.53	2.01	2.19	2.29	22.31	4.15	74.13	38.68	96.44	42.83	344.34	344.34	344.34	344.34	
YDL237W	AIM6	0.91	1.23	1.36	1.83	2.03	2.18	30.03	8.71	84.06	54.11	114.09	62.82	350.43	350.43	350.43	350.43	
YDL238C	GUD1	1.01	1.32	1.44	1.81	2.11	2.12	2.10	7.72	7.89	53.26	50.40	60.98	58.29	308.08	308.08	308.08	308.08
YDL239C	ADY3	0.95	1.33	1.54	1.94	2.10	2.20	26.77	15.37	79.98	64.18	106.75	79.55	307.86	307.86	307.86	307.86	
YDL240W	LRG1	1.02	1.42	1.67	2.08	2.23	2.31	38.12	16.37	91.51	68.02	129.53	84.39	322.84	322.84	322.84	322.84	
YDL241W	YDL241w	0.99	1.33	1.50	1.88	2.04	2.15	15.56	7.62	70.32	50.26	85.88	57.87	236.81	236.81	236.81	236.81	
YDL242W	YDL242w	1.01	1.47	1.72	2.08	2.20	2.26	17.68	6.62	68.67	47.67	86.54	54.69	280.88	280.88	280.88	280.88	
YDL243C	AAD4	1.02	1.49	1.74	2.11	2.25	2.33	23.96	8.18	75.24	51.58	99.20	59.76	320.61	320.61	320.61	320.61	
YDL301C	NTTH1	1.01	1.32	1.52	1.94	2.14	2.26	23.03	9.11	6.63	46.53	65.19	53.16	317.43	317.43	317.43	317.43	
YDR030C	RCP2	0.98	1.27	1.41	1.88	2.05	2.12	23.03	6.58	72.72	50.37	82.36	55.93	316.83	316.83	316.83	316.83	
YDR030W	YDR030w-a	0.87	1.18	1.30	1.58	1.77	1.81	10.65	3.48	59.23	29.27	92.88	32.74	308.34	308.34	308.34	308.34	
YDR030W	RAD57	0.91	1.34	1.53	1.80	2.00	2.16	20.83	11.92	76.19	64.14	96.22	76.06	316.86	316.86	316.86	316.86	
YDR030C	MAF1	0.91	1.22	1.34	1.71	1.90	2.06	23.36	4.16	76.38	38.88	97.74	42.74	255.43	255.43	255.43	255.43	
YDR030C	SOK1	0.83	1.29	1.45	1.87	2.03	2.15	28.66	8.49	81.90	54.07	110.58	62.55	284.39	284.39	284.39	284.39	
YDR007W	TRP1	0.88	1.18	1.30	1.82	1.83	1.93	3.36	5.35	35.29	40.72	38.65	46.06	304.93	304.93	304.93	304.93	
YDR008C	YDR018c	0.98	1.39	1.57	1.70	1.91	2.06	11.28	1.74	57.07	19.13	68.35	20.87	312.35	312.35	312.35	312.35	
YDR009W	YDR009c	0.89	1.23	1.30	1.81	1.83	1.98	20.56	13.52	45.41	64.48	78.00	46.89	232.29	232.29	232.29	232.29	
YDR030C	RAD28	0.90	1.35	1.51	1.96	2.16	2.29	6.26	2.16	34.98	20.96	54.84	40.35	233.39	233.39	233.39	233.39	
YDR031W	YDR031w	0.97	1.27	1.44	1.82	2.01	2.13	2.27	4.27	41.72	34.77	59.21	34.77	201.56	201.56	201.56	201.56	
YDR031W	YDR031w	0.97	1.27	1.44	1.82	2.01	2.13	14.90	6.57	62.71	42.06	83.50	55.06	146.30	146.30	146.30	146.30	
YDR032W	ERG1	0.90	1.21	1.32	1.76	1.92	2.02	12.61	2.41	62.71	42.06	74.58	38.87	150.44	150.44	150.44	150.44	
YDR034C	YDR042c	0.90	1.27	1.41	1.82	2.01	2.13	2.27	4.27	41.72	32.37	45.63	37.25	229.28	229.28	229.28	229.28	
YDR043C	NRG1	1.04	1.54	1.68	1.96	2.10	2.14	49.49	32.28	163.40	86.94	152.89	119.21	222.41	222.41	222.41	222.41	
YDR046C	BAP2	0.92	1.35	1.47	1.88	2.07	2.15	4.14	3.34	45.29	37.67	49.43	41.02	184.11	184.11	184.11	184.11	
YDR048C	YDR048c	1.00	1.09	1.19	1.58	1.88	1.93	2.20	26.69	1.82	83.44	34.23	110.04	36.05	269.10	269.10	269.10	269.10
YDR049W	YDR049w	0.99	1.10	1.31	1.84	2.02	2.16	2.94	0.17	40.09	7.41	43.03	7.58	312.39	312.39	312.39	312.39	
YDR050C	TRI1	0.90	1.07	1.19	1.77	2.01	2.19	18.85	2.71	64.79	52.74	83.64	63.45	266.88	266.88	266.88	266.88	
YDR051C	DET1	0.86	1.18	1.37	1.73	1.88	2.04	1.43	1.66	12.51	14.84	13.94	16.50	268.68	268.68	26		

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YDR122W	KIN1	0.97	1.28	1.50	2.00	2.16	2.29	1.65	0.72	25.55	17.38	27.20	18.10	280.31		
YDR123C	INO2	0.82	0.90	0.86	0.90	0.92	0.89	122.29	119.20	124.07	120.08	246.36	239.29	533.74		
YDR123C	INO2	0.87	1.04	1.03	0.92	0.97	1.14	46.17	15.31	99.48	71.22	145.65	86.53	300.99		
YDR124W	YDR124w	0.88	1.16	1.33	1.78	1.85	2.00	1.23	1.20	3.71	0.79	4.94	1.99	244.96		
YDR126W	SWF1	1.06	1.32	1.52	1.93	2.04	2.18	23.21	17.62	70.06	62.48	93.27	80.10	340.62		
YDR127W	ARTO	0.56	1.11	1.11	1.03	1.13	1.02	108.98	102.86	141.08	142.05	250.06	244.91	470.18		
YDR128W	MTC5	0.88	1.30	1.49	1.87	2.01	2.17	25.30	16.37	74.66	64.31	99.57	88.68	365.49		
YDR130C	FIM1	0.84	1.24	1.24	1.98	2.08	2.14	18.82	14.82	82.25	74.59	102.59	92.55	332.55		
YDR131C	YDR131c	0.97	1.37	1.62	2.05	2.19	2.31	6.53	7.55	46.71	49.45	53.24	57.00	399.01		
YDR132C	YDR132c	1.01	1.41	1.43	1.86	2.05	2.18	10.27	8.96	58.04	49.58	68.31	57.54	316.83		
YDR133C	YDR133c	0.95	1.26	1.43	1.86	2.05	2.19	0.77	0.14	18.75	8.46	19.52	8.60	290.81		
YDR134C	YDR134c	0.99	1.29	1.49	1.95	2.11	2.24	14.75	3.87	62.13	36.09	76.88	39.96	322.12		
YDR135C	YCF1	0.90	1.07	1.14	1.52	1.71	1.84	17.69	5.65	68.58	45.98	86.27	51.63	341.02		
YDR136C	VPS61	0.99	1.24	1.39	1.86	2.03	2.12	18.91	2.61	64.00	33.07	82.91	35.67	343.41		
YDR137W	RGP1	0.97	1.16	1.32	1.84	2.11	2.26	0.53	0.10	17.34	1.37	17.87	1.47	300.27		
YDR138W	HPR1	0.92	1.14	1.13	1.00	1.02	1.05	108.08	89.17	132.77	121.80	240.85	219.97	445.92		
YDR138W	HPR1	0.84	1.06	1.05	1.03	1.21	1.35	29.77	12.71	83.74	59.62	113.51	72.33	353.69		
YDR139C	RUB1	0.98	1.27	1.47	1.87	2.02	2.16	17.52	6.93	66.97	47.29	84.49	54.22	341.47		
YDR140W	MTO2	1.01	1.35	1.57	1.96	2.10	2.23	16.08	4.32	63.30	40.87	79.39	45.19	343.68		
YDR142C	PEX7	0.99	1.23	1.39	1.86	1.99	2.14	8.72	5.71	52.60	42.19	61.32	47.90	307.49		
YDR143C	SAN1	1.03	1.41	1.64	2.08	2.22	2.34	15.28	9.84	67.30	57.01	82.58	66.85	309.55		
YDR144C	MKC7	0.97	1.17	1.34	1.93	2.15	2.30	3.15	1.98	36.87	30.18	40.02	32.16	304.16		
YDR146C	SW5	1.00	1.37	1.58	2.02	2.23	2.34	0.80	0.72	23.35	16.97	24.14	17.69	286.61		
YDR147W	EKI1	1.02	1.22	1.38	1.94	2.11	2.26	9.41	7.61	54.30	49.12	63.72	56.72	289.93		
YDR147W	EKI1	0.99	1.35	1.52	1.93	2.09	2.23	2.85	1.02	32.47	7.16	35.32	8.18	271.60		
YDR148C	KGD2	1.04	1.12	1.10	1.07	1.10	1.32	103.46	95.51	125.26	122.74	228.72	219.25	424.10		
YDR149C	YDR149c	1.02	1.21	1.31	1.68	1.87	2.07	34.15	14.98	83.62	62.99	117.77	77.98	326.94		
YDR150W	NUM1	1.08	1.47	1.66	2.01	2.16	2.32	7.41	1.38	49.40	16.39	56.81	17.78	282.54		
YDR151C	CTH1	0.93	1.18	1.39	1.86	2.02	2.18	10.82	1.80	58.43	22.66	69.25	24.66	313.78		
YDR152W	GIR2	1.03	1.27	1.47	1.90	2.06	2.23	5.32	2.33	44.23	27.76	49.56	30.09	249.78		
YDR153C	ENT5	0.91	1.12	1.29	1.85	2.08	2.05	2.96	1.10	35.95	10.39	38.91	11.49	252.96		
YDR154C	YDR154c	1.08	1.37	1.52	1.91	2.08	2.22	2.76	1.12	38.93	13.31	41.69	14.43	288.47		
YDR155C	CPR1	0.91	1.12	1.26	1.71	1.89	2.08	2.40	1.15	34.78	12.66	37.18	14.02	290.35		
YDR156W	RPAT1	1.02	1.24	1.35	1.81	2.02	2.21	9.82	2.38	57.36	30.34	67.19	32.73	258.45		
YDR157W	YDR157w	0.93	1.30	1.50	1.74	2.00	2.14	3.59	1.78	39.71	17.17	43.30	18.94	272.52		
YDR158C	YDR158c	0.97	1.09	1.28	1.42	1.62	1.78	33.62	19.71	53.69	119.51	171.11	131.29	331.29		
YDR159W	SAC3	0.96	1.18	1.36	1.62	1.80	1.97	78.29	61.65	119.63	122.84	192.92	173.85	446.08		
YDR159W	SAC3	0.82	1.01	0.99	0.97	0.99	1.17	67.49	51.31	104.89	127.30	145.92	139.14	308.66		
YDR161W	YDR161w	0.98	1.18	1.37	1.70	1.91	2.01	17.45	13.78	70.33	62.45	87.79	76.22	313.97		
YDR162C	NBP2	0.99	1.53	1.88	2.18	2.29	2.33	14.21	5.67	69.70	38.53	74.91	43.61	377.47		
YDR163W	CWC15	0.95	1.21	1.42	1.90	2.07	2.25	7.52	1.25	48.92	11.57	56.43	12.81	314.80		
YDR165W	TRM62	1.04	1.36	1.55	2.01	2.17	2.31	16.47	3.71	66.07	36.20	82.54	38.91	302.31		
YDR169C	STB3	0.91	1.04	1.14	1.55	1.72	1.92	8.88	2.75	55.31	33.43	64.19	36.19	283.91		
YDR169C-A	YDR169c-a	0.92	1.22	1.46	1.85	2.05	2.18	9.81	4.57	56.78	32.09	66.59	36.66	308.32		
YDR171W	HSP42	0.90	1.09	1.21	1.77	1.79	1.93	3.55	1.33	36.81	15.36	40.16	16.69	273.17		
YDR173C	ARG62	0.93	1.32	1.40	1.84	2.00	2.16	1.31	1.14	12.95	3.42	14.25	4.56	246.83		
YDR174W	HMO1	0.86	0.90	0.93	1.11	1.34	1.59	57.35	43.36	98.60	90.77	155.95	134.14	299.43		
YDR175C	RSM24	0.88	1.05	1.06	0.91	0.90	0.85	127.20	37.90	142.85	100.36	270.04	138.26	517.55		
YDR176W	NGG1	0.94	1.07	1.08	1.35	1.51	1.73	38.63	22.99	94.29	77.44	132.93	100.43	363.21		
YDR178W	SDH4	0.92	1.13	1.27	1.71	1.89	2.09	2.41	4.18	28.22	38.07	30.63	42.25	273.46		
YDR179C	CNS9	0.98	1.15	1.27	1.80	2.00	2.20	11.02	10.80	58.70	55.32	69.72	66.12	294.54		
YDR179W-A	YDR179w-a	0.90	1.11	1.23	1.73	1.92	2.10	4.40	2.26	41.46	28.76	45.86	31.03	284.33		
YDR181C	SAS4	0.96	1.13	1.20	1.57	1.78	2.01	3.15	1.35	34.57	13.85	37.72	15.21	288.47		
YDR182W	YDR182w	0.91	1.20	1.37	1.75	1.96	2.12	1.75	1.51	46.89	37.75	51.82	44.74	273.43		
YDR183W	PLP1	1.07	1.43	1.64	2.08	2.22	2.33	13.39	2.97	62.00	31.89	75.39	34.87	311.74		
YDR184C	ATC1	1.08	1.34	1.48	1.96	2.07	2.22	23.77	5.68	79.20	44.13	102.97	49.81	329.81		
YDR185C	UPS3	1.05	1.39	1.58	1.99	2.14	2.26	21.30	4.38	74.10	40.13	95.39	44.50	333.27		
YDR186C	YDR186c	0.94	1.11	1.22	1.54	1.79	2.02	0.95	1.12	2.75	0.51	3.70	1.63	257.64		
YDR191W	HST4	0.90	1.10	1.21	1.61	1.78	2.00	3.15	1.35	34.57	14.72	37.72	15.21	288.47		
YDR192C	NUP42	0.91	1.10	1.21	1.63	1.82	2.02	1.23	0.99	40.25	17.75	41.48	16.26	247.81		
YDR193W	YDR193w	0.97	1.18	1.30	1.68	1.86	2.08	25.05	11.68	77.65	57.84	102.90	82.82	332.26		
YDR196C	YDR196c	0.93	1.12	1.28	1.69	1.87	2.05	21.89	5.88	77.57	65.13	99.46	77.25	278.93		
YDR200C	YPS84	1.08	1.22	1.25	1.50	1.66	1.88	21.89	3.63	3.63	37.09	9.33	40.73	10.47	275.52	
YDR202C	RAV2	0.90	1.09	1.30	1.75	1.88	2.09	37.79	30.97	87.19	82.27	124.98	115.24	308.66		
YDR203W	YDR203w	1.10	1.47	1.70	2.09	2.20	2.29	43.86	27.17	98.22	80.81	142.17	107.98	355.15		
YDR204W	COD4	0.82	0.98	0.99	0.84	0.84	0.81	128.52	128.81	140.38	261.72	270.17	151.77	519.44		
YDR205C	MSC2	0.94	1.16	1.39	1.84	2.03	2.19	15.74	9.30	62.62	50.92	78.35	69.22	328.76		
YDR206W	EB51	0.97	1.20	1.33	1.76	1.96	2.17	0.99	1.27	3.74	4.44	4.73	5.71	261.42		
YDR22																

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE			FRUCTOSE			TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YDR277C	MTH1	0.94	1.47	1.60	2.00	2.13	2.23	7.44	9.81	46.77	48.93	54.21	58.74	307.82	307.82	307.82	307.82	
YDR278C	YDR278c	0.92	1.44	1.51	1.94	2.10	2.21	6.87	11.55	40.64	50.15	47.51	61.70	303.69	303.69	303.69	303.69	
YDR279W	RNH202	0.98	1.49	1.58	2.05	2.22	2.34	4.88	8.37	34.55	40.02	39.43	48.40	294.15	294.15	294.15	294.15	
YDR281C	PHM6	0.94	1.46	1.48	1.86	2.03	2.15	10.38	10.88	51.40	50.54	61.79	61.43	316.51	316.51	316.51	316.51	
YDR283C	GCR2	0.91	1.12	1.27	1.65	1.86	1.98	7.02	-0.01	54.64	4.89	61.66	4.88	301.58	301.58	301.58	301.58	
YDR284C	DPT1	0.91	1.41	1.50	1.91	2.06	2.10	5.81	7.17	41.54	43.48	47.35	50.65	279.63	279.63	279.63	279.63	
YDR285W	ZIP1	0.91	1.49	1.54	1.93	2.03	2.18	8.67	8.89	48.93	42.30	57.68	49.10	312.14	312.14	312.14	312.14	
YDR286C	YDR286c	0.92	1.45	1.54	1.92	2.03	2.18	2.26	2.28	32.04	34.61	32.02	32.02	303.69	303.69	303.69	303.69	
YDR287W	IMR2	0.92	1.45	1.51	1.91	2.06	2.22	3.06	3.41	42.46	16.83	31.92	20.03	288.63	288.63	288.63	288.63	
YDR289C	RTT103	0.92	1.41	1.32	1.78	1.95	2.08	11.68	12.15	52.25	54.88	63.93	66.83	319.73	319.73	319.73	319.73	
YDR290W	YDR290w	1.00	1.51	1.54	1.97	2.09	2.22	19.35	24.76	63.22	70.89	82.58	95.65	344.70	344.70	344.70	344.70	
YDR291W	HRO1	0.94	1.45	1.60	2.00	2.15	2.25	4.92	11.86	33.78	48.31	38.70	60.18	297.11	297.11	297.11	297.11	
YDR293C	SSDI	0.97	1.51	1.60	2.00	2.15	2.29	3.72	3.84	21.56	19.68	25.28	23.52	277.65	277.65	277.65	277.65	
YDR294C	DPL1	0.96	1.49	1.56	2.00	2.15	2.28	3.53	6.11	21.20	33.34	24.74	39.45	286.04	286.04	286.04	286.04	
YDR295C	HDA2	0.88	1.22	1.30	1.73	1.96	2.10	2.18	3.27	32.38	12.51	34.56	13.88	211.93	211.93	211.93	211.93	
YDR296W	MHR1	0.93	1.04	1.03	0.92	0.93	0.91	124.00	123.76	151.41	116.87	275.41	246.64	418.55	418.55	418.55	418.55	
YDR297W	SUR2	0.96	1.55	1.72	2.03	2.14	2.21	3.56	3.28	19.01	13.01	22.57	16.28	302.02	302.02	302.02	302.02	
YDR298C	ATP5	0.85	1.19	1.27	1.50	1.66	1.75	10.25	2.78	57.95	31.61	68.20	34.39	249.16	249.16	249.16	249.16	
YDR300C	PRO1	1.00	1.07	1.11	1.01	1.01	0.99	125.83	123.74	145.57	111.99	271.40	235.74	436.62	436.62	436.62	436.62	
YDR304C	CPR5	0.93	1.47	1.63	1.97	2.09	2.18	7.19	5.19	42.23	30.49	49.42	39.24	304.43	304.43	304.43	304.43	
YDR305C	HNT2	0.90	1.42	1.58	1.94	2.08	2.15	6.79	5.10	45.03	33.49	51.82	38.59	299.91	299.91	299.91	299.91	
YDR306C	YDR306c	0.98	1.45	1.46	1.93	2.11	2.20	6.25	3.27	42.50	12.27	48.74	15.55	300.91	300.91	300.91	300.91	
YDR307W	YDR307w	0.94	1.49	1.61	1.98	2.12	2.22	5.28	3.68	36.52	24.89	41.80	28.56	293.72	293.72	293.72	293.72	
YDR309C	GIC2	0.92	1.45	1.57	2.01	2.15	2.26	3.29	3.49	16.13	19.43	19.43	22.92	279.94	279.94	279.94	279.94	
YDR310C	SUM1	0.95	1.44	1.48	1.87	2.01	2.15	9.07	6.44	47.53	37.29	56.60	43.73	303.32	303.32	303.32	303.32	
YDR312W	SSF2	0.93	1.43	1.54	1.94	2.08	2.20	7.93	9.80	47.20	49.46	55.14	59.26	308.30	308.30	308.30	308.30	
YDR313C	PIB1	0.94	1.49	1.61	2.03	2.16	2.26	4.47	4.59	31.76	50.28	33.59	58.64	38.35	38.35	38.35	38.35	
YDR314C	RAD34	1.02	1.56	1.71	2.11	2.27	2.36	3.74	5.23	24.43	32.03	28.18	37.26	282.75	282.75	282.75	282.75	
YDR315C	IPK1	1.00	1.48	1.46	1.79	2.02	2.22	23.54	27.88	67.85	73.94	91.39	101.83	323.61	323.61	323.61	323.61	
YDR316W	OMS1	0.95	1.49	1.63	2.02	2.18	2.27	3.24	3.89	14.10	17.75	17.35	23.64	258.02	258.02	258.02	258.02	
YDR317W	HIM1	0.94	1.42	1.44	1.87	2.03	2.15	5.40	6.07	38.44	39.86	43.84	45.93	291.56	291.56	291.56	291.56	
YDR318W	MCM21	0.91	1.31	1.27	1.55	1.75	1.91	25.80	24.33	77.12	77.80	102.92	102.23	333.51	333.51	333.51	333.51	
YDR319W	MCM21	1.02	1.34	1.38	1.69	1.90	2.11	26.09	7.92	64.85	46.05	110.94	53.97	327.90	327.90	327.90	327.90	
YDR319C	YDR319c	0.88	1.38	1.49	1.86	2.03	2.13	8.36	7.76	50.28	33.59	58.64	38.35	304.43	304.43	304.43	304.43	
YDR320C	SWA2	0.91	1.34	1.19	1.62	2.11	2.17	27.30	17.54	73.62	64.09	100.92	81.83	319.25	319.25	319.25	319.25	
YDR321W	ASD1	0.91	1.34	1.51	1.62	2.05	2.24	5.83	3.11	39.65	12.30	45.52	15.47	297.44	297.44	297.44	297.44	
YDR322C-A	LM11	0.88	1.28	1.35	1.68	1.97	2.03	9.04	1.48	53.24	16.87	62.28	16.34	308.12	308.12	308.12	308.12	
YDR323C	MRPL35	0.89	1.46	1.62	0.99	0.97	1.03	85.91	85.71	131.89	131.51	226.91	219.15	409.47	409.47	409.47	409.47	
YDR323C	PEP7	0.88	1.42	1.41	1.80	2.05	2.25	28.38	34.12	74.13	32.85	120.52	117.07	329.62	329.62	329.62	329.62	
YDR326C	YSP2	0.97	1.24	1.47	1.87	2.03	2.18	13.01	1.17	55.69	18.43	68.70	16.59	311.19	311.19	311.19	311.19	
YDR328C	PEX3	0.84	1.49	1.58	1.95	2.09	2.19	11.94	9.08	56.94	49.17	68.88	58.25	299.88	299.88	299.88	299.88	
YDR329W	UBX5	0.94	1.46	1.57	1.98	2.12	2.24	7.04	4.92	45.71	33.07	52.75	37.90	304.43	304.43	304.43	304.43	
YDR329W	IRC3	0.98	1.51	1.55	1.99	2.15	2.28	4.17	5.05	26.67	31.09	36.84	36.14	283.06	283.06	283.06	283.06	
YDR333C	YDR333c	0.96	1.52	1.63	1.98	2.11	2.23	3.83	4.03	25.52	25.95	29.35	29.98	284.33	284.33	284.33	284.33	
YDR334W	SWR1	0.91	1.32	1.25	1.62	1.80	1.94	19.28	25.11	67.40	67.50	86.67	92.61	322.47	322.47	322.47	322.47	
YDR335W	MSN5	0.90	1.39	1.44	1.84	2.02	2.14	2.91	3.96	13.75	21.70	16.66	25.66	283.33	283.33	283.33	283.33	
YDR336W	YDR336w	0.93	1.43	1.52	1.87	2.04	2.19	11.62	4.05	52.44	34.02	58.69	46.52	311.92	311.92	311.92	311.92	
YDR337W	MRPS28	0.89	1.41	1.53	1.88	2.04	2.16	4.88	4.84	5.48	32.44	1.86	296.76	296.76	296.76	296.76		
YDR338C	MRPS28	0.91	1.26	1.33	1.71	1.89	2.09	33.44	15.50	63.23	62.78	116.87	78.28	351.91	351.91	351.91	351.91	
YDR339W	SEMA1	0.99	1.23	1.33	1.71	1.89	2.09	33.44	15.50	63.23	62.78	116.87	78.28	351.91	351.91	351.91	351.91	
YDR340W	CDC40	0.96	1.18	1.25	1.60	1.85	2.05	7.53	7.53	73.53	63.34	102.63	12.57	404.14	404.14	404.14	404.14	
YDR341W	TPR1	0.91	1.31	1.32	1.97	2.12	2.29	14.08	5.21	61.51	41.89	75.57	47.10	348.19	348.19	348.19	348.19	
YDR343C	XRS2	0.99	1.14	1.19	1.48	1.66	1.88	47.82	42.46	94.67	90.57	142.48	130.93	410.31	410.31	410.31	410.31	
YDR370C	YDR370c	0.95	1.30	1.47	1.85	2.00	2.12	10.30	2.84	57.76	26.35	68.06	29.19	311.33	311.33	311.33	311.33	
YDR371W	CT23	0.92	1.33	1.55	1.99	2.15	2.32	0.46	0.06	22.78	7.54	23.24	7.48	312.96	312.96	312.96	312.96	
YDR372C	WST4	0.90	1.32	1.49	1.84	2.07	2.10	6.10	5.22	51.42	34.05	49.90	36.73	56.87	40.96	307.67	307.67	
YDR374C	YDR374c	0.91	1.42	1.62	2.08	2.21	2.34	1.48	0.92	23.44	24.92	24.92	24.92	293.86	293.86	293.86	293.86	
YDR375C	BCS1	0.97	1.04	1.0														

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE			FRUCTOSE			TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YDR447C	RPS17b	0.97	1.29	1.48	1.89	2.14	2.26	2.02	0.95	25.23	9.86	27.25	10.81	305.47	305.47	305.47	305.47	
YDR448W	ADA2	1.01	1.32	1.41	1.64	1.83	1.88	53.67	37.34	101.27	85.17	154.94	122.52	360.88	360.88	360.88	360.88	
YDR448W	ADA2	0.92	1.17	1.24	1.46	1.65	1.81	33.45	24.53	85.45	57.99	118.90	82.52	326.11	326.11	326.11	326.11	
YDR450W	RPS18a	0.95	1.38	1.57	1.90	2.14	2.27	6.12	0.92	39.25	7.40	45.38	8.32	328.90	328.90	328.90	328.90	
YDR451C	YHP1	0.95	1.37	1.59	2.04	2.26	2.38	2.13	0.86	25.70	2.34	27.83	3.19	286.01	286.01	286.01	286.01	
YDR452W	PPN1	0.93	1.31	1.50	1.91	2.22	2.19	9.42	0.88	50.40	5.88	59.82	6.80	304.16	304.16	304.16	304.16	
YDR453C	TSZ2	1.04	1.41	1.63	2.07	2.25	2.35	1.98	0.83	15.46	3.63	17.44	4.76	290.62	290.62	290.62	290.62	
YDR454C	YDR455c	1.07	1.43	1.60	1.93	2.17	2.19	18.88	8.15	71.18	52.51	90.95	60.76	373.23	373.23	373.23	373.23	
YDR454W	YDR455c	1.04	1.48	1.60	2.09	2.18	2.24	25.49	12.45	78.40	60.91	104.04	72.99	373.23	373.23	373.23	373.23	
YDR455W	YDR455c	0.89	1.15	1.25	1.71	2.03	2.13	12.64	6.77	57.35	9.174	89.99	343.94	343.94	343.94	343.94		
YDR456C	HEH2	0.97	1.35	1.60	1.97	2.22	2.30	1.69	0.77	24.53	4.88	28.22	5.65	298.68	298.68	298.68	298.68	
YDR456C	PEF1	1.05	1.45	1.65	2.03	2.23	2.32	0.89	0.75	13.69	2.07	14.58	2.82	301.18	301.18	301.18	301.18	
YDR456C	MFA1	0.89	1.08	1.27	1.66	1.85	2.02	6.73	1.97	42.15	22.24	48.88	24.22	301.03	301.03	301.03	301.03	
YDR462W	MFB28	0.97	1.28	1.50	1.91	2.13	2.28	22.22	18.02	7.15	64.92	47.58	82.93	54.73	320.37	320.37	320.37	320.37
YDR463W	STP1	0.95	1.28	1.44	1.82	2.05	2.14	5.26	2.93	42.18	32.22	47.43	35.15	270.29	270.29	270.29	270.29	
YDR465C	RM72	0.97	1.34	1.53	1.86	2.10	2.21	4.88	2.76	43.70	29.31	48.57	32.07	308.31	308.31	308.31	308.31	
YDR466W	PKH3	0.91	1.23	1.39	1.75	1.98	2.06	15.66	2.75	62.96	30.89	78.61	33.64	304.34	304.34	304.34	304.34	
YDR467C	YDR467c	0.97	1.48	1.74	2.10	2.25	2.31	13.02	1.17	58.62	14.82	71.64	15.99	323.66	323.66	323.66	323.66	
YDR469W	SDC1	0.95	1.41	1.69	2.04	2.22	2.31	15.85	1.00	62.01	12.27	77.86	13.37	325.44	325.44	325.44	325.44	
YDR470C	UGO1	0.97	1.14	1.19	1.38	1.66	1.82	38.12	27.79	90.86	80.59	128.98	108.38	347.95	347.95	347.95	347.95	
YDR470C	UGO1	1.10	1.42	1.44	1.51	1.77	2.04	54.89	43.01	109.87	99.07	164.76	142.08	368.25	368.25	368.25	368.25	
YDR471W	RPL27b	1.05	1.30	1.47	1.97	2.17	2.28	16.69	5.59	63.12	42.84	79.81	48.43	337.59	337.59	337.59	337.59	
YDR474C	YDR474c	0.98	1.32	1.52	1.98	2.18	2.26	20.66	5.44	69.37	43.11	90.04	48.54	335.93	335.93	335.93	335.93	
YDR475C	JIP4	0.89	1.21	1.37	1.74	1.93	2.11	2.06	0.87	29.76	7.08	31.82	7.95	284.79	284.79	284.79	284.79	
YDR475C	JIP4	0.95	1.29	1.50	1.94	2.13	2.20	18.51	9.04	66.69	50.94	85.21	59.97	330.76	330.76	330.76	330.76	
YDR476C	YDR476c	0.97	1.28	1.50	1.97	2.20	2.27	3.72	1.03	38.23	17.46	41.94	18.49	302.53	302.53	302.53	302.53	
YDR477W	SNF1	0.91	1.04	1.10	1.44	1.73	2.03	24.45	3.81	71.77	37.12	96.23	40.94	307.51	307.51	307.51	307.51	
YDR479C	PEX29	0.94	1.29	1.51	1.94	2.06	2.12	4.92	0.95	41.62	12.32	46.55	13.28	302.05	302.05	302.05	302.05	
YDR480W	DIG2	0.96	1.46	1.68	2.02	2.12	2.25	2.95	0.88	29.41	7.29	32.36	8.09	276.85	276.85	276.85	276.85	
YDR481C	PHOB	0.96	1.33	1.54	1.93	2.14	2.23	11.99	2.90	58.19	33.88	70.18	36.78	295.40	295.40	295.40	295.40	
YDR482C	CWC21	0.87	1.08	1.22	1.61	1.87	1.97	10.46	5.81	57.11	41.65	67.57	47.46	293.15	293.15	293.15	293.15	
YDR483W	KRE2	0.93	1.17	1.37	1.79	1.98	2.14	10.27	4.11	51.82	30.89	62.09	35.00	309.77	309.77	309.77	309.77	
YDR484W	VPS22	0.99	1.42	1.62	1.92	2.05	2.09	6.66	1.72	44.45	21.20	51.11	22.92	261.03	261.03	261.03	261.03	
YDR485C	VPS7	0.95	1.25	1.37	1.75	1.98	2.05	24.09	6.39	72.62	41.43	96.70	47.82	329.54	329.54	329.54	329.54	
YDR486C	VPS6	0.94	1.26	1.42	1.82	2.17	2.15	1.68	0.86	25.91	6.22	27.59	7.08	253.42	253.42	253.42	253.42	
YDR488C	PAC11	0.91	1.22	1.36	1.71	1.92	2.05	12.29	3.98	60.20	38.67	72.50	42.85	291.70	291.70	291.70	291.70	
YDR489C	PRK1	0.95	1.31	1.43	1.95	2.16	2.23	19.36	5.31	63.64	31.21	69.42	24.72	333.38	333.38	333.38	333.38	
YDR490C	YDR491c	1.08	1.46	1.63	2.14	2.44	2.52	14.61	2.25	58.83	24.63	73.43	26.88	331.91	331.91	331.91	331.91	
YDR492W	ZHI1	0.95	1.26	1.48	1.69	2.17	2.27	15.12	2.17	63.00	44.58	78.11	26.79	321.61	321.61	321.61	321.61	
YDR493W	MBM1	0.92	1.20	1.43	2.20	2.23	2.29	4.99	1.24	37.79	13.17	42.37	14.41	288.63	288.63	288.63	288.63	
YDR494W	RS408	0.93	1.21	1.35	1.84	1.95	2.03	15.29	3.63	60.53	24.40	78.92	37.93	319.49	319.49	319.49	319.49	
YDR495C	VPS3	0.91	1.21	1.35	1.86	1.98	2.06	5.20	1.60	60.29	28.32	65.49	29.44	353.12	353.12	353.12	353.12	
YDR496C	PUF6	0.97	1.25	1.37	1.75	2.03	2.17	6.93	1.32	49.24	20.04	56.17	21.36	280.46	280.46	280.46	280.46	
YDR497C	ITR1	0.83	1.07	1.26	1.79	2.02	2.11	11.49	14.51	61.53	32.97	73.02	78.48	307.68	307.68	307.68	307.68	
YDR500C	RPL37b	0.89	1.16	1.26	1.69	1.98	2.11	2.14	0.86	29.31	9.50	31.45	10.37	229.95	229.95	229.95	229.95	
YDR506C	YDR506c	0.96	1.38	1.61	2.04	2.21	2.31	1.05	1.05	20.72	16.30	21.76	17.35	273.38	273.38	273.38	273.38	
YDR507C	YDR507c	0.98	1.22	1.31	1.74	1.97	2.03	1.07	0.99	127.93	135.31	125.31	132.02	253.24	253.24	253.24	253.24	
YDR507C	GIN4	1.02	1.16	1.16	1.80	1.95	2.06	6.93	1.69	45.66	24.46	52.59	24.46	306.81	306.81	306.81	306.81	
YDR508C	GIN1	0.91	1.27	1.35	1.76	1.99	2.08	4.19	1.14	34.84	14.03	39.03	15.17	230.05	230.05	230.05	230.05	
YDR511W	EUD1	0.95	1.11	1.23	1.65	1.86	1.95	5.64	0.98	41.67	10.21	46.51	11.20	226.48	226.48	226.48	226.48	
YDR515W	EUDG	0.93	1.28	1.48	1.84	2.01	2.13	4.59	2.71	44.16	27.48	49.14	30.18	262.16	262.16	262.16	262.16	
YDR519W	YER2	0.98	1.28	1.36	2.06	2.22	2.28	2.78	2.28	32.78	13.77	35.56	32.19	299.17	299.17	299.17	299.17	
YDR520C	URC2	0.94	1.49	1.74	2.16	2.30	2.40	0.83	0.88	2.03	1.46	2.86	2.26	253.94	253.94	253.94	253.94	
YDR521W	YDR521w	0.94	1.04	1.07	1.18	1.40	1.58	36.05	22.71	82.85	82.89	118.98	91.71	319.89	319.89	319.89	319.89	
YDR522C	SPS2	0.96	1.38	1.61	2.06	2.23	2.33	0.94	0.76	8.06	5.96	9.00	6.76	236.81	236.81	236.81	236.81	
YDR523C	SPS1	0.95	1.14	1.13	0.99	1.02	1.07	1.04	0.73	123.89	121.20	126.87	249.46	534.46	534.46	534.46	534.46	
YDR524C	AGE1	0.99	1.33	1.55	1.96	2.17	2.26	1.04	0.73	23.19	10.35	24.23	11.08	286.19	286.19	286.19	286.19	
YDR524C-B	YDR524-b	0.92	1.21	1.35														

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YEL047C	YEL047c	1.03	1.45	1.65	2.08	2.22	2.32	4.67	0.11	38.31	9.98	42.99	10.09	295.94		
YEL048C	TCA17	0.94	1.27	1.46	1.91	2.07	2.17	3.51	0.05	38.94	5.16	42.45	5.20	292.33		
YEL049W	PA02	0.92	1.27	1.48	1.96	2.11	2.22	2.02	0.12	29.31	7.60	31.33	7.73	289.45		
YEL050C	RLM2	0.88	1.12	1.18	1.35	1.53	1.60	19.34	6.00	68.63	44.36	87.97	50.37	313.70		
YEL051W	VMA9	0.87	1.06	1.06	1.05	1.34	1.62	35.61	24.03	67.42	64.62	123.03	108.85	321.67		
YEL052W	AFG1	0.90	1.19	1.37	1.81	2.05	2.15	3.43	0.61	38.60	19.12	42.03	19.73	293.23		
YEL053C	MAK10	0.94	1.23	1.37	1.69	2.05	2.15	5.66	2.62	47.41	33.49	53.07	36.49	295.45		
YEL054C	RPL12a	0.82	1.18	1.20	1.16	1.93	2.13	13.30	3.82	61.87	36.17	75.18	39.99	328.12		
YEL055C	AT52	1.13	1.49	1.50	1.79	1.95	2.08	5.60	1.24	44.61	24.47	52.47	22.47	301.37		
YEL056C	YEL057c	1.03	1.38	1.60	2.06	2.20	2.30	5.60	1.18	41.63	23.33	47.23	24.43	293.45		
YEL059C-A	SOM1	0.90	1.09	1.07	0.96	0.96	0.94	100.25	80.16	131.69	98.63	231.95	178.79	423.68		
YEL059W	YEL059w	0.88	1.16	1.16	1.40	1.61	1.84	70.37	59.19	111.70	101.07	182.07	160.26	465.96		
YEL060C	PBB1	1.03	1.39	1.58	2.01	2.14	2.14	10.26	2.27	56.11	34.54	66.37	36.81	305.20		
YEL061C	CIN8	0.93	1.16	1.29	1.81	2.01	2.16	3.03	0.69	31.09	13.00	34.12	13.09	255.44		
YEL062W	NPR2	0.94	1.18	1.31	1.82	2.00	2.12	7.13	1.02	54.89	23.91	62.02	24.93	279.94		
YEL063C	CAN1	0.91	1.18	1.32	1.81	1.99	2.11	2.83	0.32	37.43	16.55	40.26	16.86	239.29		
YEL064C	AVT2	0.91	1.20	1.39	1.90	2.04	2.16	9.76	0.98	55.80	24.47	65.56	25.46	312.67		
YEL065W	SIT1	0.94	1.24	1.44	1.92	2.10	2.22	3.48	0.29	39.10	16.71	42.58	17.01	300.39		
YEL066W	HPA3	0.94	1.19	1.33	1.88	2.08	2.18	5.97	3.73	48.46	38.80	54.43	42.53	308.79		
YEL067C	YEL067c	0.96	1.27	1.49	1.99	2.14	2.25	7.40	7.77	48.76	46.88	56.17	54.65	304.21		
YEL068C	YEL068c	0.94	1.22	1.41	1.93	2.09	2.18	13.14	8.44	57.31	46.62	70.45	55.06	326.91		
YEL071W	DLD3	0.97	1.27	1.47	1.97	2.13	2.22	10.41	4.73	55.23	38.21	65.64	42.94	312.62		
YEL072W	RMD6	0.96	1.33	1.56	2.06	2.20	2.31	2.87	-0.16	35.70	4.81	38.57	4.64	300.49		
YER001W	MNN1	0.94	1.26	1.48	1.97	2.12	2.22	10.43	0.72	54.12	20.58	64.55	21.30	303.13		
YER002W	NP16	1.00	1.29	1.49	2.03	2.18	2.27	15.72	0.35	61.69	15.21	77.41	15.56	334.49		
YER004W	FMP52	1.00	1.33	1.53	1.99	2.14	2.26	11.52	2.66	55.96	24.84	67.47	26.10	323.02		
YER005W	YND1	0.96	1.26	1.44	1.87	2.04	2.17	16.36	6.23	62.15	45.02	78.50	51.25	328.92		
YER007C-A	TMA20	0.95	1.19	1.38	1.94	2.15	2.27	19.06	3.71	65.13	37.37	84.19	41.08	330.62		
YER007W	PAC2	0.94	1.24	1.46	1.95	2.11	2.22	16.35	7.58	61.45	45.75	77.80	53.33	322.22		
YER007W	PAC2	0.87	1.19	1.32	1.68	1.89	2.02	2.23	2.23	35.93	23.80	38.17	31.06	263.35		
YER007W	PAC2	0.93	1.23	1.33	1.65	1.86	2.01	11.81	3.14	58.86	28.82	70.67	31.96	306.48		
YER010C	YER010c	0.93	1.17	1.38	1.90	2.09	2.21	20.38	12.11	64.05	57.68	84.43	69.79	335.84		
YER011W	TIR1	1.01	1.23	1.46	1.99	2.17	2.28	10.63	12.30	55.50	56.88	66.13	69.19	308.34		
YER014C	BUD25	0.87	1.06	1.02	0.90	0.91	0.87	124.35	120.82	142.21	106.32	266.56	227.14	475.43		
YER014W	HEM14	1.01	1.11	1.12	1.05	1.05	1.04	126.23	120.23	123.81	119.78	250.05	240.01	286.37		
YER015W	FAA2	0.90	1.24	1.39	1.71	1.97	2.14	11.51	3.73	57.15	5.10	40.86	5.95	273.06		
YER016W	BIM1	1.02	1.31	1.51	1.85	2.07	2.25	16.59	3.34	63.39	31.92	79.99	35.26	315.20		
YER016W	YER016c	0.96	1.24	1.41	1.93	2.09	2.18	16.35	7.58	61.45	45.75	77.80	53.33	322.22		
YER017C	AFG3	0.86	1.13	1.18	1.66	1.84	1.94	14.41	8.67	72.52	14.28	121.74	19.53	344.44		
YER018C-A	YER018c-a	0.96	1.24	1.46	1.98	2.13	2.22	20.49	3.22	67.11	35.12	87.51	33.33	321.44		
YER019W	ISC1	0.93	1.20	1.37	1.88	2.05	2.16	17.15	3.43	61.57	31.24	78.72	34.67	329.49		
YER020W	CP42	0.99	1.29	1.50	2.03	2.20	2.23	10.93	2.46	49.49	31.04	59.53	32.59	343.15		
YER024W	YAT2	0.81	1.40	1.65	2.10	2.34	2.33	16.80	7.06	64.26	44.31	81.16	51.36	330.44		
YER024W	GAL83	0.91	1.06	1.24	1.76	1.97	2.14	11.51	3.85	61.14	37.24	72.64	41.09	305.76		
YER028C	MIG2	0.93	1.17	1.37	1.74	1.90	2.07	19.35	0.97	65.18	37.73	84.53	12.72	337.43		
YER030W	CHZ1	1.02	1.33	1.53	2.01	2.17	2.28	26.33	8.95	74.38	53.86	97.71	62.80	356.60		
YER031C	YPT31	0.99	1.59	1.77	2.16	2.29	2.37	4.78	4.78	28.62	41.68	33.40	51.43	284.42		
YER032W	FIR1	0.92	1.19	1.41	1.89	2.06	2.21	19.42	1.55	65.40	28.65	84.82	30.20	338.33		
YER033C	ZRB8	0.91	1.20	1.39	1.88	2.07	2.20	5.80	1.46	41.17	25.89	46.78	27.35	305.99		
YER034W	YER034w	0.92	1.15	1.35	1.82	2.00	2.13	7.06	4.37	50.92	38.88	57.98	43.65	305.37		
YER035W	EDC2	1.00	1.23	1.43	1.95	2.15	2.27	10.33	7.50	56.56	46.63	66.90	54.13	301.22		
YER037W	PHMB	0.92	1.15	1.42	1.89	2.07	2.21	2.46	1.40	30.73	19.01	33.19	20.41	259.32		
YER038W-A	Unknown	1.02	1.38	1.62	2.04	2.19	2.30	20.00	2.43	64.75	31.27	84.75	33.70	338.05		
YER039C	HVG1	0.96	1.27	1.47	1.94	2.09	2.21	19.08	3.66	64.98	35.33	84.06	38.99	344.75		
YER039C-A	YER039c-a	0.97	1.23	1.33	1.79	1.98	2.15	13.38	4.42	57.29	30.15	70.68	34.57	326.19		
YER040W	GLN3	0.98	1.20	1.32	1.79	1.99	2.14	23.38	6.24	71.52	42.36	94.90	48.59	364.21		
YER041W	YEN1	0.97	1.31	1.53	1.96	2.11	2.23	20.19	1.30	65.19	23.32	85.38	24.62	349.36		
YER042W	MXR1	0.91	1.16	1.35	1.83	2.01	2.12	20.33	3.78	67.35	35.03	87.68	38.81	346.05		
YER050C	RHM1	0.99	1.28	1.51	1.97	2.15	2.26	23.06	0.64	71.15	19.74	94.21	20.38	357.20		
YER052W	YER052w	1.02	1.22	1.41	1.71	1.96	2.10	21.05	4.59	68.70	48.75	90.53	55.76	324.46		
YER053C	HGM3	0.90	1.13	1.33	1.72	1.98	2.11	11.69	3.35	54.59	36.20	65.48	32.76	384.38		
YER053C	PIC2	0.99	1.29	1.45	1.90	2.07	2.18	18.63	1.33	61.96	24.32	80.00	22.64	350.23		
YER054C	GP2	0.93	1.26	1.47	1.96	2.12	2.22	18.25	2.15	65.87	28.53	82.19	30.68	343.20		
YER055C	HIS1	0.84	1.08	1.05	0.89	0.95	0.87	18.20	1.03	62.80	12.62	115.68	254.82	513.41		
YER056C	FCY2	0.96	1.29	1.51	1.98	2.13	2.21	32.39	9.68	74.44	53.31	106.83	62.99	376.68		
YER056C-A	YER056a	0.97	1.21	1.38	1.83	1.99	2.09	25.34	6.80	70.27	46.18	95.61	52.98	383.83		
YER057C	HMF1	0.98	1.23	1.40	1.92	2.09	2.21	11.90	0.27	55.62	10.37	67.52	10.65	327.88		
YER059W	PC16	0.92	1.20	1.37	1.82	1.97	2.08	28.49	5.66	75.27	43.79	103.77	49.45	348.36		
YER060W	FCY21	0.92	1.18	1.37												

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE			FRUCTOSE			TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YER111C	SW14	0.97	1.10	1.28	1.74	1.99	2.15	19.13	1.64	70.33	21.30	89.47	22.94	324.26				
YER113C	TMN3	1.02	1.22	1.45	1.96	2.11	2.23	20.33	1.59	71.94	24.03	92.27	25.62	336.19				
YER114C	BO12	0.98	1.22	1.45	1.92	2.12	2.25	14.84	5.64	65.25	45.10	80.09	50.73	322.79				
YER115C	SPR6	1.00	1.23	1.44	1.84	2.06	2.19	13.40	9.36	62.68	55.18	76.29	64.55	306.23				
YER116C	SLX8	0.90	1.07	1.18	1.54	1.79	1.99	28.96	26.58	66.47	64.36	115.43	110.93	322.18				
YER117W	RPL23b	0.89	1.08	1.26	1.71	1.93	2.09	9.87	6.75	59.50	46.62	69.37	53.57	302.76				
YER118C	SHO1	0.98	1.13	1.32	1.73	1.91	2.14	5.64	4.85	51.15	23.85	59.76	25.50	310.44				
YER119C	AVT6	0.88	1.06	1.19	1.61	1.87	2.01	8.61	7.45	59.40	16.12	44.43	17.33	327.15				
YER120C	YER119c-a	0.88	1.07	1.20	1.63	1.83	1.99	17.74	9.53	52.88	49.99	62.81	32.81	329.42				
YER120W	SOS2	0.93	1.11	1.28	1.62	1.83	2.01	2.11	5.89	4.28	47.94	40.03	53.63	44.31	255.00			
YER122C	GE03	0.99	1.11	1.12	1.92	1.93	0.99	125.63	123.51	139.90	108.97	265.53	232.49	344.21				
YER123W	YCK3	0.97	1.17	1.34	1.69	1.98	2.14	12.18	0.92	61.76	13.21	73.92	14.13	336.47				
YER124C	DSE1	0.92	1.08	1.27	1.69	1.93	2.10	20.82	1.26	77.56	19.84	98.38	21.11	339.47				
YER128W	YER128w	1.02	1.16	1.39	1.86	2.09	2.21	18.34	1.45	66.57	23.07	84.91	24.52	328.48				
YER129W	SAK1	1.02	1.30	1.53	1.96	2.14	2.26	20.04	6.66	75.19	44.63	95.23	50.69	329.09				
YER130C	YER130c	0.95	1.18	1.34	1.70	1.95	2.11	5.86	7.93	46.59	51.34	52.45	59.28	301.31				
YER131W	PPS26	0.99	1.17	1.35	1.81	2.07	2.20	13.65	12.34	61.22	57.86	74.87	70.20	323.21				
YER132C	PM1	1.01	1.22	1.44	1.81	2.03	2.15	21.73	14.73	76.23	65.37	97.96	80.10	317.16				
YER134C	YER134c	0.93	1.23	1.47	1.95	2.12	2.25	16.24	9.58	64.99	52.36	81.22	61.94	330.68				
YER135C	YER135c	1.00	1.25	1.44	1.85	2.03	2.15	18.59	13.75	68.23	59.51	86.82	73.26	336.23				
YER137C	YER137c	0.97	1.10	1.24	1.66	1.89	2.09	15.32	13.03	66.33	60.64	81.65	73.67	326.18				
YER139C	RT1	0.91	1.05	1.14	1.58	1.86	2.10	21.86	15.36	76.50	65.61	98.36	80.97	320.52				
YER140W	YER140w	0.90	1.10	1.23	1.66	1.88	2.00	24.78	11.63	76.64	60.64	101.42	72.27	343.17				
YER141W	COK15	0.97	1.13	1.27	1.63	1.94	2.11	19.74	1.59	71.14	25.61	90.88	27.20	333.49				
YER142C	MAG1	0.96	1.16	1.40	1.85	2.06	2.18	16.22	1.10	66.82	17.77	83.04	18.87	332.84				
YER143W	DD11	1.04	1.19	1.41	1.88	2.10	2.24	17.23	3.01	68.20	38.63	85.43	41.64	333.40				
YER144C	UBP5	0.95	1.33	1.38	1.84	2.00	2.12	17.59	7.67	68.91	47.69	86.50	55.36	333.18				
YER145C	FTR1	0.97	1.17	1.36	1.73	1.99	2.14	5.91	1.64	50.94	28.28	56.84	29.92	312.17				
YER149C	PEA2	0.98	1.30	1.53	1.87	2.08	2.21	0.72	0.74	9.36	8.29	10.08	9.03	271.67				
YER150W	SP11	0.94	1.23	1.44	1.86	2.06	2.19	4.82	4.23	42.25	39.58	47.07	43.81	291.21				
YER151C	UBP3	0.93	1.09	1.22	1.67	1.94	2.17	1.58	2.00	24.95	28.90	26.53	30.90	276.10				
YER152C	YER152c	0.93	1.14	1.33	1.76	1.97	2.13	6.73	5.55	51.50	45.63	58.23	51.18	302.25				
YER153C	PET122	0.97	1.02	1.03	0.96	0.96	0.92	135.67	136.96	133.66	134.58	269.34	271.54	321.64				
YER154W	OXA1	0.98	1.02	1.03	0.95	0.95	0.92	132.20	141.30	127.69	138.06	260.89	278.36	311.35				
YER155C	BEM2	0.99	1.16	1.30	1.74	1.95	2.16	5.60	4.20	18.49	18.67	18.97	13.19	290.02				
YER156C	BEM2	0.89	1.19	1.24	1.59	1.88	2.18	0.49	1.32	18.49	11.67	21.28	21.60	212.79				
YER158C	YER158c	0.88	1.08	1.20	1.58	1.83	2.03	3.29	1.25	23.29	19.28	36.74	21.60	335.37				
YER159C	YER159c	0.99	1.18	1.38	1.94	2.06	2.18	16.03	1.75	63.10	17.62	79.12	19.68	337.11				
YER161C	SP72	0.91	1.20	1.35	1.78	2.05	2.26	25.85	20.88	78.19	70.23	103.24	99.32	369.51				
YER163C	RAD4	0.93	1.16	1.38	1.79	1.97	2.11	7.49	2.23	54.05	23.57	61.54	34.79	307.24				
YER163C	YER163c	0.94	1.29	1.53	1.97	2.16	2.29	7.51	2.83	53.59	36.28	61.10	38.10	303.35				
YER164W	CHD1	0.99	1.16	1.31	1.72	1.96	2.14	16.04	10.45	68.77	59.71	84.81	70.16	311.40				
YER166W	DD11	0.89	1.17	1.31	1.67	1.87	2.07	7.02	5.40	54.03	46.79	61.04	52.19	296.48				
YER167W	BCK2	1.00	1.20	1.35	1.69	1.85	1.94	42.85	35.51	91.11	82.11	133.96	117.63	369.69				
YER169W	RPH1	0.93	1.05	1.06	0.98	0.99	0.96	149.34	147.33	143.82	144.09	293.16	291.41	308.26				
YER170W	ADK2	1.00	1.20	1.35	1.84	2.05	2.20	15.46	11.86	57.55	52.75	80.90	69.61	323.45				
YER173W	RAD24	0.97	1.10	1.25	1.64	1.86	2.01	12.93	7.27	65.00	49.12	77.93	56.39	322.39				
YER174C	GRX4	0.95	1.16	1.37	1.79	1.97	2.11	7.49	2.23	54.05	23.57	61.54	34.79	327.26				
YER175C	MTT1	0.90	1.07	1.19	1.58	1.80	1.93	14.34	1.30	56.37	19.49	70.70	20.79	285.43				
YER175W-A	YER175w-a	0.88	1.12	1.25	1.60	1.80	1.94	14.47	2.79	63.13	24.09	77.60	26.88	322.35				
YER176W	ECM32	0.90	1.10	1.24	1.65	1.86	2.00	9.66	2.43	49.34	26.47	59.00	28.90	304.34				
YER177W	BMH1	0.85	1.00	1.06	1.29	1.49	1.67	1.75	0.79	28.12	3.78	29.87	4.57	297.58				
YER178W	PD1A	0.93	1.15	1.31	1.73	1.93	2.07	10.11	0.94	48.57	9.36	58.67	10.30	302.73				
YER179W	DMC1	0.85	1.05	1.11	1.43	1.63	1.87	0.71	0.80	13.82	1.96	14.53	2.76	265.91				
YER180C	ISC10	0.98	1.24	1.43	1.84	2.04	2.20	11.89	1.94	53.87	23.34	65.76	25.28	287.27				
YER180W	YER180w	0.94	1.21	1.31	1.88	2.03	2.20	1.67	1.25	22.97	2.38	24.63	3.63	284.64				
YFL011W	HXT10	0.93	1.33	1.50	1.96	2.04	2.19	1.67	1.25	26.55	4.78	30.50	5.619	287.53				
YFL012W	YFL012w	0.94	1.18	1.38	1.77	1.94	2.10	7.62	1.83	47.58	47.64	55.20	25.46	303.09				
YFL013C	IEST1	0.93	1.21	1.42	1.75	1.94	2.10	0.79	0.74	6.44	1.24	7.23	1.97	242.86				
YFL023W	Unknown	0.95	1.28	1.53	1.90	2.09	2.22	1.21	3.84	16.08	31.34	17.29	35.18	272.38				
YFL014W	HSP12	0.98	1.27	1.50	1.87	2.06	2.20	1.00	2.13	8.29	23.30	9.29	25.43	218.39				
YFL015C	WMM1	0.97	1.31	1.55	1.91	2.11	2.32	0.79	0.78	4.59	0.57	18.06	2.08	280.83				
YFL016C	AU1A	0.99	1.13	1.31	1.68	1.89	2.03	1.35	1.07	6.69	16.16	26.64	2.39	346.40				
YFL016C	AU1A	0.96	1.21	1.41	1.78	1.95	2.07	10.62	8.87	49.33	42.32	59.95	51.19	326.46				
YFL017W	HAC1	0.91	1.28	1.45	1.86	2.06	2.18	9.34	5.47	42.								

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YFR018C	YFR018c	0.98	1.45	1.64	2.00	2.15	2.25	10.57	10.49	50.20	47.44	60.77	57.93	321.78		
YFR019W	FAB1	0.92	1.12	1.30	1.64	1.80	1.91	14.25	12.61	63.69	58.55	77.94	71.17	296.95		
YFR020W	YFR020w	1.03	1.58	1.79	2.09	2.27	2.24	3.45	3.91	21.07	24.07	24.51	27.97	290.13		
YFR021W	ATG18	0.96	1.39	1.57	1.91	2.05	2.19	17.61	16.06	61.12	54.35	78.73	70.41	327.17		
YFR022W	ROG3	1.03	1.55	1.77	2.10	2.24	2.31	9.35	10.42	48.01	49.83	57.36	60.24	313.33		
YFR023W	PES4	0.92	1.26	1.41	1.74	1.98	2.12	7.14	8.05	39.76	40.01	46.90	48.06	310.32		
YFR024W	YFR024c	0.98	1.14	1.35	1.81	1.97	2.08	7.35	1.59	48.68	27.00	56.02	28.98	287.31		
YFR024C-A	LSD1	0.88	1.27	1.38	1.76	1.95	2.03	13.64	9.67	53.1	44.77	68.61	54.84	321.42		
YFR025C	YFR025c	0.98	1.23	1.38	1.49	1.64	1.81	7.85	8.49	51.49	52.89	52.84	53.13	313.33		
YFR026C	ULF1	0.88	1.23	1.38	1.79	2.02	2.14	9.26	10.44	44.65	11.50	53.91	14.60	344.44		
YFR030W	MET10	0.93	1.15	1.38	1.85	2.08	2.24	0.99	0.67	17.11	2.25	18.09	2.92	298.89		
YFR031C-A	RPL2a	1.00	1.39	1.54	1.96	2.14	2.24	16.78	10.81	57.12	48.21	73.89	58.22	358.73		
YFR032C	RPL5	1.00	1.32	1.47	1.89	2.04	2.20	13.49	3.75	61.05	23.25	74.54	27.09	329.05		
YFR032C-B	RPL29	0.99	1.28	1.39	1.79	1.96	2.15	16.73	3.74	66.29	23.78	83.01	27.52	331.35		
YFR032C-B	YFR032c-b	0.85	1.17	1.28	1.62	1.85	2.00	9.66	1.73	54.96	21.52	64.63	23.25	305.33		
YFR033C	QCR6	1.02	1.44	1.66	2.06	2.20	2.31	6.91	3.20	44.86	8.21	51.76	11.41	297.90		
YFR034C	PHO4	0.95	1.19	1.30	1.82	2.00	2.18	9.90	3.50	51.07	19.03	60.97	22.53	308.99		
YFR035C	YFR035c	1.01	1.39	1.52	1.92	2.08	2.25	4.25	3.18	28.85	9.35	33.09	12.52	294.35		
YFR036W	CDC26	1.05	1.39	1.51	1.93	2.07	2.24	9.70	3.77	53.40	21.17	63.10	24.93	311.55		
YFR038W	IRC5	1.09	1.46	1.61	2.04	2.19	2.31	14.87	4.07	59.39	25.68	74.26	29.75	330.77		
YFR038W	IRC5	0.92	1.29	1.46	1.86	2.05	2.19	3.80	1.83	37.97	18.05	41.77	19.88	269.43		
YFR039C	YFR039c	0.86	1.01	1.19	1.63	1.86	2.04	2.75	0.92	35.22	15.34	37.97	16.26	291.12		
YFR040W	SAP155	0.99	1.30	1.41	1.88	2.01	2.18	11.11	5.93	52.32	40.06	63.44	46.00	307.90		
YFR041C	ERJ5	0.97	1.24	1.37	1.87	2.04	2.21	15.79	5.57	60.24	35.53	76.03	41.10	328.38		
YFR043C	IRC6	1.08	1.54	1.64	2.04	2.16	2.27	6.50	3.21	40.10	12.92	46.60	16.13	300.17		
YFR044C	DUG1	0.96	1.20	1.30	1.74	1.94	2.15	6.70	2.85	42.09	10.50	48.79	13.35	305.18		
YFR045W	YFR045w	1.00	1.31	1.49	1.92	2.09	2.26	3.34	2.92	16.63	3.70	19.97	6.62	296.40		
YFR045W	YFR045w	0.91	1.27	1.49	1.89	2.05	2.20	6.13	2.04	45.41	18.70	51.54	20.75	293.67		
YFR046C	CNN1	1.00	1.22	1.25	1.66	1.91	2.15	29.49	11.25	84.85	61.32	114.34	72.57	353.43		
YFR047C	BNA6	0.94	1.23	1.36	1.79	1.95	2.13	10.60	3.08	55.41	14.03	66.02	17.11	320.25		
YFR048W	RMD8	1.04	1.45	1.68	2.07	2.18	2.30	20.09	4.54	66.19	30.59	86.27	35.13	309.63		
YFR049W	YMR31	0.92	1.14	1.26	1.79	2.00	2.21	3.34	3.08	12.44	2.46	15.78	5.54	287.00		
YFR053C	HXK1	0.96	1.29	1.44	1.87	2.02	2.19	6.84	3.54	72.43	60.75	79.27	64.29	322.69		
YFR054C	YFR054c	0.97	1.30	1.46	1.91	2.05	2.22	9.15	3.43	48.72	16.12	57.86	19.55	311.48		
YFR055W	IRC7	0.97	1.25	1.37	1.79	1.96	2.15	9.22	3.47	49.14	18.93	58.37	22.40	316.05		
YFR056C	YFR056c	1.04	1.35	1.48	1.94	2.09	2.25	11.00	4.47	51.28	29.57	62.27	34.04	318.42		
YFR057W	YFR057w	0.95	1.22	1.32	1.81	2.00	2.19	10.13	3.49	51.10	31.56	61.23	35.95	315.20		
YGL002W	ERP5	1.00	1.56	1.72	2.06	2.26	2.36	1.68	2.04	5.48	4.16	36.37	4.23	307.44		
YGL002W	ERP5	0.97	1.35	1.54	1.73	1.92	2.08	26.71	15.55	73.81	54.82	97.07	75.52	339.44		
YGL003C	YGL003c	0.94	1.23	1.33	1.73	1.92	2.08	1.68	1.26	4.25	2.33	32.51	2.51	307.44		
YGL007W	YGL007w	0.91	1.19	1.30	1.50	1.59	1.68	48.82	40.69	88.85	82.62	137.37	123.31	411.90		
YGL010W	YGL010w	0.93	1.22	1.38	1.76	1.95	2.02	6.91	4.09	40.31	25.22	47.22	29.31	303.44		
YGL012W	ERG4	0.93	1.36	1.49	1.84	2.00	2.09	8.42	1.84	52.37	16.09	60.79	17.73	337.35		
YGL012W	ERG4	0.83	1.06	1.18	1.31	1.60	1.71	24.82	6.96	72.35	42.17	97.17	49.13	360.63		
YGL013C	PDR1	0.97	1.25	1.39	1.81	1.99	2.14	6.37	3.24	39.14	24.43	45.51	24.99	302.27		
YGL014W	PUF4	0.92	1.16	1.25	1.64	1.83	1.98	10.46	4.16	51.05	28.66	61.51	32.82	308.28		
YGL015C	YGL015c	0.92	1.06	1.01	1.10	1.40	1.55	44.26	51.30	88.87	90.24	133.12	141.54	381.37		
YGL016W	KAP122	0.93	1.29	1.38	1.77	1.96	2.12	5.19	3.09	20.33	9.18	25.53	12.28	274.54		
YGL017W	ATE1	0.95	1.34	1.47	1.88	1.99	2.12	11.25	6.71	49.12	48.81	60.37	45.32	298.95		
YGL019W	CKB1	1.01	1.30	1.37	1.76	1.91	2.07	20.60	13.11	63.71	52.35	84.31	65.46	327.97		
YGL020C	GET1	0.98	1.20	1.37	1.76	2.00	2.19	1.03	0.72	1.74	3.97	2.77	1.11	294.10		
YGL021W	ALK1	0.97	1.35	1.50	1.88	2.04	2.18	5.80	3.42	36.42	17.28	42.22	20.70	296.75		
YGL023C	PIB2	0.90	1.20	1.40	1.74	1.96	2.15	1.15	0.79	5.39	6.53	6.55	2.42	279.16		
YGL023C	PIB2	1.02	1.42	1.56	1.89	2.04	2.17	23.48	16.47	67.61	58.64	91.09	75.11	350.83		
YGL024W	YGL024w	0.97	1.34	1.47	1.81	1.98	2.13	20.38	13.47	63.73	50.24	84.10	63.71	347.63		
YGL025C	PGD1	0.89	1.15	1.23	1.56	1.74	1.92	5.78	3.41	37.97	19.68	43.75	23.09	301.48		
YGL026C	TRP5	0.95	1.43	1.60	1.90	2.07	2.17	2.50	1.12	31.64	5.78	34.14	6.90	271.60		
YGL026C	TRP5	0.82	0.97	0.95	0.79	0.91	1.21	16.08	5.18	69.75	40.61	85.84	45.80	316.83		
YGL027C	CWH41	0.91	1.16	1.24	1.46	1.88	2.06	2.06	2.11	32.11	21.21	42.72	18.08	276.72		
YGL028W	SCW11	0.91	1.16	1.24	1.60	1.79	1.94	11.79	4.86	52.69	31.03	64.48	35.89	318.91		
YGL029W	YGL029w	0.93	1.05	1.06	1.86	2.01	2.11	11.80	11.56	118.99	116.56	236.49	232.12	517.53		
YGL031C	CHC1	0.86	1.33	1.48	1.86	2.01	2.11	27.36	13.52	75.23	58.14	102.59	71.65	360.45		
YGL041C	YGL041c	0.91	1.33	1.43	1.83	2.00	2.12	13.85	8.22	58.38	44.99	72.23	53.21	335.47		
YGL041C-B	YGL041c-b	0.91	1.15	1.23	1.51	1.81	2.04	26.77	14.16	83.67	50.99	110.44	65.15	329.02		
YGL042C	YGL042c	0.92	1.32	1.40	1.80	1.98	2.09	36.40	30.93	93.35	79.37	119.92	110.30	375.47		
YGL043W	YST1	0.99	1.40	1.43	1.80	2.00	2.14	4.21	3.80	31.52	12.69	35.73	16.49	311.67		
YGL045W	RIM8	0.97	1.52	1.55	2.02	2.16	2.29	10.39	3.76	51.05	27.25	61.44	31.01	339.04		
YGL046W	RIM8	0.92	1.29	1.32	1.68	1.86	1.98	8.11	3.37	52.66	20.30	62.34	23.67	317.56		
YGL046W	RIM8	0.87	1.32	1.40	1.7											

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YGL117W	YGL117w	0.91	1.20	1.35	1.73	1.96	2.07	3.10	0.77	34.50	8.56	37.60	9.33	286.46		
YGL118C	YGL118c	0.91	1.25	1.43	1.86	2.08	2.17	2.92	0.85	33.68	11.10	36.60	11.95	284.11		
YGL119W	ABC1	0.97	1.11	1.12	0.99	0.98	0.98	123.82	120.93	138.49	104.25	262.30	225.17	496.00		
YGL121C	GPG1	1.01	1.35	1.58	2.02	2.19	2.27	4.38	0.94	39.73	14.68	44.11	15.62	288.09		
YGL124C	MONT	0.96	1.27	1.43	1.79	1.97	2.02	21.72	18.04	60.73	76.16	102.45	94.20	330.21		
YGL125W		0.90	1.11	1.18	1.56	1.92	2.11	0.93	0.70	17.18	4.71	18.11	5.41	294.26		
YGL126W	MET13	0.95	1.18	1.24	1.49	1.83	2.06	15.89	2.14	70.29	28.57	86.18	30.17	322.16		
YGL126G	SCS2	0.88	1.14	1.30	1.75	2.03	2.09	6.24	3.48	45.98	35.16	52.22	38.64	296.25		
YGL127W	YGL127w	0.96	1.18	1.24	1.46	1.74	2.00	16.21	2.06	92.46	12.24	128.27	33.14	334.14		
YGL128W	RSM23	0.97	1.04	1.08	0.98	1.01	0.98	133.89	145.14	128.93	138.82	382.83	232.57	298.89		
YGL131C	SNT2	0.92	1.10	1.28	1.74	1.99	2.13	18.79	10.87	71.48	54.87	90.25	62.74	316.16		
YGL132W	YGL132w	0.93	1.16	1.38	1.90	2.16	2.33	0.80	0.79	5.12	1.83	5.82	2.62	249.84		
YGL133W	ITC1	0.91	1.11	1.33	1.79	1.99	2.15	22.67	12.00	80.70	65.79	103.37	77.79	323.37		
YGL134W	PCL10	0.87	1.06	1.17	1.54	1.83	2.03	21.79	6.53	76.55	37.84	98.34	44.38	341.73		
YGL135W	RPL1b	0.97	1.32	1.61	2.04	2.21	2.33	0.83	0.81	8.18	6.67	9.01	7.27	227.45		
YGL136C	MRF2	0.90	1.09	1.21	1.62	1.86	2.06	11.78	8.74	56.99	48.45	68.76	57.29	269.61		
YGL138C	YGL138c	0.94	1.18	1.38	1.82	2.03	2.19	4.68	2.12	41.46	30.10	46.13	32.21	280.03		
YGL139W	FLC3	0.90	1.11	1.30	1.79	1.97	2.10	19.26	13.51	68.17	59.54	87.44	73.05	329.30		
YGL140C	YGL140c	0.92	1.15	1.35	2.00	1.93	2.05	19.30	8.20	68.39	46.70	87.68	54.91	329.75		
YGL141W	HUL5	0.91	1.08	1.24	1.70	1.92	2.03	28.81	8.97	81.17	51.34	109.98	60.32	331.60		
YGL143C	MRF1	0.91	1.00	1.01	0.95	0.95	0.91	136.80	143.42	132.90	140.43	269.70	283.85	250.21		
YGL144C	ROG1	0.88	1.06	1.23	1.75	1.97	2.08	11.53	1.81	56.48	30.65	68.01	32.46	310.54		
YGL146C	RRT6	0.96	1.17	1.37	1.80	2.05	2.14	18.57	7.85	70.38	51.10	88.94	58.95	323.84		
YGL147C	RPL5a	0.94	1.09	1.23	1.67	1.96	2.12	22.82	4.30	73.38	38.31	96.20	42.61	332.46		
YGL148W	ARO2	0.95	1.01	1.04	1.00	1.03	1.03	137.13	155.22	135.10	150.31	272.23	305.53	235.10		
YGL149W	YGL149w	0.93	1.14	1.39	1.80	2.03	2.15	12.52	3.24	62.14	31.68	74.66	34.92	314.00		
YGL151W	NUT1	0.92	1.06	1.21	1.75	1.99	2.12	27.64	19.22	78.90	71.60	106.54	90.82	323.68		
YGL152C	YGL152c	0.91	1.22	1.47	1.93	2.12	2.22	15.31	15.43	60.99	60.74	76.30	76.17	301.99		
YGL153W	PEX14	0.90	1.17	1.38	1.82	2.02	2.15	16.88	9.71	69.42	51.25	86.30	60.95	290.27		
YGL154C	LYSS	0.91	1.15	1.34	1.77	1.95	2.11	16.11	6.93	63.97	32.12	80.08	52.05	332.51		
YGL155W	AMS1	0.91	1.11	1.28	1.71	1.91	2.06	21.98	11.90	71.53	57.41	93.51	69.31	336.16		
YGL157W	ARI1	0.93	1.15	1.34	1.76	1.97	2.09	19.15	6.69	66.50	44.36	85.64	51.06	338.30		
YGL158W	RCK1	0.88	1.04	1.20	1.62	1.85	1.98	13.58	8.86	61.69	48.33	75.27	57.19	323.59		
YGL159W	YGL159w	0.88	1.06	1.22	1.71	1.89	2.01	13.89	8.09	62.24	50.10	76.13	58.20	318.50		
YGL160W	AMT4	0.92	1.04	1.22	1.60	1.87	2.03	24.02	1.43	78.01	22.21	102.03	23.84	331.10		
YGL161C	YIP5	0.92	1.16	1.39	1.87	2.07	2.21	21.39	1.01	69.02	47.47	90.41	15.47	355.26		
YGL162W	UTT1	0.93	1.15	1.35	1.76	1.96	2.10	19.34	5.18	73.13	44.84	92.47	49.94	327.61		
YGL163C	RAT44	0.90	1.10	1.24	1.60	1.85	2.03	31.95	14.33	68.65	42.42	124.54	80.27	344.39		
YGL164W	YGL164w	0.91	1.12	1.31	1.76	1.97	2.08	12.68	9.61	59.82	52.22	72.68	61.89	302.66		
YGL165C	YGL165c	0.92	1.14	1.34	1.64	1.84	2.01	39.23	30.44	92.10	80.31	131.33	120.75	342.42		
YGL166W	CUP2	0.93	1.27	1.52	1.88	1.99	2.09	31.77	23.89	67.08	50.50	118.85	104.31	339.56		
YGL167C	PMH1	0.89	0.99	1.19	1.57	1.81	1.99	19.95	9.74	68.32	53.81	87.38	63.58	314.46		
YGL168W	HUR7	0.94	1.01	1.11	1.51	1.80	2.00	20.79	10.88	72.18	60.32	92.97	71.30	317.10		
YGL170C	SPOT4	0.94	1.13	1.30	1.74	1.93	2.06	20.04	17.08	71.21	66.24	91.24	83.33	325.18		
YGL173C	KEM1	0.92	0.99	1.02	1.09	1.30	1.53	39.27	27.27	75.71	128.33	102.97	293.88			
YGL174W	BUD13	0.92	1.05	1.16	1.62	1.90	2.05	11.53	10.60	58.77	54.84	70.30	65.44	300.70		
YGL175C	SAE2	0.94	1.10	1.25	1.65	1.90	2.04	18.24	1.21	61.81	21.27	88.05	22.48	335.86		
YGL176C	YGL176c	0.91	1.05	1.24	1.69	1.92	2.09	28.89	1.54	82.85	24.43	111.84	25.96	352.35		
YGL177W	YGL177w	0.92	1.08	1.28	1.71	1.93	2.14	11.25	0.90	55.65	5.54	66.90	6.44	332.02		
YGL178W	MPTS	0.95	1.17	1.31	1.70	1.99	2.19	6.88	2.27	48.21	22.59	55.09	24.86	293.14		
YGL179C	TOS3	0.91	1.11	1.33	1.76	1.97	2.12	12.60	6.19	63.80	45.59	76.40	52.78	320.45		
YGL180W	ATG1	0.93	1.13	1.32	1.78	2.03	2.17	16.59	15.43	66.69	61.78	83.29	77.21	302.55		
YGL181W	GTS1	0.97	1.22	1.44	1.83	2.06	2.19	15.45	12.88	66.82	57.34	82.27	70.22	316.81		
YGL184C	STR3	0.96	1.24	1.50	1.98	2.17	2.30	3.92	4.60	41.64	32.36	45.55	36.95	279.99		
YGL185C	YGL185c	0.89	1.15	1.37	1.82	2.07	2.24	1.25	1.24	13.62	14.14	14.87	15.38	257.08		
YGL185C	YGL185c-a	0.92	1.13	1.20	1.43	1.67	1.87	2.05	2.21	54.09	30.54	122.22	80.43	314.86		
YGL186C	SIP2	1.02	1.36	1.47	1.89	2.06	2.21	9.85	4.94	55.91	43.40	65.76	48.34	302.42		
YGL189W	MIG2	0.98	1.38	1.56	2.03	2.20	2.35	0.89	0.51	7.95	9.37	8.84	10.49	269.67		
YGL201W	YPT32	0.89	1.16	1.27	1.67	1.87	2.05	1.00	1.69	22.60	21.75	23.60	23.44	289.32		
YGL211W	NCS6	0.91	1.15	1.26	1.76	2.02	2.24	1.21	2.20	19.93	29.50	21.14	31.70	275.52		
YGL212W	NCS6	0.98	1.32	1.49	1.89	2.05	2.19	4.47	1.28	38.55	38.81	7.32	8.30	278.31		
YGL213C	VAM7	0.89	1.10	1.17	1.45	1.60	1.83	0.72	1.01	28.88	9.23	30.55	23.15	302.26		
YGL213C	SK8	1.01	1.26	1.36	1.87	2.08	2.24	24.73	10.36	74.73	53.08	99.46	63.44	317.67		
YGL214W	YGL214w	1.07	1.33	1.44	1.91	2.11	2.29	28.00	22.42	80.10	76.45	108.10	98.88	321.43		
YGL214W	YGL214w	0.93	1.29	1.48	1.89	2.07	2.20	12.82	8.24	49.89	39.08	62.71	47.32	308.25		
YGL215W	CLG1	1.03	1.26	1.37	1.85	2.05	2.20	15.39	6.22	64.23	46.22	79.62	52.44	314.48		
YGL216W	KIP3	0.99	1.20	1.31	1.87	2.08	2.27	14.39	6.90	64.89	47.24	79.28	54.14	288.72		
YGL216W	KIP3	0.94	1.22	1.33	1.74	1.95	2.10	5.16	9.54	46.32	43.36	51.48	52.90	269.07		
YGL217C	YGL217c	0.94	1.13													

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YGL261C	PAU11	0.96	1.25	1.38	1.79	1.97	2.14	12.19	4.59	60.35	40.01	72.54	44.60	315.26		
YGL262W	YGL262w	1.05	1.34	1.52	1.96	2.12	2.28	8.78	3.84	53.83	39.58	62.61	43.42	305.11		
YGL263W	COST12	1.10	1.48	1.69	2.00	2.15	2.31	1.49	1.10	19.35	12.53	20.84	13.63	283.30		
YGR0001C	YGR001c	1.15	1.52	1.71	2.12	2.24	2.35	15.34	6.04	73.64	44.74	88.98	50.78	322.76		
YGR003W	CUL3	1.10	1.54	1.75	2.12	2.24	2.35	11.85	4.72	62.12	38.30	73.97	43.02	316.09		
YGR004W	PEX1	1.05	1.41	1.64	2.01	2.22	2.34	10.01	2.43	55.45	27.90	75.46	30.33	279.48		
YGR006W	PRF1B	0.93	1.08	1.08	1.01	1.13	1.13	10.46	1.13	12.56	12.50	23.62	215.75	476.41		
YGR007W	YGR007	1.08	1.47	1.58	2.08	2.28	2.42	6.81	2.88	50.05	35.05	56.96	38.03	282.24		
YGR008W	S21	1.03	1.49	1.60	1.65	1.89	2.08	1.20	0.97	16.29	12.03	12.19	8.51	320.00		
YGR010W	NM423	1.06	1.32	1.40	1.90	2.10	2.19	11.37	3.63	67.85	34.42	79.22	37.45	307.75		
YGR011W	YGR011w	1.05	1.32	1.44	1.89	2.09	2.28	1.20	0.97	10.00	3.68	11.20	4.65	284.43		
YGR011W	YGR011w	0.94	1.33	1.48	1.79	1.95	2.05	15.01	5.37	53.67	45.10	68.68	56.32	362.30		
YGR012W	YGR012w	1.01	1.24	1.36	1.82	2.04	2.25	1.27	0.97	13.62	4.50	14.89	5.47	280.07		
YGR014W	MSB2	0.98	1.24	1.32	1.68	1.85	2.05	8.26	2.22	55.06	25.10	87.00	63.33	31.07	310.42	
YGR015C	YGR015c	1.10	1.40	1.58	2.07	2.22	2.37	6.04	2.07	50.17	26.41	56.20	28.48	284.71		
YGR016W	YGR016w	1.15	1.41	1.54	1.94	2.14	2.32	2.00	1.42	27.21	19.33	29.21	20.75	295.37		
YGR017W	YGR017w	1.05	1.36	1.52	1.94	2.10	2.27	4.88	4.04	41.38	37.27	46.26	41.31	285.89		
YGR018C	YGR018c	1.04	1.31	1.47	1.81	1.97	2.16	15.68	7.24	67.65	51.01	83.33	58.26	285.29		
YGR018C	YGR018c	1.04	1.58	1.79	2.13	2.27	2.34	3.32	3.30	17.79	13.93	21.10	17.23	295.54		
YGR019W	UGA1	0.97	1.16	1.24	1.68	1.88	2.10	12.56	3.38	64.21	37.94	76.77	41.32	298.41		
YGR020C	VMA7	1.11	1.35	1.31	1.54	1.76	2.01	60.53	44.06	115.38	100.89	175.91	144.95	341.38		
YGR020C	VMA7	0.86	1.17	1.17	1.15	1.45	1.71	50.87	25.10	112.36	87.00	163.23	112.10	337.03		
YGR021W	YGR021w	1.05	1.41	1.53	2.00	2.21	2.38	19.48	7.00	71.76	51.26	91.23	58.25	328.01		
YGR022C	YGR022c	1.05	1.23	1.28	1.66	1.87	2.07	29.52	21.94	87.01	84.07	116.54	106.01	335.83		
YGR022C	YGR022c	0.94	1.37	1.55	1.90	2.06	2.17	5.31	3.81	31.83	21.94	37.13	25.75	298.57		
YGR023W	MTL1	1.05	1.31	1.45	1.90	2.10	2.24	10.05	1.83	51.83	24.57	61.88	26.40	314.83		
YGR025W	YGR025w	1.00	1.25	1.40	1.88	2.05	2.22	5.66	2.20	47.41	30.30	53.06	32.50	295.69		
YGR025W	YGR025w	0.94	1.37	1.55	1.91	2.05	2.18	8.96	4.58	45.23	28.91	54.20	33.49	307.20		
YGR026W	YGR026w	1.03	1.41	1.65	2.10	2.26	2.41	0.99	0.88	11.14	6.25	12.13	7.13	282.83		
YGR027C	RPS25a	1.01	1.31	1.45	1.92	2.12	2.20	8.26	1.91	58.07	29.57	66.32	31.48	339.26		
YGR028W	MSP1	0.91	1.08	1.29	1.72	1.93	2.11	7.50	1.99	50.06	27.18	57.56	29.17	283.50		
YGR031W	YGR031w	0.98	1.36	1.59	2.06	2.21	2.29	0.96	0.87	17.31	4.67	18.27	5.54	273.10		
YGR032W	SGC2	1.00	1.30	1.57	1.99	2.15	2.28	11.17	2.32	56.71	28.93	67.88	31.24	322.23		
YGR033C	TM21	1.00	1.41	1.63	2.08	2.24	2.31	0.80	1.22	12.65	4.24	13.44	5.16	277.63		
YGR040W	RPL26b	0.94	1.22	1.35	1.79	2.04	2.20	0.57	0.88	6.01	0.22	1.18	1.10	260.61		
YGR045C	YGR045c	0.93	1.21	1.41	1.89	2.09	2.17	1.05	1.07	19.66	8.44	20.71	9.40	279.20		
YGR055W-A	YGR055w-a	0.96	1.20	1.38	1.74	2.04	2.19	8.14	8.42	52.66	14.85	60.79	15.87	300.80		
YGR068C	CAP1	1.06	1.37	1.56	1.74	2.04	2.23	16.89	4.53	11.58	4.20	12.70	8.84	47.44	298.48	
YGR069C	AGD1	1.03	1.30	1.56	2.10	2.28	2.37	0.58	0.65	8.07	3.45	9.45	6.10	254.46		
YGR070C	CBP1	0.96	1.05	1.09	1.44	1.35	1.48	39.96	24.14	86.12	57.47	121.26	116.11	307.45		
YGR070W	ORM1	0.94	1.15	1.36	1.84	2.03	2.18	11.05	1.88	57.78	26.45	68.82	28.25	329.21		
YGR030W	YGR030w	0.98	1.26	1.44	1.91	2.09	2.18	13.62	6.83	63.67	48.80	77.29	55.63	343.44		
YGR040W	KSS1	0.92	1.16	1.41	1.84	2.02	2.17	4.85	3.36	43.01	35.05	47.86	38.41	306.46		
YGR041W	BUD9	0.98	1.35	1.58	2.04	2.21	2.29	7.96	2.65	51.19	30.81	59.14	32.66	320.63		
YGR042W	YGR042w	0.99	1.39	1.62	2.07	2.22	2.30	5.93	3.35	47.08	34.72	53.01	38.08	323.75		
YGR043C	NOM1	0.98	1.28	1.44	1.91	2.12	2.22	3.37	1.17	38.17	18.97	41.54	20.13	316.74		
YGR044C	RME1	0.94	1.28	1.45	1.87	2.05	2.14	2.02	0.81	29.02	4.29	31.04	5.10	313.47		
YGR045C	YGR045c	1.06	1.44	1.67	2.16	2.32	2.39	1.74	1.00	27.30	9.23	29.04	10.22	308.56		
YGR049W	SCM4	0.99	1.37	1.61	2.08	2.23	2.29	1.21	1.03	17.56	7.93	18.77	8.96	285.90		
YGR050C	YGR050c	0.90	1.05	1.26	1.70	1.92	2.12	1.08	0.83	14.44	7.41	15.52	8.25	270.41		
YGR051C	YGR051c	1.02	1.43	1.63	2.02	2.22	2.32	2.30	0.70	6.54	0.25	8.84	0.95	263.35		
YGR052W	FMP48	0.99	1.33	1.53	1.99	2.19	2.28	1.11	1.10	8.01	4.01	9.11	5.11	269.95		
YGR053C	YGR053c	0.91	1.07	1.27	1.73	1.95	2.13	4.64	5.50	42.09	27.60	68.93	11.52	23.08	303.63	
YGR054W	YGR054w	0.93	1.16	1.31	1.73	1.96	2.04	1.63	0.94	26.91	14.60	28.54	15.54	300.60		
YGR055W	MUP1	0.98	1.15	1.31	1.87	2.09	2.18	3.64	1.58	37.30	21.95	40.94	23.54	292.34		
YGR056W	RSC1	1.02	1.22	1.32	1.78	2.00	2.13	21.17	8.93	75.56	53.56	96.74	62.48	330.47		
YGR057C	LST7	1.06	1.51	1.72	2.13	2.30	2.41	2.85	1.30	30.80	11.04	33.65	12.34	333.36		
YGR058W	PEF1	1.01	1.41	1.62	2.05	2.26	2.30	4.09	1.21	24.31	16.17	46.40	17.38	319.54		
YGR058C	SPR3	0.99	1.33	1.49	1.89	2.06	2.13	2.38	1.11	33.37	8.15	35.75	9.26	313.55		
YGR061C	ADE6	0.85	1.12	1.12	1.06	1.15	1.14	109.85	105.65	124.77	119.13	234.62	224.78	500.28		
YGR062C	COX18	1.10	1.22	1.23	1.50	1.80	1.97	27.60	11.29	86.93	64.99	114.52	76.28	332.81		
YGR062C	COX18	0.89	1.06	1.08	0.93	0.92	0.94	123.61	116.33	130.91	96.69	254.52	215.02	507.04		
YGR063C	SPT4	0.93	0.96	1.00	1.23	1.47	1.74	48.31	38.62	96.25	91.32	144.56	129.95	367.02		
YGR064W	YGR044w	1.05	1.29	1.41	1.84	2.06	2.18	1.05	0.85	43.44	27.27	48.15	35.40	308.34		
YGR065C	ART5	1.06	1.47	1.69	2.10	2.26	2.33	1.17	0.97	23.74	10.72	24.91	11.69	282.47		
YGR069A	YGR069w	1.06	1.53	1.79	2.08	2.23	2.33	0.88	0.83	20.94	12.08	23.00	11.26	270.15		
YGR070C	RMP1	1.11	1.47	1.71	2.21	2.32	2.39	0.68	0.63	22.13	10.63	23.09	11.26	270.15		
YGR070W	URP3	0.98	1.32	1.53	2.01	2.19	2.28	3.22	3.64	30.64	8.45	32.97	5.51	288.18		
YGR070C	MRP12	0.94</														

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YGR153W	YGR153w	1.03	1.44	1.74	2.18	2.30	2.41	0.05	0.07	5.96	-0.15	6.02	-0.09	261.85		
YGR154C	GTO1	1.04	1.47	1.72	2.14	2.26	2.38	1.36	0.21	27.92	9.47	29.29	9.68	282.90		
YGR155W	CYS4	0.86	1.08	1.04	0.90	0.93	0.86	120.27	121.11	124.32	120.05	244.59	241.16	535.24		
YGR157W	CHO2	0.97	1.36	1.63	2.09	2.23	2.35	0.16	0.14	6.81	-0.99	6.97	-0.85	233.10		
YGR159C	NSR1	1.00	1.19	1.27	1.82	2.12	2.31	22.42	5.20	74.19	45.22	96.62	50.42	323.19		
YGR160W	YGR160w	0.95	1.17	1.27	1.74	2.06	2.27	21.38	1.03	71.61	25.81	92.98	26.84	334.07		
YGR161C	RTS1	0.89	0.98	0.97	1.22	1.46	1.68	1.60	0.81	18.88	2.51	19.46	2.68	260.86		
YGR161W-C	YGR161w-c	0.88	1.19	1.36	1.67	1.88	2.03	1.68	0.81	18.84	2.51	19.46	2.68	258.44		
YGR162C	YGR162c	1.07	1.44	1.50	1.74	1.86	1.99	3.24	0.09	56.41	1.46	56.46	1.46	228.48		
YGR163W	GT22	1.02	1.44	1.72	2.19	2.32	2.41	4.95	0.09	27.64	9.32	29.59	9.41	284.22		
YGR164W	YGR164w	0.97	1.32	1.54	2.00	2.16	2.29	4.65	-0.19	42.88	3.76	47.33	3.57	301.09		
YGR165W	MBS33	0.93	1.12	1.10	1.91	1.15	1.41	98.89	88.85	134.59	125.81	233.57	214.68	389.34		
YGR166W	KRE11	0.96	1.23	1.41	1.89	2.06	2.19	5.54	0.23	43.36	2.42	49.10	2.65	300.75		
YGR167W	CLC1	0.85	1.10	1.09	0.98	1.15	1.30	25.31	8.67	80.09	52.97	105.49	61.64	311.48		
YGR168C	YGR168c	0.94	1.36	1.42	1.78	2.03	2.17	16.38	5.24	66.28	38.43	82.66	43.67	322.58		
YGR169C	PUSE	1.03	1.44	1.69	2.08	2.23	2.35	1.42	0.35	27.38	10.72	28.79	11.07	286.54		
YGR169C-A	YGR169c-a	0.91	1.12	1.21	1.61	1.80	1.97	16.83	8.32	70.29	24.18	87.12	50.55	345.16		
YGR170W	PSD2	1.03	1.36	1.56	2.03	2.20	2.33	0.36	0.84	14.57	15.38	14.93	16.22	279.61		
YGR171C	MSM1	0.94	1.10	1.07	0.94	0.95	0.92	122.98	123.29	131.80	131.82	254.79	255.11	531.35		
YGR173W	RBG2	1.05	1.52	1.75	2.21	2.34	2.45	0.05	5.20	-0.14	5.26	-0.08	265.06			
YGR174C	CBP4	1.01	1.39	1.67	2.19	2.33	2.43	0.43	0.37	15.53	11.82	15.96	12.19	264.55		
YGR174W-A	YGR174w-a	0.94	1.37	1.58	1.98	2.13	2.24	10.82	7.79	57.89	20.03	68.71	21.81	307.98		
YGR176W	YGR176w	0.91	1.23	1.44	1.96	2.16	2.27	0.13	0.09	9.90	-0.02	10.03	0.08	281.25		
YGR177C	ATF2	0.96	1.31	1.52	2.06	2.26	2.37	0.11	-0.05	0.97	-0.37	1.08	-0.92	263.81		
YGR178C	PBP1	0.98	1.35	1.62	2.12	2.28	2.38	4.66	-0.02	39.77	3.36	44.44	3.34	299.53		
YGR180C	RNR4	0.90	1.15	1.22	1.51	1.77	1.96	19.82	10.73	71.38	5.16	91.20	6.39	320.16		
YGR180C	RNR4	0.96	1.08	1.05	0.92	0.90	0.90	127.50	124.35	135.64	130.94	263.14	255.29	535.39		
YGR182C	RNR4	0.88	1.16	1.14	0.99	0.99	0.96	115.96	90.90	121.52	109.54	237.48	200.44	480.98		
YGR181W	TIM13	1.00	1.29	1.46	1.93	2.11	2.24	19.07	0.93	65.30	23.23	84.36	24.08	341.52		
YGR182C	YGR182c	0.99	1.27	1.44	1.99	2.20	2.31	2.09	0.74	28.67	6.10	30.96	6.83	277.32		
YGR183C	CCR9	0.92	1.08	1.07	0.93	0.94	0.90	117.36	106.80	134.45	122.88	251.81	229.68	499.93		
YGR184C	UBR1	1.01	1.36	1.56	1.98	2.15	2.28	0.58	0.13	15.45	4.73	16.04	4.86	273.47		
YGR187C	HGH1	1.03	1.43	1.66	2.10	2.26	2.37	0.21	-0.01	5.55	0.63	5.76	0.02	218.98		
YGR188C	BUB1	0.91	1.33	1.44	1.74	1.87	1.97	32.52	21.20	62.37	68.60	114.89	89.80	343.46		
YGR189C	CRH1	0.95	1.28	1.49	1.97	2.14	2.27	2.17	2.07	36.49	29.35	38.66	31.42	293.06		
YGR192C	TDH2	0.94	1.24	1.45	1.94	2.10	2.25	1.14	0.69	25.99	5.66	27.13	6.95	296.91		
YGR193C	PDX1	1.01	1.33	1.55	2.01	2.17	2.30	1.46	2.24	27.47	30.18	28.93	32.42	256.15		
YGR194C	XKL1	0.96	1.25	1.43	1.68	2.00	2.12	0.11	0.63	16.65	0.45	16.76	0.65	281.13		
YGR195C	ITF4	0.93	1.19	1.33	1.84	2.00	2.17	6.76	0.18	50.45	12.74	51.96	12.96	301.47		
YGR197C	SMG1	0.98	1.26	1.38	2.13	2.30	2.39	0.76	0.30	18.68	0.71	19.42	0.74	287.65		
YGR199W	PMT1	0.91	1.46	1.69	2.12	2.25	2.39	-0.02	0.71	2.42	4.58	2.39	1.21	268.46		
YGR200C	EPU2	0.94	1.15	1.28	1.71	1.93	2.11	0.14	0.16	9.23	0.11	9.77	0.15	255.43		
YGR201C	YGR201c	0.94	1.36	1.57	1.97	2.16	2.30	3.24	2.29	12.49	5.73	15.73	8.93	286.45		
YGR202C	PTC1	1.06	1.32	1.48	2.03	2.21	2.34	11.36	0.64	54.95	18.31	66.31	18.95	305.37		
YGR203W	YCH1	1.06	1.55	1.83	2.24	2.36	2.45	0.17	0.08	5.52	-0.64	5.69	-0.56	262.89		
YGR204C-A	YGR204c-a	0.91	1.25	1.39	1.73	1.90	2.03	8.57	1.48	55.11	18.19	63.67	19.67	299.42		
YGR204W	ADE3	0.83	1.14	1.14	1.07	1.16	1.20	103.87	104.13	119.22	113.86	223.09	217.99	488.50		
YGR205W	YGR205w	1.03	1.45	1.71	2.14	2.28	2.39	0.26	0.18	11.88	2.26	12.13	0.44	279.39		
YGR206W	MV12	0.97	1.22	1.36	1.78	2.02	2.18	0.16	0.09	13.98	0.65	14.14	0.74	266.89		
YGR207C	YGR207c	1.04	1.42	1.61	2.07	2.22	2.35	0.43	0.10	19.60	6.80	20.03	6.90	289.25		
YGR208W	SER2	0.85	1.19	1.33	1.67	1.92	2.07	2.35	1.53	21.93	8.05	24.28	9.57	258.15		
YGR209C	TRX2	0.93	1.16	1.29	1.81	2.01	2.15	0.09	0.09	5.66	0.12	5.74	0.21	258.46		
YGR210C	YGR210c	0.98	1.53	1.72	2.07	2.23	2.32	8.52	3.30	42.27	14.19	50.79	17.50	301.24		
YGR212W	SLJ1	0.90	1.19	1.40	1.96	2.13	2.24	0.11	0.03	10.33	1.66	10.44	1.69	273.25		
YGR213C	RTA1	0.94	1.36	1.61	2.01	2.23	2.30	3.14	2.00	36.66	9.76	39.79	9.96	305.22		
YGR214W	RPS9a	0.95	1.20	1.33	1.93	2.19	2.31	1.87	0.32	31.88	3.76	33.75	4.08	291.99		
YGR215W	RSM27	0.88	1.09	1.12	1.30	1.57	1.73	1.98	0.91	53.90	9.12	51.42	8.82	287.98		
YGR217W	CCH1	0.96	1.29	1.50	2.01	2.20	2.31	0.66	0.26	41.26	12.69	47.92	15.95	304.31		
YGR217W	DIE2	0.99	1.33	1.53	1.96	2.12	2.26	5.04	3.18	34.85	2.12	39.90	1.38	297.19		
YGR220C	YGR220w	0.93	1.36	1.58	2.03	2.21	2.32	3.84	2.37	23.77	5.74	27.44	6.53	322.11		
YGR221C	SMH1	0.93	1.12	1.17	1.53	1.70	1.82	2.07	0.69	32.54	8.08	35.51	8.77	285.13		
YGR220W	BWS1	0.98	1.27	1.36	1.68	1.82	2.02	5.55	3.16	35.82	11.29	41.38	14.45	301.24		
YGR221C	PHB2	1.04	1.30	1.39	1.76	1.94	2.11	12.97	4.17	58.91	27.35	71.87	31.52	322.66		
YGR222W	NAS8	0.97	1.27	1.39	1.85	2.04	2.22	6.79	3.57	51.04	20.26	58.83	22.83	311.44		
YGR223C	PHO81	0.90	1.10	1.14	1.75	1.84	1.86	10.59	6.25	51.58	43.23	62.08	49.48	311.95		
YGR224W	YBH1	1.02	1.48	1.69	2.03	2.15	2.25	9.69	3.95	38.68	10.70	42.81	9.88	308.30		
YGR225C	YGR25c	1.08	1.34	1.45	1.93	2.07	2.15	4.95	2.24	32.17	9.42	37.12	9.45	280.23		
YGR224C	LSC2	0.93	1.28	1.39	1.88	2.04	2.19	8.26	4.02	46.85	27.78	55.11	31.80	308.78		
YGR224C	LSC2	0.92	1.25	1.34	1.73	1.92	2.07	8.21	4.02	51.20	16.09	59.41	17.41	292.6		

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YHL010C	ETP1	0.98	1.35	1.58	2.06	2.23	2.34	0.02	-0.04	6.57	0.04	6.59	0.01	275.01		
YHL011C	PRS3	1.01	1.40	1.44	1.75	1.93	2.10	6.93	4.07	47.01	36.29	53.94	40.36	358.54		
YHL012W	YHL012w	1.02	1.34	1.53	2.03	2.20	2.31	3.02	0.07	36.27	9.95	39.30	10.02	295.00		
YHL013C	OTU2	1.03	1.44	1.63	2.06	2.25	2.36	-0.01	-0.06	2.98	-0.24	2.97	-0.31	284.71		
YHL014W	YLF2	1.00	1.38	1.60	2.05	2.23	2.34	0.02	0.02	5.66	-0.21	5.68	-0.19	267.87		
YHL015W-A	YHL015w-a	0.93	1.27	1.51	1.89	2.04	2.18	8.55	1.74	54.69	18.95	63.24	20.70	299.09		
YHL016C	DUR2	0.98	1.22	1.40	1.91	2.08	2.23	0.65	0.62	8.21	0.90	8.27	0.51	252.40		
YHL017W	YHL017w	1.00	1.35	1.56	1.98	2.14	2.30	0.64	-0.01	3.83	-0.55	3.87	-0.67	253.35		
YHL018C	YHL018w	0.98	1.21	1.33	1.98	2.08	2.23	0.24	0.23	5.23	0.24	5.26	0.22	225.00		
YHL019C	CPR1	0.98	1.21	1.33	1.98	2.10	2.21	0.12	-0.02	10.24	-0.26	10.36	-0.28	276.51		
YHL021C	AHM17	0.94	1.23	1.39	1.90	2.10	2.21	0.12	-0.02	5.77	5.57	-8.56	-8.56	283.56		
YHL022C	SPO11	0.97	1.37	1.57	2.02	2.15	2.29	0.12	0.14	5.44	-0.27	5.57	-0.28	285.42		
YHL023C	NPR2	0.97	1.21	1.33	1.89	2.09	2.20	6.83	0.43	54.98	21.23	61.81	21.66	285.42		
YHL024W	RIM4	0.91	1.35	1.53	1.90	2.06	2.19	8.88	3.81	43.22	22.48	52.10	26.29	296.37		
YHL025W	SMF6	0.93	1.27	1.39	1.71	1.90	2.07	13.39	3.78	57.96	25.17	71.35	28.95	367.51		
YHL026C	YHL026c	0.93	1.23	1.42	1.89	2.08	2.23	0.08	-0.01	3.95	-0.58	4.03	-0.58	273.74		
YHL027W	RIM101	0.98	1.18	1.34	1.77	1.95	2.09	2.22	0.16	27.76	5.98	29.98	6.13	288.98		
YHL028W	WSC4	1.00	1.33	1.48	1.95	2.13	2.27	0.29	0.02	17.95	3.18	18.24	3.20	285.81		
YHL029C	OC45	1.05	1.55	1.78	2.16	2.33	2.35	0.03	-0.07	-0.11	-1.00	-0.08	-1.07	237.92		
YHL030W	ECM29	0.86	1.09	1.21	1.45	1.68	1.87	1.32	1.36	4.56	1.76	5.87	3.11	293.16		
YHL031C	GOS1	0.97	1.28	1.50	1.99	2.17	2.33	0.56	0.04	15.38	2.25	15.94	2.29	279.67		
YHL032C	GUT1	1.05	1.50	1.72	2.13	2.29	2.43	0.24	0.04	0.27	-0.93	0.51	-0.89	271.07		
YHL033C	RPL8a	1.03	1.30	1.49	1.98	2.18	2.28	0.12	0.06	0.73	-0.75	0.84	-0.69	257.06		
YHL034C	SBP1	1.02	1.27	1.41	1.91	2.13	2.23	6.31	0.66	42.80	11.00	49.11	11.66	312.24		
YHL035C	VMR1	0.93	1.21	1.40	1.89	2.09	2.20	0.15	0.33	11.09	1.35	11.24	1.68	281.51		
YHL036W	MUP3	0.96	1.28	1.44	1.91	2.09	2.21	0.00	0.00	10.07	-0.38	10.07	-0.38	287.90		
YHL037C	YHL037c	1.03	1.44	1.66	2.17	2.34	2.41	1.20	-0.01	26.19	1.87	27.39	1.86	296.39		
YHL038C	CBP2	0.93	1.14	1.11	0.97	1.02	0.94	124.63	124.69	131.63	135.21	256.26	259.90	538.63		
YHL039W	EPM1	0.99	1.50	1.63	1.99	2.13	2.21	7.80	5.96	40.24	32.14	48.03	38.11	304.31		
YHL040C	ARN1	0.89	1.19	1.30	1.72	1.90	2.05	0.02	0.13	1.05	-0.90	1.07	-0.76	275.56		
YHL041W	YHL041w	1.03	1.44	1.62	2.04	2.21	2.36	0.00	0.01	0.11	-1.04	0.11	-1.02	263.44		
YHL042W	YHL042w	0.98	1.35	1.55	2.01	2.16	2.30	0.79	0.02	24.23	2.65	25.02	2.67	293.11		
YHL043W	ECM34	0.96	1.30	1.51	2.01	2.15	2.29	2.51	0.07	34.63	6.36	37.14	6.42	294.02		
YHL044W	YHL044w	0.94	1.27	1.45	1.90	2.10	2.26	0.08	0.09	1.22	-0.88	1.29	-0.80	276.18		
YHL045W	YHL045w	0.95	1.28	1.47	1.96	2.13	2.31	0.10	0.05	3.21	-0.64	3.31	-0.59	264.14		
YHL046C	PAU13	1.03	1.60	1.74	2.11	2.28	2.34	0.04	0.02	0.43	-0.99	0.47	-0.97	262.46		
YHL047C	ARN2	1.06	1.39	1.62	2.06	2.33	2.43	0.85	0.87	27.24	2.74	28.18	7.34	296.59		
YHL001W	OS27	1.01	1.25	1.39	1.70	1.96	2.15	0.05	0.03	4.11	1.37	45.56	15.24	285.35		
YHL002W-A	CGT70	0.98	1.30	1.55	2.02	2.18	2.32	0.07	-0.01	3.15	5.81	3.74	0.12	226.42		
YHL003W	YHL003c	1.01	1.47	1.60	1.96	2.10	2.21	11.19	12.69	53.70	49.80	64.89	62.59	315.62		
YHL004C	NEML1	0.97	1.44	1.57	1.89	2.02	2.08	22.44	17.83	65.23	58.71	78.68	77.74	347.40		
YHL005C	GPM1	0.98	1.19	1.42	1.81	2.02	2.18	7.80	0.87	48.45	6.95	54.28	7.82	295.96		
YHL006W	STP2	1.00	1.53	1.68	2.08	2.23	2.32	4.77	4.49	28.65	24.91	33.42	29.41	275.41		
YHL007C-A	YHL022c-a	0.93	1.27	1.48	1.84	2.02	2.15	11.46	3.97	61.50	30.34	72.96	34.31	308.58		
YHL008C	SOD2	0.97	1.45	1.60	1.96	2.12	2.21	7.21	3.69	41.29	10.97	48.50	14.06	302.43		
YHL009C	YHL009c	0.96	1.54	1.72	2.08	2.21	2.29	3.73	3.00	18.45	2.62	22.19	5.63	284.69		
YHL010W	RPL27a	0.88	1.07	1.06	1.08	1.36	1.73	41.93	25.78	102.97	68.92	144.90	94.70	319.31		
YHL011W	DI44	0.92	1.23	1.32	1.86	2.12	2.30	14.14	4.30	63.34	37.36	77.48	41.66	265.74		
YHL012W	VPS29	0.94	1.25	1.39	1.74	1.88	2.04	0.83	-0.13	26.24	5.44	27.07	5.31	332.43		
YHL013C	ARD1	1.01	1.29	1.49	2.01	2.18	2.27	1.81	-0.01	18.65	5.75	20.45	5.74	243.62		
YHL014W	SPO13	0.98	1.35	1.56	2.00	2.17	2.30	-0.09	-0.17	8.26	0.84	8.17	0.67	265.95		
YHL015W	MIP6	0.97	1.26	1.41	1.86	2.07	2.22	-0.09	-0.04	1.72	-0.78	1.63	-0.82	260.43		
YHL016C	YSC84	0.86	1.21	1.30	1.58	1.75	1.91	8.96	3.13	43.90	12.02	52.86	15.15	263.17		
YHL017W	YSC83	0.85	1.26	1.37	1.73	1.92	2.05	7.35	3.72	36.49	18.18	43.84	21.90	302.34		
YHL018C	ARG4	0.88	1.10	1.13	1.30	1.53	1.74	16.02	3.54	73.92	38.97	89.94	42.51	304.57		
YHL019C	RPS27b	0.84	1.09	1.10	1.37	1.66	1.90	1.98	0.29	33.65	6.57	35.62	6.86	264.16		
YHL021W-A	ECM12	0.93	1.37	1.53	1.96	2.16	2.21	13.08	2.12	59.50	24.69	69.57	34.85	326.21		
YHL022C	YHL022c	0.92	1.16	1.32	1.86	2.06	2.19	-0.03	-0.07	6.86	0.00	6.83	-0.07	276.83		
YHL034C	PHT1	1.00	1.29	1.44	1.95	2.16	2.30	7.35	0.86	46.91	13.06	54.25	13.93	289.88		
YHL035A	YHL035w	0.94	1.27	1.58	2.04	2.21	2.35	4.82	3.63	36.03	4.25	41.26	2.70	279.77		
YHL036W	PFT2	1.01	1.33	1.53	1.97	2.16	2.29	3.97	2.65	37.63	16.71	41.69	16.97	327.40		
YHL038W	RPSF1	0.91	1.28	1.48	2.00	2.17	2.28	9.98	2.92	55.08	34.86	65.06	37.90	269.45		
YHL039C	MSC7	0.98	1.36	1.60	2.11	2.28	2.37	1.25	0.69	19.28	-0.05	20.53	0.64	260.59		
YHL039C-B	YHL039c-B	0.91	1.12	1.12	1.14	1.44	1.72	46.37	26.59	104.56	85.17	150.94	111.86	324.42		
YHL041C	DGQ1	0.98	1.30	1.47	1.96	2.17	2.32	1.97	-0.02	21.95	0.78	23.93	0.78	245.05		
YHL045W	YHL045w	0.95	1.33	1.53	2.03	2.20	2.33	1.09	0.50	15.78	1.33	16.87	1.83	255.07		
YHL046C	INMT	0.88	1.14	1.33	1.84	2.05	2.18	0.63	4.67	3.21	46.09	3.84	50.76	273.36		
YHL047C	AAP1	0.99	1.34	1.56	2.06	2.25	2.36	5.98	1.29	45.31	19.70	51.29	20.99	272.32		
YHL048W	YHK8	0.95	1.24	1.48	2.00	2.17	2.28	3.14	1.31	26.90	26.70	30.03	29.21	298.03		
YHL049C-A	YHL049c-a	1.03	1.34	1.53</												

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE			FRUCTOSE			TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YHR123W	EP11	0.88	1.12	1.24	1.71	1.93	2.02	1.14	1.12	9.63	8.26	10.77	9.38	287.48				
YHR124W	ND780	0.87	1.12	1.28	1.82	2.14	2.18	127.68	1.05	124.78	5.88	252.46	6.93	295.54				
YHR125W	YHR125w	0.91	1.17	1.35	1.92	2.13	2.21	20.65	1.19	73.24	8.94	93.88	10.13	285.49				
YHR126C	ANST	0.89	1.13	1.25	1.82	2.04	2.19	139.28	1.08	135.28	7.92	274.56	9.00	299.30				
YHR127W	YHR127w	0.98	1.51	1.75	2.14	2.25	2.32	19.49	6.61	59.62	32.57	79.30	39.18	343.89				
YHR129C	ARP1	0.95	1.14	1.19	1.66	1.89	2.08	3.31	12.69	35.36	7.20	38.67	84.89	314.85				
YHR130C	YHR130c	1.00	1.29	1.47	1.86	2.07	2.29	11.08	1.13	57.94	12.49	69.02	13.62	360.27				
YHR131C	YHR131c	1.42	1.58	1.79	2.17	2.29	2.37	6.60	3.14	35.71	7.63	41.71	10.33	361.31				
YHR132W	A02	0.90	1.06	1.22	1.78	1.99	2.16	3.71	6.65	38.42	4.63	42.13	4.88	297.74				
YHR133C	NSG1	0.97	1.33	1.54	2.06	2.24	2.38	27.17	1.09	80.81	1.02	107.98	2.11	278.80				
YHR134W	WSS1	0.95	1.20	1.33	1.80	2.00	2.14	122.41	2.26	128.09	29.48	242.51	31.17	326.16				
YHR135C	YCK1	0.88	1.31	1.49	2.02	2.25	2.36	4.64	1.20	40.18	5.20	44.82	6.41	274.38				
YHR136C	SPL2	0.89	1.19	1.30	1.67	1.88	2.03	13.81	4.54	60.18	29.53	73.99	34.07	312.38				
YHR137W	AR09	0.92	1.21	1.41	1.96	2.15	2.23	1.10	1.32	5.22	7.39	6.32	8.71	304.45				
YHR138C	YHR138c	0.89	1.18	1.34	1.84	2.05	2.20	1.14	1.05	9.33	0.19	10.47	1.24	275.74				
YHR139C	SPS700	0.91	1.23	1.40	1.97	2.16	2.26	1.32	1.89	12.83	1.41	14.16	2.51	286.10				
YHR139c-a	YHR139c-a	0.96	1.24	1.41	1.91	2.10	2.22	1.48	1.28	18.15	10.25	19.63	11.53	298.89				
YHR140W	YHR140w	0.92	1.28	1.51	1.93	2.10	2.20	9.58	3.39	46.51	20.89	56.09	24.28	311.21				
YHR141C	RPL42b	0.93	1.15	1.17	1.20	1.45	1.88	36.33	16.00	95.96	57.37	132.29	73.37	332.04				
YHR142W	CHST	0.90	1.13	1.28	1.86	2.08	2.19	3.30	1.33	33.40	16.34	36.70	17.73	312.04				
YHR143W	DSE2	0.97	1.18	1.31	1.92	2.14	2.27	4.46	1.08	38.69	11.37	43.15	12.45	306.52				
YHR146W	CRP1	0.97	1.23	1.50	1.98	2.13	2.26	4.97	0.86	42.25	10.66	47.21	11.52	311.83				
YHR147C	MRLP6	0.92	1.17	1.16	1.16	1.51	1.91	40.15	24.37	95.78	71.76	135.93	96.13	301.79				
YHR149C	SKG6	0.92	1.29	1.45	1.87	2.04	2.17	5.59	1.30	44.87	13.99	50.46	15.29	289.27				
YHR150W	PEX28	0.97	1.25	1.44	1.95	2.14	2.24	8.49	2.04	52.54	25.30	61.02	27.34	309.75				
YHR151C	MTC6	0.96	1.27	1.48	1.99	2.14	2.21	4.14	1.45	36.75	16.25	40.89	17.70	324.51				
YHR152W	SP012	0.96	1.26	1.47	1.99	2.15	2.26	3.74	1.14	37.51	5.58	41.25	6.71	308.44				
YHR153C	SP016	0.91	1.17	1.34	1.90	2.09	2.20	0.93	1.21	4.97	7.12	5.90	8.33	314.35				
YHR154W	RTT107	0.97	1.19	1.31	2.07	2.05	1.99	1.00	1.15	7.93	6.05	8.94	7.20	310.60				
YHR155W	YSP1	0.95	1.27	1.49	2.00	2.15	2.25	1.12	0.99	0.54	8.48	1.66	9.47	310.88				
YHR156C	LIN1	0.90	1.17	1.33	1.84	2.05	2.19	0.88	1.17	4.53	0.18	5.41	1.34	293.55				
YHR157W	REC104	0.90	1.19	1.37	1.92	2.08	2.18	1.27	1.03	4.44	8.08	5.71	9.11	316.80				
YHR158C	KEL7	0.88	1.08	1.19	1.92	2.20	2.30	1.10	0.92	17.40	2.50	18.14	3.43	288.51				
YHR159W	YHR159w	0.90	1.14	1.28	1.84	2.05	2.18	4.97	1.23	39.67	14.33	44.64	15.55	314.48				
YHR160C	PEX18	0.95	1.17	1.33	1.92	2.13	2.26	122.42	0.60	126.25	7.05	247.68	1.35	313.85				
YHR161C	YAP1801	0.98	1.21	1.34	1.86	2.10	2.26	116.27	2.64	118.71	7.70	234.98	10.34	335.35				
YHR162W	YHR162w	0.93	1.33	1.50	1.91	2.15	2.29	13.56	6.85	52.53	3.43	67.17	4.17	327.47				
YHR163W	SCD1	0.97	1.28	1.45	2.01	2.12	2.35	15.43	1.64	61.70	4.66	71.13	1.59	284.71				
YHR167W	MP2	0.96	1.22	1.33	1.81	1.99	2.13	126.39	7.47	121.76	56.86	245.05	64.33	378.46				
YHR168W	MTC6	0.91	1.20	1.19	1.55	1.84	2.07	13.64	6.55	56.64	37.97	70.28	44.51	308.29				
YHR171W	ATG7	0.91	1.21	1.43	1.93	2.13	2.26	28.89	4.94	77.83	4.20	103.72	4.74	311.16				
YHR175W-A	YHR175w-a	0.93	1.12	1.08	0.97	0.97	0.95	120.68	122.10	135.34	104.12	256.02	226.22	472.78				
YHR176W	FMO1	0.88	1.23	1.36	1.68	1.87	1.98	12.84	3.79	57.39	3.25	70.23	3.64	314.89				
YHR177W	YHR177w	0.90	1.21	1.33	1.74	1.98	2.08	4.48	1.70	35.79	13.10	40.27	14.89	316.52				
YHR178W	STB5	0.91	1.16	1.28	1.93	2.01	2.21	1.14	1.05	9.43	2.94	10.56	3.98	284.51				
YHR179W	OYE2	0.91	1.19	1.36	1.90	2.08	2.19	1.74	0.81	23.62	7.78	25.36	8.58	293.68				
YHR180W	YHR180w	0.99	1.54	1.76	2.14	2.26	2.33	7.12	3.20	40.81	9.69	47.92	12.89	304.24				
YHR181W	SP26	0.99	1.44	1.60	2.00	2.19	2.32	9.92	4.57	46.32	27.50	56.23	32.07	305.41				
YHR182W	YHR182w	0.96	1.32	1.53	2.05	2.20	2.31	4.56	0.44	42.74	2.32	47.30	2.76	280.66				
YHR183W	GND1	0.92	1.19	1.39	1.72	1.93	2.07	1.58	1.51	14.63	3.16	16.22	4.66	276.39				
YHR184W	SSP1	0.92	1.17	1.35	1.89	2.10	2.23	21.37	0.96	71.37	6.56	92.74	7.52	315.76				
YHR185C	PFS1	1.00	1.50	1.70	2.15	2.29	2.36	5.49	3.34	33.00	13.02	38.48	16.35	275.71				
YHR187W	IK1	1.00	1.29	1.42	1.78	1.94	2.06	27.66	5.87	80.22	35.66	107.88	41.53	333.74				
YHR189W	PTH1	0.95	1.17	1.33	1.95	2.14	2.27	27.31	1.05	75.91	2.16	103.22	3.20	312.69				
YHR191C	CTFB	0.94	1.25	1.25	1.47	1.73	1.98	43.33	3.20	83.76	7.72	127.08	10.62	394.10				
YHR192W	YHR192w	0.98	1.31	1.48	1.85	2.03	2.18	4.64	1.01	39.08	7.63	43.71	8.63	276.44				
YHR193C	EGD2	0.89	1.27	1.40	1.81	2.01	2.16	9.15	3.47	46.37	17.96	55.52	21.44	238.15				
YHR205W	SKN7	0.95	1.22	1.39	1.91	2.12	2.24	5.58	0.98	42.73	9.44	48.24	10.42	322.14				
YHR206C	SET1	0.95	1.12	1.26	1.74	1.96	2.11	16.41	3.51	67.50	36.99	83.23	40.51	339.39				
YHR209W	CBS1	0.97	1.25	1.51	1.95	2.13	2.24	13.31	1.60	60.32	38.23	73.63	14.98	348.45				
YHR210C	YIL010w	0.89	1.08	1.22	1.72	1.96	2.06	16.68	1.08	70.82	30.40	86.91	32.35	357.60				
YIL001W	YIL001w	0.99	1.52	1.51	1.98	2.15	2.27	3.66	3.09	21.33	8.51	24.89	11.61	269.70				
YIL002C	INPS1	0.98	1.54	1.63	2.03	2.14	2.23	5.76	4.22	37.21	2.91	42.97	30.23	320.95				
YIL002W-a	YIL002w-a	0.97	1.42	1.49	1.81	1.95	2.07	6.32	3.07	38.88	2.44	44.44	2.74	288.19				
YIL004C	EPS1	0.89	1.41	1.56	1.93	2.07	2.16	3.15	3.07	7.75	4.73	10.90	7.80	256.59				

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YIL074C	SER33	0.89	1.24	1.36	1.71	1.86	2.01	11.70	11.29	51.44	48.22	63.14	59.52	297.19		
YIL076W	SEC24	0.87	1.15	1.24	1.66	1.87	1.98	10.39	6.28	50.52	38.22	60.91	44.50	309.37		
YIL077C	YIL077c	0.92	1.33	1.51	1.90	2.09	2.21	5.07	3.20	31.50	19.51	36.57	22.72	300.21		
YIL079C	AIR1	0.86	1.22	1.39	1.76	1.92	2.05	3.34	2.92	20.46	12.61	23.80	15.53	272.95		
YIL084C	SDS2	0.85	1.25	1.46	1.82	1.98	2.08	3.62	3.24	22.72	21.47	26.34	24.71	270.01		
YIL085C	KTR7	0.92	1.16	1.32	1.75	1.97	2.12	4.32	1.56	37.41	21.38	41.73	22.94	299.27		
YIL086C	YIL086c	0.88	1.25	1.43	1.81	1.98	2.08	4.10	4.16	27.44	23.17	31.60	29.32	282.30		
YIL087C	ADM10	0.81	1.27	1.42	1.80	1.98	2.08	6.84	11.02	40.67	47.62	46.61	58.84	384.24		
YIL088C	AVT1	0.88	1.24	1.41	1.74	1.91	2.08	4.10	3.85	34.95	24.59	35.89	26.51	330.14		
YIL089W	YIL089w	0.88	1.29	1.45	1.85	2.03	2.17	4.68	5.17	30.35	28.19	35.63	33.36	310.40		
YIL090W	ICE2	0.86	1.18	1.28	1.82	1.79	1.92	15.63	9.35	60.59	49.23	76.23	58.57	337.72		
YIL092W	YIL092w	0.91	1.26	1.38	1.73	1.84	1.98	9.86	5.71	46.68	33.09	56.54	38.79	335.57		
YIL093W	RSM25	0.88	1.28	1.49	1.86	2.02	2.12	6.10	4.86	35.03	27.68	41.13	32.54	294.30		
YIL094C	LYS12	0.93	1.24	1.46	1.82	2.01	2.15	0.60	0.57	1.33	0.81	1.93	1.18	298.64		
YIL095W	PBK1	0.86	1.18	1.31	1.68	1.85	1.95	17.41	10.13	57.51	46.60	74.92	56.73	341.10		
YIL096C	YIL096c	0.88	1.23	1.41	1.83	1.99	2.12	11.01	4.46	47.96	27.04	58.97	31.51	327.93		
YIL097W	FYV10	0.97	1.33	1.50	1.94	2.08	2.23	7.10	3.71	41.42	22.21	48.53	25.92	309.39		
YIL098C	FMC1	0.96	1.20	1.43	1.86	2.05	2.21	1.35	0.59	16.27	1.17	17.62	1.75	281.87		
YIL099W	SGA1	0.90	1.23	1.40	1.73	1.93	2.08	8.02	1.28	50.82	26.49	58.85	28.31	307.28		
YIL100W	YIL100w	0.89	1.22	1.37	1.61	1.79	1.92	27.45	10.98	77.70	56.61	105.14	67.59	316.65		
YIL101C	XBP1	0.84	1.16	1.30	1.62	1.82	1.99	0.92	0.67	17.46	14.76	18.38	6.39	361.04		
YIL102C	YIL102c	1.01	1.29	1.50	1.92	2.10	2.15	10.30	1.46	51.22	16.94	61.52	18.40	304.19		
YIL103W	DPH1	0.92	1.21	1.29	1.51	1.74	1.92	14.19	3.47	61.04	36.88	75.22	40.35	323.09		
YIL105C	SLM1	0.92	1.24	1.39	1.77	1.97	2.13	15.54	4.22	64.83	39.25	80.37	43.48	273.78		
YIL107C	PPK26	0.93	1.28	1.43	1.73	1.91	2.06	8.67	1.38	54.58	24.10	63.24	25.48	324.87		
YIL108W	YIL108w	0.91	1.20	1.31	1.70	1.92	2.10	13.04	3.30	61.22	37.11	74.26	40.41	343.38		
YIL109W	MNI1	0.89	1.19	1.29	1.70	1.93	2.13	5.84	1.05	48.06	19.21	53.89	20.25	323.40		
YIL111W	Cox5b	0.95	1.29	1.54	1.91	2.06	2.23	7.22	0.80	43.64	3.54	50.87	4.34	308.62		
YIL112W	HOS4	0.96	0.95	0.96	1.01	1.09	1.03	136.70	122.58	141.96	130.99	278.66	253.58	187.08		
YIL113W	SDP1	1.00	1.13	1.29	1.89	2.11	2.28	5.57	7.06	47.24	51.05	52.81	58.10	161.36		
YIL114C	POR2	1.06	1.22	1.42	1.99	2.15	2.32	9.20	10.36	61.08	59.81	70.28	70.17	188.48		
YIL116W	HIS5	0.84	1.09	1.05	0.93	0.95	0.86	123.52	124.45	122.57	123.63	246.09	248.07	530.06		
YIL117C	PRM5	0.99	1.13	1.31	1.84	2.09	2.24	12.08	7.06	65.86	51.29	77.94	58.34	207.74		
YIL119C	RPT1	0.99	1.18	1.41	1.84	2.13	2.31	0.60	0.39	2.48	0.30	3.08	0.89	218.05		
YIL120W	QDR1	0.90	1.02	1.23	1.81	2.03	2.19	10.42	0.58	59.78	7.47	70.20	8.05	184.06		
YIL121W	QDR2	0.92	1.14	1.43	1.92	2.11	2.23	12.05	4.77	59.57	19.29	71.62	42.06	185.87		
YIL122W	POG1	0.92	1.16	1.44	1.81	2.01	2.14	16.35	1.67	59.03	13.96	75.39	14.81	308.48		
YIL123W	SM1	0.88	1.16	1.38	1.88	2.08	2.28	20.38	0.38	63.97	7.37	64.64	6.66	147.43		
YIL124W	VPR1	0.89	1.15	1.35	1.82	2.03	2.17	13.51	4.66	62.59	20.40	76.10	14.86	149.01		
YIL125W	KGD1	0.84	1.12	1.30	1.87	2.00	2.26	9.48	2.07	52.37	31.68	61.85	34.65	134.30		
YIL132C	RDT14	0.87	1.14	1.33	1.75	1.96	2.08	5.24	0.85	48.45	9.26	50.98	9.37	284.44		
YIL138W	MET18	0.84	1.03	1.18	1.73	1.93	2.15	17.22	3.46	58.89	36.86	86.22	49.32	114.36		
YIL139W	ASG1	0.95	0.95	0.96	1.00	1.05	1.02	86.14	138.02	121.42	142.00	207.55	280.02	228.88		
YIL131C	FKH1	0.86	1.19	1.41	1.83	2.01	2.18	11.05	3.35	52.68	28.77	63.74	32.13	300.41		
YIL132C	CSM2	1.03	1.17	1.34	1.87	2.14	2.36	2.07	0.70	4.13	4.49	6.20	5.19	231.90		
YIL133C	RPL16a	0.92	0.97	1.11	1.70	1.94	2.15	2.41	2.95	30.89	3.75	33.70	34.70	257.22		
YIL134C-A	YIL134c-a	0.91	1.29	1.43	1.74	1.89	2.01	20.70	2.69	77.17	26.60	97.87	29.29	328.34		
YIL134W	FLX1	1.04	1.13	1.29	1.84	2.06	2.23	3.65	4.04	36.48	44.06	40.13	262.68			
YIL135C	VHS2	1.02	1.10	1.29	1.57	1.69	2.00	0.85	1.15	12.45	9.29	13.30	10.44	253.80		
YIL136W	OM45	0.93	1.24	1.50	1.91	2.10	2.24	3.62	0.95	30.38	9.87	34.01	10.82	296.06		
YIL137C	TMA108	0.98	1.17	1.33	1.75	1.92	2.12	3.00	0.73	35.90	6.05	38.90	6.78	242.74		
YIL138C	TPM2	0.94	1.03	1.22	2.01	2.05	2.19	19.38	1.61	70.73	24.02	90.11	25.63	192.30		
YIL139C	REV7	0.96	1.12	1.33	1.87	2.04	2.20	23.30	10.45	73.21	55.04	96.51	65.49	213.20		
YIL140W	AXL2	1.00	1.12	1.31	1.86	2.05	2.20	31.39	15.86	86.02	66.47	117.41	82.33	184.55		
YIL141W	YIL141w	0.96	1.06	1.23	1.78	2.01	2.18	31.73	13.59	88.22	66.27	119.95	79.86	246.05		
YIL145C	PAN6	0.90	0.97	1.13	1.62	1.84	2.03	26.65	1.27	82.69	59.16	109.34	71.73	239.83		
YIL146C	PAN6	0.91	1.35	1.52	1.89	2.07	2.20	3.00	2.12	34.94	22.84	37.94	24.97	289.60		
YIL157W	AM20	0.95	1.21	1.43	1.87	2.02	2.15	14.99	3.33	30.10	32.10	74.09	35.44	305.74		
YIL158W	BHR1	0.97	1.20	1.43	1.89	2.03	2.15	7.53	1.17	48.14	17.90	55.67	19.67	220.40		
YIL160C	DTT1	0.92	1.08	1.24	1.73	1.92	2.03	8.17	0.89	49.45	15.11	53.62	16.89	242.42		
YIL161W	YIL161w	0.93	1.07	1.28	1.79	1.99	2.17	5.99	1.20	50.47	17.19	56.46	18.28	244.43		
YIL162W	SUC2	0.92	1.12	1.34	1.86	2.07	2.22	6.97	1.33	50.20	21.10	57.17	22.43	243.56		
YIL163C	YIL163c	0.96	1.00	1.10	1.71	1.94	2.04	8.89	6.37	57.21	53.32	68.11	59.69	240.40		
YIL164C	NIT1	0.98	1.09	1.28	1.92	2.14	2.32	7.38	5.29	54.25	46.72	61.64	52.01	249.39		
YIL165C	YIL165c	0.92	1.23	1.49	2.06	2.23	2.35	1.15	1.76	18.02	22.69	19.18	24.45	227.14		
YIL166C	YIL166c	0.93	1.06	1.27	1.84	2.04	2.21	3.77	3.85	41.29	40.41	45.06	44.26	239.67		
YIL167W	SDL1	0.94	1.08	1.27	1.89	2.01	2.20	14.06	11.14	51.02	36.04	57.16	38.81	206.81		
YIL168W	YIL168w	0.99	1.16	1.36	1.89	2.07	2.23	10.43	15.10	71.67	56.97	96.04	72.07	344.20		
YIL202C	YIL202c	1.00	1.52	1.69	2.24	2.15	2.10	13.4								

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE			FRUCTOSE			TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YJL024C	APS3	0.90	1.27	1.40	1.76	1.93	2.03	21.28	1.04	81.39	22.09	102.67	23.13	299.67				
YJL027C	YJL027c	0.88	0.95	0.97	0.88	0.84	131.68	122.18	156.50	114.09	288.18	236.28	412.15					
YJL028W	YJL028w	0.86	1.02	1.18	1.59	1.81	1.97	4.96	1.07	46.77	14.85	51.73	15.92	288.49				
YJL029C	VPS33	0.88	1.23	1.35	1.60	1.78	1.87	9.64	1.49	59.50	23.45	69.14	24.94	282.84				
YJL030W	MAD2	1.04	1.68	1.85	2.15	2.24	2.32	5.00	8.24	32.58	42.27	37.58	50.51	282.42				
YJL036W	SNX4	1.00	1.59	1.77	2.13	2.21	2.29	9.00	7.74	44.68	37.46	53.68	45.21	322.76				
YJL037A	IRC16	1.08	1.59	1.76	2.08	2.20	2.33	9.89	5.57	48.60	31.13	58.50	37.11	321.62				
YJL038C	LCH1	1.14	1.58	1.73	2.05	2.18	2.27	10.92	9.36	51.05	44.74	61.87	54.11	312.31				
YJL043W	MPM1	0.87	1.22	1.30	2.03	2.07	2.17	3.41	3.24	32.69	32.99	33.67	33.78	302.28				
YJL043W	YJL043w	0.93	1.31	1.46	1.93	2.12	2.23	6.60	3.56	38.91	21.56	46.51	25.96	292.27				
YJL044C	GVP2	0.89	1.31	1.45	1.94	2.12	2.23	14.18	4.87	53.84	27.78	67.94	22.65	336.19				
YJL045W	YJL045w	0.94	1.41	1.55	1.93	2.10	2.19	10.05	10.34	47.82	48.10	57.67	58.44	306.61				
YJL046W	AHM2	0.94	1.46	1.58	2.00	2.22	2.20	13.99	9.17	50.62	41.75	64.61	59.93	327.14				
YJL047C	RTT101	0.93	1.42	1.55	1.95	2.12	2.22	20.17	12.11	68.04	49.55	88.21	61.66	333.01				
YJL047C-A	YJL047c-a	1.00	1.35	1.54	1.92	2.10	2.22	6.23	1.57	48.90	20.24	55.13	21.81	301.88				
YJL048C	UBX6	0.94	1.48	1.67	2.05	2.18	2.28	3.11	2.29	12.99	20.04	15.40	23.89	271.07				
YJL049W	YJL049w	0.99	1.51	1.69	2.10	2.25	2.32	3.91	7.44	24.31	38.93	28.22	46.37	277.08				
YJL051W	IRC8	0.99	1.56	1.74	2.11	2.22	2.30	7.88	4.97	44.11	28.56	51.99	33.52	297.39				
YJL052W	TDH1	0.98	1.55	1.76	2.15	2.26	2.34	4.04	6.72	25.27	34.08	29.30	40.80	294.04				
YJL053W	PEP8	0.98	1.49	1.63	1.92	2.05	2.14	12.30	10.17	54.77	46.77	67.07	56.94	342.46				
YJL055W	YJL055w	1.03	1.58	1.75	2.08	2.18	2.30	5.20	3.27	32.00	15.21	37.20	18.48	293.88				
YJL056C	ZAP1	1.00	1.36	1.33	1.39	1.51	1.69	56.33	35.89	103.33	84.56	159.67	120.46	348.73				
YJL057C	IKS1	0.96	1.25	1.37	1.86	2.07	2.22	19.02	12.19	59.13	49.26	78.15	61.45	348.12				
YJL058C	BT161	0.87	1.27	1.39	1.78	1.96	2.12	13.66	9.00	52.92	42.13	66.58	51.13	336.02				
YJL059W	YHC3	0.93	1.38	1.58	1.98	2.14	2.25	14.46	9.66	54.28	42.19	68.74	51.86	340.79				
YJL060W	BNA3	0.94	1.37	1.55	1.96	2.11	2.23	4.71	3.76	31.06	21.25	35.77	25.33	313.40				
YJL062W	LAS21	1.00	1.48	1.69	2.06	2.21	2.29	3.34	3.08	14.71	7.69	18.05	10.78	277.13				
YJL062W-A	RRG10	0.95	1.06	1.07	0.92	0.91	0.90	120.41	125.11	137.40	105.07	257.80	230.18	488.23				
YJL063C	MRPL8	0.98	1.20	1.17	1.08	1.09	1.04	101.20	99.82	105.81	105.35	209.70	205.27	529.95				
YJL064W	YJL064w	0.98	1.50	1.72	2.01	2.15	2.24	11.29	16.41	48.93	56.33	60.22	72.74	305.82				
YJL065C	DLS1	0.95	1.34	1.49	1.84	2.01	2.14	12.96	16.61	54.64	57.04	67.60	73.65	316.32				
YJL066C	MPM1	0.97	1.30	1.48	1.80	1.99	2.10	6.17	11.16	39.73	47.33	45.90	58.49	304.51				
YJL067W	YJL067w	0.99	1.42	1.58	1.92	2.10	2.21	8.49	9.17	44.34	42.84	52.82	52.11	311.78				
YJL068C	YJL068c	0.95	1.32	1.53	1.87	2.04	2.13	21.07	15.72	60.55	53.33	81.62	69.06	369.49				
YJL070C	YJL070c	0.93	1.15	1.38	1.81	2.01	2.15	3.57	5.39	37.31	40.88	47.41	52.34					
YJL071W	ARG2	0.92	1.28	1.38	1.64	1.85	2.10	1.71	1.21	4.06	1.74	5.17	2.95	260.36				
YJL073W	JET1	0.94	1.30	1.55	1.93	2.06	2.18	2.03	1.53	44.84	9.32	54.07	42.16	68.55	51.47	336.41		
YJL074C	YJL074c	0.95	1.19	1.30	1.88	2.01	2.17	1.23	1.23	4.68	44.85	2.48	42.18	3.15	223.64			
YJL075C	ASF1	0.95	1.25	1.28	1.35	1.57	1.79	5.57	5.46	34.49	32.49	40.06	37.95	304.46				
YJL076C	NC43	0.95	1.33	1.49	1.65	2.06	2.21	1.32	1.20	4.54	52.28	47.48	52.33	336.96				
YJL077W-B	YJL077w-b	0.94	1.28	1.47	1.83	2.14	2.23	9.66	3.58	56.92	27.32	66.58	52.82	303.77				
YJL078C	PYV2	0.99	1.28	1.54	1.96	2.14	2.24	1.43	1.49	4.88	28.11	39.95	26.53	44.74	237.19			
YJL079C	PBY1	0.90	1.50	1.69	2.03	2.17	2.28	10.78	4.85	49.26	27.71	60.04	32.56	321.43				
YJL080C	SCP160	0.99	1.34	1.43	1.73	1.90	2.05	21.78	13.71	65.86	52.55	87.64	66.26	298.30				
YJL082W	IM2	0.94	1.34	1.46	1.76	1.94	2.12	5.22	3.15	28.32	10.41	33.54	13.55	296.85				
YJL083W	TAX4	0.97	1.33	1.47	1.79	1.96	2.11	5.57	5.46	34.49	32.49	40.06	37.95	304.46				
YJL084C	ALY2	0.96	1.44	1.65	1.97	2.12	2.22	10.20	7.75	48.31	40.02	58.52	47.78	327.76				
YJL088W	ARG3	0.85	1.15	1.27	1.65	1.86	2.02	5.85	4.61	15.67	52.26	16.99	16.99	286.49				
YJL089W	ARG3	0.89	1.24	1.28	1.61	1.88	2.10	4.47	1.34	45.50	11.78	49.97	13.15	300.38				
YJL090W	SIP4	0.99	1.47	1.65	1.98	2.14	2.23	15.51	9.28	57.09	43.18	72.60	52.46	343.48				
YJL092W	SRS2	0.92	1.24	1.38	1.62	1.79	1.91	19.36	14.25	59.75	53.33	79.11	67.58	356.34				
YJL093C	TOK1	0.98	1.39	1.63	1.96	2.14	2.21	22.33	14.63	59.99	51.48	82.32	66.11	363.85				
YJL094C	KHA1	0.93	1.17	1.39	1.83	2.03	2.17	0.81	1.38	5.47	19.83	6.27	21.21	5.23	253.59			
YJL095W	BC1K	1.02	1.33	1.40	1.89	2.10	2.27	26.18	16.18	69.72	58.84	95.89	75.02	334.92				
YJL096W	MRPL49	0.96	1.18	1.14	1.38	1.84	2.11	27.78	14.88	64.66	53.11	92.44	67.98	337.35				
YJL096W	MRPL49	0.99	1.15	1.17	1.05	1.05	1.04	121.61	117.17	137.96	101.96	259.57	219.13	505.61				
YJL097W	SAP165	0.97	1.46	1.65	2.03	2.17	2.27	15.61	5.42	56.33	31.35	71.94	36.77	345.35				
YJL099W	CHS6	0.98	1.41	1.53	1.83	1.99	2.12	15.65	7.68	54.00	37.85	70.05	45.54	337.26				
YJL100W	LSB6	0.93	1.34	1.45	1.79	1.96	2.11	12.03	5.79	51.86	36.30	63.89	38.70	330.15				
YJL101C	GSH1	0.90	1.14	1.25	1.58	1.77	1.89	8.52	2.95	46.07	27.56	54.60	30.51	309.53				
YJL102W	GSH1	0.87	1.18	1.28	1.63	1.85	2.02	8.23	1.84	48.98	19.56	57.21	21.40	295.67				
YJL103C	GSM1	0.95	1.27	1.38	1.89	2.08	2.18	11.46	4.44	49.59	28.66	61.04	33.10	289.89				
YJL104W	MEF2	0.96	1.15	1.14	1.01	1.03	1.09	102.27	10.41	21.60	45.07	67.47	27.07	307.27				
YJL105W	SET4	0.88	1.02	1.16	1.48	1.70	1.89	1.64	1.58	2.58	25.71	28.52	27.35	31.10	294.86	</		

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YJL171C	YJL171c	1.04	1.42	1.64	2.10	2.24	2.41	1.22	1.25	13.39	7.35	14.61	8.59	282.04		
YJL172W	CPS1	1.02	1.42	1.64	2.08	2.19	2.33	3.40	1.41	34.93	14.25	38.33	15.66	292.21		
YJL172W	CPS1	0.88	1.13	1.23	1.65	1.90	2.08	8.14	2.17	51.48	24.94	59.62	27.10	316.40		
YJL175W	YJL175w	0.93	1.11	1.29	1.71	1.90	2.07	8.44	1.88	54.42	29.94	62.86	31.83	376.58		
YJL175W	YJL175w	0.97	1.33	1.51	2.00	2.04	2.12	2.59	2.42	35.53	27.68	38.12	30.10	342.55		
YJL176C	SW3	1.02	1.13	1.18	1.48	1.59	1.72	21.71	10.57	80.39	64.04	102.10	74.91	412.64		
YJL177W	RPL17b	0.88	0.95	1.07	1.25	1.78	1.99	1.23	0.80	21.58	6.38	22.61	7.19	292.45		
YJL178C	ATG27	1.08	1.38	1.61	2.09	2.21	2.32	16.34	8.92	63.49	54.10	81.83	63.02	308.72		
YJL179W	ATP9	1.07	1.37	1.43	1.80	1.94	2.04	19.63	10.79	82.03	72.53	124.34	84.72	344.43		
YJL180C	ATP12	0.89	1.05	1.07	0.93	0.92	0.89	130.67	121.09	133.38	131.34	264.05	251.43	529.01		
YJL181W	YJL181w	1.02	1.29	1.47	1.91	2.07	2.25	6.73	1.22	47.61	13.91	54.34	15.13	315.89		
YJL182C	YJL182c	1.05	1.40	1.60	2.03	2.17	2.31	10.63	6.70	55.50	45.74	66.13	52.44	318.31		
YJL183W	MNH1	0.96	1.15	1.31	1.73	1.90	2.14	13.64	4.19	60.81	35.95	74.25	49.14	261.16		
YJL184W	GON7	0.83	1.08	1.06	1.63	1.20	1.41	37.35	19.61	86.67	70.46	124.02	90.07	322.48		
YJL185C	YJL185c	1.02	1.41	1.62	2.03	2.16	2.30	6.81	1.40	46.01	15.76	52.63	17.16	311.38		
YJL186W	MNH5	1.02	1.38	1.57	1.95	2.10	2.26	7.00	3.09	49.35	31.08	56.34	34.16	311.38		
YJL187C	SWE1	1.04	1.34	1.52	1.93	2.08	2.27	3.79	1.31	36.35	14.57	40.14	15.88	296.29		
YJL188C	BUD19	1.01	1.10	1.14	1.54	1.84	2.15	39.36	22.89	93.96	74.08	133.33	96.98	347.78		
YJL189W	RPL39	1.02	1.38	1.61	2.06	2.19	2.33	10.20	2.31	56.77	27.41	66.97	29.72	309.26		
YJL190C	RPS22a	0.88	1.10	1.11	1.11	1.33	1.56	11.47	3.12	63.41	34.49	76.89	37.62	258.82		
YJL191W	RPS14b	1.02	1.27	1.43	1.97	2.12	2.24	1.29	1.08	19.41	5.68	20.70	6.77	292.16		
YJL191W	RPS14d	0.97	1.14	1.32	1.81	2.01	2.19	11.22	3.31	57.82	33.33	69.04	36.64	304.93		
YJL192C	SOP4	0.99	1.15	1.28	1.86	1.95	2.15	5.57	1.81	44.65	23.79	50.22	25.61	300.45		
YJL193W	YJL193w	1.04	1.34	1.55	1.96	2.10	2.27	5.60	2.24	45.10	28.48	50.70	30.71	301.79		
YJL196C	EL01	1.03	1.35	1.56	1.99	2.14	2.29	0.89	1.06	9.54	3.32	10.44	4.38	257.19		
YJL196C	EL01	0.91	1.08	1.26	1.68	1.87	2.04	7.60	1.73	50.30	25.79	57.90	27.02	300.46		
YJL197W	UBP12	0.97	1.26	1.45	1.87	2.03	2.22	4.64	0.97	39.92	15.83	44.57	16.80	288.72		
YJL198W	PHO90	0.97	1.27	1.47	1.87	2.00	2.18	2.29	1.69	27.71	19.60	30.01	21.29	280.66		
YJL199C	YJL199c	0.96	1.25	1.45	1.86	2.05	2.25	1.69	1.14	23.35	9.31	25.04	10.46	292.84		
YJL200C	ACO2	0.97	1.28	1.49	1.91	2.07	2.28	1.00	1.06	8.57	1.47	9.56	2.53	277.48		
YJL200C	ACO2	0.89	1.06	1.24	1.73	1.92	2.11	3.06	0.77	33.22	8.26	36.28	9.04	309.69		
YJL201W	ECM25	0.99	1.19	1.34	1.75	1.90	2.12	13.66	5.41	61.53	43.08	75.19	48.50	322.61		
YJL204C	RCY1	0.98	1.23	1.36	1.86	1.99	2.15	18.22	8.84	65.57	47.77	84.19	56.60	367.64		
YJL206C	YJL206c	1.04	1.28	1.48	2.00	2.15	2.29	9.90	6.91	55.67	47.52	65.77	54.43	313.04		
YJL206C	YJL206c-A	0.91	1.08	1.24	1.73	1.93	2.12	9.48	1.70	55.17	24.37	64.65	26.07	302.43		
YJL207C	LAP1	0.93	1.14	1.23	1.65	1.79	2.02	2.22	2.34	5.24	1.62	42.73	19.01	47.97	20.83	
YJL208C	YJL208c	0.97	1.13	1.24	1.68	1.87	2.08	1.70	1.79	15.59	23.15	60.50	31.10	311.30		
YJL209C	CGU1	0.97	1.13	1.24	1.68	1.87	2.08	21.76	5.14	63.91	47.52	71.52	34.07	340.07		
YJL209W	CP91	0.85	1.12	1.25	1.71	1.89	2.07	1.76	1.77	35.55	37.69	31.59	14.13	256.56		
YJL210W	PEK2	0.90	1.13	1.24	1.68	1.87	2.08	4.97	1.16	41.73	22.26	47.67	24.03	270.76		
YJL211C	YJL211c	0.96	1.20	1.38	1.84	2.02	2.21	5.61	2.29	44.79	28.85	50.41	20.94	279.47		
YJL212C	OPT1	0.93	1.30	1.50	2.00	2.17	2.33	1.84	1.11	28.12	17.29	29.96	18.40	266.80		
YJL213W	YJL213w	0.84	0.94	1.07	1.50	1.75	2.21	1.93	0.90	13.81	5.24	14.83	6.15	297.40		
YJL214W	HXT8	0.92	1.11	1.24	1.70	1.90	2.11	5.26	1.54	44.12	22.87	49.38	24.42	294.13		
YJL215C	YJL215c	0.95	1.17	1.31	1.77	1.94	2.14	4.64	1.28	41.19	18.46	45.83	19.74	284.10		
YJL216C	YJL216c	0.88	1.04	1.16	1.59	1.76	1.98	3.33	2.78	35.76	36.33	39.09	39.11	283.78		
YJL217W	REE1	0.98	1.22	1.38	1.81	1.97	2.16	4.44	1.11	40.36	14.70	44.79	15.81	282.45		
YJL218W	YJL218w	0.92	1.25	1.41	1.89	2.07	2.26	3.69	1.23	37.22	15.81	40.92	17.04	291.33		
YJL201W	AVT1	0.90	1.25	1.36	1.68	1.89	2.01	15.27	2.92	71.29	36.79	86.56	39.71	301.98		
YJL003C	YJL003c	0.93	1.12	1.31	1.80	1.98	2.15	8.04	1.91	55.11	26.64	63.15	28.55	311.01		
YJL004C	SAG1	0.93	1.01	1.04	0.94	0.90	0.94	129.95	127.64	146.64	110.57	276.59	238.21	396.08		
YJL005C-A	YJL005c-a	0.96	1.21	1.26	1.65	1.89	2.04	5.37	0.88	46.77	10.80	52.14	11.68	301.70		
YJL005W	APL1	0.89	1.24	1.36	1.66	1.85	1.98	7.78	1.56	52.92	26.87	60.71	28.43	285.43		
YJL008W	YJL008w	0.90	1.22	1.38	1.78	1.97	2.09	8.20	1.29	58.29	23.99	66.48	25.28	271.67		
YJL009C	TDH2	0.87	1.16	1.29	1.61	1.81	1.97	9.80	5.15	59.70	52.02	69.49	57.18	239.98		
YJL010C-A	SPC1	1.00	1.49	1.66	2.06	2.21	2.28	5.48	2.87	47.89	36.91	53.37	39.78	248.57		
YJL010W	MET3	0.98	1.49	1.67	2.05	2.20	2.29	3.68	1.68	42.34	28.46	46.03	30.15	289.41		
YJL011C	YJL011c	0.95	1.35	1.48	1.94	2.12	2.23	13.74	4.93	71.86	51.25	85.60	56.18	299.08		
YJL014W	TMA22	0.98	1.42	1.59	2.00	2.17	2.25	13.74	0.68	69.71	8.78	83.45	9.47	311.54		
YJL015W	YJL015w	0.96	1.46	1.68	2.06	2.22	2.28	9.48	0.62	58.91	9.47	68.39	10.09	307.84		
YJL018W	TEST1	0.94	1.18	1.18	1.23	1.42	1.54	78.35	56.32	131.77	116.59	210.12	172.91	345.65		
YJL020W	YJL020w	0.95	1.33	1.48	1.86	1.99	2.10	6.26	0.74	50.47	16.42	64.22	16.73	309.08		
YJL021C	REC107	0.90	1.28	1.43	1.80	1.99	2.07	11.25	2.05	64.25	21.21	75.48	33.58	318.32		
YJL024C	MDE1	0.92	1.21	1.32	1.65	1.81	2.01	4.00	1.28	41.58	10.37	48.48	11.22	217.41		
YJL025W	BN1A	0.94	1.11	1.19	1.52	1.85	2.11	11.70	0.90	54.89	8.43	66.59	9.33	305.10		
YJL025W	HT1	0.96	1.11	1.19	1.57	1.81	2.01	4.00	1.28	41.58	9.93	45.58	11.22	283.59		
YJL026C	YJL026c	0.89	1.09	1.13	1.27	1.53	1.81	4.00	1.28	24.65	10.51	77.91	45.89	192.57	349.25	
YJL026C	YJL026c	0.86	1.17	1.26	1.65	1.83	2.05	2.24	1.17	30						

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE			FRUCTOSE			TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YJR099W	YU1H	0.97	1.31	1.48	1.87	2.06	2.19	6.41	4.72	34.66	24.35	41.07	29.06	306.67	306.67	306.67	306.67	
YJR100C	AIM25	0.99	1.36	1.55	1.99	2.15	2.23	20.88	18.91	55.08	70.76	75.96	89.67	163.14	163.14	163.14	163.14	
YJR101W	RSM26	0.89	1.04	1.03	0.91	0.91	0.89	126.21	120.06	136.65	99.92	262.86	219.97	496.45	496.45	496.45	496.45	
YJR102C	VPS25	0.99	1.46	1.57	1.83	1.94	2.02	20.77	8.96	72.19	58.00	92.96	66.96	386.57	386.57	386.57	386.57	
YJR103W	URA6	1.03	1.56	1.73	2.10	2.22	2.31	11.40	3.13	48.56	9.70	59.96	12.83	320.89	320.89	320.89	320.89	
YJR104C	SOD1	1.00	1.33	1.45	1.89	2.11	2.23	28.52	14.43	74.24	64.20	102.76	78.62	232.40	232.40	232.40	232.40	
YJR105W	ADG1	1.00	1.38	1.47	1.77	1.95	2.11	28.68	4.86	76.59	25.85	99.07	30.70	299.42	299.42	299.42	299.42	
YJR106W	ECM27	0.97	1.47	1.72	2.13	2.26	2.33	16.68	7.47	62.49	59.72	78.17	67.20	272.19	272.19	272.19	272.19	
YJR107W	YU1H	0.97	1.41	1.54	1.45	1.54	1.62	2.43	1.21	25.18	1.42	25.59	5.74	224.4	224.4	224.4	224.4	
YJR107W	YU1H	0.95	1.47	1.69	2.08	2.21	2.28	15.39	1.27	26.95	45.73	76.35	53.00	297.77	297.77	297.77	297.77	
YJR108W	ABM1	1.08	1.60	1.77	2.11	2.24	2.34	4.83	3.04	33.05	4.45	37.88	7.50	310.80	310.80	310.80	310.80	
YJR109C	CPA2	0.95	1.32	1.40	1.71	1.84	2.09	1.30	1.35	12.77	2.88	14.07	4.23	267.98	267.98	267.98	267.98	
YJR110W	YMR1	1.14	1.64	1.81	2.19	2.31	2.42	3.08	3.02	6.37	2.14	9.45	5.17	271.91	271.91	271.91	271.91	
YJR111C	YJR111c	0.96	1.50	1.67	2.05	2.21	2.31	3.27	3.06	14.77	2.47	18.04	5.53	278.15	278.15	278.15	278.15	
YJR112W-A	YU1H	0.90	1.20	1.38	1.81	2.09	2.02	6.48	1.92	48.34	10.49	54.81	11.51	297.25	297.25	297.25	297.25	
YJR113C	RSM7	0.97	1.33	1.42	1.82	2.03	2.18	34.09	28.65	78.26	84.17	112.36	112.82	309.37	309.37	309.37	309.37	
YJR113C	RSM7	0.91	1.19	1.24	1.47	1.65	1.77	51.09	37.47	104.72	95.90	155.81	132.37	335.88	335.88	335.88	335.88	
YJR114W	YJR114w	0.89	1.05	1.05	0.92	0.92	0.90	123.87	118.21	135.62	98.79	259.49	217.00	502.10	502.10	502.10	502.10	
YJR115W	YU1H	1.02	1.55	1.72	2.06	2.20	2.28	7.40	3.06	43.43	6.91	50.83	9.98	305.13	305.13	305.13	305.13	
YJR116W	YU1H	0.95	1.45	1.67	2.05	2.19	2.27	14.75	11.56	58.20	55.81	72.95	67.17	281.98	281.98	281.98	281.98	
YJR117W	STE24	0.93	1.42	1.60	2.05	2.18	2.28	4.77	4.70	38.65	43.84	43.42	51.24	267.31	267.31	267.31	267.31	
YJR118C	ILM1	0.94	1.21	1.33	1.68	1.91	2.14	1.17	0.93	0.88	0.15	2.04	1.08	232.28	232.28	232.28	232.28	
YJR119C	JHD2	0.92	1.30	1.42	1.88	2.07	2.20	28.56	15.57	71.60	64.41	100.16	79.98	205.73	205.73	205.73	205.73	
YJR120W	YU1H	0.89	1.17	1.26	1.60	1.79	1.95	14.29	2.92	68.51	35.89	82.80	38.82	326.78	326.78	326.78	326.78	
YJR121W	ATP2	0.96	1.28	1.40	1.93	2.11	2.23	17.63	10.05	68.25	56.13	85.88	66.18	231.22	231.22	231.22	231.22	
YJR122W	IBA57	0.90	1.20	1.28	1.69	1.94	2.08	14.56	7.06	66.57	47.78	81.13	54.84	303.19	303.19	303.19	303.19	
YJR124C	YU1H	0.95	1.32	1.56	1.95	2.11	2.22	29.83	15.99	80.84	68.56	110.67	84.55	257.68	257.68	257.68	257.68	
YJR125C	ENT3	0.98	1.40	1.66	2.05	2.21	2.30	21.96	7.20	74.17	52.64	96.13	60.85	233.20	233.20	233.20	233.20	
YJR126C	VPS70	1.03	1.46	1.69	2.10	2.24	2.31	22.03	13.24	73.25	64.18	95.28	77.42	281.53	281.53	281.53	281.53	
YJR127C	RSF2	0.97	1.45	1.66	2.08	2.24	2.31	9.06	3.04	48.37	6.67	57.43	9.72	302.65	302.65	302.65	302.65	
YJR128W	YU1H	0.95	1.43	1.65	2.06	2.20	2.28	8.18	4.16	44.92	49.84	18.21	18.21	302.24	302.24	302.24	302.24	
YJR129C	YU1H	0.88	1.25	1.39	1.82	2.01	2.12	8.44	3.60	47.40	20.45	55.84	24.05	298.24	298.24	298.24	298.24	
YJR130C	STR2	1.00	1.39	1.53	1.98	2.17	2.27	12.39	4.67	57.03	34.25	69.42	38.92	321.47	321.47	321.47	321.47	
YJR131W	MNV1	0.95	1.35	1.54	1.95	2.12	2.23	22.83	13.94	69.63	60.56	92.46	74.49	275.44	275.44	275.44	275.44	
YJR133W	XPT1	0.99	1.43	1.71	2.11	2.25	2.33	17.16	7.23	62.39	29.32	79.55	32.26	265.06	265.06	265.06	265.06	
YJR134C	SGM1	0.93	1.31	1.46	1.83	2.02	2.16	23.92	16.16	71.76	61.80	95.68	77.96	314.26	314.26	314.26	314.26	
YJR135C	MCM22	1.06	1.43	1.48	1.88	2.10	2.23	32.87	16.85	75.27	63.94	108.14	80.80	371.58	371.58	371.58	371.58	
YJR136C	YU1H	0.98	1.24	1.35	1.82	2.05	2.16	12.65	4.28	6.89	44.63	44.21	111.00	111.00	279.45	279.45	279.45	279.45
YJR137C	MET6	0.97	1.44	1.61	2.02	2.20	2.29	7.67	3.05	41.02	8.78	49.17	11.75	311.18	311.18	311.18	311.18	
YJR140C	HR3	0.90	1.26	1.36	1.77	1.97	2.06	36.31	13.62	82.39	63.49	119.74	72.11	211.43	211.43	211.43	211.43	
YJR142W	YU1H	0.91	1.25	1.38	1.86	2.06	2.20	29.29	18.69	76.96	68.13	106.25	86.82	181.37	181.37	181.37	181.37	
YJR143C	PMT4	0.91	1.16	1.37	1.81	1.99	2.10	8.89	1.38	53.22	18.51	61.11	16.89	398.44	398.44	398.44	398.44	
YJR144W	MGH101	0.88	1.24	1.28	1.67	2.00	2.17	2.27	4.90	1.27	47.09	11.16	52.00	12.43	244.98	244.98	244.98	244.98
YJR145C	RPS4a	0.94	1.26	1.38	1.87	2.04	2.14	39.89	26.84	88.24	82.71	128.13	109.35	219.35	219.35	219.35	219.35	
YJR146W	YU1H	0.97	1.50	1.70	2.08	2.21	2.29	7.79	3.08	45.68	9.71	53.46	12.80	309.68	309.68	309.68	309.68	
YJR147W	HMS2	1.00	1.54	1.72	2.08	2.21	2.31	6.21	3.05	39.34	7.25	45.55	10.31	302.07	302.07	302.07	302.07	
YJR148W	BAT2	0.95	1.26	1.48	1.87	2.06	2.17	22.69	20.18	78.64	72.91	101.34	93.10	213.43	213.43	213.43	213.43	
YJR149W	YU1H	0.91	1.52	1.75	2.12	2.25	2.32	4.87	3.03	32.99	5.87	37.85	8.89	278.38	278.38	278.38	278.38	
YJR150C	DAN1	1.00	1.30	1.51	1.88	2.06	2.17	2.17	2.17	26.81	14.07	78.60	62.91	105.21	76.98	257.45	257.45	257.45
YJR151C	DAN4	0.94	1.30	1.51	1.90	2.05	2.20	4.69	2.64	45.10	34.25	49.79	36.89	297.45	297.45	297.45	297.45	
YJR151C	MAE1	0.97	1.23	1.41	1.89	2.05	2.20	19.22	15.32	72.51	68.49	91.72	83.80	329.51	329.51	329.51	329.51	
YJR152W	HCS1	1.03	1.34	1.57	2.07	2.23	2.32	5.52	5.67	43.93	43.50	49.45	49.17	298.51	298.51	298.51	298.51	
YJR152C-A	YU1H	1.00	1.37	1.57	1.89	2.09	2.28	7.40	3.18	53.74	34.11	61.14	37.29	315.86	315.86	315.86	315.86	
YJR153W	SP72	0.98	1.30	1.51	1.97	2.09	2.28	7.40	3.18	53.74	34.11	61.14	37.29	316.02	316.02	316.02	316.02	
YKL009W	YKL031w	1.02	1.30	1.52	1.95	2.11	2.24	2.51	1.43	31.36	21.12	33.87	22.55	290.45	290.45	290.45	290.45	
YKL032C	IXR1	1.00	1.41	1.70	2.17	2.31	2.44	1.06	0.97	12.72	8.27	3.38	2.38	252.02	252.02	252.02	252.02	
YKL033W-A	YKL033w-a	0.95	1.34	1.48	1.97	1.98	2.08	15.37	5.79	62.38	37.49	77.75	43.28	314.34	314.34	314.34	314.34	
YKL03																		

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YKL099W	MTC2	0.95	1.30	1.52	1.98	2.11	2.17	6.87	3.00	46.12	29.48	52.99	32.48	314.76		
YKL100C	YKL100c	1.01	1.39	1.62	2.06	2.21	2.30	2.94	1.23	33.78	14.81	36.72	16.04	299.30		
YKL101W	HSL1	0.92	1.22	1.30	1.64	1.89	2.07	12.59	5.45	60.06	40.56	72.64	46.01	293.33		
YKL102C	YKL102c	0.94	1.25	1.42	1.92	2.11	2.23	0.72	0.95	11.15	3.38	11.88	4.33	275.22		
YKL102c	LAP4	0.97	1.25	1.43	1.90	2.09	2.20	1.64	1.16	23.02	19.02	24.66	20.18	272.72		
YKL105c	YKL105c	1.00	1.22	1.38	1.89	2.08	2.19	4.98	5.16	47.24	45.04	52.21	50.20	293.35		
YKL106c-A	YKL106c-a	1.00	1.30	1.50	1.99	2.14	2.27	4.13	1.62	42.41	25.43	46.54	27.25	298.46		
YKL106c	AATF1	1.00	1.30	1.50	1.99	2.14	2.27	4.13	1.62	42.41	25.43	46.54	27.25	298.46		
YKL107W	YKL107w	0.94	1.22	1.41	1.95	2.12	2.22	6.04	1.43	49.53	24.01	52.00	23.11	311.41		
YKL109W	HAP2	0.94	1.23	1.47	2.01	2.19	2.28	2.58	1.28	31.85	14.01	34.43	15.29	293.03		
YKL110C	KTH12	1.01	1.34	1.54	2.03	2.19	2.28	10.38	7.19	61.03	49.99	71.41	57.18	306.92		
YKL113C	RAD27	0.90	1.07	1.13	1.50	1.70	1.84	14.11	10.46	69.57	52.05	83.69	72.51	322.39		
YKL114C	APN1	1.00	1.31	1.53	2.06	2.22	2.31	2.32	1.89	30.64	24.73	32.97	26.72	291.66		
YKL115C	YKL115c	0.90	1.13	1.37	1.92	2.12	2.29	1.15	1.26	13.89	4.21	15.04	5.47	274.04		
YKL116C	PBR1	0.93	1.19	1.33	1.82	2.03	2.15	2.06	1.16	28.35	11.74	30.41	12.90	296.44		
YKL117W	SB41	0.98	1.36	1.58	2.12	2.26	2.33	1.25	1.11	15.16	5.64	16.41	6.75	270.15		
YKL118W	YKL118w	0.89	1.08	1.09	1.10	1.42	1.73	40.01	18.93	96.63	76.84	136.64	95.77	333.75		
YKL119C	VPH2	0.92	1.08	1.10	1.06	1.28	1.56	40.84	25.82	99.46	84.19	140.30	110.00	327.23		
YKL120W	OAC1	0.90	1.12	1.27	1.76	1.95	2.08	1.49	0.97	22.36	17.92	23.85	18.89	290.90		
YKL121W	DGR2	0.96	1.16	1.28	1.76	1.97	2.10	4.64	3.20	45.67	37.88	50.31	41.08	289.83		
YKL123W	YKL123w	1.01	1.40	1.62	2.07	2.22	2.31	1.49	1.28	21.37	6.41	22.85	7.69	289.22		
YKL124W	SSH4	0.85	1.03	1.14	1.66	1.88	2.02	1.20	1.11	12.27	6.56	13.47	7.67	271.26		
YKL126W	YPK1	0.89	1.15	1.29	1.75	1.93	2.02	7.61	2.36	51.47	19.70	59.08	31.41	309.54		
YKL127W	PGM1	0.95	1.26	1.47	1.92	2.08	2.16	2.67	2.26	31.23	29.18	33.90	31.44	284.46		
YKL128C	PMM1	0.90	1.19	1.39	1.87	2.03	2.13	1.37	1.58	17.24	20.85	18.61	22.43	284.54		
YKL129C	MYO3	0.93	1.19	1.37	1.89	2.07	2.16	2.31	1.45	30.83	18.75	33.14	20.20	283.15		
YKL130C	SHE2	0.92	1.23	1.46	1.98	2.15	2.24	2.14	1.17	28.92	15.42	31.06	16.59	289.68		
YKL131W	YKL131w	0.87	1.06	1.18	1.60	1.79	1.90	3.01	1.09	36.76	9.20	39.77	10.29	295.52		
YKL132C	RM1A1	0.96	1.24	1.42	1.83	2.00	2.10	1.51	1.01	22.28	8.69	23.79	9.10	291.92		
YKL133C	YKL133c	0.85	1.04	1.14	1.74	1.95	2.08	4.74	1.05	46.59	18.03	51.33	19.08	292.55		
YKL134C	OCT1	0.88	1.01	0.99	1.03	1.05	1.01	151.35	96.27	154.08	117.90	305.43	214.17	523.91		
YKL135C	APL2	0.96	1.13	1.17	1.60	1.86	2.03	14.10	8.19	70.94	55.91	85.04	64.11	340.27		
YKL136W	YKL136w	0.93	1.16	1.34	1.81	1.99	2.12	3.79	0.92	41.86	10.56	45.85	11.49	297.66		
YKL137W	CMT1	0.95	1.25	1.45	1.92	2.10	2.19	3.14	1.72	36.42	22.18	39.55	23.91	299.44		
YKL137W	CMT1	0.92	1.21	1.34	1.67	1.87	2.03	4.72	0.94	41.76	7.57	46.48	8.51	279.16		
YKL138C	MRPL11	0.92	1.19	1.37	1.84	2.02	2.11	2.80	2.95	32.64	33.37	35.44	36.32	283.33		
YKL139W	CTKT	0.87	0.67	0.88	0.98	1.11	1.24	81.67	72.19	112.92	107.07	194.80	178.90	386.46		
YKL140W	TGL1	0.93	1.22	1.36	1.92	2.05	2.28	1.26	1.28	43.33	11.67	15.53	12.87	290.86		
YKL142W	MRP29	0.82	1.07	1.26	1.66	2.11	2.28	1.25	1.28	23.06	12.07	31.25	16.54	285.44		
YKL143W	LTIV	0.93	1.18	1.22	1.44	1.78	2.08	1.41	1.21	24.22	12.79	25.53	14.00	223.03		
YKL149W	AVT3	0.98	1.18	1.35	1.77	1.95	2.05	1.28	1.25	15.76	16.27	17.04	17.51	275.12		
YKL147C	YKL147c	0.97	1.27	1.43	1.87	2.08	2.20	1.43	1.29	13.26	4.66	14.39	5.06	273.42		
YKL148C	SDH7	0.99	1.29	1.50	1.97	2.14	2.24	1.54	1.27	21.74	6.57	23.27	7.54	271.94		
YKL149C	DRB1	0.94	1.21	1.36	1.94	2.12	2.22	11.67	11.41	60.55	59.96	72.22	71.37	312.31		
YKL150W	MCR1	0.94	1.22	1.37	1.79	1.94	2.03	1.58	1.17	23.35	21.01	24.93	22.18	288.18		
YKL151C	YKL151c	0.99	1.21	1.35	1.86	2.08	2.20	3.56	2.48	39.45	31.35	43.01	33.82	277.58		
YKL152C	RSM22	0.87	1.05	1.01	0.90	0.90	0.86	120.43	119.97	139.90	106.93	260.33	228.89	468.67		
YKL156W	RPS27a	0.93	1.21	1.36	1.88	2.09	2.22	1.82	2.74	31.12	21.63	33.94	20.95	281.38		
YKL157W	APE2	0.96	1.35	1.55	2.00	2.14	2.24	2.52	2.04	30.88	25.88	33.49	28.02	288.44		
YKL157W	APE2	0.98	1.37	1.58	1.97	2.11	2.22	7.47	1.41	52.15	16.77	59.62	18.18	301.28		
YKL158W	YKL158w	0.93	1.25	1.46	2.06	2.13	2.08	2.75	2.03	32.71	27.05	35.47	29.08	292.45		
YKL159C	RCN1	0.98	1.37	1.56	2.02	2.17	2.25	2.32	1.99	30.16	19.33	32.48	21.32	292.70		
YKL160W	ELF1	1.04	1.40	1.59	2.07	2.25	2.38	0.92	0.93	7.62	2.09	8.54	3.02	290.78		
YKL161C	KDX1	0.95	1.24	1.42	1.87	2.04	2.15	1.87	1.33	24.57	19.64	26.44	20.97	286.57		
YKL162C	YKL162c	0.96	1.31	1.56	2.05	2.20	2.30	3.58	1.09	38.52	9.44	42.10	10.54	303.57		
YKL162C-A	YKL162c-a	1.02	1.49	1.73	2.11	2.23	2.30	8.99	3.96	50.53	24.64	59.52	28.60	323.28		
YKL163W	PIR3	0.91	1.17	1.38	1.89	2.07	2.17	3.29	1.25	34.82	14.27	38.04	17.93	303.90		
YKL164C	PIR1	0.95	1.20	1.40	1.91	2.08	2.18	3.37	1.97	37.34	21.80	40.70	29.91	296.62		
YKL165C	STZ1	0.91	1.10	1.21	1.73	2.00	2.04	3.27	1.53	36.53	30.33	36.74	32.00	322.32		
YKL166C	YKL166c	0.98	1.22	1.36	1.78	1.98	2.08	6.98	5.98	48.72	47.49	55.78	52.40	312.24		
YKL168W	LOTS	0.93	1.10	1.22	1.81	2.05	2.19	1.23	1.13	13.87	11.63	15.09	12.76	288.80		
YKL169W	SPET1	1.02	1.35	1.60	2.10	2.27	2.35	5.35	4.09	45.44	10.23	45.89	11.33	316.16		
YKL183W	ASH1	0.94	1.26	1.46	1.84	2.11	2.20	5.41	1.94	40.89	22.23	46.21	24.16	309.45		
YKL187C	YKL187c	1.02	1.42	1.63	2.04	2.19	2.28	17.57	5.07	66.84	42.39	74.47	33.80	338.30		
YKL188C	PYK2	0.99	1.38	1.61	2.07	2.22	2.30	5.90	1.38	48.01	17.81	53.91	20.09	310.23		
YKL190W	CNB1	0.94	1.20	1.35	1.79	1.99	2.06	9.47	1.51	49.52	35.94	60.32	33.30	328.76		
YKL191W	DPH2	1.03	1.47	1.57	1.93	2.09	2.22	6.26	3.04	43.56	20.16	49.82	23.20	308.02		
YKL194C	MS71	1.04	1.23	1.25	1.90	2.10	2.19	12.22	12.17	133.86	132.65	260.08	254.35	541.76		
YKL197C	PEX1	0.92	1.38	1.48	1.93	2.08	2.19	14.84								

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YKR035C	Unknown	0.91	1.29	1.31	1.52	1.64	1.71	23.43	16.32	70.31	62.58	93.74	78.90	368.80		
YKR035W-A	DID2	1.00	1.47	1.69	2.05	2.16	2.23	8.89	5.52	51.44	40.38	60.33	45.89	343.49		
YKR036C	CAF4	0.93	1.16	1.39	1.95	2.15	2.30	1.17	0.70	21.45	4.66	22.63	5.36	296.04		
YKR039W	GAPT	0.86	1.00	1.17	1.69	1.88	2.06	9.76	1.66	54.32	21.85	64.09	23.51	312.20		
YKR040C	YKR040c	0.89	1.05	1.23	1.73	1.97	2.17	0.87	0.54	15.10	1.90	15.97	2.43	287.76		
YKR041W	YKR041w	1.05	1.31	1.54	2.05	2.18	2.32	18.30	1.12	69.06	20.38	87.37	21.50	342.12		
YKR042W	UTHT	0.91	1.28	1.48	2.05	2.08	2.17	21.31	14.48	66.53	56.97	87.84	71.45	334.63		
YKR043C	YKR043c	0.88	1.44	1.62	2.04	2.18	2.28	22.55	14.05	68.23	55.07	90.78	69.12	350.02		
YKR044C	YKR044c	0.95	1.56	1.49	1.63	1.93	2.04	15.40	1.65	62.64	51.11	62.27	52.22	321.14		
YKR045C	YKR045c	0.95	1.38	1.52	1.88	2.03	2.13	16.78	6.64	60.98	38.76	77.76	46.60	338.24		
YKR046C	PET10	0.96	1.18	1.33	1.86	2.03	2.19	11.49	1.85	58.17	28.17	69.66	30.02	317.80		
YKR047W	YKR047w	1.01	1.46	1.62	2.08	2.23	2.33	15.32	7.48	58.25	40.73	73.57	48.21	325.67		
YKR048C	NAP1	0.91	1.30	1.36	1.72	1.89	2.02	27.41	15.16	67.90	52.76	95.31	67.93	357.49		
YKR049C	FMP46	0.94	1.40	1.52	1.91	2.07	2.19	8.53	3.31	46.85	20.42	55.38	23.93	315.15		
YKR050W	TRK2	0.93	1.40	1.50	1.86	2.00	2.13	13.09	7.12	56.09	41.07	69.18	48.19	320.75		
YKR051W	YKR051w	0.98	1.46	1.57	1.95	2.13	2.25	13.03	9.58	54.18	46.22	67.21	55.80	328.72		
YKR052C	MRS4	0.93	1.32	1.42	1.93	2.09	2.21	9.00	5.72	48.50	34.63	57.50	40.35	308.14		
YKR053C	YSR3	0.96	1.18	1.40	1.87	2.04	2.21	10.92	1.01	56.85	19.73	67.76	20.74	302.66		
YKR054C	DYN1	0.99	1.42	1.49	1.89	2.05	2.18	18.37	13.41	65.76	56.56	84.12	69.97	327.15		
YKR054C	DYN1	0.99	1.51	1.24	1.54	1.74	1.90	19.37	5.59	80.87	38.06	100.24	43.65	306.44		
YKR055W	RHO4	1.03	1.48	1.67	1.98	2.10	2.17	5.78	3.19	35.67	11.88	41.46	15.07	338.29		
YKR056W	TRM2	0.99	1.43	1.57	1.94	2.12	2.22	22.58	13.57	66.74	53.33	89.32	66.90	364.26		
YKR057W	RPS21a	0.99	1.41	1.53	2.01	2.00	2.21	17.96	8.28	63.20	43.27	81.16	51.55	335.26		
YKR058W	GLG1	0.96	1.43	1.57	1.94	2.09	2.21	11.65	5.10	52.61	33.25	64.26	38.35	325.36		
YKR059W	TIF1	0.93	1.39	1.52	1.88	2.06	2.18	13.18	4.41	53.55	28.89	66.72	33.29	329.10		
YKR060W	UTP30	1.09	1.55	1.71	2.11	2.25	2.35	12.57	4.49	55.77	30.84	68.34	35.33	327.82		
YKR061W	KTR2	0.95	1.36	1.49	1.91	2.06	2.16	11.66	3.97	50.82	26.36	62.48	30.34	320.25		
YKR064W	OAF3	0.98	1.47	1.63	2.04	2.17	2.27	12.76	5.05	55.05	33.39	67.81	38.44	323.64		
YKR065C	PAM17	0.98	1.46	1.59	1.94	2.07	2.17	7.98	4.50	44.78	29.62	52.76	34.12	302.96		
YKR066C	CCP1	0.98	1.45	1.64	1.98	2.13	2.23	15.84	9.12	46.62	27.46	72.97	55.74	317.34		
YKR067W	GPT2	1.01	1.48	1.66	2.04	2.17	2.28	12.55	6.14	54.22	37.72	66.77	43.86	333.49		
YKR069W	MET1	0.99	1.53	1.70	2.06	2.17	2.28	9.45	3.65	57.91	35.91	56.73	42.26	328.81		
YKR070W	YKR070w	1.05	1.57	1.80	2.17	2.28	2.36	14.53	8.53	57.12	40.18	71.65	48.70	341.72		
YKR072C	SIS2	0.94	1.43	1.59	1.93	2.05	2.16	19.83	9.76	68.09	43.78	87.92	53.54	349.42		
YKR073C	YKR073c	0.97	1.44	1.62	1.96	2.12	2.23	15.78	8.62	59.51	44.21	75.30	52.23	345.59		
YKR074W	AM29	0.98	1.31	1.42	1.96	1.98	2.10	28.59	18.33	79.54	63.61	108.13	82.15	360.38		
YKR075C	YKR075c	0.98	1.36	1.52	1.97	2.06	2.20	9.42	4.36	47.53	13.62	56.17	25.50	329.98		
YKR076W	ECA1	0.92	1.33	1.52	1.97	2.04	2.24	9.42	4.36	47.53	13.62	56.17	25.50	329.40		
YKR077W	YKR077w	0.97	1.38	1.63	1.98	2.14	2.24	12.27	4.72	56.49	27.94	62.75	32.65	324.48		
YKR078W	MTD1	1.00	1.47	1.65	2.01	2.15	2.26	15.86	5.71	57.34	48.27	73.21	66.33	337.88		
YKR083W	NUP123	1.03	1.29	1.33	1.88	1.92	2.03	24.12	18.88	81.21	73.14	122.33	95.32	359.40		
YKR084C	BHS1	0.97	1.35	1.49	1.85	2.01	2.15	24.42	18.50	66.34	63.78	90.75	82.37	357.43		
YKR085C	MBS1	0.89	1.10	1.06	0.95	0.92	11.94	121.00	141.39	109.76	260.83	230.77	474.90			
YKR087C	CM41	0.97	1.41	1.59	1.88	2.14	2.20	8.32	3.07	45.20	9.31	53.52	12.38	301.44		
YKR088C	TVP38	0.96	1.40	1.54	1.90	2.07	2.15	9.02	3.05	48.27	10.58	57.29	13.63	315.81		
YKR089C	TGL4	0.90	1.38	1.54	1.94	2.11	2.19	6.77	3.21	39.96	14.30	46.73	17.52	303.46		
YKR090W	PXL1	0.96	1.36	1.55	1.96	2.12	2.20	9.44	3.70	48.67	21.97	58.11	25.67	307.01		
YKR091W	SRL3	0.95	1.39	1.58	2.04	2.22	2.31	4.14	3.01	24.98	8.96	29.12	11.96	295.92		
YKR091W	SRL3	0.90	1.26	1.51	1.88	2.05	2.18	6.25	1.58	31.81	9.18	34.79	10.12	277.80		
YKR092C	SRP40	0.97	1.41	1.54	1.98	2.17	2.27	19.69	9.51	57.60	44.22	77.28	53.73	353.67		
YKR093W	BAS1	1.05	1.55	1.64	2.02	2.22	2.35	5.74	2.88	37.50	7.70	43.24	8.58	317.80		
YKR094C	PTR2	1.04	1.57	1.70	2.20	2.27	2.37	8.90	3.21	47.13	14.62	56.03	17.83	295.73		
YKR094C	RPL40b	0.98	1.17	1.33	1.81	1.95	1.95	23.54	7.48	74.41	49.58	97.95	57.06	367.14		
YKR095W	MLP1	0.90	1.07	1.26	1.74	1.96	2.15	1.03	0.75	16.15	2.36	17.18	3.11	279.82		
YKR096W	YKR096w	0.89	1.04	1.20	1.61	1.87	2.07	1.02	0.77	14.55	3.52	15.57	4.30	270.13		
YKR097W	PKC1	0.96	1.43	1.57	1.78	2.01	2.19	12.47	3.81	46.62	20.94	57.11	15.60	289.67		
YKR098C	YKR098c	0.90	1.22	1.47	1.83	1.97	2.17	1.95	1.20	19.19	8.01	21.14	2.01	261.13		
YKR099C	PSR1	0.89	1.10	1.28	1.60	1.75	1.96	3.19	1.13	30.02	2.36	33.22	3.49	192.13		
YKR100C	YEH1	0.91	1.21	1.42	1.74	1.87	2.09	2.55	1.17	24.63	1.68	27.18	2.25	165.15		
YKR103C	PUF3	0.93	1.14	1.32	1.63	1.78	2.01	5.30	1.20	41.36	1.72	46.66	2.92	251.44		
YKR104W	EMC6	0.90	1.14	1.37	1.70	1.85	2.07	2.29	1.19	22.12	1.00	24.41	2.19	239.96		
YKR105W	BPT1	0.89	1.16	1.38	1.72	1.85	2.06	2.23	1.37	24.53	2.03	26.76	3.40	224.91		
YKR106W	SDC25	0.91	1.13	1.33	1.65	1.80	2.02	2.74	2.02	26.43	19.06	29.18	2.17	178.12		
YKR107W	YLO07w	0.91	1.19	1.39	1.74	1.91	2.13	1.35	1.05	9.74	2.05	11.09	3.10	206.48		
YKR108C	COK17	0.90	1.34	1.47	1.83	2.01	2.24	16.06	10.79	57.68	50.12	73.74	60.91	317.37		
YKR109C	KNST1	0.87	1.28	1.45	1.89	2.10	2.24	14.40	4.55	63.20	38.87	77.60	43.42	313.70		
YKR109C	YLO20c	0.96	1.22	1.46	1.81	1.98	2.20	2.74	1.40	47.27	7.19	27.52	8.59	258.04		
YKR109C	SP2A	0.95	1.32	1.58	1.90	2.03	2.21	1.54	1.22	6.94	4.73	8.48	5.94	187.80		
YKR109C	POM33	0.91	1.20	1.43	1.79	1.93	2.13	1.53	1.12	8.13	1.70</					

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YLRO17W	MEU1	1.09	1.42	1.61	1.97	2.11	2.28	1.29	2.52	12.61	21.64	13.90	24.16	148.84		
YLRO18C	POM34	0.98	1.23	1.44	1.82	1.99	2.19	1.14	1.48	10.44	15.50	11.57	16.98	164.75		
YLRO19W	PSR2	0.99	1.36	1.59	1.95	2.08	2.28	1.01	1.22	5.50	11.35	6.51	12.57	132.61		
YLRO20C	YEH2	1.02	1.34	1.54	1.92	2.06	2.24	1.90	6.45	22.51	40.16	24.41	46.60	140.93		
YLRO21W	IRC25	1.02	1.36	1.58	1.95	2.10	2.29	0.97	3.32	7.33	25.37	8.30	28.68	200.61		
YLRO23C	ZDH3	1.08	1.48	1.72	2.02	2.21	2.36	1.39	1.14	-1.60	1.51	-0.21	2.64	121.68		
YLRO24C	UBR2	1.05	1.29	1.48	1.84	2.08	2.27	1.04	1.18	9.43	17.44	10.47	8.52	193.57		
YLRO25W	SKE7	1.07	1.30	1.50	1.76	1.89	2.12	3.63	1.31	50.22	23.89	53.85	25.20	325.44		
YLRO26C	ATP2	1.04	1.34	1.56	1.87	2.04	2.22	6.04	4.61	31.33	43.05	35.94	284.13			
YLRO28C	ADE16	1.02	1.34	1.56	1.84	2.04	2.21	1.63	1.53	7.51	55.19	57.61	65.92	220.46		
YLRO30W	YLRO30w	0.88	1.31	1.51	1.93	2.12	2.23	3.44	3.05	17.04	4.29	20.49	7.34	295.21		
YLRO31W	YLRO31w	0.93	1.33	1.55	1.96	2.12	2.22	17.31	5.48	60.59	34.29	77.89	38.68	341.75		
YLRO32W	RAD5	0.90	1.33	1.46	1.81	2.09	2.21	9.24	3.62	44.12	18.98	53.37	22.61	342.75		
YLRO34C	SMF3	0.95	1.51	1.67	2.05	2.17	2.25	15.09	5.13	53.98	29.74	69.07	34.88	330.32		
YLRO35C	MHL2	1.01	1.52	1.71	2.12	2.27	2.35	17.49	5.88	59.37	35.58	76.86	41.47	358.32		
YLRO36C	YLRO36c	0.94	1.42	1.55	1.92	2.09	2.20	17.85	5.86	60.55	35.38	78.40	41.24	350.21		
YLRO37C	PAU23	0.92	1.41	1.52	1.89	2.06	2.17	8.88	3.65	43.93	21.24	52.82	24.89	321.06		
YLRO38C	CDX12	0.96	1.37	1.53	1.93	2.11	2.23	3.72	3.16	21.67	11.13	25.39	14.29	274.24		
YLRO39C	RIC1	0.87	1.29	1.43	1.76	1.92	2.02	17.85	8.17	55.79	41.58	73.64	49.75	344.13		
YLRO40C	YLRO40c	0.90	1.33	1.46	1.82	2.01	2.12	11.33	5.27	50.54	34.03	61.87	39.30	323.85		
YLRO41W	YLRO41w	0.90	1.32	1.48	1.81	1.99	2.10	8.27	3.13	41.60	14.70	49.88	17.83	314.48		
YLRO42C	YLRO42c	1.09	1.47	1.70	2.06	2.18	2.34	3.29	3.09	31.71	26.08	35.00	29.17	242.34		
YLRO43C	TRX1	1.00	1.24	1.38	1.64	1.81	2.04	1.01	1.13	0.63	-0.20	1.64	0.93	199.41		
YLRO44C	PDC1	0.94	1.11	1.26	1.79	1.77	2.07	1.45	1.91	12.26	16.34	13.71	18.25	183.92		
YLRO46C	YLRO46c	0.94	1.18	1.35	1.68	1.81	2.02	1.89	2.32	19.50	20.51	21.39	22.83	174.64		
YLRO47C	FRE8	1.00	1.30	1.44	1.72	1.83	2.04	1.47	1.47	14.38	14.57	15.84	16.04	182.48		
YLRO48W	RPS0b	0.99	1.12	1.22	1.68	1.96	2.14	2.22	3.01	7.39	10.05	8.60	13.05	212.72		
YLRO49C	YLRO49c	1.00	1.38	1.63	2.01	2.18	2.37	0.88	1.12	-1.60	-2.26	-0.72	-1.14	139.65		
YLRO50C	YLRO50c	0.93	1.40	1.62	2.00	2.15	2.22	11.02	5.32	46.92	33.06	57.95	38.38	323.80		
YLRO52W	iES3	0.87	1.26	1.40	1.79	1.97	2.05	16.18	11.38	55.15	49.52	71.33	60.90	342.94		
YLRO53C	YLRO53c	0.95	1.22	1.43	1.84	2.01	2.22	1.02	1.06	-0.07	2.95	0.95	4.04	150.67		
YLRO54C	OSW2	0.98	1.23	1.47	1.92	2.08	2.26	4.91	1.98	36.92	20.20	41.82	22.18	209.64		
YLRO54C	OSW2	0.96	1.28	1.46	1.86	2.03	2.18	9.57	1.90	56.45	21.11	66.02	23.02	303.16		
YLRO55C	SP78	1.02	1.15	1.30	1.68	1.88	2.16	18.12	13.86	74.56	61.52	92.68	75.37	314.68		
YLRO56W	ERG3	0.98	1.16	1.31	1.52	1.60	1.75	25.70	18.42	74.04	59.73	99.73	78.15	366.29		
YLRO57W	YLRO57w	0.95	1.21	1.44	1.80	1.96	2.17	3.46	2.68	31.73	21.61	35.19	23.90	232.42		
YLRO58C	SHM2	0.89	1.11	1.14	1.39	1.54	1.74	20.31	9.57	61.61	54.68	81.93	64.25	302.13		
YLRO59C	REX2	0.92	1.29	1.49	1.68	1.86	2.05	0.89	1.62	0.51	-0.49	1.41	0.53	158.28		
YLRO61W	RLP22a	0.96	1.28	1.49	1.65	1.76	2.04	1.68	0.93	40.03	-1.52	1.63	-0.68	240.46		
YLRO62C	BUD20	1.00	1.09	1.14	1.49	1.70	1.70	1.03	1.06	-0.11	2.64	0.83	1.33	177.66		
YLRO63W	YLRO63w	0.99	1.31	1.55	1.97	2.10	2.33	1.01	1.19	1.64	3.63	2.83	3.83	156.36		
YLRO64W	PER23	0.93	1.14	1.31	1.72	1.88	2.09	1.90	1.65	18.55	16.84	20.48	18.60	194.47		
YLRO65C	YLRO65c	0.91	1.03	1.11	1.45	1.63	1.87	1.96	2.84	23.23	29.38	25.19	32.23	188.17		
YLRO67C	PET309	0.88	1.05	1.03	0.91	0.98	1.08	121.57	127.81	115.57	138.63	237.14	266.44	513.32		
YLRO68W	FIV7	0.97	1.13	1.31	1.77	1.95	2.16	15.67	4.97	61.94	37.95	77.61	42.91	304.28		
YLRO69C	MEF1	0.87	1.02	1.02	0.90	0.92	0.97	108.66	113.11	112.03	137.85	220.69	259.97	472.66		
YLRO70C	XYL2	1.05	1.45	1.70	2.04	2.15	2.34	7.03	1.09	49.75	8.84	56.78	9.94	306.34		
YLRO72W	YLRO72w	0.98	1.34	1.61	1.95	2.08	2.25	2.51	2.10	25.15	10.59	27.66	11.69	286.21		
YLRO73C	RFU1	1.02	1.32	1.52	1.84	1.99	2.18	2.50	1.91	23.72	20.84	26.21	22.75	274.35		
YLRO74C	BUD20	0.99	1.14	1.26	1.64	1.89	2.17	11.92	9.13	58.87	50.43	70.80	59.56	281.45		
YLRO77W	FMP25	1.04	1.34	1.57	1.94	2.08	2.27	2.54	2.54	16.32	5.87	18.86	7.04	185.56		
YLRO79W	SIC1	1.02	1.17	1.31	1.62	1.80	2.04	0.97	1.01	7.89	-0.26	8.87	0.75	248.36		
YLRO80W	EMP46	0.98	1.19	1.37	1.77	1.94	2.13	1.60	1.47	14.59	16.19	16.19	16.43	184.24		
YLRO81W	GAL2	0.91	1.08	1.27	1.69	1.88	2.08	1.55	1.32	15.97	11.43	17.52	12.74	179.01		
YLRO82C	SLR2	0.95	1.23	1.43	1.83	2.03	2.27	0.96	1.24	-0.51	-1.58	0.45	-0.34	193.10		
YLRO83C	EMP70	0.91	1.17	1.35	1.74	1.93	2.16	1.06	0.95	0.98	0.38	2.04	1.33	180.88		
YLRO84C	RAX2	0.95	1.16	1.35	1.80	1.99	2.22	9.04	1.24	55.22	6.59	64.26	7.83	309.26		
YLRO85C	ICD2	0.93	1.21	1.40	1.63	1.83	2.04	2.29	2.05	0.95	-1.64	-0.53	-0.69	-1.70	292.39	
YLRO96W	KIN2	0.96	1.21	1.40	1.75	1.91	2.15	1.95	1.26	1.28	9.58	10.80	10.80	12.09	217.13	
YLRO97C	HLA1	0.98	1.28	1.48	1.82	2.00	2.20	1.67	1.77	12.26	7.44	23.51	8.53	227.11		
YLRO98C	ICB44	0.86	1.18	1.30	1.65	1.82	2.00	2.23	2.36	16.77	59.69	12.26	16.46	124.43	341.24	
YLRO99C	CTT1	1.01	1.38	1.58	2.01	2.15	2.33	18.86	1.66	66.89	12.86	28.75	13.92	338.84		
YLRO102C	APC9	0.96	1.33	1.55	1.97	2.12	2.33	19.45	2.55	69.18	24.99	88.63	27.45	333.56		
YLRO104W	YLRO14w	0.91	1.30	1.58	1.98	2.13	2.31	10.38	2.43	53.05	25.64	63.43	28.07	311.82		
YLRO107W	REX2	0.96	1.18	1.37	1.68	1.86	2.09	4.55	2.31	41.91	23.04	46.46	25.34	289.81		
YLRO108C	YLRO108c	1.00	1.23	1.44	1.84	2.02	2.24	2.42	2.17	25.58	19.84	28.00	21.35	286.48		
YLRO109W	SRN2	0.98	0.98	1.44	1.78	1.94	2.15	1.58	1.53	-1.45	-1.42	0.13	0.12	314.46		
YLRO120C	YPS1	1.09	1.09	1.75	2.10	2.23	2.36	1.51	1.50	-1.45	-1.48	0.07	0.02	304.26		
YLRO121C	YPS3	1.01	1.01	1.62	1.99	2.16	2.26	1.20	1.46	-0.15	-1.34	1.05	0.12	269.43		
YLRO122C	YLRO122c	0.99	0.99	1.50	1.86</											

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE			FRUCTOSE			TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YLR190W	MMR1	0.96	1.20	1.34	1.75	1.96	2.09	5.00	2.95	46.23	36.82	51.23	39.78	337.68	337.68	337.68	337.68	
YLR191W	PEX13	1.04	1.32	1.51	2.03	2.22	2.32	4.92	3.71	44.38	37.74	49.29	41.45	272.02	272.02	272.02	272.02	
YLR192C	HCR1	0.92	1.09	1.26	1.76	1.97	2.16	1.42	1.06	21.47	9.42	22.88	10.48	265.75	265.75	265.75	265.75	
YLR193C	UPS1	1.07	1.44	1.64	2.03	2.20	2.25	11.29	8.70	60.55	53.50	71.84	62.20	341.42	341.42	341.42	341.42	
YLR194C	YLR194c	0.98	1.24	1.45	1.99	2.20	2.34	5.97	3.86	45.12	25.47	51.10	27.34	322.81	322.81	322.81	322.81	
YLR195C	PBL1	0.94	1.18	1.37	1.89	1.99	2.14	8.80	2.35	57.65	24.53	66.45	26.88	277.42	277.42	277.42	277.42	
YLR196C	YKE2	1.02	1.37	1.46	2.02	2.38	2.32	18.99	5.87	72.56	50.12	81.55	56.09	337.40	337.40	337.40	337.40	
YLR198W	C009	0.98	1.17	1.34	0.91	0.97	0.92	12.82	3.84	120.93	123.27	254.79	254.79	323.94	323.94	323.94	323.94	
YLR202C	YLR202c	0.98	1.08	1.11	0.92	0.93	0.88	124.81	123.85	130.83	126.20	255.66	255.66	331.03	331.03	331.03	331.03	
YLR203C	MSS51	0.98	1.19	1.17	1.06	1.08	1.07	120.44	120.83	132.79	128.19	253.23	249.84	320.11	320.11	320.11	320.11	
YLR204W	Q515	0.99	1.27	1.46	1.85	2.05	2.21	1.03	1.04	9.23	3.60	10.26	4.63	275.62	275.62	275.62	275.62	
YLR207W	HBD3	0.98	1.27	1.46	1.85	2.05	2.21	1.03	1.04	9.23	3.60	10.26	4.63	275.19	275.19	275.19	275.19	
YLR209C	PNP1	1.02	1.30	1.50	2.03	2.22	2.35	1.28	1.48	18.31	10.15	19.59	11.63	281.47	281.47	281.47	281.47	
YLR210W	CLB4	0.96	1.25	1.48	1.98	2.14	2.26	1.38	1.15	16.62	14.05	18.01	15.21	288.66	288.66	288.66	288.66	
YLR211C	YLR211c	0.96	1.25	1.44	1.98	2.18	2.29	9.67	3.62	59.50	38.75	69.17	42.37	323.93	323.93	323.93	323.93	
YLR211C	YLR211c	0.90	1.19	1.37	1.78	1.95	2.08	8.17	2.04	52.00	21.54	60.17	23.58	284.77	284.77	284.77	284.77	
YLR213C	CR11	0.98	1.23	1.36	1.83	2.07	2.22	2.83	1.28	35.84	17.83	38.68	19.11	296.78	296.78	296.78	296.78	
YLR214W	FRE1	0.96	1.19	1.32	1.86	2.09	2.24	1.32	0.79	19.43	10.55	20.74	11.34	293.33	293.33	293.33	293.33	
YLR216C	CPR6	0.97	1.23	1.42	2.00	2.22	2.36	1.26	0.79	22.03	9.61	23.29	10.36	294.44	294.44	294.44	294.44	
YLR217W	YLR217w	0.99	1.14	1.22	1.76	2.02	2.19	9.30	2.95	56.47	36.07	65.77	39.02	301.71	301.71	301.71	301.71	
YLR218C	YLR218c	0.89	1.23	1.43	1.78	2.00	2.16	1.41	1.47	2.59	0.34	4.00	1.81	223.54	223.54	223.54	223.54	
YLR219W	MSC3	0.98	1.22	1.38	1.88	2.07	2.21	1.88	0.92	29.51	5.10	31.39	6.02	291.21	291.21	291.21	291.21	
YLR220W	CCC1	0.99	1.24	1.43	1.89	2.08	2.24	4.43	1.31	41.94	12.69	46.37	13.99	290.28	290.28	290.28	290.28	
YLR221C	RSA3	0.97	1.22	1.40	1.84	2.04	2.19	1.12	1.05	14.46	1.57	15.58	2.62	292.37	292.37	292.37	292.37	
YLR224W	YLR224w	0.99	1.28	1.50	2.01	2.18	2.30	4.23	1.55	42.68	21.91	46.91	23.46	261.17	261.17	261.17	261.17	
YLR225C	YLR225c	0.99	1.23	1.37	1.93	2.15	2.30	1.46	1.21	20.66	12.66	22.12	13.87	294.68	294.68	294.68	294.68	
YLR226W	BUR2	0.87	1.06	1.03	0.99	1.11	1.26	82.62	73.66	130.35	98.75	212.97	172.41	395.14	395.14	395.14	395.14	
YLR227C	ADY4	0.99	1.31	1.51	2.01	2.19	2.32	1.11	1.03	10.00	5.69	11.11	6.72	286.33	286.33	286.33	286.33	
YLR228C	ECM22	0.99	1.18	1.32	1.82	2.01	2.22	3.38	1.43	34.26	9.18	37.65	20.63	284.43	284.43	284.43	284.43	
YLR231C	BNA5	0.97	1.19	1.35	1.68	1.81	2.00	8.19	2.98	54.43	32.77	62.62	35.75	301.63	301.63	301.63	301.63	
YLR232W	YLR232w	0.96	1.24	1.41	1.83	2.00	2.18	10.84	2.49	61.46	28.58	72.30	31.07	315.79	315.79	315.79	315.79	
YLR233C	ES71	0.95	1.12	1.21	1.56	1.72	1.97	12.40	5.00	67.55	45.97	79.95	50.97	309.24	309.24	309.24	309.24	
YLR234W	TOP3	0.95	1.10	1.19	1.56	1.77	2.04	7.70	2.52	52.72	32.44	60.42	34.96	344.83	344.83	344.83	344.83	
YLR235C	YLR235c	0.95	1.16	1.30	1.84	2.06	2.36	7.38	3.33	54.19	33.79	61.57	37.12	303.70	303.70	303.70	303.70	
YLR236C	YLR236c	0.95	1.17	1.28	1.68	1.81	2.05	9.17	2.58	56.53	32.94	65.70	35.88	308.46	308.46	308.46	308.46	
YLR237W	YLR237w	0.97	1.17	1.36	1.62	1.81	2.05	16.65	1.14	65.89	17.82	72.25	15.25	317.44	317.44	317.44	317.44	
YLR238W	FAR10	0.93	1.30	1.42	1.96	2.10	2.26	9.24	3.95	64.69	44.26	73.93	48.22	229.67	229.67	229.67	229.67	
YLR239C	UP2	0.99	1.28	1.47	1.90	2.06	2.24	4.66	2.08	38.70	24.34	43.37	26.41	296.88	296.88	296.88	296.88	
YLR240W	VPS24	0.98	1.09	1.21	0.93	0.97	0.98	5.23	3.74	114.48	81.46	168.00	110.51	311.48	311.48	311.48	311.48	
YLR241W	YLR241w	1.04	1.30	1.48	1.95	2.12	2.29	11.65	3.62	60.37	43.04	72.02	47.83	318.71	318.71	318.71	318.71	
YLR242C	YLR242c	1.05	1.13	1.10	1.11	1.14	1.18	138.63	132.44	134.79	130.71	273.42	265.15	353.42	353.42	353.42	353.42	
YLR243C	APV1	0.93	1.16	1.16	1.04	1.09	1.06	122.62	124.65	134.26	130.45	256.88	250.10	530.10	530.10	530.10	530.10	
YLR244C	APV1	0.83	1.16	1.16	0.89	0.90	0.92	118.75	119.24	124.22	117.58	242.97	238.81	525.44	525.44	525.44	525.44	
YLR246W	ERF2	0.94	1.14	1.34	1.77	1.93	2.07	37.75	8.94	98.03	55.62	135.78	64.56	320.03	320.03	320.03	320.03	
YLR247C	IRC20	0.96	1.07	1.10	1.50	1.70	1.96	21.70	14.98	63.81	47.31	96.01	80.78	297.71	297.71	297.71	297.71	
YLR248W	RCK2	0.94	1.15	1.28	1.74	1.95	2.14	4.84	1.71	40.17	22.99	45.01	24.69	319.37	319.37	319.37	319.37	
YLR250W	SSP120	0.96	1.15	1.29	1.71	1.89	2.10	6.41	1.41	50.74	18.29	57.15	19.70	304.84	304.84	304.84	304.84	
YLR251W	SYMF	0.98	1.24	1.42	1.85	2.02	2.20	3.38	1.39	39.48	16.73	42.86	18.12	294.51	294.51	294.51	294.51	
YLR252W	SYMF	0.85	1.10	1.21	1.62	1.84	2.00	7.91	1.14	51.15	24.50	59.06	13.28	291.04	291.04	291.04	291.04	
YLR253W	SEC22	0.90	1.09	1.21	1.56	1.73	1.94	5.45	1.98	44.13	24.63	49.58	26.61	308.61	308.61	308.61	308.61	
YLR254C	NDL1	0.94	1.15	1.28	1.63	1.83	2.00	6.49	2.80	50.28	33.49	56.77	36.29	311.25	311.25	311.25	311.25	
YLR255C	YLR255c	1.00	1.23	1.38	1.78	1.93	2.13	11.12	3.65	53.62	26.56	61.51	25.81	295.06	295.06	295.06	295.06	
YLR257W	YLR271w	1.10	1.47	1.70	1.83	2.02	2.21	1.54	0.86	24.61	5.65	26.15	6.51	289.44	289.44	289.44	289.44	
YLR278C	PIG1	0.99	1.20	1.34	1.73	1.93	2.11	20.47	5.34	76.28	46.56	96.75	51.90	324.02	324.02	324.02	324.02	
YLR278C	YLR278c	1.02	1.21	1.34	1.87	2.07	2.28	5.18	1.05	45.76	10.89	50.94	11.94	301.29	301.29	301.29	301.29	
YLR279W	YLR279w	1.07	1.30	1.45	1.97	2.17	2.32	4.88	1.11	46.58	15.22	51.46	16.32	284.91	284.91	284.91	284.91	
YLR280C	YLR280c	0.98	1.11	1.16	1.61	1.85	2.06	19.89	11.79	67.65	61.01	87.54	72.81	324.60	324.60	324.60	324.60	
YLR281C																		

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YLR342W-A	YLR342w-a	0.91	1.21	1.34	1.72	1.93	2.10	2.82	6.87	36.61	37.93	39.42	44.80	265.83		
YLR343W	GAS2	1.01	1.49	1.72	2.07	2.21	2.30	8.83	3.57	53.83	27.74	62.66	31.32	284.96		
YLR344W	RPL26a	1.03	1.32	1.45	1.93	2.07	2.21	10.64	3.72	55.35	35.84	65.99	39.56	321.28		
YLR345W	YLR345w	1.04	1.38	1.57	2.06	2.22	2.34	2.64	1.23	24.15	13.19	26.79	14.42	293.93		
YLR346C	YLR346c	0.90	1.05	1.22	1.72	1.91	2.10	2.06	0.81	31.14	3.85	33.21	4.66	290.63		
YLR348C	DIC1	1.03	1.29	1.44	1.91	2.08	2.24	8.58	2.21	49.88	28.00	58.46	30.22	314.45		
YLR349W	YLR349w	1.05	1.39	1.53	2.02	2.15	2.28	7.26	2.22	50.74	30.03	58.00	32.25	313.25		
YLR350W	CIR2	1.01	1.32	1.53	1.97	2.14	2.26	12.14	5.17	58.57	48.10	70.41	45.26	308.19		
YLR351G	YLR351g	1.17	1.28	1.35	2.04	2.16	2.30	2.48	0.45	29.03	2.71	31.14	2.71	311.41		
YLR352W	YLR352w	1.01	1.28	1.35	2.00	2.15	2.27	2.65	1.25	31.12	14.02	33.77	15.28	288.28		
YLR353W	BUD8	1.01	1.15	1.28	1.72	1.94	2.12	9.41	2.29	54.56	29.38	63.97	31.67	309.32		
YLR354C	TAL1	0.99	1.28	1.48	1.87	2.04	2.20	10.10	3.53	56.16	33.82	66.25	37.35	312.36		
YLR356W	ATG33	0.96	1.17	1.33	1.83	2.03	2.21	13.42	0.92	63.51	8.88	76.93	9.00	316.67		
YLR357W	RSC2	0.97	1.09	1.11	1.32	1.48	1.69	59.57	41.57	104.49	92.40	164.06	132.08	351.47		
YLR357W	RSC2	0.88	1.14	1.15	1.27	1.48	1.72	22.24	8.08	78.03	59.66	100.15	67.74	333.36		
YLR358C	YLR358c	0.91	0.95	0.97	1.16	1.34	1.57	65.24	50.30	111.20	103.23	176.44	153.53	331.08		
YLR360W	VPS38	0.97	1.21	1.35	1.65	1.78	1.95	2.32	0.90	38.21	12.88	40.53	13.78	318.11		
YLR361C	DCR2	0.87	0.97	1.09	1.59	1.81	2.01	5.85	1.34	46.46	2.07	52.31	22.41	305.94		
YLR361C	YLR361c-a	0.93	1.25	1.38	1.81	2.00	2.15	2.47	1.08	35.04	45.11	37.52	56.19	268.83		
YLR362W	STE11	0.91	1.23	1.41	1.78	1.94	2.10	3.87	0.96	38.37	8.08	42.25	9.04	283.81		
YLR363C	NMD4	0.94	1.27	1.49	1.82	1.97	2.11	15.05	10.19	64.85	53.37	79.90	63.56	298.86		
YLR363W-A	YLR363w-a	0.93	1.23	1.38	1.83	2.03	2.17	0.72	5.60	16.02	33.84	16.73	39.44	254.05		
YLR364W	GRX8	0.93	1.16	1.33	1.78	1.95	2.12	17.40	1.34	69.64	19.60	87.04	20.93	314.10		
YLR365W	YLR365w	0.90	1.06	1.13	1.59	1.81	2.04	13.76	2.43	66.13	28.50	79.88	30.94	321.45		
YLR366W	YLR366w	0.90	1.10	1.23	1.71	1.89	2.09	8.26	1.02	53.63	10.42	61.88	11.44	305.31		
YLR367W	RPS22b	0.99	1.15	1.27	1.78	1.98	2.17	13.76	8.84	65.39	52.06	79.15	60.90	318.69		
YLR368W	MDM30	0.92	1.06	1.15	1.64	1.85	2.04	13.63	2.78	64.07	30.62	77.70	33.39	263.78		
YLR369W	SSQ1	0.88	0.98	0.99	0.89	0.91	0.92	88.79	75.06	134.15	98.90	222.94	175.96	404.46		
YLR370C	ARC18	0.88	1.18	1.30	1.84	1.92	1.99	6.82	4.85	48.15	33.91	54.97	38.76	290.82		
YLR371W	ROM2	0.99	1.28	1.39	1.72	1.91	2.07	1.41	0.81	19.90	2.63	21.31	3.45	253.64		
YLR371W	ROM2	0.98	1.20	1.41	1.83	1.99	2.15	18.92	7.12	70.98	47.84	89.90	54.96	330.02		
YLR372W	SUR4	0.94	1.20	1.37	1.74	1.87	2.05	19.42	8.55	66.74	48.53	86.17	57.08	325.53		
YLR373C	VID22	0.98	1.19	1.34	1.74	1.94	2.18	7.36	5.37	49.00	40.08	56.37	45.45	233.24		
YLR374C	YLR374c	0.93	1.31	1.41	1.75	1.93	2.13	1.11	0.89	9.00	2.20	10.10	3.09	269.70		
YLR375W	STP3	1.01	1.29	1.49	1.86	2.02	2.19	16.99	2.51	65.31	29.02	82.29	31.53	326.29		
YLR376C	PSY3	0.94	1.22	1.39	1.76	1.94	2.15	1.34	1.06	15.96	10.55	17.30	11.61	283.30		
YLR376C	FBP1	0.99	1.25	1.43	1.86	2.03	2.19	16.62	6.68	64.10	26.25	80.92	28.33	330.35		
YLR377C	CSRT	0.99	1.30	1.44	1.86	2.03	2.19	5.73	2.85	47.11	8.62	52.84	11.77	293.35		
YLR378W	YLR378w	0.93	1.03	1.02	1.21	1.42	1.62	50.88	2.13	104.34	61.04	155.1	106.17	305.83		
YLR379C	CTT3	0.98	1.20	1.35	1.67	1.90	2.05	0.59	0.62	8.59	2.16	9.24	2.82	267.44		
YLR380C	NAM2	0.90	1.11	1.26	1.67	1.87	2.05	1.23	1.04	14.27	5.67	65.81	45.18	78.28	50.85	
YLR384C	IK3	0.98	1.17	1.33	1.82	1.99	2.18	14.81	2.95	66.14	32.97	80.98	35.92	311.06		
YLR384C	SHC7	1.00	1.17	1.26	1.79	2.00	2.19	16.82	6.78	66.28	48.70	82.89	52.49	322.21		
YLR388W	VAC14	0.91	1.21	1.35	1.77	1.98	2.21	22.27	9.49	78.74	5.71	101.01	66.99	342.43		
YLR387C	REB1	0.93	1.14	1.31	1.71	1.90	2.13	1.99	0.88	12.13	4.18	13.22	5.07	281.99		
YLR388W	RPS29a	0.94	1.07	1.13	1.58	1.85	2.10	12.53	5.84	63.58	44.61	76.12	59.45	317.07		
YLR389C	STE23	0.90	1.11	1.27	1.61	1.89	2.09	0.86	0.49	14.27	5.00	15.14	5.40	289.37		
YLR390W	ECM19	0.90	1.09	1.20	1.58	1.77	1.94	12.47	5.67	65.81	45.18	78.28	50.85	315.91		
YLR390W-A	CGW14	0.93	1.26	1.54	1.96	2.14	2.25	5.35	2.52	43.77	30.48	49.12	33.00	292.97		
YLR391W	YLR391W	0.87	1.03	1.16	1.78	1.70	1.83	2.43	0.28	33.66	7.40	36.09	7.68	277.10		
YLR392C	ART10	0.98	1.23	1.41	1.83	2.01	2.20	2.55	1.21	29.81	15.23	32.16	16.44	280.30		
YLR393W	ATP10	1.08	1.50	1.74	2.09	2.21	2.34	3.84	1.06	39.84	11.04	43.68	12.10	271.84		
YLR394W	CST9	0.88	1.04	1.21	1.66	1.84	2.02	8.98	2.98	52.85	35.74	61.84	38.72	309.71		
YLR395C	CDX8	1.07	1.39	1.54	1.93	2.07	2.24	11.67	2.98	61.28	34.84	72.96	37.82	314.99		
YLR396C	VPS33	0.98	1.05	1.07	0.97	0.99	0.95	124.32	134.48	145.89	118.02	270.21	252.50	393.16		
YLR398C	SK2	0.93	1.29	1.45	1.88	2.06	2.25	7.66	2.74	50.80	34.22	61.86	24.17	297.94		
YLR399C	BDI1	0.90	1.00	1.01	0.91	0.92	0.88	118.68	118.68	145.78	113.13	264.46	231.81	412.39		
YLR400W	YLR400w	0.94	1.14	1.27	1.88	2.07	2.33	4.72	6.52	65.32	42.31	78.65	47.03	319.01		
YLR401C	DUS3	1.03	1.41	1.52	1.94	2.08	2.25	12.09	9.21	61.64	52.18	73.73	61.39	315.06		
YLR402W	YLR402w	1.02	1.35	1.57	1.99	2.15	2.31	1.01	0.93	3.68	0.67	4.70	1.60	256.33		
YLR403W	SFP1	0.99	1.19	1.28	1.47	1.75	1.97	0.74	0.45	3.51	0.74	4.25	1.19	221.86		
YLR404W	FLD1	1.02	1.24	1.40	1.86	2.04	2.21	12.99	4.38	61.51	38.70	74.50	43.08	315.53		
YLR405W	DUS4	1.00	1.27	1.48	1.90	2.07	2.22	2.04	0.96	26.74	7.60	28.78	8.56	289.25		
YLR406C	RPL31b	0.99	1.15	1.30	1.86	2.03	2.23	10.64	5.41	60.18	43.14	70.82	46.55	318.36		
YLR406C	YLR406c-a	0.89	1.15	1.26	1.86	2.05	2.20	10.81	2.43	57.78	31.68	68.59	34.10	313.53		
YLR407W	YLR407w	0.95	1.20	1.39	1.80	1.96	2.14	21.70	1.20	118.68	59.45	71.32	51.10	329.76		
YLR408C	Unknown	1.01	1.25	1.39	1.74	1.93	2.08	13.11	5.92	64.64	46.05	74.22	40.56	325.66		
YLR420W	TSR2	0.92	1.20	1.26	1.58	1.84	2.05	23.14	15.02	63.55	51.35	86.69	66.37	338.74		
YLR420W	ECM30	0.95	1.31	1.50	1.90											

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YML018C	YML018c	0.98	0.98	1.51	1.88	2.03	2.21	11.75	1.31	2.36	-1.38	14.11	-0.08	254.15		
YML019W	CST6	0.97	0.97	1.42	1.82	1.98	2.18	1.57	1.60	-1.46	-1.48	0.11	0.13	265.97		
YML020W	YML020w	0.95	0.95	1.43	1.89	2.08	2.31	1.23	1.73	-1.38	-1.13	-0.16	0.60	252.68		
YML021C	UNG1	0.91	1.25	1.43	1.80	1.99	2.16	11.29	5.60	47.19	31.50	58.48	37.10	341.53		
YML022W	APT1	0.95	1.16	1.23	1.42	1.61	1.76	2.48	1.28	26.94	6.82	29.42	8.10	326.86		
YML024W	RPS17a	0.92	0.92	1.02	1.36	1.67	2.01	5.40	1.62	5.16	-1.37	10.56	0.25	305.50		
YML026C	RPS16b	0.89	0.89	1.15	1.61	1.81	2.11	9.13	1.56	2.37	-1.38	11.50	0.17	266.90		
YML027W	YOKT	0.91	1.03	1.52	1.98	2.07	2.29	1.62	0.93	14.55	1.74	15.57	2.66	276.33		
YML029W	YAL1	0.94	0.94	1.37	1.64	1.78	2.15	10.67	1.59	1.28	-1.28	12.84	0.11	302.40		
YML030W	USA1	0.94	0.94	1.37	1.79	1.96	2.14	9.97	1.22	1.64	-1.43	11.61	-0.21	299.73		
YML030W	AH431	0.95	1.05	1.42	1.79	1.96	2.14	11.56	1.54	3.22	-1.45	14.78	0.10	351.47		
YML032C	RAD52	1.01	1.01	1.28	1.81	1.83	2.04	10.86	1.62	-0.21	-1.54	10.64	0.09	274.19		
YML033W	YML033w	0.99	0.99	1.38	1.74	1.91	2.09	10.86	1.58	-2.95	-1.49	14.12	0.09	290.62		
YML034W	SRC1	1.01	1.01	1.51	1.90	2.06	2.22	17.07	1.58	1.27	-1.54	14.09	0.09	326.60		
YML034W	SRC1	0.91	1.22	1.37	1.72	1.91	2.08	5.49	1.39	49.04	16.08	54.54	17.47	292.33		
YML035C	AMD1	1.00	1.00	1.55	2.01	2.17	2.34	10.67	1.72	3.33	-1.51	14.20	0.21	278.10		
YML035C	AMD1	0.92	1.07	1.21	1.62	1.83	2.03	1.19	0.58	21.40	4.48	22.58	5.07	298.14		
YML035C-A	Unknown	1.00	1.00	1.45	1.88	2.05	2.22	15.75	1.50	-0.40	-1.25	15.35	0.25	304.75		
YML036W	CG121	0.92	1.05	1.21	1.72	1.92	2.09	22.56	1.75	7.34	24.85	95.80	26.61	315.13		
YML037C	YML037c	1.03	1.03	1.60	2.03	2.20	2.38	1.26	1.43	-1.43	-1.45	14.08	0.09	255.30		
YML038C	YMD8	0.99	1.22	1.40	1.86	2.04	2.18	7.89	0.95	50.74	6.60	58.63	7.55	306.42		
YML041C	VPS71	0.97	1.04	1.12	1.49	1.69	1.87	43.52	14.07	97.28	61.25	140.80	75.32	332.60		
YML042W	CAT2	0.98	1.19	1.41	1.91	2.09	2.24	4.54	0.89	42.92	10.90	47.46	11.79	292.20		
YML047C	PRM6	0.87	0.99	1.14	1.61	1.81	1.98	5.31	2.18	46.14	28.53	51.45	30.71	299.29		
YML048W	GSF2	0.89	0.89	1.16	1.49	1.66	1.90	6.24	1.70	4.08	-1.64	10.32	0.05	286.83		
YML049W	Unknown	0.92	1.26	1.40	1.82	2.04	2.16	17.49	3.68	64.93	35.95	82.42	39.62	324.57		
YML050W	AIM22	0.93	0.93	1.43	1.85	2.02	2.24	9.12	1.38	1.82	-1.51	10.94	-0.13	262.87		
YML051W	GAL80	0.93	0.93	1.27	1.61	1.78	1.99	11.53	1.81	2.83	-1.42	14.36	0.40	332.20		
YML052W	SUR7	0.93	0.93	1.20	1.48	1.63	1.85	9.74	1.44	2.02	-1.43	11.76	0.01	276.43		
YML053C	YML053c	0.97	1.37	1.52	1.97	2.40	2.37	22.94	10.99	75.09	53.02	98.04	64.01	345.88		
YML054C	CYB2	1.00	1.00	1.41	1.82	2.00	2.20	10.97	1.43	2.23	-1.40	13.21	0.03	262.66		
YML054C-A	YML054c-a	0.88	1.10	1.26	1.70	1.94	2.10	12.33	4.93	63.68	33.51	76.01	38.45	307.59		
YML055W	SPC2	0.89	1.26	1.44	1.88	2.13	2.32	1.02	0.82	12.84	2.66	13.86	3.69	255.91		
YML056C	IMD4	1.06	1.06	1.57	1.91	2.10	2.30	14.33	3.78	-0.78	-2.10	13.56	35.88	268.84		
YML057W	CMP2	0.98	1.20	1.35	1.67	1.90	2.04	11.79	1.51	54.70	14.31	66.49	15.81	301.92		
YML058-C	Unknown	0.97	0.97	1.56	1.94	2.10	2.26	13.07	1.55	1.00	-1.37	14.08	0.58	285.56		
YML059W	SM17	1.00	1.00	1.40	1.81	2.00	2.22	1.61	1.43	-1.35	-1.42	0.26	0.02	296.52		
YML060W	HOG1	0.88	1.18	1.34	1.77	2.00	2.17	11.04	4.59	56.94	30.41	69.98	34.90	306.83		
YML061W	ITTC1	0.96	1.06	1.41	1.88	2.06	2.24	1.68	1.44	-1.44	-1.53	0.15	0.28	226.45		
YML062W	OGG1	0.93	1.03	1.24	1.65	1.84	2.07	6.98	1.44	2.97	0.93	9.85	0.10	308.48		
YML061C	PIF1	0.91	0.91	1.25	1.66	1.87	2.10	9.32	1.57	0.85	-1.45	10.17	0.12	279.01		
YML063C	MST1	0.93	0.93	1.30	1.69	1.84	2.04	11.61	1.46	0.94	-1.35	12.55	0.11	346.29		
YML053W	RPS1b	0.98	1.27	1.38	1.82	2.03	2.27	20.07	6.05	72.55	44.29	92.63	50.35	326.42		
YML066C	SM2	0.98	1.07	1.24	1.62	1.88	2.07	1.59	1.11	28.89	16.56	30.47	17.67	311.62		
YML067C	ERV41	1.00	1.49	1.70	2.07	2.21	2.32	13.88	7.75	62.40	51.71	76.29	59.46	306.62		
YML068W	ITTI	0.91	1.29	1.44	1.87	2.04	2.16	16.69	7.84	70.50	48.86	87.19	56.70	325.04		
YML070W	DAK1	0.98	1.54	1.75	2.07	2.21	2.30	12.65	4.87	49.72	27.57	62.38	32.45	336.64		
YML071C	COGB	0.93	1.41	1.58	1.91	2.00	2.11	20.31	9.62	62.88	46.62	83.19	56.24	359.57		
YML072C	TCB3	0.93	1.33	1.50	1.96	2.13	2.25	21.10	9.35	52.02	32.72	62.37	340.31			
YML073C	RPL6a	1.01	1.27	1.44	1.91	2.12	2.25	8.43	1.84	45.16	20.25	53.59	22.10	289.95		
YML074C	FPR3	0.99	1.57	1.78	2.13	2.24	2.34	9.29	4.90	44.39	25.91	53.68	30.81	309.37		
YML075C	HMG1	0.84	0.94	1.07	1.57	1.77	1.97	2.59	0.92	31.64	14.71	34.23	15.62	290.09		
YML076C	WAR1	0.91	1.12	1.33	1.76	1.94	2.05	10.72	3.13	58.20	35.36	68.92	38.50	322.35		
YML078W	CPR3	0.97	0.97	1.43	1.79	1.96	2.16	9.62	1.65	2.19	-1.51	11.80	0.14	332.71		
YML079W	YML079w	0.97	0.97	1.26	1.58	1.74	1.95	10.83	1.90	1.85	-1.55	12.69	0.35	310.13		
YML080W	DUS1	1.01	1.01	1.47	1.88	2.04	2.22	12.55	1.23	1.78	-1.28	14.32	-0.04	304.93		
YML081C	ATP18	0.97	1.33	1.53	1.96	2.16	2.32	17.03	8.11	57.56	38.76	74.59	46.87	336.06		
YML081W	YML081w	1.01	1.01	1.55	1.93	2.10	2.26	12.68	2.07	2.02	-1.29	14.70	0.77	281.80		
YML082W	YML082w	0.98	0.98	1.39	1.71	1.88	2.11	10.10	1.53	2.11	-1.35	12.21	0.18	272.40		
YML083C	YML083c	0.99	0.99	1.40	1.80	1.99	2.21	6.77	1.40	4.52	-1.40	11.29	0.00	283.85		
YML084W	YML084w	1.03	1.03	1.56	1.96	2.12	2.30	1.74	1.37	-1.52	-1.26	0.22	0.11	308.98		
YML086C	ALOT	0.97	0.97	1.37	1.85	2.03	2.23	1.34	1.17	-1.48	-1.38	-0.14	-0.22	278.00		
YML086C	ALOT	0.88	1.06	1.24	1.70	1.92	2.10	5.60	2.87	2.98	-1.43	18.88	0.28	310.79		
YML088W	UF07	0.90	0.90	1.33	1.77	1.95	2.19	1.31	1.34	-0.99	-1.38	0.52	-0.04	295.02		
YML089C	YML089c	0.93	1.35	1.48	1.89	2.04	2.24	3.85	3.07	23.60	6.93	27.45	10.00	276.81		
YML101C	CUE4	0.98	1.50	1.68	2.00	2.16	2.27	3.32	3.16	16.41	9.83	19.73	12.99	293.37		
YML102C-A	Unknown	0.99	1.63	1.85	2.18	2.28	2.35	14.58	5.27	57.57	32.02	78.95	36.45	322.73		
YML102W	CAC2	0.97	1.47	1.68	2.00	2.13	2.25	8.86	0.80	7.32	39.81	18.56	48.41	22.27	270.78	
YML103C	NUP188	0.94	1.30	1.39	1.76	1.92	2.02	50.74	41.21	92.62	86.58	143.37	127.80	371.58		
YML104C	MDM1	0.97	1.44	1.62	1.97	2.11										

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YMR032W	HOF1	0.95	1.16	1.20	1.42	1.65	1.85	22.85	8.60	71.18	36.70	94.03	45.30	269.70		
YMR034C	YMR034c	0.96	0.96	1.38	1.86	2.07	2.30	1.03	0.96	-1.54	-0.78	-0.52	0.18	295.80		
YMR035W	IMP2	0.98	1.04	1.02	1.01	1.04	1.09	123.13	121.79	123.03	131.15	246.16	252.94	520.44		
YMR036C	M1H1	1.00	1.23	1.44	1.90	2.09	2.27	2.34	1.79	35.91	18.19	38.25	19.98	295.96		
YMR037C	MSN2	0.90	1.06	1.24	1.74	1.99	2.17	1.41	0.75	22.37	5.14	23.78	5.89	280.36		
YMR038C	CCST1	0.92	1.13	1.29	1.68	1.86	1.97	3.26	1.62	33.56	7.41	36.81	8.43	276.24		
YMR039C	SUB1	0.98	1.25	1.45	1.91	2.11	2.30	2.64	0.48	32.27	14.77	34.31	15.23	313.31		
YMR040W	YEL1	1.04	1.30	1.54	1.99	2.15	2.30	12.37	1.56	57.63	31.91	70.26	33.87	326.63		
YMR041C	YAP1	1.08	1.28	1.44	1.93	2.07	2.30	2.50	1.56	47.48	2.55	51.26	2.28	228.26		
YMR042W	ARG30	1.00	1.20	1.41	1.90	2.09	2.28	10.38	1.66	47.68	26.53	66.06	27.52	399.11		
YMR044W	IOC4	0.96	1.11	1.28	1.74	1.94	2.16	8.89	14.00	55.20	68.82	64.08	82.82	300.20		
YMR048W	CMS3	0.90	0.94	0.98	1.25	1.51	1.82	24.59	14.36	79.32	73.30	103.91	87.65	338.88		
YMR052C-A	YMR052c-a	0.98	1.26	1.48	1.88	2.05	2.16	11.90	2.34	54.59	25.74	66.49	28.08	313.65		
YMR052W	FAR3	1.02	1.26	1.42	1.78	1.98	2.17	10.07	2.17	49.91	27.14	59.97	29.30	308.38		
YMR053C	STB2	0.94	1.18	1.38	2.17	1.93	2.00	17.69	1.22	64.10	14.14	81.70	15.36	328.65		
YMR054W	STV1	0.95	1.26	1.49	1.86	2.02	2.14	9.87	2.81	49.23	26.18	59.10	28.98	318.57		
YMR055C	BUB2	0.96	1.31	1.52	1.90	2.06	2.19	9.11	3.32	47.37	27.86	56.48	31.18	310.04		
YMR056C	AAC1	0.96	1.22	1.47	1.94	1.94	2.14	7.67	1.83	42.20	18.73	49.88	20.55	296.79		
YMR057C	YMR057c	0.97	1.23	1.43	1.83	2.01	2.15	7.27	3.75	43.18	30.28	50.44	34.03	290.95		
YMR058W	FET3	0.95	1.18	1.38	1.75	1.95	2.05	0.85	0.96	8.19	0.78	9.04	1.75	278.14		
YMR060C	SAM37	1.02	1.31	1.42	1.82	2.05	2.22	24.70	14.57	66.15	52.90	90.85	67.47	327.15		
YMR062C	ARG7	0.97	1.44	1.62	1.99	2.14	2.26	6.24	4.69	36.51	26.46	42.74	31.15	287.76		
YMR063W	RIM9	0.97	1.41	1.59	2.09	2.15	2.19	15.13	6.40	57.99	37.03	73.11	43.44	334.90		
YMR064W	AEP1	0.88	0.94	0.96	0.87	0.87	0.83	126.75	125.89	150.91	113.85	277.66	238.75	432.00		
YMR065W	KAR5	0.94	1.36	1.51	1.88	2.03	2.17	11.91	4.99	56.22	31.76	68.13	36.75	331.82		
YMR066W	SOV1	0.98	1.05	1.07	0.99	0.99	0.96	127.78	122.73	148.38	112.33	276.16	235.06	438.83		
YMR067C	UBX4	0.98	1.50	1.71	2.06	2.20	2.28	19.53	21.31	68.09	55.41	87.62	67.73	324.16		
YMR068W	AVO2	0.96	1.41	1.61	2.11	2.08	2.24	16.53	10.38	62.35	45.84	78.88	56.22	330.18		
YMR069W	NAT4	0.92	1.32	1.47	1.84	2.03	2.19	3.60	3.73	21.03	18.43	24.63	22.16	292.44		
YMR070W	MOT3	0.92	1.41	1.65	2.07	2.20	2.30	5.61	3.12	37.01	10.67	42.62	13.79	298.95		
YMR071C	TPV18	1.01	1.41	1.58	2.03	2.17	2.26	13.45	8.31	55.10	44.40	68.55	52.71	296.04		
YMR072W	ABF2	1.02	1.20	1.21	1.16	1.15	1.10	103.03	116.23	112.42	124.54	215.44	240.76	526.71		
YMR073C	IRC21	0.99	1.45	1.68	1.99	2.12	2.19	11.21	9.77	55.50	50.44	66.71	60.21	318.79		
YMR074C	YMR074c	0.98	1.18	1.35	1.83	2.03	2.13	9.92	8.89	58.01	46.52	67.93	53.40	302.27		
YMR075W	RCG1	0.96	1.37	1.56	1.91	2.09	2.21	9.75	12.46	48.44	52.16	58.19	64.62	313.10		
YMR077C	VPS20	0.99	1.43	1.60	1.86	2.07	2.16	20.05	12.27	76.66	68.00	96.71	80.27	377.96		
YMR078C	C16B	0.95	1.53	1.73	1.90	2.08	2.24	13.47	7.61	49.57	32.33	47.71	37.17	396.10		
YMR079C	YAM7	0.94	1.38	1.58	1.83	2.00	2.10	11.41	8.66	54.11	38.96	62.53	48.52	338.86		
YMR080C	ISF1	0.93	1.37	1.53	1.85	2.03	2.18	3.28	3.39	17.84	13.54	21.22	18.83	292.20		
YMR082C	YMR082c	0.94	1.41	1.59	1.96	2.12	2.19	4.64	3.37	31.99	19.31	36.63	22.88	299.09		
YMR083W	ADH3	0.97	1.44	1.54	1.85	2.03	2.15	3.30	3.49	39.19	20.11	43.99	21.91	255.27		
YMR084W	YMR084w	0.93	1.30	1.43	1.93	2.11	2.24	14.43	7.61	57.61	39.46	72.03	47.06	325.56		
YMR085W	YMR085w	0.94	1.31	1.47	1.92	2.10	2.23	18.37	10.11	66.26	48.84	84.63	58.95	329.88		
YMR086C-A	YMR086c-a	1.04	1.47	1.66	1.86	2.09	2.21	17.72	9.13	62.25	44.77	79.97	53.90	355.44		
YMR086W	YMR086w	0.97	1.46	1.70	2.06	2.20	2.27	13.77	11.73	58.97	50.25	72.74	61.99	319.44		
YMR087W	YMR087w	0.98	1.44	1.61	1.98	2.11	2.23	10.38	9.89	50.76	48.80	61.14	58.49	316.87		
YMR088C	VBA1	0.99	1.44	1.56	1.93	2.06	2.17	6.83	10.87	40.20	50.41	47.03	61.27	306.46		
YMR089C	YTA12	1.02	1.42	1.58	2.07	2.20	2.28	9.39	10.70	49.76	46.65	59.35	57.34	255.51		
YMR090W	YMR090w	0.99	1.43	1.61	1.94	2.09	2.21	16.56	9.52	54.37	45.39	70.92	54.90	342.84		
YMR091W	MUB1	0.94	1.43	1.56	1.88	2.03	2.20	3.88	3.40	42.88	10.46	28.75	13.86	321.54		
YMR101C	SR1	1.00	1.52	1.69	2.05	2.18	2.27	12.50	7.01	48.16	35.48	60.66	42.50	319.27		
YMR102C	YMR102c	1.04	1.43	1.62	2.04	2.18	2.27	11.33	5.05	54.62	28.95	65.95	34.00	308.39		
YMR103C	YMR103c	0.93	1.41	1.59	1.91	2.01	2.17	12.56	7.04	53.67	48.84	66.24	45.88	329.06		
YMR104C	YPK2	0.90	1.05	1.25	1.71	1.91	2.09	5.08	0.80	44.05	8.71	49.08	9.51	306.36		
YMR105C	PGM2	1.00	1.50	1.68	1.99	2.07	2.37	24.17	10.17	51.60	45.55	50.06	35.15	325.26		
YMR105W-A	YMR105w-a	0.89	1.15	1.32	1.72	1.97	2.15	1.20	2.14	15.62	19.69	17.01	21.83	251.96		
YMR106C	YKU80	0.99	1.48	1.61	1.94	2.09	2.21	5.85	3.64	34.51	19.38	40.36	23.01	304.05		
YMR107W	SPG4	0.91	1.42	1.61	1.95	2.11	2.22	10.09	4.29	44.26	22.96	54.55	27.25	311.44		
YMR109W	MYO5	0.99	1.48	1.69	2.04	2.18	2.28	10.95	5.24	50.73	32.44	61.69	38.34	294.30		
YMR110C	HRD1	0.98	1.45	1.65	2.05	2.18	2.22	4.22	4.74	42.47	24.47	51.36	31.12	320.68		
YMR111C	YMR111c	0.97	1.51	1.68	2.03	2.17	2.24	14.11	7.68	57.34	36.52	71.46	43.69	322.48		
YMR114C	YMR114c	0.97	1.51	1.68	2.03	2.17	2.24	18.15	12.20	61.72	51.63	70.87	63.83	344.60		
YMR115W	MGR3	0.95	1.54	1.75	2.05	2.16	2.24	36.75	7.88	75.65	116.63	103.89	317.42			
YMR116C	ASC1	0.98	1.37	1.42	1.72	1.99	2.05	36.75	28.24	79.88	75.55	116.63	103.89	317.42		
YMR118C	YMR118c	1.00	1.33	1.54	2.02	2.19	2.27	17.70	1.70	68.64	24.25	88.34	29.95	368.93		
YMR119C	AS1	0.92	1.06	1.23	1.69	1.89	2.09	10.06	2.83	56.28	29.46	66.34	32.29	316.13		
YMR119W-A	YMR119w-a	0.94	1.40	1.59	1.84	2.06	2.23	1.23	1.21	41.81	3.60	45.45	1.47	253.58		
YMR120C	ADE17	0.98	1.44	1.63	2.02	2.17	2.29	11.36	6.35	51.45	37.44	61.96	43.78	319.22		
YMR121C	RPL15b	0.88	1.28</													

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YMR176W	ECM5	0.95	1.08	1.19	1.63	1.89	2.15	1.68	0.77	31.42	18.94	33.10	19.70	291.91		
YMR177W	MMT1	0.98	1.19	1.37	1.81	2.02	2.23	2.34	0.49	32.43	13.49	34.77	13.98	299.40		
YMR178W	YMR178w	0.95	1.11	1.23	1.64	1.85	2.07	5.11	0.69	46.86	19.51	51.97	20.21	303.83		
YMR179W	SPT21	0.97	1.16	1.36	1.86	2.05	2.23	9.01	2.73	54.98	36.18	63.99	38.91	303.43		
YMR180C	CTL1	0.91	1.09	1.28	1.75	1.96	2.16	1.78	2.05	33.90	11.88	35.68	13.93	296.67		
YMR181C	YMR181c	0.99	1.43	1.60	2.05	2.21	2.30	13.64	6.21	52.39	34.74	66.03	40.95	327.84		
YMR182C	RGR15	0.98	1.20	1.40	1.68	2.17	2.32	1.79	0.18	8.52	0.81	10.30	1.15	285.47		
YMR182W-A	YMR182w-a	0.98	1.23	1.39	1.84	2.03	2.18	3.07	0.62	36.99	33.62	39.96	41.84	270.84		
YMR184W	AD037	0.93	1.14	1.11	1.65	1.76	1.91	14.41	1.49	56.53	12.63	72.03	10.44	334.44		
YMR188W	HSC22	0.93	1.00	1.11	1.36	1.58	1.83	0.59	2.77	16.24	10.71	18.63	13.48	293.45		
YMR187C	YMR187c	0.99	1.22	1.43	1.85	2.04	2.24	0.40	0.55	9.29	6.66	9.69	7.21	281.06		
YMR188C	MPR517	0.96	1.10	1.30	1.87	2.08	2.27	4.84	0.69	56.82	22.63	65.30	23.32	306.00		
YMR189W	GCV2	0.95	1.20	1.39	1.82	2.05	2.26	0.08	0.03	1.72	-0.34	1.80	-0.81	278.96		
YMR190C	SGS1	0.99	1.20	1.39	1.97	2.20	2.38	7.14	5.51	52.50	51.90	59.64	57.40	308.33		
YMR191W	SPGS	1.00	1.15	1.29	1.74	1.99	2.26	0.72	0.60	14.75	7.22	15.48	7.82	264.70		
YMR191W	SPGS	0.94	1.15	1.34	1.83	2.02	2.18	0.92	0.80	14.28	0.90	15.21	1.70	287.79		
YMR192W	GY1	1.02	1.26	1.49	1.94	2.12	2.30	3.43	0.33	41.46	14.05	44.89	14.39	278.34		
YMR193C-A	YMR193c-a	0.96	1.16	1.36	1.85	2.04	2.22	0.64	0.37	21.45	14.25	22.09	14.62	271.40		
YMR193W	MPR124	0.91	1.19	1.38	1.81	2.02	2.23	1.25	0.66	26.71	19.39	27.95	20.05	284.61		
YMR194C-A	YMR194c-a	0.98	1.34	1.52	1.86	2.04	2.17	17.85	6.82	56.28	32.15	74.13	38.98	338.86		
YMR194C-B	CMC4	0.92	1.21	1.32	1.73	1.94	2.08	2.37	6.22	30.61	36.43	32.98	42.65	263.50		
YMR194W	RPL36a	0.97	1.13	1.31	1.81	2.02	2.24	1.60	0.15	30.95	8.41	32.56	8.56	284.17		
YMR195W	ICY1	0.94	1.19	1.43	1.92	2.11	2.26	2.64	0.43	26.04	14.45	28.68	14.87	285.46		
YMR196W	YMR196c	0.97	1.17	1.36	1.83	2.04	2.26	0.61	0.34	22.07	15.04	22.68	15.38	287.70		
YMR198W	CIK1	0.96	1.06	1.10	1.31	1.50	1.74	49.96	4.12	102.12	10.70	104.58	152.93	446.70	407.03	
YMR198W	CIK1	0.92	0.98	1.01	1.25	1.44	1.71	43.37	2.43	104.12	8.75	147.49	112.10	348.81		
YMR199W	CLN1	0.95	1.16	1.37	1.86	2.03	2.27	0.06	0.00	5.55	1.12	5.61	1.12	280.53		
YMR201C	RAD14	0.93	1.06	1.21	1.72	1.95	2.19	5.19	2.02	47.80	26.83	52.99	28.85	286.33		
YMR202W	ERG2	1.02	1.23	1.42	1.85	2.05	2.25	6.86	0.46	52.34	12.27	59.20	12.74	308.78		
YMR202W	ERG2	0.88	1.09	1.14	1.20	1.27	1.34	40.08	29.56	93.46	63.45	133.54	93.01	407.20		
YMR204C	INP1	1.04	1.30	1.55	2.00	2.19	2.37	0.55	2.22	8.41	2.65	8.96	2.87	274.96		
YMR205C	PFK2	1.01	1.23	1.43	1.93	2.11	2.29	4.30	0.49	44.91	16.84	49.22	17.32	291.63		
YMR206W	YMR206w	1.01	1.24	1.41	1.80	2.03	2.25	0.40	0.38	13.54	5.34	13.94	5.71	271.78		
YMR207C	HFAT1	0.95	1.21	1.41	1.83	2.05	2.28	0.10	0.15	11.79	1.19	11.88	1.33	272.85		
YMR209C	YMR209c	0.92	1.36	1.53	1.94	2.10	2.19	18.63	8.16	57.70	39.95	76.53	48.10	351.91		
YMR210W	YMR210w	0.95	1.14	1.35	1.85	2.05	2.25	0.58	0.57	18.87	4.19	19.45	4.25	280.84		
YMR214W	SCD1	0.99	1.24	1.44	1.87	2.07	2.27	0.10	0.16	4.79	1.45	4.89	1.41	263.73		
YMR215C	GA35	0.88	1.10	1.29	1.56	1.76	1.95	10.13	2.67	69.22	14.84	70.35	14.71	300.00		
YMR216C	CKV1	0.99	1.17	1.42	1.89	2.07	2.23	2.62	0.36	38.36	13.51	40.88	13.87	315.88		
YMR219C	ESCI1	0.95	1.15	1.38	1.84	2.10	2.48	0.72	0.26	20.81	9.95	21.52	10.21	278.48		
YMR221C	FMD2	0.98	1.20	1.44	1.89	2.06	2.24	1.09	0.28	38.85	20.23	37.58	21.42	287.68		
YMR222C	FSH2	1.00	1.14	1.35	1.84	2.06	2.27	2.67	0.62	41.22	23.65	43.89	24.27	296.07		
YMR223W	UBP8	0.97	1.14	1.30	1.73	1.95	2.17	5.46	0.22	44.83	11.38	50.39	11.59	296.12		
YMR224C	MRE11	0.96	1.06	1.10	1.33	1.49	1.72	61.83	48.16	110.03	11.58	171.85	150.73	413.00		
YMR225C	MPR244	0.97	1.20	1.42	1.85	2.03	2.21	2.44	0.15	34.19	7.80	36.63	7.95	304.15		
YMR226C	YMR226c	0.98	1.25	1.49	1.94	2.10	2.28	4.20	1.77	45.16	15.62	49.36	17.39	304.58		
YMR228C	YMR228c	0.86	1.15	1.27	1.61	1.80	1.90	11.62	4.72	58.29	38.16	69.92	42.87	311.67		
YMR229W	RP510b	0.95	1.07	1.19	1.64	1.90	2.15	0.76	0.05	24.87	6.10	25.63	6.15	282.16		
YMR230W-A	YMR230w-a	0.89	1.21	1.36	1.75	1.94	2.08	1.23	2.21	14.71	2.52	15.94	2.73	251.43		
YMR230W-A	YMR230w-a	0.92	1.24	1.39	1.81	2.01	2.16	9.99	5.10	50.36	36.29	60.35	41.39	301.70		
YMR231W	PEPS	0.98	1.21	1.42	1.89	2.10	2.29	1.13	0.01	21.19	1.37	22.32	1.38	292.46		
YMR232W	FUS2	0.91	1.06	1.24	1.71	1.94	2.18	0.68	0.01	11.41	1.98	12.09	1.99	275.61		
YMR233W	TRI1	0.95	1.14	1.34	1.79	2.03	2.25	0.14	0.09	8.97	1.29	9.11	1.38	281.16		
YMR234W	RNH1	0.93	1.08	1.26	1.73	1.97	2.00	0.02	0.13	6.01	2.43	6.03	2.56	271.45		
YMR237W	BCH1	0.94	1.07	1.23	1.67	1.91	2.16	0.92	0.18	24.94	8.84	25.87	9.01	298.78		
YMR238W	DGF5	0.97	1.07	1.21	1.63	1.83	2.07	2.95	0.29	40.66	10.78	43.60	14.06	271.12		
YMR241W	YHM2	0.91	1.08	1.29	1.78	2.00	2.27	6.24	1.27	47.30	25.66	53.54	26.93	314.88		
YMR242C	RPL20a	0.98	1.10	1.18	1.63	1.91	2.18	10.69	3.09	61.34	41.37	72.03	44.46	312.94		
YMR242W-A	YMR242w-a	0.90	1.11	1.15	1.41	1.78	2.03	3.55	8.32	41.22	43.29	44.76	51.61	235.74		
YMR243C	ZRC1	0.92	1.07	1.33	1.83	2.03	2.23	4.04	2.09	41.42	21.38	45.46	23.48	295.87		
YMR245C	YMR245c	0.95	1.08	1.22	1.62	1.85	2.10	0.19	0.03	17.35	5.87	17.55	5.90	286.16		
YMR246W	YMR246w	0.96	1.07	1.23	1.73	1.99	2.23	5.73	0.45	49.48	17.02	55.21	17.47	299.83		
YMR247C	FAA4	1.07	1.28	1.48	1.91	2.13	2.33	0.77	0.08	26.86	10.11	27.63	10.19	305.18		
YMR247C	RKR1	0.93	1.04	1.18	1.64	1.88	2.14	0.86	0.04	10.69	3.50	10.61	3.53	285.32		
YMR247W-A	ZDS1	0.99	1.31	1.56	1.97	2.13	2.29	7.80	0.58	49.29	12.32	57.09	12.90	318.17		
YMR247C	CE1	0.96	1.32	1.57	1.95	2.15	2.42	8.50	5.48	51.03	41.55	59.53	47.03	316.50		
YMR251C	G100	0.98	1.28	1.50	1.98	2.08	2.25	3.76	0.58	43.16	17.38	46.52	17.72	304.46		
YMR252C	YMR252c	0.92	1.31	1.52	1.95	2.13	2.26	12.30	1.38	60.20	12.54	72.50	15.52	334.57		
YMR253C	YMR253c	1.01	1.31	1.52	1.95	2.14	2.29	15.29	0.80	61.93	21					

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YMR319C	FET4	0.92	1.38	1.60	1.99	2.13	2.24	11.90	4.77	51.43	27.98	63.33	32.75	318.48		
YMR320W	YMR320w	0.92	1.42	1.61	2.00	2.14	2.25	7.17	2.68	41.15	9.95	48.32	12.84	303.44		
YMR322C	SNO4	0.87	1.01	1.17	1.68	1.88	2.07	6.09	0.78	48.31	5.16	54.41	5.94	304.03		
YMR326C	YMR326c	0.91	1.29	1.50	1.87	2.03	2.16	7.79	3.68	34.79	13.88	42.58	17.75	318.37		
YNL001W	DGM34	0.91	1.20	1.36	1.74	1.89	2.03	28.79	21.78	86.12	79.25	114.91	101.03	366.35		
YNL003A	PE18	0.89	1.11	1.34	1.75	1.95	2.11	1.09	0.71	5.72	2.27	6.81	2.98	282.49		
YNL004W	HRR1	0.91	1.23	1.44	1.81	1.99	2.11	20.66	10.16	78.60	56.30	99.26	66.46	340.78		
YNL005C	MKP7	0.91	1.18	1.32	1.71	1.89	2.02	8.38	2.88	54.59	30.57	62.86	33.47	254.42		
YNL006C	ASP2	0.92	1.26	1.44	1.83	2.00	2.13	8.44	2.96	53.60	30.33	64.56	32.12	312.28		
YNL009C	UP2	0.93	1.26	1.48	1.86	2.04	2.17	15.10	5.92	67.30	48.33	82.40	54.14	336.09		
YNL010W	YL010w	0.92	1.20	1.38	1.77	1.97	2.10	20.89	6.13	77.37	49.44	98.18	55.57	348.28		
YNL011C	SP01	0.97	1.31	1.52	1.90	2.10	2.27	1.37	0.59	10.04	2.31	11.31	3.10	256.38		
YNL013W	SPO1	0.92	1.23	1.44	1.82	2.01	2.13	19.56	5.94	70.05	44.23	88.61	50.17	321.85		
YNL013C	YL013c	0.93	1.26	1.48	1.86	2.03	2.15	18.44	5.24	69.68	44.40	88.12	49.64	339.80		
YNL014W	HEF3	0.96	1.32	1.56	1.94	2.10	2.20	8.39	1.19	45.74	18.11	54.13	19.30	309.15		
YNL015W	PB2	0.98	1.26	1.44	1.74	1.92	2.06	13.35	5.54	65.04	47.65	78.39	53.19	333.33		
YNL016W	PUB1	0.95	1.47	1.68	2.06	2.22	2.30	1.86	3.21	22.88	31.57	24.74	34.79	266.59		
YNL020C	ARK1	0.91	1.24	1.43	1.80	1.98	2.12	21.66	8.31	75.17	54.90	96.83	63.41	314.90		
YNL021W	HDA1	0.91	1.25	1.44	1.86	2.07	2.22	4.85	1.66	39.34	20.00	44.19	21.66	319.21		
YNL022C	YNL022c	0.95	1.35	1.58	1.93	2.10	2.20	11.87	6.75	59.44	47.99	71.31	54.74	338.62		
YNL023C	FAP1	0.93	1.25	1.43	1.76	1.94	2.07	22.85	12.21	75.54	59.44	98.39	71.65	347.85		
YNL024C	YNL024c	0.97	1.33	1.55	1.91	2.08	2.21	13.08	5.96	61.66	46.76	74.74	52.72	332.18		
YNL025C	SSN8	0.91	1.13	1.20	1.41	1.66	1.91	2.08	10.83	2.79	55.79	29.82	89.91	76.07	122.68	98.25
YNL027W	CRZ1	0.90	1.21	1.39	1.74	1.93	2.08	24.25	18.11	81.82	70.59	106.07	88.70	307.79		
YNL028W	YL028w	0.91	1.26	1.46	1.79	1.97	2.10	7.38	4.10	50.85	39.10	58.23	43.20	306.76		
YNL029C	KTR5	0.93	1.24	1.43	1.82	2.01	2.14	23.96	11.58	82.41	63.25	106.37	74.83	295.68		
YNL030W	HHF2	0.91	1.17	1.28	1.58	1.80	1.97	32.90	7.14	91.71	71.01	124.60	88.44	253.19		
YNL031C	HHT2	0.95	1.25	1.44	1.84	2.02	2.14	18.49	7.27	72.38	49.54	90.87	56.81	257.17		
YNL032W	SW14	0.99	1.38	1.54	1.88	2.06	2.20	16.65	5.81	77.61	53.08	94.26	58.89	322.53		
YNL034W	YL034w	0.95	1.31	1.50	1.91	2.08	2.19	15.21	10.00	64.37	55.13	79.57	65.13	339.94		
YNL035C	YL035c	0.91	1.25	1.40	1.71	1.92	2.08	10.83	2.79	55.79	29.82	66.62	32.60	335.95		
YNL037C	IDH1	0.96	1.34	1.55	1.88	2.04	2.15	19.42	9.18	69.40	53.98	88.83	63.16	349.16		
YNL040W	YL040w	0.94	1.36	1.55	1.94	2.10	2.25	3.20	2.77	33.94	25.99	37.14	28.76	274.64		
YNL041C	COG6	0.94	1.24	1.41	1.68	1.85	1.98	20.75	9.58	76.05	52.37	96.80	61.95	344.62		
YNL042W-B	YNL042w-b	0.93	1.20	1.37	1.83	2.07	2.22	7.94	1.80	55.60	22.55	63.74	24.35	293.35		
YNL043C	YNL043c	0.98	1.42	1.68	2.01	2.15	2.24	16.94	8.14	75.55	48.00	92.49	56.14	325.84		
YNL044W	YID1	0.94	1.29	1.49	1.82	2.00	2.13	11.36	7.48	64.21	48.44	75.57	55.91	315.79		
YNL045W	LAP2	0.96	1.30	1.48	1.84	2.02	2.16	20.24	10.55	74.02	57.05	97.00	59.91	321.42		
YNL046W	YL046w	0.93	1.25	1.42	1.77	1.96	2.05	23.65	13.34	75.02	68.36	98.21	73.64	338.44		
YNL047C	SLM2	1.02	1.24	1.40	1.99	2.13	2.27	0.52	3.08	26.27	26.19	20.79	22.27	288.42		
YNL049C	SFB2	0.87	1.13	1.26	1.58	1.79	1.96	27.18	12.55	84.21	88.82	111.39	71.37	336.78		
YNL050C	YL050c	0.97	1.22	1.33	1.71	1.91	2.08	20.73	7.39	83.09	72.23	123.82	99.32	281.16		
YNL051W	COG5	0.99	1.14	1.36	1.95	2.14	2.28	10.09	1.76	60.30	28.52	70.39	30.28	273.92		
YNL052W	COX5a	1.03	1.07	1.16	1.67	1.84	1.96	9.46	1.17	57.24	21.89	66.70	23.06	300.32		
YNL053W	MSG5	1.01	1.27	1.44	1.97	2.20	2.35	0.52	0.02	20.58	2.97	21.09	2.98	279.89		
YNL054W	VAC7	0.88	1.18	1.36	1.75	1.96	2.14	9.41	1.32	57.81	29.68	67.22	31.60	292.71		
YNL055C	POR1	0.92	1.21	1.32	1.74	1.96	2.13	4.53	1.20	41.77	5.09	46.30	6.29	284.13		
YNL056W	OCA2	1.06	1.22	1.44	2.02	2.25	2.42	2.04	0.67	11.08	3.34	13.12	4.01	288.75		
YNL057W	YL057w	0.96	1.23	1.43	2.01	2.00	2.10	5.06	1.82	46.10	28.42	51.15	30.24	283.22		
YNL058C	YL058c	0.91	1.22	1.45	1.79	1.95	2.12	12.27	2.40	59.70	30.04	71.96	32.44	297.32		
YNL059C	ARP5	0.92	1.17	1.28	1.60	1.83	2.03	0.99	0.50	7.14	0.74	8.12	1.24	273.84		
YNL063W	MTQ1	0.92	1.25	1.38	1.77	1.95	2.10	20.87	8.00	71.36	51.58	92.24	59.59	314.36		
YNL064C	YDJ1	0.89	1.14	1.19	1.47	1.83	2.18	27.64	14.94	84.65	70.62	112.29	85.55	316.91		
YNL065W	AOR1	1.01	1.13	1.31	1.89	2.10	2.28	2.90	1.24	34.64	9.05	37.54	20.29	294.68		
YNL066W	SUN4	0.99	1.09	1.28	1.92	2.13	2.29	8.35	3.70	54.60	41.12	62.96	44.81	295.59		
YNL067W	RPL9b	0.90	1.18	1.39	1.80	2.02	2.27	2.84	0.77	38.65	10.73	41.49	11.50	296.53		
YNL067W-B	YL067w-b	0.94	1.22	1.39	1.76	2.02	2.20	0.93	0.63	9.12	2.41	10.25	3.04	261.19		
YNL068C	FKH2	0.93	1.31	1.51	1.95	2.15	2.29	1.86	1.28	20.75	9.97	22.62	11.25	268.58		
YNL069C	RPL16b	0.92	1.21	1.29	1.55	1.75	1.92	18.10	6.75	69.11	49.26	87.20	56.00	298.14		
YNL069C	RPL16b	0.89	1.12	1.19	1.51	1.74	1.89	13.54	3.20	68.11	26.91	81.65	29.83	330.70		
YNL070W	TOM7	0.89	0.99	1.15	1.78	1.92	1.98	0.91	0.68	10.11	1.59	11.02	2.27	281.37		
YNL071W	LAT1	0.91	0.98	1.13	1.70	1.92	2.13	11.80	0.86	67.74	13.51	79.54	14.37	291.04		
YNL072W	RNH201	0.99	1.14	1.32	1.89	2.06	2.24	15.52	1.45	74.30	33.89	151.56	124.19	308.78		
YNL073W	MSK1	0.98	1.17	1.16	1.67	1.90	2.12	45.27	3.40	108.29	89.80	151.56	124.19	288.44		
YNL074C	SLM2	0.93	1.15	1.34	1.73	1.94	2.10	1.15	0.89	8.10	1.36	10.24	2.27	270.80		
YNL075W	MKT1	0.96	1.13	1.32	1.86	2.02	2.18	2.06	1.03	52.91	20.10	59.97	21.16	300.42		
YNL079W	YP73	0.98	1.13	1.29	1.79	1.98	2.19	2.04	0.77	35.59	15.53	37.63	16.29	292.66		
YNL094W	APP1	0.89	1.23	1.44	1.84	2.04	2.21	0.82	0.70	11.22	6.65	12.04	7.35	292.59		
YNL095C	YL095c	1.06	1.23	1.23	1.77	1.98	2.14									

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YNL154C	YCK2	0.97	1.47	1.59	1.95	2.11	2.23	12.30	4.49	48.80	25.77	61.10	30.26	332.03		
YNL155W	YNL155w	0.98	1.40	1.49	1.89	2.08	2.21	12.70	3.97	53.69	29.18	66.38	33.14	326.03		
YNL156C	NSG2	0.95	1.38	1.51	1.90	2.05	2.17	21.77	6.95	66.41	40.10	88.18	47.05	360.16		
YNL157W	IGO1	1.01	1.49	1.64	2.01	2.15	2.26	10.05	5.30	49.65	35.76	59.70	41.06	321.85		
YNL159C	AS12	1.00	1.49	1.66	2.01	2.16	2.26	23.94	13.89	65.59	54.37	89.53	68.25	371.28		
YNL160W	YGP1	0.98	1.22	1.20	1.03	1.02	0.98	136.27	130.04	143.40	142.64	279.67	274.89	534.34		
YNL162W	RPL42a	0.98	1.52	1.72	2.01	2.17	2.17	27.72	25.58	71.64	72.85	99.36	98.43	362.93		
YNL162W	RPL42a	0.93	1.31	1.34	1.69	1.91	2.05	12.74	1.36	59.88	17.73	72.63	19.09	291.20		
YNL164A	YAL164w-a	0.94	1.44	1.61	1.88	1.99	2.09	1.64	1.64	12.41	12.65	22.63	22.63	22.63		
YNL164C	IBD2	0.94	1.37	1.40	1.88	2.07	2.18	9.85	6.13	49.38	38.49	59.04	44.61	397.61		
YNL165W	YNL165w	0.91	1.39	1.48	1.90	2.07	2.18	18.69	11.74	60.14	50.80	78.75	62.54	339.91		
YNL166C	BW1	0.86	1.16	1.11	1.27	1.78	1.98	22.87	16.88	55.58	55.58	88.66	72.26	346.78		
YNL167C	SKO1	0.92	1.36	1.43	1.77	1.92	2.05	23.52	16.43	67.36	57.22	90.88	73.65	366.17		
YNL168C	FMP41	0.94	1.39	1.47	1.86	2.05	2.20	10.14	5.87	48.16	32.52	58.30	38.39	322.09		
YNL169C	PSD1	0.98	1.58	1.73	2.08	2.19	2.30	7.68	3.14	40.50	13.14	48.18	16.28	301.27		
YNL170W	YNL170w	0.96	1.28	1.24	1.18	1.19	1.23	91.89	88.37	108.49	105.18	209.38	193.55	471.44		
YNL171C	YNL171c	0.96	1.37	1.38	1.53	1.70	1.83	69.80	63.81	100.39	95.06	170.19	158.87	456.17		
YNL173C	MDG1	0.96	1.48	1.60	1.99	2.15	2.28	11.63	5.05	50.78	28.56	62.42	33.61	326.88		
YNL175C	NOP13	0.91	1.36	1.46	1.83	2.01	2.16	3.24	3.02	19.07	10.81	22.32	13.83	285.07		
YNL176C	YNL176c	0.97	1.43	1.43	1.95	2.10	2.19	21.31	14.56	66.17	55.93	87.49	70.49	351.67		
YNL177C	MPL22	0.93	1.22	1.29	1.33	1.75	1.23	47.26	30.44	104.15	88.07	151.41	119.52	303.11		
YNL179C	YNL179c	1.00	1.32	1.40	1.82	2.01	2.18	6.87	2.99	44.63	4.65	51.50	7.64	316.91		
YNL183C	NPR1	0.95	1.40	1.54	1.99	2.15	2.27	3.32	2.93	21.60	1.83	24.92	4.76	318.23		
YNL184C	YNL184c	0.98	1.29	1.14	1.15	1.15	1.11	121.36	119.38	130.58	129.17	251.94	249.55	529.12		
YNL187W	SWT21	0.95	1.45	1.54	1.92	2.06	2.19	20.29	10.13	69.88	47.30	90.17	57.43	344.91		
YNL190W	YNL190w	0.95	1.43	1.59	1.99	2.12	2.25	7.57	4.22	41.79	27.69	49.36	31.91	329.70		
YNL191W	DUG3	0.92	1.33	1.42	1.82	1.98	2.15	8.95	3.65	46.00	20.90	54.96	24.56	325.97		
YNL192W	CHS1	1.05	1.51	1.64	2.03	2.19	2.31	21.39	10.17	68.17	52.16	89.56	62.33	331.46		
YNL193W	YNL193w	0.94	1.46	1.58	1.92	2.09	2.23	4.79	2.92	32.95	2.26	37.73	5.18	311.67		
YNL194C	YNL194c	0.93	1.44	1.52	1.86	2.02	2.13	15.79	3.46	61.03	20.17	76.82	23.63	327.98		
YNL195C	SLZ1	0.94	1.29	1.40	1.73	1.88	1.92	33.95	17.24	81.66	61.59	115.61	78.83	380.80		
YNL197C	WH3	0.98	1.38	1.47	1.88	2.08	2.22	15.13	9.80	60.52	46.46	75.65	56.26	296.71		
YNL198C	YNL198c	0.93	1.37	1.48	1.93	2.14	2.29	9.47	4.43	51.53	29.32	61.00	33.74	311.14		
YNL199C	GCR2	0.97	1.45	1.54	2.03	2.21	2.34	12.58	7.91	55.39	43.66	67.97	51.77	327.46		
YNL200C	YNL200c	1.00	1.54	1.64	2.06	2.20	2.34	6.43	4.01	40.93	17.37	47.37	25.38	319.72		
YNL201C	PSY2	0.98	1.46	1.64	2.00	2.11	2.22	12.17	8.58	52.61	45.46	64.78	54.04	328.43		
YNL202W	SAP29	0.94	1.38	1.61	1.69	2.01	2.29	8.75	5.45	48.37	33.16	57.13	38.74	327.48		
YNL203W	YNL203w	0.96	1.56	1.62	1.99	2.11	2.25	13.70	5.73	53.98	34.60	67.06	40.71	345.21		
YNL204C	SPB16	0.91	1.41	1.52	1.82	2.00	2.15	2.63	1.65	6.21	1.65	9.04	1.74	287.16		
YNL205C	YNL205c	0.98	1.60	1.75	2.14	2.25	2.31	2.98	3.02	13.35	3.99	16.33	7.60	283.39		
YNL206C	RTT69	0.95	1.49	1.59	2.09	2.15	2.26	4.49	2.92	14.36	7.30	38.94	19.32	291.16		
YNL208W	YNL208w	1.01	1.58	1.72	2.08	2.21	2.34	2.94	3.26	8.57	10.73	11.50	13.94	282.43		
YNL209W	SSB2	0.91	1.35	1.49	1.88	2.07	2.22	4.43	1.16	42.23	14.91	46.67	16.07	286.88		
YNL211C	YNL211c	0.91	1.33	1.47	1.88	2.05	2.22	3.20	3.68	18.73	11.81	21.93	14.89	296.38		
YNL212W	VID27	0.95	1.38	1.44	1.91	2.07	2.21	15.93	9.31	62.41	50.99	78.35	60.29	334.68		
YNL213C	RRG9	0.93	1.26	1.19	1.12	1.10	1.14	111.64	109.72	121.84	121.66	233.48	231.39	525.71		
YNL214W	PEX17	0.89	1.39	1.47	1.84	1.99	2.13	21.63	12.84	68.92	56.86	90.56	71.90	341.38		
YNL215W	IES2	0.93	1.35	1.34	1.70	1.81	2.02	37.63	28.00	88.27	79.33	125.90	107.33	360.75		
YNL217W	YNL217w	0.93	1.38	1.49	1.87	2.01	2.17	11.45	6.82	54.62	38.03	66.07	30.82	322.35		
YNL218W	MGS1	0.88	1.14	1.20	1.58	1.78	1.95	13.14	1.30	63.82	11.35	76.97	12.65	311.84		
YNL219C	ALG9	1.09	1.56	1.76	2.17	2.31	2.40	18.63	11.07	65.72	53.99	84.35	65.06	342.62		
YNL220W	ADE12	1.12	1.12	1.29	1.38	1.44	1.52	1.51	1.56	-1.31	-1.28	0.20	0.28	411.15		
YNL220W	ADE12	1.03	1.03	1.17	1.24	1.29	1.38	1.26	1.45	-1.51	-1.19	-0.25	0.27	512.96		
YNL220W	ADE12	0.89	0.97	0.99	1.04	1.08	1.17	198.72	153.66	148.55	140.86	347.27	294.52	490.72		
YNL220W	ADE12	0.94	1.00	1.04	1.03	1.03	1.04	116.64	117.78	134.85	132.83	254.49	250.61	519.42		
YNL220W	ADE12	0.94	1.01	1.04	1.03	1.04	1.05	117.35	132.94	130.39	141.68	247.74	274.62	512.38		
YNL220W	ADE12	0.93	1.01	1.04	1.03	1.04	1.05	113.67	130.92	130.00	140.18	243.67	272.21	519.70		
YNL220W	ADE12	0.91	1.01	1.12	1.21	1.24	1.32	118.49	121.89	121.89	121.89	252.18	250.03	527.84		
YNL220W	ADE12	1.08	1.09	1.11	1.11	1.19	1.24	1.31	1.31	130.05	128.78	133.41	142.21	263.27	270.99	
YNL220W	ADE12	1.00	1.15	1.15	1.25	1.29	1.35	1.41	1.41	122.46	122.22	134.90	137.00	257.36	260.22	
YNL220W	ADE12	1.07	1.19	1.21	1.30	1.37	1.41	141	141	123.46	123.46	125.87	125.87	319.25		
YNL220W	ADE12	0.94	1.02	1.01	1.08	1.13	1.15	1.15	1.15	105.68	127.11	114.48	127.45	220.43	230.32	
YNL220W	ADE12	0.95	1.07	1.08	1.17	1.23	1.24	115.75	114.60	119.06	119.59	234.81	234.19	504.47		
YNL220W	ADE12	1.02	1.17	1.17	1.29	1.35	1.35	119.33	127.67	126.21	133.38	245.54	261.05	523.09		
YNL220W	ADE12	0.98	1.06	1.07	1.05	1.15	1.17	114.38	115.30	117.95	122.30	232.34	232.34	520.45		
YNL220W	ADE12	1.00	1.15	1.16	1.25	1.32	1.33	116.20	124.42	126.82	132.68	243.01	249.68	566.78		
YNL220W	ADE12	0.97	1.04	1.03	1.04	1.15	1.16	112.01	126.77	126.56	126.56	243.01	249.85	524.34		
YNL220W	ADE12	1.05	1.16	1.16	1.24	1.28</td										

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YNL220W	ADE12	0.85	1.10	1.12	1.08	1.15	1.17	103.75	92.37	128.09	123.32	231.84	215.69	288.35		
YNL220W	ADE12	0.92	1.08	1.17	1.11	1.16	1.12	122.88	120.67	129.80	129.80	252.68	250.47	423.99		
YNL220W	ADE12	0.92	0.93	0.97	1.07	1.10	1.09	145.34	124.43	151.62	130.12	296.96	254.55	208.24		
YNL220W	ADE12	1.05	1.09	1.14	1.26	1.31	1.30	147.34	145.19	151.06	149.83	298.40	295.02	384.68		
YNL220W	ADE12	0.92	0.96	0.99	1.03	1.08	1.08	128.43	121.12	133.00	128.48	261.43	249.60	371.77		
YNL220W	ADE12	0.98	1.01	1.04	1.11	1.13	1.08	119.57	117.61	129.06	125.25	246.83	242.86	424.76		
YNL220W	ADE12	0.92	1.00	1.02	1.03	1.08	1.08	127.10	155.63	122.67	120.89	259.97	312.01	343.62		
YNL220W	ADE12	1.08	1.08	1.14	1.16	1.19	1.19	137.63	122.85	141.94	131.88	279.54	254.74	433.22		
YNL220W	ADE12	1.00	1.08	1.14	1.18	1.26	1.20	123.24	122.55	130.93	131.57	257.51	251.44	411.41		
YNL220W	ADE12	1.00	1.08	1.14	1.18	1.24	1.21	127.57	128.69	131.74	132.24	259.31	261.93	435.00		
YNL220W	ADE12	1.05	1.25	1.29	1.30	1.35	1.34	120.30	117.45	125.32	121.70	245.62	238.15	478.01		
YNL220W	ADE12	0.96	1.09	1.15	1.13	1.21	1.19	137.62	138.52	137.26	138.57	274.87	278.09	408.44		
YNL220W	ADE12	0.98	1.10	1.17	1.16	1.23	1.21	136.25	142.12	138.09	144.59	274.34	286.71	458.80		
YNL220W	ADE12	0.95	1.06	1.09	1.07	1.11	1.12	120.16	111.74	123.07	116.49	243.23	228.24	468.06		
YNL220W	ADE12	1.02	1.20	1.29	1.59	1.81	1.98	34.05	13.90	82.16	80.37	116.20	74.27	382.38		
YNL220W	ADE12	0.98	1.10	1.18	1.14	1.17	1.17	116.08	112.48	137.49	106.21	253.57	219.69	496.51		
YNL220W	ADE12	0.98	1.09	1.14	1.10	1.15	1.16	117.55	114.25	146.64	110.71	264.19	224.95	478.50		
YNL220W	ADE12	0.93	1.15	1.14	1.10	1.14	1.15	118.79	115.24	147.26	114.19	266.04	228.44	512.85		
YNL220W	ADE12	0.92	1.12	1.17	1.11	1.14	1.14	115.67	33.22	140.37	9.43	256.04	42.66	513.37		
YNL220W	ADE12	0.94	1.14	1.15	1.08	1.11	1.12	116.85	116.74	124.63	93.95	241.48	212.09	517.28		
YNL220W	ADE12	0.92	1.08	1.13	1.02	1.06	1.08	125.36	120.59	106.79	140.74	232.16	261.33	497.01		
YNL220W	ADE12	0.91	1.05	1.11	1.01	1.03	1.06	119.29	122.78	124.68	138.94	243.97	261.72	509.86		
YNL220W	ADE12	0.97	1.15	1.23	1.15	1.21	1.20	126.62	118.81	145.66	143.57	272.29	262.39	511.57		
YNL220W	ADE12	0.90	1.25	1.25	1.13	1.16	1.16	110.12	120.88	132.20	142.97	242.32	263.85	507.09		
YNL220W	ADE12	0.94	1.28	1.29	1.18	1.22	1.21	132.00	120.46	145.87	141.62	277.88	262.08	518.36		
YNL220W	ADE12	0.94	1.18	1.20	1.13	1.16	1.18	123.24	121.26	146.62	142.07	269.85	263.33	516.38		
YNL220W	ADE12	0.88	1.21	1.24	1.20	1.25	1.25	115.23	111.55	126.86	129.47	242.09	241.02	498.93		
YNL220W	ADE12	0.97	1.14	1.15	1.10	1.15	1.15	113.76	115.47	134.67	131.80	248.44	247.28	518.41		
YNL220W	ADE12	0.89	1.12	1.14	1.07	1.11	1.12	118.66	111.55	135.57	130.02	254.23	241.58	513.07		
YNL220W	ADE12	0.89	1.26	1.26	1.23	1.28	1.28	118.23	131.11	132.96	124.73	251.19	237.84	496.02		
YNL220W	ADE12	0.92	1.17	1.18	1.13	1.20	1.17	121.36	116.79	136.37	132.53	257.73	248.32	508.11		
YNL220W	ADE12	0.98	1.19	1.24	1.14	1.23	1.22	116.62	114.82	135.36	131.63	251.99	246.15	521.01		
YNL220W	ADE12	0.91	1.17	1.21	1.15	1.18	1.18	112.61	114.82	129.00	125.68	241.62	240.51	508.30		
YNL220W	ADE12	0.93	1.15	1.17	1.08	1.13	1.09	110.93	119.48	123.61	119.59	239.54	240.07	517.53		
YNL220W	ADE12	0.91	1.17	1.17	1.11	1.14	1.14	114.52	115.22	135.22	123.70	253.79	242.56	523.75		
YNL220W	ADE12	0.87	1.13	1.15	1.07	1.08	1.08	119.48	119.68	120.30	123.45	247.79	241.41	506.00		
YNL220W	ADE12	0.92	1.16	1.19	1.13	1.16	1.12	121.35	116.97	121.45	120.40	242.80	237.37	507.28		
YNL220W	ADE12	0.90	1.18	1.20	1.11	1.16	1.16	120.98	113.43	131.70	118.91	252.66	222.34	519.44		
YNL220W	ADE12	0.87	1.07	1.07	0.95	1.05	1.00	114.74	117.12	119.46	118.76	234.31	238.85	507.24		
YNL220W	ADE12	0.88	1.07	1.08	0.98	1.04	0.96	113.23	119.55	120.19	125.23	233.42	244.78	516.31		
YNL220W	ADE12	0.94	1.15	1.16	1.11	1.18	1.12	120.51	114.82	130.51	118.17	251.02	232.98	521.42		
YNL220W	ATG4	0.93	1.37	1.50	1.87	2.04	2.15	19.33	3.93	66.72	25.80	86.05	29.71	326.31		
YNL224C	SOS1	0.96	1.40	1.52	1.86	2.03	2.16	10.05	3.48	56.90	19.94	66.95	23.43	339.12		
YNL225C	CMM67	0.96	1.04	1.07	1.02	1.09	1.08	153.44	41.36	113.39	81.20	167.33	122.56	338.76		
YNL226W	YNL226w	0.98	1.41	1.51	1.93	2.10	2.25	3.11	3.21	12.07	7.56	15.18	10.77	273.67		
YNL227C	JUJ1	0.93	1.37	1.44	1.82	1.97	2.13	19.08	3.26	66.54	13.32	85.62	16.58	345.34		
YNL228W	YNL228w	0.92	1.32	1.40	1.81	1.98	2.13	9.69	7.70	52.93	43.67	62.62	51.37	307.56		
YNL229C	URE2	0.96	1.42	1.46	1.86	2.14	2.25	15.76	7.39	64.26	46.09	80.02	53.49	327.45		
YNL230C	ELA1	0.96	1.45	1.59	1.98	2.13	2.25	14.93	5.48	59.22	35.14	74.15	40.62	342.93		
YNL231C	PDR16	0.93	1.46	1.58	1.95	2.10	2.25	3.93	3.05	25.05	8.04	28.98	11.09	303.32		
YNL233W	BIN4	0.95	1.44	1.54	1.99	2.15	2.29	17.77	11.48	65.56	54.11	83.33	65.59	306.29		
YNL234W	YNL234w	0.95	1.43	1.54	1.91	2.04	2.17	18.60	12.15	62.00	49.27	80.60	61.42	329.87		
YNL235C	YNL235c	1.00	1.52	1.64	2.11	2.25	2.37	11.53	6.29	55.38	40.37	66.90	46.65	327.81		
YNL236C	MPA43	0.98	1.36	1.61	1.96	2.17	2.27	2.45	2.47	56.15	40.91	60.01	30.02	305.33		
YNL236W	IST1	0.95	1.42	1.69	2.02	2.20	2.29	17.46	1.94	56.15	45.17	62.78	43.24	309.31		
YNL237W	BOR1	0.98	1.50	1.74	2.01	2.20	2.29	1.99	0.07	32.06	3.25	34.05	3.33	282.00		
YNL237C	YLP1	0.98	1.30	1.47	1.89	2.09	2.25	3.04	-0.24	40.64	1.31	43.68	1.06	319.19		
YNL238W	YLP1	0.97	1.23	1.49	1.93	2.10	2.22	1.93	4.25	28.35	37.73	30.28	41.98	279.81		
YNL239W	BSC4	0.92	1.39	1.56	1.95	2.15	2.30	1.14	-0.07	5.32	1.19	6.47	1.11	286.93		
YNL239W	CUS2	1.01	1.29	1.53	1.97	2.15	2.30	7.10	0.92	50.95	24.15	58.05	25.08	314.43		
YNL239W	CFA40	1.02	1.39	1.65	2.03	2.18	2.31	8.90	0.32	56.58	15.65	65.28	15.97	330.06		
YNL239W	PCY1	1.04	1.41	1.68	2.09	2.24	2.34	6.86	0.30	49.04	10.31	55.89	10.61	298.85		
YNL239W	MDB1	0.98	1.30	1.51	1.95	2.14	2.26	13.59	4.58	60.75	40.69	74.34	45.26	317.22		
YNL239W	MPG1	1.00	1.27	1.43	1.92	2.14	2.28	8.40	0.07	54.82	24.67	63.22	22.64	322.42		
YNL239W	MSB3	1.00	1.27	1.43	1.92	2.14	2.28	5.08	0.15	46.56	5.14	51.64	8.80	295.12		
YNL239W	RIM21	0.98	1.23	1.45	1.90	2.09	2.24	2.38	0.06	-0.07	10.01	0.84	10.07	0.77	268.26	
YNL239W	YNL239w	0.97	1.26	1.50	1.98	2.16	2.26	2.33	-0.06	-0.07						

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YNL339W	SN2Z	1.06	1.35	1.59	2.02	2.17	2.33	10.48	0.77	57.34	21.61	67.82	22.38	317.17		
YNL334C	SN02	0.98	1.16	1.37	1.80	2.00	2.23	3.61	0.55	37.13	6.95	40.74	7.50	299.03		
YNL335W	DD13	1.05	1.35	1.55	1.96	2.12	2.30	6.44	0.59	49.49	18.14	55.93	18.73	305.99		
YNL336W	COST	1.01	1.32	1.59	1.96	2.11	2.29	16.99	0.69	65.05	18.02	82.05	18.72	337.29		
YNL339W	YNL338w	1.01	1.32	1.58	1.96	2.11	2.30	9.82	0.32	54.74	12.96	64.56	13.27	317.55		
YNL336E	YRF1-6	0.99	1.28	1.57	1.97	2.14	2.30	9.7	0.22	56.00	8.21	65.78	8.44	317.70		
YNR001C	CIIA	0.98	1.30	1.48	1.86	2.06	2.19	8.87	0.19	54.70	24.09	63.58	25.86	333.77		
YNR002C	AT02	0.93	1.29	1.49	1.89	2.05	2.17	26.06	1.65	84.34	59.30	110.40	70.95	349.75		
YNR003C	SW02	1.14	1.44	1.54	1.79	2.01	2.16	26.44	1.25	82.24	51.35	110.35	70.86	339.50		
YNR005C	YNR005c	0.93	1.28	1.46	1.72	1.86	1.99	21.10	0.63	93.09	114.19	79.88	114.19	359.50		
YNR009W	VPS27	0.97	1.28	1.42	1.85	1.77	1.90	24.56	7.49	91.42	64.89	115.98	72.38	377.33		
YNR007C	ATG3	0.91	1.25	1.43	1.77	1.98	2.11	30.09	1.97	97.73	68.67	98.98	137.76	89.71	360.03	
YNR008W	LR01	0.90	1.23	1.42	1.79	1.98	2.11	14.45	4.13	67.20	39.51	81.65	43.64	308.86		
YNR009W	NRM1	0.91	1.23	1.38	1.75	1.93	2.07	21.73	1.80	76.26	99.32	94.86	121.56	321.56		
YNR010W	CSE2	0.91	1.10	1.14	1.20	1.47	1.74	61.84	40.35	119.14	98.04	180.99	138.39	330.22		
YNR010W	CSE2	0.86	1.20	1.37	1.82	2.00	2.16	3.71	1.47	32.54	14.08	36.25	15.55	265.81		
YNR012W	URK1	0.90	1.21	1.39	1.75	1.95	2.11	14.37	7.69	65.64	46.69	80.01	53.78	323.58		
YNR013C	PHO31	0.97	1.25	1.43	1.87	2.07	2.21	10.47	7.22	63.83	53.70	74.30	60.92	324.42		
YNR014W	YNR014w	0.96	1.33	1.50	1.83	2.02	2.17	9.18	2.68	54.43	31.03	63.62	33.71	302.24		
YNR015W	SMM1	0.90	1.23	1.39	1.77	1.95	2.09	19.47	7.46	72.83	52.60	92.30	60.06	349.70		
YNR018W	AIM38	0.88	1.20	1.40	1.76	1.94	2.07	14.23	4.60	62.66	40.02	76.89	44.62	342.42		
YNR019W	ARE2	0.92	1.30	1.49	1.84	2.04	2.17	17.17	9.49	73.58	58.93	90.74	68.42	342.52		
YNR020C	ATP23	0.96	1.37	1.57	1.88	2.04	2.17	9.53	5.75	59.36	44.12	68.88	49.87	315.27		
YNR021W	YNR021w	0.96	1.28	1.45	1.72	1.89	2.05	26.78	13.68	86.05	65.27	112.82	78.95	337.34		
YNR022C	MRLP150	0.91	1.25	1.46	1.81	1.99	2.11	17.44	7.86	71.53	52.43	88.97	60.29	351.23		
YNR024W	MPP6	0.94	1.26	1.42	1.76	1.94	2.09	10.46	5.24	60.43	42.84	70.89	48.07	302.94		
YNR025C	YNR025c	0.87	1.11	1.22	1.58	1.83	2.00	17.26	10.28	73.66	63.03	90.92	73.31	304.92		
YNR027W	BUD17	0.92	1.25	1.42	1.76	1.96	2.10	15.96	9.31	69.80	54.14	85.76	63.45	307.52		
YNR028W	CPR8	0.92	1.24	1.41	1.78	1.97	2.10	16.78	7.27	66.12	51.34	82.90	58.61	313.16		
YNR029C	YNR029c	0.95	1.21	1.32	1.71	1.94	2.12	7.99	3.53	56.21	38.59	64.20	42.13	313.30		
YNR030W	ALG12	0.99	1.30	1.39	1.70	1.89	2.06	16.57	6.69	67.34	45.11	83.91	51.80	297.53		
YNR031C	SSK2	0.91	1.24	1.34	1.70	1.89	2.06	18.10	6.58	75.30	48.26	93.40	54.84	329.14		
YNR033C-A	HUB1	0.91	1.32	1.53	1.87	2.02	2.15	9.11	4.76	39.28	22.97	48.39	27.73	292.77		
YNR032W	PPGT	0.93	1.27	1.39	1.72	1.92	2.08	29.71	16.41	93.06	80.32	122.77	96.73	358.51		
YNR033W	AB21	0.98	1.22	1.41	1.86	2.05	2.23	1.38	0.99	22.41	13.68	23.79	14.86	271.32		
YNR034W	SOL7	0.92	1.22	1.39	1.73	1.92	2.07	22.71	10.76	77.36	55.18	100.06	65.95	339.28		
YNR036C	MSPS12	0.90	1.28	1.45	1.74	1.96	2.00	1.36	1.64	12.21	3.62	13.58	4.06	272.47		
YNR037C	RIS9	0.91	1.23	1.38	1.74	1.94	2.07	14.68	4.56	46.51	34.39	51.61	38.06	183.59		
YNR038C	ZRG17	0.98	1.30	1.58	1.93	2.01	2.16	10.47	7.16	53.71	32.93	68.18	38.15	228.44		
YNR039C	YHR240w	0.95	1.20	1.30	1.71	1.91	2.04	2.23	5.43	2.07	62.29	24.14	20.21	25.84	229.44	
YNR041C	COQ2	0.92	1.34	1.51	1.86	2.03	2.16	4.16	3.51	44.81	40.65	48.97	44.16	262.62		
YNR042W	YHR242c	0.98	1.17	1.34	1.75	1.96	2.10	24.00	6.98	76.23	50.22	101.13	69.99	284.43		
YNR043W	PET484	0.90	1.28	1.45	1.77	1.97	2.12	6.06	2.35	45.75	24.05	51.81	26.29	314.45		
YNR047W	FPK1	0.98	1.26	1.42	1.72	1.91	2.09	16.90	8.84	70.84	61.11	87.74	69.94	266.12		
YNR048W	YNR048w	0.98	1.39	1.57	1.94	2.12	2.23	26.81	17.82	73.90	72.90	105.87	90.73	247.49		
YNR049C	MS01	0.89	1.25	1.40	1.76	1.95	2.10	22.33	10.02	80.87	65.80	103.29	82.82	191.72		
YNR050C	LYS9	0.91	1.27	1.37	1.63	1.82	2.00	1.37	2.29	8.07	1.45	9.44	2.74	303.86		
YNR051C	BRE5	0.93	1.33	1.51	1.91	2.09	2.22	12.47	4.91	49.22	27.18	61.69	32.10	332.32		
YNR052C	POP2	0.91	1.12	1.16	1.34	1.76	1.76	63.11	52.22	108.34	76.06	171.46	128.28	391.34		
YNR052C	POP2	0.87	1.11	1.15	1.32	1.56	1.86	42.09	34.80	101.02	93.50	143.11	128.31	347.86		
YNR055C	HOL1	0.91	1.06	1.26	1.75	1.94	2.10	3.10	1.18	36.35	17.39	39.54	18.57	305.50		
YNR056C	BIO5	0.99	1.41	1.61	1.97	2.12	2.24	23.49	9.89	61.25	43.85	84.74	53.74	364.17		
YNR057C	BIO4	1.01	1.33	1.49	1.89	2.07	2.27	11.66	4.45	47.52	27.99	59.18	32.44	320.21		
YNR058W	BI03	0.99	1.42	1.62	1.98	2.18	2.31	1.78	1.34	21.96	2.74	23.74	4.08	258.70		
YNR059W	MNT9	0.94	1.30	1.47	1.83	2.03	2.18	8.25	5.10	41.15	30.59	49.40	35.69	309.26		
YNR060W	FRE4	0.90	1.25	1.42	1.80	2.00	2.18	7.65	4.04	37.35	19.77	45.00	23.80	304.43		
YNR061C	YNR061c	0.91	1.30	1.49	1.83	2.03	2.18	10.12	3.10	44.57	27.71	57.19	24.32	243.78		
YNR062C	YNR062c	0.92	1.29	1.45	1.83	2.01	2.14	10.67	7.12	56.31	33.50	72.38	40.62	336.93		
YNR063W	YNR036w	0.97	1.38	1.57	1.94	2.10	2.20	13.71	9.18	53.70	42.51	67.40	51.69	326.45		
YNR064C	YHR243c	0.98	1.38	1.54	1.95	2.03	2.20	10.21	5.21	48.55	38.49	58.76	45.28	336.96		
YNR065W	AIF1	1.03	1.36	1.56	2.01	2.27	2.30	15.28	7.58	48.57	30.21	45.78	47.79	311.47		
YNR067W	COS10	1.01	1.61	1.88	2.16	2.27	2.33	6.21	7.58	39.57	40.21	45.78	27.87	278.77		
YOL019W	PHO80	0.94	1.15	1.24	1.85	2.14	2.33	21.34	14.49	83.76	68.72	105.19	83.21	362.49		
YOL020C	LAG2	0.93	1.41	1.67	2.07	2.22	2.36	1.07	0.96	27.68	24.38	28.75	25.34	295.29		
YOL020C	PLB3	0.98	1.39	1.62	2.04	2.19	2.34	6.21	4.09	44.05	23.00	50.26	23.69	301.40		
YOL021C	HTZ1	1.04	1.19	1.29	1.72	1.93	2.10	38.74	20.65	88.92	70.57	127.65	91.23	370.43		
YOL023C	HRD1	0.95	1.35	1.61	1.98	2.08	2.23	5.89	0.22	47.29	14.57	53.18	14.80	304.60		
YOL019W-A	YOL019w-a	0.98	1.44	1.63	1.98	2.14	2.22	8.97	1.39	38.69	10.34	57.78	70.60	308.47		
YOL019W-B	YOL019w-b	0.9														

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YOL063C	CRT10	0.96	1.36	1.62	2.00	2.13	2.36	13.97	0.16	60.49	10.20	74.46	10.37	340.37		
YOL064C	MET22	0.80	0.98	1.14	1.71	1.88	2.04	23.72	4.99	74.50	33.83	98.22	38.82	321.92		
YOL065C	INP54	0.95	1.20	1.40	1.88	2.03	2.18	15.83	0.16	64.90	4.87	80.73	5.03	331.88		
YOL067C	RTGT	0.94	1.18	1.37	1.94	2.11	2.26	17.99	0.21	72.61	8.97	90.60	9.19	348.46		
YOL068C	RTGT	0.98	1.23	1.41	1.82	2.07	2.23	4.55	1.55	39.04	14.96	43.59	16.51	287.97		
YOL069C	NBT1	0.96	1.33	1.57	1.95	2.11	2.25	17.06	-0.22	67.85	8.41	84.91	8.19	331.85		
YOL070C	NBT1	0.94	1.30	1.53	1.99	2.14	2.27	13.85	0.18	62.44	5.85	76.29	6.13	324.92		
YOL071W	EMR5	0.83	1.18	1.39	1.92	2.06	2.21	16.32	0.23	68.54	7.42	84.75	8.11	332.48		
YOL072C	MP07	0.96	1.24	1.47	1.94	2.14	2.29	18.29	0.21	49.91	7.94	52.02	7.72	400.26		
YOL073C	MP07	0.90	1.22	1.41	1.76	1.94	2.07	3.24	0.59	34.59	9.66	37.83	10.59	277.05		
YOL075C	YOL075c	1.01	1.31	1.55	1.99	2.14	2.29	19.19	6.16	65.30	11.00	84.48	11.15	347.41		
YOL076W	MDM20	1.00	1.15	1.22	1.89	1.98	2.26	20.03	2.22	74.03	36.82	94.05	38.94	258.78		
YOL077W-A	ATP19	0.89	1.09	1.21	1.63	1.90	2.09	5.63	3.29	48.84	28.43	54.47	31.72	295.27		
YOL079W	YOL079w	0.96	1.22	1.43	1.87	2.02	2.18	10.76	0.32	58.45	13.32	69.22	13.65	321.45		
YOL080C	REX4	0.98	1.19	1.39	1.94	2.10	2.28	14.50	0.33	66.42	16.53	80.92	16.87	323.77		
YOL081W	IR42	0.98	1.29	1.46	1.87	2.03	2.31	-0.02	0.02	8.33	-0.67	8.31	-0.65	341.87		
YOL082W	ATG19	0.83	1.16	1.40	1.87	2.02	2.18	4.84	-0.11	45.58	0.97	50.42	0.86	312.47		
YOL083W	YOL083w	0.93	1.21	1.46	2.06	2.04	2.15	12.53	0.15	67.27	5.92	79.81	6.06	336.71		
YOL084W	PHM7	0.91	1.18	1.40	1.88	2.02	2.18	15.17	0.02	64.87	4.68	80.04	4.70	335.19		
YOL085C	YOL085c	0.94	1.24	1.47	1.94	2.07	2.21	15.84	0.12	62.45	5.58	78.29	5.70	335.02		
YOL086C	ADH1	0.92	1.18	1.15	1.02	1.01	0.97	124.37	132.15	140.34	143.68	264.71	275.83	525.56		
YOL086C	ADH1	0.86	1.02	1.02	0.92	1.02	1.02	107.56	104.63	134.67	126.48	242.24	231.11	497.87		
YOL086C	ADH1	0.87	1.10	1.07	0.97	1.10	1.27	110.13	101.83	123.77	115.61	233.90	217.44	499.47		
YOL086W-A	YOL086w-a	0.93	1.15	1.31	1.67	1.92	2.11	3.57	2.28	40.42	15.60	43.99	16.88	280.33		
YOL087C	YOL087c	0.98	1.51	1.75	2.15	2.27	2.35	15.07	4.21	53.52	15.21	68.60	19.42	314.26		
YOL088C	MPD2	0.88	1.30	1.47	1.87	2.01	2.04	10.86	7.46	50.83	40.28	61.69	47.73	319.89		
YOL089C	HAL9	0.93	1.40	1.57	1.96	2.11	2.20	14.93	7.58	57.52	42.81	72.24	50.39	342.43		
YOL090W	MSH2	0.91	1.45	1.62	1.96	2.09	2.18	18.15	8.43	60.72	45.23	78.87	53.66	346.23		
YOL091W	SP021	0.94	1.49	1.67	2.04	2.15	2.25	17.90	10.15	64.08	49.37	81.98	59.52	346.52		
YOL092W	YOL092w	0.97	1.50	1.69	2.06	2.17	2.26	17.64	7.02	58.96	37.27	76.60	44.29	347.30		
YOL093W	TRM10	1.00	1.47	1.63	2.04	2.18	2.28	14.96	8.70	58.38	47.86	73.33	56.56	334.63		
YOL095C	HMT1	0.95	1.37	1.52	1.93	2.11	2.24	6.44	2.98	38.47	10.91	44.92	13.89	297.73		
YOL096C	COQ3	0.82	1.00	0.97	1.15	1.52	1.79	51.96	36.49	88.70	84.96	140.66	121.45	360.92		
YOL097W-A	YOL097w-a	0.92	1.18	1.31	1.73	1.97	2.11	10.90	4.02	60.45	30.86	71.34	34.88	298.64		
YOL098C	YOL098c	0.94	1.44	1.60	2.02	2.18	2.28	3.80	2.97	21.67	6.00	25.67	8.98	265.22		
YOL099C	YOL099c	0.88	1.29	1.41	1.81	1.99	2.13	8.87	3.02	42.68	12.90	51.75	16.92	316.31		
YOL100W	PKH2	0.85	1.09	1.08	1.21	1.51	1.72	12.21	9.61	53.56	45.77	56.23	55.54			
YOL101C	ZTH4	0.94	1.46	1.64	2.04	2.20	2.31	3.35	2.93	44.65	2.63	18.00	5.43	230.40		
YOL102W	YOL102w	0.98	1.30	1.48	1.88	2.04	2.14	12.10	9.26	49.63	33.92	51.85	43.18	324.40		
YOL104C	NDJ1	0.94	1.46	1.62	2.07	2.20	2.30	6.67	3.43	40.97	18.21	49.64	17.65	305.43		
YOL105C	WSCI	0.88	1.27	1.42	1.83	2.01	2.11	15.61	5.24	31.54	6.60	68.20	36.78	339.36		
YOL106W	YOL106w	0.98	1.26	1.35	1.95	2.13	2.21	14.21	9.07	52.46	44.81	68.67	54.77	311.93		
YOL107W	YOL107w	0.89	1.28	1.51	1.92	2.09	2.21	12.54	3.03	49.70	15.39	62.24	18.42	332.43		
YOL108C	IM04	0.90	1.11	1.12	1.23	1.72	1.84	28.54	14.68	69.73	56.18	98.27	70.86	259.19		
YOL108C	IM04	0.87	1.14	1.16	1.32	1.60	1.71	27.77	15.88	67.95	70.98	115.73	86.86	241.64		
YOL109W	ZFO1	0.93	1.36	1.52	1.91	2.09	2.22	12.64	3.91	48.79	24.14	61.43	20.05	331.15		
YOL110W	SHR5	0.96	1.41	1.56	1.98	2.17	2.30	3.69	3.01	18.91	4.07	22.60	7.08	298.43		
YOL111C	MDY2	0.90	1.35	1.51	1.94	2.10	2.19	11.08	4.21	42.95	22.79	54.03	27.00	322.04		
YOL112W	MSB4	0.91	1.48	1.63	2.00	2.15	2.25	6.67	3.29	36.42	13.19	43.09	16.48	314.81		
YOL113W	SKM1	0.92	1.38	1.53	1.95	2.13	2.25	12.98	5.86	51.02	33.19	64.00	39.04	325.95		
YOL114C	YOL114c	0.97	1.50	1.73	2.16	2.28	2.35	5.06	3.20	29.15	10.55	34.22	13.74	286.55		
YOL115W	PAP2	0.92	1.37	1.56	1.95	2.12	2.23	15.39	4.21	54.32	26.36	69.70	30.57	339.69		
YOL116W	MSN1	0.87	1.28	1.48	1.88	2.07	2.20	4.00	3.15	22.14	9.68	26.14	12.83	284.33		
YOL117W	RR12	0.88	1.32	1.56	1.96	2.13	2.24	6.61	3.05	38.03	6.76	44.64	9.81	316.86		
YOL118C	YOL118c	0.88	1.28	1.46	1.89	2.08	2.21	5.01	3.16	30.15	8.06	35.16	11.22	303.10		
YOL119C	MCH4	0.89	1.31	1.56	1.97	2.13	2.24	3.34	3.17	17.60	8.26	20.93	11.43	291.20		
YOL121C	RPS19a	0.92	1.20	1.33	1.78	2.02	2.13	10.67	4.09	46.42	16.82	56.28	22.05	311.33		
YOL122C	SMF1	0.86	1.21	1.40	1.66	2.06	2.18	9.57	3.67	46.00	21.14	55.57	24.81	312.98		
YOL123W	RTCT	0.85	1.23	1.40	1.74	1.95	2.08	18.10	4.96	60.07	32.52	78.53	37.48	342.36		
YOL124W	ARG5	0.92	1.37	1.57	1.72	1.99	2.09	8.25	3.21	53.53	14.14	61.56	15.22	300.80		
YOL124W	PEX11	0.88	1.31	1.47	1.91	2.07	2.17	10.59	4.54	48.49	24.51	58.88	29.46	317.41		
YOL145C	CTR9	0.93	1.14	1.18	1.43	1.69	1.93	41.88	26.70	100.90	68.63	142.88	92.33	326.93		
YOL147C	PEK11	0.92	1.14	1.36	1.76	1.98	2.14	4.94	5.08	41.20	40.89	46.14	45.69	269.47		
YOL148C	SPT20	0.87	1.08	1.05	0.98	1.10	1.21	6.31	46.33	108.78	93.52	171.95	139.85	409.79		
YOL150C	YOL150c	0.93	1.33	1.47	1.90	2.09	2.20	7.17	3.16	40.52	14.97	47.69	18.13	306.41		
YOL151W	GRE2	0.97	1.36	1.54	1.96	2.10	2.23	7.76	4.24	43.28	24.75	51.04	28.98	317.03		
YOL152W	GRE2	0.89	1.22	1.41	1.84	2.06	2.25	5.51	6.11	49.63	66.96	85.07	83.44	335.38		
YOL152W	FRET	0.88	1.05	1.00	1.25	1.51	1.76	35.29	33.75	82.43	82.74	117.72	116.49	343.01		
YOL152W	FRET	0.95	1.17	1.39	1.85	2.										

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YOR044W	IRC23	0.98	1.37	1.60	2.02	2.19	2.32	0.12	2.13	8.68	2.75	8.81	4.89	284.47		
YOR045W	TOM6	0.98	1.37	1.56	1.94	2.14	2.30	-0.01	-0.04	1.72	-0.56	1.71	-0.61	281.73		
YOR047C	STD1	1.00	1.38	1.61	1.99	2.17	2.31	0.33	1.62	12.47	4.06	12.81	5.68	274.39		
YOR049C	RSBT	0.96	1.28	1.53	1.91	2.08	2.25	4.15	2.25	43.04	10.75	47.19	13.00	305.33		
YOR050C	YOR050c	0.96	1.36	1.61	2.00	2.15	2.29	0.37	0.28	16.79	4.29	17.16	4.57	280.11		
YOR051C	YOR051c	0.99	1.29	1.55	2.00	2.15	2.31	4.44	1.96	39.65	13.35	44.09	15.32	297.26		
YOR052C	YOR052c	0.95	1.36	1.62	2.00	2.14	2.37	1.19	0.16	25.67	9.41	26.66	9.58	273.44		
YOR053W	YOR053w	0.97	1.26	1.50	1.92	2.07	2.22	7.34	3.63	48.97	38.37	56.40	42.80	309.68		
YOR054W	YOR054w	0.97	1.27	1.51	1.98	2.13	2.28	3.34	1.49	34.49	12.85	34.74	12.99	301.21		
YOR055W	YOR055w	0.99	1.32	1.55	1.88	2.03	2.20	2.72	1.01	33.45	10.82	36.17	10.99	300.77		
YOR056C	ASE1	0.94	1.28	1.55	1.97	2.13	2.28	2.36	0.49	33.66	16.82	38.02	17.31	292.97		
YOR059C	YOR059c	1.02	1.34	1.55	2.03	2.22	2.36	5.07	1.02	48.78	26.97	53.85	27.98	277.78		
YOR061W	CK2A	1.01	1.41	1.58	1.88	2.08	2.25	0.02	2.56	3.97	3.80	3.99	5.49	297.43		
YOR062C	YOR062c	0.99	1.33	1.51	1.91	2.12	2.28	0.15	-0.01	4.99	-0.15	5.14	-0.17	273.27		
YOR064C	YNG1	0.99	1.42	1.65	1.98	2.15	2.27	0.26	-0.01	15.73	7.66	15.98	7.65	278.04		
YOR065W	CYT1	0.88	1.18	1.17	0.97	0.96	0.96	129.75	122.89	129.94	125.04	259.69	247.93	524.36		
YOR066W	MSA1	1.04	1.48	1.72	2.08	2.25	2.36	0.16	-0.02	8.37	2.61	9.02	1.98	267.02		
YOR067C	ALG8	1.02	1.32	1.48	1.88	2.07	2.24	1.13	0.24	25.80	8.82	26.93	9.06	276.26		
YOR068C	VAM10	0.94	1.28	1.49	1.83	2.06	2.31	4.04	0.21	46.69	11.82	50.74	12.03	338.02		
YOR069W	VPSS	0.94	1.28	1.44	1.71	1.86	2.01	-0.02	-0.01	4.01	-0.47	3.99	-0.48	320.46		
YOR070W	VPSS	0.95	1.24	1.35	1.58	1.71	1.80	27.30	11.96	84.12	48.52	111.43	69.48	390.30		
YOR070C	GYP1	1.03	1.46	1.72	2.08	2.20	2.32	4.08	4.41	43.38	40.37	47.46	44.78	308.64		
YOR071C	NRT1	0.96	1.27	1.44	1.88	2.07	2.27	-0.16	0.07	1.82	-0.53	1.67	-0.46	269.46		
YOR072W	YOR072w	0.99	1.37	1.62	2.07	2.22	2.35	2.56	2.01	35.76	12.18	38.32	34.20	289.54		
YOR073W	SGO1	1.00	1.24	1.38	1.91	2.15	2.33	2.87	0.67	39.53	21.32	42.40	21.99	282.28		
YOR076C	SKT1	0.98	1.29	1.48	1.88	2.08	2.21	3.09	3.46	38.36	28.44	41.45	31.90	294.74		
YOR078W	BUD21	1.05	1.38	1.50	1.98	2.22	2.31	16.90	8.94	71.49	6.31	88.39	7.25	322.33		
YOR079C	ATX2	1.03	1.49	1.70	2.11	2.28	2.38	1.26	2.09	5.67	3.82	6.94	5.91	269.36		
YOR080W	DIA2	1.04	1.27	1.29	1.44	1.66	1.90	55.16	26.76	103.87	86.91	159.03	115.67	374.59		
YOR081C	TGL5	0.99	1.31	1.50	1.89	2.11	2.27	0.02	-0.21	9.44	2.5	9.47	2.04	279.24		
YOR082C	YOR082c	1.00	1.37	1.58	2.03	2.23	2.34	1.82	0.02	32.78	9.51	34.60	9.53	253.21		
YOR083W	WH5	0.98	1.31	1.48	1.92	2.13	2.30	0.12	-0.05	16.30	3.33	16.42	3.28	270.58		
YOR084W	LPL1	0.97	1.33	1.57	1.96	2.14	2.26	1.43	0.47	30.30	9.67	31.73	20.13	282.06		
YOR085W	OS73	1.00	1.33	1.53	1.98	2.18	2.33	2.14	0.31	29.76	14.01	31.90	14.32	288.02		
YOR086C	TCB1	1.04	1.47	1.70	2.11	2.26	2.34	1.02	0.58	27.76	19.69	28.77	20.48	295.34		
YOR087W	YCT1	1.10	1.49	1.71	2.15	2.35	2.46	0.54	2.1	52.75	22.72	60.46	25.02	301.36		
YOR088W	YOR088w	0.98	1.27	1.41	1.66	2.01	2.19	19.44	5.34	63.59	48.63	64.27	341.44			
YOR089C	YOR089c	1.06	1.37	1.54	1.89	2.02	2.21	10.73	0.29	78.46	46.05	80.18	46.95	328.36		
YOR090C	PTC5	1.06	1.34	1.59	1.88	2.04	2.24	2.00	11.79	1.05	60.02	24.70	71.88	22.75	331.33	
YOR091W	TM446	0.99	1.28	1.48	1.87	2.01	2.17	13.66	0.07	60.44	10.14	73.51	10.21	328.30		
YOR093W	EC44	1.03	1.28	1.59	2.01	2.16	2.29	14.37	0.21	61.30	6.71	76.58	6.91	337.13		
YOR093C	YOR093c	1.03	1.29	1.51	1.96	2.10	2.28	15.25	0.74	61.75	13.93	77.00	14.67	340.70		
YOR094W	ARP3	1.00	1.34	1.55	1.95	2.09	2.24	5.09	2.68	43.58	13.93	48.67	16.01	301.92		
YOR096W	RPS7a	0.96	1.38	1.51	1.83	1.90	2.20	11.98	4.66	43.64	5.48	48.24	6.27	310.14		
YOR097C	YOR097c	0.95	1.24	1.41	1.91	2.10	2.16	8.39	1.57	52.98	19.89	61.38	21.45	319.31		
YOR099W	KTR1	1.01	1.41	1.65	2.07	2.23	2.28	7.06	1.20	50.63	14.84	57.69	16.05	317.86		
YOR100C	CRC1	1.11	1.53	1.76	2.19	2.36	2.43	1.32	1.08	20.10	3.61	21.42	4.69	295.69		
YOR101W	RAS1	0.99	1.33	1.54	1.97	2.14	2.21	2.81	2.47	32.04	6.80	34.85	9.27	291.06		
YOR104W	PIN2	1.02	1.38	1.61	2.04	2.22	2.28	3.30	0.74	35.84	8.09	39.14	8.83	298.04		
YOR105W	YOR105w	1.00	1.36	1.58	2.01	2.19	2.25	1.98	1.02	26.31	5.68	28.29	6.70	292.02		
YOR106W	VAM3	1.04	1.32	1.47	1.83	2.00	2.06	46.73	26.31	108.31	88.89	154.83	115.20	419.47		
YOR107W	RGS2	1.04	1.41	1.63	2.06	2.23	2.32	4.60	0.80	48.44	5.48	48.24	6.27	310.14		
YOR108W	LEU9	1.06	1.34	1.57	2.08	2.25	2.33	17.05	1.78	67.29	24.63	84.34	26.40	348.50		
YOR109W	INP53	0.97	1.26	1.44	1.92	2.12	2.21	17.89	1.33	67.33	16.38	85.22	17.71	371.70		
YOR111W	YOR111w	1.03	1.41	1.64	2.06	2.21	2.28	9.46	1.06	54.79	11.34	64.25	12.40	328.91		
YOR112W	CEX1	1.00	1.34	1.54	2.01	2.18	2.25	7.55	1.38	51.83	15.83	60.21	17.22	323.44		
YOR113W	AZF1	0.97	1.24	1.41	1.89	2.10	2.21	11.90	1.55	62.00	15.76	73.90	17.31	360.66		
YOR117W	RGS1	0.97	1.25	1.44	1.94	2.13	2.33	102.56	92.21	133.00	32.44	206.41	42.42	332.42		
YOR119C	YOR119c	1.02	1.34	1.51	2.05	2.22	2.28	11.42	2.15	52.14	12.53	70.57	22.62	332.33		
YOR120C	YOR120c	1.01	1.32	1.43	1.86	2.05	2.25	3.25	3.40	36.25	8.64	39.65	8.69	307.33		
YOR131C	YOR131c	1.01	1.35	1.57	2.05	2.24	2.29	5.88	1.97	43.43	22.87	49.31	24.85	393.92		
YOR132W	VPS17	1.03	1.32	1.52	1.88	2.10	2.14	24.64	14.57	81.87	6.76	106.51	7.33	387.81		
YOR134W	BAG7	0.99	1.32	1.50	1.94	2.14	2.24	1.08	0.96	17.89	4.96	19.07	5.02	297.48		
YOR135C	YOR135c	1.11	1.52	1.73	2.14	2.30	2.39	5.51	1.36	50.42	20.37	55.94	21.73	327.20		
YOR136W	IDH2	1.00	1.25	1.44	1.95	2.15	2.22	16.62	5.48	66.05	42.05	82.67	47.53	343.79		
YOR137C	SIA1	1.08	1.38	1.63	2.09	2.27	2.33	19.79	10.79	71.76	52.46	91.55	63.25	342.79		
YOR138C	RUP1	0.95	0.93	0.99	1.06	1.12	1.14	41.33	18.31	98.55	76.80	139.88	95.11	341.71		
YOR139C	YOR139c	0.99	1.04	1.05	1.10	1.14	1.19	108.84	98.54	135.51	134.64	245.35	233.19	476.36		
YOR140W	SFL1	0.93	0.98	0.99	1.06	1.19	1.20	50.60	26.35	105.96	66.39	156.76</td				

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YOR209C	NPT1	0.85	1.21	1.32	1.62	1.84	1.98	54.83	49.98	102.01	100.76	156.84	150.74	292.67		
YOR211C	MGM1	1.00	1.16	1.18	1.08	1.24	1.64	39.70	22.35	97.48	81.00	137.18	103.35	259.85		
YOR212W	STE4	0.90	0.99	1.00	1.29	1.51	1.67	35.55	8.32	91.94	57.28	127.49	65.60	356.09		
YOR213C	SASS	0.95	1.07	1.13	1.67	1.94	2.09	33.83	12.68	98.52	66.15	132.35	78.84	356.41		
YOR214C	YOR214c	0.88	0.98	1.01	1.41	1.74	1.88	19.71	12.63	73.95	64.83	93.67	77.46	338.10		
YOR215C	AIM41	0.88	1.00	1.02	1.33	1.57	1.74	19.17	2.29	81.93	32.15	101.10	34.45	333.92		
YOR216C	RGD3	0.91	1.10	1.11	1.59	1.80	1.91	22.56	7.72	67.27	55.47	109.83	63.19	335.41		
YOR218C	STE13	0.83	1.05	1.06	1.37	1.55	1.86	21.96	8.53	75.52	53.54	97.48	64.09	332.48		
YOR219C	MCT1	0.83	1.25	1.44	1.78	2.00	2.15	1.80	1.29	78.26	63.71	92.74	53.24	314.46		
YOR220W	OOC2	0.95	1.24	1.48	1.95	2.12	2.18	4.07	0.98	41.68	7.46	45.75	8.44	303.93		
YOR223W	YOR223w	0.97	1.12	1.17	1.53	1.74	1.85	19.76	3.75	75.65	39.79	95.41	43.54	312.44		
YOR225W	YOR225w	0.96	1.06	1.09	1.42	1.71	1.89	44.85	16.33	98.37	72.72	142.32	89.05	363.38		
YOR226C	ISU2	0.90	0.97	0.99	1.17	1.27	1.33	24.65	10.06	85.12	61.76	109.77	71.82	337.90		
YOR227W	HER1	0.97	1.19	1.20	1.40	1.53	1.63	23.31	10.94	80.38	61.90	103.68	72.84	336.56		
YOR228C	YOR228c	0.90	1.02	1.02	1.11	1.34	1.69	13.07	9.38	64.86	59.52	77.93	68.89	319.42		
YOR229W	WTM2	0.92	1.05	1.08	1.42	1.61	1.72	18.84	4.94	78.38	38.42	97.22	43.38	333.67		
YOR230W	WTM1	0.95	1.06	1.07	1.35	1.56	1.68	25.14	4.45	88.19	10.20	113.33	44.65	347.44		
YOR231W	MKK1	0.94	1.03	1.04	1.27	1.46	1.54	23.48	3.27	80.82	35.26	104.30	38.52	342.96		
YOR233W	KIN4	1.00	1.12	1.17	1.59	1.84	1.97	15.08	2.95	74.70	35.26	89.78	38.20	310.53		
YOR234C	RPL33b	1.02	1.12	1.13	1.37	1.56	1.64	18.81	7.67	75.05	56.93	93.66	64.61	319.61		
YOR235W	YOR235w	1.00	1.10	1.10	1.32	1.54	1.65	20.34	1.26	79.86	24.02	100.20	25.29	328.88		
YOR237W	HES1	0.97	1.08	1.10	1.39	1.59	1.65	32.60	9.24	92.59	59.30	125.18	68.54	336.22		
YOR238W	YOR238w	0.99	1.09	1.11	1.27	1.40	1.43	94.77	24.85	129.56	80.63	224.33	105.48	450.25		
YOR239W	ABP140	0.94	0.99	1.01	1.19	1.30	1.38	44.23	13.41	97.68	7.07	141.91	84.27	371.22		
YOR240W	ABP140	1.04	1.44	1.62	2.02	2.17	2.28	10.71	1.59	58.94	19.34	69.65	20.92	311.80		
YOR241W	MET17	0.90	1.20	1.24	1.44	1.69	1.85	45.53	24.14	101.74	87.47	147.27	111.61	363.28		
YOR242C	SSP2	0.99	1.30	1.26	1.51	1.71	1.81	33.67	11.76	94.32	70.25	128.00	82.01	343.74		
YOR243C	PUT3	0.93	1.03	1.04	1.26	1.43	1.53	14.38	5.38	68.34	52.44	82.72	57.81	315.48		
YOR245C	DGA1	1.02	1.17	1.24	1.68	1.91	2.04	19.69	5.47	73.88	45.37	93.57	50.84	332.61		
YOR246C	YOR246c	1.02	1.17	1.27	1.75	1.99	2.09	15.98	3.49	68.61	39.21	84.59	42.69	322.82		
YOR247W	SLR1	1.01	1.18	1.31	1.89	2.09	2.15	14.87	6.43	67.83	51.06	82.71	57.48	304.89		
YOR248W	YOR248w	0.95	1.06	1.08	1.38	1.58	1.62	26.67	14.85	84.67	77.10	111.34	91.95	340.79		
YOR251C	TUM1	1.07	1.21	1.25	1.56	1.78	1.93	32.66	7.33	89.65	52.54	122.32	59.87	337.20		
YOR252W	TUM16	0.95	1.07	1.07	1.31	1.52	1.63	30.09	5.63	88.65	48.14	116.74	53.77	343.75		
YOR253W	NAT5	0.97	1.07	1.07	1.24	1.45	1.56	34.09	9.24	92.43	72.43	127.11	72.73	352.41		
YOR254W	OSM1	0.97	1.07	1.07	1.37	1.57	1.68	30.03	9.03	84.34	61.03	102.33	44.86	334.36		
YOR255W	YOR255w	1.01	1.13	1.16	1.32	1.45	1.57	74.11	43.67	117.42	100.63	188.54	144.09	337.87		
YOR256W	PNT1	0.93	1.21	1.44	1.70	1.91	2.06	8.50	0.73	54.86	6.88	63.36	7.71	306.76		
YOR257C	HWK1	0.93	1.20	1.40	1.89	2.05	2.15	61.02	34.72	115.58	98.34	176.60	133.06	316.91		
YOR258C	YOR258c	0.92	1.13	1.30	1.79	1.98	2.13	9.11	1.72	58.88	25.74	68.08	27.46	265.28		
YOR259W	PAC1	0.89	1.04	1.18	1.63	1.83	2.00	14.41	5.84	68.02	47.93	82.43	53.77	314.80		
YOR270C	VPH1	0.93	1.08	1.22	1.73	1.94	2.04	5.39	2.01	52.90	34.71	58.28	36.72	293.44		
YOR271C	F5F1	0.88	1.06	1.26	1.82	2.01	2.15	6.23	10.96	50.06	56.28	56.29	67.25	302.19		
YOR273C	TP04	0.88	1.01	1.14	1.62	1.90	2.07	10.71	10.76	59.26	56.64	69.97	67.40	308.90		
YOR274W	MQD5	0.90	1.03	1.21	1.84	1.86	1.93	5.93	0.70	39.06	6.19	44.99	6.89	268.75		
YOR275C	RIM20	0.94	1.18	1.41	1.84	2.06	2.34	14.27	1.12	60.44	18.47	74.71	19.60	277.95		
YOR276W	CAF20	0.88	1.02	1.19	1.62	1.85	2.00	8.28	1.87	49.76	4.76	55.16	5.85	261.95		
YOR277C	YOR277c	0.89	1.04	1.09	1.56	1.81	1.90	25.89	7.74	84.76	55.93	110.74	63.67	332.08		
YOR279C	RFM1	1.04	1.20	1.28	1.75	2.01	2.15	31.06	8.64	90.66	54.72	121.72	63.36	332.72		
YOR280C	FSH3	0.97	1.08	1.12	1.47	1.67	1.79	27.78	13.13	86.62	68.05	114.40	81.18	342.91		
YOR283W	YOR283w	0.98	1.12	1.15	1.51	1.78	1.93	16.84	5.03	78.46	53.05	95.31	58.08	320.04		
YOR284W	HUA2	1.00	1.11	1.12	1.29	1.48	1.54	30.50	7.03	91.19	59.40	121.69	66.43	353.32		
YOR285W	RDL1	0.97	1.07	1.07	1.29	1.52	1.63	40.76	8.23	58.35	141.33	66.59	363.06			
YOR286W	RDL2	1.03	1.13	1.13	1.35	1.53	1.60	48.79	22.83	98.20	78.94	146.99	101.77	373.59		
YOR288C	MPD1	0.97	1.15	1.29	1.77	1.96	2.16	9.28	2.65	52.59	31.54	61.87	34.19	307.90		
YOR289W	YOR289w	0.98	1.25	1.42	1.82	1.99	2.16	14.28	0.37	58.72	16.76	73.00	17.13	328.97		
YOR290C	SNF2	0.99	1.17	1.31	1.64	1.76	1.92	44.17	30.95	95.70	89.98	139.87	120.93	439.08		
YOR291W	YPK3	0.96	1.22	1.42	1.83	1.99	2.21	0.50	-0.01	17.75	3.94	18.25	3.93	293.28		
YOR292C	YOR292c	1.03	1.24	1.35	1.61	1.80	1.93	1.99	2.21	2.25	2.25	2.25	2.25	342.23		
YOR293W	YOR293w	1.01	1.37	1.60	1.90	2.10	2.26	0.77	1.81	12.62	21.49	13.39	23.31	289.01		
YOR295W	RAX1	1.02	1.32	1.50	1.92	2.10	2.27	0.84	0.13	25.29	5.20	26.14	5.33	293.51		
YOR302W	YOR302w	0.87	1.07	1.27	1.72	1.95	2.13	0.99	0.70	18.23	5.99	19.22	6.69	276.68		
YOR302W	CPA1	0.97	1.27	1.53	1.98	2.09	2.23	2.52	0.01	34.51	5.03	37.03	5.05	289.54		
YOR303W	CPA1	1.01	1.35	1.60	2.01	2.14	2.27	3.04	0.13	32.82	2.04	35.56	10.37	286.89		
YOR304C	YOR304c-a	1.00	1.29	1.54	1.97	2.11	2.25	11.42	2.06	60.06	30.37	71.48	23.43	323.64		
YOR304W	ISW2	0.95	1.22	1.44	1.87	2.02	2.17	2.15	7.54	75.43	53.86	96.87	61.40	330.78		
YOR305W	RRG7	1.01	1.21	1.43	1.92	2.11	2.28	23.73	8.33	72.08	55.39	95.81	63.73	327.70		
YOR306C	MCH5	1.00	1.46	1.60	1.85	2.07	2.21	8.44	13.59	57.85	62.81	66.28	76.40	324.26		
YOR306C	MCH5	1.06														

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE			FRUCTOSE			TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YOR364W	YOR364w	0.93	1.18	1.38	1.79	1.97	2.08	1.01	0.76	17.01	4.17	18.02	4.93	281.93				
YOR364W	YOR364w	0.90	1.01	1.14	1.66	1.87	2.05	5.59	1.54	44.77	24.10	50.36	25.65	295.64				
YOR365C	YOR365c	1.04	1.37	1.60	2.00	2.13	2.26	0.40	0.19	14.62	8.38	15.02	8.57	272.02				
YOR366W	YOR366w	0.88	1.16	1.31	1.68	1.88	2.03	18.55	9.59	74.10	53.29	92.65	62.88	335.91				
YOR367W	SCP1	1.04	1.36	1.60	1.99	2.20	2.16	0.46	0.32	17.52	7.22	17.98	7.55	279.11				
YOR368W	RAD17	1.03	1.35	1.55	1.94	2.12	2.27	0.50	0.31	17.70	21.69	18.21	22.50	276.90				
YOR369C	RPS12	0.89	1.15	1.60	1.49	1.75	1.96	24.96	9.31	63.45	45.97	108.41	55.28	319.43				
YOR370W	GFB1	1.00	1.40	1.60	2.01	2.16	2.33	0.11	0.07	0.12	0.58	0.23	0.64	247.46				
YOR374W	ADP1	1.07	1.29	1.51	1.95	2.16	2.31	3.45	1.29	36.69	24.25	42.02	25.10	223.23				
YOR375C	CDF1	1.01	1.33	1.57	1.97	2.16	2.51	5.34	4.67	46.12	38.43	51.46	43.10	278.98				
YOR376W	YOR376w-a	1.00	1.35	1.58	1.98	2.13	2.27	2.15	1.45	33.48	26.27	35.63	27.72	295.29				
YOR376W	AIF1	0.86	1.15	1.26	1.85	1.87	1.99	7.44	2.19	54.74	24.32	62.19	26.51	288.37				
YOR377W	ATF1	0.99	1.20	1.43	1.98	2.13	2.20	7.83	5.84	52.09	44.79	59.93	50.73	308.71				
YOR378W	YOR378w-a	1.07	1.45	1.68	2.08	2.23	2.36	16.40	6.32	64.53	43.38	80.93	49.69	341.31				
YOR379C	YOR379c	0.92	1.23	1.42	1.79	1.98	2.12	17.29	10.56	68.51	56.27	85.71	66.83	348.34				
YOR380W	RDR1	1.00	1.41	1.65	2.04	2.18	2.30	17.83	1.33	73.27	29.35	91.10	31.28	343.43				
YOR381W	FBE3	1.04	1.41	1.63	2.01	2.15	2.29	1.81	1.32	31.32	29.86	33.23	31.88	285.62				
YOR382W	FIT2	1.05	1.44	1.66	2.02	2.14	2.28	0.42	0.39	16.32	24.70	16.74	25.69	271.43				
YOR383C	FIT3	1.01	1.39	1.59	1.98	2.14	2.26	0.46	0.37	17.40	13.25	17.86	13.62	268.36				
YOR384W	FRE5	0.97	1.29	1.50	1.90	2.05	2.20	0.61	0.00	20.59	7.50	21.20	7.50	272.95				
YOR385W	FRE5	1.02	1.37	1.59	1.96	2.09	2.23	0.21	0.16	12.99	11.95	13.20	12.12	277.47				
YOR386W	PHR1	0.99	1.34	1.58	1.99	2.12	2.27	0.08	0.05	4.41	5.06	4.49	5.11	271.53				
YPL001W	HAT1	0.91	1.13	1.24	1.67	1.84	2.04	13.07	5.34	55.27	32.43	68.35	37.77	319.47				
YPL002C	SNF8	0.89	1.14	1.24	1.49	1.58	1.76	20.95	10.66	72.15	59.50	93.11	70.16	369.01				
YPL003W	ULA1	0.92	1.24	1.35	1.69	1.81	2.02	9.95	4.78	48.74	28.14	58.69	32.92	314.19				
YPL004C	LSP1	0.98	1.24	1.47	1.91	2.09	2.22	0.92	1.50	17.15	27.57	18.07	29.07	265.39				
YPL005W	AEP3	0.83	0.97	0.97	1.34	1.54	1.76	7.75	6.82	44.28	36.16	52.03	42.99	244.54				
YPL006W	NCR1	0.93	1.33	1.46	1.79	1.95	2.12	3.21	3.47	4.18	3.91	7.38	7.39	253.80				
YPL008W	CHL1	0.89	1.13	1.17	1.63	1.55	1.78	7.52	8.67	44.68	44.68	52.20	53.35	288.95				
YPL009C	YPL009c	1.14	1.76	2.01	2.19	2.29	2.31	3.97	4.31	25.47	24.62	29.44	28.93	286.93				
YPL013C	MRP516	1.05	1.39	1.59	2.05	2.18	2.27	5.59	5.41	38.28	33.33	43.88	38.74	235.45				
YPL014W	YPL014w	0.94	1.26	1.36	1.70	1.88	2.11	3.21	3.33	6.98	3.72	10.18	7.05	254.96				
YPL015C	HST2	0.85	1.08	1.19	1.57	1.72	1.89	3.23	3.30	13.80	5.37	17.03	8.67	275.38				
YPL017C	IRC15	0.87	0.96	1.01	1.27	1.51	1.78	11.63	9.36	66.09	59.46	77.71	68.81	288.92				
YPL018W	CTF19	0.85	1.00	0.98	1.14	1.30	1.53	24.49	14.39	74.03	62.35	98.53	76.74	326.31				
YPL019C	VTC3	0.89	1.16	1.28	1.65	1.80	1.98	14.95	5.74	58.16	34.89	73.11	40.63	324.93				
YPL021W	ECM23	0.96	1.28	1.45	1.88	2.03	2.20	13.78	5.82	56.57	34.17	70.46	39.99	320.13				
YPL022W	RAD1	0.92	1.21	1.34	1.74	1.88	2.01	7.48	3.78	42.20	21.79	49.64	23.58	306.42				
YPL023W	YPL023w	0.85	1.31	1.48	1.87	2.02	2.26	4.36	3.29	32.89	13.75	35.82	25.28	225.28				
YPL024W	RMH1	0.83	1.11	1.22	1.65	1.84	2.00	15.37	9.13	74.14	45.79	89.51	44.91	311.14				
YPL025C	YPL025c	0.82	1.24	1.38	1.79	1.97	2.17	3.16	3.38	10.65	10.96	13.80	14.34	269.01				
YPL028C	SKS1	0.93	1.26	1.38	1.87	1.98	2.03	1.77	1.14	32.55	48.97	37.33	55.19	278.32				
YPL029W	SM1	0.94	1.19	1.42	1.84	2.02	2.17	4.08	6.04	39.13	41.51	43.21	47.55	289.37				
YPL029W	SUV3	0.90	1.28	1.31	1.49	1.85	2.01	25.64	14.68	84.82	68.35	110.47	83.04	267.61				
YPL030W	TRM44	0.92	1.25	1.45	1.91	2.16	2.27	4.37	5.41	26.86	31.75	31.23	37.16	275.65				
YPL031C	PHOB5	0.93	1.07	1.18	1.59	1.78	1.72	32.53	18.82	101.54	64.93	134.06	83.75	416.11				
YPL032C	SVL3	0.87	1.12	1.21	1.54	1.70	1.88	3.44	4.37	16.11	24.76	19.55	29.13	262.45				
YPL033C	SR14	0.86	1.11	1.24	1.66	1.86	2.05	3.09	3.20	6.41	2.98	9.50	6.18	250.96				
YPL034W	YPL034w	0.91	1.12	1.32	1.72	1.90	2.04	5.83	6.83	46.21	45.07	52.04	51.90	286.88				
YPL035C	YPL035c	0.84	1.07	1.17	1.56	1.71	1.89	3.45	3.59	18.03	6.23	21.48	9.73	263.91				
YPL036W	PMA2	0.84	0.94	1.03	1.38	1.58	1.70	6.01	3.49	46.58	36.11	52.59	39.60	289.92				
YPL037C	EGD1	0.91	1.15	1.24	1.58	1.74	1.91	10.39	6.11	46.03	33.09	56.42	39.20	316.82				
YPL038W	MET31	0.91	1.12	1.22	1.63	1.80	1.99	11.57	7.63	51.78	41.15	63.35	48.77	320.34				
YPL038W	YPL038w-a	0.92	1.17	1.34	1.75	2.01	2.12	1.02	0.00	3.01	4.37	6.74	23.77	9.75	242.18			
YPL039W	YPL039w	0.94	1.31	1.48	1.87	2.02	2.19	12.44	6.17	51.10	36.56	63.54	42.73	311.14				
YPL040C	ISM1	0.94	1.30	1.46	1.86	2.00	2.17	5.22	6.83	47.33	38.27	52.55	45.10	301.09				
YPL041C	YPL041c	0.86	1.09	1.17	1.54	1.79	1.99	4.98	3.03	11.26	9.33	49.31	36.90	60.57	267.94			
YPL042C	SSN3	0.92	1.07	1.04	1.15	1.26	1.46	26.83	25.06	70.18	76.25	97.02	101.32	309.87				
YPL045W	KTR6	0.87	1.08	1.17	1.53	1.68	1.84	2.14	10.62	11.28	51.21	48.60	61.83	59.89	307.51			
YPL055C	LG1	0.92	1.11	1.18	1.62	1.79	2.01	7.48	8.62	46.95	46.19	54.03	54.52	273.13				
YPL056C	YPL056c	0.90	1.19	1.33	1.70	1.87	2.06	4.48	4.65	18.19	21.59	40.98	36.05	407.17				
YPL057C	SUR1	0.88	1.16	1.31	1.66	1.81	1.98	4.05	4.23	28.02	41.37	32.06	49.60	266.65				
YPL058C	PDR12	0.87	1.14	1.25	1.63	1.82	2.01	3.07	3.98	6.85	22.68	11.72	26.66	255.68				
YPL059W	GRX5	0.96	1.05	1.07	1.09	1.00	1.02	1.01	1.01	12.95	11.84	11.99	11.99	237.78				
YPL060W	LEP10	0.87	1.20	1.38	1.71	1.81	1.93	3.28	3.36	16.41	21.41	24.77	24.77	25				

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE			FRUCTOSE			TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YPL127C	HH01	0.99	1.24	1.40	1.90	2.07	2.25	9.32	1.92	56.47	32.58	65.79	34.50	330.16	330.16	330.16	330.16	
YPL129W	TAF14	0.96	1.06	1.03	1.06	1.09	1.18	110.25	95.44	129.45	120.48	239.70	215.92	458.89	458.89	458.89	458.89	
YPL130W	SPO19	0.89	1.06	1.15	1.63	1.83	2.05	0.33	0.07	18.69	3.22	19.02	3.30	313.89	313.89	313.89	313.89	
YPL132W	COK11	0.87	1.16	1.21	1.39	1.69	1.90	14.12	5.94	61.35	41.51	75.47	47.46	277.08	277.08	277.08	277.08	
YPL133C	RDS2	0.90	1.14	1.32	1.87	2.02	2.18	3.96	0.94	41.96	22.89	45.93	23.83	297.66	297.66	297.66	297.66	
YPL134C	QDC1	0.99	1.50	1.71	2.04	2.17	2.29	7.62	3.72	37.63	17.12	45.24	20.83	295.89	295.89	295.89	295.89	
YPL135W	ISU	1.00	1.40	1.58	2.08	2.28	2.40	0.12	0.19	4.41	-0.09	4.53	0.10	274.43	274.43	274.43	274.43	
YPL136W	YPL19w	0.88	1.27	1.45	1.91	2.07	2.24	2.33	0.20	34.97	9.83	37.28	10.13	306.62	306.62	306.62	306.62	
YPL138G	SPH1	0.88	1.33	1.44	1.52	1.89	2.07	6.46	0.59	44.97	12.04	51.65	12.52	275.25	275.25	275.25	275.25	
YPL139C	SPP1	0.88	1.30	1.53	1.98	2.10	2.25	4.71	0.71	43.59	19.99	48.30	19.80	255.36	255.36	255.36	255.36	
YPL139C	UME1	1.01	1.40	1.61	2.00	2.14	2.28	5.56	0.91	52.91	31.35	58.47	32.28	293.10	293.10	293.10	293.10	
YPL140C	MKK2	0.95	1.33	1.51	2.02	2.17	2.36	0.35	0.05	9.28	-0.62	8.57	-0.57	277.43	277.43	277.43	277.43	
YPL141C	FBB1	1.01	1.41	1.57	1.98	2.09	2.27	0.05	0.07	0.14	-0.97	0.18	-0.91	292.14	292.14	292.14	292.14	
YPL144W	POC4	1.13	1.52	1.73	2.14	2.24	2.34	24.43	18.51	73.66	65.81	98.09	84.33	335.71	335.71	335.71	335.71	
YPL145C	KES1	1.00	1.37	1.56	2.03	2.17	2.29	2.44	0.09	18.18	3.88	20.62	3.96	299.62	299.62	299.62	299.62	
YPL147W	PWA1	1.02	1.43	1.65	2.13	2.27	2.41	1.30	1.20	14.83	1.41	16.13	0.61	288.92	288.92	288.92	288.92	
YPL148C	PBT2	0.94	1.16	1.39	1.88	2.10	2.27	1.18	0.66	9.38	1.08	10.57	1.25	268.10	268.10	268.10	268.10	
YPL149W	ATG5	1.08	1.26	1.34	1.79	2.00	2.19	14.66	3.47	71.81	39.10	86.27	42.57	340.77	340.77	340.77	340.77	
YPL150W	YPL150w	0.89	1.10	1.25	1.75	1.92	2.12	4.08	0.10	39.42	9.58	43.51	9.68	297.89	297.89	297.89	297.89	
YPL152W	RRD2	0.94	1.22	1.37	1.85	2.03	2.23	0.05	0.09	1.73	-0.03	1.79	0.05	282.53	282.53	282.53	282.53	
YPL152W-A	YPL152w-a	0.95	1.25	1.39	1.78	2.01	2.14	5.71	1.20	48.19	17.42	53.90	18.62	283.76	283.76	283.76	283.76	
YPL154C	PEP4	0.98	1.14	1.25	1.81	2.04	2.23	6.94	0.75	58.47	26.86	65.40	27.61	298.61	298.61	298.61	298.61	
YPL155C	KIP2	0.97	1.24	1.42	1.97	2.15	2.34	1.23	0.16	27.62	6.84	28.85	7.01	291.69	291.69	291.69	291.69	
YPL156C	PRM4	0.95	1.18	1.32	1.84	2.01	2.21	16.39	6.83	69.69	46.98	86.08	53.81	324.92	324.92	324.92	324.92	
YPL157W	TGS1	0.95	1.33	1.55	2.00	2.10	2.25	18.82	9.14	74.83	55.47	93.66	64.61	309.56	309.56	309.56	309.56	
YPL158C	AIM44	0.99	1.15	1.10	1.19	1.36	1.46	26.64	36.73	78.14	82.74	104.78	119.47	324.61	324.61	324.61	324.61	
YPL159C	PET20	1.01	1.35	1.53	2.00	2.15	2.32	4.49	0.10	40.55	8.93	45.04	9.07	300.18	300.18	300.18	300.18	
YPL161C	BEM4	1.03	1.43	1.67	2.11	2.22	2.35	7.16	2.41	48.37	32.85	55.53	35.27	299.16	299.16	299.16	299.16	
YPL162C	YPL162c	1.09	1.52	1.72	2.22	2.36	2.49	0.47	0.16	18.59	2.84	19.05	3.01	301.67	301.67	301.67	301.67	
YPL163C	SV1	1.02	1.42	1.62	2.09	2.23	2.36	0.74	0.02	16.40	3.79	17.14	3.80	299.25	299.25	299.25	299.25	
YPL164C	MHL3	0.98	1.38	1.57	2.08	2.22	2.37	0.12	-0.01	7.79	0.29	7.90	0.28	285.04	285.04	285.04	285.04	
YPL165C	SET6	0.91	1.19	1.31	1.65	1.83	1.98	15.87	3.35	68.60	29.37	84.47	32.71	300.38	300.38	300.38	300.38	
YPL165C	SET6	0.84	1.13	1.26	1.69	1.86	1.98	18.80	8.08	70.67	47.94	89.57	56.02	308.41	308.41	308.41	308.41	
YPL166W	ATG29	0.90	1.08	1.19	1.69	1.87	2.10	6.05	2.29	53.17	32.05	59.23	33.34	299.22	299.22	299.22	299.22	
YPL167C	REV3	0.91	1.14	1.27	1.78	1.98	2.20	0.27	0.13	2.05	-0.17	2.32	-0.03	271.03	271.03	271.03	271.03	
YPL168W	YPL168w	1.04	1.36	1.54	2.01	2.19	2.34	11.66	3.64	56.67	34.30	68.34	37.94	320.21	320.21	320.21	320.21	
YPL169W	DAP1	0.99	1.32	1.52	2.02	2.14	2.30	2.10	0.26	32.84	17.17	34.93	17.72	296.26	296.26	296.26	296.26	
YPL170W	YPL170c	1.01	1.30	1.51	2.04	2.17	2.32	2.03	0.23	19.19	21.76	22.46	2.48	293.10	293.10	293.10	293.10	
YPL172C	COK20	0.85	1.12	1.30	0.92	0.92	0.92	122.69	123.00	141.63	123.90	246.53	246.80	332.75	332.75	332.75	332.75	
YPL173W	MRP140	0.93	1.23	1.44	1.01	1.02	1.14	123.59	124.26	132.82	124.59	252.33	252.33	322.40	322.40	322.40	322.40	
YPL174C	NIP100	1.03	1.34	1.47	1.82	1.97	2.12	16.19	10.88	67.59	8.78	83.78	6.94	292.60	292.60	292.60	292.60	
YPL178C	TRF1	1.03	1.43	1.63	2.08	2.21	2.35	11.43	3.31	57.05	9.84	68.48	4.49	334.47	334.47	334.47	334.47	
YPL177C	CUP9	0.86	1.33	1.52	1.97	2.10	2.25	5.83	0.51	40.70	18.62	46.53	19.13	303.59	303.59	303.59	303.59	
YPL178W	CB2C	1.01	1.33	1.08	1.29	1.61	1.92	45.14	26.88	93.87	74.79	139.02	101.77	300.62	300.62	300.62	300.62	
YPL179W	PP01	1.04	1.50	1.70	2.17	2.29	2.42	-0.28	-0.30	3.86	3.75	-0.03	-0.03	278.37	278.37	278.37	278.37	
YPL180W	TCO99	0.91	1.15	1.29	1.73	1.88	2.05	28.57	24.99	77.55	72.43	106.12	97.42	420.88	420.88	420.88	420.88	
YPL181W	CT16	1.04	1.33	1.46	1.98	2.16	2.36	0.24	0.17	17.41	8.18	17.66	8.35	296.34	296.34	296.34	296.34	
YPL182C	YPL182c	0.87	1.11	1.23	1.68	1.82	2.05	0.24	0.16	11.48	2.27	11.72	2.43	315.88	315.88	315.88	315.88	
YPL183C	RTT10	0.92	1.09	1.26	1.78	1.98	2.14	14.60	2.15	50.56	20.62	56.79	22.39	278.22	278.22	278.22	278.22	
YPL184W	POSS	0.91	1.26	1.44	1.97	2.12	2.28	4.77	1.26	49.22	14.92	51.80	16.66	358.06	358.06	358.06	358.06	
YPL185W	YR1	0.94	1.25	1.44	1.86	2.04	2.20	0.22	-0.02	0.07	0.13	11.59	0.71	280.18	280.18	280.18	280.18	
YPL186W	YPL186w	0.94	1.22	1.39	1.86	2.02	2.21	5.30	0.02	46.78	6.30	52.08	6.31	297.36	297.36	297.36	297.36	
YPL187W	YPL187w	0.93	1.17	1.37	1.86	2.02	2.19	2.88	0.02	45.00	4.00	49.06	4.00	321.46	321.46	321.46	321.46	
YPL188W	YPL188w	0.94	1.20	1.36	1.82	2.00	2.21	-0.18	-0.12	0.03	-0.05	-0.05	-0.05	260.85	260.85	260.85	260.85	
YPL189W	SSO1	0.82	1.18	1.43	1.88	2.02	2.20	0.00	-0.15	5.46	-0.50	5.46	-0.50	260.45	260.45	260.45	260.45	
YPL190C	YPL190c	1.01	1.19	1.20	1.38	1.59	1.83	38.57	17.19	94.03	80.22	132.59	97.41	332.15	332.15	332.15	332.15	
YPL194C	TFP1	0.94	1.14	1.17	1.30	1.63	1.89	37.94	26.98	91.05	90.61	128.99	117.60	322.18	322.18	322.18	322.18	
YPL195C	YPL195c	0.94	1.20	1.39	1.86	2.02	2.19	1.08	-0.07	27.79								

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YPR005C	HAL1	0.89	1.16	1.27	1.58	1.70	1.88	4.18	3.79	26.97	19.53	31.15	23.32	278.99		
YPR006C	ICL2	0.89	1.30	1.48	1.79	1.83	2.10	16.57	8.86	57.95	43.19	74.52	52.05	345.32		
YPR007C	REC8	0.96	1.23	1.48	1.93	2.11	2.25	13.87	3.68	61.49	32.46	75.35	35.54	311.83		
YPR008W	HAAT	0.96	1.29	1.57	1.98	2.14	2.24	14.03	5.89	71.41	53.42	85.44	59.31	309.86		
YPR009W	SU12	0.97	1.38	1.52	1.84	1.98	2.10	15.22	7.66	56.26	38.88	71.48	46.53	340.51		
YPR011C	YPR011c	1.04	1.26	1.49	1.97	2.13	2.23	12.63	2.43	53.51	21.96	65.54	24.39	321.80		
YPR012A	YPR012w	1.05	1.25	1.47	1.97	2.08	2.19	14.70	8.92	54.29	45.45	68.99	54.57	330.37		
YPR013C	YPR013c	0.92	1.20	1.43	1.85	2.04	2.17	6.48	1.73	48.86	27.31	55.32	29.89	315.20		
YPR014C	YPR014c	1.08	1.28	1.49	1.98	2.08	2.23	3.44	1.71	49.18	22.22	52.22	22.00	321.00		
YPR015C	YPR015c	1.08	1.24	1.49	1.85	2.21	2.31	2.41	6.01	7.79	48.19	41.92	510.21			
YPR017C	DSS4	1.03	1.53	1.73	2.07	2.18	2.28	15.82	7.48	55.79	40.23	71.61	47.71	331.91		
YPR018W	REF2	1.00	1.49	1.69	2.03	2.17	2.27	14.99	8.59	54.42	40.60	69.41	49.20	330.37		
YPR020W	ATP20	1.02	1.48	1.67	2.03	2.17	2.28	9.54	4.50	43.43	24.33	52.98	28.83	318.82		
YPR021C	AGC1	1.03	1.38	1.61	2.02	2.17	2.33	0.35	0.16	16.40	8.59	16.74	8.75	284.97		
YPR022C	YPR022c	0.85	0.99	1.16	1.58	1.78	1.95	3.62	1.05	37.17	16.23	40.79	17.28	295.05		
YPR023C	EAF3	0.93	1.10	1.25	1.73	1.98	1.95	6.17	1.10	47.14	18.91	53.31	20.02	318.80		
YPR024W	YME1	0.93	1.09	1.27	1.69	1.87	2.03	4.08	0.79	42.75	9.90	46.84	10.69	336.34		
YPR026W	ATH1	0.99	1.22	1.46	1.88	2.08	2.23	2.14	0.79	30.55	7.17	32.69	7.97	282.64		
YPR027C	YPR027c	1.03	1.49	1.71	2.06	2.19	2.28	15.40	8.02	56.64	41.70	72.04	49.72	335.76		
YPR028W	YOP1	1.00	1.54	1.77	2.10	2.21	2.30	13.55	6.45	52.75	35.94	66.30	42.39	335.45		
YPR029C	APL4	1.00	1.48	1.69	2.08	2.23	2.32	15.00	6.47	58.54	34.93	73.54	41.40	333.85		
YPR030W	CSR2	0.93	1.29	1.51	1.89	2.07	2.18	12.61	5.47	51.54	32.54	64.15	38.01	338.48		
YPR031W	NT01	0.93	1.13	1.38	1.81	2.01	2.17	5.32	0.91	42.15	10.63	47.47	11.54	311.43		
YPR032W	SRO7	0.92	1.35	1.61	1.92	2.06	2.16	8.07	2.02	38.25	23.22	46.31	27.44	309.63		
YPR036W	VMA13	0.94	1.24	1.38	1.75	1.99	2.15	5.71	4.43	35.79	17.70	41.50	22.13	321.16		
YPR037C	ERV2	0.98	1.27	1.56	2.01	2.16	2.29	4.48	0.90	41.26	11.17	45.74	12.07	301.78		
YPR038W	YPR038w	0.93	1.31	1.48	1.82	2.00	2.14	13.64	5.52	52.70	32.92	66.34	38.44	349.05		
YPR039W	YPR039w	0.91	1.38	1.58	1.95	2.11	2.24	5.38	3.36	29.52	16.29	34.90	19.64	287.35		
YPR040W	TII41	0.99	1.40	1.61	1.95	2.11	2.25	4.11	2.86	24.31	7.51	28.41	10.37	293.85		
YPR042C	PUF2	0.88	1.31	1.49	1.85	2.02	2.17	3.05	2.88	18.67	5.39	21.72	8.26	304.58		
YPR043W	RPL43a	0.90	1.02	1.12	1.50	1.78	2.02	0.75	0.57	6.29	2.58	7.03	3.14	236.53		
YPR044C	YPR044c	0.91	1.21	1.25	1.61	1.89	2.13	4.36	2.67	33.01	8.70	37.37	11.58	274.36		
YPR045C	YPR045c	0.91	1.22	1.27	1.63	1.91	2.14	4.47	3.36	33.50	12.30	37.98	15.66	271.23		
YPR046W	MCM16	0.90	1.16	1.21	1.50	1.72	1.95	15.88	15.08	69.07	28.52	97.90	72.80	363.85		
YPR047W	MSF1	1.01	1.24	1.20	1.13	1.14	1.08	10.12	100.00	105.88	106.54	208.00	201.10	529.10		
YPR049C	ATG11	0.98	1.31	1.35	1.73	1.97	2.15	35.10	26.70	79.64	72.03	114.75	98.72	335.19		
YPR050W	YPR050c	0.90	1.10	1.32	1.86	2.00	2.14	2.09	2.74	27.45	18.23	29.54	19.47	267.85		
YPR051W	MAK3	0.93	1.36	1.56	1.95	2.05	2.13	13.65	8.03	53.02	41.51	68.67	49.52	328.46		
YPR052C	YPR052c	0.92	1.26	1.42	1.82	2.01	2.17	19.98	11.88	57.85	47.16	77.03	58.92	344.44		
YPR053C	YPR053c	0.95	1.36	1.56	1.96	2.02	2.12	13.16	6.24	51.77	34.36	65.55	40.31	335.37		
YPR054W	SMK1	0.98	1.34	1.51	1.93	2.12	2.28	26.20	13.27	67.85	52.40	83.05	65.67	355.29		
YPR055W	BRR1	0.98	1.26	1.41	1.75	1.95	2.11	5.44	2.94	31.54	14.42	38.97	17.82	284.18		
YPR056W	YMC1	0.97	1.39	1.58	1.97	2.13	2.26	15.59	8.00	51.87	38.38	67.46	47.38	341.93		
YPR058C	YMC1	0.98	1.39	1.58	1.97	2.13	2.26	15.59	8.00	51.87	38.38	67.46	47.38	343.08		
YPR059C	YPR059c	0.90	1.34	1.49	1.84	2.00	2.12	14.60	12.66	53.80	49.83	68.50	62.59	302.72		
YPR060C	ARO7	0.90	1.19	1.16	1.03	1.03	1.03	12.45	1.75	50.00	10.00	52.27	12.00	527.91		
YPR061C	JID1	0.91	1.31	1.45	1.73	1.94	2.07	9.71	4.37	45.32	24.90	55.03	29.27	337.76		
YPR062W	FCY1	0.89	1.35	1.52	1.87	2.02	2.14	17.83	8.24	57.47	39.02	75.30	47.26	350.48		
YPR063C	YPR063c	0.97	1.44	1.67	2.10	2.16	2.27	18.39	7.90	59.55	37.91	79.44	45.81	355.67		
YPR064W	YPR064w	0.92	1.08	1.26	1.70	1.94	2.10	2.41	1.71	33.25	26.33	35.66	28.04	293.44		
YPR065W	ROX1	0.96	1.35	1.47	1.87	2.04	2.17	17.81	9.79	57.07	44.55	74.67	54.34	343.98		
YPR066W	UBA3	1.00	1.37	1.63	1.98	2.13	2.23	20.39	12.36	62.21	49.26	82.61	61.62	363.67		
YPR067W	IS2A	0.94	1.17	1.19	1.37	1.72	1.94	25.26	12.15	78.76	63.29	104.02	75.44	314.96		
YPR068W	IS2A	0.85	1.05	1.04	1.15	1.48	1.68	22.85	10.41	72.35	49.53	95.20	59.94	302.31		
YPR068C	HOS1	0.99	1.40	1.59	1.90	2.06	2.17	18.39	11.17	58.00	47.91	76.39	59.07	342.39		
YPR069C	SP3	1.00	1.43	1.67	2.03	2.17	2.27	15.38	8.04	55.55	41.46	70.94	49.50	335.59		
YPR070W	MED1	1.04	1.25	1.29	1.43	1.70	1.92	45.35	36.01	88.11	78.31	133.46	114.31	388.60		
YPR071W	YPR071w	0.95	1.35	1.57	1.92	2.07	2.19	18.38	1.61	56.35	42.95	74.72	52.16	353.50		
YPR072C	NOT5	0.90	1.08	1.19	1.56	1.78	1.96	2.06	26.77	19.18	67.01	59.83	93.78	79.01	338.10	
YPR073C	LTP1	0.93	1.27	1.42	1.77	1.94	2.09	16.89	7.47	53.65	38.07	72.54	45.24	342.24		
YPR074C	TKL1	0.94	1.10	1.11	1.09	1.14	1.65	17.00	2.17	52.00	3.05	53.26	1.61	351.21		
YPR075C	OPV2	0.92	1.32	1.47	1.78	1.94	2.08	19.37	9.40	54.95	43.91	74.32	53.31	358.25		
YPR076W	YPR076w	0.93	1.40	1.59	1.98	2.12	2.24	14.15	6.09	52.84	33.07	66.99	39.17	333.54		
YPR077C	YPR077c	0.93	1.32	1.50	1.88	2.04	2.16	22.05	13.70	62.91	50.14	84.95	63.84	361.81		
YPR078C	YPR078c	0.93	1.13	1.33	1.77	2.00	2.16	8.25	1.48	52.26	21.42	60.52	22.91	303.60		
YPR079W	MRL1	0.98	1.34	1.46	1.91	2.10	2.22	17.86	4.90	50.50	31.54	74.36	36.44	281.08		
YPR080W	YPR080w	0.92	1.21	1.35	1.71	1.96	2.05	3.03	3.19	5.97	2.57	9.00	5.76	279.68		
YPR081W	YPR081w	0.91	1.26	1.43	1.82	1.97	2.15	6.31	3.38	38.02	14.92	45.32	18.70	285.23		
YPR082W	SCD6	1.01	1.46	1.69												

ORF.name	Strain	Optical density at 600 nm at regular intervals (h)						GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	102 h
YPR166C	MRP2	0.90	1.22	1.33	1.80	1.96	2.11	15.91	7.25	64.74	44.70	80.65	51.95	335.42	
YPR167C	MET16	0.93	1.31	1.40	1.85	2.03	2.19	2.97	3.32	13.09	3.70	16.06	7.01	281.34	
YPR170C	YPR170c	0.87	1.20	1.21	1.63	1.83	2.00	7.27	3.50	45.42	15.05	52.69	18.54	303.36	
YPR171W	BSP1	0.88	1.23	1.30	1.75	1.93	2.08	7.63	3.40	46.70	14.61	54.32	18.00	304.48	
YPR172W	YPR172w	0.88	1.21	1.28	1.73	1.94	2.11	3.07	3.40	18.05	5.49	21.12	8.68	288.03	
YPR173c	VPS4	0.95	1.34	1.42	1.76	1.86	1.95	25.70	18.28	77.56	74.98	103.26	93.26	383.40	
YPR174C	YPR174c	0.89	1.33	1.38	1.71	1.89	2.13	7.50	3.47	46.76	20.06	54.26	23.53	310.40	
YPR179C	HDA3	0.88	1.27	1.29	1.82	2.02	2.17	5.76	3.33	37.19	16.89	42.94	20.22	292.18	
YPR184W	CDC42	0.98	1.29	1.30	1.91	1.98	2.12	12.46	3.35	48.71	19.25	51.61	19.25	333.28	
YPR185W	ATG13	0.94	1.26	1.27	1.77	1.99	2.18	5.31	3.04	34.93	8.49	40.23	12.52	398.25	
YPR188C	MLC2	0.98	1.37	1.38	1.93	2.10	2.23	12.24	9.41	55.15	46.79	67.39	56.19	330.49	
YPR189W	SKO3	0.97	1.24	1.30	1.71	1.90	2.08	15.82	5.85	58.08	37.40	73.90	43.25	323.44	
YPR191W	OCR2	0.93	1.20	1.30	1.66	1.83	2.02	11.46	4.78	55.17	33.11	66.64	37.89	315.58	
YPR192W	A0Y1	0.97	1.25	1.36	1.79	1.98	2.18	6.89	3.25	41.71	10.91	48.61	14.16	309.43	
YPR193C	HPA2	0.91	1.15	1.21	1.57	1.75	1.96	4.92	3.17	33.57	6.32	38.49	9.49	303.36	
YPR194C	OPT2	0.93	1.24	1.31	1.57	1.69	1.84	25.33	11.93	74.95	66.11	100.28	78.04	376.41	
YPR195C	YPR195c	1.04	1.36	1.50	1.96	2.12	2.29	6.00	3.51	37.98	15.74	43.98	19.25	311.05	
YPR196W	YPR196w	0.90	1.15	1.22	1.61	1.80	2.02	8.67	3.62	47.05	19.59	55.72	23.21	317.87	
YPR197C	YPR197c	1.05	1.52	1.73	2.13	2.25	2.36	3.47	3.19	14.61	5.59	18.08	8.78	277.44	
YPR198W	SGE1	0.93	1.24	1.36	1.84	2.01	2.20	4.22	3.28	27.45	9.21	31.66	12.49	294.32	
YPR199C	ARR1	0.95	1.22	1.33	1.82	2.01	2.21	8.05	3.30	45.90	11.96	53.95	15.26	307.28	
YPR200C	ARR2	0.92	1.19	1.27	1.77	1.97	2.17	11.73	6.91	52.20	37.49	63.93	44.39	323.28	
YPR201W	ARR3	0.95	1.30	1.37	1.91	2.09	2.21	20.38	13.24	66.15	57.99	86.53	71.23	348.02	

Appendix 2. Micro-fermentation data of WYDL_HN (all candidates).

ORF	Optical density at 600 nm at time intervals (h)							GLUCOSE	FRUCTOSE	TOTAL Sugars(g/L)	TOTAL AMINO ACIDS (mg/L)
	0	20	27	42	67	43 h	67 h				
YAL002W	0.95	1.70	2.00	2.28	2.36	20.46	4.99	71.52	32.33	91.98	37.31
YAL007C	0.92	1.52	1.73	1.97	2.03	30.19	4.89	78.00	30.22	108.20	35.11
YAL012W	0.88	1.39	1.65	2.05	2.23	38.63	6.81	107.69	41.10	146.32	47.91
YAL013W	0.89	1.60	1.88	2.22	2.34	29.74	4.16	91.73	28.84	121.47	33.00
YAL014C	0.86	1.51	1.80	2.17	2.26	28.41	3.11	85.17	27.79	113.58	29.90
YAL015C	0.88	1.61	1.89	2.23	2.31	16.93	1.76	87.42	5.65	104.35	7.41
YAL030W	0.98	1.76	2.00	2.21	2.29	31.39	2.22	106.81	21.09	138.19	23.31
YAL031C	0.98	1.79	2.09	2.35	2.40	23.58	2.49	85.61	22.76	109.20	25.25
YAL036C	0.95	1.88	2.16	2.39	2.43	21.65	2.64	81.50	21.65	106.18	24.25
YAL042C	0.92	1.55	1.81	2.07	2.08	41.00	5.73	98.32	32.42	129.49	36.41
YAL044C	0.95	1.68	1.95	2.26	2.32	23.44	3.53	76.45	24.89	99.85	28.42
YAL044W	0.95	1.90	2.00	2.40	2.48	10.84	1.64	51.44	1.56	62.07	1.59
YAL046C	1.00	1.37	1.50	2.02	2.04	9.11	1.87	147.66	77.64	238.78	95.11
YAL051W	0.94	1.54	1.74	2.13	2.38	41.86	4.54	107.01	38.91	148.87	43.46
YAL055W	0.89	1.46	1.72	2.09	2.29	53.94	6.37	124.72	49.05	178.66	55.42
YAL059W	1.01	1.68	2.00	2.21	2.35	14.04	2.34	78.69	30.42	92.73	32.76
YAL062W	0.91	1.45	1.73	2.05	2.18	53.72	5.61	121.74	37.56	175.46	43.17
YAR000C	0.92	1.45	1.70	2.07	2.33	32.02	4.12	93.74	45.92	125.76	51.04
YAR015W	0.96	1.81	2.05	2.22	2.34	24.12	2.35	80.40	17.29	104.52	19.64
YAR020C	0.98	1.40	1.69	1.90	2.07	36.79	2.53	118.88	25.84	155.57	28.37
YAR027W	0.91	1.59	1.87	2.20	2.27	21.74	2.42	75.08	20.23	95.82	22.64
YAR029W	0.95	1.65	1.97	2.26	2.32	22.57	2.19	78.99	19.56	101.56	21.75
YAR033W	0.89	1.61	1.89	2.24	2.30	29.26	3.56	92.78	26.62	122.04	30.18
YAR046W	0.92	1.65	1.96	2.24	2.31	22.61	2.41	77.14	21.46	99.75	23.87
YAR073W	0.88	1.39	1.68	2.01	2.09	26.77	3.38	79.75	28.88	106.52	32.25
YBL003C	0.90	1.33	1.50	1.80	1.92	25.51	2.89	74.14	23.05	99.65	25.94
YBL005W	0.98	1.60	1.95	2.10	2.24	27.39	2.06	104.89	24.26	132.28	26.31
YBL007C	0.93	1.57	1.94	2.24	2.32	26.10	3.73	84.08	27.60	110.18	31.32
YBL008W	0.90	1.53	1.84	2.17	2.23	46.80	10.42	112.58	49.68	159.38	60.10
YBL013W	0.91	1.49	1.82	2.16	2.27	36.43	4.27	104.73	39.20	141.16	43.47
YBL024W	0.87	1.32	1.66	2.12	2.26	41.77	2.47	105.07	24.06	146.84	26.53
YBL027W	0.86	1.51	1.80	2.12	2.20	31.25	2.66	101.33	23.08	132.58	25.74
YBL028C	0.98	1.89	2.16	2.31	24.3	22.06	2.51	82.17	16.14	104.25	18.64
YBL037W	1.06	1.79	2.10	2.34	2.45	24.55	2.06	83.12	21.43	107.67	23.50
YBL043W	1.02	1.75	2.04	2.32	2.38	25.33	2.03	87.02	19.10	112.35	21.13
YBL054W	0.98	1.67	1.95	2.25	2.33	30.80	2.81	95.41	24.75	126.22	27.56
YBL082C	1.03	1.76	2.02	2.27	2.37	25.53	2.48	79.63	23.71	103.17	26.19
YBR016W	1.00	1.52	1.85	2.09	2.21	31.45	2.77	112.88	28.00	144.34	30.78
YBR033W	0.96	1.35	1.51	1.82	2.01	30.01	3.27	101.11	33.79	137.12	37.06
YBR047W	1.01	1.68	2.04	2.28	2.43	31.29	3.53	85.53	30.51	117.86	30.82
YBR059W	0.98	1.61	1.90	2.20	2.34	28.02	2.72	88.97	22.22	116.58	28.49
YBR076C	0.97	1.41	1.71	2.10	2.25	37.29	3.69	116.20	33.36	149.28	33.65
YBR101C	1.19	1.83	2.08	2.35	2.49	33.19	4.67	89.93	36.65	122.73	41.22
YBR103W	0.96	1.26	1.58	1.96	2.12	28.62	4.48	102.44	37.62	131.26	42.10
YBR108W	0.98	1.42	1.75	2.08	2.22	28.03	2.38	110.63	30.74	138.66	33.12
YBR114W	1.08	1.86	2.10	2.37	2.41	11.17	1.52	68.80	6.21	75.98	7.72
YBR125C	1.04	1.84	2.17	2.36	2.51	33.42	4.94	79.55	37.33	112.37	42.27
YBR126C	0.93	1.47	1.78	2.23	2.31	25.96	2.06	90.69	15.91	116.65	17.97
YBR132C	0.97	1.18	1.50	2.04	2.11	30.33	3.48	95.47	29.41	125.80	32.89
YBR149W	0.97	1.48	1.84	2.15	2.38	55.66	8.36	122.79	56.42	178.44	64.79
YBR150C	0.92	1.37	1.88	2.19	2.24	25.62	3.32	74.53	24.61	100.14	27.92
YBR162W	0.98	1.54	1.81	2.17	2.24	17.55	1.36	75.92	22.70	93.48	24.06
YBR164C	0.95	1.64	1.95	2.25	2.34	27.51	1.78	95.19	23.62	120.91	25.39
YBR171W	1.03	1.43	1.79	2.24	2.35	40.15	0.63	105.03	19.36	145.18	19.99
YBR176W	1.10	1.83	2.13	2.41	2.45	22.55	1.20	79.12	22.38	101.66	23.58
YBR180W	0.98	1.63	1.99	2.23	2.31	40.19	3.29	105.09	34.31	145.28	37.60
YBR183W	1.12	1.80	2.08	2.41	2.47	25.29	0.76	94.46	17.40	119.75	18.16
YBR186W	0.91	1.22	1.53	2.00	2.05	23.48	3.04	90.82	25.50	114.30	26.63
YBR187W	1.06	1.87	2.17	2.39	2.47	24.48	1.95	82.93	19.99	107.42	21.94
YBR188C	1.08	1.66	1.92	2.14	2.24	21.55	1.25	83.48	18.34	105.03	19.58
YBR194W	1.04	1.54	1.85	2.25	2.33	27.51	1.78	95.19	23.62	120.91	25.39
YBR195C	1.03	1.85	2.14	2.33	2.43	26.13	2.68	88.45	28.00	114.58	30.68
YBR197C	0.90	1.23	1.47	1.90	2.04	26.86	1.67	92.03	26.60	118.91	28.27
YBR201W	0.96	1.40	1.74	2.12	2.24	32.83	0.25	113.35	16.79	146.18	3.51
YBR203W	0.87	1.18	1.51	2.04	2.11	30.33	3.48	95.47	29.41	125.80	33.09
YBR204C	0.91	1.61	1.80	2.14	2.21	15.61	2.81	65.14	22.18	80.75	25.00
YBR204C	1.15	1.14	1.54	2.12	2.16	106.59	1.55	115.29	183.63	190.09	205.38
YBR205W	1.04	1.75	2.04	2.32	2.39	17.57	1.36	80.20	23.98	97.77	25.34
YBR207W	0.95	1.60	1.91	2.23	2.33	22.88	2.95	93.29	31.99	116.17	34.94
YBR217W	1.04	1.64	1.91	2.27	2.37	23.31	6.75	95.85	40.71	122.22	55.45
YBR218W	1.04	1.28	1.69	2.11	2.18	24.21	2.80	117.53	19.47	147.20	24.26
YBR219C	1.03	1.33	1.62	2.24	2.34	23.47	3.27	118.83	15.54	126.12	51.66
YBR219C	0.98	1.48	1.79	2.17	2.25	27.76	2.37	94.64	22.62	115.29	21.99
YBR217W	1.08	1.65	1.97	2.34	2.38	34.44	1.85	111.44	25.50	145.88	32.20
YBR219W	1.08	1.91	2.20	2.41	2.48	26.20	3.61	82.83	21.24	109.03	35.35
YBR222C	0.98	1.64	1.95	2.24	2.34	26.77	3.12	81.47	27.02	108.23	30.14
YBR223C	0.99	1.52	1.77	2.14	2.22	23.23	1.48	96.01	24.26	119.24	25.74
YBR227C	1.09	1.60	1.92	2.34	2.41	24.52	2.55	88.34	32.49	112.86	35.04
YBR228W	1.09	1.80	2.07	2.37	2.45	20.75	4.19	85.89	47.42	110.64	51.60
YBR230C	0.99	1.45	1.73	2.15	2.24	27.67	2.00	99.53	23.78	127.20	25.78
YBR231C	0.88	1.11	1.29	1.70	1.84	51.68	1.13	119.38	20.58	171.06	21.71
YBR233W	0.96	1.41	1.73	2.17	2.27	26.73	0.48	92.06	15.82	118.79	16.35
YBR236C	0.99	1.40	1.76	2.17	2.36	24.95	1.73	84.01	17.39	137.20	3.54
YBR239C	0.97	1.64	1.96	2.25	2.32	25.32	5.45	92.09	41.59	127.29	47.04
YBR240C	1.03	1.76	2.04	2.31	2.42	21.54	3.42	63.87	30.13	85.41	33.55
YBR241C	1.00	1.66	2.01	2.24	2.38	39.89	9.49	99.34	52.51	139.23	62.00
YBR242W	0.88	1.38	1.88	2.03	2.12	22.74	2.53	94.4			

ORF	Optical density at 600 nm at time intervals (h)										GLUCOSE		FRUCTOSE		TOTAL Sugars(g/L)		TOTAL AMINO ACIDS (mg/L)	
	0	20	27	42	67	43 h	67 h	43 h	67 h	43 h	67 h	43 h	67 h	43 h	67 h	48 h		
YCD032W	0.65	1.08	1.39	2.03	2.03	37.45	3.52	93.32	29.06	126.32	31.75	52.23	52.23	304.19	304.19			
YCR034W	1.07	1.59	1.61	2.31	2.35	25.21	2.45	66.02	15.89	114.23	18.34	32.58	32.58	203.35	203.35			
YCD034W	0.91	1.18	1.70	2.17	2.24	22.80	2.68	69.20	17.67	117.76	20.81	32.58	32.58	232.05	232.05			
YCR035C	0.88	1.17	1.44	1.94	2.00	28.93	2.66	88.83	18.15	117.76	20.81	32.58	32.58	232.05	232.05			
YCR036C	0.94	1.41	1.75	2.19	2.24	24.57	2.33	97.67	20.90	122.23	23.23	31.21	31.21	256.56	256.56			
YCR034W	0.84	0.97	1.08	1.36	1.36	60.86	16.08	110.28	59.53	171.13	75.66	354.71	354.71	30.10	30.10			
YCR075W-A	0.86	0.85	0.86	0.98	0.87	142.4	128.30	160.86	195.93	303.00	334.23	521.41	521.41	324.23	324.23			
YCR106W	0.92	1.25	1.54	2.06	2.14	31.60	3.30	93.87	25.80	125.46	30.10	354.71	354.71	30.10	30.10			
YDL001W	0.91	0.90	0.90	1.00	0.91	104.91	108.18	123.63	166.17	278.55	274.35	523.06	523.06	274.35	274.35			
YDL012C	0.97	1.45	1.81	2.26	2.30	22.87	4.58	59.47	29.78	82.33	34.36	358.05	358.05	34.36	34.36			
YDL022W	0.81	1.24	1.58	2.11	2.18	21.82	4.26	70.57	39.78	92.38	44.04	286.68	286.68	44.04	44.04			
YDL039C	0.87	1.10	1.35	1.88	1.95	28.93	3.97	75.59	28.80	105.51	32.76	253.12	253.12	32.76	32.76			
YDL076C	0.96	1.64	1.95	2.25	2.34	19.53	0.65	80.73	17.28	100.26	17.93	164.02	164.02	17.93	17.93			
YDL078C	0.93	1.73	2.04	2.28	2.37	21.43	0.48	83.66	16.83	105.08	17.30	138.52	138.52	17.30	17.30			
YDL081C	0.92	0.96	1.05	1.58	1.92	86.70	26.88	159.26	101.47	245.96	128.35	328.89	328.89	128.35	128.35			
YDL082W	1.01	1.98	2.06	2.16	2.24	8.90	0.50	65.79	4.99	74.69	5.50	74.07	74.07	5.50	5.50			
YDL083C	0.93	1.05	1.05	1.08	1.02	183.51	163.40	167.18	152.56	350.69	315.96	522.99	522.99	315.96	315.96			
YDL085C-A	0.95	1.04	1.03	0.96	0.98	181.55	183.36	183.45	165.12	345.59	348.48	517.58	517.58	348.48	348.48			
YDL085W	0.92	1.18	1.44	2.03	2.25	57.08	3.83	124.19	37.38	181.26	41.20	155.28	155.28	41.20	41.20			
YDL086W	0.92	1.45	1.75	2.13	2.26	20.72	0.52	77.94	17.80	98.66	18.32	110.66	110.66	18.32	18.32			
YDL088C	0.85	1.30	1.70	2.00	2.16	25.00	0.45	85.18	14.24	110.18	14.69	99.42	99.42	14.69	14.69			
YDL091C	0.98	1.72	1.97	2.24	2.31	10.99	-0.12	64.50	9.64	75.49	9.53	109.82	109.82	9.53	9.53			
YDL093W	0.95	1.72	2.01	2.27	2.35	15.24	1.26	68.90	11.21	84.14	12.46	140.60	140.60	12.46	12.46			
YDL096C	1.10	1.54	1.88	2.28	2.37	32.04	3.28	100.55	35.04	132.59	38.32	146.48	146.48	38.32	38.32			
YDL099W	0.98	1.60	1.89	2.23	2.29	13.16	-0.02	71.18	11.83	84.34	11.81	127.32	127.32	11.81	11.81			
YDL100C	0.98	1.56	1.84	2.19	2.31	22.04	1.84	91.53	13.58	113.57	13.73	207.02	207.02	13.73	13.73			
YDL109C	1.03	1.80	2.06	2.27	2.39	19.16	0.14	81.72	14.20	100.89	14.35	133.69	133.69	14.35	14.35			
YDL112W	0.98	1.76	2.07	2.33	2.42	15.84	0.30	75.45	15.75	91.30	16.04	158.84	158.84	16.04	16.04			
YDL114W	0.96	1.61	1.89	2.21	2.31	16.45	0.15	74.84	13.61	91.29	13.76	112.14	112.14	13.76	13.76			
YDL117W	0.96	1.55	1.86	2.18	2.31	22.41	2.02	75.53	10.62	97.95	12.65	282.31	282.31	12.65	12.65			
YDL120C	0.94	1.52	1.80	2.14	2.28	16.29	0.24	80.44	9.68	90.78	10.05	10.05	10.05	9.68	9.68			
YDL127C	0.95	1.47	1.76	2.17	2.28	7.69	0.02	74.41	1.74	77.16	1.76	238.36	238.36	1.76	1.76			
YDL128W	0.91	1.39	1.70	2.10	2.19	23.23	0.17	81.37	22.32	89.93	25.60	155.53	155.53	25.60	25.60			
YDL130W	0.98	1.41	1.71	2.09	2.19	23.23	0.17	88.00	13.63	131.22	13.74	134.08	134.08	13.74	13.74			
YDL130W-A	0.95	1.20	1.44	1.93	2.03	32.18	3.57	120.70	24.74	134.29	28.32	111.59	111.59	28.32	28.32			
YDL144C	1.02	1.80	2.08	2.30	2.38	13.44	-0.11	75.72	9.84	89.16	9.73	72.23	72.23	9.73	9.73			
YDL149W	0.94	1.33	1.66	2.15	2.18	33.50	6.38	100.34	43.31	133.84	49.69	208.71	208.71	49.69	49.69			
YDL154W	0.98	1.34	1.69	2.10	2.23	32.31	0.06	82.60	15.10	101.92	15.17	83.83	83.83	15.17	15.17			
YDL154W	0.92	1.76	2.04	2.28	2.36	13.69	-0.05	65.94	10.68	75.63	10.63	78.35	78.35	10.63	10.63			
YDL159W	1.01	1.80	2.08	2.30	2.38	13.44	-0.16	74.20	8.98	88.18	8.82	105.49	105.49	8.82	8.82			
YDL161W	0.99	1.73	2.17	2.15	2.24	15.73	0.46	71.03	13.07	86.77	13.53	243.95	243.95	13.53	13.53			
YDL169W	0.97	1.63	2.24	1.99	2.00	13.41	0.03	68.22	12.88	81.63	13.01	146.26	146.26	13.01	13.01			
YDL180C	0.90	1.49	1.83	2.02	2.14	21.96	1.80	80.14	19.13	102.10	20.93	80.94	80.94	20.93	20.93			
YDL194W	0.91	1.60	1.81	2.07	2.21	16.48	2.31	67.84	20.83	84.32	23.14	75.77	75.77	23.14	23.14			
YDL198C	1.03	1.88	2.16	2.38	2.44	20.54	1.40	85.56	23.29	106.10	24.69	87.39	87.39	24.69	24.69			
YDL200C	0.95	1.69	1.97	2.22	2.32	11.02	0.32	67.17	11.85	78.19	12.17	107.24	107.24	12.17	12.17			
YDL201W	1.03	1.73	2.06	2.34	2.43	23.19	0.19	89.50	18.78	116.69	19.67	88.93	88.93	19.67	19.67			
YDL203C	0.92	1.53	1.79	2.13	2.26	14.52	0.11	79.20	7.92	93.72	8.03	168.07	168.07	8.03	8.03			
YDL211C	0.90	1.47	1.78	2.10	2.24	19.78	0.76	83.66	14.64	103.46	15.40	149.18	149.18	15.40	15.40			
YDL213C	0.93	1.55	1.84	2.19	2.31	16.40	1.02	69.13	16.88	85.53	17.91	135.34	135.34	17.91	17.91			
YDL215C	0.93	1.43	1.71	2.09	2.20	19.32	1.13	78.55	18.25	97.88	19.38	151.57	151.57	19.38	19.38			
YDL219W	0.87	1.38	1.69	2.08	2.19	18.04	1.05	73.55	19.05	91.60	20.11	88.37	88.37	20.11	20.11			
YDL222C	0.85	1.24	1.77	1.93	2.05	15.34	1.06	72.20	19.79	87.55	20.85	73.02	73.02	20.85	20.85			
YDL226C	0.91	1.59	1.81	2.15	2.21	24.02	4.29	75.05	23.73	99.08	34.02	152.09	152.09	34.02	34.02			
YDL227C	0.91	1.56	1.77	2.17	2.26	17.88	3.50	63.37	20.49	86.23	31.96	126.73	126.73	31.96	31.96			
YDL233W	1.05	1.15	1.13	1.62	1.68	100.50	160.54	100.50	160.54	317.80	331.56	518.46	518.46	331.56	331.56			
YDL234W	1.04	1.44	2.11	2.34	2.38	17.65	2.31	62.57	25.25	101.93	20.39	200.57	200.57	20.39	20.39			
YDL234C	1.00	1.83	2.10	2.33	2.37	14.14	0.57	68.07	14.73	83.21	15.30	76.84	76.84	15.30	15.30			
YDL239C	0.96	1.76	2.05	2.28	2.37	13.33	0.79	73.38	18.87	85.70	18.66	119.62	119.62	18.66	18.66			
YDL243C	0.97	1.79	2.05	2.28	2.34	15.28	0.35	73.82	12.07	89.10	12.42	116.25	116.25	12.42	12.42			
YDL245C	0.87	1.65	1.98	2.25	2.30	19.11	1.75	74.45	18.04	92.56	19.79	150.51	150.51	19.79	19.79			
YDR003W	0.98	1.67	1.98	2.22	2.31	16.38	0.38	74.84	9.62	91.20	10.01	195.95	195.95	10.01	10.01			
YDR004W	0.98	1.54	1.79	2.09	2.24	28.38	1.75	73.99	23.85	122.37	25.60	146.65	146.65	25.60	25.60			
YDR005C	0.93	1.37	1.64	2.02	2.17	21.05	1.18	87.23	17.56	108.28	18.75	207.06	207.06	18.75	18.75			
YDR009W	0.89</																	

ORF	Optical density at 600 nm at time intervals (h)										GLUCOSE		FRUCTOSE		TOTAL Sugars(g/L)		TOTAL AMINO ACIDS (mM/L)	
	0	20	27	42	67	43 h	67 h	43 h	67 h	43 h	67 h	48 h	67 h	43 h	67 h	48 h		
YDR281C	0.49	1.18	1.49	1.57	2.04	27.0	3.56	93.7	29.47	120.55	37.0	113.59	37.0	113.59	37.0	113.59		
YDR282C	0.50	1.12	1.40	1.52	2.03	23.00	3.11	97.30	25.21	110.30	26.72	104.23	26.72	104.23	26.72	104.23		
YDR283C	0.89	1.27	1.46	1.56	2.03	26.89	3.57	83.11	25.45	120.26	26.02	122.26	26.02	122.26	26.02	122.26		
YDR284C	0.96	1.42	1.48	2.16	2.22	28.53	4.12	107.61	25.82	136.33	26.94	130.08	26.94	130.08	26.94	130.08		
YDR291W	1.00	1.83	2.09	2.32	2.37	16.17	0.48	72.26	21.96	88.43	22.45	80.15	22.45	80.15	22.45	80.15		
YDR298W	0.94	1.70	1.98	2.24	2.29	15.22	1.87	76.88	19.36	91.88	21.23	79.53	21.23	79.53	21.23	79.53		
YDR299W	0.88	1.13	1.47	2.00	2.09	30.92	3.68	100.30	27.06	131.22	30.74	104.99	30.74	104.99	30.74	104.99		
YDR299W	0.97	1.06	1.04	0.98	0.98	207.77	175.47	194.06	166.30	491.84	341.78	526.16	341.78	526.16	341.78	526.16		
YDR304C	0.95	1.04	1.02	0.98	0.97	180.10	174.81	187.77	168.83	367.87	343.65	522.67	343.65	522.67	343.65	522.67		
YDR305C	0.94	1.51	1.79	2.14	2.25	17.69	0.74	78.09	25.04	95.78	25.78	113.59	25.78	113.59	25.78	113.59		
YDR306C	0.90	1.18	1.56	2.10	2.18	29.42	3.31	101.95	30.05	131.37	33.37	109.80	33.37	109.80	33.37	109.80		
YDR307W	0.94	1.63	1.86	2.13	2.28	15.27	0.17	70.18	17.40	85.45	17.57	79.53	17.57	79.53	17.57	79.53		
YDR309C	0.91	1.28	1.67	2.15	2.23	29.78	3.32	102.38	30.67	132.16	33.99	98.59	33.99	98.59	33.99	98.59		
YDR312W	0.96	1.43	1.81	2.18	2.27	20.25	2.73	72.26	19.34	92.54	22.07	114.24	22.07	114.24	22.07	114.24		
YDR313C	0.88	1.37	1.68	2.06	2.22	15.75	0.24	69.80	18.86	85.61	19.09	130.58	19.09	130.58	19.09	130.58		
YDR316W	0.87	1.22	1.48	1.90	2.11	24.83	0.87	90.10	28.83	114.93	29.69	99.97	29.69	99.97	29.69	99.97		
YDR320C	1.06	1.91	2.14	2.31	2.41	10.49	3.09	66.20	8.22	76.69	11.31	87.09	11.31	87.09	11.31	87.09		
YDR321W	1.09	1.80	2.05	2.28	2.39	22.27	1.90	87.11	34.43	109.38	36.33	124.84	36.33	124.84	36.33	124.84		
YDR326C	1.00	1.81	2.07	2.31	2.36	13.92	0.44	75.78	18.75	89.70	19.19	99.44	19.19	99.44	19.19	99.44		
YDR329C	0.93	1.49	1.86	2.22	2.31	27.49	3.03	105.04	20.92	132.53	23.95	101.45	23.95	101.45	23.95	101.45		
YDR330W	0.86	1.22	1.57	2.03	2.13	22.95	2.91	74.32	21.01	97.27	23.92	106.09	23.92	106.09	23.92	106.09		
YDR332W	0.86	1.15	1.45	1.96	2.07	20.42	2.36	83.20	21.11	103.62	23.46	224.93	23.46	224.93	23.46	224.93		
YDR333C	0.96	1.27	1.61	2.10	2.20	36.06	4.26	112.64	32.87	148.69	37.13	190.14	37.13	190.14	37.13	190.14		
YDR336W	0.87	1.07	1.34	1.85	1.96	27.17	2.52	114.44	24.13	141.61	26.65	99.11	26.65	99.11	26.65	99.11		
YDR342C	0.90	1.62	1.99	2.24	2.34	25.51	2.96	82.95	23.36	108.47	25.34	103.58	25.34	103.58	25.34	103.58		
YDR343C	0.86	1.62	2.01	2.24	2.38	23.97	2.96	73.49	19.65	97.47	22.61	130.51	22.61	130.51	22.61	130.51		
YDR345C	0.87	1.08	1.37	1.93	2.02	20.32	4.03	67.77	27.86	88.10	31.89	147.96	31.89	147.96	31.89	147.96		
YDR351W	0.88	1.11	1.43	1.97	2.06	26.71	2.67	110.46	22.32	137.17	24.99	277.40	24.99	277.40	24.99	277.40		
YDR354W	1.00	1.59	1.73	2.03	2.22	26.34	3.37	92.27	38.43	118.61	41.81	176.00	41.81	176.00	41.81	176.00		
YDR357C	1.01	1.71	1.99	2.28	2.39	23.23	1.16	88.22	26.57	111.48	27.85	100.91	27.85	100.91	27.85	100.91		
YDR358W	0.84	1.49	1.71	2.06	2.16	11.04	1.33	56.05	10.51	87.87	11.63	86.48	11.63	86.48	11.63	86.48		
YDR360C	1.01	1.21	1.45	1.51	1.51	46.46	5.71	93.26	16.86	114.26	18.19	240.35	18.19	240.35	18.19	240.35		
YDR363W	0.96	1.06	1.10	1.03	0.99	170.36	15.06	180.77	154.47	351.13	307.33	531.17	307.33	531.17	307.33	531.17		
YDR364W-A	1.03	1.55	1.81	2.19	2.26	11.86	2.16	58.05	10.79	63.79	12.05	195.01	12.05	195.01	12.05	195.01		
YDR365C	0.98	1.61	1.90	2.26	2.34	22.82	1.63	64.58	31.81	117.10	33.44	124.84	33.44	124.84	33.44	124.84		
YDR366W	0.96	1.50	1.73	2.14	2.22	11.06	1.61	47.70	15.69	58.76	17.70	288.28	17.70	288.28	17.70	288.28		
YDR370C	0.98	1.61	1.94	2.29	2.32	16.27	1.82	65.87	15.22	82.15	17.04	181.55	17.04	181.55	17.04	181.55		
YDR372C	0.92	1.50	1.82	2.16	2.27	17.07	0.49	74.37	21.26	91.44	21.67	98.49	21.67	98.49	21.67	98.49		
YDR375C	0.96	1.50	1.82	2.16	2.25	16.42	0.50	70.80	19.79	87.22	20.29	116.26	20.29	116.26	20.29	116.26		
YDR377W	0.90	1.49	1.90	2.14	2.33	25.57	2.99	69.77	25.86	92.34	28.95	131.51	28.95	131.51	28.95	131.51		
YDR378C	0.97	1.61	1.90	2.21	2.31	18.98	1.33	75.34	25.84	94.32	28.87	137.41	28.87	137.41	28.87	137.41		
YDR379C-A	0.97	1.74	2.04	2.30	2.33	10.16	0.84	76.63	20.56	95.98	21.41	164.24	21.41	164.24	21.41	164.24		
YDR379W	0.93	1.44	1.78	2.19	2.23	20.04	2.25	72.55	14.81	82.38	17.07	277.84	17.07	277.84	17.07	277.84		
YDR382W	0.93	1.11	1.29	1.95	2.23	60.69	14.45	121.14	16.69	181.84	19.14	175.21	19.14	175.21	19.14	175.21		
YDR383C	1.04	1.64	1.98	2.27	2.33	20.86	2.00	69.06	17.19	89.92	19.20	245.56	19.20	245.56	19.20	245.56		
YDR385W	1.01	1.39	2.08	2.06	2.20	27.63	5.27	68.23	13.56	95.87	18.83	278.51	18.83	278.51	18.83	278.51		
YDR386W	1.06	1.50	1.77	2.19	2.32	26.34	4.24	65.78	12.48	92.12	16.73	394.64	16.73	394.64	16.73	394.64		
YDR388W	0.97	1.41	1.68	2.18	2.28	22.25	2.61	64.16	18.31	86.41	20.93	384.09	20.93	384.09	20.93	384.09		
YDR389W	1.00	1.36	1.82	2.31	2.37	38.37	9.75	110.90	48.02	149.27	57.59	264.62	57.59	264.62	57.59	264.62		
YDR391C	0.99	1.43	1.80	2.27	2.34	38.60	5.27	89.93	32.92	128.53	38.19	280.44	38.19	280.44	38.19	280.44		
YDR420W	0.86	1.31	1.78	2.00	2.23	26.23	2.78	80.75	23.88	107.05	26.61	97.87	26.61	97.87	26.61	97.87		
YDR423C	0.92	1.65	1.88	2.11	2.24	17.51	1.75	67.46	15.84	84.97	18.64	97.03	18.64	97.03	18.64	97.03		
YDR424C	0.92	1.65	1.88	2.10	2.28	18.87	1.22	69.92	26.45	85.01	30.05	311.67	30.05	311.67	30.05	311.67		
YDR425C	0.96	1.56	1.82	2.10	2.28	23.55	2.51	76.71	20.51	103.60	22.15	216.55	22.15	216.55	22.15	216.55		
YDR426C	0.95	1.56	1.87	2.24	2.31	25.23	2.87	78.38	22.27	103.60	32.15	142.17	32.15	142.17	32.15	142.17		
YDR426W	1.05	1.56	1.87	2.24	2.31	25.23	3.18	73.55	26.69	99.78	28.87	141.62	28.87	141.62	28.87	141.62		
YDR427C	0.93	1.64	1.98	2.26	2.36	23.32	3.18	71.85	34.86	95.17	38.04	166.77	38.04	166.77	38.04	166.77		
YDR427W	0.94	1.57	1.86	2.20	2.28	23.16	1.52	72.00	28.95	95.16	30.47	196.88	30.47	196.88	30.47	196.88		
YDR479C	0.91	1.40	1.74	2.12	2.23	27.53	2.81	80.43	32.04	107.97	34.86	149.23	34.86	149.23	34.86	149.23		
YDR482C	0.97	1.94	2.82	2.20	2.28	19.13	1.20	70.56	23.87	89.69	25.07	84.58	25.07	84.58	25.07	84.58		
YDR485C	0.98	1.57																

ORF	Optical density at 600 nm time intervals (h)										GLUCOSE		FRUCTOSE		TOTAL Sugars(g/L)		TOTAL AMINO ACIDS (mg/L)	
	0	20	27	42	67	43 h	67 h	43 h	67 h	43 h	67 h	48 h	67 h	43 h	67 h	48 h		
YER072W	1.62	1.58	1.88	2.05	2.33	23.0	2.90	61.70	70.00	85.20	20.80	148.54	52.42	101.57	116.11	122.42		
YER075C	1.62	1.55	1.98	2.36	2.43	24.5	5.53	7.29	38.87	55.54	44.50	18.44	83.87	20.76	20.76	20.76		
YER080W	1.00	1.53	1.55	2.30	2.31	26.7	1.77	102.49	102.49	129.25	129.25	189.44	270.33	115.31	154.49	154.49		
YER081W	1.62	1.58	1.86	2.24	2.32	21.86	2.14	65.45	16.29	88.25	18.94	116.11	188.49	23.45	34.69	34.69		
YER085C	0.98	1.24	1.48	1.94	2.08	27.44	4.27	106.48	30.42	143.92	34.69	301.32	119.57	125.19	125.19	125.19		
YER088C	1.02	1.62	1.92	2.28	2.34	23.89	1.88	95.63	21.57	129.52	23.45	144.32	22.27	133.38	22.27	22.27		
YER090W	1.02	1.62	1.95	2.28	2.35	27.88	1.64	105.50	20.63	108.82	21.06	94.98	125.19	21.59	21.59	21.59		
YER090C	0.98	1.48	1.79	2.18	2.27	30.83	1.65	105.33	20.56	136.16	22.22	101.41	125.19	21.59	21.59	21.59		
YER115C	0.92	1.03	0.99	0.96	0.93	163.53	187.40	171.69	177.78	335.22	365.18	517.05	177.78	177.78	177.78	177.78		
YER116C	1.01	1.37	1.61	1.59	2.23	66.03	11.94	131.42	66.04	197.45	77.98	226.91	116.11	116.11	116.11	116.11		
YER118C	1.06	1.55	1.80	2.17	2.22	13.98	1.75	52.83	9.87	66.80	11.61	188.49	125.19	125.19	125.19	125.19		
YER119C	1.08	1.81	2.09	2.34	2.39	23.07	0.80	101.81	15.86	124.89	16.45	90.65	19.93	19.93	19.93	19.93		
YER122C	1.07	1.92	2.17	2.39	2.42	24.17	1.28	102.02	18.64	126.19	19.93	144.32	22.08	22.08	22.08	22.08		
YER123W	1.05	1.63	1.83	2.24	2.31	21.06	2.63	68.80	20.37	89.86	23.00	221.93	125.19	125.19	125.19	125.19		
YER124C	1.06	1.88	2.16	2.38	2.42	24.37	1.13	100.82	21.06	104.55	23.45	111.26	22.22	22.22	22.22	22.22		
YER128W	1.02	1.66	1.96	2.29	2.35	24.48	1.33	102.34	20.94	126.82	22.28	87.42	125.19	21.59	21.59	21.59		
YER129W	1.03	1.70	1.96	2.29	2.32	17.90	2.28	56.64	13.71	76.55	15.99	322.22	177.78	177.78	177.78	177.78		
YER134C	0.99	1.56	1.88	2.22	2.29	27.77	1.10	98.88	19.86	126.66	20.96	89.83	125.19	21.59	21.59	21.59		
YER137C	0.98	1.50	1.85	1.92	2.02	17.15	1.71	73.59	13.67	90.73	15.38	197.28	125.19	125.19	125.19	125.19		
YER144C	1.06	1.81	2.06	2.34	2.37	28.36	2.55	84.89	18.43	113.25	20.97	262.87	125.19	125.19	125.19	125.19		
YER145C	1.06	1.55	1.84	2.21	2.25	22.70	2.39	71.02	19.69	93.72	22.08	329.15	125.19	125.19	125.19	125.19		
YER150W	1.00	1.66	1.98	2.31	2.34	19.25	2.71	67.30	18.13	82.95	20.84	224.07	125.19	125.19	125.19	125.19		
YER151C	1.02	1.68	1.99	2.29	2.36	28.96	1.66	110.45	23.87	139.41	25.53	102.26	125.19	21.59	21.59	21.59		
YER152C	1.06	1.67	1.99	2.33	2.37	20.24	2.19	72.58	16.20	92.82	20.39	220.36	125.19	125.19	125.19	125.19		
YER153C	1.02	1.52	1.84	2.26	2.32	25.46	2.94	72.47	23.49	97.93	26.43	248.00	125.19	125.19	125.19	125.19		
YER155C	0.98	1.59	1.84	2.22	2.29	16.76	1.93	68.68	13.23	85.45	15.16	178.11	125.19	125.19	125.19	125.19		
YER156C	0.98	1.55	2.08	2.26	2.34	24.15	2.04	75.88	15.41	101.04	17.45	189.39	125.19	125.19	125.19	125.19		
YER161C	1.17	1.96	2.18	2.44	2.48	14.77	1.58	61.55	12.16	76.32	13.74	143.44	125.19	125.19	125.19	125.19		
YER162C	1.02	1.68	1.96	2.30	2.34	23.69	2.25	70.11	19.55	87.06	21.89	256.29	125.19	125.19	125.19	125.19		
YER163C	1.05	1.63	1.94	2.30	2.34	24.83	2.66	73.58	16.26	126.30	17.42	96.29	125.19	125.19	125.19	125.19		
YER164W	1.05	1.63	2.08	2.39	2.42	18.93	2.00	63.60	15.77	87.53	17.78	125.46	125.19	125.19	125.19	125.19		
YER165W	1.06	1.64	2.07	2.36	2.39	23.48	1.91	64.32	16.41	81.18	18.32	133.53	125.19	125.19	125.19	125.19		
YER167W	1.03	1.77	1.81	2.23	2.28	19.42	2.26	64.64	15.89	84.15	18.16	111.50	125.19	125.19	125.19	125.19		
YER174C	0.98	1.63	1.90	2.19	2.26	13.69	0.29	68.90	10.69	82.58	9.80	137.37	125.19	125.19	125.19	125.19		
YER180C	0.94	1.05	1.00	0.99	0.95	137.58	184.80	145.07	170.33	282.64	355.13	513.81	125.19	125.19	125.19	125.19		
YER180C-A	0.93	1.08	1.18	1.93	1.90	167.12	177.18	167.25	163.08	334.37	340.26	519.66	125.19	125.19	125.19	125.19		
YER181C-A	0.96	1.02	1.01	0.97	0.96	166.42	210.94	165.42	169.95	331.85	380.89	515.91	125.19	125.19	125.19	125.19		
YER183C	0.98	1.72	2.01	2.27	2.32	17.94	0.47	73.14	15.52	91.08	15.98	96.81	125.19	125.19	125.19	125.19		
YER184C	1.01	1.54	1.76	2.14	2.19	15.45	1.87	62.62	12.35	78.08	15.13	226.95	125.19	125.19	125.19	125.19		
YER185W	0.97	1.67	1.92	2.22	2.29	12.40	0.30	62.92	15.26	75.32	15.56	76.13	125.19	125.19	125.19	125.19		
YER007W	0.92	1.58	1.88	2.18	2.25	23.64	1.32	82.95	16.97	106.60	18.29	101.96	21.42	100.55	21.42	100.55		
YFL010W-A	0.96	1.62	1.90	2.21	2.28	19.54	0.89	69.39	18.30	88.54	19.19	116.45	125.19	21.59	21.59	21.59		
YFL011W	1.03	1.51	1.77	2.19	2.24	24.69	3.07	73.34	22.80	98.03	25.87	249.33	125.19	125.19	125.19	125.19		
YFL012W	0.93	1.59	1.89	2.20	2.26	21.03	2.80	69.51	30.69	92.54	33.49	168.86	125.19	125.19	125.19	125.19		
YFL019C	0.98	1.37	1.61	2.06	2.10	16.93	2.22	68.58	18.31	85.48	20.53	121.15	125.19	21.59	21.59	21.59		
YFL020C	1.06	1.54	1.73	2.12	2.18	13.87	2.03	55.35	20.24	69.21	22.27	170.15	125.19	125.19	125.19	125.19		
YFL021W	1.03	1.64	1.87	2.24	2.28	19.96	2.26	69.74	19.02	89.70	21.28	141.57	125.19	125.19	125.19	125.19		
YFL023W	0.99	1.42	1.75	2.17	2.27	22.99	2.03	73.96	19.39	100.55	21.42	342.01	125.19	125.19	125.19	125.19		
YFL026W	0.94	1.04	1.00	0.99	0.95	141.85	208.94	139.01	163.25	280.56	372.19	520.71	125.19	125.19	125.19	125.19		
YFL031W	1.01	1.62	1.81	2.10	2.13	34.95	5.29	88.08	37.45	124.03	42.74	131.79	125.19	125.19	125.19	125.19		
YFL036W	0.91	1.01	0.97	0.96	0.92	138.05	194.09	143.05	166.05	281.09	360.14	518.32	125.19	125.19	125.19	125.19		
YFL040W	0.92	1.40	1.69	2.07	2.16	23.27	0.69	76.01	22.68	99.28	23.37	93.87	125.19	125.19	125.19	125.19		
YFL042C	0.94	1.49	1.80	2.19	2.21	15.61	2.69	73.59	18.02	89.20	20.70	219.64	125.19	125.19	125.19	125.19		
YFL047W	0.96	1.11	1.06	1.06	1.01	143.41	247.78	135.49	166.59	278.99	314.37	519.56	125.19	125.19	125.19	125.19		
YFL048C	0.96	1.34	1.56	1.96	2.01	23.54	2.59	74.32	22.73	92.86	25.32	285.71	125.19	125.19	125.19	125.19		
YFL066C	1.06	1.77	2.04	2.35	2.38	23.16	1.62	87.34	11.47	110.50	12.80	202.90	125.19	125.19	125.19	125.19		
YFL067C	0.99	1.48	1.74	2.17	2.24	31.03	1.65	83.74	11.47	114.78	12.24	114.08	125.19	125.19	125.19	125.19		
YFL068W	0.98	1.47	1.76	2.17	2.24	31.03	1.65	83.74	11.47	114.78	12.24	114.08	125.19	125.19	125.19	125.19		
YFL069C	0.96	1.44	1.62	1.99	2.09	24.98	1.91	81.93	14.78	98.84	24.78	216.66	125.19	125.19	125.19	125.19		
YFL070C	0.97	1.73	2.04	2.34	2.38	16.64	2.08	61										

ORF	Optical density at 600 nm time intervals (h)										GLUCOSE		FRUCTOSE		TOTAL Sugars(g/L)		TOTAL AMINO ACIDS (mg/L)	
	0	2h	27	42	67	43 h	67 h	43 h	67 h	43 h	67 h	48 h	67 h	43 h	67 h	48 h		
YGL13TC	0.95	1.47	1.76	1.56	2.22	1.78	1.75	60.75	61.04	55.51	55.47	145.5	145.4	145.5	145.4	145.5		
YGL134W	0.97	1.58	1.92	2.21	2.26	2.16	1.8	56.14	57.03	104.01	104.01	141.58	141.58	141.58	141.58	141.58		
YGL135C	0.94	1.55	1.73	2.10	2.19	2.13	2.08	59.99	59.71	142.29	114.09	116.68	116.68	116.68	116.68	116.68		
YGL139C	0.94	1.53	1.79	2.13	2.21	1.98	1.98	57.61	53.22	6.21	92.49	15.28	256.91	256.91	256.91	256.91		
YGL141W	0.95	1.55	2.10	2.43	2.38	2.32	1.96	78.50	16.90	102.02	18.85	61.88	194.11	194.11	194.11	194.11		
YGL157W	1.11	1.79	2.08	2.35	2.40	2.27	2.43	2.42	2.32	105.09	25.69	160.41	160.41	160.41	160.41	160.41		
YGL158W	0.96	1.65	1.59	2.03	2.12	19.70	1.56	82.43	15.68	102.12	17.24	144.42	144.42	144.42	144.42	144.42		
YGL162W	0.95	1.45	1.70	2.07	2.15	24.19	1.52	92.06	18.08	116.25	15.61	162.23	15.56	15.56	15.56	15.56		
YGL164C	0.89	1.30	1.52	1.86	1.96	22.16	1.42	95.02	15.89	117.38	17.31	148.30	148.30	148.30	148.30	148.30		
YGL169W	1.02	1.42	1.66	1.98	2.09	36.18	5.55	111.79	58.64	147.97	64.19	489.23	489.23	489.23	489.23	489.23		
YGL175C	0.87	1.18	1.41	1.87	2.02	35.24	1.55	113.94	18.55	148.78	20.10	127.64	127.64	127.64	127.64	127.64		
YGL178W	1.10	1.76	2.07	2.35	2.42	16.77	1.80	67.96	11.54	84.73	13.34	332.81	332.81	332.81	332.81	332.81		
YGL181W	1.00	1.50	1.73	2.18	2.26	22.89	1.90	86.31	22.07	109.21	23.96	185.62	185.62	185.62	185.62	185.62		
YGL191W	0.85	1.27	1.56	1.92	2.04	23.69	0.61	103.72	12.19	127.40	12.81	147.07	147.07	147.07	147.07	147.07		
YGL192W	1.01	1.59	1.87	2.25	2.32	28.39	2.90	85.97	24.04	114.35	26.94	185.35	185.35	185.35	185.35	185.35		
YGL194C	1.03	1.58	1.94	2.28	2.33	43.26	6.91	118.59	48.75	162.23	55.56	177.22	177.22	177.22	177.22	177.22		
YGL205W	0.99	1.51	1.86	2.25	2.33	33.85	0.62	113.76	14.67	147.61	15.29	108.44	108.44	108.44	108.44	108.44		
YGL208W	0.97	1.62	1.90	2.07	2.23	18.36	0.26	90.16	14.11	108.52	14.37	113.98	113.98	113.98	113.98	113.98		
YGL210W	0.93	1.49	1.73	2.03	2.09	30.17	2.34	105.71	24.53	135.88	26.87	149.42	149.42	149.42	149.42	149.42		
YGL211W	0.98	1.54	1.79	2.12	2.22	31.40	1.19	113.73	19.75	145.14	20.95	82.81	82.81	82.81	82.81	82.81		
YGL212W	0.95	1.46	1.68	2.01	2.09	22.96	0.50	95.14	18.17	118.09	18.68	143.58	143.58	143.58	143.58	143.58		
YGL213C	0.92	1.34	1.51	1.86	2.02	26.84	1.15	103.51	14.38	130.35	15.53	172.11	172.11	172.11	172.11	172.11		
YGL219C	0.95	1.64	1.89	2.19	2.25	22.13	1.04	94.43	14.24	116.56	15.28	119.62	119.62	119.62	119.62	119.62		
YGL222C	0.96	1.61	1.85	2.15	2.22	22.87	1.32	90.51	13.92	113.37	15.24	144.15	144.15	144.15	144.15	144.15		
YGL227W	0.97	1.44	1.75	2.00	2.29	38.14	0.45	142.23	13.99	180.37	14.36	220.02	220.02	220.02	220.02	220.02		
YGL228W	0.95	1.56	1.82	2.12	2.18	43.92	5.71	122.28	39.31	166.21	45.02	180.28	180.28	180.28	180.28	180.28		
YGL230C	0.92	1.32	1.55	1.98	2.13	41.28	1.26	137.75	17.74	179.00	19.00	169.16	169.16	169.16	169.16	169.16		
YGL231C	0.94	1.70	1.94	2.17	2.21	14.83	0.82	83.48	9.68	98.31	10.50	196.78	196.78	196.78	196.78	196.78		
YGL233W	0.93	1.40	1.67	2.02	2.15	25.73	1.18	98.42	15.98	124.15	17.17	160.97	160.97	160.97	160.97	160.97		
YGL236C	0.87	1.31	1.58	1.94	2.06	26.42	1.05	104.01	15.01	130.43	17.12	113.38	113.38	113.38	113.38	113.38		
YGL242C	0.98	1.38	1.73	2.05	2.15	22.96	0.28	98.76	12.68	126.65	12.65	164.14	164.14	164.14	164.14	164.14		
YGL243C	0.98	1.37	1.55	2.00	2.20	16.87	0.47	12.63	6.24	100.87	9.37	160.58	160.58	160.58	160.58	160.58		
YGL249W	0.98	1.67	1.95	2.20	2.26	16.53	0.50	121.31	11.51	98.84	12.02	170.27	170.27	170.27	170.27	170.27		
YGL250W	0.98	1.67	1.90	2.20	2.26	16.64	16.45	125.45	16.23	131.46	16.45	188.18	188.18	188.18	188.18	188.18		
YGL257C	1.04	1.17	1.13	1.10	1.07	16.41	10.0	175.53	16.62	161.85	33.72	337.38	337.38	337.38	337.38	337.38		
YGL262C	0.95	1.35	1.66	1.98	2.13	42.73	1.72	142.08	17.85	184.61	19.58	189.58	189.58	189.58	189.58	189.58		
YGL263W	0.92	1.23	1.53	1.96	2.05	2.18	0.39	136.75	10.49	180.65	15.93	166.08	166.08	166.08	166.08	166.08		
YGL267C	0.98	1.55	1.80	2.11	2.21	20.98	0.64	94.29	12.09	115.25	12.73	148.82	148.82	148.82	148.82	148.82		
YGL268C	0.97	1.64	1.93	2.22	2.28	17.72	0.47	85.32	11.47	104.04	11.94	115.53	115.53	115.53	115.53	115.53		
YGL269C	0.94	1.47	1.72	2.02	2.11	22.40	1.00	100.70	15.84	123.30	16.59	126.58	126.58	126.58	126.58	126.58		
YGL270W	0.92	1.37	1.55	1.88	2.03	24.33	0.94	99.35	15.23	123.68	16.17	128.92	128.92	128.92	128.92	128.92		
YGR007W	1.02	1.75	2.00	2.27	2.35	18.33	0.39	69.35	17.19	87.69	17.58	148.67	148.67	148.67	148.67	148.67		
YGR023W	0.99	1.54	1.94	2.32	2.37	43.30	4.06	131.95	11.46	175.24	15.52	275.04	275.04	275.04	275.04	275.04		
YGR027C	0.92	1.03	0.99	1.05	1.09	157.71	162.18	161.91	150.44	319.81	312.63	515.21	515.21	515.21	515.21	515.21		
YGR014W	0.98	1.44	1.79	2.20	2.26	32.80	0.73	125.36	6.21	158.16	6.94	98.15	98.15	98.15	98.15	98.15		
YGR015C	0.96	1.69	1.99	2.25	2.32	23.41	0.49	100.29	12.70	123.70	13.19	175.24	175.24	175.24	175.24	175.24		
YGR016W	0.93	1.65	1.95	2.21	2.27	14.24	-0.14	69.45	10.44	83.69	9.90	67.62	67.62	67.62	67.62	67.62		
YGR017W	0.92	1.48	1.72	2.02	2.11	17.42	0.97	73.64	18.06	91.06	19.05	139.93	139.93	139.93	139.93	139.93		
YGR019W	1.01	1.45	1.81	2.28	2.35	38.84	1.86	118.54	10.34	157.38	12.20	259.71	259.71	259.71	259.71	259.71		
YGR021W	1.00	1.72	2.00	2.27	2.35	18.33	0.39	69.35	17.19	87.69	17.58	148.67	148.67	148.67	148.67	148.67		
YGR040W	1.07	1.62	1.96	2.24	2.38	11.41	0.48	83.44	9.52	154.24	15.24	154.24	154.24	154.24	154.24	154.24		
YGR041W	0.98	1.32	1.69	2.22	2.30	40.56	2.24	127.99	5.76	168.85	8.01	180.67	180.67	180.67	180.67	180.67		
YGR043C	0.96	1.25	1.56	2.09	2.11	34.42	1.82	106.05	5.09	139.48	7.90	277.17	277.17	277.17	277.17	277.17		
YGR044C	0.98	1.34	1.64	2.05	2.18	31.28	2.41	102.88	10.26	134.18	10.46	310.35	310.35	310.35	310.35	310.35		
YGR045C	1.00	1.41	1.63	2.02	2.11	31.02	0.77	127.72	10.22	161.48	14.49	246.09	246.09	246.09	246.09	246.09		
YGR071C	0.97	1.50	1.79	2.27	2.31	12.23	0.88	117.16	10.77	167.48	11.45	111.04	111.04	111.04	111.04	111.04		
YGR072W	0.98	1.48	1.76	2.16	2.23	28.95	0.42	121.06	15.06	159.31	16.83	264.93	264.93	264.93	264.93	264.93		
YGR073W	0.98	1.55	1.71	2.23	2.30	12.23	0.75	128.33	5.24	168.30	6.58	141.33	141.33	141.33	141.33	141.33		
YGR070W	0.95	1.53	1.85	2.17	2.24	24.60	1.61	82.60	18.48	107.20	20.09	116.13	116.13	116.13	116.13	116.13		
YGR072W	0.97	1.06	1.03	1.09	1.23	37.34												

ORF	Optical density at 600 nm at time intervals (h)										GLUCOSE		FRUCTOSE		TOTAL Sugars(g/L)		TOTAL AMINO ACIDS (mg/L)	
	0	20	27	42	67	43 h	67 h	43 h	67 h	43 h	67 h	43 h	67 h	43 h	67 h	48 h		
YGR206W	1.26	1.56	1.90	1.45	2.26	16.36	0.35	83.44	102.25	15.16	16.78	164.78	164.78	164.78	164.78			
YGR207C	1.01	1.78	2.03	2.29	2.38	21.16	0.96	88.40	98.26	101.66	103.32	158.33	158.33	158.33	158.33			
YGR207W	1.00	1.68	1.93	2.20	2.27	18.74	1.06	77.89	16.93	95.62	18.40	137.29	137.29	137.29	137.29			
YGR210C	0.91	1.41	1.90	2.05	2.17	18.04	1.08	78.71	16.85	96.74	24.73	70.54	70.54	70.54	70.54			
YGR212W	0.89	1.41	1.65	1.97	2.04	12.97	0.38	56.33	13.11	69.30	13.46	149.71	149.71	149.71	149.71			
YGR214W	0.96	1.46	1.71	2.06	2.13	14.44	1.12	54.22	17.51	69.66	15.63	171.27	171.27	171.27	171.27			
YGR220	0.86	1.23	1.37	1.87	2.09	20.18	4.07	73.14	37.50	93.93	41.57	83.14	83.14	83.14	83.14			
YGR221C	0.94	1.49	1.76	2.09	2.17	14.90	1.44	71.30	21.93	86.20	23.37	122.30	122.30	122.30	122.30			
YGR222W	0.96	1.34	1.43	1.80	2.05	120.10	39.80	171.28	99.23	291.38	139.03	380.74	380.74	380.74	380.74			
YGR223C	0.93	1.50	1.84	2.18	2.30	30.36	3.61	91.71	35.61	122.07	39.22	118.79	118.79	118.79	118.79			
YGR224W	1.02	1.78	2.05	2.23	2.28	28.75	1.74	106.76	21.57	135.51	23.31	141.77	141.77	141.77	141.77			
YGR225W	0.95	1.27	1.44	1.87	1.97	23.84	1.33	94.08	26.11	117.32	27.45	168.78	168.78	168.78	168.78			
YGR226	0.88	1.39	1.71	2.08	2.16	23.83	0.74	92.71	23.04	116.53	23.78	133.68	133.68	133.68	133.68			
YGR227W	0.88	1.30	1.61	1.95	2.07	30.25	0.37	112.33	23.06	142.58	23.43	193.20	193.20	193.20	193.20			
YGR230W	1.03	1.64	1.99	2.35	2.39	32.97	2.39	109.24	23.85	142.25	26.24	201.09	201.09	201.09	201.09			
YGR231C	0.93	2.06	2.22	2.41	2.47	13.07	0.13	65.15	15.64	78.17	15.77	144.01	144.01	144.01	144.01			
YGR232W	0.91	1.50	1.76	2.05	2.14	16.72	2.55	80.40	22.34	97.12	24.90	143.82	143.82	143.82	143.82			
YGR233C	0.97	1.53	1.78	2.10	2.19	26.67	2.40	99.15	25.11	125.82	27.51	148.32	148.32	148.32	148.32			
YGR234W	0.99	1.56	1.74	2.03	2.13	52.68	14.19	123.25	67.41	175.93	81.59	203.59	203.59	203.59	203.59			
YGR235C	0.94	1.63	1.90	2.18	2.24	14.06	0.76	69.36	21.48	83.43	22.24	158.03	158.03	158.03	158.03			
YGR236	0.94	1.51	1.75	2.04	2.12	16.00	0.70	74.06	18.10	90.08	18.80	131.49	131.49	131.49	131.49			
YGR237C	0.94	1.51	1.77	2.10	2.18	18.47	0.48	75.28	15.53	94.75	16.01	199.91	199.91	199.91	199.91			
YGR238C	1.06	1.57	1.87	2.28	2.36	36.28	1.90	129.87	24.86	166.15	26.75	144.95	144.95	144.95	144.95			
YGR239C	0.97	1.58	1.97	2.32	2.35	42.05	1.99	136.19	27.84	178.25	29.83	101.46	101.46	101.46	101.46			
YGR240C	0.97	1.09	1.05	1.04	1.00	188.57	161.05	179.66	151.20	368.23	312.25	510.36	510.36	510.36	510.36			
YGR240C-A	0.93	1.44	1.76	2.16	2.24	27.35	1.76	91.50	26.97	118.86	30.73	118.17	118.17	118.17	118.17			
YGR243W	0.98	1.55	1.86	2.21	2.27	16.72	0.95	72.15	24.33	88.86	25.31	125.43	125.43	125.43	125.43			
YGR244C	1.02	1.67	1.96	2.20	2.32	12.82	2.44	65.52	27.25	78.34	29.69	68.82	68.82	68.82	68.82			
YGR250	1.05	1.18	1.19	1.02	1.19	1.68	16.64	2.03	200.26	18.07	382.90	20.10	527.69	527.69	527.69	527.69		
YGR250W	1.05	1.63	1.93	2.23	2.30	24.74	7.56	94.83	54.61	119.57	62.11	99.56	99.56	99.56	99.56			
YGR253C	1.02	1.69	1.93	2.25	2.35	20.65	6.35	87.44	103.78	109.26	109.26	84.60	84.60	84.60	84.60			
YGR255C	0.98	1.43	1.70	2.08	2.10	22.87	1.63	72.23	21.95	109.50	22.59	85.20	85.20	85.20	85.20			
YGR260C	0.88	1.34	1.54	1.93	2.09	26.26	2.10	109.05	20.68	129.40	31.68	209.60	209.60	209.60	209.60			
YGR262C	0.94	1.54	1.71	2.11	2.15	21.17	1.26	84.68	13.86	128.35	15.16	117.46	117.46	117.46	117.46			
YGR266W	0.99	1.39	1.84	2.22	2.26	45.81	1.80	158.28	28.57	203.88	30.37	220.10	220.10	220.10	220.10			
YGR268C	0.94	1.64	1.94	2.19	2.26	19.98	4.08	83.57	16.51	102.25	31.31	147.62	147.62	147.62	147.62			
YGR271C-A	0.95	1.20	1.36	1.82	2.17	95.95	12.00	172.01	63.29	267.96	76.09	281.73	281.73	281.73	281.73			
YGR271W	1.01	1.63	1.92	2.23	2.31	18.59	0.17	75.31	14.40	93.90	14.57	173.90	173.90	173.90	173.90			
YGR273C	1.02	1.58	1.88	2.24	2.29	27.98	0.20	125.33	16.25	153.31	16.45	101.48	101.48	101.48	101.48			
YGR276	0.97	1.59	1.79	2.05	2.27	19.12	3.05	83.52	26.24	102.64	29.29	143.60	143.60	143.60	143.60			
YGR282C	0.96	1.52	1.80	2.11	2.20	21.29	1.29	78.09	21.73	97.38	22.79	191.81	191.81	191.81	191.81			
YGR286C	0.95	1.53	1.79	2.12	2.21	20.38	0.63	80.93	21.05	101.32	21.68	74.60	74.60	74.60	74.60			
YGR288C	0.93	1.44	1.56	2.08	2.12	16.74	3.69	65.38	28.20	82.12	31.89	54.19	54.19	54.19	54.19			
YGR292W	0.96	1.55	1.85	2.16	2.26	19.44	1.70	77.97	30.21	97.41	31.91	140.57	140.57	140.57	140.57			
YGR294W	0.96	1.06	1.04	1.01	1.00	0.99	208.79	137.02	178.84	131.89	387.63	268.91	511.86	511.86	511.86	511.86		
YGR296W	0.96	1.89	1.90	2.01	2.10	8.33	0.47	56.92	8.49	65.25	8.96	163.40	163.40	163.40	163.40			
YHL006C	0.95	1.23	1.56	1.73	2.00	43.77	5.67	111.59	49.82	155.36	55.49	169.41	169.41	169.41	169.41			
YHL008C	1.00	1.77	2.06	2.23	2.37	15.76	1.45	70.52	13.43	86.28	14.88	109.78	109.78	109.78	109.78			
YHL011C	0.96	1.06	1.03	1.02	0.98	193.03	17.67	161.79	151.48	354.82	32.51	508.25	508.25	508.25	508.25			
YHL014C	0.94	1.74	2.06	2.30	2.41	22.29	1.52	85.39	12.82	107.68	14.35	116.66	116.66	116.66	116.66			
YHL014C	0.96	1.08	1.04	1.04	0.99	197.80	167.57	160.77	153.91	358.37	32.14	511.60	511.60	511.60	511.60			
YHL015W-A	0.94	1.04	1.01	1.01	0.98	177.14	163.44	148.29	144.14	325.43	30.48	509.57	509.57	509.57	509.57			
YHL018W	1.00	1.10	1.07	1.03	1.03	155.81	159.45	140.32	134.91	296.13	29.43	511.32	511.32	511.32	511.32			
YHL020C	0.93	1.07	1.03	0.99	0.96	171.62	162.85	145.33	142.41	316.96	30.52	513.20	513.20	513.20	513.20			
YHL021C	0.96	1.10	1.06	1.02	0.98	192.40	161.90	152.89	140.44	345.29	30.33	512.68	512.68	512.68	512.68			
YHL022C	0.97	1.26	1.47	1.71	1.87	21.21	2.02	82.81	13.47	104.37	13.77	226.73	226.73	226.73	226.73			
YHL023C	0.96	1.32	1.41	1.92	2.08	33.73	3.60	161.81	35.40	135.54	39.00	182.70	182.70	182.70	182.70			
YHL030C	0.89	1.57	2.04	2.09	2.18	51.18	7.37	92.29	51.61	107.47	58.98	53.47	53.47	53.47	53.47			
YHL034C	1.04	1.59	1.90	2.10	2.31	19.55	3.41	85.97	42.25	105.52	45.66	76.02	76.02	76.02	76.02			
YHR006W	0.97	1.46	1.78	2.21	2.26	27.87	0.35	107.06	16.90	134.92	17.25	109.39	109.39	109.39	109.39			
YHR007C-A	1.00	1.58	1.88	2.01	2.22	26.07	3.09	95.61	33.44	121.68	35.53	84.41	84.41	84.41	84.41			
YHR008C	1.02	1.56	1.88	2.04	2.23	29.45	2.08	98.33	29.73	127.78	31.81	92.65	92.65	92.65	92.65			
YHR009C	1.04	1.77	2.01	2.20	2.26	35.96	2.08	105.68	30.74	141.64	33.12	205.42	205.42	205.42	205.42			
YHR009C																		

ORF	Optical density at 600 nm at time intervals (h)							GLUCOSE		FRUCTOSE		TOTAL Sugars(g/L)		TOTAL AMINO ACIDS (mg/L)	
	0	20	27	42	67	43 h	67 h	43 h	67 h	43 h	67 h	43 h	67 h	48 h	
YHR147C	1.05	1.06	1.04	1.02	0.99	164.56	161.66	149.49	152.30	307.23	295.59	51.52	51.52	51.52	
YHR148C	1.04	1.07	1.07	2.23	2.36	1.76	3.52	76.46	37.20	90.41	46.92	116.08	116.08	116.08	
YHR149C	0.94	1.23	1.49	1.98	2.00	34.99	1.51	104.60	33.44	138.19	36.33	279.87	279.87	279.87	
YHR150C	0.94	1.52	1.78	2.09	2.18	14.43	0.75	58.28	20.63	73.31	7.39	214.95	214.95	214.95	
YHR154W	0.94	1.46	1.68	2.02	2.11	16.72	0.90	58.26	22.75	74.98	23.66	294.07	294.07	294.07	
YHR155W	0.94	1.60	2.04	2.14	2.20	15.42	0.79	55.19	16.43	70.51	17.22	308.66	308.66	308.66	
YHR156C	1.00	1.32	1.81	1.88	1.59	38.68	2.32	121.58	31.20	169.26	33.52	116.81	116.81	116.81	
YHR157W	0.93	1.63	1.85	2.15	2.22	14.74	1.07	54.46	22.47	69.20	23.53	347.18	347.18	347.18	
YHR158C	0.97	1.43	1.77	2.25	2.31	36.10	2.09	117.33	27.12	153.23	29.20	135.81	135.81	135.81	
YHR159W	0.98	1.58	1.85	2.17	2.24	17.65	0.45	65.32	20.10	82.97	20.55	127.90	127.90	127.90	
YHR160C	0.95	1.69	1.90	2.19	2.30	17.32	0.40	60.55	14.67	77.87	15.08	246.51	246.51	246.51	
YHR161C	0.93	1.48	1.75	2.09	2.17	22.77	3.38	69.74	31.03	92.51	34.41	274.68	274.68	274.68	
YHR163W	0.94	1.59	2.10	2.04	2.15	20.20	2.23	69.54	29.83	89.74	32.06	241.13	241.13	241.13	
YHR167W	0.93	1.34	1.60	1.95	2.08	34.09	7.26	81.02	49.73	115.11	56.99	212.18	212.18	212.18	
YHR168W	0.93	1.81	2.02	2.21	2.27	16.32	0.09	65.67	15.07	82.00	15.15	147.60	147.60	147.60	
YHR175W-A	0.94	1.61	1.88	2.17	2.22	13.07	0.45	58.84	16.54	71.94	16.99	257.95	257.95	257.95	
YHR176W	0.98	1.66	1.89	2.16	2.24	15.20	1.03	59.95	19.88	75.16	20.91	203.65	203.65	203.65	
YHR178W	0.96	1.63	1.90	2.22	2.28	30.05	-0.09	124.57	11.89	154.62	11.80	116.64	116.64	116.64	
YHR179W	0.94	1.61	1.79	2.08	2.21	19.02	1.67	61.66	26.22	80.68	27.89	285.84	285.84	285.84	
YHR181W	0.93	1.22	1.56	2.08	2.18	38.73	2.54	129.34	39.24	168.07	41.78	275.99	275.99	275.99	
YHR185C	0.94	1.48	1.71	2.03	2.12	14.46	2.16	51.79	21.19	66.25	23.35	212.95	212.95	212.95	
YHR189W	0.92	1.49	1.72	2.03	2.13	16.19	1.10	61.13	16.31	77.32	17.40	260.89	260.89	260.89	
YHR191C	0.87	1.60	1.82	2.07	2.13	10.33	1.80	52.11	22.49	62.44	24.29	241.38	241.38	241.38	
YHR193C	0.93	1.47	1.70	2.04	2.14	18.13	1.56	71.00	17.08	89.13	18.58	199.51	199.51	199.51	
YHR194W	0.93	1.52	1.73	2.06	2.17	34.25	12.49	83.45	60.17	117.70	72.66	346.82	346.82	346.82	
YHR195W	0.98	1.58	1.81	2.13	2.21	20.93	1.16	75.07	20.11	95.01	21.26	220.91	220.91	220.91	
YHR198C	0.94	1.48	1.71	2.04	2.15	21.19	0.94	73.29	20.80	94.48	21.74	279.30	279.30	279.30	
YHR199C	0.90	1.38	1.59	1.90	2.03	23.79	1.01	78.55	19.97	102.35	20.98	239.29	239.29	239.29	
YHR202W	0.90	1.49	1.75	2.07	2.18	21.90	2.41	87.55	33.59	100.46	36.00	183.82	183.82	183.82	
YHR203C	0.95	1.28	1.55	2.01	2.14	22.36	0.39	82.81	18.93	106.19	19.32	240.56	240.56	240.56	
YHR204W	0.90	1.44	1.68	1.93	2.06	21.63	0.90	64.91	20.60	81.34	20.65	204.54	204.54	204.54	
YHR205W	1.00	1.67	1.99	2.04	2.23	21.63	0.32	97.21	25.64	111.3	28.64	119.13	119.13	119.13	
YHR210C	0.93	1.59	1.83	2.17	2.26	16.45	-0.19	72.92	8.84	85.67	8.65	314.61	314.61	314.61	
YHR210W	0.93	1.67	1.80	2.21	2.28	16.38	1.67	73.23	24.28	106.68	24.75	242.80	242.80	242.80	
YHR214W-A	0.98	1.59	1.88	2.14	2.27	23.92	0.14	103.69	19.00	127.61	19.14	253.16	253.16	253.16	
YHR219W	0.94	1.54	1.79	2.10	2.17	17.17	1.64	71.00	19.00	88.17	20.64	194.84	194.84	194.84	
YLR023W-A	0.92	1.62	1.91	2.20	2.26	20.28	1.82	77.07	21.84	97.35	23.65	222.10	222.10	222.10	
YLR005W	0.94	1.56	1.89	2.29	2.17	17.57	2.00	68.43	28.47	85.00	30.47	126.03	126.03	126.03	
YLR006W	0.91	1.48	1.76	2.07	2.16	17.97	0.72	72.59	18.34	90.55	19.06	309.83	309.83	309.83	
YLR007C	0.93	1.61	1.85	2.16	2.21	17.56	0.11	62.77	16.13	74.33	16.24	154.70	154.70	154.70	
YLR008W	0.91	1.40	1.65	1.98	2.10	2.08	1.03	69.31	26.01	89.38	27.04	165.06	165.06	165.06	
YLR009C	0.93	1.55	1.84	2.04	2.13	14.46	-0.11	53.40	12.41	66.86	12.30	249.05	249.05	249.05	
YLR009W	0.94	1.54	1.82	2.15	2.22	15.63	0.32	73.82	13.49	89.44	13.81	217.65	217.65	217.65	
YLR010W	0.93	1.78	2.09	2.22	2.29	8.99	0.11	49.80	7.29	58.79	7.40	352.74	352.74	352.74	
YLR011W	0.91	1.22	1.54	2.06	2.14	46.53	4.85	143.16	44.56	189.59	49.41	134.74	134.74	134.74	
YLR014C-A	0.91	1.36	1.65	2.06	2.11	24.29	1.18	93.53	19.99	117.82	21.17	110.85	110.85	110.85	
YLR015W	0.90	1.52	1.79	2.10	2.17	19.13	1.80	74.37	23.83	93.50	25.63	227.60	227.60	227.60	
YLR016W	0.95	1.60	1.86	2.14	2.21	16.83	1.93	72.18	19.58	89.01	21.50	205.63	205.63	205.63	
YLR017C	0.92	1.42	1.65	1.93	2.02	19.76	1.75	72.03	19.76	89.21	21.52	233.62	233.62	233.62	
YLR018W	1.02	1.69	1.95	2.16	2.29	25.28	5.42	98.44	19.84	121.72	24.44	44.96	44.96	44.96	
YLR020C	0.93	1.55	1.80	2.15	2.23	20.08	1.37	76.13	21.95	98.21	23.32	210.84	210.84	210.84	
YLR023C	1.01	1.44	1.74	2.16	2.23	31.68	0.73	116.98	21.93	148.67	22.66	119.17	119.17	119.17	
YLR024C	0.95	1.53	1.89	2.27	2.31	34.14	1.18	120.50	29.56	154.63	30.74	170.55	170.55	170.55	
YLR027C	0.93	1.74	1.96	2.18	2.24	10.06	0.25	56.07	11.89	66.13	12.14	379.41	379.41	379.41	
YLR029C	0.91	1.55	1.86	2.05	2.20	17.93	0.73	61.49	16.39	73.41	17.60	51.56	51.56	51.56	
YLR030C	0.92	1.52	1.82	2.08	2.16	17.03	0.03	61.44	16.34	71.97	16.97	88.65	88.65	88.65	
YLR034C	0.94	1.45	1.74	2.08	2.14	18.00	0.63	63.54	15.44	75.55	15.44	276.95	276.95	276.95	
YLR044C	0.92	1.50	1.86	2.05	2.05	12.42	0.24	65.56	12.01	77.97	12.23	210.74	210.74	210.74	
YLR049W	0.98	1.58	1.80	2.10	2.17	27.31	3.75	97.81	35.95	125.01	35.97	161.53	161.53	161.53	
YLR050W	0.97	1.00	1.02	1.96	2.06	34.78	4.03	119.76	15.12	143.03	36.88	524.68	524.68	524.68	
YLR055C	0.91	1.56	1.80	2.08	2.14	8.47	0.25	34.40	5.26	42.87	5.51	271.19	271.19	271.19	
YLR056W	0.96	1.42	1.58	1.93	2.13	29.42	7.24	111.80	46.62	141.22	55.85	110.21	110.21	110.21	
YLR064W	0.91	1.44	1.68	2.00	2.09	16.17	0.70	65.03	18.70	81.20	19.39	211.61	211.61	211.61	
YLR069C	0.95	1.47	1.78	2.16	2.27	17.14	2.42	75.72	31.61	101.86	34.03	111.31	111.31	111.31	
YLR070C	0.91	1.34	1.56	2.05	2.05	21.67	1.07	77.97	24.16	99.64	25.23	252.23	252.23	252.23	
YLR071C	0.89	1.53	1.81	2.10	2.19	17.43	0.98	70.77	26.99	88.20	27.97	97.49	97.49	97.49	
YLR073C	1.02	1.49	1.68	2.02	2.20	31.80	4.99	110.67	44.30	142.47	49.28	137.96	137.96	137.96	
YLR076W	0.94	1.54	1.84	2.14	2.25	12.75	5.89	59.20	20.33	71.95	26.22	85.24	85.24	85.24	
YLR077C	0.96	1.49	1.78	2.											

ORF	Optical density at 600 nm at time intervals (h)							GLUCOSE		FRUCTOSE		TOTAL Sugars(g/L)		TOTAL AMINO ACIDS (mM/L)	
	0	20	27	42	67	43 h	67 h	43 h	67 h	43 h	67 h	43 h	67 h	48 h	
YJL051W	1.37	1.73	2.03	2.32	2.36	18.30	11.2	81.24	24.42	102.44	24.44	116.35	24.44	116.35	
YJL053W	1.10	1.41	1.78	2.36	2.45	18.12	0.6	64.16	14.42	82.92	14.58	207.38	14.58	207.38	
YJL054W	1.01	1.63	1.93	2.27	2.31	16.72	1.63	80.79	23.17	97.51	24.80	140.33	24.80	140.33	
YJL062W	1.03	1.44	1.78	2.26	2.31	16.55	0.66	78.94	18.16	96.20	18.81	202.47	18.81	202.47	
YJL070C	1.03	1.54	1.84	2.18	2.22	16.39	1.00	81.37	17.19	97.76	18.19	155.20	18.19	155.20	
YJL079C	0.98	1.65	1.95	2.28	2.34	16.40	1.03	76.71	20.72	93.11	21.76	213.50	21.76	213.50	
YJL083W	1.05	1.54	1.84	2.18	2.23	18.65	1.15	81.22	20.56	98.88	21.71	249.82	21.71	249.82	
YJL095W	1.13	1.64	1.94	2.32	2.40	21.82	1.83	91.04	22.50	112.86	24.33	305.04	24.33	305.04	
YJL098W	1.06	1.58	1.92	2.32	2.35	23.34	2.05	90.27	24.42	113.51	25.47	153.69	25.47	153.69	
YJL099W	1.00	1.65	1.99	2.29	2.39	31.84	2.78	110.72	26.11	142.56	28.89	53.51	28.89	53.51	
YJL100W	0.96	1.41	1.98	2.20	2.27	22.07	1.55	88.06	23.80	108.12	25.35	216.41	25.35	216.41	
YJL108C	0.96	1.34	1.69	2.14	2.20	24.15	1.62	96.13	26.58	120.28	28.20	166.80	28.20	166.80	
YJL110C	1.06	1.75	2.02	2.34	2.37	15.91	1.60	77.97	21.75	93.88	23.35	98.65	23.35	98.65	
YJL134W	1.02	1.79	2.08	2.35	2.36	12.29	1.04	71.41	21.86	83.70	22.90	103.32	22.90	103.32	
YJL139C	1.06	1.68	2.01	2.34	2.36	16.39	0.68	80.02	19.50	98.41	20.18	101.48	20.18	101.48	
YJL142C	0.96	1.87	2.08	2.34	2.34	6.08	0.24	52.52	6.10	58.61	6.34	129.16	6.34	129.16	
YJL146W	0.97	1.64	1.95	2.28	2.29	13.31	0.66	71.38	14.82	84.70	15.47	259.00	15.47	259.00	
YJL157C	0.94	1.69	1.97	2.28	2.35	11.85	0.56	66.29	14.53	78.13	15.09	103.75	15.09	103.75	
YJL159W	0.99	1.69	1.99	2.34	2.35	10.47	0.19	62.66	9.38	73.13	9.54	111.95	9.54	111.95	
YJL165C	0.97	1.89	2.10	2.31	2.33	9.55	0.08	58.12	8.31	67.67	8.38	66.42	8.38	66.42	
YJL175W	1.02	1.30	1.51	1.81	1.91	46.51	13.11	123.22	61.26	169.73	74.37	224.04	74.37	224.04	
YJL187C	1.01	1.46	1.71	2.14	2.19	22.15	2.65	90.98	29.58	113.13	32.23	55.50	32.23	55.50	
YJL191W	0.96	1.32	1.62	2.07	2.15	25.13	2.29	93.61	27.21	118.74	29.50	236.88	29.50	236.88	
YJL201W	0.93	1.42	1.79	2.21	2.27	23.14	2.30	83.96	27.35	107.12	29.65	104.94	29.65	104.94	
YJL206C	0.98	1.50	1.79	2.24	2.24	15.83	1.85	75.84	27.59	91.67	29.44	112.41	29.44	112.41	
YJL210W	1.02	1.65	1.94	2.30	2.34	14.50	1.58	73.90	22.86	88.40	24.44	63.89	24.44	63.89	
YJL214W	1.01	1.72	1.99	2.31	2.35	12.27	1.37	71.97	20.33	84.24	21.70	108.10	21.70	108.10	
YJR001W	1.05	1.64	1.98	2.30	2.31	17.03	1.21	84.27	16.33	101.32	17.59	69.32	17.59	69.32	
YJR009C	0.93	1.41	1.60	1.94	1.99	10.88	2.18	65.64	21.64	75.44	23.82	131.32	23.82	131.32	
YJR019C	0.95	1.67	1.95	2.25	2.29	16.16	1.59	83.25	24.46	99.41	26.05	191.52	26.05	191.52	
YJR025C	0.99	1.55	1.77	2.04	2.13	30.82	0.16	88.02	18.70	118.70	18.43	155.37	18.43	155.37	
YJR030C	1.07	1.43	1.62	1.84	1.91	13.66	0.62	76.67	17.87	93.34	18.14	233.03	18.14	233.03	
YJR032W	1.03	1.44	1.62	1.83	1.87	62.88	16.59	132.65	63.15	185.63	78.65	432.24	78.65	432.24	
YJR035C	0.98	1.31	1.59	1.74	1.84	18.73	15.69	121.05	15.58	165.63	18.25	213.68	18.25	213.68	
YJR039W	0.94	1.31	1.63	2.04	2.16	33.16	3.17	70.78	23.68	123.94	26.85	142.24	26.85	142.24	
YJR039W	1.00	1.14	1.18	1.85	1.92	180.43	17.35	137.49	14.67	318.33	31.42	520.88	31.42	520.88	
YJR036C	1.07	1.74	2.09	2.27	2.35	21.46	1.20	69.69	21.43	91.15	22.63	194.30	22.63	194.30	
YJR040W	1.09	1.69	1.96	2.23	2.24	22.48	0.71	67.43	21.75	89.91	22.46	186.12	22.46	186.12	
YJR043C	0.92	1.43	1.72	2.05	2.16	46.43	9.49	97.71	53.02	137.15	62.51	181.70	62.51	181.70	
YJR044C	0.98	1.60	1.87	2.14	2.28	23.05	0.17	88.19	15.10	111.25	15.27	176.10	15.27	176.10	
YJR051W	1.02	1.44	1.72	2.17	2.26	20.61	0.19	85.99	15.80	107.61	16.00	110.25	16.00	110.25	
YJR053W	0.95	1.57	1.90	2.26	2.30	20.63	1.20	91.74	23.67	111.77	24.88	86.41	24.88	86.41	
YJR056C	0.97	1.51	1.86	2.26	2.36	23.88	0.48	79.20	19.49	105.34	19.97	28.94	19.97	28.94	
YJR058C	0.91	1.22	1.51	2.01	2.09	30.58	2.52	112.97	29.72	143.53	32.24	107.52	32.24	107.52	
YJR069W	1.06	1.74	2.02	2.32	2.35	21.46	1.20	69.69	21.43	91.15	22.63	194.30	22.63	194.30	
YJR061W	0.93	1.47	1.75	2.04	2.22	17.82	2.19	61.06	26.19	78.86	28.37	176.15	28.37	176.15	
YJR062C	0.95	1.56	1.91	2.22	2.31	24.26	0.68	80.64	17.40	104.90	18.06	29.55	18.06	29.55	
YJR069C	0.97	1.60	1.92	2.21	2.32	33.37	2.71	91.48	32.44	124.85	35.15	183.99	35.15	183.99	
YJR070C	0.87	1.35	1.69	2.09	2.23	23.05	1.74	73.64	25.70	95.69	27.44	162.38	27.44	162.38	
YJR073C	1.11	1.87	1.99	2.21	2.28	22.49	-0.25	78.41	7.27	100.90	7.03	160.05	7.03	160.05	
YJR075W	0.98	1.42	1.63	2.03	2.09	18.70	1.23	81.66	28.25	100.36	29.48	116.80	29.48	116.80	
YJR079W	1.05	1.74	1.97	2.27	2.32	19.62	0.59	68.22	17.44	87.84	18.03	115.23	18.03	115.23	
YJR080C	1.05	1.74	1.96	2.21	2.31	16.10	1.50	61.72	23.33	77.82	24.83	163.62	24.83	163.62	
YJR084W	1.06	1.52	1.79	2.17	2.21	10.67	0.68	68.31	15.00	78.98	15.67	88.59	15.67	88.59	
YJR090W	1.03	1.48	1.74	2.33	2.36	15.51	1.72	82.10	21.28	97.62	23.01	148.84	23.01	148.84	
YJR094C	1.02	1.56	1.72	2.06	2.13	15.04	1.26	85.64	28.75	108.10	30.10	163.68	30.10	163.68	
YJR109W	1.06	1.58	1.82	2.29	2.36	17.99	1.19	90.83	21.96	108.82	23.07	160.39	23.07	160.39	
YJR111C	0.88	1.35	1.66	2.01	2.15	25.01	1.02	82.51	27.44	107.52	28.45	168.67	28.45	168.67	
YJR115W	1.04	1.66	1.87	2.07	2.33	33.27	0.00	95.11	12.37	128.38	12.37	270.11	12.37	270.11	
YJR116W	0.98	1.63	1.87	2.12	2.21	14.95	1.52	62.69	22.88	77.64	24.40	321.43	24.40	321.43	
YJR118C	1.01	1.14	1.11	1.05	1.02	150.85	182.31	136.49	154.29	287.34	336.60	525.95	336.60	525.95	
YJR119C	1.02	1.77	2.29	2.30	2.37	16.45	1.73	67.58	21.10	84.03	22.63	190.18	22.63	190.18	
YJR120W	1.03	1.02	1.03	0.93	0.93	14.33	1.44	77.50	13.46	102.28	13.61	311.72	13.61	311.72	
YJR124C	1.02	1.81	2.06	2.25	2.38	16.45	2.71	65.64	25.75	82.09	28.45	225.72	28.45	225.72	
YJR125C	1.02	1.79	2.29	2.32	2.37	17.32	2.03	74.45	22.34	102.58	24.59	162.62	24.59	162.62	
YJR129C	0.99	1.67	1.98	2.26	2.38	22.27	1.92	80.26	24.34	102.53	25.25	179.76	25.25	179.76	
YJR130C	0.98	1.62	1.96	2.24	2.36	16.88	1.42	64.56	23.58	81.24	25.00	92.90	25.00	92.90	
YJR133W	1.03	1.67	1.86	2.13	2.22	19.93	0.14	76.16	12.33	96.09	13.38	98.70	13.38	98.70	
YJR134C	1.01	1.58	1.82	2.10	2.21	21.84	2.07	82.81							

ORF	Optical density at 600 nm at time intervals (h)										GLUCOSE		FRUCTOSE		TOTAL Sugars(g/L)		TOTAL AMINO ACIDS (mM/L)	
	0	20	27	42	67	43 h	67 h	43 h	67 h	43 h	0	20	67	0	20	48 h		
YKL120W	0.93	1.73	1.99	2.05	2.31	21.00	0.37	80.7	78.0	101.73	1.26	1.26	1.26	1.26	1.26	2.51		
YKL121W	0.96	1.92	2.19	2.38	2.45	18.44	1.25	13.49	15.68	11.93	1.42	1.42	1.42	1.42	1.42	32.13		
YKL122W	1.08	1.72	1.81	1.20	1.07	1.13	195.66	160.20	117.48	152.53	107.5	320.73	519.83					
YKL123W	1.00	1.63	1.68	2.17	2.27	25.79	1.20	68.03	14.44	111.82	15.17	209.60						
YKL128C	1.05	1.42	1.59	1.89	2.02	19.81	1.59	71.30	23.31	90.9	24.90	109.06						
YKL132C	1.00	1.83	2.13	2.37	2.43	21.75	3.14	77.38	21.52	99.13	24.66	85.86						
YKL133C	1.02	1.16	1.15	1.98	1.07	209.88	161.60	162.93	143.13	372.51	304.73	528.01						
YKL134C	1.03	1.58	1.88	2.20	2.30	27.57	0.57	92.12	18.01	119.69	19.58	174.19						
YKL135C	1.00	1.35	1.13	1.07	1.04	182.29	191.50	164.54	163.82	346.74	355.33	519.93						
YKL137W	1.04	1.76	1.90	2.15	2.24	13.71	0.58	69.67	20.87	83.38	21.45	144.16						
YKL138C	0.99	1.42	1.61	1.91	2.12	88.98	29.20	138.08	74.25	227.07	103.44	277.50						
YKL139C	0.95	1.09	1.06	0.98	0.97	135.12	133.88	164.33	126.41	299.45	260.29	526.17						
YKL140W	0.99	1.74	1.96	2.21	2.32	19.00	2.23	75.37	24.84	94.37	27.07	232.15						
YKL146W	0.97	2.27	2.36	1.93	2.04	18.36	1.80	76.78	28.50	95.14	30.31	160.47						
YKL148C	1.01	1.90	2.15	2.21	2.31	16.39	0.72	74.34	19.13	90.73	19.85	161.99						
YKL150C	1.04	1.67	1.96	2.19	2.34	19.34	2.06	92.39	21.43	111.75	23.50	70.87						
YKL161C	1.06	1.81	2.09	2.30	2.40	23.64	2.48	94.37	23.71	118.01	26.19	262.23						
YKL162C	0.93	1.47	1.73	1.88	1.99	18.09	0.61	73.74	17.53	91.82	18.14	133.37						
YKL164C	1.00	1.19	1.37	1.73	1.76	12.29	0.88	65.60	20.13	79.19	21.01	120.25						
YKL166C	0.95	1.69	1.97	2.19	2.29	22.39	0.73	80.40	22.90	102.79	23.63	137.99						
YKL167C	0.94	1.56	1.85	2.02	2.13	22.57	3.28	80.40	32.56	102.97	35.84	191.47						
YKL168C	1.01	1.49	1.70	2.04	2.10	14.93	0.96	77.10	18.73	92.03	19.69	72.63						
YKL171W	0.94	1.51	1.72	2.09	2.14	18.52	1.35	91.74	21.52	110.26	22.86	177.85						
YKL174C	0.97	1.64	1.95	2.24	2.35	22.40	3.06	79.03	32.06	101.43	35.12	165.10						
YKL175W	0.96	1.51	1.81	2.25	2.32	25.91	3.05	95.31	34.45	122.22	37.49	54.93						
YKL176C	1.00	1.75	2.07	2.31	2.42	33.45	7.20	91.77	45.76	125.22	52.96	171.30						
YKL178C	1.02	1.54	1.86	2.17	2.26	26.63	3.31	88.40	30.51	113.03	33.82	130.17						
YKL188C	0.96	1.52	1.88	2.19	2.31	23.34	1.45	83.10	23.11	106.48	24.56	145.11						
YKL197C	0.96	1.55	1.94	2.20	2.39	23.97	3.31	90.84	34.93	120.12	38.25	163.93						
YKL206C	1.04	1.67	1.93	2.16	2.29	23.95	2.14	88.13	26.75	112.08	26.88	179.18						
YKL210W	0.91	1.72	1.99	2.03	2.10	2.50	0.87	78.51	18.24	98.70	19.65	176.05						
YKL212C	1.00	1.75	2.01	2.03	2.13	16.59	0.95	76.5	8.12	87.61	8.21	312.08						
YKL214C	0.98	1.68	1.83	2.19	2.28	22.68	2.04	66.14	18.93	101.59	19.87	158.45						
YKL215C	0.99	1.72	2.01	2.26	2.38	24.03	2.38	73.71	28.41	105.74	21.84	139.16						
YKL216W	1.02	1.07	1.06	1.13	1.19	169.06	199.00	185.13	173.86	355.18	363.86	511.70						
YKL218C	0.95	1.72	2.02	2.27	2.36	18.17	0.87	77.24	18.39	95.41	19.26	154.41						
YKL220C	0.99	1.86	2.23	2.30	2.38	13.03	2.27	85.72	23.22	99.74	25.49	205.72						
YKL221W	0.94	1.60	1.57	1.91	2.01	20.08	0.96	78.71	20.06	98.79	21.02	132.42						
YKL222C	0.91	1.56	1.85	2.23	2.29	19.66	1.33	90.01	29.58	109.67	30.91	144.21						
YKL0001C	0.98	1.50	1.68	2.00	2.18	20.45	4.67	73.64	36.55	94.08	41.22	132.82						
YKL0002C	0.95	1.65	1.93	2.21	2.28	22.58	0.67	79.79	16.31	102.37	16.98	170.55						
YKL0006W	0.96	1.62	1.95	2.21	2.32	24.18	1.67	83.78	22.41	107.96	24.08	154.51						
YKL0007W	0.96	1.53	1.83	2.19	2.28	47.59	18.99	112.00	67.19	159.59	86.18	237.45						
YKL0009C	0.91	1.53	1.88	2.20	2.29	23.47	2.42	83.20	31.39	105.67	33.81	172.95						
YKL0010C	0.91	1.07	1.05	0.88	0.98	154.80	140.99	179.22	122.60	334.02	263.59	517.58						
YKL0011C	0.98	1.63	1.80	2.07	2.15	17.14	0.86	78.33	17.96	95.47	18.82	162.98						
YKL0013W	1.00	1.73	2.00	2.23	2.31	18.39	0.70	73.82	15.86	92.21	16.56	175.48						
YKL0015C	0.96	1.60	1.86	2.13	2.21	15.47	1.62	72.33	22.05	87.80	23.68	185.71						
YKL0016C	0.90	1.43	1.72	2.09	2.20	20.79	2.90	78.63	23.56	102.42	22.46	234.39						
YKL0017W	0.91	1.54	1.72	2.21	2.32	23.20	1.83	84.43	22.22	111.63	31.05	151.22						
YKL0018W	0.91	1.75	2.05	2.33	2.33	21.83	0.78	193.00	18.06	124.53	18.81	173.38						
YKL004W	0.99	1.61	1.90	2.24	2.28	27.80	7.37	114.98	52.59	141.69	59.96	106.76						
YKL008W	0.85	0.98	0.95	0.92	0.87	164.84	154.64	185.20	141.41	350.04	296.05	532.55						
YKL0082W	1.00	1.94	1.84	2.31	2.23	15.64	1.95	72.23	19.99	87.87	21.94	82.95						
YKL004W	0.95	1.62	1.82	2.14	2.30	21.13	4.94	80.55	37.33	101.69	42.27	150.44						
YKL005W	0.98	1.68	1.86	2.05	2.26	23.84	0.30	73.46	23.85	102.54	24.84	166.32						
YKL006W	0.98	1.49	1.75	2.17	2.27	17.44	0.44	75.77	17.36	93.76	20.56	209.50						
YKL007W	0.99	1.76	1.89	2.14	2.22	17.19	1.24	76.57	19.32	93.76	20.56	209.50						
YKL007C	0.99	1.76	1.89	2.14	2.22	17.19	1.24	76.57	19.32	93.76	20.56	209.50						
YKL0072C	0.94	1.56	1.91	2.32	2.35	19.46	2.51	83.66	36.51	103.11	36.02	6.36						
YKL0076W	0.94	1.60	1.96	2.29	2.38	20.25	1.22	80.66	25.43	101.11	27.34	111.85						
YKL0077C	0.95	1.49	1.75	2.24	2.30	21.09	1.56	91.04	31.90	112.33	33.46	100.30						
YKL0078W	0.95	1.57	1.84	2.20	2.26	16.07	1.67	78.16	16.31	82.53	19.95	96.48						
YKL0079W	0.95	1.57	1.84	2.20	2.26	16.07	1.67	78.16	16.31	82.53	19.95	96.48						
YKL0080W	0.95	1.38	1.62	2.17	2.14	29.02	1.26	101.85	32.92	130.87	35.69	188.42						
YKL0083C	0.99	1.75	1.89	2.12	2.19	15.58	0.52	84.28	15.44	99.86	15.96	179.30						
YKL0093C	1.00	1.40	1.69	2.00	2.07	17.10	0.17	95.56	26.29	116.65	26.46	71.76						
YKL0045C	1.00	1.52	1.82	2.18	2.30	37.22	2.53	110.13	30.98	147.35	33.51	210.27						
YKL0046C	0.99	1.70</																

ORF	Optical density at 600 nm at time intervals (h)										GLUCOSE		FRUCTOSE		TOTAL Sugars(g/L)		TOTAL AMINO ACIDS (mg/L)	
	0	20	27	42	67	43 h	67 h	43 h	67 h	43 h	67 h	48 h	67 h	67 h	43 h	67 h		
YLR072W	0.49	1.42	1.75	2.14	3.74	1.56	1.56	108.74	26.53	188.52	26.53	16.24	26.53	16.24	16.24	16.24	16.24	
YLR073C	0.68	1.41	1.78	2.09	2.21	2.57	9.6	100.56	25.25	188.56	23.19	16.57	23.19	16.57	16.57	16.57	16.57	
YLR074W	0.91	1.43	1.76	2.18	2.23	1.16	3.44	82.28	27.79	95.41	21.23	58.91	21.23	58.91	58.91	58.91	58.91	
YLR084C	0.91	1.38	1.70	2.14	2.19	1.74	2.61	1.35	34.88	91.01	37.48	134.81	37.48	134.81	134.81	134.81	134.81	
YLR088S	0.89	0.93	0.91	0.97	0.90	168.29	227.07	170.67	174.28	339.16	491.35	520.04	491.35	520.04	520.04	520.04	520.04	
YLR090W	1.01	1.74	2.04	2.24	2.35	16.78	0.49	83.04	16.29	98.82	16.59	16.43	16.59	16.43	16.43	16.43	16.43	
YLR095C	0.99	1.72	2.00	2.22	2.31	12.67	-0.07	77.12	11.03	89.78	10.95	206.35	10.95	206.35	206.35	206.35	206.35	
YLR096W	0.87	1.15	1.47	1.57	2.06	33.27	2.27	109.19	31.00	142.46	33.26	177.15	33.26	177.15	177.15	177.15	177.15	
YLR097C	0.95	1.53	1.81	1.93	2.00	23.46	1.00	103.91	19.61	127.37	20.61	182.41	20.61	182.41	182.41	182.41	182.41	
YLR121C	0.94	1.43	1.78	2.22	2.30	26.82	6.18	99.03	51.38	125.85	57.56	201.57	57.56	201.57	201.57	201.57	201.57	
YLR128W	1.01	1.59	1.69	1.99	2.22	16.96	1.15	81.55	26.69	98.51	27.84	127.64	27.84	127.64	127.64	127.64	127.64	
YLR130C	1.02	1.79	1.98	2.22	2.28	15.03	1.36	66.38	28.50	81.41	28.87	97.77	28.87	97.77	97.77	97.77	97.77	
YLR144C	0.98	1.64	1.92	2.16	2.26	23.63	0.41	104.33	15.84	127.96	16.25	229.82	16.25	229.82	229.82	229.82	229.82	
YLR149C	0.99	1.65	1.96	2.20	2.31	20.35	0.78	87.64	17.89	108.00	18.67	216.92	18.67	216.92	216.92	216.92	216.92	
YLR150W	0.98	1.73	2.05	2.28	2.37	22.87	2.14	91.99	28.30	114.86	30.45	180.41	30.45	180.41	180.41	180.41	180.41	
YLR151C	0.99	1.60	1.93	2.18	2.33	24.74	1.98	92.64	29.72	117.43	31.70	183.03	31.70	183.03	183.03	183.03	183.03	
YLR152C	0.97	1.70	1.99	2.20	2.36	21.92	1.51	92.80	24.77	114.72	26.27	183.28	26.27	183.28	183.28	183.28	183.28	
YLR164W	0.97	1.70	2.02	2.26	2.36	27.08	1.30	118.44	14.81	145.52	16.11	174.62	16.11	174.62	174.62	174.62	174.62	
YLR165C	0.93	1.49	1.90	2.07	2.16	31.06	1.94	119.47	19.10	150.53	21.04	186.89	21.04	186.89	186.89	186.89	186.89	
YLR170C	1.02	1.67	1.92	2.19	2.29	22.71	2.22	102.68	20.83	125.39	23.05	215.79	23.05	215.79	215.79	215.79	215.79	
YLR172C	1.01	1.68	2.13	2.08	2.26	40.82	2.70	120.79	25.58	173.61	28.28	179.73	28.28	179.73	179.73	179.73	179.73	
YLR173W	0.99	1.72	2.02	2.27	2.34	29.90	2.35	117.61	18.07	147.51	20.42	177.15	20.42	177.15	177.15	177.15	177.15	
YLR174W	0.95	1.54	1.83	2.14	2.24	33.58	2.51	111.35	21.12	144.93	23.63	174.60	23.63	174.60	174.60	174.60	174.60	
YLR178C	1.06	1.78	2.09	2.35	2.42	41.67	2.96	126.47	21.09	168.14	24.05	226.31	24.05	226.31	226.31	226.31	226.31	
YLR179C	0.96	1.11	1.08	1.03	1.01	254.94	1.53	153.29	19.85	176.44	452.45	329.74	452.45	329.74	329.74	329.74	329.74	
YLR181C	1.03	1.66	1.94	2.12	2.21	26.10	3.11	94.99	21.26	121.09	24.37	124.72	24.37	124.72	124.72	124.72	124.72	
YLR185W	0.98	1.27	1.45	1.93	2.23	64.71	10.93	143.09	48.88	207.81	59.81	132.20	59.81	132.20	132.20	132.20	132.20	
YLR187W	1.01	1.70	1.91	2.18	2.28	17.47	0.42	76.73	15.84	93.60	16.25	235.49	16.25	235.49	235.49	235.49	235.49	
YLR188W	0.97	1.57	1.84	2.15	2.24	39.18	2.86	121.20	23.77	160.38	26.63	102.13	26.63	102.13	102.13	102.13	102.13	
YLR189C	0.99	1.59	1.89	2.10	2.30	44.95	3.49	142.05	28.95	187.00	32.44	222.09	32.44	222.09	222.09	222.09	222.09	
YLR191W	1.02	1.76	1.86	2.18	2.25	36.07	1.48	118.76	16.59	155.37	18.40	213.49	18.40	213.49	213.49	213.49	213.49	
YLR193C	1.06	1.71	1.97	2.22	2.31	6.07	2.32	103.84	20.24	170.84	20.65	194.65	20.65	194.65	194.65	194.65	194.65	
YLR195C	1.02	1.71	1.74	2.06	2.16	29.53	1.53	100.22	27.71	125.10	30.55	216.41	30.55	216.41	216.41	216.41	216.41	
YLR196W	1.02	1.68	1.89	2.11	2.24	32.68	2.06	126.16	38.11	207.28	32.63	216.41	32.63	216.41	216.41	216.41	216.41	
YLR197C	1.02	1.63	1.82	2.06	2.24	32.68	2.06	126.16	38.11	207.28	32.63	216.41	32.63	216.41	216.41	216.41	216.41	
YLR198C	0.95	1.65	1.82	2.18	2.27	37.57	2.46	126.79	22.62	184.36	25.08	276.22	25.08	276.22	276.22	276.22	276.22	
YLR202C	0.97	1.69	1.97	2.25	2.33	1.84	2.99	121.84	16.88	163.48	20.67	199.99	20.67	199.99	199.99	199.99	199.99	
YLR207W	1.05	1.79	2.02	2.37	2.37	22.95	2.59	89.52	28.86	112.47	31.45	178.70	31.45	178.70	178.70	178.70	178.70	
YLR211C	1.02	1.78	2.08	2.39	2.40	25.72	2.62	97.08	26.67	122.81	28.29	191.45	28.29	191.45	191.45	191.45	191.45	
YLR213C	1.00	1.72	2.03	2.29	2.37	1.30	3.54	121.36	31.48	156.75	35.02	154.79	35.02	154.79	154.79	154.79	154.79	
YLR214W	1.00	1.80	1.72	2.07	2.23	19.56	1.75	73.93	26.52	93.49	30.27	138.23	30.27	138.23	138.23	138.23	138.23	
YLR218C	0.98	1.67	2.01	2.30	2.38	33.73	2.88	125.04	25.03	158.77	27.91	179.45	27.91	179.45	179.45	179.45	179.45	
YLR219W	0.94	1.50	1.79	2.09	2.20	54.50	2.72	125.44	25.15	229.94	27.87	211.29	27.87	211.29	211.29	211.29	211.29	
YLR220W	0.99	1.64	1.98	2.25	2.35	53.04	3.25	167.57	26.88	220.81	26.93	180.33	26.93	180.33	180.33	180.33	180.33	
YLR221C	0.99	1.90	1.71	2.08	2.17	38.10	1.84	136.53	37.47	174.74	42.39	154.20	42.39	154.20	154.20	154.20	154.20	
YLR224W	1.02	1.68	1.94	2.23	2.30	34.13	2.02	116.55	20.37	150.88	20.67	206.77	20.67	206.77	206.77	206.77	206.77	
YLR225C	0.98	1.56	1.84	2.14	2.22	43.18	2.98	127.82	26.88	170.99	29.66	202.14	29.66	202.14	202.14	202.14	202.14	
YLR227C	0.99	1.36	1.42	1.74	1.93	34.77	1.42	107.27	29.18	141.97	31.80	187.99	31.80	187.99	187.99	187.99	187.99	
YLR229W	0.98	1.56	1.81	2.12	2.20	55.71	2.95	149.52	35.54	205.22	38.50	175.98	38.50	175.98	175.98	175.98	175.98	
YLR300W	0.98	1.31	1.48	1.99	2.16	18.86	2.71	82.48	29.27	101.34	31.98	147.70	31.98	147.70	147.70	147.70	147.70	
YLR304C	0.93	1.38	1.67	2.12	2.19	19.69	2.29	69.07	27.39	88.16	29.68	165.55	29.68	165.55	165.55	165.55	165.55	
YLR307C	0.95	1.44	1.72	2.13	2.23	19.33	2.88	71.68	34.57	97.01	37.45	147.84	37.45	147.84	147.84	147.84	147.84	
YLR312C	0.93	1.56	1.87	2.19	2.26	27.52	2.97	82.97	33.30	110.50	36.27	193.62	36.27	193.62	193.62	193.62	193.62	
YLR313C	0.93	1.47	1.73	2.07	2.17	32.48	4.90	93.80	47.12	126.28	52.02	203.10	52.02	203.10	203.10	203.10	203.10	
YLR319C	0.97	1.62	1.49	1.84	1.96	28.74	4.11	89.27	43.00	118.01	47.11	181.39	47.11	181.39	181.39	181.39	181.39	
YLR325C	0.98	1.36	1.52	1.91	1.96	17.90	2.13	87.96	30.10	105.86	32.23	151.97	32.23	151.97	151.97	151.97	151.97	
YLR326W</																		

ORF	Optical density at 600 nm at time intervals (h)										GLUCOSE		FRUCTOSE		TOTAL Sugars(g/L)		TOTAL AMINO ACIDS (mg/L)	
	0	20	27	42	67	43 h	67 h	43 h	67 h	43 h	67 h	48 h	67 h	43 h	67 h	48 h		
YMR160W	0.26	1.80	2.04	2.05	2.29	16.30	17.5	75.63	21.00	93.47	20.30	250.32	20.30	107.88	20.30	250.32		
YMR161W	0.34	1.41	1.83	2.16	2.23	39.24	3.69	102.42	46.63	127.46	50.33	194.11	50.33	102.42	50.33	194.11		
YMR162C	0.35	1.58	1.98	2.16	2.37	37.48	4.43	109.28	50.29	116.76	54.82	207.13	54.82	116.76	54.82	207.13		
YMR169C	0.34	1.67	1.94	2.17	2.23	37.05	4.33	107.19	51.62	133.24	17.00	274.36	17.00	108.99	17.00	274.36		
YMR171C	0.35	1.50	1.77	2.10	2.22	18.03	1.26	81.87	8.02	99.99	9.27	308.01	9.27	81.87	9.27	308.01		
YMR173W	1.02	1.58	1.84	2.21	2.29	53.38	0.92	92.50	19.85	107.88	20.56	100.40	20.56	107.88	20.56	100.40		
YMR174C	0.98	1.54	1.83	2.14	2.24	27.08	2.65	96.78	28.60	123.86	31.26	150.87	31.26	96.78	31.26	150.87		
YMR176W	0.94	1.47	1.77	1.93	2.05	27.04	3.12	87.52	36.61	114.56	39.73	163.91	39.73	87.52	39.73	163.91		
YMR181C	0.91	1.42	1.89	2.13	2.29	31.88	2.31	104.31	29.47	136.39	31.77	170.51	31.77	104.31	31.77	170.51		
YMR182C	0.95	1.41	1.62	1.95	1.99	25.09	2.60	107.26	32.97	132.35	35.57	91.12	35.57	107.26	35.57	91.12		
YMR189W	0.87	1.24	1.75	2.06	2.19	31.85	2.46	103.50	33.21	135.36	35.67	134.88	35.67	103.50	35.67	134.88		
YMR198W	1.03	1.82	1.93	2.22	2.29	20.41	1.34	101.73	23.11	122.13	24.46	107.26	24.46	101.73	24.46	107.26		
YMR199W	1.01	1.68	1.99	2.26	2.37	23.22	2.17	92.60	32.88	115.82	35.15	157.12	35.15	92.60	35.15	157.12		
YMR201C	0.97	1.61	1.90	2.18	2.28	22.17	1.96	88.82	29.18	108.99	31.14	161.97	31.14	88.82	31.14	161.97		
YMR202W	1.03	1.81	2.00	2.31	2.35	21.92	1.92	91.56	24.32	113.48	26.24	187.37	26.24	91.56	26.24	187.37		
YMR204C	0.92	1.47	1.74	2.08	2.18	34.06	2.58	114.88	37.61	148.93	40.23	190.28	40.23	114.88	40.23	190.28		
YMR205C	0.97	1.52	1.75	2.11	2.17	29.80	1.58	119.97	24.51	149.77	26.09	94.14	26.09	119.97	26.09	94.14		
YMR214W	0.86	1.12	1.29	1.68	1.73	17.17	1.39	81.63	23.24	98.80	24.62	32.02	24.62	81.63	24.62	32.02		
YMR215W	0.90	1.17	1.33	1.73	1.81	16.05	2.13	75.87	20.80	91.91	22.93	150.66	22.93	75.87	22.93	150.66		
YMR216C	0.91	1.26	1.33	1.73	1.89	26.93	3.64	90.51	34.95	117.44	38.59	25.35	38.59	90.51	38.59	25.35		
YMR221C	0.93	1.29	2.23	2.07	2.20	23.82	2.79	104.83	23.44	128.65	26.22	134.67	26.22	104.83	26.22	134.67		
YMR222C	0.94	1.59	2.37	2.05	2.23	48.09	3.29	130.29	39.55	178.38	42.84	187.10	42.84	130.29	42.84	187.10		
YMR225C	0.97	1.59	1.88	2.19	2.28	35.53	3.14	119.73	42.46	155.25	45.60	180.12	45.60	119.73	45.60	180.12		
YMR226C	0.99	1.68	1.98	2.24	2.33	33.97	2.45	106.46	34.45	140.43	36.90	197.00	36.90	106.46	36.90	197.00		
YMR228W	0.98	1.58	1.91	2.20	2.29	41.25	3.30	133.74	38.91	174.99	42.21	172.49	42.21	133.74	42.21	172.49		
YMR230W	0.98	1.18	1.56	2.05	2.24	62.78	5.55	147.52	64.08	210.30	69.63	156.11	69.63	147.52	69.63	156.11		
YMR232W	0.87	1.32	1.57	1.86	1.98	21.30	1.52	91.23	22.99	112.53	24.51	211.85	24.51	91.23	24.51	211.85		
YMR233W	0.93	1.59	1.88	2.14	2.24	18.22	2.42	77.98	25.38	95.21	27.80	228.85	27.80	77.98	27.80	228.85		
YMR234W	0.92	1.45	1.68	1.91	2.02	24.88	1.99	88.87	17.81	113.77	19.79	248.69	19.79	88.87	19.79	248.69		
YMR235W	0.94	1.54	1.83	2.07	2.20	23.65	2.81	91.95	29.21	115.60	32.08	186.52	32.08	91.95	32.08	186.52		
YMR245W	0.94	1.26	1.58	1.76	1.85	21.6	2.05	101.26	21.6	130.87	30.71	119.48	30.71	101.26	30.71	119.48		
YMR246W	0.69	1.12	1.29	1.71	1.85	35.13	2.34	120.73	31.41	155.36	34.36	152.65	34.36	120.73	34.36	152.65		
YMR250C	0.91	1.12	1.66	2.14	2.21	21.62	2.54	84.48	23.31	107.48	25.65	187.22	25.65	84.48	25.65	187.22		
YMR252C	0.88	1.46	1.71	2.04	2.14	20.28	2.58	81.93	27.68	102.16	30.33	203.19	30.33	81.93	30.33	203.19		
YMR254C	1.00	1.60	1.82	2.17	2.11	27.02	0.16	76.28	11.53	86.00	11.69	83.35	11.69	76.28	11.69	83.35		
YMR259W	0.88	1.35	1.57	1.84	1.94	16.42	1.94	78.14	16.05	94.57	17.99	196.23	17.99	78.14	17.99	196.23		
YMR267C	0.85	0.90	0.88	0.84	0.84	151.84	20.77	181.00	183.76	332.55	399.93	527.98	399.93	181.00	399.93	527.98		
YMR269C	0.86	1.34	1.62	1.94	2.06	24.92	2.19	93.34	31.14	118.26	34.33	142.01	34.33	93.34	34.33	142.01		
YMR270C	0.92	1.44	1.70	2.04	2.17	29.92	1.76	104.79	13.54	134.71	15.30	200.16	15.30	104.79	15.30	200.16		
YMR271C	0.92	1.38	1.54	1.81	1.92	29.50	2.68	99.74	22.90	129.24	25.59	179.93	25.59	99.74	25.59	179.93		
YMR272W	0.96	1.61	1.89	2.13	2.25	23.02	3.04	84.97	32.52	107.98	36.07	158.65	36.07	84.97	36.07	158.65		
YMR273C	1.01	1.65	1.65	1.99	2.11	20.47	2.04	96.42	25.41	111.33	27.52	171.56	27.52	96.42	27.52	171.56		
YMR275C	1.01	1.58	1.84	2.14	2.24	25.53	2.03	104.21	39.69	129.74	43.73	193.92	43.73	104.21	43.73	193.92		
YMR276W	0.94	1.27	1.55	1.87	2.05	75.57	2.87	159.58	93.77	228.05	121.94	188.73	121.94	159.58	93.77	188.73		
YMR279C	0.88	1.27	1.56	1.92	2.13	25.77	1.94	99.58	17.29	125.43	19.23	199.22	19.23	99.58	19.23	199.22		
YMR280W	0.92	1.51	1.56	1.87	1.98	21.06	2.04	87.89	22.70	108.94	24.75	184.94	24.75	87.89	24.75	184.94		
YMR285C	0.86	1.48	1.82	2.13	2.21	23.49	3.91	88.28	30.79	109.77	34.70	217.11	34.70	88.28	34.70	217.11		
YMR289W	0.87	1.32	1.54	1.81	1.92	29.50	2.68	99.74	22.90	129.24	25.59	179.93	25.59	99.74	25.59	179.93		
YMR291W	0.99	1.63	1.65	1.99	2.10	25.33	2.79	121.73	54.19	147.06	56.97	94.02	56.97	121.73	54.19	94.02		
YMR294W	0.88	1.37	1.64	1.96	2.06	26.55	1.97	94.51	16.81	121.06	20.89	243.67	20.89	94.51	20.89	243.67		
YMR300C	0.88	1.07	1.09	1.21	1.26	16.04	1.45	104.75	19.47	162.75	16.75	346.37	16.75	104.75	16.75	346.37		
YMR302C	0.89	1.38	1.68	2.02	2.14	25.28	2.24	86.13	18.54	111.41	20.82	151.01	20.82	86.13	20.82	151.01		
YMR304W	0.88	1.53	1.79	2.08	2.17	24.84	2.72	85.87	25.64	118.71	28.36	188.94	28.36	85.87	28.36	188.94		
YNL029C	0.87	1.35	1.72	1.88	26.78	3.61	103.14	31.08	139.92	34.69	194.56	34.69	103.14	34.69	194.56			
YNL030W	0.90	1.32	1.56	1.93	2.09	31.86	2.91	90.82	26.91	122.38	28.82	194.82	28.82	90.82	28.82	194.82		
YNL031C	0.86	1.20	1.43	1.73	1.87	25.31	2.61	90.97	28.89	116.27	31.50	123.64	31.50	90.97	31.50	123.64		
YNL032W	0.92	1.24	1.42	1.83	1.94	26.33	2.43	110.89	35.18	137.22	37.62	105.69	37.62	110.89	37.62	105.69		
YNL034W	0.88	1.41	1.76	2.10	2.20	25.87	2.57	91.13	24.20	117.00	26.77	157.93	26.77	91.13	26.77	157.93		
YNL051W	1.07	1.69	1.93	2.21	2.28	24.96	0.77	101.52	20.33	120.98	21.10	191.43	21.10	101.52	21.10	191.43		
YNL052W	0.99	1.52	1.67	2.00	2.12	33.49	3.32	113.41	34.56	146.30	37.95	227.75	37.95	113.41	37.95	227.75		
YNL054W	0.87</																	

ORF	Optical density at 600 nm at time intervals (h)										GLUCOSE		FRUCTOSE		TOTAL Sugars(g/L)		TOTAL AMINO ACIDS (mg/L)	
	0	20	27	42	67	43 h	67 h	43 h	67 h	43 h	67 h	43 h	67 h	43 h	67 h	48 h		
YNL147W	0.51	1.53	2.45	2.20	20.65	1.91	65.53	77.61	107.18	15.60	2.54	1.53	1.53	22.68	23.61	145.11		
YNL148W	0.55	1.41	2.00	1.01	0.96	162.47	172.33	161.51	151.79	326.28	330.12	52.69	52.69	405.30	405.30	512.28		
YNL149W	0.90	1.31	1.43	1.76	1.02	1.62	1.82	2.16	2.47	1.73	1.73	1.73	1.73	1.73	1.73	1.73		
YNL149W	0.95	1.01	0.99	1.01	0.96	166.44	197.07	167.44	174.04	323.68	374.64	51.52	51.52	88.82	88.82	100.81		
YNL149C-A	0.87	1.14	1.62	1.86	2.01	16.14	1.71	78.41	23.59	95.55	25.30	12.01	12.01	12.01	12.01	12.01		
YNL149C	0.87	1.12	1.58	1.88	1.85	19.12	1.45	84.73	19.53	103.84	20.98	10.82	10.82	10.82	10.82	10.82		
YNL148C	0.90	1.25	1.72	2.03	2.18	30.98	4.54	98.92	43.95	129.89	48.09	14.05	14.05	14.05	14.05	14.05		
YNL157W	0.91	1.59	1.83	2.22	2.22	18.01	1.67	75.23	14.93	94.24	16.60	12.47	12.47	12.47	12.47	12.47		
YNL157W	0.97	1.00	0.98	1.00	0.94	151.37	198.88	169.46	169.66	320.84	366.54	516.32	516.32	516.32	516.32	516.32		
YNL160W	0.88	1.41	1.72	1.88	1.99	21.12	2.21	74.14	21.41	95.26	23.61	12.69	12.69	12.69	12.69	12.69		
YNL160W	1.01	1.05	1.03	1.07	1.01	148.43	161.14	169.41	150.87	317.83	312.02	515.12	515.12	515.12	515.12	515.12		
YNL164C	0.88	1.16	1.62	1.85	2.00	16.82	1.94	70.31	26.65	87.13	28.59	94.11	94.11	94.11	94.11	94.11		
YNL165W	0.82	0.95	1.39	1.73	1.31	32.01	4.72	101.59	46.86	133.50	51.38	118.07	118.07	118.07	118.07	118.07		
YNL167C	0.92	1.61	1.94	2.19	2.33	23.51	4.43	79.27	32.58	102.78	37.01	142.13	142.13	142.13	142.13	142.13		
YNL167C	1.04	1.09	1.07	1.11	1.05	161.44	183.06	167.88	165.57	329.12	348.62	515.04	515.04	515.04	515.04	515.04		
YNL173C	0.91	1.49	1.86	2.19	2.35	23.74	2.42	81.89	20.80	105.62	23.22	108.82	108.82	108.82	108.82	108.82		
YNL173C	1.06	1.10	1.08	1.12	1.06	149.83	194.14	159.21	167.07	309.04	361.21	516.93	516.93	516.93	516.93	516.93		
YNL176C	0.76	1.02	1.51	1.78	1.91	13.47	1.22	77.50	11.47	90.97	12.69	85.56	85.56	85.56	85.56	85.56		
YNL180C	0.89	1.58	1.95	2.09	2.23	80.86	50.83	153.32	125.36	234.18	176.19	209.72	209.72	209.72	209.72	209.72		
YNL180C	0.93	0.98	0.95	0.98	0.92	144.17	176.05	167.65	155.85	311.82	331.89	520.40	520.40	520.40	520.40	520.40		
YNL183C	0.91	1.50	1.71	1.98	2.13	24.53	2.16	97.97	23.25	122.50	25.40	104.52	104.52	104.52	104.52	104.52		
YNL183C	0.94	0.99	0.96	0.99	0.93	151.81	198.52	150.28	166.89	302.10	365.41	521.77	521.77	521.77	521.77	521.77		
YNL187W	0.86	1.43	1.87	2.16	2.30	23.24	2.36	81.06	26.52	104.30	30.88	104.61	104.61	104.61	104.61	104.61		
YNL190W	0.89	1.47	1.67	1.96	2.14	20.16	1.61	87.77	15.85	107.92	17.46	123.35	123.35	123.35	123.35	123.35		
YNL190W	0.97	1.02	0.99	1.02	0.96	156.00	185.20	161.59	163.54	317.59	348.74	519.76	519.76	519.76	519.76	519.76		
YNL191W	0.81	1.20	1.50	1.76	1.92	25.33	3.43	87.93	41.83	113.26	45.28	107.17	107.17	107.17	107.17	107.17		
YNL193W	0.79	1.23	1.54	1.85	1.97	23.17	2.83	86.64	37.01	109.81	39.85	134.88	134.88	134.88	134.88	134.88		
YNL194C	0.87	1.52	1.70	2.04	2.18	19.58	1.78	81.73	19.39	101.30	21.17	108.41	108.41	108.41	108.41	108.41		
YNL194C	1.03	1.10	1.07	1.09	1.03	148.69	177.92	162.17	151.04	310.86	328.95	519.87	519.87	519.87	519.87	519.87		
YNL195C	0.85	1.29	1.60	1.94	2.12	21.32	2.42	84.12	32.84	105.45	35.25	103.60	103.60	103.60	103.60	103.60		
YNL196C	0.72	1.67	1.74	1.84	1.76	24.95	2.72	86.88	36.55	115.62	38.81	105.04	105.04	105.04	105.04	105.04		
YNL196C	0.68	1.21	1.36	1.89	1.92	31.63	3.23	81.52	10.11	157.55	37.02	123.34	123.34	123.34	123.34	123.34		
YNL196C	0.97	1.33	1.00	1.02	0.97	161.64	161.45	161.25	154.67	327.19	316.03	514.90	514.90	514.90	514.90	514.90		
YNL200C	0.81	1.22	1.55	1.88	2.07	18.86	2.92	81.15	31.16	109.23	21.28	121.51	121.51	121.51	121.51	121.51		
YNL204W	0.81	1.17	1.63	1.85	2.04	17.85	2.11	83.64	26.47	101.50	28.58	81.01	81.01	81.01	81.01	81.01		
YNL204C	0.88	1.48	1.77	2.12	2.17	17.45	1.80	86.29	20.08	86.74	21.88	105.69	105.69	105.69	105.69	105.69		
YNL204C	0.98	1.04	1.01	1.02	0.97	147.23	188.11	162.70	161.40	309.93	369.51	518.11	518.11	518.11	518.11	518.11		
YNL206C	0.79	1.15	1.61	1.86	1.95	18.96	1.97	85.73	25.49	104.69	27.48	94.43	94.43	94.43	94.43	94.43		
YNL208W	0.74	0.98	1.39	1.57	1.68	20.10	2.36	86.76	33.18	106.36	35.55	105.25	105.25	105.25	105.25	105.25		
YNL211C	0.70	1.05	1.05	1.51	1.73	21.31	2.59	85.91	32.87	107.22	35.45	105.61	105.61	105.61	105.61	105.61		
YNL214W	0.69	1.25	1.58	1.72	1.88	21.88	2.83	85.95	37.24	108.84	40.07	92.25	92.25	92.25	92.25	92.25		
YNL219W	0.65	0.98	1.24	1.49	1.63	37.15	4.23	113.37	43.29	150.51	47.52	77.21	77.21	77.21	77.21	77.21		
YNL217W	0.64	1.06	1.35	1.62	1.74	23.79	3.21	88.53	43.90	112.32	47.11	116.25	116.25	116.25	116.25	116.25		
YNL218W	0.69	1.14	1.48	1.81	1.98	22.78	3.05	87.81	38.22	110.58	41.27	104.59	104.59	104.59	104.59	104.59		
YNL219C	0.89	1.51	1.85	2.15	2.19	18.97	1.97	85.73	25.49	104.69	27.48	94.43	94.43	94.43	94.43	94.43		
YNL219C	1.01	1.08	1.05	1.07	1.01	160.53	184.58	157.49	165.03	318.02	349.61	517.10	517.10	517.10	517.10	517.10		
YNL223W	0.89	1.60	1.90	2.16	2.23	16.59	1.72	69.75	29.58	86.34	31.27	66.74	66.74	66.74	66.74	66.74		
YNL229W	0.95	1.62	1.91	2.18	2.29	17.03	1.76	69.79	28.58	86.82	30.34	161.51	161.51	161.51	161.51	161.51		
YNL230C	0.97	1.68	1.97	2.23	2.34	18.71	1.73	73.08	25.08	97.08	26.00	101.79	101.79	101.79	101.79	101.79		
YNL234W	0.93	1.56	1.96	2.21	2.31	23.71	3.11	87.08	18.46	109.67	19.88	199.55	199.55	199.55	199.55	199.55		
YNL234W	0.94	1.57	1.96	2.21	2.31	22.77	2.98	87.41	19.87	98.17	22.92	140.43	140.43	140.43	140.43	140.43		
YOL028C	0.94	1.41	1.66	2.06	2.11	18.56	1.07	81.43	20.69	99.99	21.76	146.74	146.74	146.74	146.74	146.74		
YOL042W	1.00	1.80	2.11	2.06	2.16	18.80	1.70	87.61	21.57	106.41	23.26	139.57	139.57	139.57	139.57	139.57		
YOL070C	1.00	1.61	1.84	2.22	2.30	21.69	2.30	82.02	28.97	103.70	31.27	124.79	124.79	124.79	124.79	124.79		
YOL089C	0.99	1.56	1.81	2.20	2.27	21.70	1.54	87.84	24.26	109.54	25.81	171.05	171.05	171.05	171.05	171.05		
YOL092W	0.98	1.50	1.83	2.22	2.28	21.13	1.97	82.81	26.10	103.33	28.07	123.90	123.90	123.90	123.90	123.90		
YOL098C	1.02	2.05	2.05	2.18	2.08	9.51	-0.02	67.31	5.96	76.82	5.95	48.90	48.90	48.90	48.90	48.90		
YOL100W	0.96	1.53	1.72	2.01	2.11	17.40	4.84	80.04	16.88	111.16	31.48	140.22	140.22	140.22	140.22	140.22		
YOL103W	1.06	1.65	1.69	1.91	2.12	17.54	1.20	75.66	19.95	93.21	21.15	146.55	146.55	146.55	146.55	146.55		
YOL134W	1.06	1.60	1.88	2.24	2.29													

ORF	Optical density at 600 nm at time intervals (h)							GLUCOSE		FRUCTOSE		TOTAL Sugars(g/L)		TOTAL AMINO ACIDS (mg/L)	
	0	20	27	42	67	43 h	67 h	43 h	67 h	43 h	67 h	43 h	67 h	48 h	
YOR293W	0.57	1.05	1.36	1.92	2.30	194.00	171.60	167.33	162.80	272.80	120.01	214.51	102.01	205.52	
YOR294W	0.58	1.14	1.22	1.25	2.09	111.1	31.75	171.20	120.02	261.30	157.77	246.21	102.01	205.52	
YOR295W	0.93	1.68	1.04	0.87	0.60	154.18	185.41	169.53	208.41	337.71	352.62	517.27	140.05	218.68	
YOR296W	0.98	1.63	1.90	2.21	2.28	213.1	1.15	97.01	21.19	317.32	229.34	517.27	140.05	218.68	
YOR297W	0.95	1.09	1.06	0.91	0.89	166.77	166.82	188.50	191.62	355.28	358.44	516.38	102.01	205.52	
YOR298W	0.96	1.62	1.93	2.09	2.26	22.18	2.29	81.59	22.76	103.77	25.05	218.68	102.01	205.52	
YOR299C	0.95	1.35	1.67	2.01	2.15	17.71	2.34	72.91	25.15	90.62	27.49	199.32	102.01	205.52	
YOR300W	0.98	1.44	1.75	2.04	2.22	21.45	3.00	75.51	26.04	97.96	26.04	241.88	102.01	205.52	
YOR302C	0.97	0.98	1.06	0.99	1.34	157.01	152.17	184.22	212.12	341.72	364.29	489.73	102.01	205.52	
YOR322W	1.01	1.72	2.08	2.33	2.35	35.03	2.02	105.86	31.46	140.89	33.48	402.52	102.01	205.52	
YOR334W	0.96	1.52	1.78	1.92	2.16	20.11	1.74	88.65	19.53	108.76	21.27	189.04	102.01	205.52	
YOR337W	0.96	1.58	1.80	2.08	2.13	18.94	0.68	88.23	12.44	105.18	13.12	370.65	102.01	205.52	
YOR338W	0.97	1.71	2.00	2.13	2.25	22.57	1.71	99.04	20.05	121.81	21.76	240.92	102.01	205.52	
YOR342C	0.97	1.67	1.91	2.06	2.20	25.49	1.97	100.20	20.11	125.59	22.08	268.70	102.01	205.52	
YOR344C	0.95	1.55	1.67	2.06	2.16	27.02	1.44	109.93	23.63	136.94	25.08	115.46	102.01	205.52	
YOR346W	0.96	1.03	0.99	1.01	0.96	137.13	176.05	188.35	168.67	325.49	344.72	524.92	102.01	205.52	
YOR347C	1.00	1.75	2.04	2.17	2.29	27.72	2.21	105.50	21.18	133.21	23.38	282.36	102.01	205.52	
YOR349W	0.93	1.47	1.78	1.95	2.12	24.39	2.37	81.24	24.29	105.63	26.66	256.25	102.01	205.52	
YOR355W	0.88	1.27	1.55	1.98	2.10	30.04	1.09	115.26	20.19	145.29	21.28	112.10	102.01	205.52	
YOR356W	0.94	1.54	1.86	2.09	2.20	18.42	1.73	77.06	18.96	95.47	20.69	207.68	102.01	205.52	
YOR358W	1.04	1.82	1.92	2.12	2.24	12.22	0.19	81.63	2.73	93.85	2.91	273.56	102.01	205.52	
YOR363C	0.99	1.03	1.02	1.05	0.99	147.34	223.35	176.76	187.55	324.10	410.90	506.85	102.01	205.52	
YOR365C	0.94	1.36	1.69	2.01	2.16	24.30	1.94	85.51	22.85	109.82	24.79	252.63	102.01	205.52	
YOR368W	0.96	1.30	1.58	1.94	2.14	37.09	2.76	106.20	31.48	143.29	34.24	234.32	102.01	205.52	
YOR375C	1.01	1.60	1.90	2.15	2.25	35.98	3.09	106.97	32.38	142.95	35.47	164.81	102.01	205.52	
YOR377W	1.00	1.05	1.03	1.00	1.00	145.22	163.32	176.87	165.60	322.09	328.92	513.95	102.01	205.52	
YOR380W	0.96	1.36	1.73	2.20	2.23	33.24	2.78	110.79	35.49	144.03	38.27	204.27	102.01	205.52	
YOR381W	0.94	1.53	1.41	1.75	1.89	32.50	3.81	112.77	39.53	145.26	43.34	113.26	102.01	205.52	
YOR382W	1.00	1.06	1.03	1.04	0.98	143.99	204.48	169.64	181.71	313.63	384.19	512.51	102.01	205.52	
YOR383C	0.96	1.46	1.72	2.10	2.19	24.12	0.86	113.91	19.56	138.03	20.42	158.60	102.01	205.52	
YOR385W	0.93	1.66	1.96	2.13	2.31	21.95	1.63	114.90	15.75	151.50	16.65	149.32	102.01	205.52	
YOR386W	0.97	1.51	1.88	2.23	2.36	25.29	2.40	124.51	26.11	120.10	31.81	175.69	102.01	205.52	
YPL001W	0.95	1.28	1.30	1.96	2.20	22.58	2.03	100.41	17.58	123.08	18.23	191.38	102.01	205.52	
YPL002C	0.99	1.57	1.86	1.89	2.03	29.68	0.68	124.68	18.89	156.28	18.41	491.88	102.01	205.52	
YPL004C	0.93	1.29	1.86	1.99	2.08	26.88	1.54	109.28	22.65	138.67	24.18	143.96	102.01	205.52	
YPL009W	0.95	1.58	1.81	2.12	2.16	21.64	1.76	91.41	18.04	113.05	19.80	228.73	102.01	205.52	
YPL014W	0.97	1.58	1.87	2.04	2.16	24.43	1.61	105.20	15.45	138.84	17.06	257.73	102.01	205.52	
YPL015C	0.95	1.51	1.82	1.99	2.15	29.72	1.87	112.78	19.22	142.50	21.09	219.19	102.01	205.52	
YPL017C	1.19	2.38	2.22	2.45	2.45	15.58	1.93	75.56	5.25	93.14	7.18	266.55	102.01	205.52	
YPL021W	1.02	1.09	1.07	1.08	1.04	158.18	166.10	169.32	160.59	327.50	326.69	6.36	102.01	205.52	
YPL022W	1.00	1.59	1.89	2.17	2.25	30.15	2.51	100.45	25.87	130.59	28.38	215.81	102.01	205.52	
YPL022W	0.98	1.40	1.68	2.00	2.14	25.97	1.47	97.81	11.93	123.78	13.40	314.83	102.01	205.52	
YPL030W	0.97	1.35	1.60	1.91	2.14	31.81	2.19	113.21	26.07	145.02	26.27	196.95	102.01	205.52	
YPL034W	1.13	1.83	2.12	2.40	2.49	28.18	153.30	113.79	160.61	141.97	313.91	6.44	102.01	205.52	
YPL036W	1.07	1.71	2.01	2.28	2.37	33.55	3.63	109.25	20.75	142.60	31.18	4.20	102.01	205.52	
YPL037C	1.01	1.46	1.78	2.13	2.27	36.76	2.36	113.37	24.17	150.13	26.53	289.91	102.01	205.52	
YPL038W	1.14	1.24	1.24	1.27	1.20	164.20	5.15	175.32	36.46	339.52	41.61	4.45	102.01	205.52	
YPL039W	0.96	1.50	1.85	2.17	2.30	28.06	2.87	93.84	31.51	121.91	34.38	216.22	102.01	205.52	
YPL040C	0.95	1.24	1.56	2.21	2.28	46.77	6.53	119.06	64.54	165.83	71.07	262.13	102.01	205.52	
YPL041C	0.94	1.39	1.72	2.20	2.26	31.26	2.74	113.19	30.99	144.44	33.73	207.99	102.01	205.52	
YPL046C	0.97	1.63	1.94	2.08	2.27	19.83	1.64	85.50	19.42	106.32	21.06	190.97	102.01	205.52	
YPL047W	0.93	1.58	1.89	2.06	2.18	30.83	1.56	101.27	17.60	132.10	19.16	245.87	102.01	205.52	
YPL051W	0.95	1.56	1.87	2.02	2.14	20.28	1.94	77.72	20.77	98.00	22.71	175.70	102.01	205.52	
YPL057C	0.97	1.62	1.96	2.08	2.17	31.82	2.79	107.49	26.53	139.31	29.32	249.08	102.01	205.52	
YPL068C	0.97	1.25	1.68	2.11	2.18	29.63	2.36	97.14	27.05	126.78	28.43	297.78	102.01	205.52	
YPL097W	0.94	1.41	1.72	2.19	2.27	22.67	2.88	87.14	12.90	108.81	15.77	197.57	102.01	205.52	
YPL098C	0.95	2.22	1.59	1.76	1.94	20.94	2.54	93.24	20.17	114.28	22.71	251.88	102.01	205.52	
YPL099C	0.97	1.54	1.84	1.98	2.11	21.18	1.88	84.91	21.16	108.28	24.01	213.49	102.01	205.52	
YPL100W	0.97	1.66	1.95	2.08	2.20	25.77	1.79	108.64	18.81	134.40	20.60	188.77	102.01	205.52	
YPL107W	0.96	1.46	1.79	2.18	2.22	33.99	2.47	123.99	27.05	157.99	26.52	225.83	102.01	205.52	
YPL108W	0.95	1.40	1.75	2.07	2.17	34.70	2.48	118.08	27.05	152.77	28.53	210.56	102.01	205.52	
YPL109C	1.07	1.67	2.00	2.27	2.36	28.59	1.91	94.43	20.48	123.02	22.39	149.42	102.01	205.52	
YPL111W	1.01	1.54	1.85	2.14	2.25	36.26	2.46	118.52	29.90	154.78	32.36	198.62	102.01	205.52	
YPL112C	1.01	1.43	1.69	2.05	2.22	36.26	2.47	115.12	27.54	151.38	30.01	234.47	102.01	205.52	
YPL114W	1.06	1.75	2.09	2.41	2.48	30.06	1.73	132.92	16.46	122.82	33.88	4.40	102.01	205.52	
YPL137C	0.93	1.37	1.59	2.00	2.09	14.85	1.45	88.61	12.57	107.59	14.02	99.66	102.01	205.52	
YPL138C	0.90	1.49	1.63	1.90	2.12	25.20	1.51	103.28	14.09	128.48	15.60	51.36	102.01	205.52	
YPL140C	0.97	1.66	1.98	2.30	2.34	18.86									

ORF	Optical density at 600 nm time intervals (h)										GLUCOSE		FRUCTOSE		TOTAL Sugars(g/L)		TOTAL AMINO ACIDS (mg/L)	
	0	20	27	42	67	43 h	67 h	43 h	67 h	43 h	0	20	42	67	43 h	67 h	48 h	
YPL246C	0.41	1.92	2.03	1.45	2.52	15.74	21.00	8.01	20.06	10.77	25.04	3.29	3.29	3.29	3.29	3.29	3.29	
YPL249C	0.93	1.44	1.72	1.53	2.20	18.62	18.62	7.10	14.06	9.71	17.41	137.10	137.10	137.10	137.10	137.10	137.10	
YPL251W	0.95	1.50	1.82	1.92	2.27	19.56	2.30	7.29	22.90	91.85	25.20	118.29	118.29	118.29	118.29	118.29	118.29	
YPL253C	0.95	1.33	1.81	1.97	2.25	32.64	2.39	105.20	23.13	137.63	25.52	127.94	127.94	127.94	127.94	127.94	127.94	
YPL259C	0.96	1.37	1.80	2.00	2.29	17.76	2.26	103.18	15.39	139.45	17.14	154.60	154.60	154.60	154.60	154.60	154.60	
YPL260W	0.94	1.35	1.73	2.01	2.07	13.18	1.64	69.87	17.69	83.05	19.33	102.41	102.41	102.41	102.41	102.41	102.41	
YPL263C	0.91	1.33	1.82	2.13	2.18	17.75	1.49	83.66	14.84	101.41	16.33	139.68	139.68	139.68	139.68	139.68	139.68	
YPL264C	0.97	1.50	1.96	2.24	2.30	23.47	1.86	93.30	20.77	116.77	22.63	130.98	130.98	130.98	130.98	130.98	130.98	
YPL269W	1.13	1.76	1.97	2.30	2.40	27.44	6.41	92.35	42.33	119.79	48.54	3.51	3.51	3.51	3.51	3.51	3.51	
YPL267W	0.91	1.48	1.80	2.12	2.18	20.40	1.59	85.95	19.25	106.36	20.84	129.38	129.38	129.38	129.38	129.38	129.38	
YPL268W	0.91	1.50	1.82	2.13	2.18	22.72	1.49	85.98	16.31	109.70	17.80	147.62	147.62	147.62	147.62	147.62	147.62	
YPL271W	1.07	1.63	1.91	2.28	2.38	24.78	1.72	87.58	22.38	112.36	24.09	3.72	3.72	3.72	3.72	3.72	3.72	
YPL271W	0.91	1.22	1.25	1.29	1.52	142.35	80.16	172.00	113.41	314.35	193.57	411.87	411.87	411.87	411.87	411.87	411.87	
YPL274W	1.06	1.71	2.01	2.35	2.42	29.41	1.52	107.76	18.00	136.67	19.57	3.60	3.60	3.60	3.60	3.60	3.60	
YPR0022W	1.00	1.44	1.69	2.07	2.17	28.16	1.26	98.59	20.46	126.75	21.71	3.68	3.68	3.68	3.68	3.68	3.68	
YPR003C	0.94	1.47	1.76	1.95	2.23	24.82	2.79	92.11	22.56	116.94	25.35	159.89	159.89	159.89	159.89	159.89	159.89	
YPR006C	0.93	1.41	1.75	1.88	2.18	26.36	2.89	90.06	22.76	116.42	25.65	169.95	169.95	169.95	169.95	169.95	169.95	
YPR007C	0.92	1.34	1.87	1.94	2.30	25.97	2.41	90.40	26.59	116.37	29.00	208.98	208.98	208.98	208.98	208.98	208.98	
YPR008W	1.10	1.72	2.02	2.31	2.41	37.42	0.79	120.03	15.39	157.45	16.18	3.73	3.73	3.73	3.73	3.73	3.73	
YPR009W	0.98	1.41	1.72	2.17	2.28	37.90	0.71	127.67	15.82	165.57	16.54	3.97	3.97	3.97	3.97	3.97	3.97	
YPR011C	0.93	1.20	1.58	1.76	2.14	30.25	2.31	107.79	25.38	138.04	27.69	151.26	151.26	151.26	151.26	151.26	151.26	
YPR013C	1.00	1.61	1.94	2.27	2.32	38.87	1.99	130.69	26.87	169.56	28.86	102.86	102.86	102.86	102.86	102.86	102.86	
YPR015C	0.94	1.78	2.01	2.26	2.29	9.74	2.06	59.05	23.56	68.80	25.62	102.19	102.19	102.19	102.19	102.19	102.19	
YPR016W	0.90	1.15	1.40	1.70	1.99	28.44	2.23	97.08	20.97	125.52	23.20	144.76	144.76	144.76	144.76	144.76	144.76	
YPR022C	1.04	1.77	2.07	2.37	2.40	23.74	1.78	95.05	22.75	118.78	24.53	3.10	3.10	3.10	3.10	3.10	3.10	
YPR023C	0.88	1.20	1.49	1.80	2.08	25.47	1.62	102.50	13.60	127.97	15.22	170.10	170.10	170.10	170.10	170.10	170.10	
YPR024W	0.88	1.39	1.63	1.90	2.08	60.99	17.69	123.05	123.88	123.88	123.88	211.05	211.05	211.05	211.05	211.05	211.05	
YPR024W	1.02	1.04	1.16	1.14	1.10	158.23	174.30	174.07	165.92	332.30	360.21	514.50	514.50	514.50	514.50	514.50	514.50	
YPR026W	0.95	1.08	1.26	1.60	1.73	87.91	24.35	163.48	102.35	251.46	126.70	523.23	523.23	523.23	523.23	523.23	523.23	
YPR027C	1.04	1.62	2.14	2.05	2.47	16.87	1.35	85.55	14.26	112.42	15.62	514.44	514.44	514.44	514.44	514.44	514.44	
YPR031W	0.95	1.36	1.32	1.17	1.21	178.04	13.55	178.04	13.55	181.50	15.62	292.05	292.05	292.05	292.05	292.05	292.05	
YPR032W	0.99	1.44	1.15	1.10	1.64	101.15	216.15	164.44	194.94	344.69	410.99	520.98	520.98	520.98	520.98	520.98	520.98	
YPR034W	0.98	1.14	1.15	1.06	1.01	176.25	202.22	193.22	193.22	371.58	395.51	518.76	518.76	518.76	518.76	518.76	518.76	
YPR040W	1.05	1.75	2.08	2.36	2.40	19.78	2.70	92.85	27.05	112.24	29.75	3.01	3.01	3.01	3.01	3.01	3.01	
YPR042C	0.93	1.68	1.96	2.23	2.29	21.39	2.79	69.26	23.16	91.34	25.95	83.59	83.59	83.59	83.59	83.59	83.59	
YPR042C	0.99	1.63	1.83	2.06	2.11	28.78	3.15	104.99	32.75	133.78	35.90	186.00	186.00	186.00	186.00	186.00	186.00	
YPR043W	1.00	1.72	2.04	2.23	2.41	24.08	1.90	92.88	22.85	116.96	24.75	152.19	152.19	152.19	152.19	152.19	152.19	
YPR046W	0.96	1.53	1.82	1.97	2.24	28.93	2.22	100.91	14.61	129.84	16.83	252.08	252.08	252.08	252.08	252.08	252.08	
YPR047W	0.96	1.56	1.93	1.96	2.08	28.88	2.52	102.74	22.76	131.61	25.28	128.49	128.49	128.49	128.49	128.49	128.49	
YPR049C	0.96	0.95	1.09	0.91	1.05	155.19	192.45	168.72	192.45	192.45	232.31	323.31	323.31	323.31	323.31	323.31	323.31	
YPR052C	0.88	1.13	1.55	1.75	1.88	33.68	2.51	110.37	30.36	144.05	32.87	107.76	107.76	107.76	107.76	107.76	107.76	
YPR054W	1.07	1.64	1.92	2.28	2.36	26.57	2.07	99.76	24.32	126.33	26.39	3.11	3.11	3.11	3.11	3.11	3.11	
YPR057W	0.86	1.10	1.35	1.69	1.99	35.42	3.69	119.04	35.20	154.47	38.89	110.28	110.28	110.28	110.28	110.28	110.28	
YPR058W	0.98	1.75	2.13	2.30	2.32	7.35	1.19	62.06	3.47	69.43	4.66	100.40	100.40	100.40	100.40	100.40	100.40	
YPR061C	1.00	1.74	2.02	2.25	2.35	11.96	1.56	66.66	6.95	104.22	8.51	190.19	190.19	190.19	190.19	190.19	190.19	
YPR062W	0.97	1.64	2.13	2.13	2.29	22.09	1.19	72.31	6.43	84.27	8.04	212.13	212.13	212.13	212.13	212.13	212.13	
YPR079W	0.92	1.43	1.72	2.14	2.23	20.82	1.69	95.52	23.87	116.34	25.56	-2.96	-2.96	-2.96	-2.96	-2.96	-2.96	
YPR084W	0.92	1.30	2.00	1.86	2.01	28.41	2.48	102.82	25.01	131.23	27.49	139.50	139.50	139.50	139.50	139.50	139.50	
YPR098C	1.02	1.60	1.98	2.31	2.40	35.66	4.61	117.01	38.77	152.67	43.38	3.15	3.15	3.15	3.15	3.15	3.15	
YPR100W	0.88	1.22	1.42	1.76	2.08	41.16	4.21	127.98	37.73	169.16	41.94	94.61	94.61	94.61	94.61	94.61	94.61	
YPR105W	1.06	1.74	1.99	2.27	2.37	18.26	1.27	126.34	15.08	105.05	31.41	148.39	148.39	148.39	148.39	148.39	148.39	
YPR109W	0.98	1.26	1.51	2.18	2.35	23.81	1.76	87.74	20.49	116.49	25.80	195.60	195.60	195.60	195.60	195.60	195.60	
YPR111W	1.04	1.78	2.00	2.31	2.38	13.68	1.28	103.98	12.05	124.82	13.34	124.84	124.84	124.84	124.84	124.84	124.84	
YPR119W	1.03	1.88	1.18	1.14	1.48	159.83	173.23	191.90	193.61	351.73	369.64	569.12	569.12	569.12	569.12	569.12	569.12	
YPR120C	1.04	1.69	1.97	2.20	2.30	20.89	1.56	87.93	17.64	108.83	33.19	148.51	148.51	148.51	148.51	148.51	148.51	
YPR121W	1.01	1.79	2.08	2.33	2.37	21.38	1.84	98.86	21.36	129.24	21.20	162.09	162.09	162.09	162.09	162.09	162.09	
YPR124W	1.10	1.89	2.09	2.33														

Appendix 3. Micro-fermentation data of WYDL_LN (all candidates).

ORF name	OD at 600 nm at sampling time points (hours)												GLUCOSE		FRUCTOSE		TOTAL SUGARS	
	0 h	20 h	27 h	44 h	66 h	93 h	113 h	120 h	146 h	120 h	146 h	120 h	146 h	120 h	146 h	120 h	146 h	
YNL145W	0.87	1.15	1.25	1.29	1.42	1.66	1.64	-4.51	0.06	6.29	-0.33	7.79	-0.27					
YDR388W	0.90	1.30	1.48	1.54	1.70	1.90	1.88	3.32	0.52	26.28	-0.63	26.60	-0.11					
YKL037W	0.86	1.12	1.24	1.31	1.45	1.78	1.69	0.38	0.76	9.47	0.00	9.85	0.77					
YDR388W	0.95	1.32	1.43	1.55	1.68	1.87	1.85	9.13	1.04	43.21	-0.17	52.33	0.88					
YER118C	0.95	1.47	1.60	1.67	1.81	1.99	1.98	2.55	1.58	40.32	-0.42	52.87	1.16					
YER115C	0.90	1.35	1.50	1.59	1.76	1.93	1.93	11.61	1.64	82.99	-0.42	94.60	1.22					
YDR496C	0.87	1.19	1.32	1.37	1.51	1.70	1.66	9.61	1.52	31.97	-0.25	41.58	1.27					
YER115C	0.90	1.41	1.59	1.74	1.87	2.04	2.04	10.83	1.67	77.80	-0.38	86.53	1.29					
YDR379W	0.90	1.31	1.41	1.54	1.69	1.89	1.86	1.26	1.26	33.59	0.07	45.50	1.32					
YER123W	0.97	1.25	1.49	1.57	1.71	1.89	1.85	1.44	1.65	60.55	-0.17	83.29	1.37					
YER123C	0.94	1.17	1.34	1.43	1.56	1.73	1.70	1.28	1.51	57.48	-0.86	69.66	1.45					
YDR333W-A	0.91	1.35	1.58	1.59	1.73	1.81	1.89	14.71	2.09	43.21	0.21	73.22	1.88					
YER114C	0.98	1.55	1.69	1.77	1.91	2.07	2.05	17.72	2.34	91.27	-0.29	108.99	2.05					
YER167W	1.00	1.47	1.63	1.76	1.88	2.05	2.04	17.00	2.30	93.60	-0.23	110.66	2.08					
YDR383C	0.90	1.34	1.50	1.61	1.75	1.95	1.92	15.63	2.47	36.74	-0.14	45.37	2.32					
YER027W	0.92	1.34	1.47	1.56	1.70	1.87	1.84	16.69	2.54	39.38	0.13	56.08	2.67					
YDR270C	0.90	1.39	1.58	1.66	1.80	1.98	1.97	1.70	2.35	43.32	0.53	60.02	2.88					
YEL011W	0.95	1.41	1.57	1.69	1.82	1.99	1.99	15.31	3.44	83.32	-0.23	96.63	3.21					
YEL019C	0.93	1.38	1.53	1.63	1.76	1.94	1.91	16.63	3.74	85.00	-0.21	101.83	3.53					
YGR266W	0.87	1.26	1.42	1.49	1.58	1.88	1.82	6.99	0.14	21.95	3.81	22.94	3.74					
YNL176C	0.80	1.16	1.22	1.29	1.39	1.71	1.64	1.60	0.36	16.45	3.68	18.05	4.04					
YIL156W	0.91	1.25	1.37	1.44	1.54	1.90	1.79	6.73	0.41	16.00	3.77	18.73	4.18					
YER083C	0.92	1.21	1.38	1.56	1.68	1.86	1.84	21.10	4.18	19.67	0.11	40.77	4.29					
YAL044W-A	0.90	1.48	1.60	1.64	1.85	2.02	2.00	2.51	0.49	30.77	4.24	33.27	4.73					
YER072W	0.96	1.49	1.65	1.75	1.90	2.06	2.05	16.69	4.12	24.43	0.66	41.12	4.78					
YOR113W	0.95	1.38	1.51	1.52	1.64	1.92	1.83	2.04	0.43	29.01	5.17	31.04	5.59					
YGL087C	0.87	1.27	1.41	1.45	1.59	1.89	1.81	1.16	0.83	23.12	5.27	24.28	6.10					
YMR171C	1.02	1.50	1.64	1.75	1.82	1.93	2.08	2.57	0.20	24.62	5.91	27.19	6.11					
YDR386W	0.96	1.40	1.55	1.68	1.82	1.98	1.97	19.46	5.99	29.03	0.40	48.49	6.40					
YPL084W	0.94	1.39	1.56	1.57	1.66	1.92	1.89	0.33	-0.07	27.70	6.47	28.03	6.40					
YNL318C	0.98	1.56	1.71	1.73	1.84	2.09	2.03	4.34	0.26	46.64	6.53	50.98	6.79					
YLR027C	0.90	1.34	1.50	1.54	1.58	1.74	1.72	1.83	0.50	34.90	6.32	36.73	6.82					
YPL096C-A	1.14	1.54	1.81	1.95	2.12	2.30	2.36	2.06	-0.04	23.98	7.46	32.04	7.42					
YGR040W	0.93	1.41	1.55	1.57	1.70	2.00	1.93	2.58	0.52	32.62	7.46	35.19	7.98					
YMR029C	0.94	1.58	1.75	1.79	1.87	2.13	2.06	2.90	0.12	45.50	7.88	47.90	8.01					
YLL039C	0.87	1.15	1.30	1.28	1.33	1.65	1.47	1.00	0.31	36.67	7.76	37.67	8.07					
YBR269W	0.86	1.33	1.62	1.57	1.69	2.00	2.02	1.46	0.12	28.01	7.99	28.47	8.11					
YLR413W	0.91	1.33	1.51	1.60	1.70	2.01	1.96	1.37	0.61	23.59	8.73	30.96	9.34					
YL055C	0.90	1.40	1.52	1.57	1.67	1.95	1.87	3.49	0.68	36.71	8.83	40.26	9.51					
YAL044C	0.95	1.47	1.60	1.63	1.74	2.00	1.95	1.96	0.59	35.10	8.71	37.55	9.75					
YAL020C	0.89	1.36	1.53	1.57	1.67	2.03	2.01	2.61	0.23	42.85	16.14	43.10	16.39					
YPL074C	0.97	1.58	1.63	1.71	1.87	2.11	2.08	1.45	0.04	33.43	10.77	34.65	10.76					
YEL012W	0.93	1.38	1.51	1.58	1.72	1.89	1.89	10.03	1.07	32.23	10.85	42.27	11.13					
YLR095C	1.01	1.51	1.66	1.76	1.85	2.11	2.12	3.36	0.39	31.74	10.83	35.10	11.22					
YLR203W	0.89	1.39	1.40	1.54	1.68	1.78	1.82	2.68	0.49	31.08	10.89	33.76	11.28					
YGR269W	1.07	1.66	1.80	1.85	1.96	2.19	2.18	2.93	0.63	35.44	11.00	38.38	11.63					
YDR320C	0.91	1.37	1.49	1.64	1.74	2.00	2.01	2.78	0.37	40.32	11.61	43.10	11.98					
YKL213C	0.93	1.43	1.55	1.59	1.71	2.00	1.92	4.97	0.39	33.48	11.71	44.45	12.10					
YIL035C	0.94	1.46	1.63	1.65	1.75	1.91	1.96	1.83	0.28	39.93	12.01	40.76	12.29					
YML084C	0.96	1.16	1.33	1.36	1.41	1.61	1.51	5.60	0.50	46.83	11.97	52.43	12.47					
YLN279C	0.88	1.28	1.42	1.52	1.67	1.89	1.85	6.91	0.59	40.89	12.00	47.80	12.58					
YGR112W	0.94	1.33	1.46	1.48	1.65	1.92	1.83	3.70	0.59	35.66	12.22	33.36	12.82					
YLR417W	0.87	1.16	1.28	1.29	1.39	1.60	1.55	2.85	0.59	52.23	12.27	56.08	12.85					
YGL019W	0.88	1.27	1.43	1.47	1.59	1.83	1.79	3.70	0.42	45.33	12.60	49.03	13.03					
YNL127W	0.86	1.11	1.27	1.28	1.37	1.70	1.57	4.10	0.72	42.09	13.00	46.20	13.72					
YLR080W	1.20	1.78	1.91	1.98	2.06	2.28	2.27	4.39	0.55	44.45	12.65	48.85	13.19					
YMR245C	0.94	1.34	1.47	1.53	1.62	1.91	1.89	3.85	0.12	48.17	14.31	51.02	14.44					
YLR030W	0.94	1.44	1.59	1.65	1.73	2.02	1.96	6.99	0.42	42.21	14.02	48.20	14.44					
YKR066C	0.94	1.43	1.60	1.69	1.81	2.02	2.04	5.53	1.16	38.08	13.49	45.81	14.66					
YLR062W	0.92	1.37	1.57	1.64	1.76	2.05	1.99	5.81	0.59	50.30	14.12	56.11	14.71					
YDL262W	0.86	1.11	1.28	1.34	1.44	1.61	1.60	6.11	0.23	45.26	14.75	52.62	14.75					
YOL124W	0.87	1.44	1.58	1.59	1.68	1.92	1.82	5.95	0.24	42.94	14.84	48.86	14.88					
YNL078W	0.91	1.42	1.69	1.73	1.71	1.88	1.92	6.46	0.25	58.65	14.67	65.13	14.92					
YKL003C	0.90	1.36	1.49	1.59	1.74	2.00	1.99	3.47	1.48	48.01	13.49	39.48	13.66					
YGL082W	0.91	1.43	1.56	1.60	1.72	1.99	1.93	6.21	1.11	47.79	15.24	54.00	16.35					
YER070W	0.90	1.34	1.49	1.56	1.69	1.98	1.91	4.68	1.05	45.80	15.32	50.48	16.38					
YMR255W	0.83	1.18	1.28	1.34	1.42	1.74	1.67	4.02	0.55	38.32	16.28	48.34	16.83					
YGR155W	0.93	1.31	1.47	1.59	1.68	1.92	1.81	6.20	0.62	52.27	16.54	54.48	17.16					
YLB268W																		

ORF name	OD at 600 nm at sampling time points (hours)										GLUCOSE		FRUCTOSE		TOTAL SUGARS	
	0 h	20 h	27 h	44 h	66 h	93 h	113 h	120 h	146 h	120 h	146 h	120 h	146 h	120 h	146 h	
YPL136C	0.86	1.00	1.47	1.57	1.65	1.64	1.61	1.62	2.11	3.66	1.74	3.56	19.66	33.56	19.66	
YER090C	0.85	1.28	1.44	1.52	1.65	1.64	1.61	1.62	2.11	3.66	1.74	3.57	19.67	33.57	19.67	
YBL003C	0.85	1.23	1.32	1.33	1.48	1.68	1.62	1.61	1.84	3.76	1.89	3.87	19.87	33.87	19.87	
YCR178C	0.91	1.22	1.41	1.41	1.53	1.83	1.71	1.72	1.99	3.54	1.88	3.74	19.88	33.74	19.88	
YLR093W	0.94	1.31	1.47	1.55	1.63	1.93	1.89	1.91	2.01	5.61	1.03	4.58	19.97	51.26	20.00	
YGR007W	1.00	1.52	1.66	1.68	1.78	2.06	2.01	2.06	2.11	5.76	1.14	4.63	19.93	51.53	19.93	
YNL141W	0.89	1.30	1.43	1.53	1.61	1.87	1.86	1.88	1.92	1.79	0.59	4.612	19.53	57.91	20.12	
YPL260W	0.93	1.38	1.48	1.54	1.64	1.92	1.78	1.79	1.82	1.79	0.59	4.612	19.53	57.91	20.12	
YHL028C	0.88	1.17	1.29	1.36	1.46	1.70	1.67	1.62	1.62	0.92	0.92	4.461	19.22	50.63	20.14	
YPL098C	0.96	1.43	1.57	1.72	1.81	2.04	2.02	2.04	2.04	6.51	1.02	4.81	19.19	47.32	20.20	
YGR169C	0.99	1.49	1.61	1.63	1.76	2.04	1.98	2.04	2.04	6.11	1.03	4.67	19.24	52.28	20.27	
YHR032W	0.89	1.26	1.39	1.56	1.66	1.99	1.93	1.93	1.93	7.93	0.42	5.73	19.85	67.65	20.27	
YMR232W	0.90	1.34	1.52	1.62	1.72	1.99	1.97	1.97	1.97	5.82	0.50	4.135	19.88	47.17	20.38	
YDR335W	0.88	1.16	1.34	1.45	1.61	1.82	1.80	1.80	1.80	15.60	2.87	29.73	17.51	45.33	20.39	
YGR017W	0.96	1.42	1.56	1.59	1.70	1.98	1.93	1.93	1.93	5.23	1.14	4.100	19.32	48.23	20.46	
YNL024C	0.87	1.21	1.30	1.27	1.34	1.68	1.57	1.64	1.64	8.44	1.58	4.634	19.87	54.78	20.55	
YDR438W	0.95	1.49	1.67	1.76	1.87	2.11	2.08	2.08	2.08	7.52	0.97	5.428	19.60	61.80	20.57	
YGL027C	0.91	1.32	1.48	1.54	1.65	1.85	1.80	1.80	1.80	9.29	0.90	6.94	19.75	79.13	20.65	
YGR068C	1.00	1.49	1.66	1.73	1.82	2.10	2.05	2.05	2.05	7.30	1.05	5.78	19.66	64.88	20.71	
YNL134C	0.94	1.35	1.54	1.66	1.76	1.99	1.99	1.99	1.99	12.46	0.75	6.042	19.97	72.88	20.71	
YGR004W	1.01	1.62	1.75	1.76	1.86	2.13	2.08	2.08	2.08	6.85	0.74	5.311	20.04	59.96	20.78	
YBR204C	0.89	1.31	1.42	1.51	1.61	1.84	1.83	1.83	1.83	7.55	0.89	5.222	19.91	60.76	20.80	
YGR232W	1.01	1.65	1.80	1.85	1.98	2.17	2.19	2.19	2.19	7.44	1.24	47.40	19.64	54.84	20.88	
YOL098C	0.94	1.52	1.70	1.72	1.80	2.07	2.01	2.01	2.01	6.94	0.67	5.645	20.23	63.38	20.90	
YLR452C	0.88	1.34	1.52	1.67	1.82	2.05	2.03	2.03	2.03	5.91	0.92	5.442	20.02	60.33	20.94	
YER183C	0.93	1.48	1.65	1.71	1.83	2.11	2.03	2.03	2.03	8.30	0.84	6.001	20.12	63.31	20.96	
YKL071W	0.94	1.46	1.65	1.74	1.87	2.09	2.06	2.06	2.06	5.30	1.76	4.139	19.30	46.69	21.06	
YJR103W	0.93	1.28	1.37	1.37	1.46	1.76	1.74	1.74	1.74	5.74	1.78	5.179	19.30	59.73	21.08	
YDL227C	0.91	1.37	1.53	1.61	1.71	2.00	2.03	2.03	2.03	6.49	0.68	5.629	20.43	64.78	21.11	
YMR34W	0.84	1.17	1.27	1.29	1.35	1.68	1.58	1.58	1.58	6.80	1.20	39.79	19.97	45.59	21.17	
YNL194W	0.85	1.18	1.29	1.32	1.43	1.76	1.76	1.76	1.76	6.76	0.79	5.040	20.38	56.80	21.47	
YOL025C	0.91	1.45	1.61	1.68	1.78	1.99	1.97	1.97	1.97	5.74	1.73	4.747	19.88	53.41	21.22	
YGR162W	0.92	1.50	1.72	1.85	2.00	2.25	2.24	2.24	2.24	5.87	0.92	4.677	20.42	53.24	21.34	
YLR151W	0.85	1.23	1.36	1.49	1.58	1.76	1.76	1.76	1.76	5.21	1.20	3.921	19.98	54.46	21.36	
YIL078W	0.92	1.30	1.45	1.51	1.62	1.88	1.84	1.84	1.84	5.55	1.07	37.59	20.29	42.14	21.37	
YGR015C	1.01	1.65	1.82	1.88	1.98	2.25	2.21	2.21	2.21	6.28	1.07	51.10	20.34	57.38	21.41	
YLR237C	0.92	1.33	1.47	1.50	1.61	1.80	1.76	1.76	1.76	5.296	1.02	49.88	20.42	63.88	21.48	
YGR036W	1.00	1.49	1.62	1.66	1.77	2.01	1.96	1.96	1.96	6.06	0.99	4.237	20.55	48.42	21.53	
YPR070W	0.91	1.26	1.28	1.43	1.51	1.60	1.65	1.65	1.65	6.59	0.61	4.638	20.96	51.27	21.57	
YEL017C-A	0.94	1.43	1.58	1.66	1.80	1.98	1.96	1.96	1.96	6.16	0.29	57.44	20.74	77.00	21.63	
YNL311C	0.92	1.37	1.56	1.59	1.72	1.95	1.90	1.90	1.90	8.37	0.89	6.633	20.74	62.58	21.85	
YDL257C	0.94	1.38	1.56	1.64	1.75	2.03	1.98	1.98	1.98	6.47	0.73	5.600	20.90	62.47	21.63	
YGL139W	0.85	1.28	1.43	1.49	1.65	1.95	1.88	1.88	1.88	3.06	1.19	35.74	20.48	48.79	21.67	
YNL146W	0.84	1.20	1.36	1.41	1.49	1.79	1.73	1.73	1.73	5.38	1.85	39.69	19.82	45.07	21.67	
YLR150W	0.91	1.39	1.54	1.63	1.74	2.03	1.97	1.97	1.97	6.96	1.08	42.97	20.81	49.93	21.69	
YGR180C	1.04	1.62	1.70	1.83	1.92	2.01	2.19	2.19	2.19	5.94	1.30	43.22	20.42	49.17	21.72	
YKL117W	0.94	1.37	1.53	1.63	1.74	2.00	1.96	1.96	1.96	6.56	1.58	49.19	20.15	55.75	21.72	
YNL183C	0.92	1.30	1.50	1.51	1.60	1.91	1.84	1.84	1.84	7.82	1.10	53.45	20.70	61.27	21.80	
YGR021W	1.02	1.61	1.78	1.85	1.96	2.18	2.17	2.17	2.17	7.15	1.26	54.53	20.59	62.58	21.85	
YPL046C	0.91	1.31	1.39	1.54	1.65	1.87	1.85	1.85	1.85	6.93	1.18	40.10	20.70	47.03	21.88	
YJU044C	0.86	1.19	1.31	1.31	1.40	1.74	1.74	1.74	1.74	4.39	1.51	4.272	20.90	47.10	22.41	
YGR271C	0.92	1.40	1.57	1.64	1.72	2.01	1.94	1.94	1.94	6.70	1.10	47.16	21.33	53.87	22.43	
YKR101W	0.88	1.30	1.47	1.44	1.59	1.87	1.87	1.87	1.87	10.20	1.02	61.40	21.43	71.60	22.45	
YNL146C-A	0.88	1.39	1.54	1.57	1.67	1.96	1.90	1.90	1.90	6.74	2.49	41.60	21.67	48.38	22.46	
YER559C	0.85	1.25	1.40	1.49	1.57	1.84	1.76	1.76	1.76	5.04	1.02	50.48	21.65	52.55	22.51	
YOL024C	0.88	1.12	1.21	1.22	1.30	1.56	1.49	1.49	1.49	4.74	1.41	47.48	21.78	44.95	22.59	
YPL198W	0.95	1.46	1.58	1.64	1.74	2.03	1.99	1.99	1.99	6.96	0.96	4.497	21.64	55.86	22.60	
YCR066C	1.03	1.75	1.88	1.99	2.19	2.05	2.02	2.02	2.02	6.68	1.34	54.20	21.26	69.89	22.69	
YLR032C-A	0.94	1.34	1.50	1.59	1.76	2.00	1.93	1.93	1.93	7.51	1.03	51.28	21.11	73.86	22.81	
YNL086C	0.88	1.33	1.52	1.59	1.69	2.02	1.94	1.94	1.94	6.82	1.37	37.55	21.37	49.83	22.74	
YNL042W-B	0.88	1.29	1.48	1.57	1.67	1.97	1.93	1.93	1.93	5.96	1.05	41.49	21.69	47.44	22.75	
YLM202W	0.85	1.21	1.40	1.46	1.55	1.84	1.83	1.83	1.83	5.92	1.21	40.16	21.61	46.68	22.81	
YLU041W	0.88	1.25	1.44	1.52	1.62	1.72	1.72	1.72	1.72	4.71	1.41	45.10	21.43	49.81	22.84	
YGR035W-A	0.90	1.35	1.59	1.69	1.80	2.05	2.02	2.02	2.02	6.63	0.13	47.71	22.72	54.34	22.85	
YOL008W	0.86	1.18	1.31	1.40	1.51	1.76	1.72	1.72	1.72	4.55	1.50	39.56	21.07	50.49	23.24	
YKL121W-A	0.90	1.33	1.44	1.53	1.60	1.87	1.83	1.83	1.83	7.70	1.93	47.92	21.87	55.62	23.80	
YJR130C	0.95	1.30	1.38	1.38	1.47	1.76	1.74	1.74	1.74	6.85	1.90	56.15	22.72	63.95	23.80	
YLR149W	0.93	1.40	1.49	1.54	1.65											

ORF name	OD at 600 nm at sampling time points (hours)											GLUCOSE		FRUCTOSE		TOTAL SUGARS	
	0 h	20 h	27 h	44 h	66 h	93 h	113 h	120 h	146 h	120 h	146 h	120 h	146 h	120 h	146 h	120 h	146 h
YCL001W-A	0.91	1.45	1.50	1.74	1.85	2.04	2.26	6.23	0.98	45.62	23.16	52.85	24.15	52.85	24.15	52.85	24.15
YGL077C	0.92	1.46	1.51	1.67	1.78	2.05	2.28	6.31	1.10	45.70	23.38	52.88	24.16	52.88	24.16	52.88	24.16
YKL077C	0.91	1.19	1.31	1.35	1.43	1.68	1.83	1.16	0.67	3.89	24.69	50.07	24.76	50.07	24.76	50.07	24.76
YOR169W	0.93	1.51	1.66	1.71	1.78	1.87	2.02	7.04	1.08	42.64	23.89	48.68	24.76	48.68	24.76	48.68	24.76
YKL079W	0.92	1.59	1.75	1.77	1.88	2.13	2.05	6.68	1.12	51.02	23.84	57.71	24.77	57.71	24.77	57.71	24.77
YKL034W	0.88	1.30	1.47	1.53	1.64	1.89	1.83	7.24	2.18	45.84	22.63	52.88	24.81	52.88	24.81	52.88	24.81
YKL119W	0.92	1.42	1.56	1.62	1.72	2.01	1.94	5.53	1.07	42.60	23.75	48.13	24.82	48.13	24.82	48.13	24.82
YAB054W	0.89	1.33	1.45	1.43	1.65	1.85	1.80	9.96	1.92	47.94	23.01	57.90	24.93	57.90	24.93	57.90	24.93
YKL026C	0.91	1.31	1.45	1.48	1.61	1.90	1.81	10.58	1.52	50.89	23.42	61.47	24.94	61.47	24.94	61.47	24.94
YGR181W	0.95	1.40	1.54	1.60	1.72	1.99	1.92	6.96	1.07	50.81	23.88	57.77	24.95	57.77	24.95	57.77	24.95
YKL137W	0.91	1.37	1.50	1.57	1.67	1.98	1.92	6.02	0.30	42.91	24.65	48.94	24.95	48.94	24.95	48.94	24.95
YMR269W	0.89	1.32	1.48	1.59	1.68	1.96	1.92	7.52	1.60	48.45	23.36	55.97	24.96	55.97	24.96	55.97	24.96
YFR043C	0.89	1.34	1.49	1.51	1.64	1.93	1.83	9.92	1.85	59.78	23.12	69.70	24.97	69.70	24.97	69.70	24.97
YCL009C	0.89	1.49	1.71	1.71	1.82	2.07	2.10	7.14	1.17	51.38	23.83	58.52	25.00	58.52	25.00	58.52	25.00
YFL051C	0.90	1.39	1.54	1.58	1.71	2.01	1.92	7.68	1.44	53.23	23.58	60.90	25.02	60.90	25.02	60.90	25.02
YL221C	1.00	1.45	1.60	1.71	1.82	2.06	2.04	7.68	0.72	51.61	24.33	59.29	25.05	59.29	25.05	59.29	25.05
YNL056W	0.82	1.13	1.21	1.21	1.26	1.65	1.50	5.55	2.02	38.15	23.04	43.71	25.06	43.71	25.06	43.71	25.06
YKR015C	0.94	1.41	1.54	1.56	1.69	1.93	1.88	10.35	1.18	59.53	23.95	69.88	25.13	69.88	25.13	69.88	25.13
YPR117W	0.96	1.47	1.54	1.70	1.78	2.05	2.01	4.70	0.45	49.37	24.68	54.07	25.14	54.07	25.14	54.07	25.14
YGR097C	0.97	1.39	1.56	1.65	1.71	2.03	2.02	7.58	1.15	46.81	24.01	54.39	25.16	54.39	25.16	54.39	25.16
YHR157W	0.85	1.23	1.35	1.36	1.46	1.79	1.69	5.49	2.28	42.21	22.88	47.70	25.17	47.70	25.17	47.70	25.17
YKL124W	0.90	1.35	1.47	1.50	1.68	1.87	1.86	5.96	1.69	35.10	23.48	41.06	25.17	41.06	25.17	41.06	25.17
YDR420W	0.85	1.04	1.17	1.30	1.39	1.65	1.61	8.79	0.99	56.76	23.81	64.85	24.82	64.85	24.82	64.85	24.82
YPR125W	0.98	1.30	1.38	1.51	1.61	1.67	1.73	6.20	0.91	57.70	24.30	63.91	25.21	63.91	25.21	63.91	25.21
YIL164C	0.83	1.19	1.37	1.41	1.49	2.01	1.71	7.72	0.88	68.17	24.34	75.88	25.22	75.88	25.22	75.88	25.22
YHR189W	0.86	1.26	1.40	1.43	1.55	1.86	1.75	8.51	2.11	53.76	23.16	62.27	25.27	62.27	25.27	62.27	25.27
YER044C	0.94	1.29	1.51	1.66	1.82	2.09	2.04	8.40	2.21	52.64	23.14	61.05	25.35	61.05	25.35	61.05	25.35
YGR239W	0.90	1.25	1.40	1.46	1.56	1.83	1.77	7.09	1.55	47.95	23.81	54.78	25.36	54.78	25.36	54.78	25.36
YDL178W	0.87	1.40	1.59	1.64	1.75	2.03	2.03	7.15	1.14	50.85	24.23	56.00	25.37	56.00	25.37	56.00	25.37
YHL268C	0.83	1.23	1.40	1.49	1.59	1.83	1.75	6.45	1.42	45.43	23.95	51.88	25.37	51.88	25.37	51.88	25.37
YAR026C	0.91	1.48	1.64	1.71	1.87	2.04	2.05	8.25	1.23	61.21	23.71	67.53	25.38	67.53	25.38	67.53	25.38
YHL263C	0.86	1.12	1.26	1.33	1.39	1.54	1.52	7.68	1.52	43.42	23.73	49.84	25.38	49.84	25.38	49.84	25.38
YAL023W	0.91	1.42	1.53	1.55	1.65	1.96	1.93	7.11	2.11	53.62	23.23	54.43	25.39	54.43	25.39	54.43	25.39
YKL2027W	0.89	1.29	1.42	1.43	1.57	1.87	1.80	10.59	1.41	56.04	23.89	67.03	25.39	67.03	25.39	67.03	25.39
YNL080W	0.85	1.15	1.25	1.30	1.41	1.63	1.59	9.88	1.24	51.99	24.18	60.18	25.42	60.18	25.42	60.18	25.42
YGR016W	1.02	1.67	1.82	1.88	1.98	2.20	2.19	6.53	1.26	52.22	24.17	61.84	25.44	61.84	25.44	61.84	25.44
YGR038W	0.91	1.39	1.59	1.66	1.76	2.05	1.98	7.32	1.62	45.78	23.83	53.19	25.45	53.19	25.45	53.19	25.45
YLU062C	0.91	1.26	1.41	1.55	1.64	1.94	1.88	8.21	1.69	47.44	23.77	55.65	25.47	55.65	25.47	55.65	25.47
YKL096C	1.04	1.46	1.68	1.68	1.79	2.18	2.17	6.41	0.97	50.47	24.50	56.88	25.47	56.88	25.47	56.88	25.47
YKL097W	0.94	1.46	1.63	1.72	1.84	2.05	2.04	9.22	1.32	54.38	24.15	61.60	25.48	61.60	25.48	61.60	25.48
YGR056W	0.99	1.53	1.70	1.70	1.87	2.13	2.08	7.35	1.54	49.15	24.00	56.50	25.54	56.50	25.54	56.50	25.54
YLR144C	0.95	1.41	1.56	1.67	1.74	2.01	1.99	8.46	1.19	50.85	24.51	54.12	25.70	54.12	25.70	54.12	25.70
YKL027W	0.90	1.28	1.43	1.47	1.59	1.87	1.79	9.56	1.42	57.81	24.30	67.37	25.72	67.37	25.72	67.37	25.72
YKL039W	0.98	1.40	1.47	1.50	1.60	1.88	1.78	12.00	2.95	57.60	22.78	69.60	25.73	69.60	25.73	69.60	25.73
YER024C	0.93	1.45	1.57	1.65	1.75	2.04	1.90	10.39	1.03	52.45	24.56	62.84	25.59	62.84	25.59	62.84	25.59
YGR240C-A	0.96	1.37	1.51	1.55	1.61	1.92	1.82	8.22	1.65	49.52	24.40	57.75	25.65	57.75	25.65	57.75	25.65
YLL009W	0.90	1.24	1.35	1.36	1.45	1.76	1.64	10.00	2.00	52.25	24.21	62.34	25.75	62.34	25.75	62.34	25.75
YHR143C	0.87	1.21	1.34	1.40	1.51	1.76	1.70	8.41	1.44	45.74	24.28	52.16	25.69	52.16	25.69	52.16	25.69
YHR016C	0.91	1.23	1.39	1.46	1.58	1.80	1.78	10.86	2.05	50.00	24.00	58.47	25.44	58.47	25.44	58.47	25.44
YGR235C	0.99	1.56	1.69	1.73	1.82	2.08	2.03	7.42	1.35	50.09	24.80	57.51	25.15	57.51	25.15	57.51	25.15
YDL237W	0.93	1.29	1.45	1.54	1.65	1.86	1.84	7.95	1.65	54.42	24.53	62.37	25.37	62.37	25.37	62.37	25.37
YER023C	0.84	1.21	1.27	1.27	1.33	1.69	1.55	9.04	2.13	47.34	24.27	56.38	25.37	56.38	25.37	56.38	25.37
YLR265C	1.06	1.60	1.78	1.87	1.96	2.19	2.18	8.10	1.00	53.52	25.38	61.62	25.38	61.62	25.38	61.62	25.38
YPL192C	0.85	1.16	1.25	1.28	1.35	1.41	1.48	7.30	1.24	43.41	25.15	51.61	25.39	51.61	25.39	51.61	25.39
YGR286C	0.86	1.12	1.25	1.26	1.36	1.61	1.51	7.00	1.24	48.43	25.29	55.43	25.52	55.43	25.52	55.43	25.52
YPL264C	0.91	1.37	1.39	1.53	1.63	1.94	1.79	8.46	1.35	51.99	25.16	60.44	25.53	60.44	25.53	60.44	25.53
YHR061C	1.01	1.51	1.69	1.76	1.87	2.10	2.09	9.37	1.54	53.74	25.01	65.10	25.55	65.10	25.55	65.10	25.55
YKL116W	0.98	1.50	1.66	1.74	1.85	2.03	1.97	7.56	1.55	52.62	25.08	60.66	25.63	60.66	25.63	60.66	25.63
YPL020C	0.91	1.36	1.52	1.59	1.70	1.90	1.87	7.56	1.55	52.65	25.05	61.86	25.68</td				

ORF name	OD at 600 nm at sampling time points (hours)											GLUCOSE		FRUCTOSE		TOTAL SUGARS		
	0 h	20 h	27 h	44 h	66 h	93 h	113 h	120 h	146 h	120 h	146 h	120 h	146 h	120 h	146 h	120 h	146 h	
YPL014W	0.99	1.00	1.00	1.04	1.00	2.04	2.06	6.10	1.00	4.13	2.04	25.88	51.90	27.35	25.88	51.90	27.35	
YGL124W	0.93	1.46	1.56	1.78	2.05	1.99	6.24	1.44	5.13	2.52	65.26	21.65	55.92	21.65	55.92	21.65		
YDR434C	0.98	1.15	1.34	1.45	1.55	1.79	1.77	1.16	1.53	6.03	2.55	53.85	15.59	21.38	53.85	15.59	21.38	
YLR164W	0.95	1.31	1.50	1.59	1.68	1.65	1.89	1.60	0.88	4.03	26.58	56.46	27.38	26.58	56.46	27.38		
YER019C-A	0.88	1.00	1.13	1.20	1.26	1.53	1.45	6.87	1.69	5.23	25.71	57.09	27.40	57.09	27.40	57.09	27.40	
YLR053C	0.89	1.38	1.53	1.61	1.71	2.01	1.94	8.53	1.70	4.93	25.74	58.46	27.44	25.74	58.46	27.44	25.74	58.46
YPL041C	0.89	1.34	1.60	1.69	1.79	2.04	1.99	8.55	1.60	4.81	25.85	57.67	27.45	25.85	57.67	27.45	25.85	57.67
YDR436W	0.95	1.46	1.59	1.69	1.78	2.01	2.00	8.22	1.73	5.24	25.74	61.47	27.47	25.74	61.47	27.47	25.74	61.47
YER033C	0.91	1.44	1.59	1.64	1.75	2.03	1.95	7.92	1.61	5.97	25.86	61.79	27.47	25.86	61.79	27.47	25.86	61.79
YMR234W	0.94	1.43	1.60	1.71	1.82	2.06	2.06	7.71	1.30	4.91	26.17	56.90	27.47	26.17	56.90	27.47	26.17	56.90
YGR197C	0.93	1.33	1.49	1.54	1.65	1.96	1.86	7.29	1.58	4.76	25.90	54.46	27.48	25.90	54.46	27.48	25.90	54.46
YBR129C	1.01	1.47	1.62	1.72	1.87	2.06	2.04	9.27	1.18	7.54	26.32	82.21	27.50	26.32	82.21	27.50	26.32	82.21
YBR195C	0.90	1.29	1.37	1.54	1.64	1.93	1.89	7.67	1.28	5.18	25.23	58.15	27.51	25.23	58.15	27.51	25.23	58.15
YLR214W	1.03	1.57	1.72	1.78	1.88	2.13	2.09	8.70	1.78	5.12	25.75	59.90	27.53	25.75	59.90	27.53	25.75	59.90
YDL177W	0.95	1.36	1.60	1.85	1.85	2.00	1.98	8.89	1.11	6.03	26.44	59.52	27.55	26.44	59.52	27.55	26.44	59.52
YLR151C	0.88	1.29	1.44	1.55	1.64	1.96	1.88	9.14	1.51	5.13	26.06	60.67	27.57	26.06	60.67	27.57	26.06	60.67
YMR130W	0.97	1.47	1.59	1.65	1.71	1.87	1.96	9.29	1.80	4.16	26.25	50.54	27.62	26.25	50.54	27.62	26.25	50.54
YJR111C	0.85	1.16	1.34	1.47	1.55	1.81	1.78	6.96	1.90	4.66	25.75	53.61	27.65	25.75	53.61	27.65	25.75	53.61
YHL033C	0.86	1.14	1.29	1.34	1.50	1.68	1.65	7.79	1.04	6.18	26.61	69.64	27.65	26.61	69.64	27.65	26.61	69.64
YKL133C	0.92	148	1.61	1.45	1.72	1.85	1.89	9.13	1.58	5.17	26.09	69.92	27.67	26.09	69.92	27.67	26.09	69.92
YDL226C	0.89	1.32	1.48	1.58	1.68	1.99	2.00	11.33	1.40	6.15	26.35	72.87	27.75	26.35	72.87	27.75	26.35	72.87
YPR127W	0.95	1.33	1.43	1.58	1.68	1.76	1.85	7.26	1.43	5.80	26.32	62.28	27.75	26.32	62.28	27.75	26.32	62.28
YAR029W	0.89	1.39	1.54	1.47	1.76	1.97	1.96	6.68	2.05	3.67	25.71	43.60	27.75	25.71	43.60	27.75	25.71	43.60
YNL194C	0.88	1.23	1.35	1.43	1.54	1.75	1.78	1.75	1.09	1.45	26.37	62.32	27.77	26.37	62.32	27.77	26.37	62.32
YLR221W	0.91	1.40	1.55	1.61	1.75	2.02	1.95	9.42	1.60	5.25	26.20	61.96	27.80	26.20	61.96	27.80	26.20	61.96
YIL016W	0.92	1.39	1.50	1.55	1.68	1.96	1.88	7.88	2.89	5.17	24.97	59.45	27.86	24.97	59.45	27.86	24.97	59.45
YDL144C	0.92	1.43	1.56	1.68	1.80	2.03	2.06	6.43	1.12	5.67	26.76	62.09	27.88	26.76	62.09	27.88	26.76	62.09
YEL048C	0.92	1.36	1.51	1.59	1.72	1.92	1.88	2.06	4.67	28.17	23.22	50.23	27.89	23.22	50.23	27.89	23.22	50.23
YFL053W	0.95	1.48	1.62	1.66	1.77	2.05	1.98	9.04	1.36	5.64	26.53	65.50	27.89	26.53	65.50	27.89	26.53	65.50
YKL218C	1.00	1.56	1.70	1.74	1.88	2.11	2.06	9.95	1.80	5.13	26.12	61.58	27.92	26.12	61.58	27.92	26.12	61.58
YLG021W	0.86	1.37	1.49	1.54	1.67	1.89	1.86	9.65	1.50	5.18	26.44	62.22	27.94	26.44	62.22	27.94	26.44	62.22
YDR447W	0.87	1.25	1.39	1.51	1.61	1.81	1.93	9.93	1.52	5.14	26.80	63.45	27.95	26.80	63.45	27.95	26.80	63.45
YER031W	0.87	1.28	1.45	1.54	1.67	1.86	1.84	9.44	1.50	5.87	26.47	58.21	27.97	26.47	58.21	27.97	26.47	58.21
YER032C	0.86	1.22	1.30	1.40	1.58	1.88	1.84	9.11	1.55	5.69	26.82	59.67	27.97	26.82	59.67	27.97	26.82	59.67
YER022C	0.84	1.23	1.32	1.37	1.42	1.77	1.86	8.84	2.38	4.08	25.59	52.92	27.97	25.59	52.92	27.97	25.59	52.92
YGL129C	0.91	145	1.59	1.61	1.70	1.97	1.99	7.72	1.69	5.24	25.98	68.65	27.97	25.98	68.65	27.97	25.98	68.65
YHR159W	0.81	1.34	1.49	1.53	1.65	1.93	1.85	6.91	2.30	5.14	25.73	58.31	28.03	25.73	58.31	28.03	25.73	58.31
YMR313C	0.91	1.37	1.53	1.64	1.74	2.00	1.97	7.89	1.82	4.12	26.15	52.11	28.06	26.15	52.11	28.06	26.15	52.11
YKL068W	0.83	1.61	1.66	1.76	1.83	2.00	1.97	7.89	1.54	4.27	26.50	69.85	28.08	26.50	69.85	28.08	26.50	69.85
YER221W	0.92	1.47	1.65	1.71	1.86	2.06	2.07	6.75	1.15	5.04	26.59	57.00	28.14	26.59	57.00	28.14	26.59	57.00
YPL017C	0.94	1.42	1.53	1.67	1.79	1.93	1.96	8.41	1.15	4.85	27.02	58.96	28.18	27.02	58.96	28.18	27.02	58.96
YER032C	0.89	1.48	1.62	1.65	1.76	2.02	1.98	6.00	1.73	4.87	26.45	54.56	28.18	26.45	54.56	28.18	26.45	54.56
YLL020C	0.88	1.18	1.28	1.37	1.38	1.69	1.67	9.17	2.27	5.44	26.15	63.85	28.42	26.15	63.85	28.42	26.15	63.85
YDR442W	0.96	1.39	1.57	1.61	1.76	2.02	2.02	7.66	1.81	5.18	26.61	54.49	28.43	26.61	54.49	28.43	26.61	54.49
YER054C	0.96	1.59	1.77	1.85	1.92	2.15	2.13	7.25	1.61	5.05	26.66	62.25	28.27	26.66	62.25	28.27	26.66	62.25
YPL162C	0.87	1.24	1.37	1.47	1.58	1.66	1.70	8.22	1.29	26.00	58.99	28.29	26.00	58.99	28.29	26.00	58.99	28.29
YGR076C	0.89	1.24	1.38	1.43	1.57	1.80	1.75	8.72	2.24	45.33	26.07	52.05	28.30	26.07	52.05	28.30	26.07	52.05
YPL244C	0.90	1.39	1.47	1.56	1.65	1.91	1.80	10.82	1.79	5.49	26.67	63.31	28.46	26.67	63.31	28.46	26.67	63.31
YER026C	0.88	1.22	1.33	1.48	1.56	1.81	1.76	9.02	1.97	5.10	26.72	63.60	28.46	26.72	63.60	28.46	26.72	63.60
YER032C	0.91	1.33	1.48	1.50	1.66	1.98	1.95	9.08	1.50	5.15	26.85	62.94	28.06	26.85	62.94	28.06	26.85	62.94
YER239W	0.91	1.43	1.56	1.62	1.75	1.98	1.93	11.42	2.22	5.87	26.85	62.94	28.06	26.85	62.94	28.06	26.85	62.94
YER011W	0.91	1.43	1.57	1.62	1.72	2.02	1.94	7.20	1.14	5.10	27.65	58.49	28.78	27.65	58.49	28.78	27.65	58.49
YER095C	0.84	1.64	1.81	1.86	1.93	2.01	1.94	9.18	1.51	5.24	27.34	59.59	28.79	27.34	59.59	28.79	27.34	59.59
YLR079W	1.00	1.47	1.63	1.71	1.81	2.02	1.99	7.23	2.51	45.10	28.30	52.33	28.81	28.30	52.33	28.81	28.30	52.33
YKL026C	0.85	1.11	1.19	1.20	1.30	1.51	1.51	5.91	1.32	4.26	26.55	58.40	28.75	26.55	58.40	28.75	26.55	58.40
YER016W	0.92	1.41	1.62	1.71	1.88	2.09	2.12	5.72	1.66	4.79	27.14	55.83	28.74	27.14	55.83	28.74	27.14	55.83
YLR121W	0.98	1.31	1.43	1.51	1.60													

ORF name	OD at 600 nm at sampling time points (hours)											GLUCOSE		FRUCTOSE		TOTAL SUGARS	
	0 h	20 h	27 h	44 h	66 h	93 h	113 h	120 h	146 h	120 h	146 h	120 h	146 h	120 h	146 h	120 h	146 h
YKL140C	0.97	1.50	1.69	1.72	1.81	2.13	2.05	6.61	1.00	2.91	2.13	5.35	2.69	6.33	30.19	30.19	30.19
YIL102W	0.85	1.21	1.38	1.36	1.46	1.76	1.68	6.78	2.89	5.45	2.75	6.37	3.01	5.63	32.24	32.24	32.24
YKL188C	0.95	1.13	1.30	1.33	1.45	1.73	1.68	6.63	1.63	7.94	2.86	5.63	2.97	7.14	30.25	30.25	30.25
YKL190C	1.02	1.44	1.60	1.60	1.70	1.97	1.90	5.24	1.28	6.51	2.87	6.35	2.97	7.14	30.25	30.25	30.25
YPL001W	0.94	1.45	1.54	1.70	1.82	2.04	2.03	6.76	2.01	4.73	2.25	5.49	2.36	5.49	30.26	30.26	30.26
YOR232W	0.90	1.50	1.60	1.68	1.82	1.94	2.08	6.82	1.00	3.68	2.28	4.59	2.36	4.59	30.28	30.28	30.28
YER034W	0.97	1.52	1.68	1.78	1.94	2.09	2.08	6.13	2.95	2.32	27.33	42.44	30.28	42.44	30.28	42.44	30.28
YER062W	0.98	1.39	1.54	1.54	1.63	1.91	1.80	7.19	1.16	5.68	29.12	66.78	30.28	66.78	30.28	66.78	30.28
YER025C	1.00	1.61	1.82	1.84	1.95	2.13	2.14	6.41	2.41	4.90	27.90	56.01	30.31	49.60	30.31	49.60	30.31
YPL156C	0.88	1.20	1.30	1.35	1.44	1.51	1.57	6.84	1.88	24.49	28.43	33.33	30.31	33.33	30.31	33.33	30.31
YLR172C	1.03	1.61	1.79	1.89	1.94	2.11	2.15	8.50	1.48	4.64	28.84	54.97	30.32	54.97	30.32	54.97	30.32
YNL173C	0.85	1.20	1.34	1.44	1.59	1.80	1.77	11.19	1.65	6.23	28.69	74.12	30.35	74.12	30.35	74.12	30.35
YHR226W	0.88	1.23	1.33	1.66	1.76	1.78	1.76	5.96	1.92	6.08	28.43	70.95	30.35	70.95	30.35	70.95	30.35
YMR224W	0.85	1.20	1.30	1.33	1.38	1.71	1.61	8.03	2.29	4.24	28.08	52.27	30.37	52.27	30.37	52.27	30.37
YKL077W	0.89	1.23	1.39	1.45	1.56	1.83	1.76	9.39	2.82	54.20	27.58	63.59	30.40	63.59	30.40	63.59	30.40
YHR149C	0.96	1.40	1.65	1.78	1.98	1.72	1.98	7.69	1.55	5.63	28.87	64.62	30.42	64.62	30.42	64.62	30.42
YPL137C	0.85	1.30	1.39	1.44	1.53	1.62	1.71	7.49	1.47	3.10	28.96	35.99	30.43	35.99	30.43	35.99	30.43
YKR064C	0.89	1.41	1.56	1.55	1.72	1.97	1.97	12.23	2.23	55.80	28.22	68.04	30.46	68.04	30.46	68.04	30.46
YGL134W	0.91	1.45	1.60	1.65	1.77	2.05	1.97	9.31	1.68	59.27	28.78	68.59	30.46	68.59	30.46	68.59	30.46
YJR040W	0.94	1.38	1.50	1.53	1.63	1.89	1.82	7.58	2.17	51.43	28.30	59.01	30.47	59.01	30.47	59.01	30.47
YDL188C	0.94	1.27	1.46	1.48	1.53	1.81	1.73	11.98	1.81	6.26	28.69	74.94	30.50	74.94	30.50	74.94	30.50
YPR138C	0.99	1.43	1.53	1.61	1.68	1.97	1.92	10.34	1.65	6.01	28.87	70.35	30.52	70.35	30.52	70.35	30.52
YDL093W	0.89	1.45	1.62	1.73	1.84	2.05	2.08	6.02	1.38	56.58	29.16	64.61	30.54	64.61	30.54	64.61	30.54
YJL013C	0.90	1.43	1.49	1.59	1.70	1.97	1.97	9.48	2.91	50.15	27.69	56.63	30.60	56.63	30.60	56.63	30.60
YMR226C	0.98	1.55	1.68	1.76	1.86	1.94	2.11	8.97	2.18	52.60	28.43	61.58	30.61	61.58	30.61	61.58	30.61
YLR165C	1.03	1.63	1.80	1.90	1.99	2.19	2.18	8.98	1.42	43.31	29.19	58.29	30.61	58.29	30.61	58.29	30.61
YLM053C	0.92	1.58	1.74	1.76	1.88	2.08	2.03	6.98	1.31	62.24	29.31	73.21	30.62	73.21	30.62	73.21	30.62
YJL110C	1.01	1.56	1.74	1.80	1.88	2.16	2.09	9.01	1.48	60.99	29.14	70.00	30.62	70.00	30.62	70.00	30.62
YKL097C	0.93	1.45	1.56	1.61	1.73	1.97	1.91	10.55	3.28	57.19	27.35	67.73	30.63	67.73	30.63	67.73	30.63
YKR065C	0.86	1.28	1.42	1.46	1.60	1.85	1.71	12.28	1.83	56.73	28.81	71.01	30.64	71.01	30.64	71.01	30.64
YHL022C	0.92	1.35	1.52	1.59	1.71	2.00	1.93	7.26	1.99	42.91	28.66	56.11	30.65	56.11	30.65	56.11	30.65
YPL071C	0.91	1.25	1.42	1.46	1.59	1.77	1.70	5.76	1.51	51.66	29.10	63.59	30.67	63.59	30.67	63.59	30.67
YPL082C	0.93	1.50	1.64	1.74	1.85	1.95	2.01	9.66	1.44	59.33	28.84	61.78	30.68	61.78	30.68	61.78	30.68
YKL198C	0.81	1.18	1.35	1.38	1.51	1.84	1.74	6.31	1.76	56.47	28.95	65.78	30.71	65.78	30.71	65.78	30.71
YKL083W	0.95	1.42	1.52	1.59	1.69	1.97	1.89	6.38	1.56	52.98	27.16	61.36	30.72	61.36	30.72	61.36	30.72
YGL230C	0.98	1.41	1.60	1.71	1.83	2.11	2.01	11.23	1.58	7.24	28.14	83.37	30.73	83.37	30.73	83.37	30.73
YER069C	0.85	1.21	1.38	1.47	1.56	1.85	1.79	7.53	2.30	43.63	28.45	51.17	30.74	51.17	30.74	51.17	30.74
YER134C	0.94	1.42	1.60	1.69	1.79	2.05	2.02	6.96	2.63	45.94	28.13	52.99	30.76	52.99	30.76	52.99	30.76
YDL078C	0.86	1.35	1.56	1.54	1.64	1.96	1.96	9.24	1.54	6.06	29.25	72.29	30.78	72.29	30.78	72.29	30.78
YGL232W	0.85	1.28	1.43	1.48	1.59	1.91	1.82	8.54	1.65	56.68	28.14	65.22	30.79	65.22	30.79	65.22	30.79
YIL123W	0.90	1.20	1.33	1.51	1.75	1.79	1.68	14.35	2.45	61.93	28.34	81.28	30.80	81.28	30.80	81.28	30.80
YAL036C	0.90	1.55	1.68	1.53	1.80	1.96	1.94	7.21	2.19	39.15	28.61	45.36	30.80	45.36	30.80	45.36	30.80
YDR165W	0.93	1.37	1.59	1.50	1.88	1.97	1.97	8.46	2.02	58.70	28.78	67.17	30.80	67.17	30.80	67.17	30.80
YNL309W	0.99	1.51	1.69	1.73	1.84	2.12	2.04	12.49	1.42	75.15	29.40	87.63	30.82	87.63	30.82	87.63	30.82
YNL031C	0.85	1.14	1.26	1.27	1.34	1.68	1.56	9.46	2.42	46.83	28.40	58.30	30.82	58.30	30.82	58.30	30.82
YLR270W	1.01	1.51	1.64	1.70	1.79	2.06	2.02	6.72	0.90	57.44	29.92	64.16	30.82	64.16	30.82	64.16	30.82
YRL108W	0.90	1.31	1.45	1.51	1.64	1.93	1.87	10.86	2.37	58.61	28.49	69.47	30.86	69.47	30.86	69.47	30.86
YLR251W	1.03	1.55	1.66	1.72	1.82	2.04	2.05	10.95	1.83	51.10	29.04	62.62	30.90	62.62	30.90	62.62	30.90
YDR470C	0.92	1.33	1.53	1.59	1.73	2.00	1.95	7.92	1.38	62.92	29.52	70.84	30.90	70.84	30.90	70.84	30.90
YER287W	0.93	1.32	1.40	1.41	1.49	1.69	1.69	8.77	2.40	61.80	29.34	72.01	30.95	72.01	30.95	72.01	30.95
YPR044W	0.87	1.28	1.42	1.41	1.62	1.81	1.79	7.22	1.88	41.30	29.25	45.82	31.13	45.82	31.13	45.82	31.13
YDR020W	0.95	1.36	1.53	1.56	1.66	1.94	1.84	7.70	1.32	57.60	28.86	65.29	31.16	65.29	31.16	65.29	31.16
YPL258W	0.88	1.31	1.49	1.54	1.64	1.94	1.57	7.00	2.05	54.81	29.19	64.50	31.24	64.50	31.24	64.50	31.24
YHR199C	0.97	1.27	1.44	1.52	1.64	1.97	1.74	7.22	1.42	51.02	29.42	61.52	31.26	61.52	31.26	61.52	31.26
YER073W	0.97	1.20	1.38	1.37	1.48	1.79	1.65	8.44	2.26	51.67	28.09	62.11	31.28	62.11	31.28	62.11	31.28
YER070C	0.97	1.27	1.44	1.41	1.51	1.74	1.64	8.03	2.11	52.18	29.63	59.89	31.59	59.89	31.59	59.89	31.59
YER045W	0.94	1.31	1.51	1.53	1.63	2.01	1.95	7.70	2.16	53.80	29.56	67.50	31.73	67.50	31.73	67.50	31.73
YER032C	0.94	1.45	1.59	1.63	1.76	2.01	1.95	7.70	2.16	53.80	29.56	67.50	31.73	67.50	31.73	67.50	31.73
YPR066C	0.90	1.31	1.31	1.44	1.57	1.66	1.66	11.34	2.25	58.91	29.51	70.24	31.77	70.24	31.77	70.24	31.7

ORF name	OD at 600 nm at sampling time points (hours)											GLUCOSE		FRUCTOSE		TOTAL SUGARS		
	0 h	20 h	27 h	44 h	66 h	93 h	113 h	120 h	146 h	120 h	146 h	120 h	146 h	120 h	146 h	120 h	146 h	
YHR102W	0.89	1.04	1.35	1.72	2.04	1.99	1.72	2.36	5.25	3.07	6.21	3.25	3.25	3.07	6.21	3.25	3.25	
YKR025W	0.93	1.35	1.54	1.56	1.65	1.97	1.54	1.66	1.11	6.45	3.18	7.91	3.27	3.27	6.45	7.91	3.27	3.27
YMR28C	0.97	1.26	1.49	1.43	1.43	1.50	1.86	1.81	2.27	5.13	3.39	5.14	3.23	3.39	5.14	3.23	3.23	
YPL163W	0.90	1.36	1.40	1.41	1.51	1.60	1.69	1.77	3.22	4.81	3.49	5.24	3.53	3.49	5.24	3.53	3.53	
YHR214W-A	0.90	1.24	1.42	1.45	1.64	1.94	1.84	1.85	1.04	6.98	3.19	6.73	3.24	3.19	6.73	3.24	3.24	
YIL018W	0.92	1.25	1.51	1.72	1.73	1.79	1.68	1.88	2.44	6.26	3.13	8.14	3.27	3.13	8.14	3.27	3.27	
YBL103W	0.92	1.27	1.42	1.57	1.67	1.94	1.84	1.97	2.20	6.59	3.39	7.71	3.29	3.39	7.71	3.29	3.29	
YNL070W	0.94	1.42	1.60	1.70	1.75	2.03	1.98	1.49	2.00	5.94	3.60	7.42	3.29	3.60	7.42	3.29	3.29	
YAL013W	0.90	1.34	1.49	1.52	1.71	1.90	1.89	1.52	2.29	5.49	3.30	6.50	3.26	3.30	6.50	3.26	3.26	
YLR073C	0.86	1.18	1.39	1.48	1.57	1.86	1.82	1.44	1.77	5.77	3.83	6.72	3.26	3.83	6.72	3.26	3.26	
YPR023W	1.06	1.61	1.77	1.87	1.96	2.13	2.14	1.79	1.46	5.39	3.17	6.48	3.23	3.17	6.48	3.23	3.23	
YLR032W	0.91	1.32	1.43	1.48	1.57	1.90	1.79	1.04	1.82	5.18	3.03	6.22	3.25	3.03	6.22	3.25	3.25	
YLR281C	1.04	1.67	1.82	1.89	1.99	2.17	2.18	1.80	1.95	4.93	3.07	5.15	3.26	3.07	5.15	3.26	3.26	
YNL041C	0.82	1.11	1.30	1.35	1.44	1.76	1.70	1.34	2.78	5.52	2.89	5.86	3.27	2.89	5.86	3.27	3.27	
YKL214C	0.92	1.38	1.52	1.58	1.69	1.99	1.92	1.03	2.47	5.64	3.22	6.67	3.26	3.22	6.67	3.26	3.26	
YPL008W	0.95	1.44	1.50	1.65	1.74	1.93	1.94	1.80	2.06	4.95	3.03	5.65	3.29	3.03	5.65	3.29	3.29	
YPR179C	0.91	1.33	1.43	1.47	1.56	1.86	1.79	1.26	1.80	5.58	3.05	6.42	3.25	3.05	6.42	3.25	3.25	
YMR273C	0.92	1.34	1.50	1.51	1.61	1.94	1.81	1.51	2.11	5.17	3.04	7.68	3.27	3.04	7.68	3.27	3.27	
YKL140W	0.93	1.50	1.64	1.70	1.82	2.07	2.04	1.82	1.19	5.85	3.15	6.84	3.26	3.15	6.84	3.26	3.26	
YHR163W	0.92	1.35	1.51	1.56	1.68	1.93	1.87	1.76	2.29	5.51	3.01	6.27	3.28	3.01	6.27	3.28	3.28	
YLLO056C	0.92	1.37	1.55	1.65	1.77	1.99	1.98	1.16	2.15	5.69	3.06	6.26	3.28	3.06	6.26	3.28	3.28	
YDL088C	0.82	1.22	1.42	1.29	1.42	1.74	1.70	1.66	1.41	4.97	3.10	5.63	3.28	3.10	5.63	3.28	3.28	
YLR011W	0.83	1.15	1.30	1.37	1.48	1.77	1.77	1.11	2.20	5.63	3.03	6.74	3.23	3.03	6.74	3.23	3.23	
YJR130C	0.91	1.46	1.63	1.65	1.79	2.02	2.07	1.16	2.42	4.93	3.01	5.30	3.23	3.01	5.30	3.23	3.23	
YGL222C	0.90	1.34	1.48	1.53	1.65	1.94	1.85	1.79	1.90	5.68	3.03	6.54	3.23	3.03	6.54	3.23	3.23	
YKL174C	0.91	1.43	1.60	1.69	1.81	2.09	2.05	1.82	1.54	5.17	3.13	6.13	3.24	3.13	6.13	3.24	3.24	
YMR215W	0.86	1.33	1.51	1.54	1.61	1.93	1.86	1.32	2.14	6.76	3.07	7.78	3.26	3.07	7.78	3.26	3.26	
YHR039C-A	0.86	1.18	1.35	1.51	1.61	1.95	1.90	1.77	1.98	5.86	3.09	7.65	3.27	3.09	7.65	3.27	3.27	
YLR028C	0.93	1.39	1.59	1.59	1.67	1.98	1.96	1.09	2.49	5.87	3.09	6.93	3.28	3.09	6.93	3.28	3.28	
YER041W	0.99	1.69	1.77	1.87	1.94	2.17	2.14	1.67	2.31	7.02	3.62	8.93	3.29	3.62	8.93	3.29	3.29	
YDL099W	0.88	1.41	1.59	1.61	1.71	1.97	1.95	1.77	2.31	6.23	3.17	6.81	3.24	3.17	6.81	3.24	3.24	
YMR140C	0.93	1.42	1.55	1.62	1.72	1.96	1.95	1.95	2.31	6.23	3.27	7.58	3.24	3.27	7.58	3.24	3.24	
YPL260W	0.85	1.40	1.57	1.62	1.74	1.92	1.93	1.53	2.06	6.54	3.04	7.44	3.25	3.04	7.44	3.25	3.25	
YMR144C	0.91	1.56	1.88	1.91	1.91	2.03	2.01	1.56	2.01	6.96	3.27	7.98	3.28	3.27	7.98	3.28	3.28	
YLR253W	1.05	1.59	1.75	1.87	1.97	2.17	2.17	1.62	2.54	6.84	3.36	7.86	3.30	3.36	7.86	3.30	3.30	
YBL206W	0.90	1.34	1.58	1.49	1.59	1.81	1.77	1.51	2.05	6.77	3.12	7.98	3.22	3.12	7.98	3.22	3.22	
YKL220C	0.98	1.49	1.59	1.70	1.83	2.02	2.03	1.81	2.52	7.42	3.51	7.53	3.32	3.51	7.53	3.32	3.32	
YPR015C	0.95	1.60	1.77	1.78	1.87	2.14	2.06	1.87	1.74	6.72	3.18	7.59	3.30	3.18	7.59	3.30	3.30	
YPL134W	0.91	1.36	1.49	1.60	1.68	1.96	1.90	1.25	2.05	6.97	3.13	8.98	3.23	3.13	8.98	3.23	3.23	
YLR236C	1.12	1.83	1.97	2.04	2.14	2.30	2.30	1.97	1.56	6.40	3.14	7.74	3.31	3.14	7.74	3.31	3.31	
YGL257C	0.91	1.39	1.52	1.59	1.71	1.97	1.91	1.51	2.04	6.85	3.17	7.82	3.24	3.17	7.82	3.24	3.24	
YLU001W	0.87	1.26	1.39	1.46	1.55	1.88	1.79	1.83	1.87	5.02	3.27	5.85	3.34	3.27	5.85	3.34	3.34	
YDL100C	0.91	1.38	1.59	1.58	1.70	1.95	1.97	1.20	2.08	6.88	3.19	8.09	3.31	3.19	8.09	3.31	3.31	
YNL230C	0.92	1.35	1.55	1.63	1.70	1.81	1.98	1.81	2.21	6.81	3.21	7.81	3.26	3.21	7.81	3.26	3.26	
YIL045W	0.84	1.18	1.29	1.29	1.36	1.70	1.57	1.02	2.34	5.48	3.05	6.19	3.19	3.05	6.19	3.19	3.19	
YLR319C	1.01	1.44	1.55	1.59	1.65	1.90	1.90	1.12	2.40	6.23	3.08	7.47	3.23	3.08	7.47	3.23	3.23	
YER015C-A	0.95	1.46	1.63	1.63	1.77	1.98	1.89	1.46	2.59	6.55	3.28	10.81	3.32	3.28	10.81	3.32	3.32	
YLR070C	0.87	1.43	1.65	1.67	1.78	2.09	2.09	9.97	1.40	5.16	3.26	8.08	6.21	3.26	8.08	6.21	3.26	
YDR313C	0.81	1.04	1.19	1.20	1.31	1.62	1.62	1.54	2.07	6.71	3.01	6.91	3.22	3.01	6.91	3.22	3.22	
YPL147W	0.91	1.35	1.46	1.50	1.60	1.67	1.73	1.73	2.30	6.49	3.14	7.81	3.24	3.14	7.81	3.24	3.24	
YLR044C	0.88	1.41	1.60	1.60	1.82	2.02	1.99	7.85	2.14	6.03	3.17	7.22	3.25	3.17	7.22	3.25	3.25	
YHR100C	1.00	1.52	1.67	1.86	1.96	2.15	2.21	13.97	2.47	6.89	3.09	8.28	3.36	3.09	8.28	3.36	3.36	
YKL166C	0.90	1.45	1.62	1.68	1.76	2.06	1.97	12.44	1.83	5.75	3.25	6.68	3.36	3.25	6.68	3.36	3.36	
YLR078W	0.91	1.42	1.54	1.61	1.73	1.98	1.94	1.20	2.28	6.87	3.10	7.28	3.33	3.10	7.28	3.33	3.33	
YER079C-A	0.89	1.14	1.28	1.46	1.56	1.81	1.77	1.01	2.05	6.97	3.17	8.03	3.32	3.17	8.03	3.32	3.32	
YAR033W	0.92	1.39	1.53	1.57	1.62	1.92	1.92	1.81	2.40	6.23	3.17	7.47	3.32	3.17	7.47	3.32	3.32	
YLR023C	0.93	1.50	1.64	1.64	1.76	1.94	1.86	1.73	2.31	6.84	3.22	7.81	3.34	3.22	7.81	3.34	3.34	
YMR020W	0.99	1.49	1.61	1.61	1.74	1.90	1.90	1.06	2.14	6.86	3.22	7.86	3.37	3.22	7.86	3.37	3.37	
YER046C	0.90	1.22	1.38	1.46	1.59	1.83	1.86	1.05	2.22	6.86	3.22	7.86	3.37	3.22	7.86	3.37	3.37	
YPL285C	0.95	1.35	1.51	1.58	1.78	1.98	1.92	1.83	2.56	6.84	3.21	7.86	3.37	3.21	7.86	3.37	3.37	
YMR229W	0.88	1.29	1.48	1.56	1.66	1.94	1.88	1.88	2.05	6.84	3.21	7.76	3.34	3.21	7.76	3.34	3.34	
YPL229W	1.06	1.75	1.90	1.96	2.06	2.23	2.22	2.08	1.56	6.23								

ORF name	20 h	at 60 min time points (hours)						120 h	146 h	120 h	146 h	TOTAL SUGARS
		20 h	27 h	44 h	61 h	93 h	113 h					
YMR286W	0.88	1.29	1.36	1.52	1.67	1.76	1.70	10.61	2.75	49.37	31.73	58.24
YMR262W	0.88	1.19	1.33	1.52	1.67	1.76	1.70	10.61	2.75	49.37	31.73	58.24
YGL026C	0.93	1.44	1.59	1.63	1.74	1.92	1.94	10.05	2.53	50.70	32.07	57.21
YLL023C	1.02	1.50	1.65	1.72	1.78	2.00	1.96	9.13	3.34	74.45	31.27	83.57
YGL424C	0.81	1.22	1.37	1.45	1.56	1.88	1.80	6.59	1.45	48.65	32.77	53.44
YNL045W	0.92	1.42	1.60	1.68	1.77	1.77	2.07	2.04	7.77	2.87	48.41	31.80
YLT12W	0.86	1.26	1.49	1.65	1.76	2.02	1.93	13.23	2.57	66.85	32.12	80.08
YGR259C	0.87	1.21	1.36	1.42	1.52	1.79	1.73	8.27	2.91	49.23	32.68	57.50
YDR273W	0.91	1.32	1.61	1.59	1.62	1.98	1.88	9.82	1.32	74.93	33.39	84.55
YKL146W	0.90	1.37	1.51	1.57	1.65	1.96	1.89	11.40	1.92	57.79	32.94	69.19
YHR037W	0.88	1.20	1.35	1.49	1.59	1.90	1.85	10.97	2.48	57.79	32.30	68.75
YKL032C	1.01	1.44	1.60	1.61	1.69	1.89	1.84	12.51	2.40	69.26	32.39	81.87
YKL134C	0.98	1.46	1.63	1.75	1.84	2.07	2.09	7.33	2.41	52.78	32.41	60.11
YMR227C	0.86	1.14	1.28	1.34	1.45	1.66	1.61	9.62	2.60	77.62	32.24	82.25
YLG060W	0.88	1.34	1.49	1.52	1.64	1.93	1.84	8.37	1.88	59.29	32.98	67.67
YHR080C	0.84	1.05	1.26	1.41	1.51	1.80	1.75	11.48	2.42	66.24	32.44	77.74
YER003C	1.08	1.74	1.90	1.99	2.08	2.28	2.28	8.33	1.92	62.84	32.98	71.18
YLR224W	1.06	1.60	1.70	1.76	1.85	2.07	2.05	10.74	1.88	57.34	33.03	68.08
YHL035C	0.93	1.28	1.49	1.76	1.71	1.90	1.85	11.64	2.69	67.21	32.24	78.85
YPR189W	0.90	1.29	1.39	1.44	1.51	1.83	1.75	9.22	2.32	56.42	32.62	65.65
YHR039C	0.92	1.32	1.49	1.66	1.75	2.04	1.99	13.73	2.69	69.83	32.27	83.56
YDR533C	0.86	1.26	1.43	1.48	1.59	1.91	1.81	10.16	1.65	66.35	33.34	76.52
YOL019W	0.95	1.43	1.62	1.68	1.78	2.03	1.97	12.09	1.87	77.86	33.12	89.95
YHR210C	0.87	1.27	1.40	1.44	1.56	1.85	1.78	7.87	3.08	54.01	31.92	61.88
YRN21C	0.71	1.05	1.12	1.18	1.25	1.53	1.46	8.42	2.39	46.01	32.62	54.43
YOR133W	0.95	1.44	1.60	1.64	1.78	2.04	1.97	10.90	2.13	71.31	32.89	82.21
YLD182W	0.85	1.40	1.63	1.64	1.74	2.04	2.06	7.64	1.88	55.94	33.15	63.58
YPL470C	0.99	1.53	1.71	1.74	1.86	2.10	2.06	8.50	1.98	57.50	33.08	66.00
YLR283W	1.04	1.62	1.75	1.83	1.93	2.17	2.15	10.31	2.32	48.59	32.76	58.89
YGL174W	0.62	1.32	1.46	1.54	1.68	1.96	1.89	8.13	2.29	53.83	32.61	61.97
YHR097C	0.91	1.24	1.40	1.54	1.63	1.95	1.96	15.17	2.70	71.62	32.41	87.00
YLG138C	0.89	1.46	1.61	1.67	1.79	2.01	1.99	9.11	2.32	58.73	32.81	67.84
YMR045W	0.90	1.24	1.33	1.34	1.42	1.51	1.55	9.94	2.53	55.14	32.68	65.09
YVR20W	0.86	1.38	1.53	1.56	1.60	1.88	1.82	9.53	2.53	52.65	32.86	64.15
YKL082W	0.81	1.32	1.48	1.52	1.70	1.92	1.92	7.76	2.20	60.39	33.03	70.16
YMR035W	0.88	1.15	1.35	1.37	1.49	1.76	1.68	8.46	2.03	50.43	32.87	59.89
YPL249C	1.00	1.55	1.70	1.74	1.86	2.10	2.08	12.22	2.16	65.65	33.10	77.57
YHL138C	0.88	1.13	1.23	1.29	1.45	1.82	1.72	8.94	2.10	59.27	32.11	69.21
YGR142W	0.86	1.33	1.48	1.59	1.71	2.01	1.88	8.07	1.97	64.01	33.93	72.08
YNL219W	0.64	0.81	0.93	0.97	1.03	1.35	1.22	9.66	3.18	49.74	32.15	59.39
YBR207W	1.06	1.70	1.88	1.89	1.97	2.18	2.21	10.07	1.90	66.57	33.44	76.63
YGR271C-A	0.91	1.16	1.29	1.44	1.58	1.92	1.86	13.10	2.47	69.90	32.87	83.00
YPL171C	0.90	1.44	1.50	1.57	1.71	1.73	1.83	8.31	1.97	59.84	33.38	48.16
YOR238W	0.95	1.43	1.58	1.60	1.69	1.78	1.96	9.93	3.03	47.92	32.32	57.85
YPL249W	0.93	1.41	1.43	1.57	1.66	1.86	1.83	9.36	2.08	58.50	33.29	67.86
YDR214W	0.92	1.38	1.61	1.70	1.80	2.05	2.02	11.00	1.23	81.14	34.15	92.14
YJR110W	0.84	1.13	1.30	1.35	1.46	1.75	1.66	7.33	2.75	47.11	32.64	54.44
YJR070C	0.85	1.21	1.45	1.43	1.51	1.83	1.80	9.75	2.99	50.30	32.41	60.06
YPL248C	1.13	1.77	1.92	1.95	2.07	2.19	2.21	10.11	2.36	64.58	33.04	74.69
YLR297W	0.95	1.33	1.45	1.48	1.54	1.76	1.74	12.60	1.78	60.05	33.64	72.65
YBL082C	0.90	1.32	1.36	1.50	1.58	1.85	1.83	12.08	2.53	62.06	32.91	74.14
YLR267W	0.99	1.42	1.54	1.60	1.68	1.92	1.89	8.74	2.12	47.63	33.32	56.38
YBL150W	0.99	1.43	1.61	1.71	1.81	2.11	2.06	9.68	2.24	65.61	33.24	75.29
YLR300W	0.90	1.35	1.53	1.66	1.80	2.00	2.00	11.57	1.92	71.89	33.57	83.46
YLR173W	1.02	1.58	1.73	1.81	1.88	2.12	2.10	8.07	1.68	46.77	33.82	54.85
YLG206C	0.82	1.25	1.41	1.47	1.58	1.88	1.80	9.04	2.13	56.35	33.38	65.39
YPR046W	0.94	1.28	1.33	1.37	1.43	1.74	1.65	10.81	3.00	62.78	32.53	73.58
YPL051W	0.91	1.36	1.49	1.65	1.76	1.89	1.90	9.16	2.45	46.31	33.09	55.47
YGR161W-C	1.03	1.55	1.70	1.79	1.90	2.10	2.08	7.54	2.51	50.21	33.04	57.74
YLR193C	0.98	1.47	1.61	1.71	1.78	2.09	2.07	11.15	1.95	56.68	33.64	67.83
YNL104C	0.85	1.19	1.40	1.44	1.52	1.91	1.87	8.24	3.13	53.72	32.47	61.96
YLR300W	0.93	1.32	1.34	1.47	1.54	1.83	1.69	9.39	2.44	65.16	33.35	74.55
YLR133C	0.83	1.00	1.17	1.29	1.37	1.63	1.59	10.19	3.05	63.90	32.58	74.09
YMR110C	1.00	1.45	1.59	1.63	1.90	1.84	1.88	8.98	2.11	56.06	33.53	65.04
YLR128W	0.91	1.33	1.50	1.54	1.60	1.91	1.81	7.65	2.01	60.48	33.65	65.66
YDR155C	0.93	1.37	1.57	1.64	1.77	1.99	1.93	10.30	2.37	67.03	34.39	97.33
YLO69C	0.88	1.12	1.29	1.38	1.48	1.71	1.67	9.82	3.27	57.69	32.41	67.51
YER016C	0.88	1.13	1.30	1.34	1.42	1.71	1.67	15.06	3.16	67.92	34.24	73.44
YNL095C	0.69	0.93	1.01	1.11	1.17	1.40	1.38	10.35	3.52	54.03	32.41	64.38
YMR119W	0.69	1.47	1.63	1.77	1.86	2.08	2.09	9.14	3.46	59.42	32.50	68.86
YLR313C	0.88	1.61	1.77	1.86	1.98	2.13	2.16	10.66	2.70	46.54	33.28	57.20
YDR453W	0.95	1.44	1.66	1.65	1.74	2.08	2.06	10.38	2.22	60.23	33.76	70.42
YNL160W	0.96	1.47	1.62	1.70	1.83	2.02	2.00	11.48	2.38	60.11	33.61	71.60
YER168W	0.94	1.47	1.62	1.70	1.81	2.09	2.07	11.24	2.53	89.11	33.48	106.35
YER168W	0.94	1.41	1.55	1.64	1.77	1.96	1.93	16.27	3.39	86.59	32.64	102.86
YPR100W	0.87	1.18	1.31	1.48	1.56	1.89	1.81	8.00	1.76	58.96	34.29	66.96
YIL108C	0.89	1.34	1.55	1.63	1.74	2.01	1.94	9.53	2.65	64.52	34.47	74.05
YLR219W	0.79	1.10	1.23	1.40	1.50	1.78	1.74	7.70	3.03	44.90	33.09	52.60
YFR001W	0.91	1.33	1.38	1.48	1.63	1.94	1.85	11.94	2.51	65.96	33.61	77.91
YER121W	0.89	1.09	1.25	1.37	1.46	1.77	1.71	10.40	2.95	54.27	34.20	63.33
YER133W-A	0.91	1.36	1.54	1.62	1.74	1.84	1.75	13.15	2.68	61.86	33.58	75.00
YLL024C	1.02	1.53	1.71	1.78	1.85	2.10	2.04	10.79	1.92	74.80	34.37	85.29
YKR060W	0.87	1.26	1.44	1.53	1.61	1.91	1.87	10.17	2.65	60.35	33.64	70.52
YNL030W	0.86	1.17	1.30	1.34	1.40	1.75	1.64	10.35	2.86	55.33	32.44	65.68
YOL092W	0.96	1.38	1.55	1.58	1.68	1.98	1.87	9.11	1.98	64.15	34.33	73.26
YLR191W	1.01	1.53	1.73	1.80	1.90	2.17	2.14	9.42	2.02	49.99	34.35	59.41
YBR222C	0.92	1.33	1.46	1.62	1.75	1.97	1.95	12.04	3.08	56.32	33.44	68.36
YLD290C	0.86	1.21	1.41	1.51	1.63	1.86	1.84	9.40	3.38	57.40	33.17	66.80
YDL215C	0.87	1.17	1.32	1.32	1.48	1.78	1.70	9.94	2.43	58.94	34.12	68.89
YOR365C	0.89	1.31	1.35	1.52	1.61	1.81	1.79	9.19	3.01	44.82	33.55	54.01
YMR160W	0.99	1.61	1.75	1.79	1.84	2.03	1.97	7.97	2.75	51.69	33.82	59.66
YPL115C	1.14	1.71	1.89	1.99	2.05	2.24	2.25	8.27	1.75	47.26	34.83	55.39
YER099C	0.88	1.40	1.59	1.56	1.68							

ORF name	OD at 600 nm at sampling time points (hours)											GLUCOSE		FRUCTOSE		TOTAL SUGARS	
	0 h	20 h	27 h	44 h	66 h	93 h	113 h	120 h	146 h	120 h	146 h	120 h	146 h	120 h	146 h	120 h	146 h
YGR029C	0.85	1.04	1.37	1.54	1.44	1.54	1.57	1.69	1.86	1.22	4.74	51.91	34.77	61.49	36.93		
YLR235C	0.84	1.37	1.53	1.54	1.54	1.54	1.54	1.64	1.69	2.03	52.15	34.69	67.94	36.93			
YMR238W	0.98	1.46	1.65	1.78	1.86	1.87	2.13	1.55	2.52	50.29	34.49	58.63	36.93				
YER019C-A	0.86	1.27	1.47	1.58	1.68	1.97	1.91	1.63	1.77	4.82	35.17	58.63	36.94				
YPL159W	1.08	1.55	1.74	1.81	1.91	2.11	2.07	11.51	2.20	71.69	34.77	83.11	36.96				
YNL073W	0.98	1.43	1.63	1.70	1.78	2.07	2.01	11.58	2.43	71.54	34.56	83.12	36.96				
YMR147W	1.04	1.70	1.83	1.93	2.03	2.12	2.22	10.38	2.48	47.87	34.52	52.25	37.01				
YHR128W	0.93	1.35	1.42	1.59	1.68	1.95	1.92	12.24	3.03	56.52	33.99	68.76	37.02				
YIL095W	0.89	1.24	1.40	1.58	1.68	1.95	1.85	15.23	2.92	73.00	34.11	82.24	37.03				
YGL261C	0.88	1.39	1.53	1.61	1.73	2.00	1.94	9.70	2.69	57.77	34.45	67.47	37.05				
YPR049W	0.97	1.48	1.66	1.70	1.83	2.06	1.99	9.72	1.69	65.90	35.38	76.62	37.07				
YPL212C	0.84	1.24	1.21	1.32	1.40	1.46	1.54	8.17	2.41	53.63	34.67	61.79	37.08				
YDL109C	0.87	1.38	1.58	1.52	1.65	1.95	1.96	8.38	2.09	58.96	35.00	67.34	37.09				
YPL220W	1.12	1.74	1.91	1.99	2.08	2.25	2.26	10.33	2.11	65.14	35.04	75.47	37.16				
YKR009C	0.85	1.20	1.36	1.45	1.59	1.80	1.77	9.52	3.28	48.10	33.88	57.62	37.16				
YJR036C	0.98	1.51	1.67	1.73	1.84	2.06	2.02	8.82	3.46	47.90	33.78	58.72	37.24				
YLN034W	0.82	1.11	1.25	1.29	1.36	1.66	1.58	9.91	4.34	46.91	32.91	58.82	37.25				
YPR018W	0.90	1.30	1.34	1.50	1.59	1.67	1.84	13.06	3.28	60.29	33.96	73.36	37.25				
YER049W	0.85	1.24	1.43	1.53	1.63	1.94	1.87	10.36	3.86	67.04	35.40	77.40	37.26				
YLR258W	1.08	1.45	1.57	1.65	1.69	1.88	1.88	9.51	2.32	48.59	34.99	58.69	37.31				
YDR517W	0.94	1.42	1.60	1.65	1.77	2.02	2.02	9.00	2.51	60.01	34.81	68.02	37.32				
YOL089C	0.98	1.50	1.65	1.71	1.80	2.02	1.98	10.04	2.00	67.50	35.34	77.54	37.34				
YJR009C	0.90	1.33	1.52	1.52	1.63	1.93	1.81	10.91	2.34	70.41	35.02	81.32	37.37				
YGR202C	0.86	1.14	1.29	1.32	1.45	1.69	1.63	8.68	2.55	50.65	34.83	59.34	37.38				
YMR158W	0.88	1.27	1.44	1.45	1.55	1.86	1.74	8.68	2.11	64.31	35.27	72.99	37.39				
YNL196C	0.76	0.95	1.03	1.12	1.17	1.46	1.37	9.54	3.78	52.10	33.61	61.64	37.39				
YDL222C	0.81	1.05	1.25	1.20	1.31	1.65	1.60	6.95	1.54	50.77	35.86	57.72	37.41				
YEL025C	0.90	1.42	1.62	1.68	1.76	2.07	1.99	9.81	2.70	56.04	34.73	65.85	37.42				
YPL112C	0.94	1.34	1.49	1.67	1.77	1.97	1.91	9.59	2.59	50.63	34.84	60.22	37.44				
YFL106W	0.93	1.36	1.50	1.68	1.79	1.99	1.94	8.83	2.36	43.17	35.08	52.00	37.44				
YIL154W	0.88	1.21	1.34	1.46	1.54	1.86	1.74	10.23	2.55	66.80	34.33	78.13	37.46				
YLR150C	0.89	1.15	1.26	1.32	1.43	1.71	1.68	10.25	2.55	71.68	34.84	81.32	37.46				
YLR152C	0.85	1.12	1.22	1.22	1.36	1.71	1.64	10.17	2.54	50.53	34.43	63.29	37.49				
YIL234C	0.76	1.13	1.37	1.41	1.44	1.88	1.82	10.27	2.38	64.06	35.19	74.33	37.49				
YHR101C	0.87	1.45	1.59	1.61	1.71	2.00	1.94	9.55	2.44	48.48	34.47	54.59	37.55				
YPL188C	1.08	1.55	1.71	1.80	1.89	2.09	2.09	10.48	2.43	60.01	35.17	70.49	37.59				
YIL029W	0.85	1.14	1.30	1.35	1.38	1.77	1.73	11.18	2.41	67.33	35.19	75.51	37.60				
YNR087C	0.94	1.22	1.37	1.43	1.50	1.76	1.71	10.22	2.41	58.66	35.74	67.88	37.61				
YPL189C-A	0.93	1.40	1.47	1.52	1.62	1.87	1.74	11.14	2.85	34.20	34.78	45.35	37.63				
YIL138C	0.87	1.08	1.20	1.31	1.44	1.71	1.55	12.56	3.02	68.08	34.61	80.64	37.63				
YIL122W	0.96	1.39	1.55	1.68	1.81	2.02	1.93	12.23	3.20	66.62	34.46	78.85	37.66				
YDR262C	0.88	1.38	1.57	1.61	1.72	2.01	2.02	7.76	2.02	55.51	35.65	61.28	37.67				
YGR134C	0.96	1.31	1.47	1.54	1.68	1.89	1.81	10.64	3.28	61.01	34.43	71.65	37.72				
YOR335W	0.88	1.50	1.66	1.70	1.78	1.94	1.92	8.00	2.96	48.22	34.78	58.21	37.75				
YGR232C	0.93	1.32	1.49	1.60	1.72	1.91	1.89	8.00	1.24	65.28	36.52	73.28	37.75				
YDL238C	0.87	1.33	1.55	1.60	1.60	1.89	1.93	9.12	2.07	57.48	35.78	66.60	37.85				
YFL012W	0.87	1.32	1.50	1.56	1.68	1.98	1.90	9.44	2.45	58.69	35.42	68.33	37.87				
YHR068W-A	1.01	1.37	1.47	1.60	1.66	1.93	1.95	11.45	2.24	61.47	35.63	72.92	37.87				
YMR276W	0.94	1.48	1.65	1.67	1.81	2.06	2.01	11.80	2.45	73.28	35.44	85.08	37.90				
YBL186W	0.85	1.13	1.30	1.34	1.46	1.74	1.65	14.51	2.53	81.19	35.45	95.70	37.98				
YDL239C	0.91	1.56	1.76	1.70	1.82	2.10	2.14	10.24	2.05	64.27	35.97	74.50	38.02				
YJL043W	0.88	1.21	1.35	1.41	1.50	1.82	1.75	9.84	3.97	55.61	34.05	65.45	38.02				
YRN027W	0.93	1.24	1.40	1.49	1.59	1.87	1.74	10.75	2.28	69.36	35.76	80.11	38.04				
YNL164C	0.84	1.27	1.43	1.50	1.63	1.84	1.79	10.33	2.38	55.57	34.96	69.91	38.21				
YJL037W	0.88	1.26	1.34	1.41	1.47	1.86	1.74	10.33	2.38	55.57	34.96	69.91	38.21				
YNL161W	0.85	1.11	1.23	1.24	1.34	1.70	1.61	12.53	3.28	51.53	34.90	64.06	38.18				
YPR040W	1.06	1.70	1.85	1.88	1.99	2.21	2.17	9.43	1.91	63.36	36.29	72.99	38.20				
YMR067W	0.93	1.24	1.39	1.46	1.54	1.86	1.74	10.33	2.17	59.33	35.25	69.91	38.21				
YMR161W	0.93	1.41	1.51	1.58	1.66	2.04	2.06	10.23	2.16	52.33	35.77	65.25	38.25				
YPL191C	0.93	1.32	1.39	1.43	1.53	1.86	1.73	10.78	2.25	52.25	35.52	64.83	38.27				
YHL031C	0.94	1.26	1.39	1.48	1.58	1.88	1.72	10.23	2.25	67.13	35.31	78.19	38.43				
YLR045C	0.97	1.37	1.41	1.51	1.65	1.97	1.77	10.42	2.25	51.75	35.17	61.13	38.66				
YBL013W	0.90	1.30	1.43	1.59	1.68	2.01	2.01	12.67	2.55	62.42	35.90	75.09	38.45				
YGL090W	0.87	1.36	1.51	1.54	1.69	1.97	1.88	9.14	2.47	57.83	35.59	69.97	38.46				
YDR288C	0.87	1.39	1.59	1.62	1.72	2.01	2.03	10.70	2.30	65.05	36.20	75.75	38.50				
YLR335W	0.94	1.17	1.29	1.40	1.47	1.80	1.73	11.32	3.54	58.11	35.02	69.43	38.56				
YNL068W	0.89	1.19	1.34	1.36	1.41	1.74	1.62	11.17	2.61	58.24	35.95	80.99	38.56				
YLR300W	0.94	1.45	1.60	1.66	1.81	2.03	1.97	9.87	2.69	68.67	36.07	75.54	38.77				
YIL073C	0.92	1.23	1.37	1.49	1.57	1.87	1.73	12.15	3.37	68.90	35.40	81.06	38.77				
YNL217W	0.68	1.01	1.12	1.24	1.34	1.76	1.64	11.54	3.47	71.09	35.22	83.39	39.01				
YNL214W	0.68	0.89	1.02	1.07	1.13	1.41	1.31	9.24	2.92	48.37	35.87	57.62	38.78				
YBL054W	0.87	1.38	1.54	1.55	1.75	1.94	1.8										

ORF name	OD at 600 nm at sampling time points (hours)										GLUCOSE		FRUCTOSE		TOTAL SUGARS	
	0 h	20 h	27 h	44 h	66 h	93 h	113 h	120 h	146 h	120 h	146 h	120 h	146 h	120 h	146 h	
YIL029W	0.89	1.40	1.30	1.45	1.45	1.69	1.59	1.61	1.59	3.32	5.11	3.87	6.06	3.57	5.57	
YPL274W	1.11	1.33	1.32	1.82	1.67	2.08	2.07	1.17	2.10	1.33	3.70	9.60	38.66	10.06	39.66	
YGR045C	0.94	1.26	1.44	1.53	1.65	2.16	1.85	1.55	1.71	5.93	35.20	63.99	33.91	53.91	33.91	
YLA283C	0.94	1.33	1.37	1.45	1.58	1.84	1.77	1.31	2.49	6.02	37.42	7.33	39.91	7.33	39.91	
YHR033W	0.94	1.25	1.38	1.51	1.59	1.92	1.86	1.23	3.04	7.23	36.92	86.17	39.96	86.17	39.96	
YAR028C	0.90	1.14	1.29	1.45	1.55	1.78	1.76	1.52	2.79	6.57	37.19	78.49	39.98	78.49	39.98	
YDR212W	0.96	1.36	1.57	1.66	1.77	2.03	1.97	1.26	2.08	8.05	37.90	95.79	39.98	95.79	39.98	
YBR168W	0.89	1.24	1.40	1.57	1.67	1.97	1.85	1.04	2.86	6.93	37.13	72.33	39.99	72.33	39.99	
YER119C	0.93	1.50	1.69	1.77	1.87	2.13	2.08	1.56	2.13	6.61	37.82	74.57	39.99	74.57	39.99	
YHR154W	0.89	1.28	1.41	1.45	1.56	1.85	1.74	1.12	4.02	56.19	36.01	67.32	40.03	67.32	40.03	
YER044C	0.95	1.29	1.40	1.43	1.47	1.74	1.64	1.01	2.49	72.55	37.57	83.46	40.06	83.46	40.06	
YER048W	0.91	1.36	1.50	1.54	1.66	1.93	1.85	1.25	3.53	67.46	36.56	81.71	40.09	81.71	40.09	
YPL091W	0.93	1.39	1.56	1.75	1.86	2.00	2.01	1.27	3.15	49.11	36.95	58.39	40.10	58.39	40.10	
YDR216W	1.07	1.55	1.76	1.83	1.92	2.15	2.11	1.66	1.87	8.030	38.24	89.96	40.11	89.96	40.11	
YPL141C	1.16	1.82	1.95	2.07	2.12	2.29	2.32	1.62	2.30	58.82	37.82	69.44	40.12	69.44	40.12	
YDL161W	0.94	1.50	1.71	1.75	1.87	2.06	2.10	1.05	2.73	73.00	37.51	83.15	40.24	83.15	40.24	
YER081W	0.91	1.43	1.61	1.71	1.87	2.04	2.03	1.47	3.10	22.55	37.17	37.34	40.27	37.34	40.27	
YER134C	0.90	1.37	1.54	1.65	1.74	2.03	1.97	0.93	2.45	6.232	37.82	71.75	40.27	71.75	40.27	
YLR395C	0.91	1.25	1.37	1.38	1.52	1.73	1.65	1.10	2.83	70.78	37.47	81.78	40.29	81.78	40.29	
YHR129C	0.94	1.35	1.41	1.56	1.62	1.83	1.83	1.20	4.65	70.26	35.66	89.46	40.31	89.46	40.31	
YHR142W	0.91	1.07	1.22	1.17	1.23	1.81	1.37	0.83	1.88	61.52	38.50	69.84	40.37	69.84	40.37	
YDR415C	0.99	1.46	1.64	1.74	1.83	2.10	2.05	1.32	2.79	63.78	37.59	75.50	40.38	75.50	40.38	
YDR269W	0.90	1.42	1.58	1.60	1.69	1.98	1.94	0.97	2.36	5.03	38.03	62.11	40.39	62.11	40.39	
YPR160W	0.88	1.10	1.27	1.34	1.40	1.69	1.63	0.90	2.86	12.40	38.86	75.67	40.40	75.67	40.40	
YLR262C-A	1.05	1.50	1.66	1.78	1.86	2.06	2.07	10.14	2.56	50.63	37.86	60.77	40.42	60.77	40.42	
YGR252W	0.95	1.40	1.52	1.73	1.91	2.01	2.02	13.19	3.50	73.16	36.92	83.35	40.42	83.35	40.42	
YCL071C	0.94	1.53	1.72	1.77	1.87	2.06	2.09	10.62	2.94	61.42	37.49	72.03	40.43	72.03	40.43	
YKL075W	0.96	1.46	1.64	1.67	1.75	2.03	1.94	0.85	2.15	62.24	38.29	77.09	40.44	77.09	40.44	
YHE026C	0.86	1.03	1.23	1.39	1.36	1.68	1.61	1.13	3.51	66.84	36.94	77.97	40.44	77.97	40.44	
YAR099C	0.85	1.11	1.40	1.44	1.53	1.86	1.81	1.29	3.46	62.26	37.01	80.45	40.47	80.45	40.47	
YAR101C	1.01	1.51	1.60	1.69	1.76	1.97	1.95	1.10	3.54	56.64	36.95	67.94	40.49	67.94	40.49	
YGR256W	0.93	1.46	1.54	1.69	1.74	1.90	1.84	0.98	2.70	44.63	37.80	54.52	40.50	54.52	40.50	
YMR129W	0.97	1.35	1.46	1.45	1.48	1.69	1.74	1.23	3.33	54.83	37.19	67.16	40.55	67.16	40.55	
YKL165W	0.94	1.33	1.58	1.61	1.68	2.13	2.13	1.19	2.95	58.63	37.71	76.62	40.57	76.62	40.57	
YER124C	0.91	1.50	1.68	1.76	1.86	2.11	2.07	1.30	2.49	61.17	38.10	75.55	40.58	75.55	40.58	
YGR052C	0.86	1.29	1.46	1.54	1.68	1.98	1.90	1.30	2.47	49.35	37.15	57.97	40.81	57.97	40.81	
YKL059W	0.92	1.38	1.34	1.48	1.61	1.80	1.77	1.23	2.35	72.36	37.48	85.18	40.83	85.18	40.83	
YER221W-A	0.94	1.34	1.48	1.66	1.76	2.03	2.00	1.48	2.82	75.26	37.74	89.24	40.67	89.24	40.67	
YGL117W	0.92	1.45	1.62	1.74	1.88	2.05	2.04	1.21	2.21	59.07	38.67	77.84	40.84	77.84	40.84	
YIL121W	0.93	1.34	1.59	1.83	1.87	1.85	1.83	1.37	3.49	68.83	37.19	82.20	40.68	82.20	40.68	
YDR382W	1.06	1.26	1.51	1.71	1.84	2.13	2.09	1.35	3.10	72.87	37.65	86.62	40.75	86.62	40.75	
YKR098C	0.90	1.34	1.51	1.61	1.74	1.96	1.95	1.56	2.19	51.61	37.57	61.17	40.76	61.17	40.76	
YOR347C	0.98	1.55	1.63	1.80	1.89	1.98	2.03	10.09	3.21	50.28	37.57	60.37	40.77	60.37	40.77	
YGL262W	0.84	1.20	1.32	1.37	1.50	1.79	1.70	0.98	2.45	45.41	36.66	54.36	41.00	54.36	41.00	
YPR135W	0.91	1.29	1.42	1.52	1.62	1.91	1.86	1.12	2.84	65.52	38.01	77.84	40.84	77.84	40.84	
YPR052C	0.86	1.22	1.29	1.45	1.54	1.62	1.78	0.98	2.78	59.11	38.09	68.59	40.87	68.59	40.87	
YHR167W	0.88	1.13	1.44	1.39	1.56	1.88	1.90	0.97	2.61	56.33	37.28	65.90	40.88	65.90	40.88	
YER080W	0.90	1.29	1.45	1.51	1.61	1.91	1.81	1.01	2.34	67.00	38.07	77.21	40.90	77.21	40.90	
YDR033W	0.93	1.36	1.51	1.51	1.59	1.91	1.84	0.98	2.85	58.20	38.12	68.00	40.97	68.00	40.97	
YAL012W	0.91	1.29	1.42	1.48	1.64	1.81	1.78	1.11	3.74	62.29	37.24	78.40	40.98	78.40	40.98	
YNL111C	0.85	1.26	1.34	1.37	1.44	1.70	1.69	0.95	4.35	45.41	36.66	54.36	41.00	54.36	41.00	
YER128W	0.88	1.41	1.59	1.70	1.79	2.06	2.00	0.95	2.23	64.15	38.79	73.50	41.02	73.50	41.02	
YKL059W	0.97	1.49	1.68	1.73	1.84	2.11	2.04	0.86	2.37	68.00	38.67	78.87	41.03	78.87	41.03	
YLR250C	0.95	1.47	1.61	1.67	1.76	2.07	2.03	0.22	3.24	54.63	37.80	64.85	41.04	64.85	41.04	
YNL008C	0.85	1.29	1.41	1.51	1.58	1.88	1.88	0.97	2.06	52.69	38.59	64.60	41.05	64.60	41.05	
YER167W	0.98	1.24	1.36	1.39	1.46	1.76	1.76	0.93	2.32	57.63	38.63	65.44	41.05	65.44	41.05	
YLR240C	1.13	1.78	1.85	2.08	2.13	2.21	2.27	1.51	2.04	50.09	38.59	64.61	41.34	64.61	41.34	
YMR242C	0.92	1.41	1.60	1.71	1.76	2.04	1.98	1.87	2.81	64.91	38.69	76.78	41.49	76.78	41.49	
YKL148C	0.90	1.39	1.53	1.57	1.68	1.97	1.90	0.97	2.47	59.79	38.03	61.16	41.50	61.16	41.50	
YPL022W	0.86	1.16	1.29	1.44	1.55	1.87	1.72	0.85	3.12	47.96	38.39	58.81	41.50	58.81	41.50	
YLR178C	1.07	1.62	1.78	1.86	1.95	2.12	2.13	1.25	3.03	63.66	38.50	76.18	41.53	76.18	41.53	
YDR229W	0.86	1.07	1.29	1.27	1.39	1.70	1.56	1.28	2.11	52.85	37.57	61.23	41.42	61.23	41.42	
YDR529C	0.94	1.52	1.71	1.80	1.91	2.17	2.11	10.22	2.52	61.24	39.07	71.46	41.59	71.46	41.59	
YBL251W	1.01	1.52	1.67	1.75	1.83	2.11	2.07	8.18	2.35	60.94	39.28	67.22	41.63	67.22	41.63	
YDR279W	0.85	1.18	1.40	1.44	1.52	1.82	1.72	12.35	2.21	79.08	39.42	91.42	41.63	91.42	41.63	
YLL055W	0.91	1.07	1.14	1.24	1.36	1.56	1.43	13.75	3.71	69.52	37.95	83.27	41.65	83.27	41.65	
YER233W	0.85	1.17	1.34	1.38	1.46	1.78	1.76	1.21	2.							

ORF name	OD at 600 nm at sampling time points (hours)											GLUCOSE		FRUCTOSE		TOTAL SUGARS	
	0 h	20 h	27 h	44 h	66 h	93 h	113 h	120 h	146 h	120 h	146 h	120 h	146 h	120 h	146 h	120 h	146 h
YGR101W	0.91	1.23	1.45	1.47	1.53	2.08	1.75	2.33	2.94	41.25	102.4	45.10					
YKR040W	0.89	1.33	1.57	1.70	1.92	1.97	1.99	3.82	5.06	35.38	63.04	43.20					
YIL107C	0.85	1.16	1.32	1.33	1.46	1.69	1.13	3.53	5.13	39.46	39.26	33.01	43.21				
YPR269W	1.10	1.59	1.75	1.79	1.90	2.16	2.12	16.18	2.44	58.14	40.79	62.92	43.23				
YL002W	0.87	1.12	1.26	1.30	1.43	1.78	1.71	1.56	4.09	60.01	39.24	75.08	43.24				
YFR046C	1.04	1.69	1.86	1.99	2.09	2.24	2.24	1.62	2.74	70.25	40.52	83.88	43.26				
YDL233W	0.86	1.32	1.50	1.56	1.66	1.96	2.00	8.87	2.47	6.265	40.79	71.52	43.26				
YMR139W	0.88	1.21	1.35	1.43	1.55	1.78	1.73	11.44	2.08	66.67	41.19	80.12	43.27				
YBR192W	0.92	1.36	1.50	1.57	1.65	1.97	1.89	9.27	3.33	48.37	39.97	57.64	43.39				
YHL039W	0.88	1.14	1.32	1.40	1.53	1.72	1.68	16.33	4.22	69.18	39.09	85.52	43.39				
YHR044C	0.90	1.23	1.43	1.46	1.57	1.63	1.76	14.27	3.15	89.95	40.18	104.22	43.34				
YKL168C	0.96	1.28	1.44	1.49	1.49	1.81	1.66	12.05	2.69	71.42	40.65	83.47	43.34				
YPR022C	1.10	1.77	1.92	2.01	2.10	2.25	2.26	1.01	2.51	74.72	40.84	85.72	43.35				
YOR187W	0.88	1.30	1.48	1.57	1.65	1.75	1.91	11.94	3.83	50.93	39.53	62.87	43.36				
YEL052W	0.98	1.49	1.64	1.71	1.84	2.09	2.03	11.60	3.58	65.38	39.82	76.97	43.41				
YCL032W	0.96	1.40	1.60	1.70	1.83	2.07	2.06	12.89	1.84	85.16	41.66	98.05	43.49				
YDL085W	0.84	1.15	1.34	1.20	1.40	1.68	1.68	10.52	2.48	63.88	41.03	74.40	43.51				
YDR524C	0.93	1.41	1.58	1.67	1.77	2.03	1.97	10.44	2.55	64.29	40.96	74.73	43.51				
YER064C	0.94	1.51	1.69	1.80	1.89	2.10	2.09	11.49	2.96	65.46	40.56	76.95	43.53				
YDL243C	0.87	1.27	1.43	1.38	1.50	1.88	1.77	9.48	3.39	64.99	40.14	74.47	43.53				
YKL046C	0.96	1.35	1.54	1.58	1.68	1.93	1.86	11.57	2.22	74.45	41.34	86.01	43.56				
YER161C	0.94	1.36	1.55	1.63	1.76	1.95	1.93	15.15	3.11	86.15	40.46	101.30	43.57				
YNE129W	0.74	1.05	1.13	1.17	1.23	1.56	1.45	9.75	3.84	50.75	39.76	60.50	43.60				
YDR094C	0.92	1.31	1.52	1.64	1.78	1.98	1.91	13.63	3.65	75.16	39.97	87.87	43.61				
YLR448W	0.89	1.17	1.38	1.50	1.64	1.83	1.82	12.03	3.15	69.06	40.48	81.10	43.62				
YGL121C	0.91	1.25	1.42	1.45	1.54	1.82	1.75	15.88	2.60	87.90	41.05	103.78	43.65				
YHR127W	0.86	1.09	1.29	1.38	1.47	1.80	1.76	18.09	5.45	77.13	38.24	96.22	43.69				
YDR307W	0.88	1.27	1.45	1.47	1.57	1.89	1.84	9.75	2.57	57.42	41.17	67.71	43.74				
YNE091W	0.86	1.22	1.34	1.36	1.44	1.76	1.69	9.53	3.85	46.83	39.91	56.36	43.76				
YNL049W	0.95	1.37	1.51	1.58	1.68	1.91	1.89	15.52	3.94	67.98	39.85	84.50	43.79				
YER044W	0.96	1.52	1.70	1.76	1.87	2.14	2.07	9.50	2.56	58.45	41.24	66.06	43.80				
YER205W	0.83	1.23	1.39	1.43	1.43	1.60	1.60	17.77	3.25	63.66	40.53	72.68	43.84				
YDR165C	0.84	1.13	1.39	1.46	1.64	1.74	1.69	11.23	3.10	61.68	40.89	74.39	43.84				
YER146W	0.91	1.33	1.49	1.55	1.61	1.86	1.89	6.59	2.54	33.62	41.38	76.23	43.88				
YDR034W	0.85	1.21	1.35	1.31	1.43	1.74	1.65	9.77	2.64	63.45	41.28	73.22	43.92				
YIL024C	0.93	1.47	1.70	1.78	1.90	2.16	2.10	10.33	2.20	8.27	41.74	94.69	43.94				
YKL019C	0.91	1.31	1.49	1.50	1.57	1.86	1.74	13.08	3.78	75.59	40.16	89.67	43.94				
YDR089W	0.99	1.47	1.63	1.74	1.85	2.04	2.03	13.95	2.17	88.85	41.78	103.89	43.94				
YKL048C	0.89	1.66	1.74	1.77	1.86	1.94	1.97	15.44	3.54	73.41	38.03	87.76	43.98				
YER194W	0.94	1.27	1.40	1.45	1.58	1.78	1.60	12.10	2.98	73.94	41.03	86.04	44.00				
YER281C	0.82	1.09	1.34	1.46	1.57	1.83	1.80	14.76	3.18	67.26	40.85	82.02	44.03				
YDR307C	0.91	1.38	1.44	1.60	1.70	1.79	1.86	10.42	4.05	48.17	40.03	56.60	44.08				
YOR196C	0.92	1.43	1.60	1.72	1.82	1.91	2.07	11.94	3.94	51.79	40.14	63.73	44.08				
YNE191W	0.67	0.95	1.10	1.07	1.13	1.49	1.44	9.48	4.67	52.47	39.44	61.95	44.11				
YML251W-A	0.87	1.27	1.45	1.54	1.70	1.89	1.87	9.26	2.01	65.42	42.14	74.68	44.15				
YLG057C	0.89	1.32	1.50	1.56	1.67	1.88	1.80	16.59	2.97	91.26	41.22	108.24	44.19				
YDR265W	0.96	1.39	1.58	1.68	1.80	2.01	1.97	12.52	2.46	74.05	41.73	85.57	44.19				
YLR130C	0.97	1.42	1.58	1.67	1.77	1.98	1.89	11.16	2.94	71.33	41.28	82.49	44.21				
YLR368W	0.94	1.33	1.48	1.63	1.72	1.99	1.95	12.44	2.52	73.38	39.97	79.29	44.22				
YKL178C	0.93	1.28	1.44	1.76	1.82	1.83	1.80	14.39	3.71	71.00	40.52	83.39	44.23				
YGR183C	0.88	1.10	1.24	1.39	1.56	1.79	1.66	11.48	4.88	57.19	39.35	68.67	44.24				
YJL201W	0.98	1.39	1.57	1.64	1.73	1.92	1.89	10.54	2.70	65.53	41.55	76.07	44.25				
YLG028C	0.90	1.24	1.41	1.47	1.58	1.81	1.75	13.98	2.89	73.02	41.38	87.00	44.27				
YER180W	0.87	1.09	1.23	1.39	1.48	1.79	1.71	11.96	4.28	57.09	40.03	69.04	44.31				
YLL046C	0.95	1.30	1.57	1.63	1.72	1.90	1.96	11.47	3.08	56.39	41.26	67.85	44.34				
YER162C	0.92	1.29	1.42	1.50	1.61	1.81	1.78	10.81	4.20	49.31	40.99	58.82	45.19				
YEL001C	0.102	1.65	1.82	1.91	1.98	2.00	2.05	15.80	4.77	73.16	39.97	88.95	44.47				
YBL008W-A	0.89	1.42	1.51	1.54	1.59	1.83	1.79	16.85	4.01	57.06	40.73	73.91	44.74				
YPL224C	1.06	1.72	1.89	1.95	2.04	2.24	2.27	12.23	3.11	65.49	41.64	77.72	44.74				
YDR108W	0.93	1.31	1.48	1.52	1.66	1.92	1.85	13.46	2.52	78.81	42.24	92.26	44.77				
YPL425W	0.98	1.36	1.52	1.62	1.79	1.98	1.96	11.05	3.32	65.77	41.49	76.82	44.81				
YAL037C-A	0.90	1.38	1.52	1.54	1.72	1.91	1.90	18.43	4.80	60.19	40.06	78.62	44.86				
YER161C	1.03	1.63	1.77	1.83	1.96	2.13	2.11	10.07	2.69	68.15	41.95	82.22	44.86				
YKL055W	0.88	1.21	1.37	1.44	1.57	1.80	1.75	12.23	2.65	73.61	41.93	85.83	44.88				
YDR285W	0.97	1.37	1.60	1.65	1.71	1.90	1.93	10.81	3.50	47.73	41.43	58.54	44.93				
YPL170W	0.89	1.32	1.38	1.45	1.55	1.76	1.70	12.75	4.02	36.59	40.96	43.83	44.99				
YGL053W	0.95	1.36	1.53	1.73	1.87	2.05	2.05	10.66	3.39	47.79	41.17	58.55	45.13				
YPR074C	0.95	1.36	1.53	1.73	1.87	2.01	2.01	10.70	3.39	53.66	41.96	59					

ORF name	OD at 600 nm at sampling time points (hours)										GLUCOSE		FRUCTOSE		TOTAL SUGARS	
	0 h	20 h	27 h	44 h	66 h	93 h	113 h	120 h	146 h	120 h	146 h	120 h	146 h	120 h	146 h	
YPL130W	0.90	1.29	1.30	1.37	1.41	1.73	1.59	1.54	4.24	7.04	4.21	8.95	45.42	75.84	45.41	
YER054C-A	0.87	1.12	1.33	1.65	1.73	2.00	1.93	1.54	4.24	7.04	4.21	8.95	45.42	75.84	45.41	
YHR08C	0.94	1.34	1.49	1.59	1.65	1.84	1.65	1.53	8.44	11.99	7.41	11.99	45.45	75.89	45.45	
YPR154W	1.04	1.57	1.72	1.82	1.90	2.09	1.78	1.65	3.04	6.93	4.37	7.16	45.51	75.96	45.51	
YBR246W	1.11	1.68	1.82	1.88	1.97	2.15	2.21	1.92	3.03	7.07	4.34	8.19	45.52	75.94	45.52	
YKR033W	0.88	1.09	1.23	1.36	1.45	1.73	1.60	1.48	3.83	8.05	4.20	42.72	95.94	45.55	75.94	
YML148C	1.06	1.28	1.38	1.42	1.50	1.79	1.71	1.35	5.51	55.04	41.08	68.39	45.60	75.49	45.60	
YDL175C	1.00	1.62	1.79	1.83	1.94	2.11	2.15	2.11	5.17	43.79	58.82	45.57	75.82	45.57	75.82	
YML148C	1.06	1.28	1.38	1.42	1.50	1.79	1.71	1.35	5.51	55.04	41.08	68.39	45.60	75.49	45.60	
YDL130W	0.89	1.12	1.29	1.36	1.49	1.71	1.64	1.64	3.55	8.70	43.05	104.78	45.60	75.82	45.60	
YKR053C	0.85	1.24	1.44	1.47	1.59	1.92	1.79	1.74	2.68	7.24	43.93	84.19	45.62	75.82	45.62	
YFR055W	0.88	1.33	1.49	1.60	1.74	1.92	1.89	1.62	3.19	8.98	43.60	106.60	45.79	75.82	45.79	
YER128C	0.99	1.50	1.66	1.71	1.79	2.07	1.98	1.24	3.27	7.80	43.55	92.21	45.82	75.82	45.82	
YNL046W	0.89	1.31	1.50	1.59	1.71	1.96	1.92	1.72	3.13	7.37	43.70	85.50	45.83	75.82	45.83	
YDR351W	1.00	1.14	1.40	1.36	1.42	1.79	1.66	1.82	2.38	7.44	44.48	82.24	45.86	75.82	45.86	
YBR228W	1.00	1.54	1.70	1.76	1.86	2.08	2.04	10.42	2.75	70.21	44.14	80.63	45.89	75.82	45.89	
YPR003C	0.90	1.39	1.37	1.52	1.66	1.77	1.77	1.09	3.32	48.20	43.57	57.29	45.89	75.82	45.89	
YDL213C	0.89	1.28	1.44	1.40	1.55	1.83	1.75	9.67	2.95	64.54	44.08	74.21	47.03	75.82	47.03	
YGL157W	0.93	1.35	1.52	1.55	1.64	1.92	1.83	1.24	2.58	75.65	44.48	82.20	45.76	75.82	45.76	
YL1016W	0.92	1.37	1.58	1.68	1.76	2.03	1.97	11.61	3.08	72.87	44.00	84.48	47.07	75.82	47.07	
YDR354W	0.87	1.35	1.49	1.56	1.66	1.97	1.96	1.27	3.09	62.84	44.00	73.11	47.09	75.82	47.09	
YPL036W	1.08	1.67	1.84	1.92	2.02	2.21	2.20	8.87	2.91	59.06	44.19	67.93	47.09	75.82	47.09	
YLR333C	0.93	1.39	1.54	1.70	1.80	2.07	2.11	11.63	4.38	58.42	42.75	70.05	47.13	75.82	47.13	
YKL116C	0.88	1.17	1.31	1.30	1.40	1.67	1.55	11.23	2.92	81.74	44.25	92.97	47.16	75.82	47.16	
YKL171W	1.01	1.52	1.70	1.75	1.84	2.08	2.02	16.44	2.71	75.75	44.61	92.19	47.32	75.82	47.32	
YL051C	1.00	1.54	1.72	1.80	1.90	2.13	2.09	12.86	2.83	82.28	44.50	95.15	47.33	75.82	47.33	
YDR442C	0.88	1.24	1.44	1.54	1.64	1.93	1.87	11.92	3.42	62.69	43.93	74.60	47.35	75.82	47.35	
YKR076W	0.96	1.33	1.49	1.50	1.53	1.85	1.69	10.92	2.81	67.37	44.54	75.28	47.36	75.82	47.36	
YIL131C	0.84	1.06	1.22	1.29	1.37	1.63	1.58	10.73	4.78	57.26	42.59	68.02	47.36	75.82	47.36	
YKL020C	0.90	1.17	1.32	1.27	1.31	1.63	1.46	11.40	3.07	75.13	44.38	86.52	47.45	75.82	47.45	
YBL216C	1.02	1.42	1.56	1.59	1.65	1.94	1.84	11.74	3.42	65.49	44.00	77.23	47.46	75.82	47.46	
YCL002C	0.93	1.24	1.48	1.62	1.70	1.97	1.86	12.03	3.45	66.67	44.04	72.86	47.49	75.82	47.49	
YGL151W	0.93	1.41	1.58	1.64	1.84	2.08	2.02	14.35	3.45	84.74	44.84	90.40	47.49	75.82	47.49	
YER168C	1.07	1.60	1.78	1.89	1.98	2.18	2.18	12.41	3.09	83.47	44.54	87.69	47.48	75.82	47.48	
YLR497W	1.01	1.60	1.78	1.89	1.98	2.14	2.14	9.64	2.87	93.71	44.84	116.55	47.71	75.82	47.71	
YKL191W	0.94	1.33	1.52	1.55	1.69	1.95	1.87	11.35	3.32	69.61	44.44	80.95	47.76	75.82	47.76	
YPL176C	1.04	1.53	1.71	1.74	1.88	2.11	2.07	12.86	3.64	67.25	44.12	80.11	47.76	75.82	47.76	
YLR153W	0.93	1.44	1.61	1.64	1.75	2.01	1.93	12.44	3.38	85.76	44.46	93.20	47.78	75.82	47.78	
YLR109W	0.93	1.39	1.61	1.68	1.80	2.08	2.03	1.86	2.72	69.45	45.11	81.31	47.84	75.82	47.84	
YER057C	0.95	1.49	1.66	1.76	1.87	2.13	2.08	10.62	3.54	72.27	44.38	77.88	47.92	75.82	47.92	
YER047W	1.01	1.60	1.78	1.89	1.98	2.12	2.19	10.73	3.32	56.06	44.60	68.79	47.92	75.82	47.92	
YER046W	0.98	1.63	1.81	1.91	2.02	2.25	2.20	11.51	3.29	67.02	44.69	75.52	47.98	75.82	47.98	
YCL005W-A	0.96	1.30	1.46	1.59	1.69	1.98	1.90	9.35	4.26	49.58	43.75	59.93	48.01	75.82	48.01	
YPL144W	0.92	1.41	1.52	1.58	1.70	1.97	1.81	12.21	5.02	33.72	43.01	45.93	48.04	75.82	48.04	
YBR230W	0.87	1.21	1.39	1.48	1.60	1.84	1.79	12.78	2.82	78.81	45.34	92.59	48.16	75.82	48.16	
YLN042W	0.89	1.18	1.34	1.37	1.45	1.70	1.61	11.94	3.79	67.85	44.38	87.97	48.17	75.82	48.17	
YGL253W	0.90	1.16	1.39	1.44	1.53	1.98	1.74	12.01	2.63	80.65	45.58	92.65	48.21	75.82	48.21	
YBR201W	0.92	1.36	1.56	1.64	1.75	2.02	1.97	12.11	2.78	77.29	45.56	89.40	48.34	75.82	48.34	
YFL023W	0.92	1.29	1.47	1.51	1.64	1.91	1.88	10.85	2.03	87.20	44.86	105.04	48.44	75.82	48.44	
YDR281C	0.84	1.05	1.26	1.25	1.36	1.66	1.61	12.37	2.39	87.64	46.07	100.01	48.46	75.82	48.46	
YAL062W	0.88	1.25	1.39	1.41	1.60	1.80	1.74	19.04	5.33	65.06	43.19	84.10	48.52	75.82	48.52	
YJR090C	0.95	1.37	1.53	1.54	1.68	1.92	1.82	10.33	2.93	77.16	45.64	87.49	48.57	75.82	48.57	
YFL055W	0.94	1.39	1.56	1.66	1.81	1.97	1.96	15.28	3.19	91.00	45.41	106.28	48.60	75.82	48.60	
YEL054C	0.92	1.13	1.39	1.56	1.71	1.99	1.93	12.64	3.78	68.30	44.86	89.94	48.64	75.82	48.64	
YJL079C	0.87	1.25	1.45	1.49	1.59	1.89	1.79	10.58	3.66	69.49	45.01	80.07	48.67	75.82	48.67	
YPR089C	1.08	1.70	1.88	1.95	2.06	2.25	2.23	12.11	3.40	73.22	45.35	85.33	48.75	75.82	48.75	
YGR261C	0.98	1.44	1.61	1.63	1.76	1.98	1.86	10.85	2.03	77.24	44.88	91.11	48.81	75.82	48.81	
YGR032W	0.92	1.27	1.49	1.52	1.63	1.93	1.80	10.00	2.83	80.85	45.34	94.75	48.83	75.82	48.83	
YER087C	0.98	1.45	1.67	1.70	1.81	2.02	1.97	14.93	3.30	84.87	45.51	104.89	49.29	75.82	49.29	
YER136C	0.88	1.27	1.49	1.52	1.65	1.94	1.84	14.70	4.02	86.89	45.18	101.59	49.20	75.82	49.20	
YIL070C	0.85	1.07	1.24	1.32	1.42	1.67	1.62	10.50	4.55	55.96	44.69	64.46	49.24	75.82	49.24	
YOR311C	1.02	1.60	1.64	1.82	1.91	2.02	2.06	13.43	4.31	54.59	45.24	68.03	49.55	75.82	49.55	
YDR069W	0.85	1.12	1.26	1.41	1.57	1.76	1.60	1.59	2.03	77.97	46.02	92.58	49.28	75.82	49.28	
YER019C	0.91	1.29	1.46	1.56	1.68	1.91	1.77	1.94	2.43	71.44	45.62	84.76	49.29	75.82	49.29	
YLR259W	1.04	1.55	1.72	1.83	1.90	2.15	2.11	11.07	3.26	68.13	46.10	78.20	49.37	75.82	49.37	
YER255W	1.00	1.48	1.64	1.67	1.76	2.03	1.98	12.27	3.83	64.87	45.58	78.14	49.41	75.82	49.41	
YDR318W	0.81	1.02	1.22	1.30												

ORF name	OD at 600 nm at sampling time points (hours)										GLUCOSE		FRUCTOSE		TOTAL SUGARS			
	0 h	20 h	27 h	44 h	66 h	93 h	113 h	120 h	146 h	120 h	146 h	120 h	146 h	120 h	146 h	120 h	146 h	
YGR246C	0.98	1.47	1.65	1.72	1.64	2.10	2.08	1.71	4.11	93.17	45.23	104.2	52.34	93.17	45.23	104.2	52.34	
YLR234W	0.93	1.23	1.53	1.74	1.84	2.02	2.02	1.16	4.02	53.17	48.43	64.33	52.45	53.17	48.43	64.33	52.45	
YMR075W	1.09	1.58	1.73	1.83	1.89	2.11	2.11	1.79	4.44	58.41	48.62	71.29	52.46	58.41	48.62	71.29	52.46	
YBR178W	1.19	1.70	1.85	1.82	1.83	2.18	2.17	1.12	4.17	58.07	48.29	71.29	52.47	58.07	48.29	71.29	52.47	
Y4.098W	0.88	1.19	1.40	1.41	1.51	1.83	1.70	1.57	4.14	68.65	48.46	77.43	52.81	68.65	48.46	77.43	52.81	
YBL167C	0.84	1.11	1.25	1.34	1.47	1.68	1.65	1.42	5.59	58.93	47.14	73.13	52.84	58.93	47.14	73.13	52.84	
YL214W	0.86	1.24	1.43	1.45	1.59	1.85	1.75	1.23	3.94	72.28	48.78	84.86	52.72	72.28	48.78	84.86	52.72	
YBL188C	1.04	1.54	1.70	1.78	1.90	2.15	2.10	1.22	3.91	77.96	48.88	89.87	52.79	77.96	48.88	89.87	52.79	
YHR138C	0.92	1.19	1.28	1.43	1.49	1.78	1.79	1.46	6.39	59.16	48.41	73.84	52.89	59.16	48.41	73.84	52.89	
YEL052W	0.93	1.32	1.47	1.58	1.70	1.89	1.86	1.78	4.45	86.75	48.46	103.83	52.91	86.75	48.46	103.83	52.91	
YKL056C	0.92	1.23	1.43	1.50	1.55	1.84	1.75	1.20	4.46	67.58	48.53	76.61	53.00	67.58	48.53	76.61	53.00	
YDR538W	0.92	1.41	1.63	1.72	1.82	2.08	2.05	1.38	3.96	68.65	48.18	82.46	53.14	68.65	48.18	82.46	53.14	
YLR095W	0.89	1.35	1.55	1.59	1.69	2.00	1.89	1.48	3.34	68.34	48.85	78.82	53.20	68.34	48.85	78.82	53.20	
YOL108C	0.95	1.38	1.55	1.60	1.64	1.92	1.83	1.53	4.45	83.47	48.80	94.41	53.25	83.47	48.80	94.41	53.25	
YHR089C	0.92	1.23	1.35	1.48	1.56	1.89	1.88	1.71	6.11	66.49	47.23	83.60	53.34	66.49	47.23	83.60	53.34	
YBR230C	1.08	1.65	1.80	1.89	1.95	2.11	2.13	1.30	3.90	71.34	49.52	84.44	53.42	71.34	49.52	84.44	53.42	
YPR167C	0.91	1.33	1.43	1.71	1.77	1.99	1.96	1.47	4.75	70.42	48.70	85.12	53.44	70.42	48.70	85.12	53.44	
YMR205C	0.88	1.24	1.37	1.33	1.44	1.71	1.60	1.19	5.60	96.03	47.85	115.22	53.46	96.03	47.85	115.22	53.46	
YLR099W	0.94	1.44	1.65	1.75	1.87	2.06	2.05	1.24	4.31	68.95	49.18	81.38	53.49	68.95	49.18	81.38	53.49	
YMR181C	0.98	1.40	1.54	1.62	1.66	1.76	1.91	1.34	4.26	56.95	49.28	70.49	53.54	56.95	49.28	70.49	53.54	
YLR388W	0.95	1.14	1.41	1.69	1.85	2.13	2.11	1.60	7.28	75.59	46.38	91.59	53.66	75.59	46.38	91.59	53.66	
YDR036C	0.91	1.23	1.39	1.47	1.60	1.82	1.77	1.45	4.03	87.76	48.80	102.01	53.84	87.76	48.80	102.01	53.84	
YLR021W	0.93	1.40	1.56	1.61	1.72	1.97	1.93	1.27	4.67	58.82	49.25	72.53	53.92	58.82	49.25	72.53	53.92	
YIR024C	0.93	1.39	1.53	1.57	1.64	1.91	1.87	1.42	6.07	76.43	47.85	90.55	53.93	76.43	47.85	90.55	53.93	
YJL012C	0.97	1.42	1.65	1.77	1.87	2.12	2.08	1.26	3.64	84.53	50.38	97.19	54.02	84.53	50.38	97.19	54.02	
YKR072C	1.01	1.43	1.60	1.65	1.73	1.98	1.91	1.34	3.95	82.01	50.11	95.46	54.06	82.01	50.11	95.46	54.06	
YDR119C	1.04	1.54	1.68	1.78	1.90	2.09	2.08	1.24	4.29	93.36	50.12	105.76	54.41	93.36	50.12	105.76	54.41	
YPL239W	0.86	1.22	1.27	1.38	1.48	1.55	1.72	1.46	6.10	64.01	48.37	76.61	54.47	64.01	48.37	76.61	54.47	
YLG096W	0.94	1.41	1.57	1.72	1.84	2.02	2.01	1.73	3.91	95.06	50.78	112.79	54.69	95.06	50.78	112.79	54.69	
YOR268C	0.88	1.39	1.40	1.54	1.64	1.93	1.86	1.40	6.12	59.93	48.58	65.02	54.70	59.93	48.58	65.02	54.70	
YKL175W	0.95	1.34	1.54	1.62	1.71	1.97	1.86	1.20	3.71	74.41	50.99	86.42	54.76	74.41	50.99	86.42	54.76	
YNG032C	0.88	1.42	1.62	1.62	1.62	1.82	1.77	1.25	5.66	53.46	48.65	63.95	54.71	53.46	48.65	63.95	54.71	
YHR112C	1.03	1.43	1.57	1.72	1.83	2.06	2.02	1.34	5.35	68.89	49.43	82.23	54.78	68.89	49.43	82.23	54.78	
YOR366W	0.91	1.24	1.37	1.47	1.58	1.82	1.78	1.25	3.95	74.13	51.16	105.65	55.02	74.13	51.16	105.65	55.02	
YOR346C	0.99	1.09	1.25	1.43	1.54	1.87	1.78	1.23	3.92	87.13	51.31	131.56	55.29	87.13	51.31	131.56	55.29	
YHR043C	0.92	1.32	1.54	1.66	1.68	2.03	1.87	1.50	4.84	89.77	50.49	105.27	55.33	89.77	50.49	105.27	55.33	
YDR261C	0.98	1.39	1.61	1.69	1.82	2.04	1.99	1.45	3.27	83.62	52.18	95.57	55.45	83.62	52.18	95.57	55.45	
YDR123C	0.99	1.43	1.63	1.75	1.86	2.05	2.04	1.23	4.78	97.68	50.69	120.20	55.46	97.68	50.69	120.20	55.46	
YMR230W	0.95	1.25	1.43	1.56	1.66	1.75	1.94	1.34	5.13	59.45	50.34	73.39	55.47	59.45	50.34	73.39	55.47	
YKL039C	0.96	1.51	1.72	1.84	1.96	2.13	2.14	1.27	3.23	78.91	52.47	91.62	55.70	78.91	52.47	91.62	55.70	
YIL023C	0.89	1.27	1.48	1.53	1.63	1.97	1.83	1.03	3.36	75.81	52.36	84.44	55.72	75.81	52.36	84.44	55.72	
YDL271C	0.89	1.20	1.32	1.28	1.41	1.74	1.65	1.08	4.65	62.88	51.08	73.76	55.73	62.88	51.08	73.76	55.73	
YMR140W	1.02	1.51	1.63	1.72	1.77	1.84	1.97	1.22	4.50	78.13	52.03	93.35	56.53	78.13	52.03	93.35	56.53	
YBR323W	0.87	1.00	1.14	1.37	1.67	1.83	1.78	1.02	4.84	80.93	51.79	109.11	56.65	80.93	51.79	109.11	56.65	
YER164W	0.93	1.37	1.53	1.64	1.76	1.99	1.97	1.53	4.86	90.18	51.79	109.11	56.65	90.18	51.79	109.11	56.65	
YHR126C	0.91	1.24	1.41	1.46	1.54	1.77	1.73	1.26	4.54	78.97	51.36	93.39	55.47	78.97	51.36	93.39	55.47	
YGR155C	0.92	0.96	1.01	1.03	1.04	1.03	1.03	1.21	1.80	78.89	50.30	91.57	56.78	78.89	50.30	91.57	56.78	
YGR122W	0.96	1.50	1.64	1.72	1.82	2.01	1.96	1.23	3.79	63.67	52.28	78.12	56.27	63.67	52.28	78.12	56.27	
YDR336W	0.96	1.35	1.62	1.74	1.88	2.01	1.96	1.26	4.35	86.71	50.91	103.87	57.80	86.71	50.91	103.87	57.80	
YDR276C	0.98	1.35	1.48	1.57	1.70	1.86	1.94	1.24	4.38	80.76	50.46	96.78	56.87	80.76	50.46	96.78	56.87	
YOR276W	0.93	1.25	1.50	1.50	1.59	1.90	1.85	1.26	4.03	100.66	52.92	122.21	56.95	100.66	52.92	122.21	56.95	
YDR053C	0.92	1.14	1.47	1.47	1.53	1.74	1.84	1.23	4.03	62.85	52.92	77.04	57.04	62.85	52.92	77.04	57.04	
YDR242C	0.94	1.68	1.79	1.82	1.93	1.96	1.96	1.26	4.03	62.85	52.92	77.04	57.04	62.85	52.92	77.04	57.04	
YDR242W	0.93	1.29	1.51	1.52	1.63	1.66	1.66	1.27	4.03	69.69	53.52	93.45	57.14	69.69	53.52	93.45	57.14	
YDR242C	0.94	1.31	1.51	1.51	1.61	1.72	1.81	1.21	4.03	69.69	53.52	93.45	57.14	69.69	53.52	93.45	57.14	
YGR224W	0.95	1.31	1.51	1.61	1.71	1.82	1.78	1.24	4.57	84.59	53.52	98.92	58.09	84.59	53.52	98.92	58.09	
YKL222C	0.92	1.34	1.53	1.57	1.64	1.91	1.82	1.28	4.57	73.67	54.95	94.55	58.17	73.67	54.95	94.55	58.17	
YGR059W	0.88	1.25	1.48	1.50	1.61	1.89	1.80	1.26	4.74	77.57	53.46	90.22	58.20	77.57	53.46	90.22	58.20	
YMR090C	1.01	1.23	1.52	1.72	1.84	2.09	2.05	1.37	5.64	72.59								

ORF name	OD at 600 nm at sampling time points (hours)										GLUCOSE			FRUCTOSE			TOTAL SUGARS		
	0 h	20 h	27 h	44 h	66 h	93 h	113 h	120 h	146 h	120 h	146 h	120 h	146 h	120 h	146 h	120 h	146 h		
YHR043C	0.95	1.45	1.37	1.58	1.95	1.59	1.59	1.55	1.74	2.32	9.12	10.72	57.12	85.54	66.50	66.50			
YHR044W	0.93	1.23	1.37	1.44	1.55	1.77	1.74	1.77	1.74	2.37	9.12	10.72	57.12	85.54	66.50	66.50			
YPR021W	0.92	1.35	1.50	1.55	1.68	1.87	1.85	1.85	1.85	2.37	9.46	10.60	61.25	74.58	68.66	68.66			
YPR174C	0.93	1.21	1.32	1.43	1.54	1.83	1.76	1.62	1.61	2.18	7.19	7.29	55.62	87.59	68.81	68.81			
YHR161W	0.83	1.07	1.28	1.34	1.45	1.88	1.66	1.43	1.43	5.68	8.99	63.19	103.32	68.87	68.87				
YMR281W	0.91	1.37	1.56	1.60	1.71	2.00	1.91	1.26	1.69	11.80	67.28	119.05	65.97	65.97	65.97				
X4L59W	0.91	1.31	1.53	1.59	1.71	2.02	1.88	1.80	1.80	8.46	8.67	60.85	83.47	69.31	69.31				
YFR036W	0.92	1.27	1.40	1.39	1.48	1.76	1.66	2.07	2.07	7.61	8.96	62.18	104.93	69.79	69.79				
YHR075C	0.89	1.27	1.46	1.56	1.72	2.13	1.93	1.55	1.55	5.31	85.50	64.96	101.00	70.27	70.27				
YHR092C	0.90	1.32	1.52	1.54	1.65	1.92	1.85	1.25	1.45	4.95	7.09	65.51	83.74	70.45	70.45				
YDR381W	0.97	1.45	1.61	1.65	1.73	1.97	1.92	2.17	2.17	7.27	9.12	63.40	113.69	70.67	70.67				
YNN077W	0.87	1.28	1.37	1.40	1.46	1.79	1.68	1.50	1.50	15.70	7.19	60.11	63.63	75.81	70.82				
YDR232W	0.85	1.28	1.44	1.45	1.57	1.85	1.83	1.80	1.81	6.75	7.94	64.33	97.95	71.08	71.08				
YBR245C	0.91	1.27	1.43	1.63	1.75	2.01	1.96	17.45	10.88	58.57	60.41	76.02	81.09	71.09	71.09				
YDR244W	0.92	1.25	1.45	1.48	1.60	1.86	1.77	15.58	7.44	90.49	63.74	106.06	71.18	71.18	71.18				
YGR176W	0.90	1.27	1.44	1.53	1.64	1.98	1.82	2.27	4.75	7.82	66.44	91.62	71.18	71.18	71.18				
YHR066C	0.88	1.19	1.40	1.45	1.55	1.89	1.73	15.20	5.93	87.02	65.80	102.22	71.73	71.73	71.73				
YJR032W	0.97	1.08	1.25	1.22	1.30	1.64	1.43	21.25	8.08	86.03	64.08	107.28	72.16	72.16	72.16				
YGR239C	0.91	1.27	1.50	1.53	1.63	1.80	1.80	16.13	6.77	9.20	65.55	108.21	72.32	72.32	72.32				
YPL184C	0.98	1.47	1.53	1.59	1.66	1.71	1.77	20.49	10.77	52.27	61.67	72.76	72.44	72.44	72.44				
YPR026W	0.91	1.25	1.27	1.47	1.59	1.91	1.81	27.06	8.84	80.92	63.69	101.68	72.53	72.53	72.53				
YCL025C	0.95	1.22	1.38	1.57	1.66	1.96	1.90	19.80	9.25	77.64	63.34	97.44	72.59	72.59	72.59				
YIL037C	0.92	1.30	1.51	1.58	1.69	1.85	1.86	15.34	5.50	82.24	67.62	101.59	73.11	73.11	73.11				
YIL087C	0.90	1.10	1.20	1.32	1.39	1.67	1.51	27.12	13.40	87.84	60.38	115.86	73.78	73.78	73.78				
YHR041C	0.90	1.33	1.53	1.63	1.75	1.92	1.92	13.71	5.22	87.65	68.82	101.36	74.03	74.03	74.03				
YKR064W	0.90	1.27	1.44	1.44	1.50	1.82	1.65	16.85	4.95	71.09	65.51	83.74	70.45	70.45	70.45				
YDR377C	0.91	1.37	1.56	1.53	1.61	1.93	1.93	16.55	6.96	83.98	67.57	100.63	74.54	74.54	74.54				
YDR065C	0.85	1.11	1.27	1.27	1.25	1.61	1.47	16.40	7.09	83.12	67.57	99.52	74.67	74.67	74.67				
YGR199C	0.99	1.13	1.29	1.47	1.63	1.96	1.90	19.98	10.67	81.21	64.03	100.89	74.70	74.70	74.70				
YLL028W	0.91	1.36	1.49	1.52	1.61	1.89	1.79	17.64	6.62	83.98	69.03	101.62	75.65	75.65	75.65				
YBR256W	0.96	1.25	1.42	1.46	1.59	1.93	1.91	13.35	9.75	68.18	66.82	87.53	76.58	76.58	76.58				
YBR085C	0.86	1.14	1.24	1.43	1.59	1.82	1.73	16.20	7.09	87.49	92.75	101.71	76.35	76.35	76.35				
YLL034W	0.88	1.18	1.35	1.45	1.63	1.87	1.80	23.66	16.20	103.05	68.50	126.71	78.70	78.70	78.70				
YPR023C	0.96	1.39	1.57	1.57	1.71	1.98	1.91	16.15	8.10	78.85	70.84	90.00	78.94	78.94	78.94				
YBR231C	1.11	1.52	1.68	1.77	1.82	2.02	2.00	23.82	9.60	92.17	69.79	115.99	79.39	79.39	79.39				
YDR085C	1.08	1.70	1.86	1.97	2.07	2.20	2.20	22.60	8.82	97.09	71.96	119.69	80.78	80.78	80.78				
YGR123C	0.92	1.20	1.42	1.46	1.56	2.01	1.75	20.33	8.24	92.97	72.59	113.80	80.83	80.83	80.83				
YBR261C	1.04	1.51	1.68	1.75	1.85	2.12	2.06	21.42	8.99	99.04	72.00	120.45	80.98	80.98	80.98				
YEL053C	0.88	1.23	1.42	1.53	1.65	1.94	1.85	17.59	9.03	73.34	72.12	96.93	81.16	81.16	81.16				
YBR241C	0.91	1.16	1.29	1.45	1.52	1.83	1.74	25.28	13.67	78.10	68.38	104.38	82.04	82.04	82.04				
YNE125C	0.83	1.04	1.07	1.00	1.01	1.34	1.23	19.04	13.55	62.28	68.49	81.32	82.04	82.04	82.04				
YGR135W	0.90	1.14	1.32	1.33	1.42	1.92	1.56	22.78	10.39	96.85	72.31	119.63	82.71	82.71	82.71				
YHR151C	0.88	1.20	1.40	1.44	1.56	2.08	1.74	21.14	8.48	101.59	78.19	122.73	86.68	86.68	86.68				
YOR265W	0.89	1.31	1.47	1.56	1.64	1.90	1.86	21.40	15.14	74.31	72.86	95.70	87.99	87.99	87.99				
YOR375C	0.89	1.29	1.33	1.50	1.61	1.70	1.74	17.88	13.91	59.08	74.53	78.56	88.43	88.43	88.43				
YDR247W	0.99	1.41	1.63	1.71	1.82	2.03	1.97	19.27	10.40	93.56	78.54	112.82	88.94	88.94	88.94				
YIL140W	0.91	1.15	1.43	1.61	1.73	2.03	1.94	25.90	10.68	100.95	75.57	126.85	89.26	89.26	89.26				
YEL166W	0.90	1.17	1.38	1.44	1.55	1.79	1.71	27.84	16.70	67.52	75.86	95.36	96.56	96.56	96.56				
YDR241W	0.94	1.06	1.23	1.36	1.48	1.78	1.67	26.87	13.12	104.49	83.59	134.37	96.71	96.71	96.71				
YDR174W	0.94	0.93	1.03	1.13	1.48	1.78	1.77	33.71	12.72	127.62	84.28	166.33	97.00	97.00	97.00				
YPL271W	0.93	0.99	1.17	1.22	1.37	1.51	1.44	20.45	7.93	86.45	75.79	133.90	96.02	96.02	96.02				
YPR231C	0.92	1.31	1.49	1.46	1.46	1.74	1.64	23.21	11.77	80.71	81.78	119.26	99.75	99.75	99.75				
YMR233W	0.96	1.26	1.29	1.34	1.44	1.62	1.54	23.46	22.46	77.50	77.93	106.22	102.34	102.34	102.34				
YGR223C	0.94	1.34	1.49	1.54	1.61	1.78	1.65	25.19	25.79	97.92	77.24	133.71	103.03	103.03	103.03				
YPR042C	0.90	1.20	1.30	1.37	1.47	1.74	1.67	24.47	20.01	85.87	84.42	118.34	104.44	104.44	104.44				
YLR234C	0.88	1.29	1.45	1.53	1.62	1.82	1.77	23.73	15.97	106.69	91.15	124.33	107.12	107.12	107.12				
YHR255W	0.87	1.15	1.26	1.31	1.46	2.06	1.63	25.97	12.70	104.48	94.55	136.45	108.15	108.15	108.15				
YER042W	0.88	1.01	1.17	1.23	1.33	1.58	1.55	34.43	27.05	87.66	81.67	122.09	108.72	108.72	108.72				
YDR288W	1.01	1.47	1.71	1.88	2.00	2.12	2.12	47.44	26.76	38.99	80.83	84.43	109.59	109.59	109.59				
YDR242W	0.94	1.30	1.52	1.56	1.66	1.90	1.80	30.33	15.90	112.89	94.83	143.22	110.74	110.74	110.74				
YOR233C	0.94	1.07	0.90	0.96	1.22	1.41	1.54	33.90	23.24	85.62	85.76	119.52	119.00	119.00	119.00				
YGL166W	0.90	1.13	1.27	1.26	1.38	1.69	1.60	38.99	19.44	120.10	103.75	159.08	123.20	123.20	123.20				
YLL007C-A	0.87	1.09	1.11	1.03	1.03	1.09	1.09	13.56	10.93	115.28	115.56	130.81	159.05	176.03	176.03				
YOR111W	0.93	1.11	1.16	1.08	1.70	1.77	1.77	69.01	50.60	140.61	128.11	209.62	178.71	178.71	178.71				
YOL110W	0.96	1.47	1.64	1.65	1.74	1.99	1.90	65.45	51.87	151.81	130.74	217.06	182.61	182.61	182.61				

ORF name	OD at 600 nm at sampling time points (hours)											GLUCOSE		FRUCTOSE		TOTAL SUGARS	
	0 h	20 h	27 h	44 h	66 h	93 h	113 h	120 h	146 h	120 h	146 h	120 h	146 h	120 h	146 h	120 h	146 h
YLR135C	0.91	0.91	0.91	0.94	0.88	0.95	0.90	133.44	166.43	149.39	151.63	282.45	314.95				
YNL036W	0.95	0.95	1.02	0.92	0.90	1.02	0.93	133.44	144.78	162.49	171.17	332.44	315.95				
YAR118C	0.95	0.95	0.98	0.92	0.92	1.00	0.94	134.73	166.81	148.63	156.19	283.36	315.99				
YPR036W	0.89	0.95	0.99	0.87	0.90	1.00	0.95	184.45	176.31	177.38	178.44	321.83	317.74				
YPR031W	0.89	1.00	1.01	0.85	0.83	0.94	0.91	173.53	156.93	127.56	162.06	311.09	318.10				
YJR104C	0.89	0.96	0.94	0.85	0.83	0.93	0.87	131.82	170.84	135.51	148.80	269.52	318.84				
YJR205W	0.97	0.99	1.02	0.96	0.94	1.04	0.98	122.37	162.00	143.58	156.95	275.94	318.95				
YNL160W	0.92	0.93	0.98	0.89	0.87	0.99	0.91	146.20	139.06	169.38	180.63	315.57	319.69				
YGR155W	0.95	1.01	1.02	0.93	0.90	1.04	0.95	141.93	157.48	164.44	162.53	306.37	320.01				
YNL180C	0.89	0.91	0.95	0.86	0.84	0.96	0.87	141.43	147.60	197.32	173.58	338.76	321.18				
YDR495C	0.91	0.95	0.98	0.88	0.86	0.98	0.90	158.79	150.81	171.62	174.08	330.41	324.89				
YNL094W	0.94	0.96	0.99	0.91	0.88	1.00	0.92	134.44	153.18	169.41	171.95	303.86	325.13				
YOR108W	0.84	0.87	0.91	0.81	0.78	0.91	0.82	174.19	162.29	190.07	163.63	364.25	326.59				
YNL134C	0.93	0.95	0.98	0.89	0.87	0.99	0.90	109.77	157.51	167.99	169.38	277.76	326.89				
YOR346W	0.89	0.95	0.93	0.86	0.83	0.94	0.87	176.95	161.81	184.43	166.87	361.38	328.68				
YGR087C	0.95	0.94	0.94	0.92	0.91	0.90	0.92	172.50	168.31	138.48	160.63	310.98	328.94				
YIL036W	0.92	0.98	0.97	0.89	0.87	0.95	0.91	200.59	170.67	173.72	158.31	374.31	328.98				
YGL249W	0.96	1.02	1.01	0.93	0.91	1.00	0.94	137.99	155.81	159.16	173.64	297.14	329.44				
YFL036W	0.88	0.93	0.93	0.84	0.82	0.93	0.86	132.51	161.12	158.57	168.67	291.08	329.79				
YNL173C	0.95	0.98	1.02	0.93	0.90	1.03	0.94	168.11	145.62	175.93	184.45	344.03	330.07				
YIL219C	0.92	0.95	0.98	0.89	0.86	0.99	0.90	155.11	157.18	208.18	173.87	363.30	311.05				
YPL071C	0.94	0.92	0.91	0.92	0.89	0.88	0.88	130.86	157.74	140.48	173.98	271.34	331.72				
YER180C-A	0.93	0.98	1.00	0.90	0.88	1.00	0.92	137.37	152.53	177.25	179.90	314.62	332.42				
YER180C	0.93	0.98	0.98	0.90	0.88	0.99	0.91	142.77	158.81	171.81	174.35	314.58	333.16				
YDL083C	0.87	0.98	1.04	0.83	0.81	1.08	0.94	138.30	158.71	151.81	174.84	290.10	333.63				
YNL157W	0.96	0.97	0.99	0.93	0.91	1.00	0.94	152.88	155.51	170.39	178.21	323.37	333.72				
YNL194C	0.96	0.98	1.03	0.93	0.91	1.03	0.94	144.74	147.24	181.06	186.93	325.80	334.17				
YDL085C-A	0.86	0.98	1.07	0.82	0.80	1.09	0.96	134.36	161.94	153.40	172.56	287.77	334.50				
YOR328W	0.99	1.02	1.04	0.97	0.94	1.06	0.97	151.48	154.60	198.02	180.06	349.50	334.66				
YER15C	0.82	0.84	0.87	0.79	0.77	0.86	0.81	142.22	151.82	165.57	183.75	320.78	335.57				
YMR033C	0.89	0.95	0.96	0.85	0.82	0.99	0.99	159.60	156.76	169.65	185.63	326.65	335.94				
YOR221C	0.92	0.95	0.98	0.92	0.92	0.95	0.92	124.40	167.93	128.00	168.36	241.47	333.35				
YER182W	0.95	1.00	1.12	1.04	1.00	1.13	1.05	159.58	163.57	168.24	173.70	321.62	337.27				
YER200W-A	0.93	0.95	0.98	0.91	0.93	0.93	0.93	133.48	169.88	159.59	167.27	232.62	338.26				
YGL251C	0.94	1.00	1.08	0.98	0.91	0.97	0.92	167.45	163.49	165.16	174.84	312.62	338.43				
YLA238C	1.04	1.11	1.12	1.04	1.01	1.12	1.06	148.69	154.70	159.36	162.97	299.05	340.37				
YNL182W	0.99	1.03	1.07	0.97	0.95	1.08	0.98	158.68	161.90	174.99	178.18	324.67	341.14				
YNL183C	0.92	0.94	0.98	0.88	0.86	0.98	0.90	155.00	163.95	167.04	178.67	342.05	342.62				
YFL028W	0.88	0.93	0.93	0.84	0.82	0.94	0.86	138.18	164.89	159.57	178.04	298.75	342.93				
YLR098C	0.85	0.86	0.88	0.82	0.79	0.90	0.82	161.93	159.40	180.86	185.23	342.79	344.63				
YDR414W	0.94	0.99	1.00	0.92	0.89	1.00	0.93	148.93	165.63	170.23	179.10	319.16	344.73				
YDR363C	0.92	0.94	0.96	0.89	0.87	0.98	0.90	186.64	155.55	199.89	185.77	386.54	345.33				
YPL038W	1.00	1.05	1.08	0.97	0.95	1.08	1.00	154.69	163.02	184.94	184.66	339.63	347.68				
YNL204C	0.99	1.03	1.06	0.97	0.95	1.07	0.99	141.32	162.27	195.39	186.09	336.71	348.36				
YNL145W	0.93	0.95	0.96	0.90	0.88	0.97	0.90	143.56	157.12	194.40	191.97	337.95	349.08				
YNL141W	0.94	0.98	1.00	0.92	0.90	1.01	0.93	164.20	162.18	195.82	188.86	360.02	351.13				
YER052C	0.96	0.99	1.01	0.94	0.92	1.04	0.96	158.88	167.81	170.47	183.58	334.35	351.39				
YOR317W	0.96	0.98	1.00	0.94	0.92	1.02	0.95	167.25	158.71	188.55	195.82	356.20	354.53				
YLR375W	0.89	0.93	0.97	0.85	0.84	0.97	0.87	143.10	167.11	175.30	187.50	318.40	354.61				
YDE001W	0.91	0.99	0.93	0.86	0.84	0.96	0.89	223.80	171.18	220.57	184.03	444.17	355.21				
YNL096C	0.88	0.90	0.94	0.84	0.82	0.96	0.86	173.14	154.76	203.05	200.63	376.19	353.39				
YIL036W	0.88	0.89	0.95	0.85	0.83	1.08	0.86	175.93	152.72	191.31	202.74	367.24	355.50				
YNL116W	0.94	0.97	1.01	0.91	0.89	1.02	0.93	162.64	151.84	217.80	197.95	380.44	358.80				
YNL128W	0.93	0.98	0.98	0.98	0.98	0.98	0.98	149.40	160.63	165.77	170.29	335.85	351.87				
YOR247C	0.91	0.94	0.98	0.97	0.95	0.96	0.98	166.27	175.29	181.51	189.59	326.28	352.58				
YLR236W	0.88	1.02	1.05	0.95	0.93	1.08	0.98	161.27	167.30	188.23	188.83	349.51	326.13				
YDR340C	0.84	0.89	0.92	0.81	0.78	0.95	0.87	144.48	170.14	162.51	187.37	306.99	358.51				
YDR298C	0.89	0.94	0.96	0.85	0.82	0.97	0.88	149.73	178.45	169.42	189.95	298.15	359.49				
YEL041W	0.90	0.94	0.96	0.87	0.85	0.98	0.89	154.88	168.81	171.35	182.10	326.33	368.70				
YDR488C	0.94	1.00	1.04	0.92	0.89	1.06	0.96	154.48	178.87	167.17	181.31	321.66	361.17				
YBR034C	1.13	1.13	1.18	1.12	1.10	1.25	1.17	247.01	168.42	223.88	194.39	470.99	362.81				
YIL049W	0.89	0.90	0.96	0.86	0.83	2.07	0.87	168.36	158.84	206.43	206.95	374.79	365.89				
YPR031W	0.95	1.04	1.01	0.92	0.90	1.00	0.88	155.49	185.01	146.15	181.25	361.63	366.26				
YOR288W	0.90	0.92	0.86	0.86	0.83	0.90	0.84	122.01	180.04	150.73	188.27	272.74	368.32				
YDR448W	1.02	1.07	1.10	1.01	0.98	1.12	1.03	153.85	182.84	167.62</							

Appendix 4. Shortlisted 842 deletants from the LYDL_HN based on sugars at both time points (102 and 118 h).

ORF.name	Strain	Optical density at 600 nm at regular intervals (hours)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YML009c	MRPL39	0.91	0.91	1.34	1.75	1.92	2.11	6.33	47.62	4.12	-58.41	10.45	-1.79	280.86		
YMR007W	YMR007w	0.89	0.89	1.31	1.73	1.92	2.13	6.04	-0.33	6.40	-1.57	12.44	-1.90	279.07		
YLR095C	IOC2	1.00	1.33	1.51	2.28	2.09	2.18	0.95	0.93	-1.64	-2.63	-0.69	-1.70	202.39		
YLR062C	BUD28	1.00	1.09	1.14	1.49	1.70	1.79	1.03	1.10	-0.11	-2.44	0.93	-1.33	177.56		
YLR049C	YLR049c	1.00	1.36	1.63	2.01	2.18	2.37	0.88	1.12	-1.64	-2.26	-0.72	-1.14	139.65		
YHL029C	DCA5	1.05	1.55	1.78	2.16	2.33	2.35	0.03	-0.07	-0.11	-1.00	-0.08	-1.0	237.92		
YHL041W	YHL041w	1.03	1.44	1.62	2.04	2.21	2.36	0.00	0.01	0.11	-1.04	0.11	-1.02	263.44		
YOR035C	SCE4	0.93	1.14	1.28	1.55	1.71	1.87	-0.03	0.06	0.64	-0.10	0.39	-0.29	219.19		
YPL229W	DSY1	0.84	1.20	1.44	1.62	2.00	2.21	0.18	-0.12	0.03	-0.35	-0.15	-0.37	264.78		
YHL046C	BAU13	1.03	1.60	1.74	2.11	2.28	2.34	0.04	0.02	0.43	-0.98	0.47	-0.87	362.46		
YHL008C	SHU1	0.95	1.21	1.32	1.75	1.92	2.08	0.04	-0.05	1.81	-0.88	1.85	-0.94	287.25		
YGR177C	ATF2	0.96	1.31	1.52	2.06	2.26	2.37	0.11	-0.05	0.97	-0.87	1.08	-0.92	363.81		
YBL141C	FRK1	1.01	1.41	1.57	1.98	2.09	2.27	0.05	0.07	0.14	-0.97	0.18	-0.91	292.14		
YMR145C	NDE1	1.08	1.37	1.62	2.00	2.17	2.33	0.13	0.21	-0.58	-1.19	-0.45	-0.90	275.70		
YHL022C	GUT1	1.05	1.50	1.72	2.13	2.29	2.43	0.24	0.04	0.27	-0.93	0.51	-0.89	271.07		
YGR157W	CHQ2	0.97	1.36	1.63	2.09	2.23	2.35	0.16	0.14	6.51	-0.99	6.97	-0.85	233.10		
YHR015W	MIP6	0.97	1.26	1.41	1.86	2.07	2.22	-0.09	-0.04	1.72	-0.78	1.63	-0.82	260.53		
YMR189W	GCV2	0.95	1.20	1.39	1.82	2.05	2.26	0.08	0.03	1.72	-0.84	1.80	-0.81	278.96		
YOR029W	YOR029w	0.93	1.23	1.42	1.86	2.08	2.29	0.19	0.00	0.33	-0.81	0.52	-0.81	276.11		
YHL044W	YHL044w	0.94	1.27	1.45	1.90	2.10	2.26	0.08	0.09	1.23	-0.88	1.29	-0.80	276.18		
YPL263C	KEL3	1.00	1.37	1.61	2.07	2.21	2.37	0.04	0.08	7.47	-0.87	7.51	-0.79	287.03		
YPL119C	DBP1	0.97	1.35	1.59	2.02	2.02	2.14	0.15	-0.12	5.54	-0.65	5.69	-0.78	283.77		
YPL221W	FLC1	1.04	1.52	1.73	2.17	2.28	2.42	-0.16	0.05	2.51	-0.82	2.34	-0.77	276.58		
YHL040C	ARN1	0.89	1.19	1.30	1.72	1.90	2.05	0.02	0.13	1.05	-0.90	1.07	-0.76	275.56		
YPL244C	HUT1	0.97	1.28	1.51	2.00	2.16	2.32	0.07	0.05	1.52	-0.81	1.59	-0.75	268.41		
YPL202C	ATF2	0.90	1.21	1.56	1.62	1.80	1.95	0.06	0.05	1.14	-0.79	1.20	-0.74	272.54		
YPL116W	HOS3	0.91	1.18	1.33	1.80	1.96	2.16	0.04	-0.01	1.79	-0.72	1.83	-0.73	278.05		
YDR152C	ECM18	0.94	1.18	1.34	1.84	2.07	2.22	0.57	0.07	0.84	-0.79	1.41	-0.72	267.42		
YMR024W	MRPL3	1.01	1.01	1.05	1.04	1.06	1.12	1.22	0.66	-1.52	-1.38	-0.39	-0.72	475.77		
YOR031W	CRSS1	1.00	1.34	1.52	1.98	2.20	2.37	0.39	-0.04	0.84	-0.69	1.41	-0.72	267.81		
YHR029C	VH9	0.94	1.25	1.43	1.94	2.11	2.24	-0.03	-0.25	10.11	-0.46	10.08	-0.71	270.28		
YHL033C	RPL8a	1.03	1.30	1.49	1.98	2.18	2.28	0.12	0.06	0.73	-0.75	0.84	-0.69	257.06		
YAR042W	SWH1	0.87	1.11	1.28	1.62	1.80	2.05	1.45	1.18	4.16	-1.85	5.62	-0.67	186.14		
YHL017W	YHL017w	1.00	1.35	1.56	1.98	2.14	2.30	0.04	-0.01	3.83	-0.65	3.87	-0.67	283.55		
YPL232W	SSO1	0.82	1.18	1.43	1.88	2.02	2.20	0.00	-0.15	5.46	-0.50	5.46	-0.55	260.85		
YOL081W	IRAD2	0.98	1.29	1.46	1.87	2.03	2.21	0.23	-0.02	0.02	8.33	-0.67	8.31	-0.65	341.87	
YOR045W	TOM6	0.98	1.17	1.36	1.94	2.14	2.14	0.20	-0.01	1.74	-0.56	1.71	-0.61	281.73		
YLR060W	RPL22b	1.00	1.09	1.18	1.55	1.76	2.08	1.06	0.93	-0.53	-1.52	1.03	-0.61	261.44		
YHL045W	YHL045w	1.00	1.28	1.47	1.86	2.07	2.13	0.10	0.05	3.24	-0.64	3.31	-0.59	264.14		
YHL025C	YHL025c	0.93	1.23	1.49	1.89	2.03	2.13	0.08	0.01	3.95	-0.63	3.95	-0.58	272.74		
YAR043C	Unknown	0.92	1.12	1.28	1.63	1.81	2.06	1.28	1.28	-0.38	1.85	0.87	-0.58	126.76		
YPL140C	MKK2	0.95	1.33	1.51	2.02	2.17	2.36	0.30	0.05	5.28	-0.62	5.57	-0.57	277.63		
YGR200C	ELP2	0.94	1.15	1.25	1.71	1.93	2.11	0.14	-0.16	8.63	-0.49	8.77	-0.57	255.03		
YHL022C	SPD11	0.97	1.37	1.57	2.02	2.15	2.29	0.12	0.14	5.44	-0.70	5.77	-0.56	293.06		
YGR036W	YCH1	1.06	1.55	1.83	2.24	2.36	2.45	0.17	0.08	5.52	-0.64	5.69	-0.56	262.99		
YPL28C	THP21	0.97	1.23	1.44	1.97	2.14	2.33	0.03	0.01	4.21	-0.53	4.24	-0.52	267.47		
YPL114W	YPL114w	0.88	1.11	1.27	1.74	1.93	2.14	0.14	0.07	2.74	-0.56	2.88	-0.50	278.42		
YOL057W	YOL057w	0.97	1.31	1.51	1.98	2.14	2.23	0.26	0.14	9.19	-0.62	9.44	-0.48	281.88		
YOR069W	VPS5	0.94	1.28	1.44	1.98	2.18	2.28	0.12	-0.03	4.01	-0.41	4.07	-0.48	320.46		
YOR071C	NRT1	0.96	1.27	1.44	1.88	2.07	2.27	-0.16	0.07	1.82	-0.53	1.67	-0.46	269.46		
YAR014C	BUD14	0.98	1.19	1.37	1.77	1.97	2.24	1.81	1.09	5.35	-1.50	7.16	-0.41	342.79		
YHL036W	MUP3	0.96	1.28	1.44	1.91	2.09	2.21	0.00	0.00	10.07	-0.38	10.07	-0.38	287.90		
YOR036C	PDE2	1.02	1.36	1.50	1.85	2.03	2.21	0.23	-0.04	0.79	-0.34	1.02	-0.38	268.83		
YLR080C	SLR2	0.95	1.23	1.43	1.83	2.03	2.27	0.27	0.06	1.24	-0.51	1.45	-0.45	293.10		
YOL041C	YLF2	1.00	1.38	1.60	2.05	2.23	2.34	0.24	0.02	5.66	-0.21	5.68	-0.19	267.87		
YOL153C	YOL153c	1.00	1.40	1.65	2.14	2.30	2.39	-0.02	0.02	1.57	-0.20	1.55	-0.17	262.67		
YOR062C	YOR062c	0.99	1.33	1.51	1.91	2.12	2.28	0.15	-0.01	4.99	-0.15	5.14	-0.17	273.27		
YPL187W	MF1(ALPHA1)	0.94	1.17	1.29	1.72	1.89	2.11	0.08	0.04	6.31	-0.20	6.39	-0.16	294.03		
YAL053W	AN4	0.98	1.36	1.43	1.81	2.02	2.23	0.21	0.02	5.66	-0.17	5.69	-0.16	266.72		
YAL056W	YNS26w	1.00	1.20	1.37	1.77	1.95	2.11	0.21	0.13	10.56	-0.17	10.59	-0.13	292.65		
YAL050W	AM32	0.93	1.33	1.43	1.85	2.02	2.24	0.12	0.02	5.12	-0.21	5.32	-0.17	262.87		
YMR165C	YMR165c	0.98	1.31	1.51	1.92	2.06	2.21	0.29	0.03	5.21	-0.17	5.32	-0.13	266.73		
YAL254C	RT4C	0.96	1.31	1.56	1.97	2.16	2.29	0.19	0.01	4.14	-0.19	4.28	-0.10	264.35		
YGR153W	YGR153w	1.03	1.44	1.74	2.18	2.30	2.41	0.05	0.07	5.96	-0.15	6.02	-0.09	261.85		
YGR173W	RBG2	1.05	1.52	1.75	2.21	2.34	2.45	0.05	0.05	5.20	-0.14	5.26	-0.08	265.06		
YHL008C	YH008c	1.00	1.36	1.56	2.03	2.21	2.32	-0.05	-0.13	4.32	0.04	4.27	-0.08	284.30		
YML018C	YML018c	0.98	1.28	1.51	1.88	2.03	2.21	1.17	0.27	3.86	0.27	3.58	-0.03	278.37		
YMR002W	MIC17	0.98	1.28	1.45	1.83	2.08	2.17	0.59	0.42	4.32	-0.49	4.49	-0.46	284.22		
YHR022c	YHR022c	0.92	1.16	1.32	1.86	2.06	2.29	0.20	-0.03	6.66	0.00	6.83	-0.07	278.83		
YPL103C	FEMP30	0.94	1.27	1.45	1.93	2.08	2.28	-0.04	0.23	1.54	-0.42	1.58	-0.36	303.93		

ORF.name	Strain	Optical density at 600 nm at regular intervals (hours)							GLUCOSE	FRUCTOSE	TOTAL SUGARS (g/L)	TOTAL AMINO ACIDS (mg/L)		
		0	26	33	48	59	78	102 h						
YML021C	PH1	0.97	0.97	1.25	1.87	2.16	9.27	1.57	-	-1.45	10.2	6.12		
YML019W	DST6	0.97	0.97	1.42	1.82	2.05	9.18	1.57	0.60	-1.46	8.11	5.75		
YML004C	GL01	0.94	0.94	1.44	1.87	2.02	12.73	1.58	1.70	-1.58	14.44	7.67		
YMR0207w	YMR0207w	0.98	0.98	1.45	1.88	2.01	2.22	1.33	1.47	-1.58	8.13	5.63		
YML011C	RAD33	0.96	0.96	1.33	1.77	2.08	2.19	10.81	1.46	3.80	1.32	14.81	8.13	
YML0029w	YML0029w	0.98	0.98	1.49	1.88	2.03	2.21	11.23	1.68	1.81	-1.54	12.84	8.14	
YML078W	CR92	0.97	0.97	1.43	1.79	1.96	2.16	9.62	1.65	2.19	-1.51	11.89	8.14	
YMR021C	MAC1	0.87	0.87	0.90	0.95	1.05	1.26	1.37	1.23	-0.14	-1.07	1.24	8.16	
YML008C	ER66	0.88	0.88	1.21	1.58	1.75	1.98	4.43	1.55	7.48	-1.38	11.91	8.16	
YML029C	RP516b	0.89	0.89	1.15	1.61	1.85	2.11	9.13	1.56	2.37	-1.38	11.50	8.17	
YML005W	TRM12	0.98	0.98	1.52	1.92	2.11	2.32	9.65	1.70	2.39	-1.52	12.04	8.17	
YMR014W	BL022	0.99	0.99	1.56	1.96	2.10	2.27	7.44	1.70	3.79	-1.52	11.23	8.17	
YML082W	YML082w	0.98	0.98	1.39	1.71	1.88	2.11	10.10	1.33	2.11	-1.35	12.21	8.18	
YMR034C	YMR034c	0.96	0.96	1.38	1.86	2.07	2.30	1.02	0.96	-1.54	-0.78	-0.52	0.18	
YML012W	ERV25	0.90	0.90	1.20	1.65	1.85	2.07	6.27	1.27	4.68	-1.07	10.96	8.20	
YGR209C	TRX2	0.93	1.16	1.29	1.81	2.01	2.15	0.09	0.09	5.66	0.12	5.74	2.21	
YDR351W	SBE2	0.98	1.28	1.44	1.84	1.98	2.18	0.01	-0.01	5.76	0.21	5.78	2.21	
YML089C	YML089c	0.93	0.93	1.38	1.78	1.95	2.16	1.71	1.55	0.70	-1.34	2.41	0.21	
YML035C	AMD1	1.00	1.00	1.55	2.01	2.17	2.34	10.67	1.72	3.53	-1.51	14.20	0.21	
YHL019C	APM2	0.99	1.25	1.41	1.88	2.08	2.22	0.02	-0.01	9.02	0.23	9.04	0.22	
YML024W	RP517a	0.92	0.92	1.02	1.36	1.67	2.01	5.40	1.62	5.16	-1.37	10.56	0.25	
YML035C-A	Unknown	1.00	1.00	1.45	1.88	2.05	2.22	15.75	1.50	-0.40	-1.25	15.35	0.25	
YMR015C	ERG5	0.99	0.99	1.51	1.89	2.05	2.24	5.50	1.60	4.49	-1.34	9.99	0.27	
YNL220W	ADE12	1.03	1.03	1.17	1.24	1.29	1.38	1.26	1.45	-1.51	-0.19	-0.25	0.27	
YEL003W	GIM4	0.91	1.15	1.24	1.56	1.68	1.80	0.07	0.07	7.53	0.20	7.59	0.27	
YML029W	USA1	0.94	0.94	1.37	1.79	1.96	2.15	8.47	1.66	2.66	-1.37	11.13	0.27	
YPL164C	MLH3	0.98	1.38	1.57	2.08	2.22	2.37	0.12	-0.01	7.79	0.29	7.90	0.28	
YML059C	NTE1	0.96	0.96	1.41	1.88	2.06	2.24	1.66	1.91	-1.51	-1.63	0.15	0.28	
YML087C	AIM33	0.90	0.90	1.27	1.95	1.95	2.18	1.59	1.54	-1.45	-1.26	0.14	0.28	
YML230W	ADE12	1.12	1.12	1.29	1.38	1.44	1.52	1.51	1.36	-1.31	-1.28	0.20	0.28	
YML023W	CLD1	1.00	1.00	1.53	1.70	1.90	2.06	2.23	1.87	1.94	-1.33	11.60	0.28	
YMR030W	ESF1	0.94	0.94	1.48	1.89	2.05	2.25	1.56	1.31	1.03	-1.52	-0.42	0.30	
YMR029C	ESF3	0.93	0.93	1.43	1.87	2.04	2.24	1.23	1.11	-1.55	-0.32	0.31	0.31	
YML073W	YML073w	0.98	0.98	1.28	1.59	1.88	2.12	2.39	12.89	1.84	0.77	-1.29	14.67	0.32
YLR124W	YLR124w	0.95	0.95	1.51	1.85	2.00	2.18	1.51	1.81	0.87	-1.49	2.38	0.32	
YMR025W	CS11	0.99	0.99	1.46	1.87	2.05	2.28	1.23	1.59	-1.49	-1.26	0.26	0.33	
YML079W	YML079w	0.97	0.97	1.26	1.58	1.74	1.95	18.83	1.90	1.85	-1.59	12.89	0.35	
YML001W	VP77	1.00	1.00	1.40	1.75	1.90	2.08	9.04	1.71	3.70	-1.35	12.73	0.35	
YML003W	YML003w	0.94	0.94	1.42	1.82	1.99	2.17	13.89	1.75	0.27	-1.38	14.17	0.37	
YAL051W	OAF1	0.91	1.17	1.33	1.72	1.84	2.12	1.44	1.45	8.55	-1.06	9.99	0.39	
YAL051W	GAU80	0.93	0.93	1.27	1.61	1.78	1.99	11.53	1.81	2.83	-1.42	14.36	0.40	
YLR125W	YLR125w	0.94	0.94	1.37	1.77	1.96	2.18	1.39	1.21	1.17	-0.81	2.56	0.40	
YNL319W	YNL319w	1.03	1.44	1.68	2.05	2.21	2.38	0.30	0.09	8.35	0.32	8.65	0.40	
YMR006C	PLB2	0.94	0.94	1.37	1.72	1.87	2.08	5.45	1.82	6.19	-1.40	11.64	0.42	
YGR205W	YGR205w	1.03	1.45	1.71	2.14	2.28	2.39	0.26	0.18	11.88	0.26	12.13	0.44	
YMR008C	PLB1	0.91	0.91	1.27	1.66	1.82	2.04	0.95	1.08	7.12	-0.64	8.07	0.44	
YOL036W	YOL036w	1.02	1.31	1.51	1.97	2.15	2.31	1.65	-0.04	28.02	0.50	29.67	0.46	
YAR030C	YAR030c	0.93	1.45	1.45	1.82	1.98	2.21	1.59	1.29	5.66	-0.82	7.25	0.47	
YDR061W	YDR061w	0.90	1.07	1.23	1.77	2.01	2.19	0.02	-0.05	6.07	0.52	6.09	0.47	
YMR022W	UBC7	0.95	0.95	1.40	1.66	2.10	2.26	2.38	0.22	0.06	9.49	0.53	9.71	0.59
YML020W	YML020w	0.95	0.95	1.43	1.89	2.08	2.31	1.23	1.73	-1.38	-1.13	-0.16	0.60	
YPL147W	PKA1	1.02	1.43	1.65	2.13	2.37	2.41	1.30	2.27	1.80	-0.41	16.13	0.61	
YMR032W	PKA1	0.88	0.88	1.27	1.65	1.86	2.09	1.09	1.06	-1.51	-1.23	-0.44	0.63	
YTH199C	MS57	0.98	0.98	1.65	1.87	2.11	2.25	1.37	1.25	2.56	-1.58	-0.68	20.63	
YGR032C	SPB1	0.98	0.98	1.49	1.89	2.01	2.16	2.33	0.11	0.07	0.12	1.58	0.64	
YGR194C	XK51	0.96	1.25	1.43	1.88	2.06	2.20	0.11	0.06	10.65	0.65	10.76	0.65	
YHR014W	SP019	0.98	1.36	1.56	2.00	2.17	2.30	-0.09	-0.17	8.26	0.84	8.17	0.67	
YOL093C	SS42	0.92	1.21	1.44	1.82	1.98	2.19	-1.37	-1.15	4.78	-0.48	6.15	0.67	
YOL025W	PAU17	0.91	1.20	1.44	1.82	1.97	2.18	1.33	1.11	5.97	-0.38	7.30	0.73	
YGR269W	MVB12	0.97	1.22	1.36	1.78	2.02	2.18	0.16	0.09	13.88	0.65	14.14	0.74	
YGR197C	SGN1	0.99	1.38	1.62	2.13	2.30	2.39	0.76	0.03	18.66	0.71	19.42	0.74	
YLR073W	SCI1	1.02	1.17	1.31	1.62	1.80	2.04	0.97	1.01	7.89	-0.28	8.87	0.75	
YML285W	YNL285w	0.97	1.22	1.40	1.65	1.82	2.03	2.21	1.27	1.20	-1.29	14.08	0.76	
YDR197C	THA12	0.98	1.40	1.40	1.97	2.07	2.21	0.20	0.15	1.10	-0.27	12.02	0.76	
YGR034W	RP126	0.94	1.22	1.44	1.79	2.04	2.20	1.57	1.52	1.22	-0.21	1.16	0.61	
YHR032C	YHR032c	0.98	1.20	1.41	1.93	2.11	2.38	0.99	0.34	16.34	0.11	16.73	0.66	
YCL028C	GET1	0.98	1.20	1.37	1.76	2.00	2.19	1.03	0.72	1.74	-0.39	2.77	1.11	
YBR215W	HPC2	0.93	1.21	1.37	1.89	2.11	2.27	0.16	0.12	4.15	0.98	4.32	1.11	
YML288W	BSC4	1.02	1.39	1.56	1.95	2.15	2.30	1.14	-0.07	5.32	1.19	6.47	1.11	
YMR199W	CUN1	0.95	1.16	1.37	1.80	2.03	2.32	0.06	0.09	5.55	1.12	5.61	1.12	
YAL049C	RCM1	0.98	1.20	1.40	1.88	2.11	2.32	1.79	0.18	5.82	-0.97	18.39	0.84	
YMR031W-a	YMR031w-a	0.90	0.90	1.00	1.31	1.54	1.82	1.31	1.60	-1.60	-0.44	-0.29	1.16	
YMR031C	YMR031c	0.90	0.90	1.45	1.86	2.02	2.20	1.26	1.77	-1.57	-0.61	-0.30	1.16	
YHR105W	YPT35	0.97	1.29	1.46	1.98	2.18	2.34	1.33	1.16	15.36	0.00	16.69	1.16	
YIL094C	LYS12	0.93	1.24	1.46	1.82	2.01	2.15	0.60	0.57	1.33	0.61	1.93	1.18	
YLR433W	SFP1	0.99	1.19	1.28	1.75	1.75	1.97	0.45	0.45	3.51	0.74	4.25	1.19	
YGR199W	PMT6	1.01	1.46	1.69	2.12	2.25	2.39	-0.02	0.71	2.42	0.50	2.39	1.21	
YMR214W	SCJ1	0.99	1.22	1.44	1.91	2.11	2.31	0.10	0.16	4.79	1.05	4.89	1.21	
YAL056W	GP82	0.89	1.14	1.36	1.72	1.90	2.13	1.39	0.68	13.27	0.53	1		

ORF.name	Strain	Optical density at 600 nm at regular intervals (hours)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	49	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YLR402W	YLR402w	1.02	1.35	1.57	1.99	2.15	2.31	1.01	0.93	3.68	0.67	4.70	1.60	256.33	256.33	
YLR028W	TPO1	0.87	1.09	1.31	1.68	1.84	2.06	1.81	1.15	15.11	0.46	16.92	1.61	174.43	174.43	
YLR093C	ROX3	0.92	1.16	1.26	1.56	1.79	1.99	1.22	0.85	8.73	0.76	9.96	1.61	254.69	254.69	
YGR086C	PIL1	0.92	1.26	1.48	1.89	2.09	2.26	1.21	1.12	2.07	0.50	3.28	1.62	250.67	250.67	
YDR186C	YDR186c	0.94	1.11	1.22	1.54	1.79	2.02	0.95	1.12	2.75	0.51	3.70	1.63	257.64	257.64	
YHR100C	GEP4	0.90	1.17	1.29	1.69	1.95	2.09	0.99	1.19	5.18	0.46	6.17	1.65	239.80	239.80	
YDR223W	CRE1	1.00	1.42	1.63	2.07	2.22	2.35	1.08	1.12	2.10	0.54	3.18	1.66	259.23	259.23	
YMR144W	YMR144w	0.89	0.99	1.19	1.71	1.97	2.22	-0.05	0.07	0.44	1.60	0.39	1.67	253.58	253.58	
YPL108W	YPL108w	0.93	1.19	1.37	1.88	2.08	2.26	-0.07	-0.05	0.61	1.72	0.53	1.67	254.47	254.47	
YHL035C	VMR1	0.93	1.21	1.40	1.89	2.09	2.20	0.15	0.33	11.09	1.35	11.24	1.68	281.51	281.51	
YDL111W	YDL111w	0.97	1.36	1.53	1.89	2.10	2.28	1.32	1.14	7.95	0.55	9.26	1.68	260.04	260.04	
YGR212W	SUJ1	0.90	1.19	1.40	1.96	2.13	2.24	0.11	0.03	10.33	1.66	10.44	1.69	273.25	273.25	
YMR191W	SPG5	0.94	1.15	1.34	1.83	2.02	2.18	0.92	0.80	14.28	0.90	15.21	1.70	287.79	287.79	
YJL016W	YJL016w	0.95	1.23	1.39	1.80	2.05	2.28	0.87	0.70	3.81	1.01	4.68	1.71	232.20	232.20	
YGL062W	PYC1	0.94	1.35	1.49	1.83	2.04	2.20	1.21	1.01	2.90	0.72	4.11	1.73	238.82	238.82	
YGR079W	YGR079w	1.02	1.36	1.56	2.06	2.25	2.33	1.03	1.04	4.69	0.69	5.71	1.73	270.13	270.13	
YKL010C	UDF4	0.98	1.36	1.61	2.04	2.19	2.32	1.01	0.96	6.99	0.78	7.99	1.74	292.45	292.45	
YPL148C	PPT2	0.94	1.16	1.39	1.88	2.10	2.27	1.18	0.66	9.39	1.08	10.57	1.75	268.10	268.10	
YMR058W	FEET3	0.95	1.18	1.38	1.75	1.96	2.15	0.85	0.96	8.19	0.78	9.04	1.75	278.14	278.14	
YLG096C	FMC1	0.96	1.20	1.43	1.86	2.05	2.21	1.35	0.59	16.27	1.17	17.62	1.75	281.87	281.87	
YBR213W	MET8	1.01	1.44	1.72	2.22	2.50	2.54	0.12	0.15	6.42	1.63	6.53	1.77	284.54	284.54	
YPL186C	UP4	0.89	1.16	1.37	1.92	2.09	2.26	3.36	0.06	29.51	1.72	32.81	1.78	301.54	301.54	
YLR218C	YLR218c	0.89	1.23	1.43	1.78	2.00	2.16	1.41	1.47	2.59	0.34	4.00	1.81	223.54	223.54	
YHR044C	DOG1	0.95	1.33	1.53	2.03	2.20	2.33	1.09	0.50	15.78	1.33	16.87	1.83	255.07	255.07	
YPL247C	ZPL247c	0.96	1.26	1.49	1.98	2.14	2.30	0.64	-0.04	20.73	1.67	21.37	1.83	269.96	269.96	
YMR259W	TM23	0.94	1.21	1.31	1.73	2.00	2.20	1.33	0.82	9.32	1.04	10.65	1.86	242.77	242.77	
YDR374C	YDR374c	1.01	1.42	1.62	2.08	2.21	2.34	1.48	0.02	23.44	1.84	24.92	1.86	296.76	296.76	
YHL037C	YHL037c	1.03	1.44	1.66	2.17	2.34	2.41	1.20	-0.01	26.19	1.67	27.39	1.66	296.39	296.39	
YRC016W	YRC016w	0.95	1.19	1.34	1.73	1.87	2.02	0.93	0.90	3.88	0.97	4.81	1.87	267.63	267.63	
YDZ528W	HLR1	0.86	1.10	1.26	1.85	1.87	1.95	0.88	0.88	19.35	1.19	20.70	1.87	294.05	294.05	
YOL013C	YOL013c	1.01	1.31	1.50	2.08	2.19	2.34	0.71	0.70	1.71	0.71	2.43	1.88	250.56	250.56	
YAL061W	BWD4	0.94	1.25	1.49	1.88	2.02	2.27	2.68	1.22	21.40	0.68	24.09	1.91	160.55	160.55	
YBL151W	ADP1	0.91	1.13	1.32	1.75	2.04	2.17	0.88	1.16	2.41	0.81	2.28	1.92	259.72	259.72	
YBR120C	COP9	0.92	1.07	1.24	1.55	1.78	1.99	0.74	0.77	3.62	1.17	4.36	1.94	284.42	284.42	
YFL013C	IES1	0.93	1.21	1.42	1.75	1.94	2.10	0.79	0.74	6.44	1.24	7.23	1.97	242.86	242.86	
YOR069W	MSA1	1.04	1.48	1.72	2.08	2.25	2.38	0.16	0.02	8.87	2.01	9.02	1.98	267.02	267.02	
YMR223W	FUS2	0.91	1.06	1.24	1.71	1.94	2.18	0.68	0.01	11.41	1.98	12.09	1.99	275.61	275.61	
YDR124W	YDR124w	0.88	1.16	1.33	1.78	1.85	2.00	1.23	1.29	3.71	0.79	4.94	1.99	244.96	244.96	
YJL009C	COX17	0.90	1.22	1.47	1.83	1.97	2.17	1.95	1.20	19.19	0.81	21.14	2.01	261.13	261.13	
YOR081C	GLU5	0.99	1.31	1.50	1.89	2.11	2.27	0.02	-0.21	9.44	2.25	9.47	2.04	279.24	279.24	
YLO101W-A	AIU1	0.89	1.13	1.31	1.68	1.90	2.13	1.09	0.81	16.97	1.28	18.06	2.08	280.03	280.03	
YEI044W	IES6	0.86	1.05	1.12	1.33	1.50	1.71	1.13	0.45	19.39	1.66	20.53	2.10	323.38	323.38	
YCR026C	NPP1	0.91	1.28	1.40	1.69	1.98	2.27	1.63	0.75	25.32	1.42	26.95	2.17	298.26	298.26	
YLD104W	EMC6	0.90	1.14	1.37	1.70	1.85	2.07	2.29	1.19	22.12	1.00	24.41	2.19	239.96	239.96	
YGR029W	YGR029w	0.91	1.06	1.24	1.71	2.06	2.30	1.08	0.80	2.03	1.46	2.86	2.26	253.94	253.94	
YDR520C	URC2	1.04	1.49	1.74	2.16	2.30	2.40	0.83	0.80	2.03	1.46	2.86	2.26	253.94	253.94	
YNL048C	END3	0.93	1.15	1.34	1.73	1.94	2.10	1.15	0.90	9.10	1.36	10.24	2.27	270.80	270.80	
YDL129W	YDL129w	0.95	1.33	1.48	1.94	2.00	2.10	1.24	1.19	4.28	1.08	5.52	2.27	241.31	241.31	
YNL070W	TOM7	0.89	0.99	1.15	1.78	1.92	1.98	0.91	0.68	10.11	1.59	11.02	2.27	281.37	281.37	
YGL026C	Unknown	0.98	1.34	1.54	1.98	2.18	2.28	1.82	0.79	21.28	1.50	23.10	2.29	304.01	304.01	
YHL031C	GOS1	0.97	1.28	1.50	1.99	2.17	2.33	0.56	0.04	15.28	2.25	15.94	2.29	279.67	279.67	
YLO707W	ECM33	0.92	1.16	1.38	1.80	2.01	2.18	1.11	0.74	9.19	1.62	10.30	2.36	239.21	239.21	
YDR417C	YDR417c	0.88	1.07	1.12	1.36	1.56	1.76	1.16	0.43	19.99	1.97	21.15	2.40	323.97	323.97	
YMR119W	RH1	0.91	1.21	1.42	1.74	1.87	2.09	2.55	1.17	24.63	1.08	27.18	2.25	160.15	160.15	
YMR295C	YMR295c	0.92	1.25	1.44	1.85	2.08	2.25	1.55	1.26	6.63	1.00	8.18	2.26	246.03	246.03	
YDR520C	URC2	1.04	1.49	1.74	2.16	2.30	2.40	0.83	0.80	2.03	1.46	2.86	2.26	253.94	253.94	
YNL048C	END3	0.93	1.15	1.34	1.73	1.94	2.10	1.15	0.90	9.10	1.36	10.24	2.27	270.80	270.80	
YER179W	DRC1	0.85	1.22	1.40	1.84	2.04	2.21	1.82	1.14	1.69	1.51	1.81	2.21	290.56	290.56	
YHL033C	YHL033c	0.91	1.27	1.48	1.88	2.19	2.34	-0.21	-0.18	5.47	2.83	5.27	2.65	267.97	267.97	
YJR049W	GEF1	0.96	1.16	1.30	1.78	1.98	2.08	1.45	0.77	20.81	1.88	22.28	2.66	269.46	269.46	
YML027W	YOK1	0.97	1.27	1.52	1.98	2.14	2.29	1.02	0.92	14.95	1.74	15.57	2.66	276.33	276.33	
YGL029W	CGR1	0.85	1.08	1.20	1.65	1.89	2.05	1.41	0.79	13.96	1.87	15.37	2.66	264.10	264.10	
YHL042W	YHL042w	0.98	1.35	1.55	2.01	2.16	2.30	0.79	0.02	24.23	2.65	25.02	2.67	283.11	283.11	
YOR008C-A	YOR008e-a	0.94	1.14	1.28	1.73	1.94	2.04	1.47	0.95	18.36	1.73	19.83	2.68	298.61	298.61	
YCR045C	RTT12	0.91	1.28	1.40	1.73	1.91	2.06	0.88	0.74	5.55	2.03	6.24	2.77	272.03	272.03	
YBR241C	YBR241c	1.01	1.25	1.44	1.88	2.21	2.36	0.29	0.14	16.43	2.63	16.72	2.77	300.90	300.90	
YKL062W	MSN4	1.05	1.38	1.59	2.09	2.24	2.35	0.98	0.97	5.23	1.81	6.22	2.78	275.95	275.95	
YLR414C	PUN1	0.98	1.30	1.46	1.86	2.03	2.23	1.12	1.06	8.61	1.79	9.73	2.85	288.32	288.32	
YLR266C	CTS1	0.93	1.03	1.11	1.59	1.92	1.67	1.71	0.84	16.98	2.01	18.69	2.85	248.48	248.48	
YPL220W	YPL17a	0.97	1.39	1.59	2.06	2.18	2.33	0.36	0.10	13.38	2.77	13.74	2.86	290.64	290.64	
YMR204C	IMP1	1.04	1.30	1.55	2.00	2.19	2.37	0.55	0.22	8.41	2.65	8.86	2.87	274.96	274.96	
YDR105C	TMS1	1.05	1.40	1.57	1.98	2.17	2.32	0.47	0.16	15.32	2.71	15.78	2.87	283.50	283.50	
YGR217W	CCH1	0.96	1.29	1.50	2.01	2.20	2.31	0.99	0.46	23.45	2.43	24.44	2.89	275.29	275.29	
YLD024W	DID4	0.87	1.15	1.27	1.61	1.77	1.91	1.22	0.79	13.46	2.11	14.88	2.90	284.61	284.61	
YFR032C	NAM2	0.90	1.11	1.28	1.65	1.90	2.05	0.59	0.82	8.59	2.10	9.18	2.92	267		

ORF.name	Strain	Optical density at 600 nm at regular intervals (hours)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)			
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h			
YDR013W	GRX2	0.98	1.42	1.55	2.02	2.35	0.76	0.76	4.47	2.42	5.47	2.42	2.10	2.10	2.10	2.10		
YDR017C	HMP1	0.95	1.37	1.59	2.04	2.26	2.13	0.86	25.70	2.47	27.83	3.19	260.61	260.61	260.61	260.61		
YCR006c	WSC4	1.00	1.40	1.60	2.03	2.21	2.29	0.77	0.74	5.93	2.46	5.75	3.20	255.26	255.26	255.26	255.26	
YHL028W	WSC4	1.00	1.33	1.48	1.98	2.13	2.27	0.29	0.02	1.74	5.39	2.46	5.75	3.20	285.61	285.61	285.61	285.61
YCR023C	PRO2	0.88	1.19	1.25	1.51	1.74	1.99	1.41	1.25	3.42	1.98	4.83	3.23	331.36	331.36	331.36	331.36	
YLR455W	YL455w	0.89	1.08	1.27	1.81	2.36	2.16	0.72	0.73	3.79	2.51	4.51	3.23	262.02	262.02	262.02	262.02	
YDR441C	APT2	0.89	1.34	1.55	1.93	2.15	2.24	1.37	0.59	19.53	2.67	20.91	3.27	295.07	295.07	295.07	295.07	
YDR033W	WHS5	0.98	1.31	1.48	1.92	2.13	2.30	0.12	-0.05	16.30	3.33	15.42	3.28	270.58	270.58	270.58	270.58	
YBL130W	SP019	0.89	1.06	1.15	1.63	1.83	2.06	0.33	0.07	18.69	3.22	19.02	3.30	315.89	315.89	315.89	315.89	
YGR161W-C	YGR161w-c	0.87	1.19	1.36	1.67	1.86	2.03	1.60	0.81	18.88	2.51	20.48	3.32	258.44	258.44	258.44	258.44	
YER181C	YEI81c	0.92	1.08	1.17	1.81	1.67	1.81	0.57	1.03	13.49	2.29	14.07	3.32	302.90	302.90	302.90	302.90	
YER181W	RP121a	0.89	0.97	1.06	1.46	1.67	1.90	0.81	0.99	13.76	2.34	14.58	3.33	285.71	285.71	285.71	285.71	
YCR019W	MAK32	0.90	1.14	1.30	1.78	2.00	2.14	1.25	1.14	3.77	2.22	5.01	3.36	274.96	274.96	274.96	274.96	
YKL093W	MBR1	0.97	1.31	1.48	1.88	2.06	2.20	0.88	1.06	6.10	2.30	6.98	3.36	270.38	270.38	270.38	270.38	
YKL055C	OAR1	0.96	1.31	1.56	2.02	2.17	2.27	0.58	0.58	5.46	2.79	6.05	3.38	266.98	266.98	266.98	266.98	
YLL038C	ENT4	0.88	1.05	1.26	1.64	1.82	2.05	1.35	1.09	4.44	2.28	1.79	3.38	153.84	153.84	153.84	153.84	
YKL032C	DR1	1.00	1.41	1.70	2.17	2.31	2.44	1.06	0.97	7.22	2.42	8.27	3.38	252.02	252.02	252.02	252.02	
YPR083W	MDM8	0.93	1.25	1.45	1.93	2.11	2.23	0.46	0.18	16.23	3.21	16.69	3.39	281.08	281.08	281.08	281.08	
YLL015W	BPT1	0.89	1.16	1.38	1.72	1.85	2.06	2.23	1.37	24.53	2.03	26.76	3.40	224.91	224.91	224.91	224.91	
YLR189C	ATG26	1.07	1.44	1.67	2.11	2.27	2.40	0.84	0.86	8.81	2.55	9.65	3.40	278.75	278.75	278.75	278.75	
YHR158C	KEL1	0.88	1.08	1.19	1.52	1.92	2.20	1.30	1.10	0.92	17.04	2.50	18.14	3.43	288.51	288.51	288.51	288.51
YPL216W	YPL216w	0.86	1.07	1.22	1.85	2.04	2.23	0.33	-0.10	13.32	3.53	13.66	3.43	286.14	286.14	286.14	286.14	
YBR214W	SDS24	0.94	1.19	1.38	1.86	2.07	2.21	0.15	0.07	9.02	3.38	9.17	3.44	295.25	295.25	295.25	295.25	
YLR371W	ROM2	0.99	1.28	1.39	1.72	1.91	2.07	1.41	0.81	19.90	2.63	21.31	3.45	253.64	253.64	253.64	253.64	
YOR338W	YOR338w	1.01	1.28	1.49	1.95	2.13	2.31	0.16	0.12	4.68	3.33	4.84	3.45	264.26	264.26	264.26	264.26	
YER076c	YER076c	0.94	1.32	1.52	1.93	2.12	2.26	1.24	0.85	10.95	2.60	12.19	3.46	232.79	232.79	232.79	232.79	
YLL010C	PSR1	0.89	1.10	1.28	1.60	1.75	1.96	3.19	1.13	30.02	2.36	33.22	3.49	192.13	192.13	192.13	192.13	
YCR050c	YCR050c	0.86	1.19	1.29	1.64	1.85	2.03	1.28	0.74	23.23	2.76	24.51	3.50	303.13	303.13	303.13	303.13	
YMR247C	RKR1	0.93	1.04	1.18	1.64	1.88	2.14	-0.08	0.04	10.69	3.50	10.61	3.53	285.32	285.32	285.32	285.32	
YOL047C	YOL047c	0.93	1.18	1.41	1.94	2.10	2.29	0.81	0.32	21.11	3.86	21.92	3.54	275.61	275.61	275.61	275.61	
YFL189W	SUR2	0.94	1.21	1.30	1.77	1.97	2.08	2.28	1.06	0.84	13.38	2.70	14.40	3.54	244.39	244.39	244.39	244.39
YCL004W-B	HBK1	0.88	1.16	1.34	1.75	1.93	2.03	2.20	0.50	0.21	7.41	3.37	7.50	3.54	244.64	244.64	244.64	244.64
YFL170W	YFL170w	1.08	1.53	1.73	2.08	2.26	2.34	1.01	0.97	7.13	2.69	8.14	3.65	268.04	268.04	268.04	268.04	
YPL224W	FMP40	1.04	1.42	1.63	2.04	2.20	2.34	0.49	-0.05	14.81	3.73	15.39	3.89	296.01	296.01	296.01	296.01	
YML055W	SPC2	0.89	1.26	1.44	1.88	2.13	2.32	1.02	0.82	12.84	2.86	13.86	3.69	355.91	355.91	355.91	355.91	
YML153C	GIM3	0.97	1.21	1.39	1.91	2.01	2.12	1.22	0.83	12.42	2.86	13.64	3.69	245.01	245.01	245.01	245.01	
YJL139C	BP521b	0.92	1.07	1.23	1.67	1.89	2.09	0.81	0.48	13.89	3.21	14.70	3.70	284.82	284.82	284.82	284.82	
YKL037W	AIM26	1.00	1.30	1.47	1.99	2.18	2.30	1.20	0.79	12.88	3.01	14.08	3.80	270.15	270.15	270.15	270.15	
YPL163C	SVS1	1.02	1.42	1.62	2.09	2.23	2.36	0.74	0.62	16.40	3.79	17.14	3.80	299.25	299.25	299.25	299.25	
YHR249W	MW1	0.96	1.29	1.50	1.96	2.15	2.26	1.20	0.74	19.01	3.08	20.21	3.81	312.21	312.21	312.21	312.21	
YDL035C	GPR1	0.87	1.05	1.22	1.65	1.89	2.06	0.72	0.88	5.79	3.03	6.52	3.83	272.35	272.35	272.35	272.35	
YPL262W	FUM1	1.00	1.45	1.72	2.15	2.28	2.40	0.18	1.62	10.01	2.22	10.18	3.84	272.45	272.45	272.45	272.45	
YLR34C	ACD1	0.87	1.06	1.27	1.60	1.80	2.02	1.03	0.63	15.76	3.21	16.80	3.84	266.93	266.93	266.93	266.93	
YAL030W	SNC1	0.91	1.14	1.33	1.71	1.85	2.08	3.57	1.41	23.15	2.46	26.72	3.86	236.59	236.59	236.59	236.59	
YEL043W	YEL043w	0.98	1.38	1.59	2.05	2.21	2.31	0.61	0.02	24.18	3.85	24.80	3.87	282.42	282.42	282.42	282.42	
YCR080C	YCR080c	0.90	1.19	1.36	1.61	2.00	2.15	2.29	0.37	0.28	16.79	4.27	18.06	4.30	280.72	280.72	280.72	280.72
YER177W	BHM1	0.85	1.00	1.06	1.29	1.49	1.67	1.75	0.79	28.12	3.78	28.87	4.57	297.58	297.58	297.58	297.58	
YCR050C	CIT2	0.98	1.28	1.44	1.89	2.11	2.22	1.99	1.12	29.19	3.47	31.18	4.59	305.66	305.66	305.66	305.66	
YER204W	YER204w	0.95	1.14	1.35	1.76	1.95	2.05	2.25	0.58	0.15	18.87	4.11	19.45	4.25	280.04	280.04	280.04	280.04
YHL005C	YHL005c	0.85	1.08	1.16	1.41	1.60	1.81	1.38	1.41	11.80	2.85	13.18	4.26	294.52	294.52	294.52	294.52	
YNL275W	BOR1	1.08	1.50	1.74	2.10	2.26	2.34	2.34	0.53	-0.05	17.64	4.35	18.17	4.29	269.28	269.28	269.28	269.28
YKR096W	YKR096w	0.89	1.04	1.20	1.61	1.87	2.07	1.02	0.77	14.55	3.52	15.57	4.30	270.13	270.13	270.13	270.13	
YKL102C	YKL102c	0.94	1.25	1.42	1.92	2.11	2.33	0.72	0.95	11.15	3.38	11.88	4.33	275.22	275.22	275.22	275.22	
YNL325C	FIG4	1.10	1.44	1.71	2.14	2.31	2.45	1.60	1.35	30.69	3.00	32.29	4.34	304.36	304.36	304.36	304.36	
YJL198C	ELO1	1.03	1.35	1.56	1.99	2.14	2.29	0.89	1.06	9.54	3.32	10.44	4.38	257.19	257.19	257.19	257.19	
YKL044W	YKL044w	0.99	1.27	1.47	2.04	2.24	2.36	1.23	1.08	20.36	4.61	21.47	4.69	295.69	295.69	295.69	295.69	
YER144C	YER144c	0.90	1.07	1.17	1.59	2.05	2.25	1.15	-0.03	26.53	4.74	27.88	4.71	280.53	280.53	280.53	280.53	
YKL096W-A	CWP2	0.87	1.02	1.17	1.59	1.82	1.98	1.10	0.68	16.64	4.06	18.74	4.73	287.39	287.39	287.39	287.39	
YML283C	WSC2	0.98	1															

ORF.name	Strain	Optical density at 600 nm at regular intervals (hours)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)		
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h		
YMR163W	NUP23	0.99	1.16	1.35	2.02	2.24	0.74	0.20	0.20	26.78	4.96	21.20	29.14	5.16	5.16	277.63	
YGR032C	TMA1	1.00	1.41	1.63	2.08	2.24	2.11	0.80	0.92	13.55	4.24	11.44	13.55	5.16	5.16	277.63	
Y/R110W	YMR1	1.14	1.44	1.64	2.19	2.31	2.42	0.80	0.96	5.37	2.04	4.45	5.17	271.01			
YBR209w	YBR209w	0.94	1.13	1.34	1.57	1.82	2.00	1.54	1.52	15.03	3.85	16.58	16.58	5.17	5.17	291.26	
YIL13C	CSM2	1.03	1.17	1.34	1.87	2.14	2.36	2.07	0.76	4.13	1.49	6.20	5.19	231.90			
YGR285C	ZU01	0.98	1.32	1.41	1.54	1.70	1.87	2.86	3.02	6.85	2.17	8.51	5.19	303.12			
YOL101C	I2H4	0.92	1.46	1.64	2.04	2.20	2.31	3.35	2.97	14.65	2.23	18.09	5.20	299.00			
YHR09W	TRB2	0.99	1.31	1.50	2.00	2.18	2.30	1.14	1.20	3.42	4.03	4.56	5.23	283.76			
YLR184C	MKN1	0.97	1.23	1.40	1.82	2.02	2.20	1.63	0.19	24.24	5.06	25.87	5.26	310.85			
YML25C	GIS2	0.95	1.25	1.47	1.88	2.07	2.23	0.41	0.10	14.78	5.16	15.19	5.26	264.85			
YHR072W	VP529	0.94	1.25	1.39	1.74	1.88	2.04	0.83	-0.13	26.24	5.44	27.07	5.31	332.43			
YDR31W	RAX1	1.02	1.32	1.50	1.92	2.10	2.27	0.84	0.13	25.29	5.20	26.14	5.33	293.51			
YDR234W	LYS4	0.93	1.17	1.31	1.65	1.85	2.00	1.54	1.24	22.50	4.11	24.44	5.35	296.34			
YHR199C	AIM46	0.94	1.20	1.36	1.81	2.03	2.15	1.57	2.07	20.17	4.29	21.75	5.35	326.06			
YKR036C	CAF4	0.93	1.16	1.39	1.95	2.15	2.30	1.17	0.70	21.45	4.66	22.63	5.36	296.04			
YLR176C	RFX1	1.04	1.33	1.57	2.08	2.24	2.36	1.25	0.93	18.25	4.46	19.50	5.39	286.05			
YLR389C	STE23	0.90	1.11	1.27	1.61	1.89	2.08	2.25	0.02	2.50	5.00	15.14	5.40	289.97			
YDR225W	HTA1	0.98	1.26	1.41	1.89	2.09	2.27	1.48	0.88	21.84	4.53	23.31	5.41	280.60			
YGL125W	0.90	1.11	1.18	1.56	1.92	2.11	0.93	0.70	17.18	4.71	18.11	5.41	294.26				
YLL007C	YLL007c	1.00	1.48	1.64	2.00	2.18	2.31	3.17	3.14	13.21	2.27	16.37	5.41	297.33			
YGR012W	YGR012w	1.01	1.24	1.36	1.82	2.04	2.25	1.27	0.97	13.62	4.50	14.89	5.47	280.07			
YKL115C	YKL115c	0.90	1.13	1.37	1.92	2.12	2.29	1.15	1.28	13.89	4.21	15.04	5.47	274.04			
YJR154W	YJR154w	0.96	1.43	1.59	1.97	2.15	2.26	2.93	2.99	9.95	2.50	12.89	5.48	280.77			
YJL160C	YJL160c	0.89	1.03	1.20	1.63	1.85	2.06	1.40	0.78	21.00	4.71	22.40	5.49	270.43			
YGR051W	CKA2	1.01	1.41	1.58	1.88	2.08	2.25	0.02	2.50	3.97	3.00	3.99	5.49	297.53			
YGR071C	YGR071c	0.98	1.32	1.53	2.01	2.19	2.28	0.32	0.56	30.64	4.95	32.97	5.51	288.18			
YOL058W	ARG1	0.89	1.16	1.32	1.75	1.95	2.08	1.27	1.30	7.79	4.21	9.06	5.51	226.88			
YJR111C	YJR111c	0.96	1.50	1.67	2.05	2.21	2.31	3.27	3.06	14.77	2.47	18.04	5.53	278.15			
Y/JL170C	ASG7	0.97	1.15	1.28	1.72	1.91	2.14	1.28	1.18	13.82	4.36	15.10	5.54	282.98			
YGR011W	YGR011w	0.98	1.36	1.59	2.06	2.21	2.29	0.96	0.87	17.33	4.67	18.21	5.54	273.10			
YFR049W	YFR049w	0.92	1.42	1.64	2.09	2.19	2.23	3.34	3.08	12.44	2.46	15.76	5.54	287.00			
YHR44C	SS24	0.85	1.18	1.46	1.70	1.94	2.06	1.59	1.56	16.64	2.52	18.09	5.55	290.35			
YDR232W	YDR232w	0.90	1.24	1.43	1.89	2.00	2.04	0.11	0.04	11.53	4.51	12.64	5.59	270.00			
YOL059W	CSO2	0.98	1.19	1.34	1.71	1.91	2.04	1.47	1.51	11.51	4.45	12.68	5.59	264.49			
YGR111C	YSA1	0.97	1.26	1.52	1.97	2.16	2.28	1.35	0.68	22.23	4.92	23.58	5.60	284.09			
YOL070C	RP99	0.98	1.16	1.32	1.71	2.00	2.20	1.14	0.70	12.24	4.00	13.38	5.62	286.88			
YHR099C	YHR099c	0.98	1.54	1.72	2.08	2.21	2.29	3.73	3.08	18.45	2.62	22.19	5.63	284.69			
YPL238C	YPL238c	0.94	1.20	1.39	1.86	2.02	2.19	1.08	-0.07	27.79	5.70	28.86	5.63	282.88			
Y/JR022W	BAD7	0.95	1.30	1.46	1.84	2.08	2.26	0.42	0.61	5.75	5.03	6.17	5.64	272.26			
YDR489C	HEH2	0.97	1.35	1.60	1.97	2.22	2.30	1.69	0.77	24.53	4.88	26.22	5.65	296.68			
YOR494C	STD1	1.00	1.38	1.61	1.99	2.17	2.31	0.33	0.12	12.47	4.06	12.81	5.68	274.39			
YJR019C	IRS4	0.87	0.97	1.09	1.45	1.63	1.81	1.01	0.22	17.98	4.97	19.00	5.70	310.98			
YDR226W	ADK1	0.97	1.20	1.33	1.76	1.96	2.17	0.99	1.27	3.74	4.44	4.73	5.71	261.42			
YMR206W	YMR206w	1.01	1.24	1.41	1.80	2.03	2.25	0.40	0.38	13.54	5.34	13.94	5.71	271.78			
Y/JR107W	Y/JR107w	0.87	1.11	1.15	1.45	1.71	1.92	2.35	2.1	25.18	5.42	27.53	5.74	283.14			
YHR013C	ARD1	1.01	1.29	1.49	2.01	2.18	2.27	1.81	-0.01	18.65	5.75	20.45	5.74	243.62			
YPR120C	CLB5	0.87	1.06	1.14	1.60	1.84	2.05	3.03	3.09	1.93	5.57	2.27	2.78	279.68			
YBR220c	YBR220c	0.91	1.15	1.31	1.82	2.04	2.22	0.59	0.11	20.43	5.66	21.02	5.77	293.33			
YER090W	TRP2	0.92	1.34	1.48	1.82	2.01	2.21	2.33	3.00	2.99	10.39	2.91	13.39	5.90	286.21		
YMR253C	YMR253c	1.05	1.44	1.70	2.08	2.26	2.36	2.04	0.18	28.43	5.61	30.47	5.79	300.67			
Y/JR055W	HIT1	0.94	1.03	1.11	1.57	1.86	2.13	0.87	0.81	16.23	4.99	17.10	5.80	265.23			
YBR044W	MIS1	0.93	1.20	1.43	1.88	2.05	2.19	1.15	0.66	20.97	5.17	22.12	5.83	281.70			
YKL001C	MET14	1.00	1.33	1.54	2.04	2.24	2.37	1.08	0.97	10.05	4.87	11.12	5.85	239.66			
YFL034W	YFL034w	0.92	1.25	1.35	1.69	1.90	2.08	3.00	2.96	4.63	2.90	7.63	5.86	283.14			
YOR050C	DNL4	0.92	1.14	1.29	1.60	1.82	2.02	2.20	0.80	0.01	20.11	5.67	20.90	5.88	304.11		
YMR037C	MSN2	0.90	1.06	1.24	1.74	1.91	2.08	1.76	1.22	19.00	4.86	20.00	5.89	280.36			
YMR090W	YMR090w	0.99	1.43	1.61	2.02	2.21	2.33	2.13	0.95	16.84	5.08	18.16	6.83	284.13			
YMR244C-A	YMR244c-a	0.95	1.08	1.22	1.62	1.85	2.01	1.26	0.86	24.60	5.43	24.85	5.60	284.46			
YDR221W	GTB1	1.03	1.40	1.65	2.12	2.26	2.38	1.52	0.93	22.23	5.12	23.77	6.20	260.16			
YOR093C	ATX2	1.03	1.49	1.70	2.11	2.28	2.38	1.26	2.09	5.67	3.82	6.94	5.91	269.36			
YLL021W	SP42	0.95	1.20	1.32	1.58	1.90	2.03	2.22	1.54	2.22	5.94	8.48	5.94	187.80			
YBR224W	YBR224w	1.02	1.39	1.54	1.87	2.07	2.26	2.02	1.23	1.65	5.82	19.05	5.97	286.17			
Y/JL144W	Y/JL144w	1.02	1.39	1.60	1.92	2.02	2.20	2.02	1.23	1.11	4.00	1.52	2.77	284.53			
YER088W	UTR1	0.98	1.20	1.39	1.78	1.95	2.05	2.02	2.62	2.44	2.31	5.78	2.23	285.23			
YLR213W	ARL3	0.86	1.09	1.17	1.54	1.74	1.93	3.03	3.47	5.17	2.67	8.20	6.14	254.53			
YKR161W	SIR1	0.96	1.43	1.57	2.05	2.24	2.38	3.30	2.99	13.81	5.24	14.83	6.15	297.00			
YOL131W	YOL131w	0.96	1.51	1.66	2.04	2.20	2.23	3.12	3.20	9.40	3.23	12.52	6.43	263.98			
Y/JR032W	CPR7	0.93	1.09	1.20	1.54	1.82	1.96	2.05	1.								

ORF.name	Strain	Optical density at 600 nm at regular intervals (hours)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)		TOTAL AMINO ACIDS (mg/L)	
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	102 h	118 h	
YGL135W	RPL1b	0.94	1.32	1.61	2.02	2.33	0.97	0.61	0.70	0.67	0.70	0.72	0.72	0.72	0.72	
YCH059W	CPR4	0.90	0.68	1.28	1.77	1.95	1.15	0.77	1.03	1.05	1.05	1.02	1.02	1.02	1.02	
YIL027C	CPB1	0.93	1.44	1.76	1.87	2.08	2.25	1.12	1.22	1.22	1.23	1.19	1.20	1.21	1.21	
YKL065C	YET1	0.99	1.30	1.51	1.99	2.18	2.29	0.85	1.12	1.12	1.13	1.13	1.14	1.14	1.14	
YHL047C	ARN2	1.06	1.39	1.62	2.16	2.33	2.43	0.94	0.71	0.74	0.63	0.63	0.63	0.63	0.63	
YLR023W	YLR023w	0.88	1.31	1.51	1.93	2.12	2.23	3.44	0.05	17.04	4.28	20.49	7.34	205.21	7.34	
YLR135W	SLX4	0.95	1.20	1.33	1.76	1.97	2.06	1.87	0.98	27.77	6.37	28.64	7.35	284.08	7.35	
YML049W	APP1	0.89	1.23	1.44	1.84	2.02	2.21	0.82	0.70	11.22	6.65	12.04	7.35	292.59	7.35	
YLR096W	NCR81	0.93	1.33	1.46	1.79	1.95	2.12	3.21	3.47	4.18	3.91	7.38	7.39	253.80	7.39	
YGR237C	YGR237c	0.99	1.38	1.55	1.98	2.12	2.25	3.46	3.07	19.69	4.31	23.16	7.39	300.24	7.39	
YIL035C	CKA1	0.94	1.52	1.65	2.00	2.13	2.23	2.88	2.98	7.05	4.43	9.33	7.41	270.04	7.41	
YIL111W	CAR1	0.95	1.19	1.36	1.82	2.03	2.24	2.46	-0.04	3.48	7.48	5.94	7.44	225.10	7.44	
YDR371W	CTS2	1.02	1.33	1.55	1.99	2.15	2.32	0.46	-0.06	22.78	7.54	23.24	7.48	312.96	7.48	
YDL013W	SLX5	0.85	0.98	1.09	1.41	1.65	1.84	0.77	0.85	6.54	6.64	7.31	7.49	263.85	7.49	
YOR384W	FRE5	0.97	1.29	1.50	1.90	2.06	2.20	0.61	0.00	20.59	7.50	21.20	7.50	272.95	7.50	
YDL114W	YDL114w	0.93	1.28	1.40	1.74	1.98	2.19	0.79	0.84	8.37	6.68	9.17	7.52	287.14	7.52	
YHL003C	LAG1	0.91	1.20	1.34	1.69	1.90	2.07	1.54	1.37	17.56	6.16	19.09	7.53	294.74	7.53	
YDL174C	DLD1	0.96	1.35	1.52	1.89	2.12	2.30	1.05	0.67	22.02	6.86	23.06	7.54	302.46	7.54	
YKL148C	SDH1	0.99	1.29	1.50	1.97	2.14	2.24	1.54	0.97	21.74	6.57	23.27	7.54	271.94	7.54	
YJR062C	NTA1	0.90	1.24	1.36	1.70	1.92	2.12	0.03	0.88	28.83	6.67	30.86	7.54	272.24	7.54	
YOR367W	SCP1	1.04	1.36	1.60	1.99	2.20	2.16	0.46	0.32	17.52	7.22	17.98	7.55	279.11	7.55	
YOR064C	VNG1	0.99	1.42	1.65	1.98	2.15	2.27	0.26	-0.01	15.73	7.66	15.98	7.65	278.04	7.65	
YKL124W	SSH4	0.85	1.03	1.14	1.66	1.88	2.02	1.20	1.11	12.27	6.56	13.47	7.67	271.26	7.67	
YKL123W	YKL123w	1.01	1.40	1.62	2.07	2.22	2.31	1.49	1.24	21.37	6.41	22.85	7.69	289.22	7.69	
YLR151C	PCD1	0.86	0.99	1.08	1.71	1.96	2.12	1.39	0.85	25.47	6.86	26.86	7.70	284.56	7.70	
YEL049W	PAU2	0.92	1.27	1.48	1.96	2.11	2.22	2.02	0.08	21.31	7.60	31.33	7.75	289.45	7.75	
YJR152W	DAL5	1.09	1.60	1.75	2.14	2.28	2.41	3.42	3.08	15.94	4.69	19.37	7.77	253.03	7.77	
YLN274C	GOR1	0.93	1.38	1.55	1.96	2.13	2.24	4.95	3.17	28.80	4.60	33.74	7.76	289.45	7.76	
YCR022C	YCR022c	0.92	1.26	1.42	1.88	2.03	2.12	1.20	1.14	7.62	6.64	8.82	7.76	275.82	7.76	
YIL023C	YKE4	0.89	1.41	1.56	1.93	2.07	2.15	3.15	3.07	7.75	4.73	10.90	7.80	256.59	7.80	
YIL272C	ALP1	1.00	1.45	1.61	2.03	2.19	2.28	0.58	0.07	18.40	7.74	18.99	7.81	274.71	7.81	
YML049W	SPG5	1.00	1.55	1.79	2.14	2.29	2.36	0.72	0.40	14.75	7.22	14.88	7.82	264.70	7.82	
YIL040C	CBR1	0.95	1.58	1.74	2.11	2.20	2.18	3.32	3.07	17.29	7.29	21.39	7.83	284.19	7.83	
YMR174C	PAF1	0.94	1.24	1.35	1.65	1.95	2.07	1.33	0.17	18.16	7.76	18.49	7.87	233.36	7.87	
YMR303C	ADH2	1.07	1.43	1.68	2.11	2.27	2.43	3.46	0.68	25.68	7.01	29.44	7.89	289.47	7.89	
YBR116W	PCII2	1.01	1.21	1.40	1.90	2.12	2.30	0.97	0.63	24.68	7.28	25.66	7.91	283.49	7.91	
YLR443C	ECM30	0.95	1.31	1.50	1.90	2.14	2.24	3.89	3.09	18.78	4.83	23.67	7.82	301.40	7.82	
YOR107W	MCA1	0.90	0.98	1.00	1.20	1.36	1.47	1.18	2.06	16.93	5.88	18.11	7.94	295.19	7.94	
YDR475C	JIP4	0.89	1.21	1.37	1.74	1.93	2.11	2.06	0.37	28.76	7.04	31.82	7.95	284.79	7.95	
YML129W	NRK1	0.91	1.31	1.51	1.89	2.07	2.15	1.52	0.37	20.17	7.08	21.68	7.95	266.33	7.95	
YPR026W	ATH1	0.99	1.22	1.46	1.88	2.08	2.23	2.14	0.79	30.55	7.17	32.69	7.97	282.64	7.97	
YBR095C-A	YBR095c-a	0.94	1.24	1.40	1.75	1.96	2.07	0.05	0.45	24.42	7.17	26.49	8.02	314.59	8.02	
YIL007C	STE20	0.92	1.22	1.39	1.89	2.07	2.16	1.20	-0.04	29.58	8.08	30.79	8.03	290.57	8.03	
YMR163C	INP2	0.95	1.12	1.30	1.73	1.94	2.16	1.06	0.16	25.17	7.88	26.23	8.04	296.76	8.04	
YEL100W	SEH1	0.81	1.01	1.13	1.64	1.93	2.10	0.15	0.85	1.29	7.22	1.44	8.07	253.01	8.07	
YDR480W	DIG2	0.96	1.46	1.68	2.02	2.12	2.25	2.95	0.80	29.41	7.29	32.36	8.09	276.85	8.09	
YLR338W	YLR338w	0.92	1.14	1.29	1.58	1.80	1.95	1.84	0.71	25.71	7.39	27.55	8.10	258.82	8.10	
YML022W	APT1	0.95	1.16	1.23	1.42	1.61	1.76	2.48	2.28	16.57	6.82	29.42	8.10	326.86	8.10	
YGR108W	CLB1	0.97	1.24	1.34	1.70	1.95	2.15	2.21	0.85	16.84	7.27	21.27	8.12	275.34	8.12	
YJR055W	HIT1	0.95	1.27	1.33	1.67	1.96	2.11	2.10	0.95	21.00	7.19	22.10	8.14	305.62	8.14	
YCR102w-A	YCR102w-a	0.87	1.00	1.16	1.55	1.76	1.98	1.97	0.59	29.06	7.56	31.02	8.14	285.43	8.14	
YEL033W	MTC7	0.91	1.21	1.39	1.81	1.99	2.08	1.71	0.16	29.84	8.05	31.55	8.20	265.52	8.20	
YLR097C	HRT3	0.93	1.22	1.38	1.75	1.91	2.15	2.44	0.21	22.86	7.04	25.31	8.23	292.77	8.23	
YGR050C	YGR050c	0.90	1.06	1.26	1.70	1.92	2.12	2.18	0.83	14.44	7.41	15.52	8.25	270.41	8.25	
YPR042C	PUF2	0.88	1.31	1.49	1.85	2.02	2.17	3.05	2.88	18.67	5.39	21.72	8.26	304.58	8.26	
YOR096C	TSR3	1.01	1.23	1.28	1.73	1.93	2.01	2.24	0.24	16.93	6.84	19.80	7.87	266.26	7.87	
YMR142C	YMR142c	0.92	1.24	1.34	1.70	1.95	2.12	2.22	0.74	21.39	7.35	24.61	8.59	282.04	8.59	
YLR057C	YIL057c	1.03	1.60	1.74	2.09	2.20	2.29	2.58	2.93	5.20	5.59	16.78	8.57	264.39	8.57	
YOR115C	TR533	1.08	1.51	1.72	2.15	2.29	2.36	2.51	0.98	30.85	7.63	33.36	8.53	300.33	8.53	
YMR149W	DUS4	1.00	1.27	1.48	1.90	2.07	2.24	2.04	0.96	26.74	7.60	28.78	8.56	289.25	8.56	
YOR365C	YOR365c	1.04	1.37	1.60	2.00	2.13	2.36	0.40	0.19	14.62	8.38	15.02	8.57	272.02	8.57	
YOR023C	AHC1	0.90	1.25	1.46	1.86	2.02	2.20	2.35	0.39	1.06	13.28	7.66	13.67	8.72	273.77	8.72
YOR010C	TRR2	1.00	1.49	1.67	2.01	2.16	2.28	0.44	0.12	16.99	6.35	17.43	8.74	279.69	8.74	
YLR171C	YJL171c	1.04	1.22	1.42	1.64	2.10	2.24	2.41	1.22	25.49	7.35	28.75	8.74	282.04	8.74	
YLR020C	KEX2	0.94	1.29	1.42	1.83	2.02	2.14	3.70	3.45	22.83	5.22	26.53	8.68	304.98	8.68	
YGL082W	YGL082w	0.99	1.52	1.68	2.06	2.22	2.35	3.20	3.06	17.21	5.64	20.40	8.70	309.95	8.70	
YMR041C	ARA2	1.00	1.25	1.49	1.97	2.15	2.35	0.36	0.16	17.48	6.55	17.83	8.71	291.78	8.71	
YHR171W	ARO9	0.92	1.21	1.41	1.96	2.15	2.23	1.10	1.32							

ORF.name	Strain	Optical density at 600 nm at regular intervals (hours)							GLUCOSE		FRUCTOSE		TOTAL SUGARS (g/L)	TOTAL AMINO ACIDS (mg/L)
		0	26	33	48	59	78	102 h	118 h	102 h	118 h	102 h	118 h	
YPL020C	EL23	1.10	1.47	1.62	2.10	2.24	2.36	1.94	0.95	28.50	5.84	30.46	7.75	30.46
YGR250c	EL23	0.93	1.10	1.26	1.75	1.96	1.16	1.48	0.68	20.67	9.21	22.45	9.29	274.95
YDR401W	YD2401w	0.90	1.18	1.34	1.80	1.95	2.08	1.27	0.59	23.53	6.08	27.89	8.91	264.59
YOR033C	EX01	1.07	1.50	1.79	2.16	2.35	2.42	1.12	0.74	23.39	9.78	27.11	8.92	284.27
YGR291C	YGR291c	0.98	1.21	1.38	1.74	1.92	2.08	0.97	0.90	16.58	9.09	17.55	10.88	271.27
YML103W-A	YM100w-a	1.03	1.84	1.84	2.23	2.35	2.44	3.85	3.07	23.60	6.93	27.45	10.88	276.81
YCR095C	OC44	0.87	0.99	1.15	1.63	1.91	2.14	1.10	0.71	19.70	9.32	20.89	10.83	265.67
YBR126C	TPS1	0.93	1.28	1.44	1.79	2.04	2.24	0.67	0.72	5.69	9.33	5.76	10.86	298.15
YDL171C	GLT1	0.98	1.42	1.55	1.93	2.13	2.28	0.71	0.74	8.66	9.32	9.37	10.86	280.64
YGR089W	CTT1	0.98	1.20	1.41	1.94	2.14	2.22	1.84	1.05	25.46	9.01	27.30	10.86	287.30
YBR141C	YBR141c	0.87	1.13	1.34	1.77	1.98	2.06	0.63	0.80	7.73	9.28	8.36	10.88	267.15
YDR221C	MCT1	0.93	1.25	1.44	1.78	2.00	2.15	1.80	1.38	20.68	8.71	22.49	10.89	277.00
YMR246W	FAA4	1.07	1.28	1.48	1.91	2.13	2.33	0.77	0.98	26.56	10.11	27.63	10.19	305.18
YDR504C	SPG3	0.96	1.32	1.56	2.00	2.18	2.26	1.94	0.86	26.70	9.35	28.64	10.21	284.70
YMR219W	ESC1	0.95	1.15	1.38	1.84	2.10	2.48	0.72	0.24	20.81	9.95	21.52	10.21	278.12
YGR045C	YGR045c	1.06	1.44	1.67	2.16	2.32	2.39	1.74	1.00	27.30	9.23	29.04	10.22	308.56
YAL007C	ERP2	0.95	1.18	1.43	1.79	1.95	2.16	3.16	1.56	28.96	8.67	32.12	10.23	220.25
YGL212W	VAM7	0.89	1.10	1.17	1.45	1.60	1.83	0.72	1.01	28.88	9.23	29.60	10.24	297.73
YGR085C	RPL11b	1.03	1.31	1.47	1.96	2.18	2.29	1.09	1.04	18.69	9.21	19.78	10.25	275.50
YGR089W	NVF2	0.95	1.19	1.44	1.95	2.14	2.30	3.25	0.85	20.58	9.42	23.83	10.26	287.06
YCR048W	ARE1	0.90	1.18	1.31	1.66	1.87	2.07	1.57	1.30	15.43	9.00	17.00	10.30	265.83
YDR514C	YDR514c	0.94	1.31	1.52	1.94	2.12	2.22	1.61	0.94	21.30	9.38	22.91	10.32	222.80
YLR216C	CPR6	0.97	1.23	1.42	2.00	2.22	2.36	1.26	0.75	22.03	9.61	23.29	10.36	294.54
YDR500C	RPL37b	0.89	1.16	1.26	1.69	1.98	2.11	2.14	0.86	29.31	9.50	31.45	10.37	229.55
YPR040W	TIP41	0.99	1.40	1.61	1.95	2.11	2.25	4.11	2.86	24.31	7.51	28.41	10.37	293.85
YGL260W	YGL260w	0.99	1.31	1.47	1.90	2.08	2.26	1.24	1.08	16.84	9.30	18.08	10.37	294.49
YMR237W	PRC1	1.00	1.32	1.58	1.98	2.17	2.35	0.56	2.57	17.23	7.86	17.79	10.43	290.19
YIL138C	VHS2	1.02	1.10	1.29	1.57	1.69	2.00	0.85	1.15	12.45	9.29	13.30	10.44	253.80
YOR037W	CYC2	1.03	1.45	1.72	2.13	2.27	2.40	0.42	0.42	16.68	8.14	17.10	10.45	273.09
YDR509W	YDR509w	0.98	1.37	1.56	1.87	2.07	2.12	1.19	0.78	16.12	9.67	17.31	10.45	317.70
YJL199C	YJL199c	0.96	1.25	1.45	1.80	2.05	2.21	1.69	1.14	23.33	9.31	25.04	10.47	292.84
YLR202C	FLA1	0.92	1.69	1.28	1.76	1.97	2.08	1.42	1.06	24.47	9.42	22.88	10.48	265.67
YLR208W	MIG2	0.98	1.38	1.59	2.03	2.05	2.35	0.89	0.51	7.85	9.01	8.17	10.49	266.67
YLR208C	YLR208c	1.00	1.39	1.47	1.91	2.08	2.23	1.04	0.79	18.22	9.57	18.36	10.56	308.00
YLL090C	GTT2	0.94	1.13	1.32	1.69	1.86	2.10	1.40	1.16	4.18	9.43	5.58	10.59	143.04
YCR087W	YCR087w	0.89	1.25	1.38	1.73	1.91	2.03	1.42	0.70	26.58	9.95	28.40	10.65	293.08
YGR122C-A	YGR122c-a	0.95	1.20	1.41	1.85	2.06	2.18	2.15	0.77	28.31	9.84	31.06	10.61	309.95
YOL015W	IRC10	1.00	1.36	1.58	1.98	2.14	2.31	1.59	0.17	7.07	10.49	8.66	10.65	273.25
YCR089W	YCR089w	0.89	1.25	1.38	1.73	1.91	2.03	1.42	0.70	26.58	9.95	28.40	10.65	293.08
YOR043W	WH2	1.04	1.39	1.58	2.07	2.26	2.38	0.52	-0.10	23.10	10.86	23.63	10.76	290.64
YDR193W	YDR193w	0.92	1.14	1.24	1.63	1.80	1.98	1.86	1.14	23.84	9.63	25.70	10.77	271.86
YML226W	YN122w	0.98	1.41	1.51	1.93	2.10	2.25	3.11	3.21	12.07	7.56	15.18	10.77	273.67
YJL063W	LAS21	1.00	1.48	1.69	2.06	2.21	2.29	3.34	3.08	14.71	7.69	18.05	10.78	277.13
YDR447C	RPS17b	0.97	1.29	1.48	1.89	2.14	2.26	2.02	0.95	25.23	9.86	27.25	10.81	305.47
YAR027W	UIP3	0.89	1.16	1.39	1.70	1.85	2.06	3.18	1.55	24.30	9.29	27.48	10.84	191.65

Appendix 5. Shortlisted 299 candidates from the WYDL_HN based on sugars at the (67 h) time point.

ORF	ODs (h)						GLUCOSE		FRUCTOSE		TOTAL sugars (g/L)		TOTAL AMINO ACIDS (mg/L)	
	0	20	27	42	67	43 h	67 h	43 h	67 h	43 h	67 h	48 h		
YOR13W	0.96	1.56	1.82	2.18	2.26	13.29	-0.07	70.36	1.54	83.65	1.47	180.14		
YBR299W	0.89	1.51	1.87	2.12	2.27	10.37	-0.25	74.34	2.04	84.71	1.80	68.15		
YGR259C	0.96	1.80	2.05	2.24	2.37	10.80	0.25	65.71	2.46	79.71	2.71	253.95		
YGR259W	1.04	1.62	1.62	2.24	2.27	12.03	0.10	81.63	2.17	93.85	2.11	273.93		
YMR029C	1.03	1.85	2.09	2.34	2.36	10.63	-0.03	77.63	2.98	88.28	2.95	273.96		
YPR123W_A	0.95	1.46	1.85	1.94	2.16	22.17	2.81	98.68	0.73	120.83	3.54	45.08		
YPR063C	0.98	1.92	2.16	2.34	2.36	10.59	1.49	69.63	2.20	80.22	3.69	161.27		
YLR035C	0.97	1.84	1.72	2.04	2.08	16.44	-0.26	94.19	4.00	110.55	3.74	324.06		
YGR070W	0.95	1.77	2.01	2.25	2.30	11.78	0.33	62.87	3.48	79.65	3.80	68.89		
YNL252C	1.01	1.78	1.99	2.25	2.26	9.23	0.26	69.80	3.88	79.03	4.14	96.94		
YNL078W	1.00	1.80	1.92	2.11	2.19	7.32	0.04	57.74	4.13	65.05	4.17	9.93		
YPR149W	1.10	1.80	2.07	2.40	2.47	25.90	0.45	98.39	3.86	124.29	4.31	-3.34		
YJL020C	0.96	1.63	1.90	2.21	2.25	15.48	-0.25	91.50	4.74	106.98	4.49	238.51		
YPR058W	0.98	1.75	2.13	2.30	2.32	7.35	1.19	62.08	3.47	69.43	4.66	100.40		
YGR101W	0.99	1.46	1.79	2.22	2.32	33.00	1.55	103.37	3.68	136.36	5.23	158.03		
YNL145W	0.90	1.31	1.43	1.70	1.80	29.32	1.82	99.16	3.47	128.48	5.28	405.30		
YGL016W	0.87	1.60	1.84	2.07	2.14	11.03	0.18	60.79	5.12	71.79	5.30	46.72		
YGL080W	0.95	1.80	2.02	2.24	2.28	9.71	0.30	58.67	5.17	68.38	5.47	187.65		
YDL082W	1.01	1.98	2.06	2.16	2.24	8.90	0.50	65.79	4.99	74.69	5.50	74.07		
YL055C	0.91	1.56	1.80	2.08	2.14	8.47	0.25	34.40	5.26	42.87	5.51	271.19		
YGR132C	1.00	1.86	2.07	2.25	2.32	9.98	-0.07	74.40	5.92	84.28	5.86	150.49		
YGR069C	1.02	2.05	1.82	1.95	2.08	9.51	-0.02	67.31	5.96	76.82	5.95	48.90		
YHR071W	0.94	1.55	1.83	2.17	2.18	17.94	0.50	95.52	5.71	113.46	6.21	322.72		
YJL142C	0.96	1.87	2.08	2.34	2.34	6.08	0.24	52.52	6.10	58.61	6.34	129.16		
YPL230W	0.92	1.38	1.61	1.90	1.97	9.93	1.92	58.42	4.44	68.35	6.37	180.04		
YGR109C	0.96	1.44	1.78	2.26	2.34	30.01	1.80	100.26	4.70	130.27	6.50	208.79		
YOR252W	1.03	1.88	2.17	2.32	2.39	15.52	1.09	66.13	5.45	81.64	6.54	231.81		
YGR067C	1.05	1.57	1.92	2.35	2.39	39.37	1.34	128.93	5.24	168.30	6.58	141.33		
YJR100C	1.01	1.97	2.18	2.29	2.37	10.77	0.40	54.81	6.27	65.58	6.66	292.38		
YGL159W	0.94	1.55	1.79	2.13	2.21	19.08	0.61	73.32	6.21	92.40	6.82	256.91		
YGR052W	0.98	1.48	1.76	2.25	2.23	28.95	0.42	121.96	6.43	150.91	6.85	178.92		
YGR047W	0.93	1.44	1.79	2.20	2.26	32.80	0.73	125.36	6.21	158.16	6.94	98.15		
YGR143W	0.95	1.53	1.83	2.22	2.29	12.70	0.03	70.07	6.98	82.77	7.01	355.30		
YGR153W	1.01	1.59	1.86	2.24	2.28	22.49	-0.02	78.45	7.27	84.24	7.63	160.05		
YGR070C	1.19	2.38	2.45	2.55	2.58	15.58	1.80	75.50	5.25	93.14	7.18	265.25		
YGR050W	1.02	1.58	1.67	2.24	2.34	10.90	0.50	133.38	6.79	171.09	7.28	138.73		
YL070W	0.83	1.78	2.09	2.22	2.29	8.99	0.11	49.83	7.29	58.79	7.40	352.74		
YAL018C	0.88	1.61	1.89	2.23	2.31	16.93	1.76	87.42	5.65	104.35	7.41	167.59		
YCL010C	0.97	1.61	1.89	2.21	2.35	10.91	-0.16	60.28	7.63	71.17	7.47	235.04		
YDR269W	0.96	1.50	1.73	2.14	2.22	11.06	1.61	47.70	6.09	58.76	7.70	288.28		
YBL144W	1.08	1.86	2.10	2.37	2.41	11.17	1.52	68.80	6.21	79.98	7.72	254.64		
YGR043C	0.96	1.25	1.56	2.09	2.21	33.42	1.82	105.06	6.09	139.48	7.90	277.17		
YGR041W	0.99	1.32	1.69	2.22	2.30	40.56	2.24	127.99	5.76	168.55	8.01	180.67		
YER090C	1.04	1.62	1.88	2.27	2.31	14.96	1.37	52.46	7.37	67.42	8.75	295.77		
YDL156W	1.01	1.80	2.10	2.32	2.41	13.99	-0.16	74.20	8.98	88.18	8.82	105.49		
YGR266W	0.96	1.89	2.01	2.10	2.10	8.33	0.47	56.92	8.49	65.25	8.96	163.40		
YL156W	0.96	1.64	1.91	2.17	2.26	6.96	0.21	52.93	8.82	59.89	9.02	112.36		
YKR088C	0.96	1.66	1.76	2.02	2.11	14.60	-0.05	74.94	9.19	89.55	9.14	182.15		
YKL173C	1.00	1.75	2.01	2.20	2.30	16.95	0.09	70.66	9.12	87.61	9.21	312.08		
YGR059W	0.94	1.39	1.72	2.18	2.23	33.22	1.82	119.27	7.43	152.48	9.25	231.11		
YMR171C	0.95	1.50	1.77	2.10	2.22	18.03	1.26	81.87	8.02	99.90	9.27	308.01		
YGR044C	0.93	1.23	1.54	2.05	2.20	31.28	2.74	102.90	6.66	134.18	9.40	310.35		
YOR274C	0.99	1.51	1.82	2.08	2.17	11.77	0.35	76.16	9.12	87.93	9.47	204.14		
YGR040W	1.07	2.06	2.29	2.38	2.38	11.97	0.04	72.03	9.48	83.44	9.52	154.24		
YGR048C	0.98	1.72	1.97	2.24	2.31	2.31	0.36	64.50	9.64	75.75	9.53	109.82		
YAL159W	0.99	1.69	1.89	2.24	2.35	10.47	0.16	62.80	9.36	73.13	9.54	115.55		
YHR269W	1.00	1.69	1.83	2.17	2.26	16.45	-0.19	69.22	9.84	85.67	9.65	314.61		
YGL196C	0.98	1.53	1.85	2.21	2.28	13.43	1.08	123.52	9.53	135.86	9.70	123.24		
YGR123C	0.94	1.23	1.51	2.04	2.14	35.57	2.26	112.00	6.46	149.57	9.73	169.38		
YEL007W	1.01	1.58	1.88	2.21	2.29	10.53	0.84	56.33	9.25	66.86	10.08	257.56		
YGL064C	0.96	1.60	1.72	2.01	2.10	13.27	0.23	67.55	9.84	80.82	10.27	115.23		
YGL079W	0.98	1.84	2.04	2.26	2.31	12.67	0.66	65.67	9.64	78.14	10.31	111.35		
YER174C	0.96	1.63	1.90	2.19	2.26	13.69	-0.29	68.90	10.09	82.58	9.80	137.37		
YGR016W	0.93	1.65	1.95	2.21	2.27	14.24	-0.14	64.49	10.04	83.69	9.90	67.62		
YGL246C	0.98	1.67	1.95	2.20	2.26	16.87	-0.27	92.53	10.24	109.40	9.97	80.68		
YDR030W	0.98	1.67	1.98	2.21	2.31	16.36	0.38	74.84	9.62	91.20	10.01	195.95		
YIR032C	0.98	1.89	2.10	2.25	2.35	16.18	-0.18	81.22	10.20	97.41	10.02	316.07		
YGR020W	0.91	1.48	1.79	2.21	2.24	12.92	1.18	60.47	9.50	73.39	10.68	74.87		
YGL242C	0.88	1.38	1.73	2.06	2.15	22.50	0.29	98.15	10.58	120.65	10.86	164.14		
YGL239W	0.92	1.23	1.53	2.05	2.18	43.90	0.45	136.75	10.49	180.65	10.93	166.08		
YLR095C	0.99	1.72	2.00	2.22	2.31	12.67	-0.07	77.12	11.03	89.78	10.95	206.35		
YDL178W	1.03	1.76	2.07	2.31	2.38	13.11	-0.09	70.56	11.12	83.67	11.03	142.08		
YOR101W	1.02	1.74	1.97	2.08	2.18	10.68	0.43	63.77	10.74	74.45	11.17	109.84		
YDR86C	1.07	1.65	1.93	2.20	2.26	11.38	0.29	49.88	8.97	60.66	11.26	265.45		
YDR302C	1.06	1.91	2.14	2.31	2.41	10.49	0.39	66.20	8.22	76.69	11.31	87.09		
YHR040C	0.94	1.36	1.65	2.12	2.18	14.70	0.36	82.57	10.97	97.27	11.33	151.86		
YGR020W	0.													

ORF	ODs (h)						GLUCOSE		FRUCTOSE		TOTAL sugars (g/L)		TOTAL AMINO ACIDS (mg/L)	
	0	20	27	42	67	43 h	67 h	43 h	67 h	43 h	67 h	48 h		
YGR004W	1.04	1.85	2.05	2.23	2.30	18.48	-0.26	94.84	127.3	113.32	12.47	65.83		
YDL174C	0.91	1.63	1.94	2.26	2.33	17.32	0.59	79.61	11.89	96.94	12.48	168.69		
YGR080W	1.02	1.72	1.96	2.25	2.33	15.53	0.14	80.76	12.39	96.28	12.53	120.08		
YPL121C	0.96	1.39	1.78	2.02	2.12	17.52	1.17	89.27	11.41	106.80	12.58	113.24		
YKL070W	1.00	1.66	1.76	2.04	2.13	15.68	0.29	64.56	12.34	106.80	12.62	125.42		
YKL072C	1.00	1.53	1.95	2.19	2.25	17.90	0.67	102.60	12.07	129.55	12.62	117.02		
YKL072C	1.08	1.66	1.78	2.07	2.16	16.15	0.47	69.22	12.48	85.36	12.65	106.80		
YPL141W	0.98	1.45	1.74	2.08	2.20	18.29	0.26	73.09	12.03	102.60	12.65	102.94		
YPL149C	0.98	1.76	2.05	2.28	2.32	18.32	0.79	73.28	11.87	85.70	12.66	119.62		
YKL120W	1.03	1.79	1.99	2.23	2.31	21.08	0.37	83.68	12.28	104.73	12.66	113.21		
YML176C	0.78	1.02	1.51	1.78	1.91	13.47	1.22	77.50	11.47	90.97	12.69	85.58		
YGL257C	0.98	1.55	1.80	2.11	2.21	20.96	0.64	94.29	12.09	115.25	12.73	148.82		
YIL050W	0.88	1.30	1.51	1.84	1.91	15.68	0.23	68.08	12.55	83.76	12.78	227.86		
YER016C	1.08	1.77	2.04	2.35	2.38	23.16	1.62	87.34	11.17	110.50	12.80	202.90		
YGL191W	0.85	1.27	1.56	1.92	2.04	23.69	0.61	103.72	12.19	127.40	12.81	147.07		
YFL051C	0.93	1.76	1.91	1.99	2.15	15.59	0.73	64.27	12.10	79.86	12.83	102.48		
YGL013C	0.98	1.63	1.88	2.23	2.29	17.07	1.40	80.94	11.52	98.01	12.92	243.33		
YGL129C	0.97	1.77	1.99	2.20	2.25	13.82	0.44	78.38	12.55	92.21	12.99	144.82		
YDL169W	0.97	1.63	2.24	1.99	2.00	13.41	0.03	68.22	12.98	81.63	13.01	146.26		
YFL053W	0.93	2.32	1.94	1.72	1.72	12.53	0.89	59.67	12.16	72.20	13.04	220.19		
YDR374W	0.96	1.58	1.80	2.08	2.13	18.94	0.68	86.23	12.44	105.18	13.12	370.65		
YGR015C	0.96	1.69	1.99	2.25	2.32	23.41	0.49	100.29	12.70	123.70	13.19	175.24		
YGL178W	1.10	1.76	2.07	2.35	2.42	16.77	1.80	67.96	11.54	84.73	13.34	332.81		
YPR117W	1.04	1.78	2.08	2.31	2.38	13.88	1.29	80.98	12.05	94.86	13.34	124.84		
YJR133W	1.03	1.67	1.86	2.13	2.22	19.93	0.14	76.16	13.23	96.09	13.38	98.70		
YBR269W	1.03	1.92	2.20	2.36	2.44	13.51	0.09	73.17	13.31	86.68	13.40	154.65		
YPL027W	0.98	1.40	1.68	2.00	2.14	25.97	1.47	97.81	11.93	123.78	13.40	314.83		
YJR025C	0.99	1.55	1.77	2.04	2.13	30.53	0.18	88.17	13.25	118.70	13.43	155.37		
YGR212W	0.89	1.41	1.65	1.97	2.04	12.97	0.35	56.33	13.11	69.30	13.46	149.71		
YPL202C	0.95	1.32	1.75	1.98	2.04	20.18	1.53	91.61	11.96	111.79	13.49	154.75		
YGL131C	0.96	1.47	1.76	2.18	2.22	17.69	0.75	68.00	11.74	85.69	13.49	175.66		
YLR029W	1.01	1.82	2.11	2.31	2.39	17.42	0.17	70.04	13.36	87.45	13.53	194.99		
YDL125W	0.99	1.73	2.17	2.15	2.24	15.73	0.46	71.03	13.07	86.77	13.53	243.95		
YDL100C	0.98	1.56	1.84	2.19	2.31	22.04	0.15	91.53	13.58	113.57	13.73	207.02		
YDL129W	0.90	1.41	1.71	2.09	2.19	23.23	0.11	98.09	13.63	121.32	13.74	124.96		
YER161C	1.17	1.96	2.18	2.44	2.48	14.17	0.56	61.59	12.16	76.32	13.74	143.44		
YDL144W	0.99	1.61	1.88	2.17	2.23	16.45	0.65	78.84	13.61	91.29	13.76	124.4		
YER162C	0.99	1.64	1.84	2.14	2.22	19.00	0.69	86.96	12.53	108.93	13.71	124.69		
YIL038W	1.04	1.54	1.82	2.05	2.22	16.53	0.23	10.04	12.49	89.44	13.61	217.65		
YDR374W	1.10	1.86	1.88	2.27	2.33	17.24	1.78	59.53	12.09	76.77	13.87	104.98		
YCR146C	0.98	1.87	1.95	2.24	2.29	17.22	0.26	78.38	13.64	96.11	13.89	128.87		
YHR142W	0.95	1.19	1.30	1.84	1.98	24.83	0.41	98.50	13.52	123.34	13.93	131.29		
YPR145W	1.03	1.78	2.55	2.47	2.39	22.37	1.12	100.95	12.82	123.32	13.94	80.67		
YBR264C	1.00	1.74	2.03	2.26	2.37	12.17	0.09	61.64	14.06	79.80	13.87	98.15		
YML265C	1.08	1.81	2.07	2.33	2.37	15.22	0.38	70.42	13.61	85.64	13.99	151.11		
YER137C	0.93	1.37	1.59	1.90	2.00	18.98	1.45	88.61	12.57	107.59	14.02	99.66		
YER030W	1.01	1.73	2.02	2.29	2.34	10.88	0.25	94.14	14.08	113.02	14.33	78.83		
YUH142C	0.94	1.74	2.06	2.30	2.41	22.29	1.52	85.39	12.82	107.68	14.35	116.66		
YDL109C	1.03	1.80	2.06	2.27	2.39	19.16	0.14	81.72	14.20	100.89	14.35	133.69		
YGL227W	0.97	1.44	1.75	2.20	2.29	38.14	0.16	64.18	14.42	82.30	14.58	207.38		
YDL175C	0.95	1.54	1.80	2.15	2.29	20.01	0.74	80.35	13.50	100.35	14.25	239.54		
YDR255C	1.04	1.91	2.04	2.25	2.36	10.95	-0.07	65.38	14.33	76.33	14.26	54.62		
YKR066C	0.96	1.69	1.96	2.19	2.28	14.57	0.44	75.27	13.83	89.84	14.27	254.98		
YER143W	0.96	1.53	1.79	2.15	2.22	34.02	0.60	134.29	13.68	168.31	14.28	114.99		
YER030W	1.01	1.73	2.02	2.29	2.34	18.88	0.25	94.14	14.08	113.02	14.33	78.83		
YUH142C	0.94	1.74	2.06	2.30	2.41	21.00	1.52	15.79	12.82	107.68	14.35	116.66		
YDL130W	0.97	1.44	1.75	2.20	2.29	38.14	0.45	142.23	13.92	180.37	14.36	220.02		
YGL208W	0.97	1.62	1.90	2.07	2.23	18.36	0.26	90.16	14.11	108.52	14.37	113.98		
YGR142W	0.97	1.48	1.76	2.15	2.24	33.63	0.46	133.24	13.94	166.87	14.40	143.54		
YGR045C	1.00	1.54	1.81	2.20	2.31	33.50	0.77	128.26	13.72	161.76	14.49	124.09		
YPL198W	1.05	1.73	2.13	2.31	2.39	10.18	1.50	66.65	13.00	76.83	14.50	110.28		
YER168C	1.03	1.82	2.06	2.30	2.37	16.38	0.27	72.35	14.24	88.73	14.51	128.19		
YER160C	1.01	1.81	2.09	2.34	2.39	18.88	0.25	70.42	13.61	85.64	13.91	99.69		
YDR062C	1.00	1.35	1.56	1.84	1.99	16.99	0.43	93.94	14.65	110.92	15.09	98.66		
YIL157C	0.94	1.69	1.97	2.28	2.35	11.85	0.56	66.29	14.53	78.13	15.09	103.75		
YER184C	1.01	1.54	1.76	2.14	2.19	15.45	1.87	64.74	13.25	80.88	14.44	126.41		
YGL262C	0.94	1.56	1.82	2.13	2.19	16.30	1.03	73.84	14.11	90.14	15.15	171.46		
YGR263C	0.94	1.64	1.92	2.17	2.25	21.17	1.25	128.48	13.90	105.85	15.15	177.46		
YHR169W	0.93	1.81	2.02	2.21	2.27	16.32	0.09	65.67	15.07	82.00	15.15	147.60		
YPR147C	1.03	1.68	2.10	2.32	2.40	16.84	1.36	66.43	13.69	83.27	15.05	52.92		
YHR160C	0.95	1.69	1.90	2.19	2.30	17.32	0.40	60.55	14.67	77.87	15.08	246.51		
YER155C	0.98	1.59	1.84	2.22	2.29	17.30	1.04	94.43	12.44	111.82	15.17	209.60		
YDR254W	0.88	1.30	1.59	1.98	2.16	17.83	0.09	78.09	15.12	95.92	15.21	130.75		
YPR023C	0.88	1.20	1.49	1.80	2.08	25.47	1.62	102.50	13.60	127.97	15.22	170.10		
YGL222C	0.96	1.61	1.85	2.15	2.22	22.87	1.32	90.51	13.92	113.37	15.24	144.15		
YER186W	0.93	1.78	2.02	2.12	2.28	20.3								

ORF	ODs (h)						GLUCOSE		FRUCTOSE		TOTAL sugars (g/L)		TOTAL AMINO ACIDS (mg/L)	
	0	20	27	42	67	43 h	67 h	43 h	67 h	43 h	67 h	48 h		
YIL038C	0.99	1.75	1.89	2.12	2.19	15.58	0.52	84.28	15.44	99.86	15.96	179.30		
YER183C	0.98	1.72	2.01	2.27	2.32	17.94	0.47	73.14	15.52	91.08	15.98	96.81		
YGL127C	0.98	1.61	1.85	2.14	2.22	20.70	0.87	94.17	15.12	114.87	15.99	146.60		
YKL003C	0.95	1.55	1.84	2.13	2.28	14.37	1.00	62.16	15.00	76.53	15.99	256.67		
YER129W	1.03	1.70	1.96	2.29	2.32	17.90	2.28	58.64	13.71	76.55	15.99	222.22		
YGR01W	1.02	1.44	1.72	2.17	2.26	20.61	0.19	86.59	15.80	107.61	16.00	110.25		
YGR24C	0.94	1.51	1.77	2.10	2.18	16.47	0.46	76.28	15.53	94.75	16.01	199.91		
YGR51W	0.96	1.83	1.86	2.13	2.20	12.88	0.81	65.21	15.41	70.21	16.02	199.49		
YLR019C	0.98	1.49	1.74	2.19	2.29	23.65	0.56	105.25	14.59	121.21	16.03	168.28		
YBL112W	1.09	1.76	2.07	2.33	2.42	15.84	0.39	73.46	15.75	91.30	16.04	158.84		
YOL129W	0.97	1.55	1.84	2.07	2.36	18.56	0.58	78.92	15.50	91.29	16.06	227.34		
YPL344C	0.98	1.79	1.87	1.93	2.11	18.53	1.50	91.08	14.58	110.57	16.08	169.00		
YGL117W	0.96	1.52	1.86	2.27	2.31	22.09	1.59	75.24	14.51	97.32	16.10	100.98		
YLR164W	0.97	1.70	2.02	2.26	2.36	27.08	1.30	119.44	14.81	145.52	16.11	174.62		
YGR096W	0.99	1.61	1.87	2.17	2.23	17.34	2.01	67.34	14.11	84.68	16.12	160.15		
YGR032C	0.91	1.45	1.75	2.10	2.20	30.99	0.70	100.88	15.46	131.87	16.16	139.35		
YGL263C	0.92	1.32	1.55	1.88	1.99	24.33	0.94	99.35	15.23	123.68	16.17	129.92		
YPR089W	1.10	1.72	2.02	2.31	2.41	37.42	0.79	120.03	15.39	157.45	16.18	-3.73		
YL007C	0.93	1.61	1.85	2.16	2.21	11.56	0.11	62.77	16.13	74.33	16.24	154.70		
YLR144C	0.98	1.64	1.92	2.16	2.26	23.63	0.41	104.33	15.84	127.96	16.25	229.82		
YGR144W	0.95	1.50	1.82	2.21	2.27	36.13	0.83	132.91	15.43	169.04	16.26	165.44		
YLR187W	1.01	1.70	1.91	2.18	2.28	17.47	0.42	76.13	15.84	93.60	16.26	235.49		
YGR061C	0.98	1.65	1.87	2.16	2.24	19.52	1.20	79.15	15.12	98.67	16.32	124.33		
YFR007W	0.94	1.48	1.70	2.02	2.11	17.77	1.28	65.35	15.05	83.12	16.33	111.25		
YPL283C	0.91	1.33	1.82	2.13	2.18	17.75	1.49	83.66	14.84	101.41	16.33	139.68		
YBR233W	0.96	1.41	1.73	2.17	2.27	26.73	0.48	92.06	15.88	118.79	16.35	-3.35		
YKL038W	1.01	1.46	1.71	2.06	2.09	12.57	0.37	67.67	16.00	80.23	16.37	200.04		
YFL049W	0.97	1.59	1.84	2.12	2.22	18.14	0.92	64.09	15.48	82.23	16.40	191.28		
YGR273C	1.02	1.58	1.88	2.24	2.29	27.98	0.20	125.33	16.25	153.31	16.45	101.48		
YER119C	1.08	1.81	2.09	2.34	2.39	23.07	0.80	101.81	15.66	124.89	16.46	90.65		
YMR169W-A	0.99	1.62	1.74	2.05	2.15	21.99	1.50	93.04	14.98	115.03	16.48	157.71		
YLR265C	1.04	1.93	1.84	2.13	2.23	18.42	1.47	80.40	15.01	98.82	16.48	121.68		
YLL053C	0.98	1.68	1.95	2.17	2.29	22.94	0.70	98.39	15.82	121.33	16.52	164.22		
YPR059W	0.98	1.41	1.72	2.17	2.28	37.90	0.71	127.67	15.82	165.57	16.54	-3.97		
YKR031W	1.00	1.73	2.00	2.23	2.31	18.39	0.70	73.82	15.86	92.21	16.56	175.48		
YHR132C	0.95	1.34	1.81	1.84	2.00	19.91	0.19	101.90	16.38	121.81	16.57	176.25		
YOR859W	0.93	1.66	1.96	2.25	2.31	21.50	1.63	93.68	14.96	115.19	16.58	149.32		
YL030C	0.90	1.45	1.72	2.06	2.18	15.84	0.30	69.04	16.29	84.88	16.58	116.64		
YNL157W	0.91	1.59	1.83	2.22	2.22	18.01	1.67	76.23	14.93	94.24	16.60	124.72		

Appendix 6. Shortlisted 257 deletants from the WYDL_LN based on sugars at the (146 h) time point.

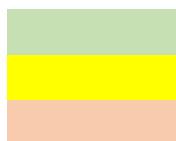
ORF	Optical density at 600 nm at time intervals (h)												GLUCOSE		FRUCTOSE		TOTAL sugars			
	0	20	27	44	67	93	113	126 h	146 h	126 h	146 h	120 h	156 h							
YMR145W	0.87	1.15	1.05	1.29	1.42	1.65	1.64	0.51	0.95	0.78	0.33	0.33	0.37	69.55	-0.27	82.99	1.37			
YDR369W	0.90	1.30	1.48	1.54	1.70	1.90	1.88	3.32	0.52	26.28	-0.63	29.60	-0.11							
YKL037W	0.86	1.12	1.24	1.31	1.45	1.78	1.69	0.38	0.76	8.47	0.00	9.85	0.77							
YDR358W	0.95	1.32	1.43	1.55	1.68	1.87	1.85	9.13	1.04	43.21	-0.17	52.33	0.88							
YER118C	0.95	1.47	1.60	1.67	1.81	1.99	1.98	12.55	1.58	40.32	-0.42	52.87	1.16							
YER105C	0.90	1.35	1.50	1.59	1.76	1.93	1.93	11.61	1.64	82.99	-0.42	94.60	1.22							
YDR496C	0.87	1.19	1.32	1.37	1.51	1.70	1.66	9.61	1.52	31.97	-0.25	41.58	1.27							
YER156C	0.90	1.41	1.59	1.74	1.87	2.04	2.04	10.83	1.67	77.80	-0.38	88.63	1.29							
YDR379W	0.90	1.31	1.47	1.54	1.69	1.89	1.88	11.91	1.26	33.59	0.07	45.50	1.32							
YER129W	0.91	1.35	1.49	1.57	1.72	1.89	1.85	13.44	1.65	69.55	-0.27	82.99	1.37							
YDR359C	0.94	1.17	1.32	1.43	1.56	1.73	1.70	12.28	1.57	56.78	-0.08	69.06	1.49							
YDR339W-A	0.91	1.35	1.58	1.58	1.73	1.91	1.89	14.71	2.09	63.21	-0.21	77.92	1.88							
YER184C	0.98	1.55	1.69	1.77	1.91	2.07	2.05	17.72	2.34	91.27	-0.29	108.99	2.05							
YER167W	1.00	1.47	1.63	1.76	1.88	2.05	2.04	17.00	2.30	93.60	-0.23	110.60	2.08							
YDR383C	0.90	1.34	1.50	1.61	1.75	1.95	1.92	15.63	2.47	30.74	-0.14	46.37	2.32							
YEL027W	0.92	1.34	1.47	1.56	1.70	1.87	1.84	16.69	2.54	39.38	0.13	56.08	2.67							
YDR370C	0.90	1.39	1.58	1.66	1.80	1.98	1.97	16.70	2.35	43.32	0.53	60.02	2.88							
YFL011W	0.95	1.41	1.57	1.69	1.82	1.99	1.99	15.31	3.44	83.32	-0.23	98.63	3.21							
YFL019C	0.93	1.38	1.53	1.63	1.76	1.94	1.91	16.83	3.74	85.00	-0.21	101.83	3.53							
YGR266W	0.87	1.26	1.42	1.49	1.58	1.88	1.82	0.99	0.14	21.95	3.61	22.94	3.74							
YNL176C	0.80	1.16	1.22	1.29	1.39	1.71	1.64	1.60	0.36	16.45	3.68	18.05	4.04							
YIL156W	0.91	1.25	1.37	1.44	1.54	1.90	1.79	0.73	0.41	16.00	3.77	16.73	4.18							
YER083C	0.92	1.21	1.38	1.56	1.68	1.86	1.84	21.10	4.18	19.67	0.11	40.77	4.29							
YAL044W-A	0.90	1.48	1.60	1.64	1.85	2.02	2.00	2.51	0.49	30.77	4.24	33.27	4.73							
YER072W	0.96	1.49	1.65	1.75	1.90	2.06	2.05	16.69	4.12	24.43	0.66	41.12	4.76							
YER173W	0.95	1.38	1.51	1.52	1.66	1.92	1.83	2.04	0.43	29.01	5.17	31.04	5.57							
YGL070C	0.87	1.21	1.41	1.45	1.59	1.89	1.81	1.16	0.83	23.12	5.27	24.28	5.60							
YMR171C	0.92	1.50	1.64	1.75	1.82	2.03	2.08	1.75	2.09	24.62	5.51	27.19	6.11							
YDR384W	0.95	1.46	1.58	1.69	1.82	1.98	1.97	19.46	5.99	29.03	0.49	48.23	6.40							
YLP084W	0.94	1.39	1.56	1.57	1.66	1.92	1.89	0.33	-0.07	17.70	2.88	26.03	6.40							
YML318C	0.98	1.56	1.71	1.73	1.84	2.09	2.03	3.44	0.26	46.64	6.53	50.98	6.79							
YLR027C	0.90	1.36	1.54	1.58	1.74	1.98	1.92	1.83	0.50	34.90	6.32	36.73	6.82							
YLP086C-A	1.14	1.54	1.81	1.98	2.12	2.30	2.36	2.06	-0.04	29.98	7.46	32.04	7.42							
YGR040W	0.93	1.41	1.55	1.57	1.70	2.00	1.93	2.58	0.52	32.62	7.46	35.19	7.98							
YMR029C	0.94	1.58	1.75	1.79	1.87	2.13	2.06	2.90	0.12	45.00	7.88	47.90	8.01							
YLL039C	0.87	1.15	1.30	1.28	1.33	1.65	1.47	1.00	0.31	36.67	7.76	37.57	8.07							
YBR269W	0.86	1.33	1.62	1.57	1.69	2.00	2.02	1.46	0.12	28.01	7.99	29.47	8.11							
YLR413W	0.91	1.33	1.51	1.60	1.77	2.01	1.96	1.37	0.61	29.59	8.73	30.96	9.34							
YIL055C	0.90	1.40	1.52	1.57	1.67	1.95	1.87	3.49	0.68	36.77	8.83	40.26	9.51							
YJU044C	0.95	1.47	1.60	1.64	1.74	2.00	1.96	3.45	0.59	34.10	9.17	37.55	9.75							
YJU020C	0.89	1.36	1.53	1.57	1.67	2.09	2.08	2.78	0.25	42.95	10.14	45.73	10.39							
YPR061C	0.96	1.58	1.62	1.78	1.87	2.11	2.08	6.43	-0.01	43.43	10.77	49.86	10.76							
YEL012W	0.93	1.38	1.51	1.58	1.72	1.90	1.89	10.03	1.07	38.23	10.05	48.27	11.13							
YLR055C	1.01	1.51	1.66	1.76	1.88	2.11	2.12	3.36	0.39	31.74	10.83	35.10	11.22							
YPL203W	0.89	1.39	1.40	1.54	1.68	1.78	1.82	2.68	0.49	31.08	10.80	33.76	11.28							
YGR266W	1.07	1.66	1.80	1.85	1.96	2.19	2.18	2.93	0.63	35.44	11.00	38.38	11.63							
YDR320C	0.91	1.37	1.49	1.64	1.74	2.00	2.01	2.78	0.37	40.32	11.61	43.10	11.98							
YKL213C	0.93	1.43	1.55	1.59	1.71	2.00	1.92	4.97	0.39	39.48	11.71	44.45	12.10							
YIL035C	0.94	1.46	1.63	1.65	1.75	1.91	1.96	1.83	0.28	38.93	12.01	40.76	12.29							
YNL084C	0.96	1.16	1.33	1.33	1.36	1.61	1.51	5.60	0.50	46.83	11.97	52.43	12.47							
YLN219C	0.88	1.28	1.42	1.52	1.67	1.89	1.85	6.91	0.59	40.89	12.00	47.80	12.58							
YGR112W	0.94	1.33	1.46	1.48	1.65	1.92	1.83	3.70	0.59	35.66	12.22	39.36	12.82							
YLR417W	0.87	1.16	1.28	1.29	1.39	1.60	1.55	2.85	0.59	53.23	12.27	56.08	12.85							
YGL019W	0.88	1.27	1.43	1.47	1.59	1.83	1.79	3.70	0.42	49.33	12.60	49.03	13.03							
YLN127W	0.86	1.11	1.27	1.28	1.41	1.70	1.67	7.11	0.19	48.79	12.94	55.90	13.12							
YGR080W	1.20	1.78	1.91	1.96	2.06	2.28	2.27	4.39	0.55	44.45	12.65	48.85	13.19							
YMR254C	0.94	1.34	1.47	1.53	1.63	1.91	1.86	3.61	0.59	46.87	13.26	53.65	13.31							
YPR056W	0.93	1.34	1.44	1.51	1.61	1.89	1.86	3.88	0.51	47.84	13.48	54.48	13.63							
YKL028W	1.02	1.56	1.73	1.76	1.89	2.11	2.06	3.81	0.44	39.18	12.92	43.06	13.36							
YGR182C	0.99	1.36	1.46	1.53	1.61	1.91	1.85	4.21	0.52	44.24	13.19	47.47	13.72							
YIL015W	0.88	1.16	1.25	1.37	1.46	1.70	1.67	4.10	0.72	42.09	13.00	46.20	13.72							
YIL085C	0.90	1.19	1.34	1.49	1.59	1.91	1.79	3.98	0.21	46.72	13.55	50.30	13.76							
YML085W	0.89	1.21	1.32	1.34	1.42	1.74	1.64	2.78	0.88	24.54	13.00	27.32	13.87							
YER145C	0.95	1.30	1.34	1.46	1.59	1.87	1.82	2.04	0.57	49.04	12.49	56.94	14.88							
YER070W	0.94	1.57	1.75	1.78	1.86	2.14	2.08	5.30	0.77	41										

ORF	Optical density at 600 nm at time intervals (h)										GLUCOSE		FRUCTOSE		TOTAL sugars	
	0	20	27	44	67	93	113	120 h	146 h	120 h	146 h	120 h	146 h	120 h	146 h	
YPR138C	0.86	1.32	1.42	1.47	1.57	1.66	1.74	5.82	0.45	29.74	19.20	35.56	19.66			
YER099C	0.88	1.28	1.44	1.52	1.65	1.84	1.81	16.92	2.17	36.69	17.64	53.61	19.81			
YBL093C	0.89	1.23	1.32	1.31	1.48	1.68	1.82	6.18	0.84	27.69	19.03	43.81	19.87			
YGR118C	0.91	1.22	1.41	1.41	1.53	1.93	1.71	7.07	1.00	65.34	18.88	72.41	19.88			
YLR039W	0.94	1.31	1.47	1.55	1.83	1.93	1.89	7.01	0.53	47.81	19.35	54.82	19.88			
YGR077W	1.00	1.52	1.66	1.68	1.78	2.06	2.01	5.61	1.03	45.64	18.97	51.26	20.00			
YML141W	0.89	1.30	1.43	1.53	1.61	1.87	1.86	8.29	0.79	52.76	19.29	61.05	20.08			
YLR269W	0.93	1.38	1.48	1.54	1.64	1.92	1.78	11.79	0.59	46.12	19.53	57.81	20.12			
YHL029C	0.88	1.17	1.29	1.36	1.46	1.70	1.67	6.02	0.92	44.61	19.22	50.63	20.14			
YBL098C	0.96	1.43	1.57	1.72	1.81	2.04	2.02	6.51	1.02	40.81	19.18	47.32	20.20			
YGR118C	0.99	1.49	1.61	1.63	1.76	2.04	1.98	6.11	1.03	46.17	19.24	52.28	20.27			
YHR032W	0.89	1.26	1.39	1.56	1.66	1.99	1.93	7.93	0.42	59.73	19.85	67.65	20.27			
YMR232W	0.90	1.34	1.52	1.62	1.72	1.99	1.97	5.82	0.50	41.35	19.88	47.17	20.38			
YDR385W	0.88	1.16	1.34	1.45	1.61	1.82	1.80	15.60	2.87	29.73	17.51	45.33	20.39			
YGR077W	0.96	1.42	1.56	1.59	1.70	1.98	1.93	5.23	1.14	41.00	19.32	46.23	20.46			
YNL024C	0.87	1.21	1.30	1.27	1.34	1.68	1.57	8.44	1.58	46.34	18.97	54.78	20.55			
YDR438W	0.95	1.49	1.67	1.76	1.87	2.11	2.08	7.52	0.97	54.28	19.60	61.80	20.57			
YGL027C	0.91	1.32	1.48	1.54	1.65	1.85	1.80	9.29	0.90	69.84	19.75	79.13	20.65			
YGR068C	1.00	1.49	1.66	1.73	1.82	2.10	2.05	7.30	1.05	57.58	19.66	64.88	20.71			
YNL134C	0.94	1.35	1.54	1.66	1.76	1.99	1.99	12.46	0.75	60.42	19.97	72.88	20.71			
YGR004W	1.01	1.62	1.75	1.76	1.86	2.13	2.08	6.85	0.74	53.11	20.04	59.96	20.78			
YBR204C	0.89	1.31	1.42	1.51	1.61	1.84	1.83	7.55	0.89	53.22	19.91	60.76	20.80			
YGR232W	1.01	1.65	1.80	1.85	1.96	2.17	2.19	7.44	1.24	47.40	19.64	54.84	20.88			
YOL098C	0.94	1.52	1.70	1.72	1.80	2.07	2.01	6.94	0.67	56.45	20.23	63.38	20.90			
YLR452C	0.88	1.34	1.52	1.67	1.82	2.05	2.03	5.91	0.92	54.42	20.02	60.33	20.94			
YER193C	0.93	1.48	1.65	1.71	1.83	2.11	2.03	8.30	0.84	60.01	20.12	68.31	20.96			
YKL077W	0.94	1.48	1.65	1.74	1.87	2.09	2.06	5.30	1.76	41.39	19.30	46.69	21.06			
YJR103W	0.93	1.28	1.37	1.37	1.46	1.76	1.64	7.94	1.76	51.79	19.30	59.73	21.08			
YDL227C	0.91	1.37	1.53	1.61	1.74	2.00	2.03	8.49	0.68	56.29	20.43	64.78	21.11			
YMR159W	0.84	1.18	1.27	1.29	1.32	1.43	1.65	6.88	1.20	39.79	19.97	46.59	21.17			
YNL120W	0.95	1.48	1.65	1.72	1.82	2.05	2.03	6.76	0.79	50.04	20.38	56.26	21.17			
YAR160C	0.91	1.45	1.63	1.69	1.78	1.93	1.87	5.41	1.73	44.67	19.49	53.41	21.22			
YGR163W	1.06	1.58	1.72	1.85	2.08	2.26	2.24	6.57	0.93	46.74	20.43	52.54	21.34			
YHR165W	0.85	1.22	1.36	1.38	1.49	1.79	1.70	5.55	1.38	42.91	19.38	48.46	21.36			
YIL076W	0.92	1.30	1.45	1.51	1.62	1.88	1.84	4.55	1.07	57.59	20.29	62.14	21.37			
YGR015C	1.01	1.65	1.82	1.88	1.98	2.25	2.21	6.28	1.07	51.18	20.34	57.38	21.41			
YGR237C	0.92	1.33	1.47	1.50	1.56	1.90	1.78	7.92	1.06	52.96	20.42	60.88	21.48			
YGR096W	1.00	1.49	1.62	1.65	1.77	2.01	1.96	6.06	0.99	42.37	20.55	48.42	21.53			
YPR070W	0.91	1.26	1.28	1.43	1.51	1.60	1.65	4.69	0.61	46.58	20.96	51.27	21.57			
YEL017C-A	0.94	1.43	1.58	1.66	1.80	1.98	1.96	16.09	2.49	38.15	19.11	54.25	21.60			
YML311C	0.92	1.37	1.56	1.59	1.72	1.95	1.90	8.37	0.89	68.63	20.74	77.00	21.63			
YDR375C	0.94	1.38	1.58	1.64	1.75	2.03	1.98	6.47	0.73	56.00	20.90	62.47	21.63			
YGL139W	0.85	1.28	1.43	1.49	1.65	1.95	1.88	3.06	1.19	35.74	20.48	38.79	21.67			
YNL146W	0.84	1.20	1.36	1.41	1.49	1.79	1.73	5.38	1.85	39.69	19.82	45.07	21.67			
YLR150W	0.91	1.39	1.54	1.63	1.74	1.86	2.05	1.97	6.96	1.06	42.97	20.61	49.93	21.69		
YBL028C	0.87	1.31	1.63	1.64	1.66	1.83	1.80	7.44	1.53	48.12	20.67	55.56	22.20			
YDR378C	0.97	1.57	1.76	1.83	1.93	2.17	2.15	7.02	0.84	57.44	21.39	64.46	22.23			
YIL007C	0.90	1.26	1.38	1.44	1.53	1.75	1.73	5.85	1.51	39.99	20.78	45.84	22.29			
YNL051W	0.93	1.30	1.42	1.43	1.50	1.81	1.71	6.26	1.17	52.55	21.16	58.80	22.33			
YJR148W	0.91	1.31	1.39	1.44	1.54	1.85	1.76	7.63	1.28	53.95	20.04	62.78	22.33			
YJU097W	0.95	1.38	1.56	1.66	1.79	2.00	1.98	7.74	1.81	49.66	20.17	57.40	21.98			
YGL211W	0.91	1.35	1.50	1.56	1.70	1.99	1.93	4.85	1.26	42.19	20.84	47.04	22.10			
YER150W	0.95	1.48	1.63	1.74	1.86	2.05	2.03	16.74	2.11	92.25	20.04	108.99	22.15			
YBL028C	0.87	1.31	1.43	1.49	1.61	1.83	1.80	7.44	1.53	48.12	20.67	55.56	22.20			
YDR202W	0.88	1.12	1.21	1.25	1.30	1.55	1.49	7.47	0.81	47.48	21.78	54.95	22.59			
YPL198W	0.96	1.46	1.58	1.64	1.74	2.03	1.90	10.00	0.96	44.97	21.64	55.86	22.60			
YGR066C	1.03	1.75	1.88	1.89	2.19	2.18	2.15	6.68	1.34	54.20	21.36	60.89	22.60			
YFR032C-A	0.94	1.34	1.50	1.58	1.70	2.00	1.93	7.51	1.03	51.28	21.58	58.79	22.61			
YNL068C	0.88	1.33	1.52	1.59	1.69	2.02	1.94	6.28	1.37	37.55	21.37	43.83	22.74			
YMD42W-B	0.88	1.29	1.48	1.57	1.67	1.97	1.93	5.96	1.05	41.49	21.69	47.44	22.75			
YME202W	0.85	1.21	1.40	1.46	1.55	1.84	1.80	5.92	1.21	40.16	21.61	46.08	22.81			
YIL021W	0.86	1.25	1.40	1.48	1.57	1.87	1.82	5.08	1.05	39.48	21.45	58.91	22.51			
YER148C	0.88	1.30	1.45	1.51	1.61	1.91	1.84	7.47	1.41	47.10	21.43	49.81	22.84			
YGR059W-A	0.90	1.35	1.59	1.69	1.80	2.05	2.02	6.63	0.13	47.71	22.72	54.34	22.85			
YAL049C	0.91	1.36	1.51	1.57	1.64	1.95	1.91	8.82	1.26	53.27	21.61	62.09	22.87			
YDL194W	0.92	1.34	1.55	1.66	1.76	2.02	2.01	7.98	0.70	59.19	22.37	67.17	23.07			
YGR169C-A	0.94	1.31	1.47	1.56	1.65	1.89	1.86	6.63	1.13	43.07	21.96	49.70	23.09			
YOR054C	0.87	1.36	1.56	1.65	1.77	2.03	1.97	8.33	1.01	65.53	22.11	73.86	23.11			
YDR329W	0.96	1.40	1.53	1.61	1.75	1.93	1.91	15.54	2.24	44.26	20.88	59.79	23.12			
YKL207W	0.91	1.39	1.54	1.58	1.67	1.96	1.90	8.19	1.15	49.52	22.02	57.71	23.17			
YIL008W	0.86	1.18	1.31	1.40	1.51	1.76	1.72	4.55	1.50	39.58	21.75	44.13	23.24			
YHR193C	0.89	1.23	1.35	1.35	1.46	1.77	1.64	8.13	1.75	50.46	21.54	58.59	23.28			

Appendix 7. Data for histogram representing the distribution of deletants (WYDL_LN, Fig. 2).

Range	deletants
0-10	40
10-20	92
20-30	390
30-40	558
40-50	350
50-60	132
60-70	57
70-80	29
80-90	14
90-100	11
100-110	7
110-120	2
120-130	6
130-140	3
140-150	0
150-160	0
160-170	4
170-180	3
180-190	3
above 190	143

Residual sugar of wildtype $48 \pm 8 \text{ g L}^{-1}$



1080 candidates (Residual sugar $< 40 \text{ g L}^{-1}$ at time point C; 146 h)

482 candidates (Residual sugar $40-60 \text{ g L}^{-1}$ at time point C; 146 h)

282 candidates (Residual sugar $> 60 \text{ g L}^{-1}$ at time point C; 146 h)

Appendix 8. Shortlisted 92 deletants based on residual sugars (<48 g L⁻¹ at 120 h and < 26 g L⁻¹ at 144 h) of the micro-fermentation.

ORF	Gene name	Gene description	TOTAL sugars	
			120 hrs	146 hrs
YNL145W	MFA2	Mating Factor A	7.79	-0.27
YDR368W	YPR1	Yeast Putative Reductase	29.60	-0.11
YKL037W	AM26	Altered Inheritance rate of Mitochondria	9.85	0.77
YDR466C	VPS60	Vacuolar Protein Sorting	41.58	1.27
YDR379W	RGA2	Rho GTPase Activating Protein	45.50	1.32
YDR383C	NKP1	Non-essential Kinetochore Protein	46.37	2.32
YGR296W	YRF1-3		22.94	3.74
YNL176C	TDAT	Topoisomerase I Damage Affected	18.05	4.04
YIL156W	UBP7	Ubiquitin-specific Protease	16.73	4.18
YER083C	GET2	Guided Entry of Tail-anchored proteins	40.77	4.29
YAL044W-A	BOL1	Bola-like protein	33.27	4.73
YER072W	VTC1	Vacuolar Transporter Chaperone	41.12	4.78
YOR113W	AZF1	Asparagine-rich Zinc-Finger	31.04	5.59
YGL087C	MMS2	Methyl Methanesulfonate sensitivity	24.28	6.10
YMR171C	EARI	Endosomal Adaptor of Rsp5p	27.19	6.11
YPL084W	BRO1	BCK1-like Resistance to Osmotic shock	28.03	6.40
YLR027C	AT2	Aspartate AminoTransferase	36.73	6.82
YPL096C-A	ERI1	ER-associated Ras Inhibitor	32.04	7.42
YGR040W	KSS1	Kinase Suppressor of Sst2 mutations	35.19	7.98
YMR029C	FAR8	Factor Arrest	47.90	8.01
YLL039C	UBI4	Ubiquitin	37.67	8.07
YBR299W	MAL32	MALtox	29.47	8.11
YLR413W	INA1	Indicator of Abscission	30.96	9.34
YIL055C			40.26	9.51
YJL044C	WPS55	Vacuolar Protein Sorting	37.55	9.75
YJL020C	BBC1	Bni1 synthetic lethal and Bee1 (fas17) Complex member	45.73	10.39
YLR095C	IOC2	Isw1 One Complex	35.10	11.22
YPL203W	TPK2	Takashi's Protein Kinase	33.76	11.28
YGR206W	MVB12	MultiVesicular Body sorting factor of 12 kilodaltons	38.38	11.63
YDR320C	SWA2	Synthetic Lethal With Arf1	43.10	11.98
YKL213C	DOA1	Degradation Of Alpha	44.45	12.10
YIL035C	CKA1	Casein Kinase Alpha subunit	40.76	12.29
YNL219C	ALG9	Asparagine-Linked Glycosylation	47.80	12.58
YGR112W	SHY1	SURF-Homolog of Yeast	39.36	12.82
YMR254C			44.21	13.22
YMR030W	RSF1	ReSpiration Factor	45.79	13.33
YKL082W	MSN4	Multicopy suppressor of SNF1 mutation	43.06	13.38
YGR132C	PHB1	Prohibitin	47.47	13.72
YIL015W	BAR1	BARrier to the alpha factor response	46.20	13.72
YNL085W	MKT1	Maintenance of K2 Killer Toxin	27.32	13.87
YPL027W	SMA1	Spore Membrane Assembly	35.75	14.39
YKR066C	CCP1	Cytochrome c Peroxidase	43.81	14.66
YKL003C	MRP17	Mitochondrial Ribosomal Protein	39.48	14.98
YKR088C	TVF38	Ttg2-Vesicle Protein	34.42	15.52
YPR071W			47.19	15.56
YGR070W	ROM1	Rho1 Multicopy suppressor	46.92	15.73
YBL024W	NCL1	NuClear protein	47.30	16.17
YDR388W	RV5167	Reduced Viability on Starvation	34.04	16.19
YMR285C			43.52	16.21
YGR105W	VMA21	Vacuolar Membrane Atpase	44.23	16.33
YMR235W	GFD1	Good For Dbs5p	40.34	16.83
YGL208W	SIP2	SNF1-Interacting Protein	42.81	17.18
YIL053W	PEP8	carboxyPEPtase Y-deficient	42.73	17.39
YJR102C	WPS25	Vacuolar Protein Sorting	47.72	17.50
YPL119C	DBP1	Dead Box Protein	41.12	17.66
YGL064C	MRH4	Mitochondrial RNA Helicase	47.37	17.94
YKL120W	OAC1	OxaloAcetate Carrier	42.88	17.96
YIL010W	DOT5	Disruptor Of Telomeric silencing	44.53	18.29
YIL017C	VID28	Vacuolar Import and Degradation	44.28	18.43
YKL070W			42.37	18.59
YNL054W	VAC7	VACUolar segregation	36.37	19.28
YPL138C	SPP1	Set1c, Phd finger Protein	35.56	19.66
YBL003C	HTA2	Histone h Two A	43.87	19.87
YPL098C	MGR2	Mitochondrial Genome Required	47.32	20.20
YMR232W	FUS2	cell FUSSION	47.17	20.38
YDR389W	EFT2	Elongation Factor Two	45.33	20.39
YGR017W			46.23	20.46
YKL071W			46.69	21.06
YMR324W	UBP15	Ubiquitin-specific Protease	46.59	21.17
YIL076W	SEC28	SECretary	42.14	21.37
YGL139W	FLC3	Flavin Carrier	38.79	21.67
YNL146W			45.07	21.67
YPL046C	ELC1	Elongin C	47.03	21.88
YGL211W	NCS6	Needs Cdc4 to Survive	47.04	22.10
YIL007C	NAS2	Non-ATPase Subunit	45.84	22.29
YIL044C	AGE2	ArfGAP Effector	47.10	22.41
YNL068C	FKH2	Fork head Homolog	43.83	22.74
YNL042W-B			47.44	22.75
YNL202W	SPS19	SPorulation-Specific	46.08	22.81
YIL008W	URM1	Ubiquitin Related Modifier	44.13	23.24
YOR334W	MRS2	Mitochondrial RNA Splicing	45.13	23.29
YBR264C	YPT10	Yeast Protein Two	46.44	23.86
YKL127W	PGM1	PhosphoGlucoMutase	47.05	24.24
YHL014C	YLF2		47.10	24.44
YAL037C	GIP4	Glc7 Interacting Protein	47.83	24.44
YKL017C	HCS1	dnab Helicase	47.74	24.57
YNL056W	OCA2	Oxidant-induced Cell cycle Arrest	43.71	25.06
YHR157W	REC104	RECombination	47.70	25.17
YKL124W	SSH4	Suppressor of Sh3 deletion	41.06	25.17
YMR233W	TRI1		47.47	25.69
YNL050C			47.29	25.69
YBR268W	MRPL37	Mitochondrial Ribosomal Protein, Large subunit	43.59	25.73

ORF – Open reading frame

Appendix 9. Shortlisted 30 candidates based on AUC and fermentation duration of 92 shortlisted strains when compared to the wildtype

Deletants	Deletants progress during fermentation (residual sugar at different sampling time points)										AUC	AUC % of parent
	24 h	49 h	71 h	95 h	107 h	119 h	131 h	151 h	181 h	205 h		
YKL120W	87.99	76.62	55.42	39.23	36.82	28.94	21.82	6.15	7.04	1.29	9516	95
YMR029C	106.67	82.71	56.68	38.21	31.06	19.76	10.06	-0.36	-0.11		9493	95
YAL044W-A	101.77	73.82	55.08	36.03	32.86	25.09	13.96	5.60	1.65		9408	94
YDR368W	105.77	92.96	51.01	31.66	28.16	5.77	5.50	1.36	-0.10		9348	93
YLL039C	106.92	84.86	47.10	32.57	27.33	20.64	10.97	3.06	0.97		9299	93
YDR320C	96.45	80.60	53.63	36.94	32.55	23.99	13.40	3.04	0.11		9298	93
YPL203W	101.75	73.26	51.17	35.44	31.43	23.73	13.50	5.49	1.40		9244	92
YKL062W	97.76	68.82	51.71	37.22	32.95	25.90	14.79	6.37	3.24	-0.01	9224	92
YKR088C	98.75	74.64	51.39	35.24	32.80	23.03	13.68	5.29	1.05		9201	92
YDR486C	99.49	69.99	53.06	33.43	32.34	29.37	14.91	3.79	0.45		9153	92
YGR112W	98.30	72.31	49.92	35.37	30.73	23.98	13.86	5.33	2.37	-0.11	9129	91
YBL024W	100.33	73.07	51.11	33.73	31.29	22.76	12.06	4.69	1.20		9110	91
YMR171C	99.38	72.14	50.36	33.69	30.81	21.73	12.49	4.42	0.23		9006	90
YMR030W	94.80	64.80	53.56	35.95	33.64	25.02	15.38	3.33	0.30		8929	89
YNL085W	101.25	76.11	50.21	32.90	27.77	17.95	8.74	0.69			8888	89
YLR027C	98.44	74.83	51.52	32.46	27.09	20.10	9.84	2.05	-0.11		8882	89
YIL035C	92.45	80.65	49.52	32.82	28.56	20.83	11.02	1.66	-0.14		8868	89
YKR066C	84.44	75.15	51.88	34.09	31.31	24.97	15.58	2.59	0.65		8812	88
YGR040W	88.03	71.67	49.86	36.10	30.76	23.84	14.70	2.41	0.10		8756	88
YLR413W	103.93	77.06	49.02	29.20	23.74	14.94	5.77	-0.75			8743	87
YGR206W	99.18	74.09	50.33	30.74	26.00	17.44	7.27	0.63			8702	87
YPL027W	97.11	71.81	48.30	31.53	26.50	16.82	7.70	0.38			8662	86
YKL003C	97.71	70.87	46.70	29.75	26.47	17.21	8.25	1.23	-0.14		8522	85
YPL084W	100.97	73.42	48.34	27.28	21.36	12.63	4.49	-0.60			8456	85
YGL087C	100.56	71.27	45.32	25.37	21.05	11.52	2.77	-0.76			8248	82
YKL037W	100.97	72.59	45.08	22.30	14.73	6.22	-0.27				8057	81
YJL020C	92.26	69.48	40.87	22.54	17.13	9.05	0.52				7746	77
YNL145W	97.44	74.02	40.43	17.46	8.29	5.52	-0.28				7725	77
YNL176C	91.89	62.39	36.11	16.19	11.11	5.32	-0.26				7228	72
YIL156W	82.79	65.17	35.89	19.15	14.63	7.03	-0.15				7185	72
Parents progress during fermentation (residual sugar at different sampling time points)												
Time	24 h	49 h	71 h	95 h	107 h	119 h	131 h	151 h	181 h	205 h		
1	100.45	88.20	57.18	40.59	38.36	32.17	19.31	11.33	8.78	2.61	10486	
2	88.64	78.36	56.43	40.52	38.27	30.81	20.46	7.75	7.48	1.89	9798	
3	102.04	77.07	56.82	39.62	36.36	30.38	17.68	9.63	5.85	0.66	9993	
4	97.31	74.64	57.17	41.13	35.60	27.58	16.65	8.36	4.73	-0.04	9717	
Avg	97	80	57	40	37	30	19	9	7	1	9999	
SD	6	6	0	1	1	2	2	2	2	1	345	



Fermentation completion of the wildtype and deletant (residual sugars <2 g L⁻¹)

Residual sugar of the deletants

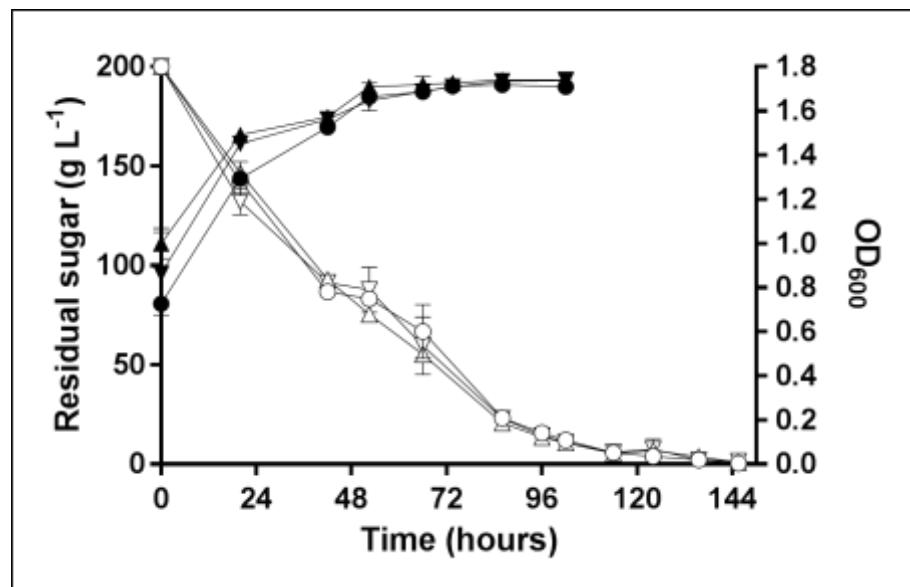
Residual sugar of the wildtype

Area under the curve calculated using the fermentation curve

Appendix 10. Enrichment of fermentation efficient genes annotated to specific GO terms from SGD using GO ToolBox GO-stats.

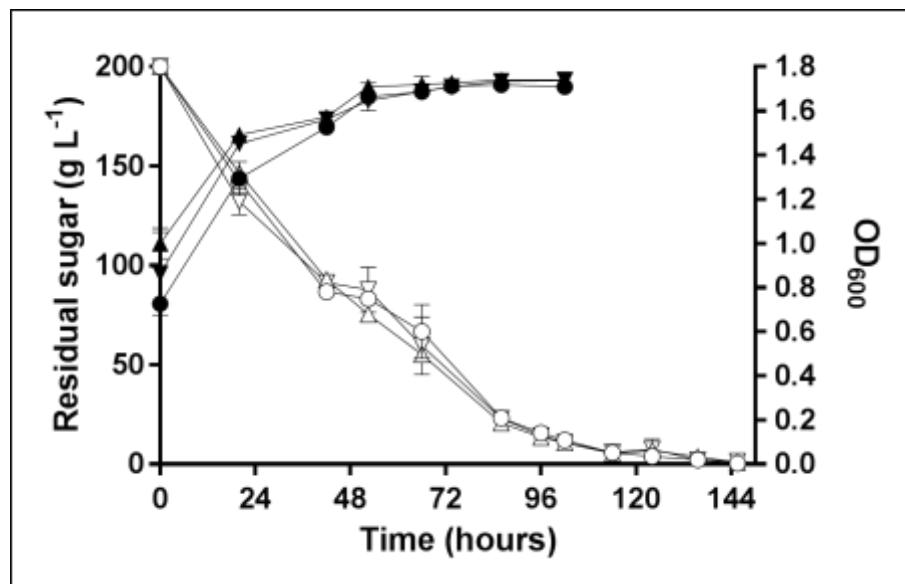
GO_ID	TERM	P-VALUE	GENES_IN_SET
GO:0070647	protein modification by small protein conjugation or removal	3.65E-06	<i>UBP7,UBI4,BRO1,MMS2,UBC13</i>
GO:0032509	endosome transport via multivesicular body sorting pathway	4.45E-06	<i>MVB12,BRO1,EAR1</i>
GO:0000209	protein polyubiquitination	7.43E-06	<i>UBI4,MMS2,UBC13</i>
GO:0016579	protein deubiquitination	8.66E-06	<i>UBP7,UBI4,BRO1</i>
GO:0070646	protein modification by small protein removal	2.11E-05	<i>UBP7,UBI4,BRO1</i>
GO:0043328	protein targeting to vacuole during ubiquitin-dependent protein catabolic process via the MVB pathway	3.85E-05	<i>BRO1,EAR1</i>
GO:0019941	modification-dependent protein catabolic process	6.76E-05	<i>UBP7,BRO1,MMS2,EAR1,UBC13</i>
GO:0043632	modification-dependent macromolecule catabolic process	7.75E-05	<i>UBP7,BRO1,MMS2,EAR1,UBC13</i>
GO:0051603	proteolysis involved in cellular protein catabolic process	8.05E-05	<i>UBP7,BRO1,MMS2,EAR1,UBC13</i>
GO:0043687	post-translational protein modification	1.12E-04	<i>UBP7,TPK2,UBI4,BRO1,MMS2,UBC13</i>
GO:0016197	endosome transport	2.42E-04	<i>MVB12,BRO1,EAR1</i>
GO:0006623	protein targeting to vacuole	2.53E-04	<i>MVB12,BRO1,EAR1</i>
GO:0044257	cellular protein catabolic process	4.51E-04	<i>UBP7,BRO1,MMS2,EAR1,UBC13</i>
GO:0032511	late endosome to vacuole transport via multivesicular body sorting pathway	4.53E-04	<i>BRO1,EAR1</i>
GO:0016567	protein ubiquitination	5.61E-04	<i>UBI4,MMS2,UBC13</i>
GO:0006301	postreplication repair	5.75E-04	<i>MMS2,UBC13</i>
GO:0006513	protein monoubiquitination	6.42E-04	<i>UBI4,UBC13</i>
GO:0043162	ubiquitin-dependent protein catabolic process via the multivesicular body sorting pathway	7.12E-04	<i>BRO1,EAR1</i>
GO:0006508	proteolysis	8.96E-04	<i>UBP7,BRO1,MMS2,EAR1,UBC13</i>
GO:0032446	protein modification by small protein conjugation	1.01E-03	<i>UBI4,MMS2,UBC13</i>
GO:0044267	cellular protein metabolic process	1.03E-03	<i>UBP7,TPK2,UBI4,BRO1,MFA2,MMS2,MKT1,EAR1,UBC13,MRP17</i>
GO:0006464	protein modification process	1.41E-03	<i>UBP7,TPK2,UBI4,BRO1,MMS2,UBC13</i>

GO:0034962	cellular biopolymer catabolic process	1.89E-03	<i>UBP7,BRO1,MMS2,EAR1,UBC13</i>
GO:0007034	vacuolar transport	1.91E-03	<i>MVB12,BRO1,EAR1</i>
GO:0019538	protein metabolic process	2.00E-03	<i>UBP7,TPK2,UBI4,BRO1,MFA2,MMS2,MKT1,EAR1,UBC13,MRP17</i>
GO:0045324	late endosome to vacuole transport	2.30E-03	<i>BRO1,EAR1</i>
GO:0030163	protein catabolic process	2.54E-03	<i>UBP7,BRO1,MMS2,EAR1,UBC13</i>
GO:0044265	cellular macromolecule catabolic process	3.70E-03	<i>UBP7,BRO1,MMS2,EAR1,UBC13</i>
GO:0006531	aspartate metabolic process	4.08E-03	<i>AAT2</i>
GO:0006532	aspartate biosynthetic process	4.08E-03	<i>AAT2</i>
GO:0044248	cellular catabolic process	4.80E-03	<i>AIM26,UBP7,BRO1,MMS2,EAR1,UBC13</i>
GO:0006511	ubiquitin-dependent protein catabolic process	5.12E-03	<i>UBP7,BRO1,EAR1</i>
GO:0043412	biopolymer modification	6.41E-03	<i>UBP7,TPK2,UBI4,BRO1,MMS2,UBC13</i>
GO:0043285	biopolymer catabolic process	6.83E-03	<i>UBP7,BRO1,MMS2,EAR1,UBC13</i>
GO:0044419	interspecies interaction between organisms	8.14E-03	<i>MKT1</i>

Appendix 11. Fermentations of L2056 and homozygous and heterozygous *MFA2* deletants

Fermentations (100 mL) were conducted in triplicate in CDGJM with 75 mg N L⁻¹, sugar consumption was monitored enzymatically and reported as total sugars (open symbols). Growth was monitored as optical density (at 600 nm (solid symbols). Data points present mean values of triplicate data with standard deviation as error bars. Circle; L2056 square; L2056 *MFA2/mfa2Δ*, inverted triangle; L2056 *mfa2Δ/mfa2Δ*.

Appendix 12. Fermentations of *MFA2* deletants in C911A/C911D background during fermentation



Fermentations (100 mL) were conducted in triplicate in CDGJM with 75 mg N L⁻¹, sugar consumption was monitored enzymatically and reported as total sugars (open symbols). Growth was monitored as optical density at 600 nm (solid symbols). Data points present mean values of triplicate data with standard deviation as error. Circle; C911A/C911D, square; C911A *mfa2Δ/C911D*, upright triangle; C911A/C911D *mfa2Δ*, inverted triangle; C911A *mfa2Δ/C911D mfa2Δ*.

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