

Functional Output Regression for Machine Learning in Materials Science

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Abstract

In recent years, there has been a rapid growth in the use of machine learning in materials science. Conventionally, a trained predictive model describes a scalar output variable, such as thermodynamic, electronic, or mechanical properties, as a function of input descriptors that vectorize the compositional or structural features of any given material, such as molecules, chemical compositions, or crystalline systems. In machine learning of materials data, on the other hand, the output variable is often given as a function. For example, when predicting the optical absorption spectrum of a molecule, the output variable is a spectral function defined in the wavelength domain. Alternatively, in predicting the microstructure of a polymer nanocomposite, the output variable is given as an image from an electron microscope, which can be represented

as a two- or three-dimensional function in the image coordinate system. In this study, we considered two unified frameworks to handle such multidimensional or functional output regressions, which are applicable to a wide range of predictive analyses in materials science. The first approach employs generative adversarial networks, which are known to exhibit outstanding performance in various computer vision tasks such as image generation, style transfer, and video generation. We also present another type of statistical modelling inspired by a statistical methodology referred to as functional data analysis. This is an extension of kernel regression to deal with functional outputs, and its simple mathematical structure makes it an effective modelling even for given data in limited supply. We demonstrate the proposed method through several case studies in materials science.

Introduction

Recently, there has been a growing trend in various domains of materials science to use machine-learning techniques to accelerate the process of designing and creating new materials. Conventionally, machine learning models are used to rapidly perform high-throughput virtual screening across millions or billions of candidate materials that span an enormous search space¹⁻⁵. In general, a model describes physicochemical, electronic, thermodynamic, or mechanical properties as a function of the input materials, which are given in various forms, such as small- or macro-molecules, crystalline systems, chemical or raw material compositions, and their mixtures. To put the task into a machine learning framework, such a non-numeric variable needs to be transformed into a fixed-length numeric vector called a *descriptor*, which represents the compositional or structural features of the given material⁵⁻¹⁶. Under the supervision of the given data, a model is trained to learn the mapping from the vectorized features to their respective properties. In this workflow, the feature representation of the input materials plays a key role in boosting the predictive power.

There is a great deal of prior work on transforming material features into numeric vectors

and constructing regression models or classifiers that represent the mapping from vectorized input materials to their output properties. A class of descriptors, referred to as molecular fingerprints, has long been studied in chemical informatics, which converts a chemical structure or molecular graph into an integer-valued vector according to the presence or absence or the number of occurrences of a particular chemical fragment, in which hundreds or thousands of fragments are considered¹⁰⁻¹⁴. Another type of molecular descriptor employs a quantitative representation of the topological or physicochemical features of a molecular system¹⁷⁻²⁰. Chemical composition can be considered as a set variable consisting of a variable number of element species and their contents. There are a large volume of previous studies on the representation of such compositional features⁵⁻⁹. A crystal structure is typically vectorized by encoding the local structural environments of each atom and the neighboring relations of constituent atoms in a unit cell^{7-9,21}. In recent years, there has also been an increasing trend in the treatment of a material structure as a graph and in modeling its properties using graph neural networks²²⁻²⁴. A natural representation of the chemical structure was created on a labeled graph. A periodic configuration of atoms in a crystalline system can also be translated into a graph called a crystal graph, which represents the coordination of constituent atoms in infinitely arranged unit cells²². In addition, when predicting the properties of a composite system from its microstructure, it is natural to treat the microstructure as an image. In the study of composite materials, scanning electron microscopy (SEM) and transmission electron microscopy (TEM) are widely used to observe the surface or interfacial structure of the fabricated materials. By treating the microstructure as an image, supervised learning can be addressed by regressing real-valued output properties onto the space of the microstructure images, as in computer vision and image recognition^{25,26}. Other representation methods have also been investigated for various material systems, such as topological feature representation of disordered material systems using persistent homology²⁷, identification of multi-component materials based on the spectral function of powder X-ray diffraction²⁸, and prediction of reaction outcomes in organic synthesis based on string

representations of product and reactant molecules^{29,30}.

As mentioned above, most previous studies have considered the ordinary problem setting of supervised learning, where an input variable is given as a relatively high-dimensional vector encoding material features, and the output is a scalar or low-dimensional real-valued vector, for example, a few sets of physicochemical properties or a class label indicating structural species or the level of physical features. On the other hand, there are many potential problem settings in materials science, where the output variable is inherently ultra-high-dimensional or multidimensional (e.g., a functional-type output), but the methodology of supervised learning in such scenarios has not been well studied. For example, in the task of predicting the ultraviolet-visible (UV-Vis) absorption spectra of molecules, the input variable can be given by a vectorized molecular structure, and the output variable is given as a function defined on the domain of wavelengths that represents the optical absorbance³¹. In the study of composite materials, it is important to qualitatively and quantitatively understand the influence of processing conditions such as temperature, pressure, and composition on the resulting microstructures. SEM and TEM are commonly used to examine microstructures. If we formulate the problem within a framework of supervised learning, the input is a real-valued vector encoding the processing condition and composition, and the output is given by an intensity matrix representing the grayscale microscopic image. This is a regression problem for multidimensional output variables. Alternatively, the problem can be reduced to an image generation task in computer vision. To solve such problems, various types of deep generative models, such as the conditional GAN (cGAN)³² and encoder-decoder networks³³, can be applied. In fact, there have been several previous studies in which the cGAN was applied to the prediction of microstructures, as described above³⁴⁻³⁶, and an encoder-decoder model for the prediction of the UV-Vis absorption spectra of organic molecules³¹. In addition, in statistical science, regression methods for functional output variables have long been studied in the context of functional data analysis³⁷, which may also be applicable to solving the aforementioned problems.

In this study, we consider two unified frameworks for multidimensional functional output regression that can cover various potential applications in materials science. The first approach employs cGAN, inspired by the study of microstructure images in Banko et al.³⁴. However, because cGAN has training instability in the adversarial learning process and weakness to limited amounts of data, we developed another framework, a statistical modeling that relies on the methodology of functional data analysis for functional output variables. The present model can be viewed as an extension of kernel regression to handle functional output variables, and has a simpler mathematical form than the cGAN model architecture. As shown later, the method exhibits outstanding predictive performance even in cases where only a small amount of data is available. We demonstrate these two methods using three case studies. In the first two case studies, the optical absorption spectra of organic molecules in two different regions of UV-Vis (170-780 nm) and near-infrared wavelength (NIR: 780–2500 nm) were predicted. The output variable was a spectral function in the wavelength domain. The number of training instances is of the order of thousands for the former case, whereas for the latter, the data are in limited supply as the number of training molecules is approximately 60. With these applications, we demonstrated the potential predictive ability of these two methods on a limited amount of training data. In particular, we compare ordinary regression, which predicts a scalar output variable with a pre-quantified spectral feature³⁸, with the present methods predicting the whole function directly, and describe the superiority of the latter and its statistical mechanisms in relation to multitask learning³⁹. The objective of the third example is to predict an electron microscopic image of the microstructure for any given composition and processing conditions in the fabrication of thin-film composite materials. Python codes used in the case studies were distributed⁴⁰.

Preliminary

The present study deals with a supervised learning problem in which the input variable $X \in \mathbb{R}^p$ is a p -dimensional descriptor vector and the output variable $Y(X, t) \in \mathbb{R}$ is given by a real-valued function of X and an additional argument $t \in \mathbb{R}^q$. Argument t corresponds to a coordinate in image space or to a wavelength at which the spectral function is defined. In the following sections, we describe potential applications in materials science.

Spectral prediction

Molecules undergo temporal transitions from their ground states to higher-energy excited electronic states in response to the absorption of light, such as UV-Vis or NIR. The absorption wavelength is proportional to the inverse of energy. The absorbance spectrum, which represents the intensity of optical absorption as a function of wavelength, is determined by the excitation energy levels and the transition probabilities of the electronic states in a molecular system. Accurately predicting molecule-specific absorbance spectra is highly beneficial for various applications, such as the design of organic light-emitting diodes^{1,41}, organic photovoltaic cells⁴², and UV filters⁴³. Usually, absorption peak wavelengths are predicted from the excited states of electrons obtained *ab initio*, for example, by performing time-dependent density functional theory (TD-DFT) calculations⁴⁴. However, owing to the high computational cost, first-principle calculations are not useful for exhaustive molecular screening. Furthermore, while the location and intensity of a peak wavelength can be estimated *ab initio*, other functional features of the full spectrum, such as the full width at half maximum and absorbance integration in a wavelength interval, cannot be determined.

Here, we address the problem of spectral prediction using a fully data-driven approach that does not rely on *ab initio* calculations (Figure 1(a)). The input variable of model f consists of a descriptor vector X and a wavelength $t \in \mathbb{R}_+^1$, where descriptor X encodes the structural and compositional features of the input molecule. The output variable is the

spectral function $Y(X, t)$ of the optical absorbance with respect to varying molecules and wavelengths. In summary, with observational noise ϵ , the model can be expressed as

$$Y(X, t) = f(X, t) + \epsilon. \quad (1)$$

For each of the n observed molecules $\{X_i | i = 1, \dots, n\}$, the absorption spectrum $Y(X, t)$ was measured over m discretized wavelengths $\{t_j | j = 1, \dots, m\}$. In this study, it is assumed that the observation points of the wavelength are common to all molecules: t_j is independent of the index i of the molecule. When the observed wavelengths vary across the molecules, one can obtain a series of complete data with the same observation points by smoothly interpolating the missing data points. The task then comes down to the regression problem for the high-dimensional vector-valued output, which is modelled by

$$\mathbf{y}(X) = \mathbf{f}(X) + \boldsymbol{\epsilon}, \quad (2)$$

where $\mathbf{y}(X)^\top = (Y(X, t_1), \dots, Y(X, t_m))$, $\mathbf{f}(X)^\top = (f(X, t_1), \dots, f(X, t_m))$ and $\boldsymbol{\epsilon}^\top = (\epsilon_1, \dots, \epsilon_m)$. As mentioned above, it is assumed that there are no missing data or that the missing data in the direction t have been interpolated beforehand based on a sufficient amount of observed data. However, the functional output kernel regression, which will be shown later, can naturally perform training with no special treatment for missing data, even if the observation points of t are quite sparse.

Machine learning for predicting the optical absorption spectra of molecular systems has not been extensively studied. To clarify the contribution of our work, we consider the recently published work of Urbina et al.³¹ that relied on Seq2Seq³³ and its variant encoder-decoder architectures with a built-in attention mechanism. Seq2Seq, which is widely used in natural language processing, was utilized to learn mapping from tokenized SMILES strings⁴⁵ or pre-defined molecular descriptors of input chemical structures to the absorption spectra. On the other hand, the present study introduces a much simpler statistical model, the

functional output kernel regression, aimed at stabilizing the learning process by reducing over-parameterization and achieving a high prediction accuracy even in cases where sufficient amounts of data are unavailable for model training. Another distinctive feature of the present method is its high degree of interpretability, which directly describes the occurrence of a peak in a specific wavelength range in relation to the presence or absence of molecular fragments encoded in the descriptor X .

In this study, we focus on the advantages of directly predicting the entire spectral function, rather than predicting a predefined univariate functional feature, such as the maximum wavelength λ_{\max} . In the experiments shown later, it was confirmed that the prediction accuracy of λ_{\max} calculated from the predicted spectral function often significantly exceeds that of the conventional univariate output regression directly trained with the pre-quantified λ_{\max} . It should be noted that high-dimensional output variables ($Y(X, t_1), \dots, Y(X, t_m)$) are closely related. Learning a single model for multivariate outputs simultaneously can be considered a type of multitask learning. In multitask learning, multiple related tasks are learned simultaneously, allowing the model to recognize common mechanisms among target tasks and consequently improve the prediction accuracy of each task. A similar learning mechanism is expected to work in regressions with high-dimensional output variables.

Microstructure image prediction

Microstructures with varying morphologies, volume fractions, and grain-size distributions can be designed by controlling the composition of the material species and processing conditions. Here, we consider the problem of predicting the microstructures for any given compositional and processing parameters. To treat the microstructure as a model output, we used images obtained by optical or electron microscopy. In the development of composite materials, SEM or TEM has widely been used to analyse the surface and morphologies. Practically, for example, we aim to improve the mechanical properties of a material by controlling the composition and temperature to obtain a finer and more homogeneous grain structure. By

defining a microstructure as an image, we can address the machine-learning task by utilizing various well-established techniques in image recognition and computer vision. Specifically, the input variable X is given by a real-valued vector representing the compositional and processing parameters, and the output $Y(X, t)$ is defined as a microscopic image that takes matrix or tensor form for a grayscale or color image, respectively. Variable t represents a two- or three-dimensional image coordinate, and its support is discretized into pixel or voxel positions. The model can therefore be written in the form of a multidimensional vector regression, in the same way as the model for the spectral function described above (Figure 1(b)). Alternatively, in the context of computer vision research, this task can be regarded as machine learning for conditional image generation.

Other potential applications

There are many other applications in materials science where the prediction of functional outputs is applicable. Many physical properties are determined by the temperature, pressure, and frequency in an external electric field. The dielectric properties of a material, that is, the dielectric constant or dielectric loss tangent, are given as a function of the frequency and temperature⁴⁶ (Figure 1(c)). In this case, t comprises two variables. In addition, various polymeric properties, such as the specific volume, linear expansion coefficient, elastic modulus, specific heat, and thermal conductivity, are also dependent on temperature. From the observed transition points in the temperature dependence curve, the glass transition temperature, crystal melting point, and crystallization temperature were calculated. For the imaging analysis of materials, a wide variety of microscopes such as SEM, TEM and optical microscopes are commonly used, depending on the size of the object to be observed. In particular, various technologies of three-dimensional measurements have been established to analyze the internal structures of materials. For example, three-dimensional TEM allows us to observe the morphology of objects ranging in size from tens to hundreds of nanometers in three dimensions⁴⁷. X-ray computed tomography (CT) is a non-destructive technique that

is often used in medical applications. With this method, we can observe microstructures ranging from a few micrometers to millimeters in size without polishing or etching the sample surface⁴⁸. In addition, X-ray CT is a non-destructive inspection method that can be used to measure the fracture process of a material and the change in the material structure in response to heating in four dimensions (three-dimensional space plus time). Furthermore, synchrotron X-ray CT using high-brilliance synchrotron radiation can noninvasively measure the inner structure of materials with a high resolution of several hundred nanometers to several micrometers, even in metals where X-ray penetration is difficult⁴⁹. In principle, the proposed regression method can be applied to a wide range of high-dimensional functional data.

Methods

In this paper, we present two different regression methodologies for multidimensional functional outputs: deep generative modeling using adversarial learning and deep kernel regression for functional outputs. The former, originally developed for image-generation tasks³⁴, was introduced to solve the two research subjects. The latter is a newly developed method for overcoming the limited learning performance of deep generative models.

Conditional GAN

We constructed a neural network with the framework for functional output regression. cGAN consists of a generative model called a generator (G) and a binary classifier called a discriminator (D). The model structure is summarized in Figure 2 (see Supplementary Note for more details). With the generator $Y = G(X, Z)$, the output vector of the m absorption values in Equation 2 or the matrix of a grayscale microstructure image is modelled as Y . The input variables consist of X in the regression model (referred to as the conditional variable in the cGAN) and random noise Z . Noise $Z \sim N(0, 1)$ is assumed to follow a normal distribution

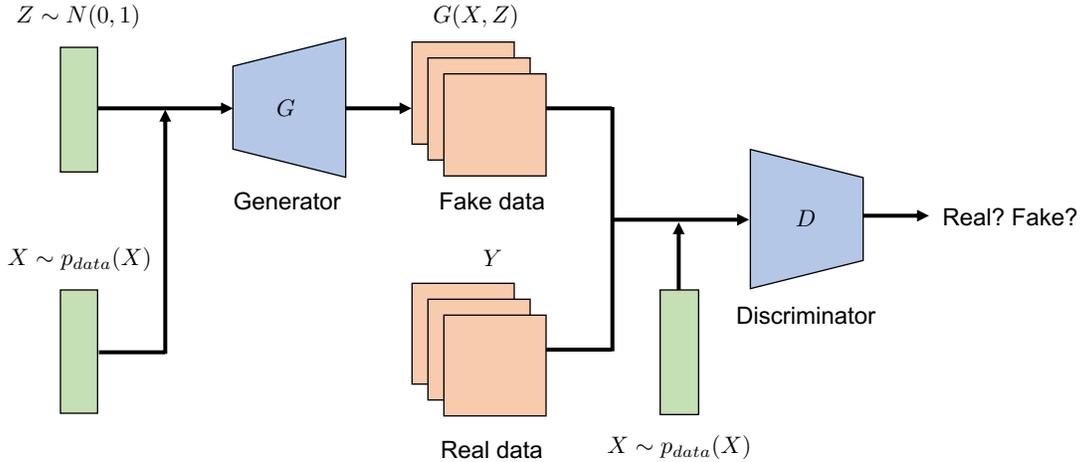


Figure 2: Architecture of the cGAN. The generator (G) consists of a fully connected embedding layer applied to the input X and repeating blocks, with each unit consisting of a convolutional layer and a leaky ReLU activation function. The number of neurons in the output layer is equal to the number of pixels in the image or observation points in the spectral function. The discriminator (D) is designed as a fully connected embedding layer or a conventional convolutional neural network. Detailed architectures are shown in the Supplementary Note.

with a mean of zero and unit variance. The input (X, Z) is transformed into an embedding vector by passing through a fully connected embedding layer with a batch normalization layer and a leaky ReLU activation function^{50,51}, and is further transformed into a vectorized spectrum or a microstructure image $Y = \mathbf{y}(X)$, as in Eq 2, by passing repeating blocks of different stacked hidden layers depending on the task. The discriminator $D(X, Y)$ is a binary classifier, in which the conditional variables X and Y are given as inputs. The discriminator judges whether object Y is a fake spectrum or an image generated from G or a real one in the given dataset. The discriminator D is modeled as a conventional fully connected neural network or a convolutional neural network⁵². The model structure, such as the number of layers and neurons in each layer, is determined based on the generalization performance in a separate validation dataset, while maintaining the basic form described here. The detailed settings for each problem are described in the Supplementary Note.

With this composite modelling, G and D are trained alternatively according to the fol-

lowing minmax strategy:

$$\min_G \max_D \mathbb{E}_{(X,Y) \sim p_{\text{data}}(X,Y)} [\log D(X, Y)] + \mathbb{E}_{Z \sim N(0,1), X \sim p_{\text{data}}(X)} [\log(1 - D(X, G(X, Z)))]. \quad (3)$$

The first term becomes larger as $D(X, Y)$ increases, that is, when the discriminator D correctly identifies the input real object as real. The second term becomes large when $1 - D(X, G(X, Z))$ becomes large, that is, when D successfully recognizes the fake $Y = G(X, Z)$ to be fake. The discriminator D is learned such that the classification error is minimized, and G is trained to reduce the second term such that D is misrecognized. By alternately training G and D , we derive G , which can produce high-quality fake spectral functions or microstructure images for any given descriptor X . For spectral data, we generated r Gaussian samples $\{Z_i | i = 1, \dots, r\}$ ($r = 100$) and used the ensemble $\mathbf{f}_X = \frac{1}{r} \sum_{i=1}^r G(X, Z_i)$ of the learned generator to improve the smoothness of the predicted spectrum.

The cGAN can be regarded as a supervised learning technique for multidimensional output variables. Because cGANs have been intensively studied, particularly for image-generation tasks, when treating images as the output variable, we can take advantage of the wealth of tips and various extended works that have been accumulated in machine-learning research. However, similar to other conventional generative adversarial networks, cGANs suffer from instability during the learning process. In particular, vulnerability to small datasets has been pointed out in many previous studies. When training data are in limited supply, the discriminator can easily overfit the data to make a perfect true/false classification, which leads to gradient vanishing and halting of the learning process before a sufficiently accurate generator, that is, a regression model, is created. The solution to stabilize the adversarial training process is to balance the learning progress of G and D , but this is not an easy task. The primary reason for this is the over-parameterization of the generator caused by the high dimensionality of the vectorized object Y . In the case studies shown below, the dimension of Y is more than 2,000 for spectral function prediction and up

to 10,000 for microstructural microscope images with a resolution of 100×100 . To describe the mapping from X to such a high-dimensional Y , it is necessary to introduce one or more transposed convolution matrices of large sizes. In applications shown later, the total numbers of model parameters in the trained generators reached the order of 10 million and 5.8 million for the spectral function prediction and microstructure image prediction, respectively.

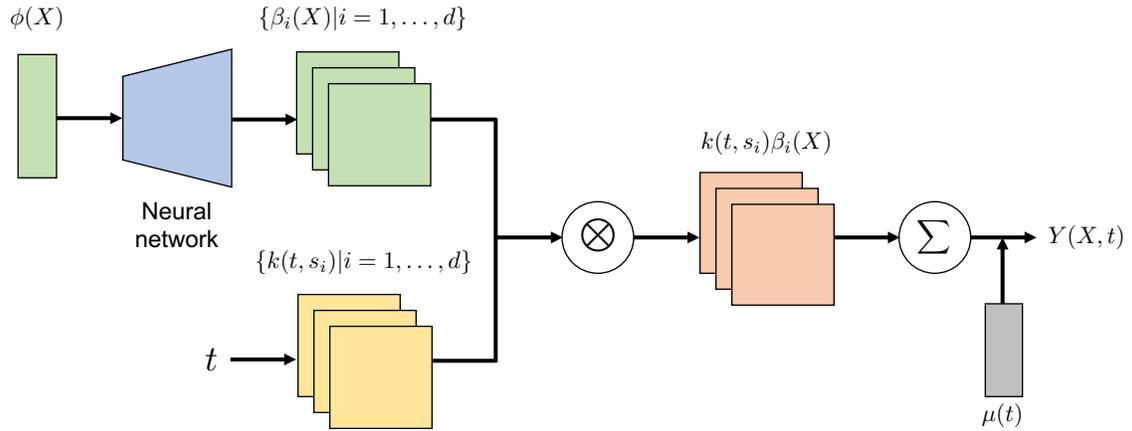


Figure 3: Model architecture of the functional output kernel regression. The coefficient functions $\{\beta_i(X) | i = 1, \dots, d\}$ for a set of d pre-defined kernels $\{k(t, s_i) | i = 1, \dots, d\}$ depend only on input X . The mapping from X to $\beta_i(X)$ is modeled by a fully connected embedding layer, followed by repeating units of a convolutional layer and a leaky ReLU activation function. The number d of kernel functions is appropriately controlled to be smaller than the number of observations for t , that is, $d \leq m$.

Kernel regression with functional outputs

In addition to cGAN, we present another model with a simple, naturally interpretable model structure. The design concept is inspired by regression models for functional outputs, which have been studied in the context of functional data analysis. The functional output $Y(X, t)$ is modeled as follows:

$$Y(X, t) = \sum_{i=1}^d k(t, s_i)\beta_i(X) + \mu(t) + \epsilon, \quad (4)$$

The first term is the weighted sum of the d kernel basis function $\{k(t, s_i)|i = 1, \dots, d\}$. The kernel centers s_i are equally spaced in the domain of wavelengths or image coordinates. In this study, we used the Gaussian radial basis function (RBF) kernel as:

$$k(t, s_i) = l \exp\left(-\frac{\|t - s_i\|^2}{2\sigma^2}\right). \quad (5)$$

The variance $\sigma^2 > 0$ and length scale $l > 0$ are hyperparameters adjusted based on the evaluation of the generalization performance. Alternatively, one may predetermine the hyperparameters based on the empirically known resolution of a measurement system or the inherent variation of $Y(X, t)$ in varying t for the physical system of interest. The regression coefficient $\beta_i(X)$ depends only on the input variable X , which is modelled by a neural network, as described below. $\mu(t)$ is a baseline function estimated by imposing smoothness as a regularizer. ϵ denotes the noise term.

The overall model represents a system in which each kernel function pre-arranged in the domain of variable t is activated or deactivated depending on the input variable X , for example, the presence or absence of a specific fragment in a fingerprinted chemical structure X . One advantage of this modelling is that the number of parameters can be reduced by controlling the number of kernels d placed in the domain t . In a cGAN generator, the number of neurons in the output layer inevitably increases because it is constrained by the dimensionality of Y .

The regression coefficients $\{\beta_i(X)|i = 1, \dots, d\}$ are modelled differently using neural networks for the two tasks (Figure 3, Supplementary Note). The modelling is arbitrary in general. Here, we used neural networks with a structure similar to that of the generator in cGAN. To predict the optical absorption spectra, the input X is a 1,024 binary vector that encodes the chemical structure of a molecule using the extended connectivity fingerprint (ECFP fingerprint¹⁰ with a radius of 3). The mapping from X to the d output coefficients is modelled by multiple blocks of stacked hidden layers, including a fully connected layer,

batch normalization layer, and leaky ReLU activation function. For microstructure image prediction, as detailed later, the input X includes six processing parameters and 100 random noises. The mapping from X to the d output coefficients consists of multiple blocks of stacked layers, including a transposed convolutional layer, batch normalization layer, and leaky ReLU activation function. Hyperparameters such as the number of convolutional layers and neurons are tuned based on a separate validation dataset. See Supplementary Note for the procedure for hyperparameter tuning.

In model training, the following empirical risk is minimized with respect to the parameters in the model of $\{\beta_i(X)|i = 1, \dots, d\}$ and the baseline function $\mu(t)$:

$$L(\beta, \mu) = \sum_{(X,t) \in \mathcal{D}_{\text{obs}}} D(Y(X,t), \hat{Y}(X,t)) + \lambda \sum_{i=1}^m \sum_{j \in \mathcal{A}_i} (\mu(t_i) - \mu(t_j))^2, \quad (6)$$

The first term involves the discrepancy D between an observed $Y(X,t)$ and its prediction counterpart $\hat{Y}(X,t) = \sum_{i=1}^d k(t, s_i) \beta_i(X) + \mu(t)$. In the experiments shown later, for both spectral prediction and microstructure prediction, D is defined by an ordinary squared loss, which is summed over all observations of X and t . The second term is the regularization term for the baseline function. Regularization induces a smooth transition between $\mu(t_i)$ and $\mu(t_j)$ for an observation point t_i and its neighborhood $t_j \in \mathcal{A}_i$. In this formulation, the observed series of $Y(X,t)$ for t is allowed to have missing values.

Results and Discussion

We highlight the potential prediction capability and learning mechanism of the proposed methods by presenting three application examples. The objective of the first two examples is to predict the function of the UV-Vis or NIR absorption spectrum of an organic molecule, taking its chemical structure as an input. In the first case, the number of samples was several thousand or less, whereas in the second case, the number of samples was only 68. The third example is to predict electron microscopy images of microstructures from the composition

Table 1: Comparison of the prediction accuracy for optical absorption spectral functions. Datasets I and II cover the UV-Vis spectra, and the USGS spectral library covers data in the NIR domain. The performance metrics shown for the LSTM and attention-based encoder-decoder models were those reported in Urbina et al.³¹.

		RMSE	R^2	MAE	RMSE derivative
Dataset I	LSTM	0.169 ± 0.132	0.626 ± 1.166	0.119 ± 0.106	0.013 ± 0.010
	Attention	0.154 ± 0.144	0.680 ± 1.230	0.091 ± 0.120	0.018 ± 0.020
	Kernel	0.111 ± 0.009	0.798 ± 0.043	0.075 ± 0.006	0.012 ± 0.001
	cGAN	0.112 ± 0.015	0.791 ± 0.056	0.072 ± 0.011	0.054 ± 0.002
Dataset II	LSTM	0.064 ± 0.062	0.710 ± 0.472	0.047 ± 0.075	0.008 ± 0.006
	Attention	0.055 ± 0.071	0.699 ± 0.259	0.044 ± 0.052	0.006 ± 0.007
	Kernel	0.093 ± 0.004	0.655 ± 0.007	0.066 ± 0.001	0.009 ± 0.000
	cGAN	0.086 ± 0.002	0.715 ± 0.017	0.058 ± 0.002	0.017 ± 0.001
USGS	Kernel	0.076 ± 0.024	0.602 ± 0.200	0.053 ± 0.009	0.022 ± 0.003
	cGAN	0.077 ± 0.004	0.495 ± 0.114	0.060 ± 0.009	0.533 ± 0.043

and processing conditions of the fabricated thin metal films.

Prediction of UV-Vis absorption spectrum

Urbina et al.³¹ produced two experimental datasets of UV-Vis spectra that encompass different chemical spaces. In the paper, these datasets are referred to as Dataset I and Dataset II. Dataset I consisted of the absorption spectra of 949 different commercial compounds measured using a high-performance liquid chromatography system. Dataset II contains the spectra of 2,222 different commercially available pharmaceutical molecules, which were measured with a spectrophotometer in a multi-well plate format. The absorption spectra were measured at 181 and 171 wavelengths equally spaced in the UV-Vis (220-400 nm) for Datasets I and II, respectively. Compared to Dataset I, Dataset II has a larger amount of data and a larger diversity of chemical structures. For model training and evaluation of prediction performance, approximately 70% of the compound set was randomly selected as the training set and the remaining approximately 15% as the test set. Partitioning of the dataset was performed according to the compound species to avoid multiple spectral profiles of the

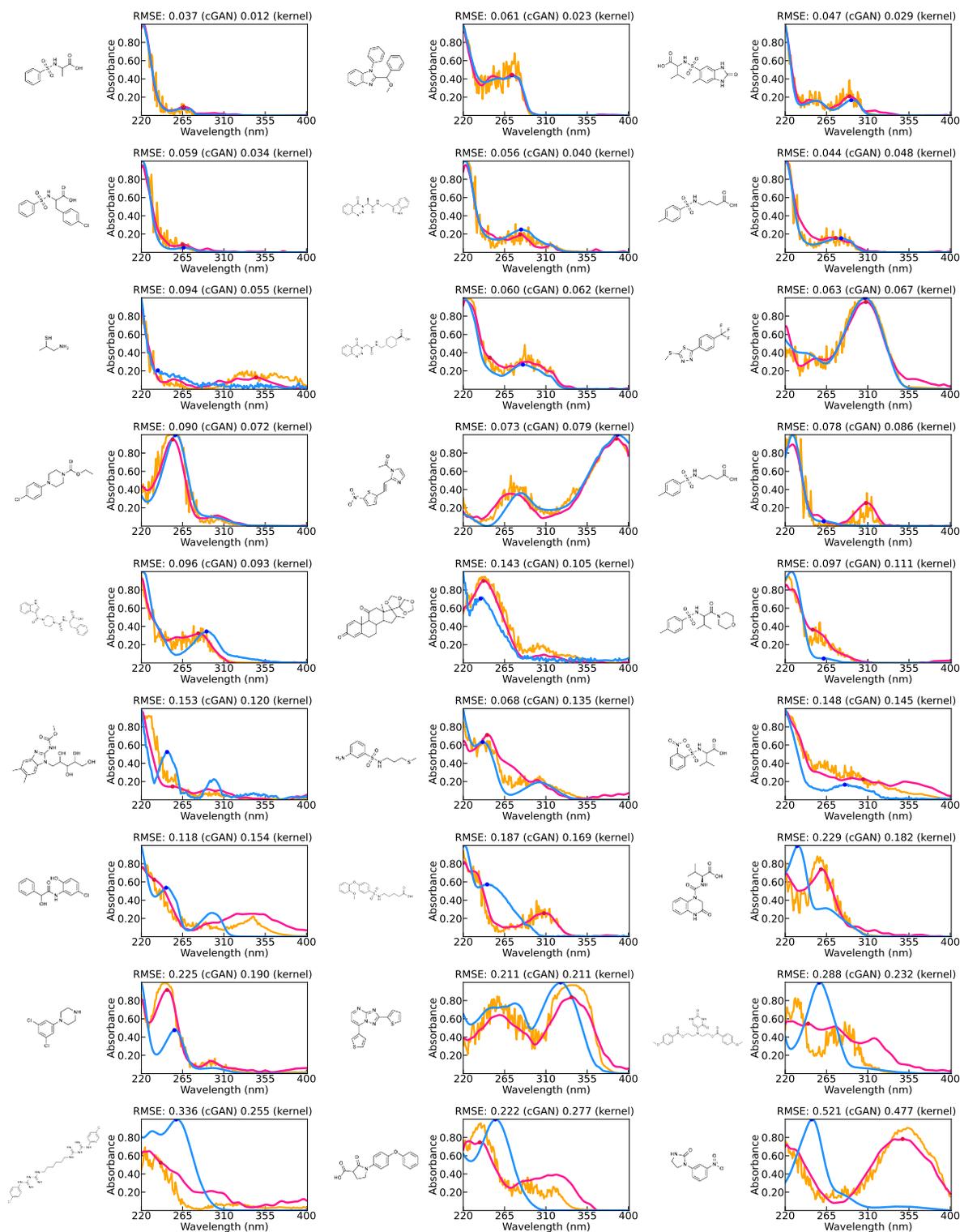


Figure 4: Prediction results of (a) the functional output kernel regression (pink) and (b) cGAN (orange) with observed UV-Vis spectra (blue) in Dataset I.

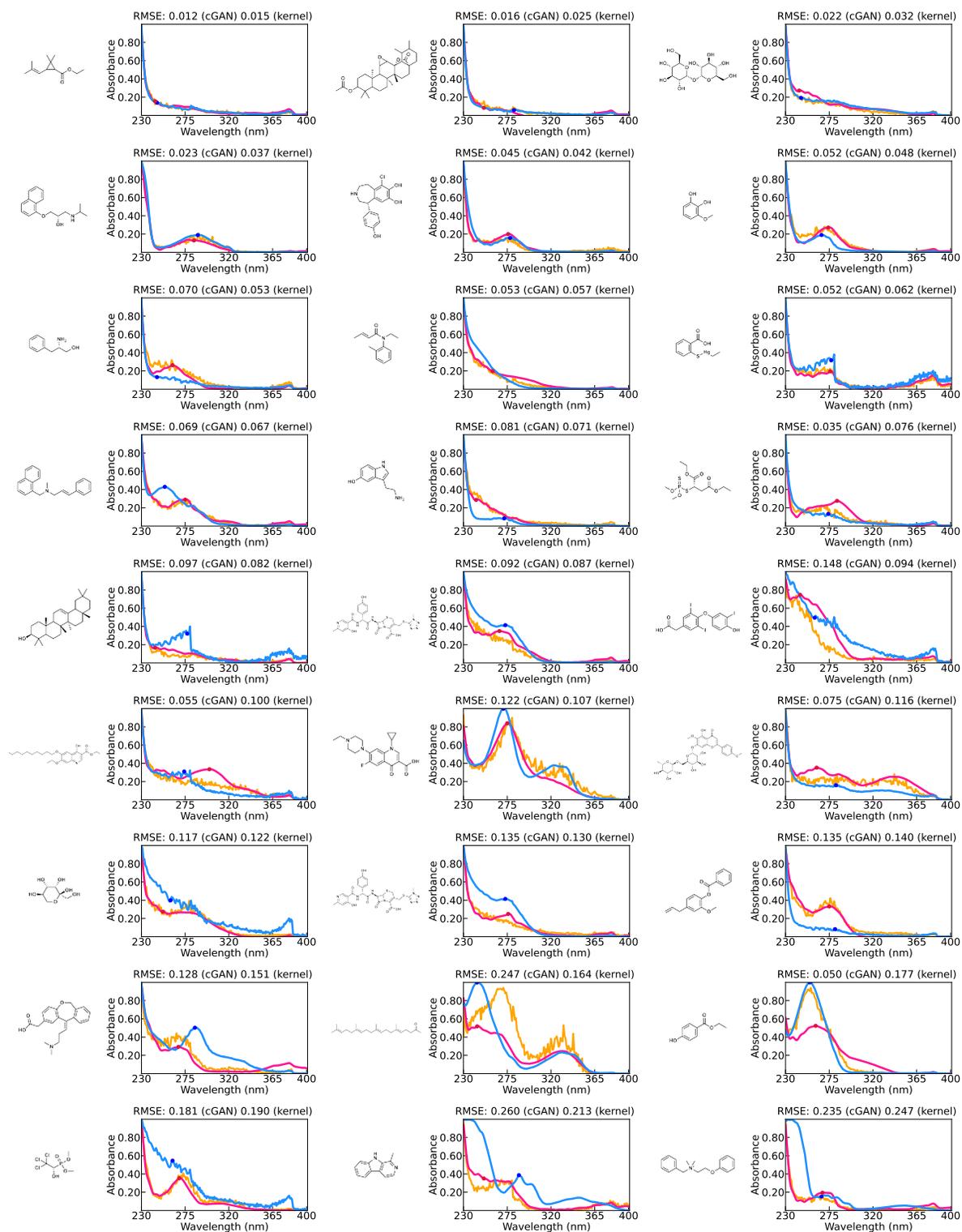


Figure 5: Prediction results of (a) the functional output kernel regression (pink) and (b) cGAN (orange) with observed UV-Vis spectra (blue) in Dataset II.

same compound leaking into the test set. In Urbina et al.³¹, the results of applying two encoder-decoder architectures with LSTM cells and an attention mechanism were reported, respectively, which were compared with the performance metrics of the present methods.

For kernel regression, the observed wavelength range 220 or 230-400 nm was divided into 128 equally spaced segments, and the kernel centers s_1, \dots, s_d ($d = 128$) were placed there. The set of hyperparameters consists of the variance σ^2 , length scale l in the RBF kernel, and the number of hidden layers in the neural network. For each σ^2 and l , three and four grid points were set as candidates. The number of hidden layer blocks was set between 1 and 4. Once the number of blocks was determined, the number of neurons in each layer was determined, as shown in Supplementary Note. Correspondingly, the total number of candidates for the hyperparameter search is 48. The generalization performance of the trained model for each candidate was measured using the root mean square error (RMSE) on the validation dataset, and the best combination was identified.

The hyperparameter of cGAN is given by the network structure of the generator and discriminator. As in kernel regression, both network structures consist of multiple blocks of stacked hidden layers, including a fully connected layer, a batch normalization layer, and a leaky ReLU activation function. The difference from kernel regression is that the output variables are directly formed by the m absorbances of the different wavelengths in the spectral function. The number of blocks to be searched was between 1 and 4, and the number of neurons in each layer was determined, as shown in the Supplementary Note. The generalization performance of the models was measured using the RMSE of the validation dataset to identify the best hyperparameter combination.

Through the process of validation, in Dataset I, the numbers of stacked hidden layers for the neural networks in the kernel regression, generator, and discriminator of the cGAN were selected to be 4, 4, and 2, respectively. In Dataset II, the number of stacked hidden layers for the neural networks in the kernel regression, generator, and discriminator of the cGAN were selected as 3, 4, and 3, respectively. The value of σ^2 was selected as 0.0005 for

both datasets, and l was selected as 0.5, or 5 for Dataset I or Dataset II, respectively. To evaluate the performance of these models, we calculated the RMSE, coefficient of determination (R^2), and mean absolute error (MAE) between the m predicted spectral values and their observations for each test molecule, and compared the median of each performance metric for the 150 and 342 test molecules. We also considered the differences in the spectral series and evaluated the gradient-level prediction performance using the same procedure. Table 1 summarizes the means and standard deviations of the performance measures for the three independent numerical experiments; the prediction accuracy of the LSTM-based and attention-based encoder-decoder models reported in Urbina et al.³¹ is also given. In Dataset I, kernel regression outperformed the other three, but the difference with cGAN was not as pronounced. The reason for the lower prediction accuracy of the difference spectrum of cGAN is that it uses an ensemble average of the randomly sampled Z in the calculation of the predicted spectra, which results in a loss of smoothness in the gradient of the spectrum. However, this problem can be solved by smoothing the predicted spectrum in the post-processing step. As the code for the encoder-decoder models is not distributed, we could not go further into the comparison of the models, but it should be concluded that in this case study, they have almost the same performance.

Here, the predicted spectral functions and their observed values are presented exhaustively to obtain a view of the predictive capability of the kernel regression and cGAN. Figures 4 and 5 show the prediction outcomes for 27 randomly selected test molecules in datasets I and II, respectively. To obtain a more comprehensive view of the prediction accuracy, Supplementary Note also provides the results of 60 randomly selected test molecules in each dataset. According to a careful visual inspection, it can be seen that the models retain a surprisingly high prediction accuracy. For a significant number of molecules, the positions of multiple peaks in the absorption spectrum and the shape of the function were almost perfectly predicted. In some cases, even features that are not visually obvious, such as plateaus or tiny peaks, can be captured appropriately. This observation suggests that the presence

or absence of chemical substructures is a major factor in the optical absorption spectra of molecules. There is no significant difference between kernel regression and cGAN in terms of capturing the broad trend of the spectral function. However, as mentioned above, cGAN requires some effort to detect the peak position because of the noise fluctuations of Z in the prediction equation. Even when smoothing is applied, unexpected false peaks can occur. Therefore, we conclude that kernel regression has an advantage in terms of spectral prediction.

Spectral prediction with limited data in the USGS spectral library

The U.S. Geological Survey (USGS) Spectral Library Version 7 contains reflectance optical spectra of over 1,000 different molecules, each of which was measured in two different sets of environments: laboratory, field, and airborne spectrometers. If the data were duplicated, we selected a set of spectral data measured at room temperature as the test case. All spectral values were recorded at 2,151 common discrete points approximately equally spaced in the wavelength range 0.35-2.5 μm from the ultraviolet to the far infrared. Compared to the number of UV-Vis data mentioned above, the sample size of 68 was quite small. The data were recorded as reflectance R . The reflectance and absorption are inversely proportional to each other. Therefore, it is difficult to express the function of a reflection spectrum as the sum of the RBF kernels, as in Eq. 4. Therefore, model training was performed after converting R to absorbance A according to equation $A = \log_{10} \frac{100}{R}$. The model structure and procedure for selecting hyperparameters for the cGAN and kernel regression were exactly the same as those for the analysis of the UV-Vis spectra described above. The numbers of stacked hidden layers for the neural networks in the kernel regression, generator, and discriminator of the cGAN were selected to be 3, 3, and 2, respectively. The values of σ^2 and l are set to 0.0005 and 1, respectively. 80%, 10%, and 10% of instances randomly selected from the entire dataset were used for model training, hyperparameter validation, and performance evaluation, respectively. In addition, data splitting was independently performed three times

to calculate the mean and variability of the performance measures.

The prediction accuracies of cGAN and kernel regression for the test instances, which were trained on the optimized hyperparameters, are summarized in Table 1. Unlike the results of the UV-Vis spectra, the kernel regression overwhelmingly outperformed the accuracy of cGAN, for example, the R^2 values for the cGAN and kernel regression were 0.495 ± 0.114 and 0.602 ± 0.200 , respectively. It is likely that for a given small dataset, the over-parameterization of the generator in the cGAN causes degradation in the predictive performance. However, the kernel regression reached a sufficiently high prediction accuracy despite being trained on only 54 samples. Figure 6 shows the predicted and true spectra for several cases (see Supplementary Note for a more comprehensive visualization of the prediction results). As in the UV-Vis cases, the peak positions and functional features of the observed spectra can generally be predicted. We also observed that, in many cases, the plateau and minor peaks could be properly captured. Kernel regression has a strong tolerance for a limited sample size, which will be further investigated later.

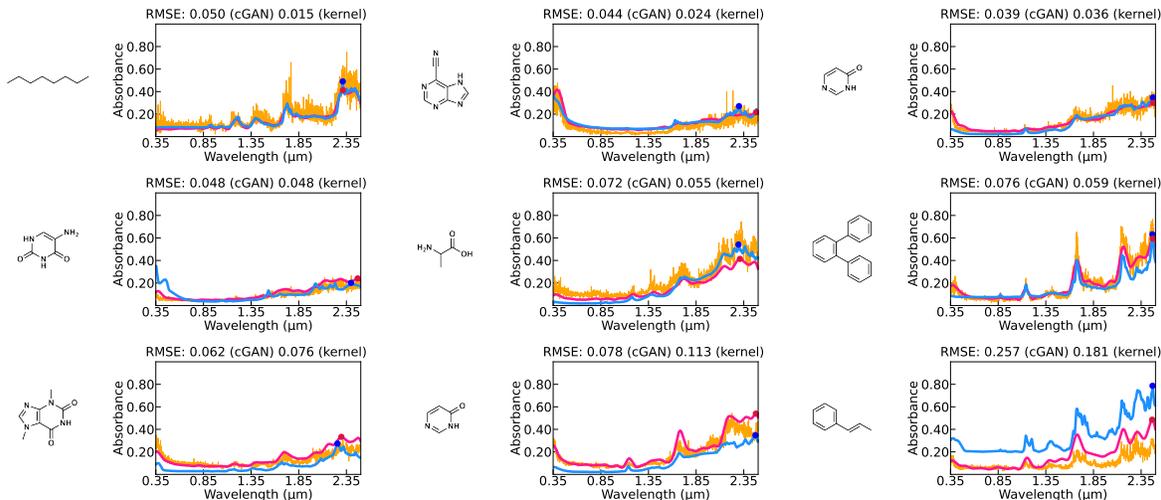


Figure 6: Prediction results of (a) the kernel regression (pink) and (b) cGAN (orange) for the optical absorbance spectra of nine test molecules with experimental profiles (blue) in the USGS spectral library.

Microstructure image prediction

A thin-film material consisting of a chromium (Cr)-based metal plate coated with aluminum (Al) was analyzed³⁴. Cr and Al metal plates were placed face to face, and argon (Ar) gas was sprayed onto the Al plate at high speed by magnetron sputtering so that the ejected Al atoms were adsorbed onto the Cr plate. The input $X \in \mathbb{R}^6$ to the model is a six-dimensional real vector representing the composition $\text{Cr}_{1-x}\text{Al}_x\text{O}_y\text{N}$ and the processing conditions.

1. Cr and Al content x
2. O content y
3. Temperature at which Al is adsorbed
4. Pressure at which Al is adsorbed
5. Average energy of the Ar ions when they are incident on the Al metal plate
6. Ionization degree of the Ar gas

Each element of the input vector X was normalized to have a mean of zero and a unit variance in the training data. The output variable Y is the SEM image of the microstructure with a resolution of 100×100 .

In functional output kernel regression, the centers of 32×32 RBF kernels ($d = 32 \times 32$) were placed at equally spaced positions in the two-dimensional image coordinate space. The modelling and training conditions were almost the same as those used to predict the spectral functions. The hyperparameter set to be explored was the network structure and the variance and length scales of the RBF kernel. The input variables were transferred to the embedding latent space via a fully connected layer. A repeating unit consisting of a transpose convolution layer and a leaky ReLU activation function was applied several times to this embedding vector, which was finally transformed into an output variable via a fully connected layer. The numbers of stacked hidden layers for the neural networks in the

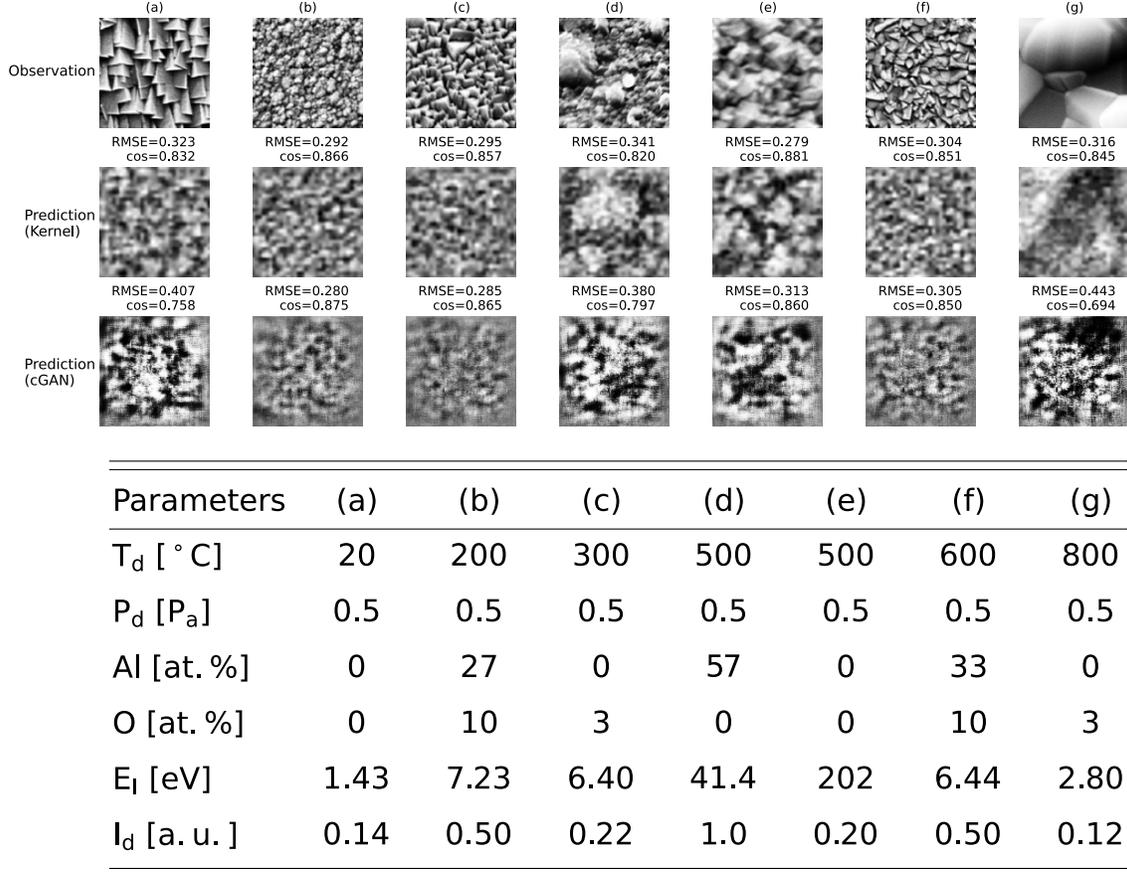


Figure 7: Results of the microstructure prediction using the functional output kernel regression and cGAN with respect to seven different SEM images.

kernel regression, generator, and discriminator of the cGAN were selected as 4, 1, and 2, respectively. The values of σ^2 and l of 0.0001 were selected as σ^2 , and five were selected as l . The cGAN modelling is also similar to the spectral function prediction in that the input variable is transformed into the output variable via a series of fully connected layers to define the feature embedding, a repeated application of the transpose convolution layer and leaky ReLU, and a fully connected layer to generate a 100×100 image Y .

The data contained 123 SEM images with their compositional and processing conditions, of which 90%, 5%, and 5% were randomly assigned to the training, validation, and test sets, respectively. Data partitioning was independently repeated thrice. The generalization performance of the model was investigated by varying the number of layers from one to four, with each repeating unit consisting of the transposed convolutional layer and leaky ReLU

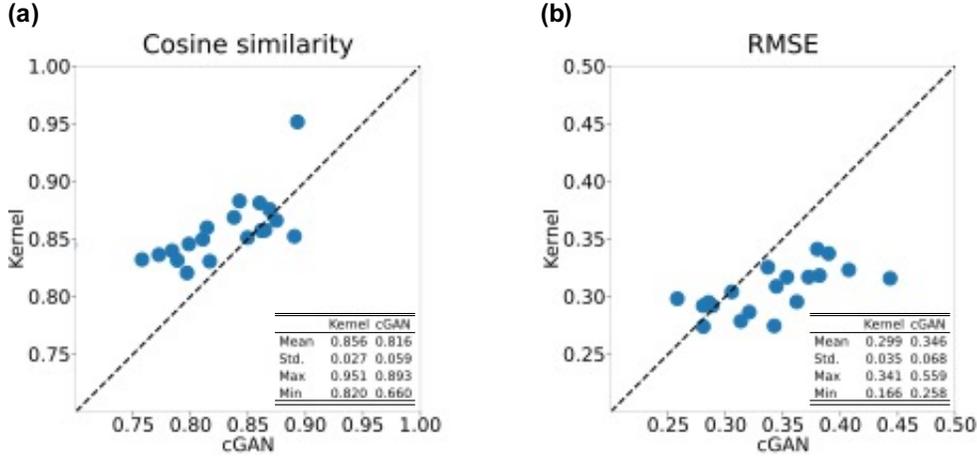


Figure 8: Cosine similarity and RMSE values of the predicted SEM images with respect to test data for the functional output kernel regression and cGAN.

activation function, while the embedding layer and output layer were fixed to one layer each. The number of neurons in each layer was designed as described in the Supplementary Note.

Figure 7 shows the experimental and predicted SEM images for the functional output kernel regression and cGAN for the seven randomly selected test conditions. To obtain a comprehensive view of the prediction performance, the Supplementary Note provides the prediction results for 14 randomly selected test conditions and training results for 60 randomly selected conditions. Comparing with the experimental SEM images, it can be seen that the predicted images of the functional output kernel regression could capture the difference in morphological features of microstructures, such as grain sizes, even though the number of training data was as small as 109. As previously known, the cGAN was unable to predict observed features of microstructures, possibly due to the limited amount of training data. We calculated the cosine similarity and RMSE as measures of image similarity. Figure 8 summarizes the comparison of cosine similarity and RMSE for the functional output kernel regression and cGAN. Obviously, the functional output kernel regression outperforms the cGAN in both cosine similarity and RMSE.

As with the spectral prediction, a high learning potential of the function output kernel regression on small data was confirmed. However, the experimental results also indicated

the weakness of the trained model as an image generator. Although the predicted images successfully captured the morphological features of the microstructures, they were less clear than those of the experimental images. The reduction in sharpness is owing to the blurring effect of using kernel functions. For functional output regression, the parameter savings using a basis set of kernel functions leads to high tolerance for small data, but at the same time, it inevitably leads to a reduction in the quality of the generated images. However, unlike image generation in computer vision, the generation of high-quality images is not as important for prediction tasks in materials science.

Tolerance to data size in function output kernel regression

We investigated the underlying mechanisms behind the success of the learning, despite the exceedingly small amount of data, that is, 54 training samples for spectral prediction and 100 or more for microstructure prediction. Functional output variable regression has a learning mechanism that is common to multi-task learning. The tasks of predicting multiple function values are not independent, but are related to each other. Joint learning of multiple related tasks is expected to be advantageous for the model to acquire common representations across tasks, if it exists. In addition, data accumulation by the simultaneous use of data from multiple tasks is generally advantageous in suppressing overlearning of task-specific noises. These mechanisms may be responsible for the success in building the model by scratching with a limited amount of data. To confirm this hypothesis, we conducted three numerical experiments using the optical absorption spectra in Dataset I.

- (a) The maximum absorption wavelength λ_{\max} and its intensity value were calculated from the spectral function predicted by the functional output kernel regression, and the accuracy of predicting λ_{\max} and the intensity was verified against their observed values.
- (b) Using λ_{\max} and its intensities as a dataset that were extracted from the observed spectra

in advance, we built random forest regressors that directly predict the extracted scalar values of λ_{\max} and the intensity from the fingerprinted chemical structure (the same to the one used in the cGAN and kernel regression).

- (c) We trained m models independently and separately that predict the spectral values of each of the m equally spaced grid points. The estimated λ_{\max} and intensity values were calculated from the predicted spectral function.

In the peak detection from a spectral profile, λ_{\max} was selected as the wavelength of the maximum intensity among the maxima that appeared 232 nm from the left end. In these three experiments, the ratios of the training, validation, and test sets were set to 649:151:150 and 100:425:425, respectively. The data partitioning was repeated ten times independently, and training and testing were performed with random forests (see Supplementary Note for details on model training and hyperparameter adjustment). For the evaluation of the accuracy of the test set, cases in which the predicted peak position was within 10 nm of the observed peak position were defined as correct, and cases in which the predicted peak intensity was within 0.1 of the observed intensity were defined as correct. As shown in Table 2, for both models obtained from the smaller and larger datasets, the prediction accuracy of the kernel regression with the entire function trained simultaneously (model (a)) overwhelmingly exceeded the performance of models (b) and (c). In particular, compared to model (c), which learned and predicted spectral values independently, model (a) showed little degradation in prediction performance when the amount of training data was reduced to 100.

This observation has implications for the methodological construction of machine learning-based material-property prediction. For example, in the prediction of a temperature-dependent property, it is often the case that the temperature range is limited to room temperature, and the model is made. However, such an approach makes the problem more difficult and reduces prediction accuracy. When functional data are available, direct prediction of the functional output variable can significantly improve prediction accuracy.

Table 2: Tolerance to small data in predicting maximum absorption wavelength λ_{\max} and peak intensity. Three cases were tested: (a) joint learning and prediction of the entire function calculated from the prediction spectrum of the function output kernel regression, (b) direct prediction of the two scalar quantities using random forest (RF), and (c) random forest regression independently trained for each wavelength. The ratios of the training, validation, and test sets were set to 649:151:150 and 100:425:425, respectively.

Method	Data Split	Position λ_{\max} (%)	Intensity of λ_{\max} (%)
(a) Kernel	649:151:150	46.3	43.9
(b) RF for λ_{\max}		38.7	36.2
(c) RF for each wavelength		43.3	33.2
(a) Kernel	100:425:425	36.1	32.6
(b) RF for λ_{\max}		31.8	23.9
(c) RF for each wavelength		30.5	25.0

Conclusions

In the past few years, a variety of machine learning techniques have been introduced in materials science. In this context, machine learning techniques for predicting the physico-chemical values of scalar quantities from input materials have matured. On the other hand, in this study, we focused on the regression problem where the output variable is given in the form of a function. We describe the potential problem setting in materials science and present two methodologies based on deep generative models and statistical functional data analysis. Because machine learning research based on this perspective is still in its infancy and not mature, there must be other promising methods besides the one proposed here. As a starting point, we have demonstrated the potential of functional output regression in two cases: the prediction of the light absorption spectral function of a molecular system and the prediction of the microstructure from experimental condition parameters.

Of particular interest is the mechanism of the high tolerance of the functional output regression to the limited availability of training data. In this study, the amount of data was much smaller than in general situations: 54 samples in the prediction of absorption spectra and 109 samples for microstructure image prediction. Despite the extremely small sample

sizes, models with sufficiently high prediction accuracies were successfully obtained. In general, machine learning prediction of physical or chemical properties is often performed by transforming the functional data into a few features and then predicting the scalar output variables, instead of directly predicting the functional data. However, the experimental results suggest that the direct prediction of functional output variables may involve a learning mechanism that favors the acquisition of higher generalization performance than the prediction of scalar variables. Intuitively, the higher dimensionality of the output variable is likely to lead to overlearning and a decrease in the generalization performance of the trained model. However, because the tasks of simultaneously predicting multiple function values are not independent but strongly related to each other, the simultaneous use of data from multiple tasks can suppress task-specific noise due to data expansion. In addition, in multi-task modeling, the complexity of the model does not increase significantly, as usually only one- or two-dimensional input variables t are added. In the trade-off between increasing model complexity and data expansion, the advantages of the latter tend to outweigh those of the former.

There are many other potential applications in material research, where the prediction of functional outputs is required. Most material properties are given as functions of temperature or frequency. The dielectric properties are defined as functions of the temperature and frequency. Polymeric properties such as the specific volume, coefficient of linear expansion, bulk modulus, specific heat, thermal conductivity, and χ parameters were also determined in a temperature-dependent manner. From these temperature-dependent curves, important properties such as the glass transition temperature, crystal melting point, and crystallization temperature can be calculated. The dependence of properties on processing conditional parameters is another typical application of functional output regression. Furthermore, in the 2D and 3D imaging of material structures, the output variable is given as a multidimensional function in the image coordinate space. Due to the lack of data availability, we have only presented a few limited applications, but there are many problem settings for functional

output regression that remain unexplored in materials research. In principle, the proposed methods are designed to handle arbitrary high-dimensional functional data. We hope that the distributed Python code can be utilized to discover more problem settings.

Supporting Information

- **Supplementary Note:** modelling details for cGAN and the functional output kernel regression, procedure of hyperparameter search, supplementary figures showing results of predicting the optical absorption spectra in the UV-Vis region and those in the USGS spectral library, and the microstructure image prediction (PDF)
- **Python code:** https://github.com/yoshida-lab/XenonPy/blob/master/samples/kernel_neural_network.ipynb

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Author contributions

Ryo Yoshida and Megumi Iwayama devised the project, main conceptual ideas, and proof outline. Megumi Iwayama implemented the machine learning algorithms and conducted the experiment with the support of Chang Liu. Megumi Iwayama and Ryo Yoshida wrote the manuscript with the support of Stephen Wu. Ryo Yoshida supervised this project.

Competing interests

The authors declare no competing interests.

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