



NIA-AA Revised Criteria for Diagnosis and Staging of Alzheimer's Disease: Impact on Diagnosis and Treatment of Underrepresented Groups

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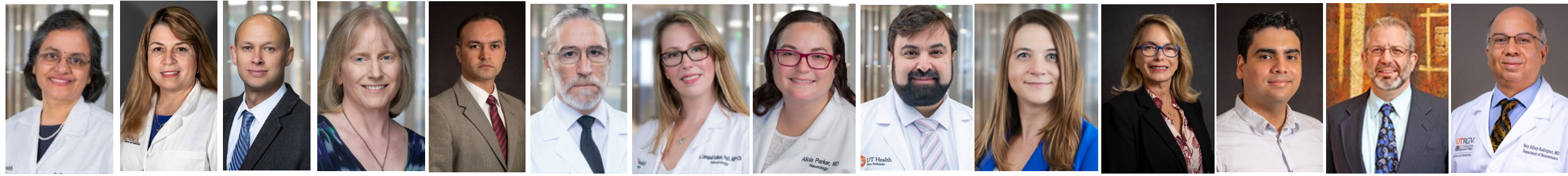
DISCLOSURES

None

FUNDING

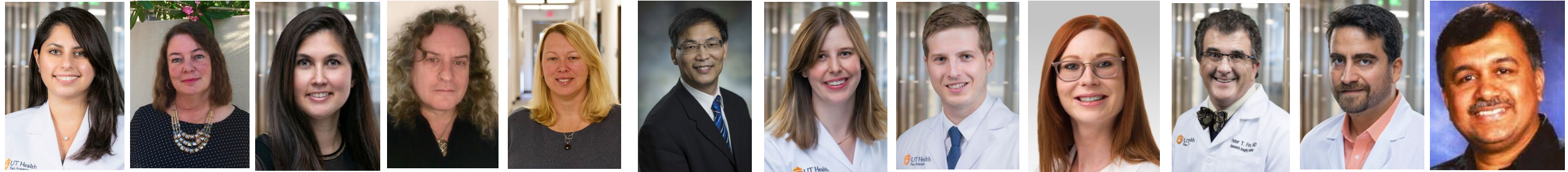
1P30AG059305-01, 1P30AG066546-01A1,
DP1AG069870

Texas Alzheimer's Disease Research and Consortium



Administrative

Clinical



Population Neuroscience

Genomics

Biomarker

Neuropathology

Imaging



OREC

Data Management

REC

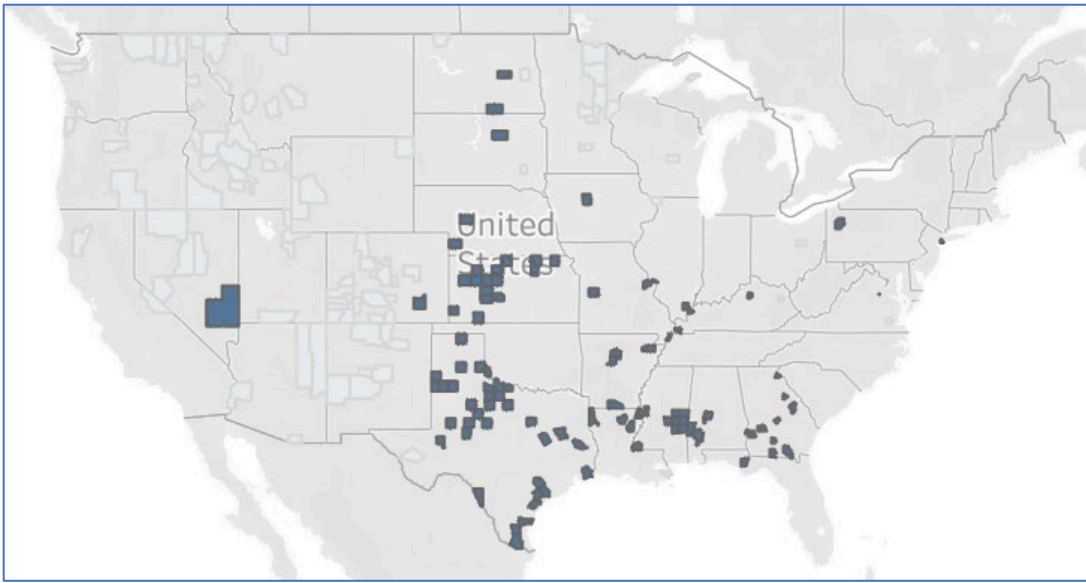
SOUTH TEXAS ALZHEIMER'S DISEASE RESEARCH CENTER



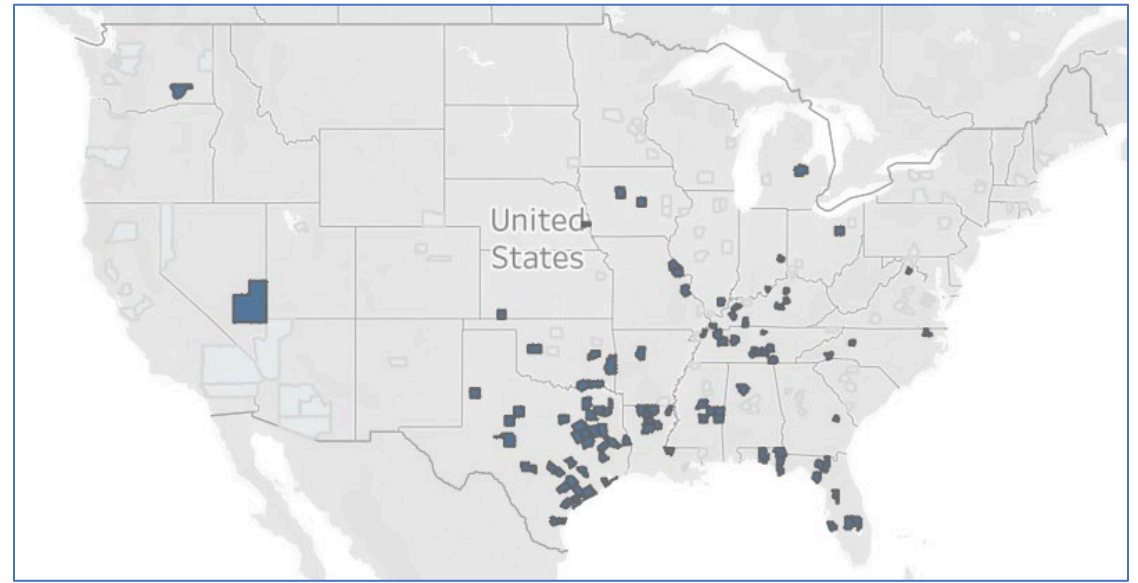
REC Scholars

Core Concepts

- **Impact:** Change.
- **Diagnostic Process:**
 - Is a complex, patient-centered, collaborative activity involving information gathering and clinical reasoning to determine a person's health situation. National Academies Press (US); 2015
<https://www.ncbi.nlm.nih.gov/books/NBK338593/>
 - cyclical process of information gathering, integration and interpretation, and forming a working diagnosis Parasuraman et al. (2000) PMID: 11760769.
- **Treatment:** provision, coordination or management of health care and related services by one or more providers.

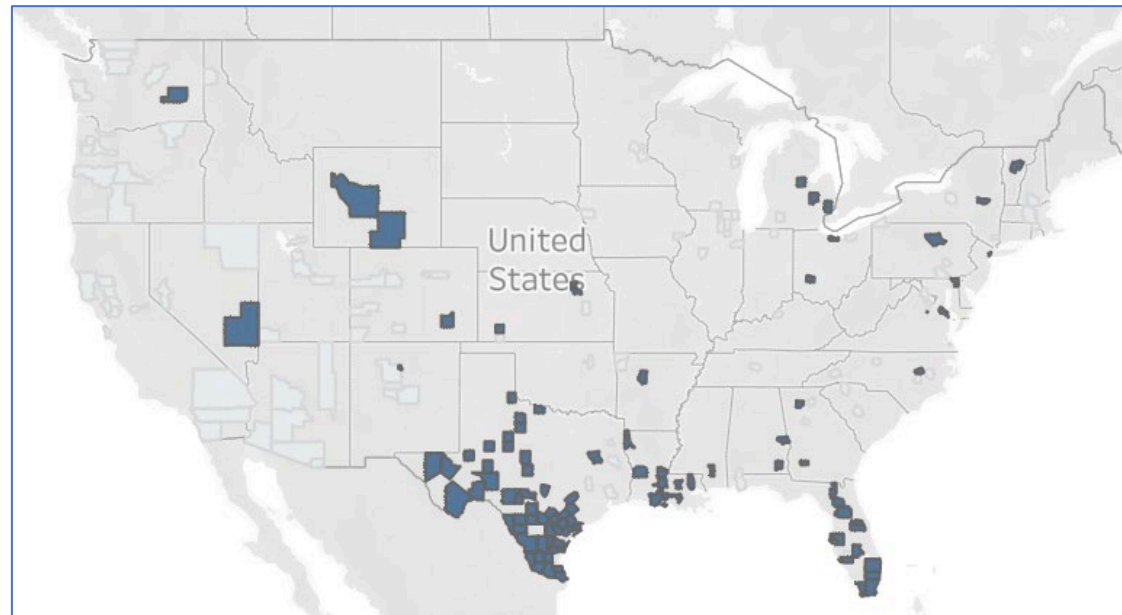


Non-Hispanic Whites



Blacks

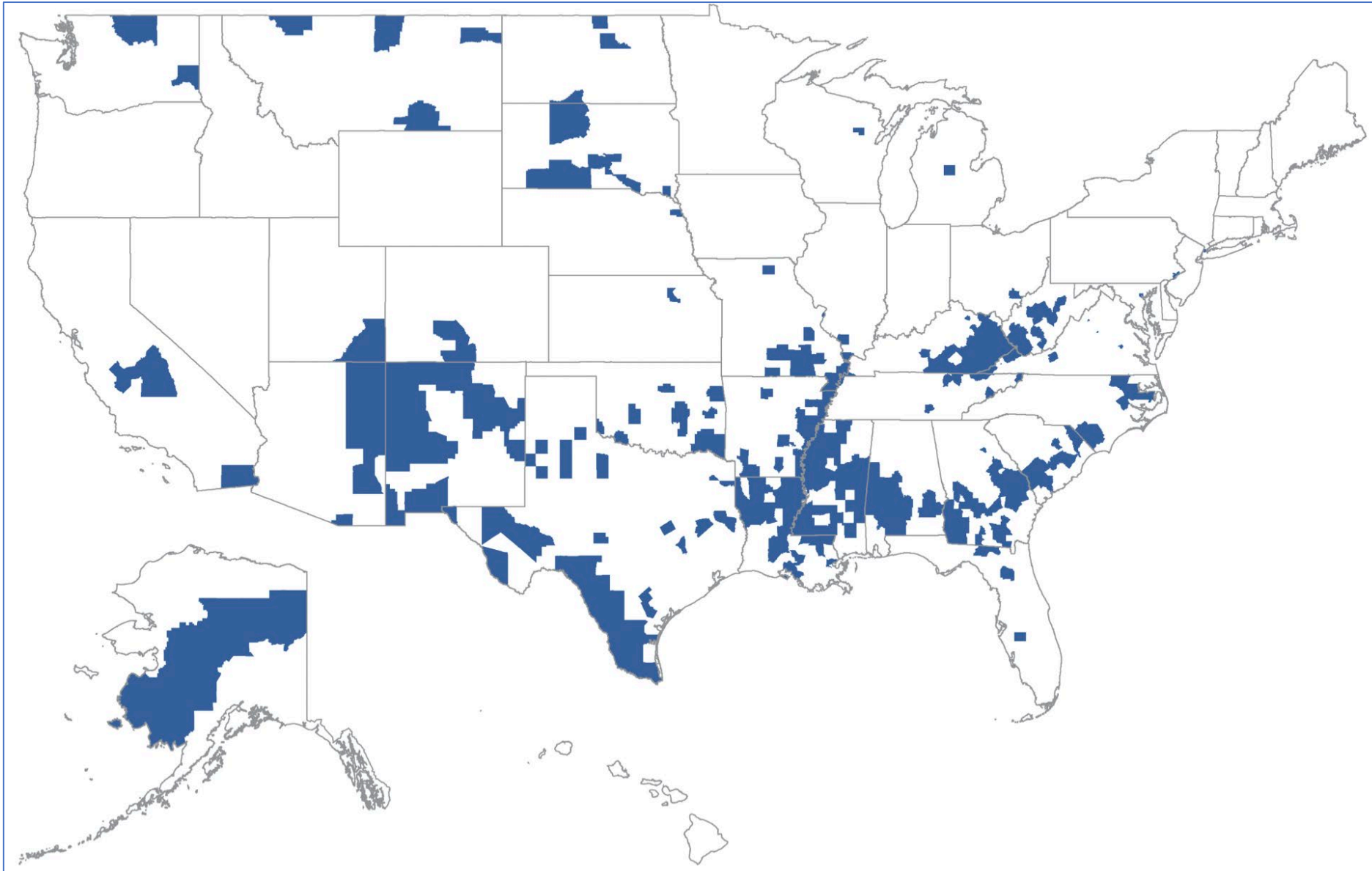
100 counties in the US with **highest** AD/DRD prevalence by race/ethnicity



Hispanic/LatinX

Race represents a social rather than a biological construct.

Persistent-Poverty Counties, as of 2017



Source: GAO analysis of Census and Small Area Income and Poverty Estimates data. | GAO-20-518

Using Population Descriptors in Genetics and Genomics Research

A New Framework for an Evolving Field

Conclusion 4-1. Race is neither useful nor scientifically valid as a measure of the structure of human genetic variation.

Conclusion 4-2. Using socially constructed groupings indiscriminately in human genetics research can be harmful.

Conclusion 4-3. Current practices often reinforce typological views of human genetic ancestry (e.g., the use of continental ancestry groups). Therefore, new models that reflect a more complex and realistic portrait of genetic ancestry are needed (e.g., genetic similarity).

Conclusion 4-4. The requirement to report participant demographics using OMB categories has perpetuated misconceptions or exacerbated typological thinking and can undermine the selection of variables that are most appropriate for a given study.

Are ATN-defined biomarkers equally prevalent among diverse populations?

Table 2. Amyloid Positivity Differences Between 1:1 Matched Participants

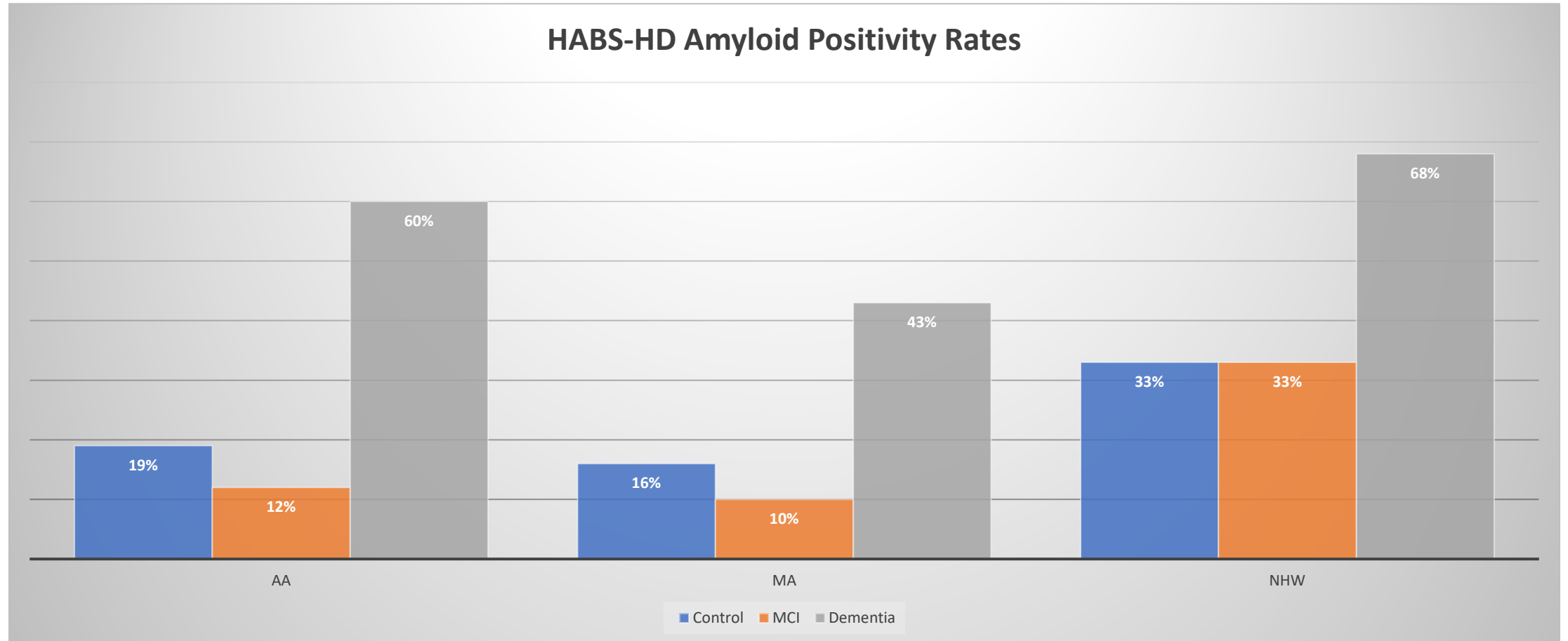
Amyloid PET scan result	Matched participants, No. (%)					
	Asian	White	Black	White	Hispanic	White
No.	313	313	615	615	780	780
MCI and dementia						
Positive, No. (%) [95% CI]	142 (45.4) [39.9-50.9]	181 (57.8) [52.3-63.2]	333 (54.1) [50.2-58.0]	359 (58.4) [54.4-62.2]	425 (54.5) [51.0-58.0]	482 (61.8) [58.3-65.1]
Negative	171 (54.6)	132 (42.2)	282 (45.9)	256 (41.6)	355 (45.5)	298 (38.2)

Biomarker	Study by	NHB vs NHW
Amyloid-PET	Gottesman et al, 2016 Morris et al., 2019 Meeker et al., 2020 Deters et al., 2021	NHB Higher No difference No difference NHB Lower
CSF AB42 & AB40	Howell et al., 2017 Garrett et al 2019 Morris et al., 2019 Deters et al., 2021	No difference NHB higher if HC NHB higher if MCI
PET-tau	Meeker et al., 2020	No difference
CSF-tau	Howell et al., 2017 Garrett et al 2019 Morris et al., 2019 Deters et al., 2021	NHB Lower

Wilkins et al., Racial and Ethnic Differences in Amyloid PET Positivity in Individuals With Mild Cognitive Impairment or Dementia *JAMA Neurol.* 2022;79(11):1139-1147.
doi:10.1001/jamaneurol.2022.3157

Gleason, CE, Zuelsdorff, M, Gooding, DC, et al. Alzheimer's disease biomarkers in Black and non-hispanic White cohorts: A contextualized review of the evidence. *Alzheimer's Dement.* 2022; 18: 1545–1564. <https://doi.org/10.1002/alz.12511>

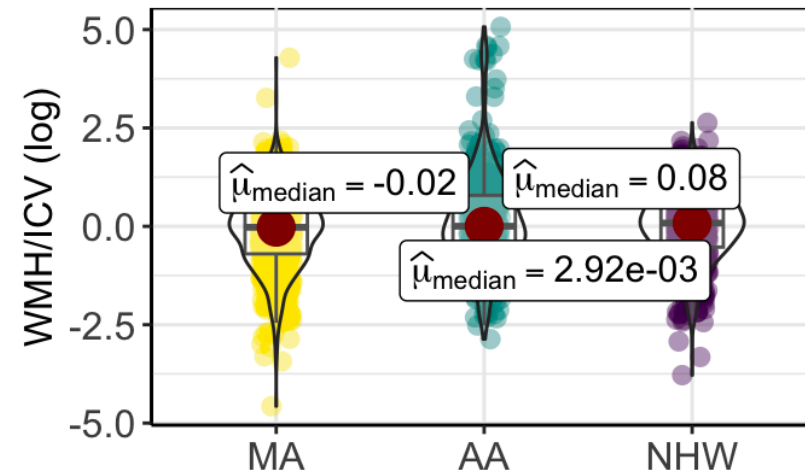
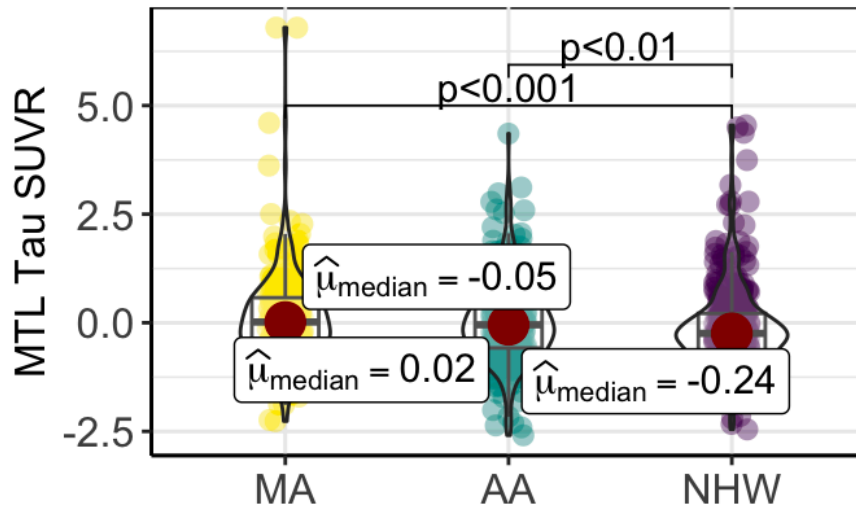
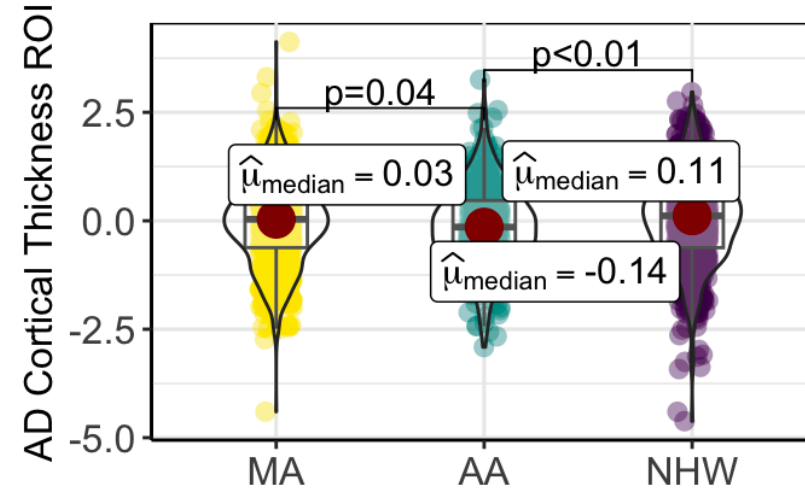
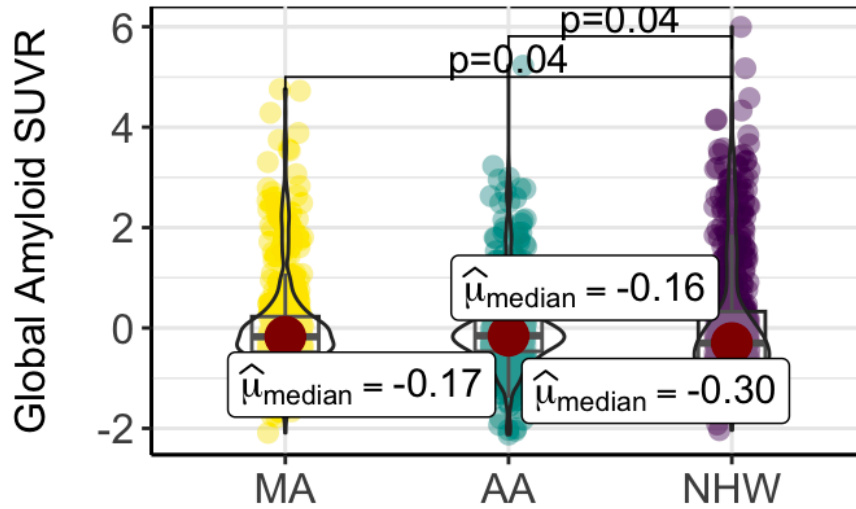
HABS-HD (formally the Health & Aging Brain study among Latino Elders, HABLE study)



Mexican American participants had significantly lower $A\beta_{40}$ ($P < .001$) and higher total tau ($P = .005$) and $A\beta_{42}/A\beta_{40}$ ratio ($P < .001$) levels.

A, T and N, but not V differ by group (With CDR)

MA = 1171
AA = 700
NHW = 1164



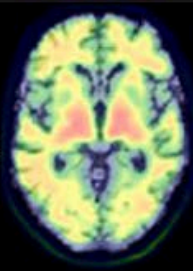
*Values = residuals after accounting for age, gender, education, and time difference between MRI and PET

Are ATN-defined biomarkers equally prevalent among diverse populations? **No**

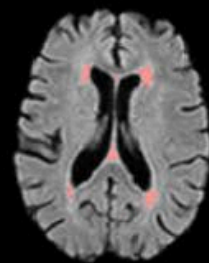
Are ATN-defined biomarkers equally related to clinical outcomes among diverse populations?

Amyloid

Amyloid PET

White Matter
Hyperintensity

Binary Label

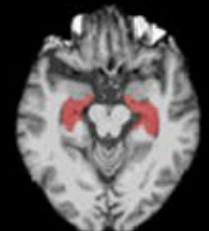


Infarct

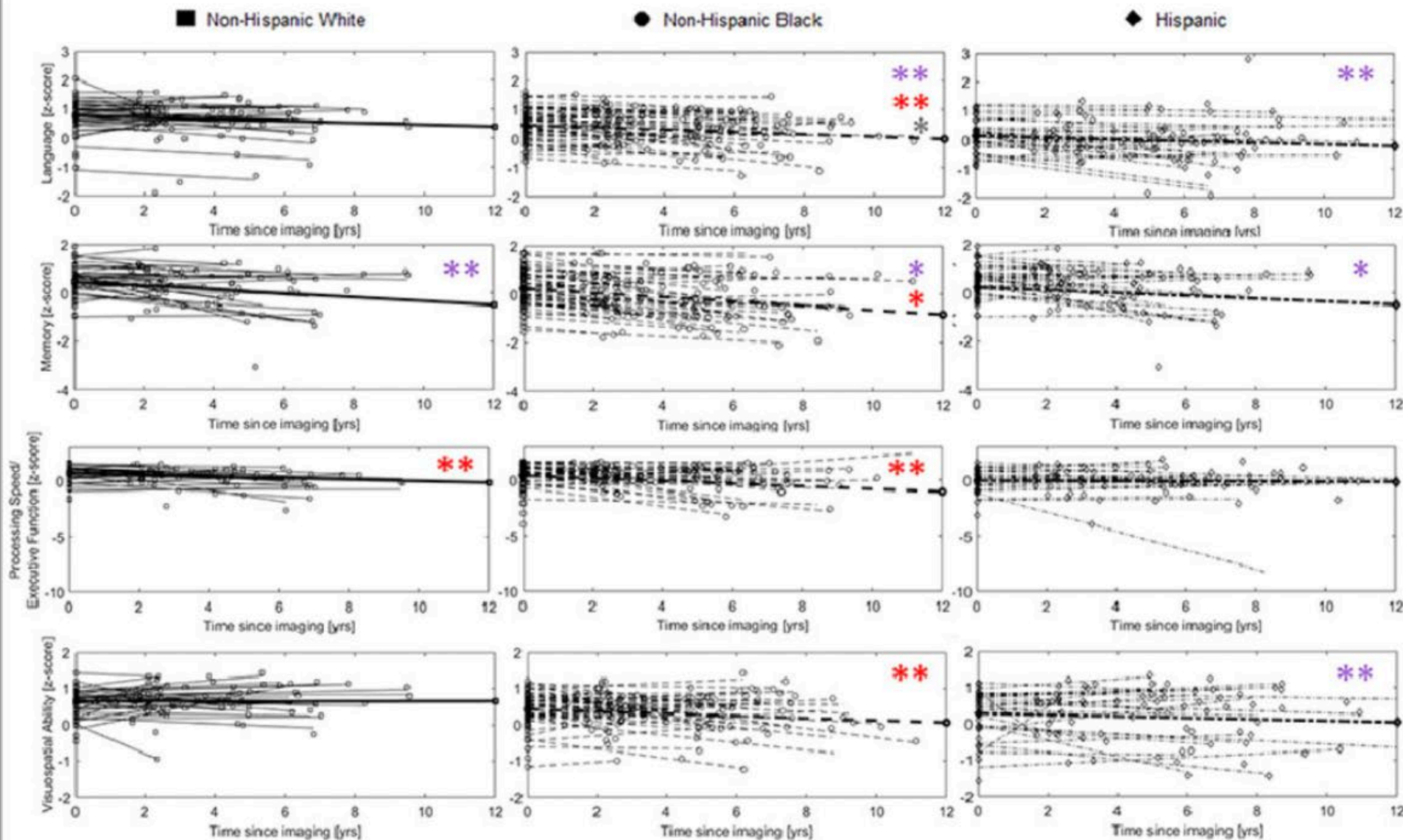
Visual Read

Cortical
ThicknessHippocampal
Volume

Binary Label



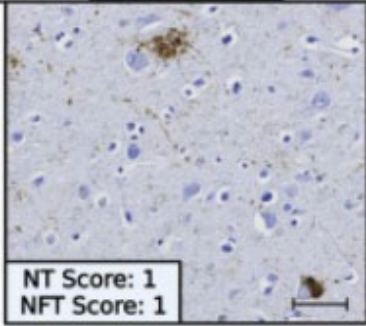
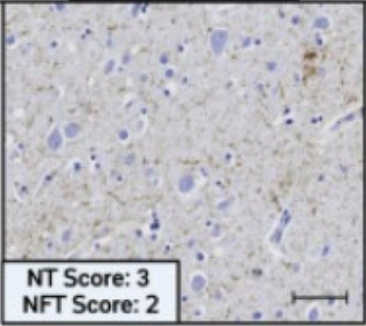
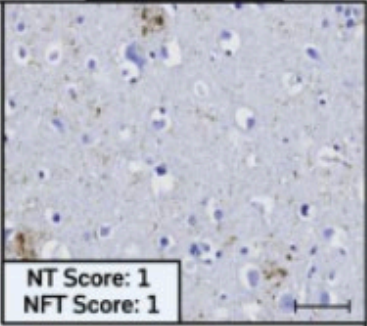
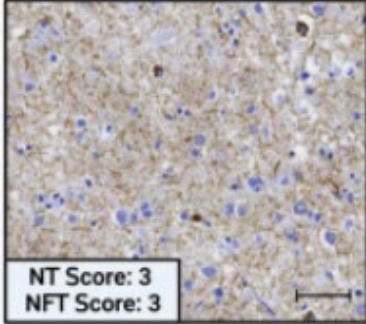
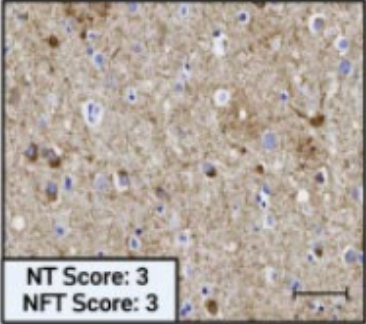
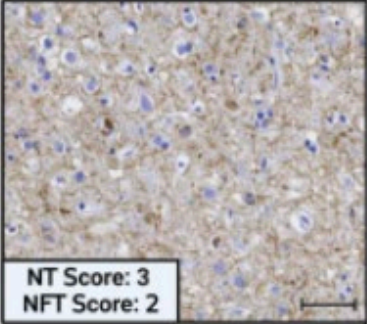
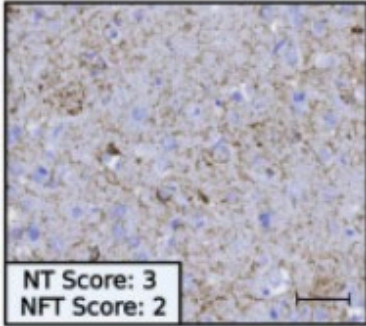
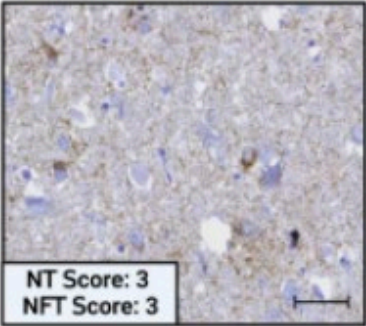
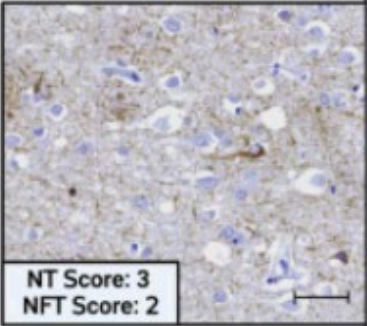
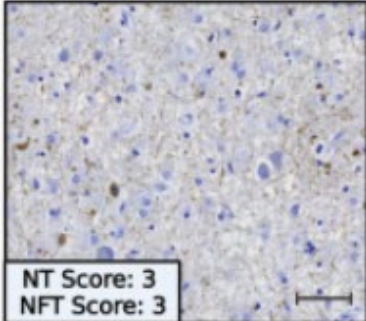
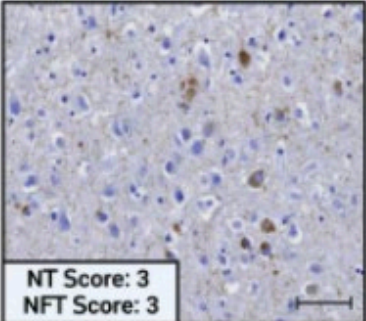
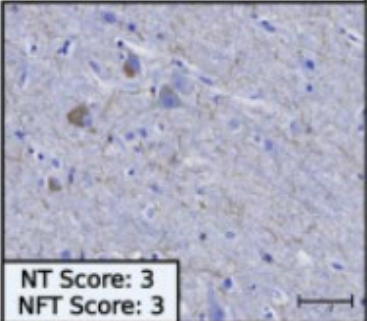
(Neurodegeneration)



Following the 2018 NIA-AA Research Framework for Alzheimer's disease, incorporation of biomarkers for amyloid, vascular disease, and neurodegeneration shows associations among baseline biomarker levels and subsequent cognitive decline over time in three major racial/ethnic groups. ** $p < 0.05$ and * $p < 0.10$, color coded by biomarker type.

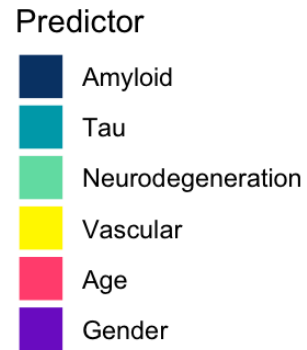
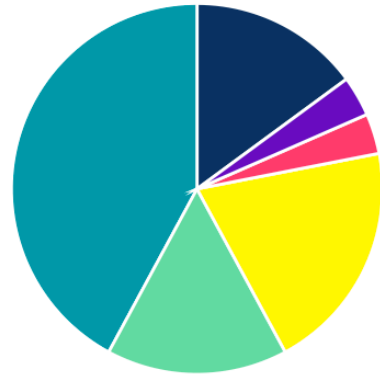
Examples of the histopathologic densities of tau-deposits and corresponding overall regional density scores (for neuropil threads: NTs, and neurofibrillary tangles: NFTs- lower left corner in each image) in three brain regions (frontal, temporal, and parietal cortices).

Scalco, R., Saito, N., Beckett, L. *et al.* The neuropathological landscape of Hispanic and non-Hispanic White decedents with Alzheimer disease. *acta neuropathol commun* **11**, 105 (2023).
<https://doi.org/10.1186/s40478-023-01574-1>

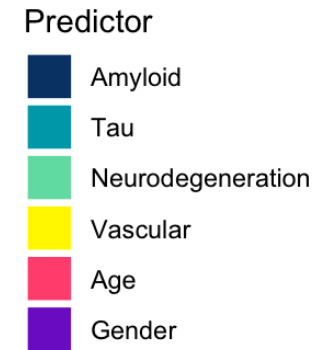
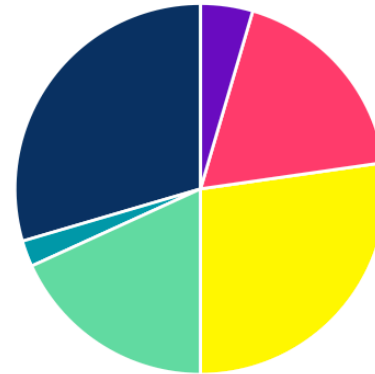
		FRONTAL	TEMPORAL	PARIETAL
NHW	Age-at-death: 70 Female ADNC: High likelihood	 NT Score: 1 NFT Score: 1	 NT Score: 3 NFT Score: 2	 NT Score: 1 NFT Score: 1
CARIBBEAN	Age-at-death: 70 Female ADNC: High likelihood	 NT Score: 3 NFT Score: 3	 NT Score: 3 NFT Score: 3	 NT Score: 3 NFT Score: 2
MEXICAN	Age-at-death: 70 Female ADNC: High likelihood	 NT Score: 3 NFT Score: 2	 NT Score: 3 NFT Score: 3	 NT Score: 3 NFT Score: 2
OTHER	Age-at-death: 76 Female ADNC: High likelihood	 NT Score: 3 NFT Score: 3	 NT Score: 3 NFT Score: 3	 NT Score: 3 NFT Score: 3

Importance of AT(N)-V imaging markers differs by group when predicting cognitive impairment

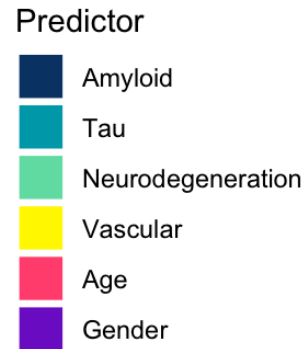
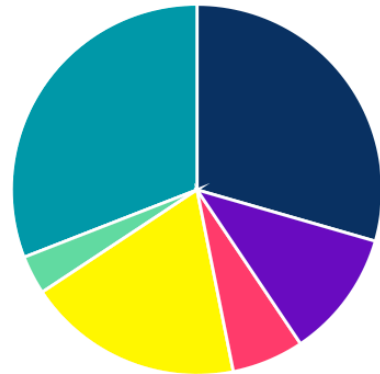
A: All



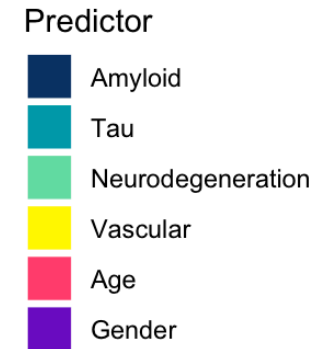
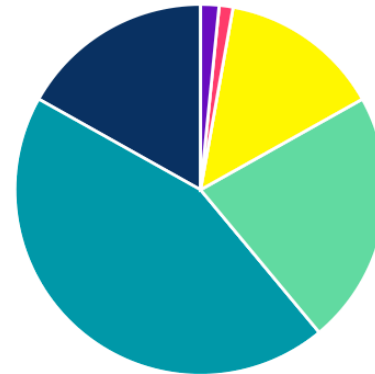
B: MA



C: AA



D: NHW



Are ATN-defined biomarkers equally prevalent among diverse populations? **No**

Are ATN-defined biomarkers equally related to clinical outcomes among diverse populations? **No**

Are modifiers of ATN-defined biomarkers the same across diverse populations? **■**

Standardized Beta Results From Regression Models Examining Associations Between Demographic and Cardiovascular Risk Factors With Serum GFAP Stratified by Ethnicity

Variables	Overall		Cognitively unimpaired		MCI		Dementia	
	MA	NHW	MA	NHW	MA	NHW	MA	NHW
Age	0.604‡	0.392‡	0.483‡	0.469‡	0.639‡	0.479‡	0.306‡	0.298‡
Women	-0.184‡	-0.132‡	-0.198‡	-0.133	-0.164‡	-0.116	-0.172*	-0.113*
BMI	-0.184‡	-0.132‡	-0.198‡	-0.133	-0.164‡	-0.116	-0.172*	-0.036*
SBP	0.037	-0.028	0.079*	-0.029	-0.038	-0.157*	0.134	-0.017
Diabetes	-0.023	-0.160‡	-0.085*	-0.134*	-0.124‡	-0.211‡	-0.075	-0.190‡
Tabaco use	-0.064‡	-0.029	-0.124‡	0.027	-0.003	-0.085	-0.032	-0.039
APOE e-4	0.108‡	0.151‡	0.037	0.146*	0.156‡	0.173*	0.186*	0.024
Cognitive impairment	0-0.88‡	0.237‡						

APOE = apolipoprotein E; MA = Mexican American; NHW = non-Hispanic White. Standardized beta values derived from regression models examining associations with serum GFAP in the whole sample and by the diagnostic group with stratification for ethnicity.
 *P<0.05; †P<0.01; ‡P<0.001

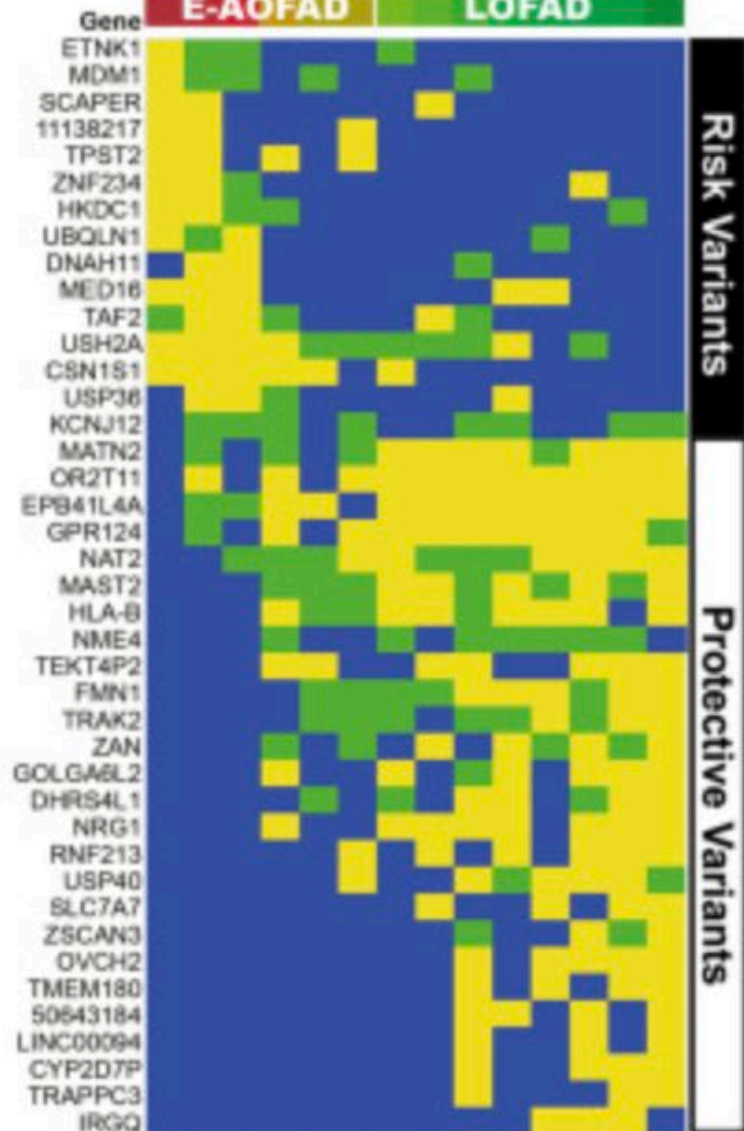
a

Age of Onset in years

37 39 40 48 49 49 52 53 54 55 56 58 58 62

E-AOFAD

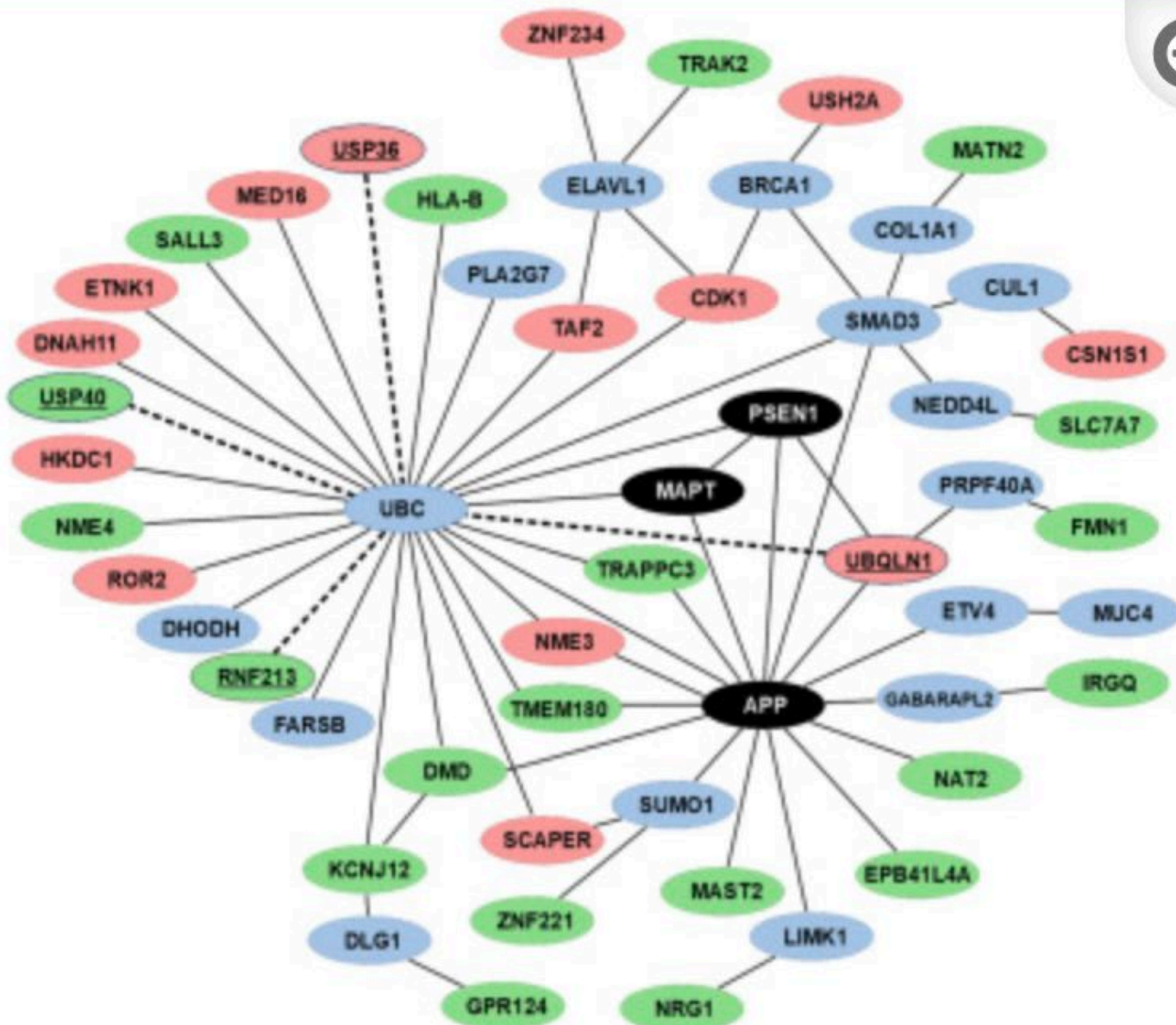
LOFAD



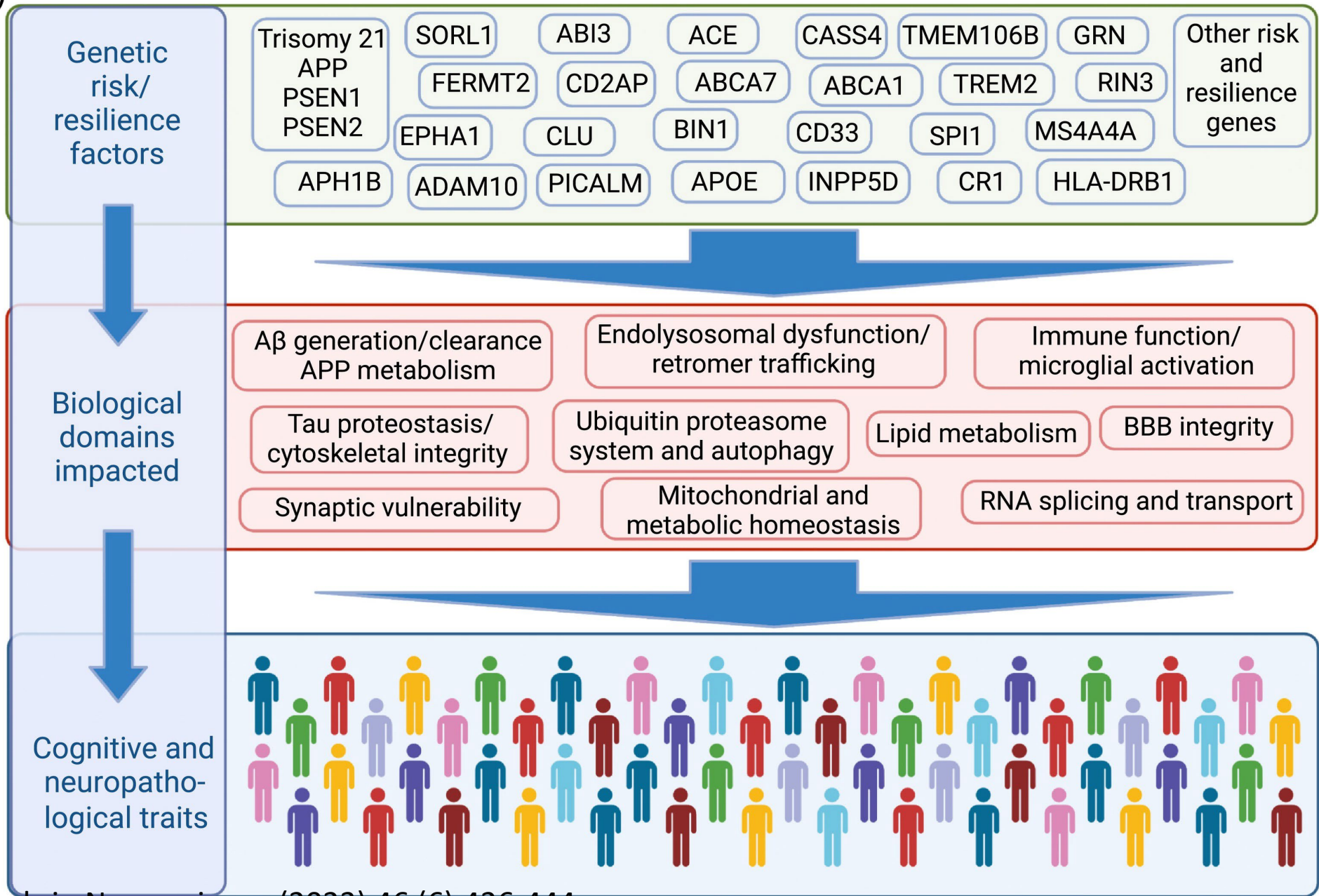
Risk Variants

Protective Variants

b



(A)

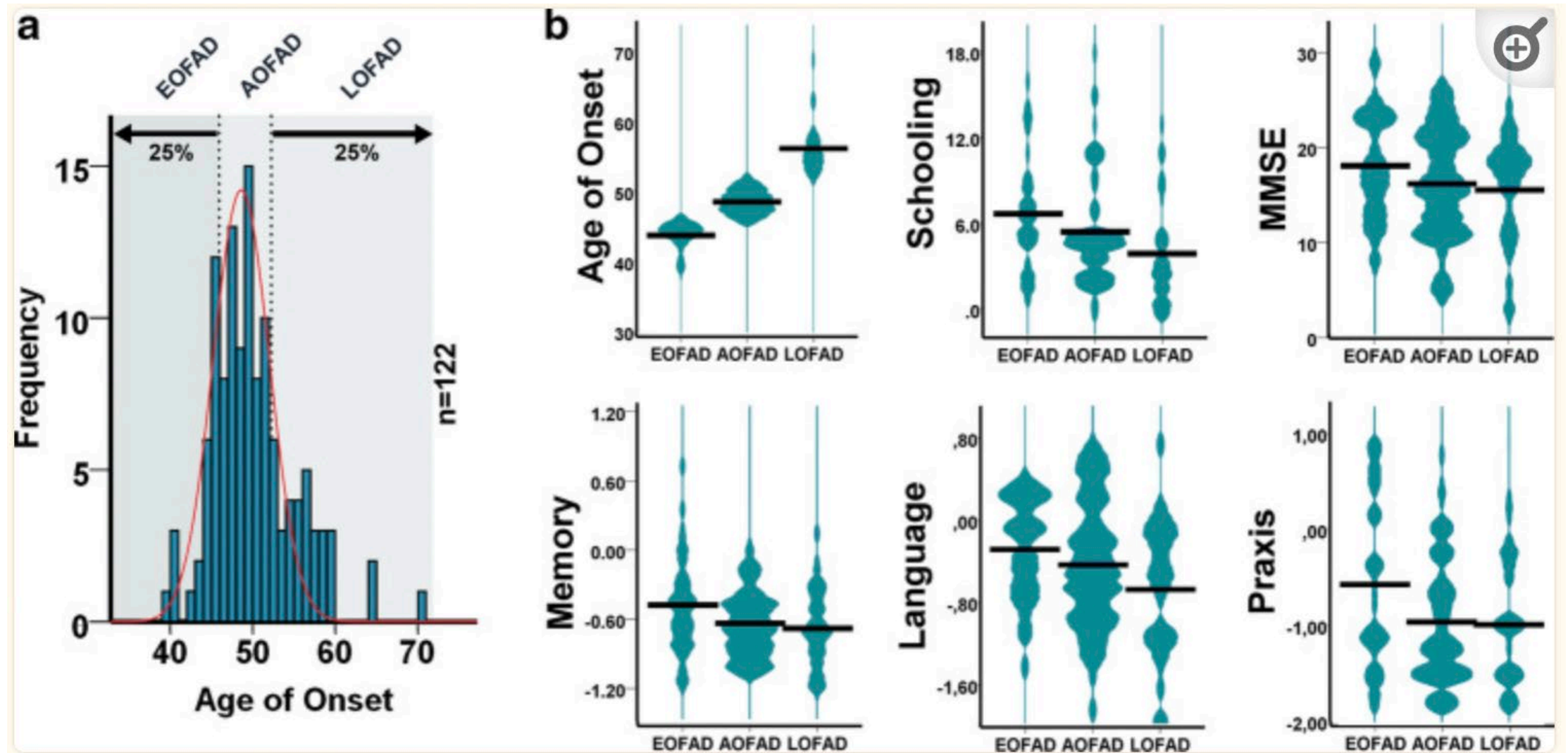


Are ATN-defined biomarkers equally prevalent among diverse populations? **No**

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Are modifiers of ATN-defined biomarkers the same across diverse populations? **No** ■

Age of onset in PSEN 1 E280A dementia patients in Colombia



Falla-Sepulveda D, et al, A multifactorial model of pathology for age of onset heterogeneity in familial Alzheimer's disease. *Acta Neuropathol.* 2021

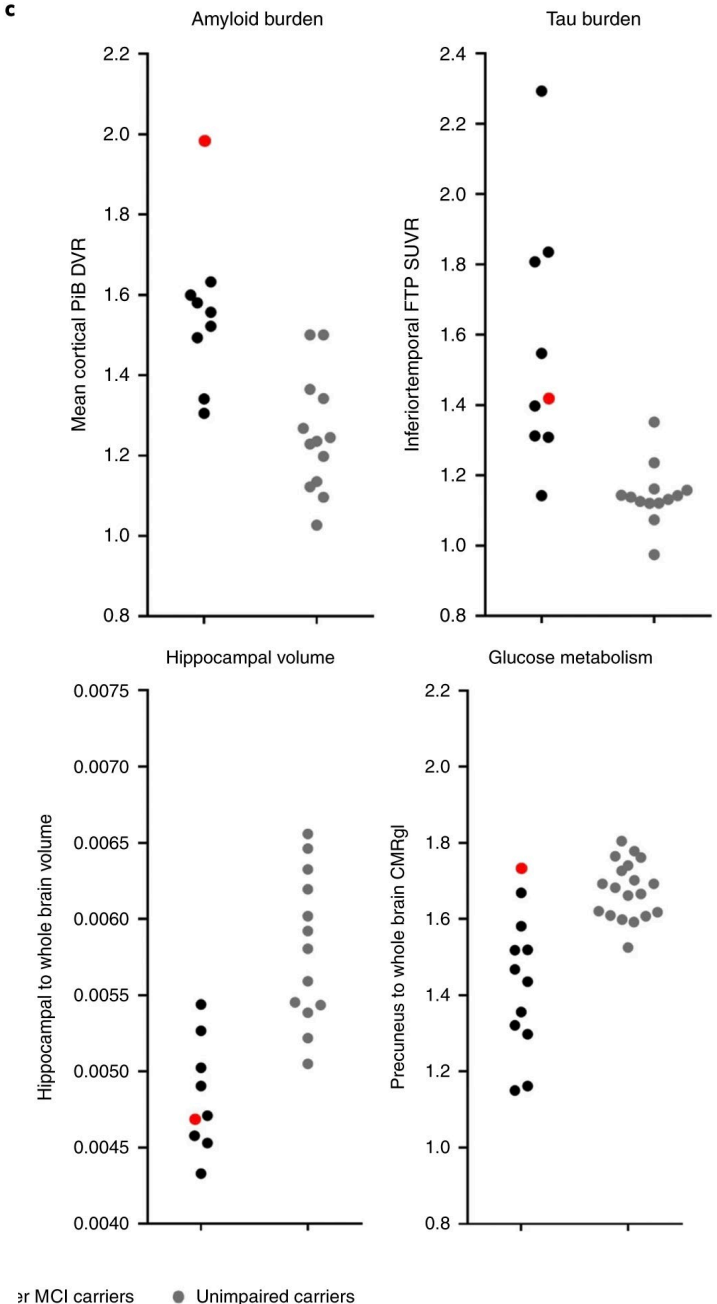
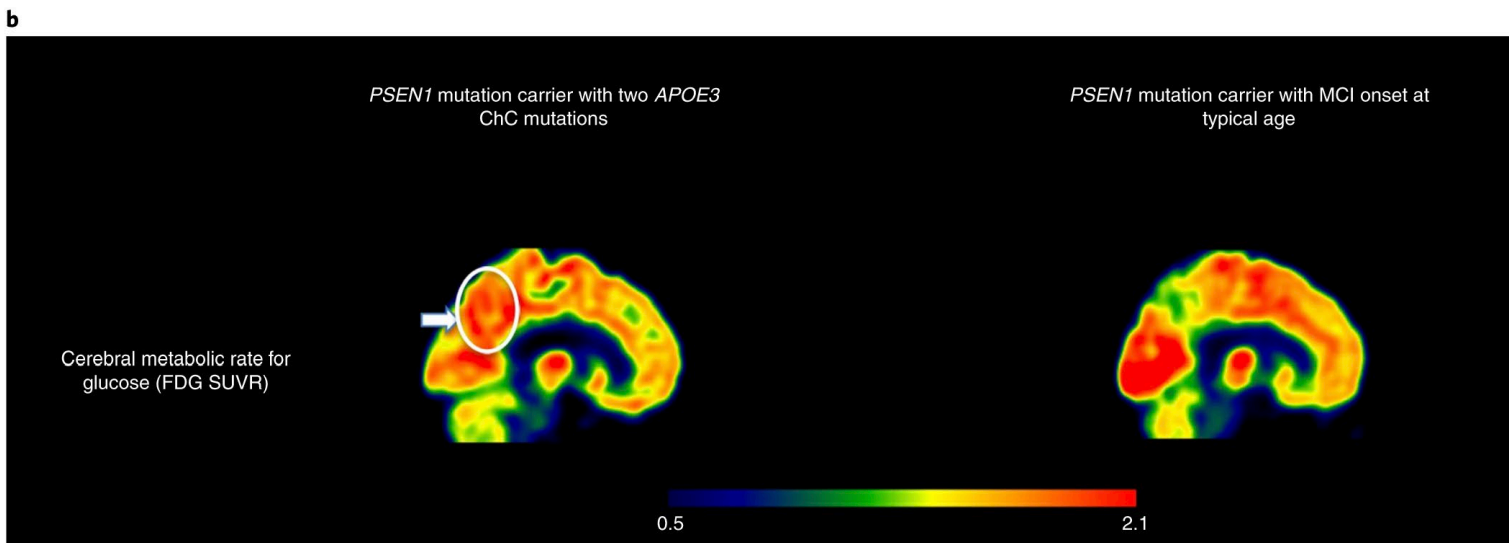
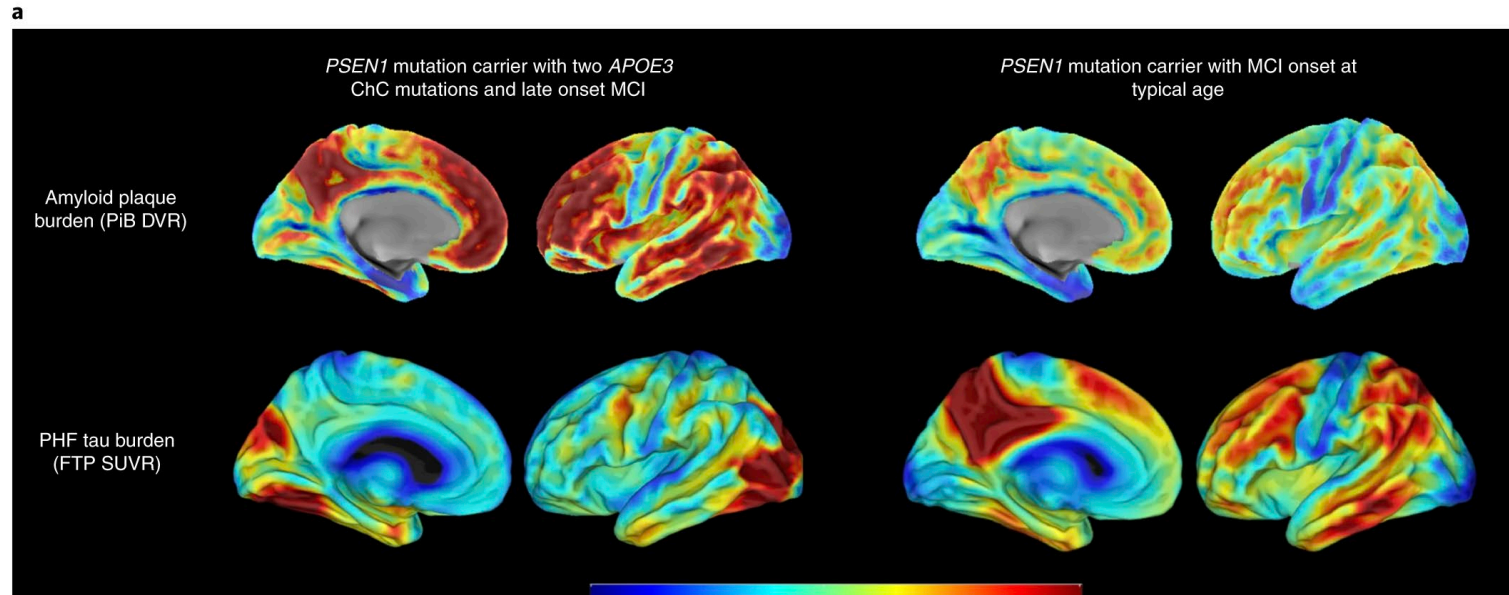
Colombian woman with PS1 E280A mutation without cognitive impairment for 73 years. She died with MCI from melanoma at 78y.



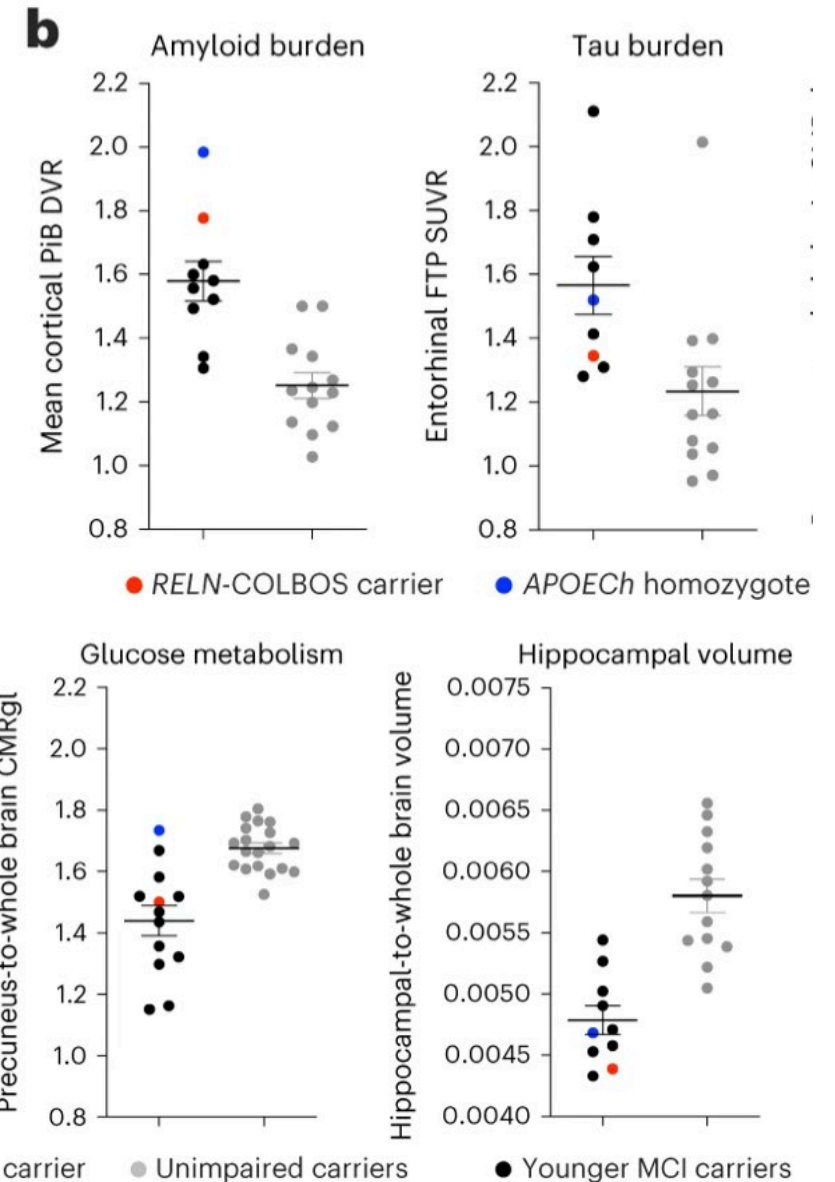
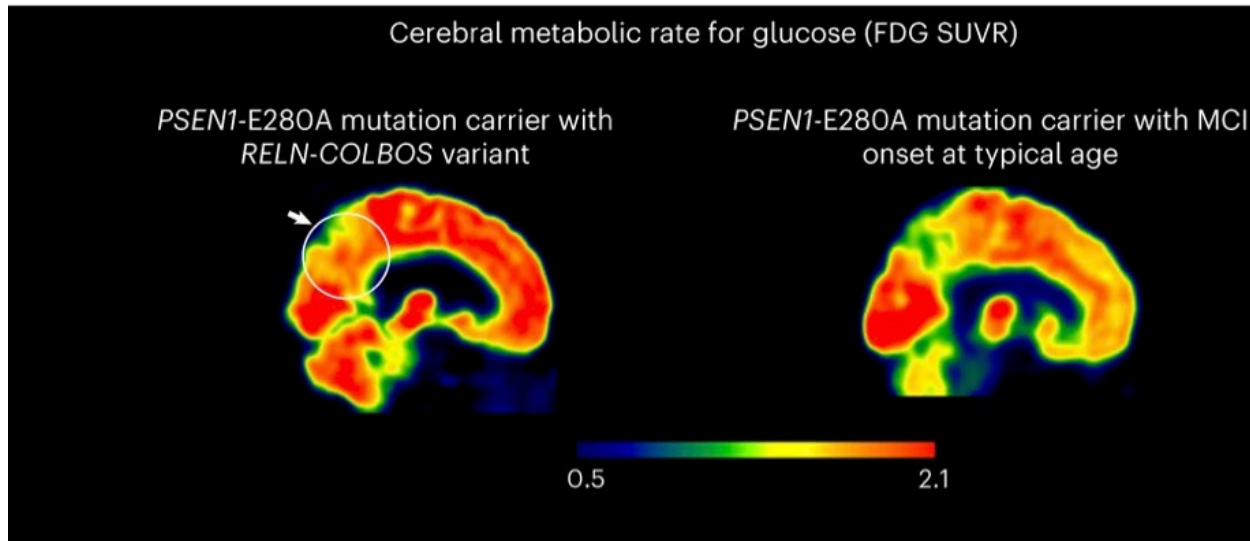
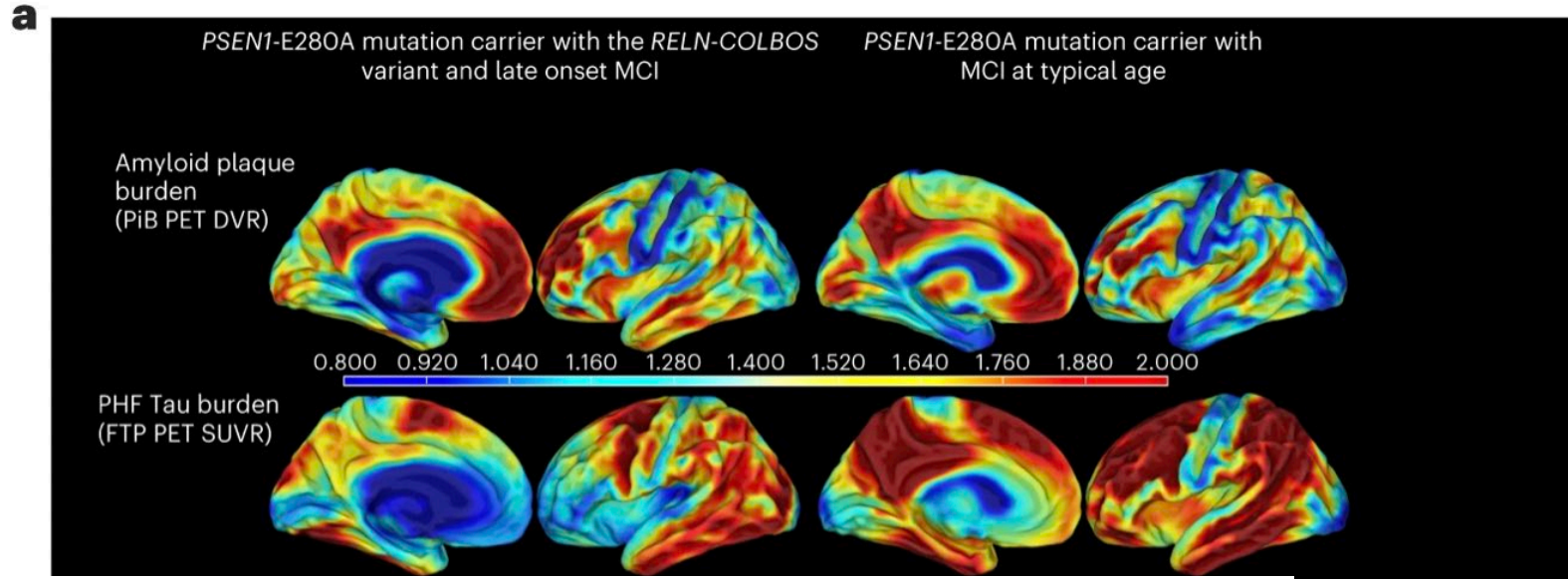
In Life, She Defied Alzheimer's. In Death, Her Brain May Show How.

A woman in Colombia with a rare genetic mutation recently made the ultimate donation to science.

Arboleda-Velasquez, J.F., Lopera, F., O'Hare, M. et al. **Resistance** to autosomal dominant Alzheimer's disease in an APOE3 Christchurch hon^c
 Nat Med 25, 1680–1683 (2019). <https://doi.org/10.1038/s41591-019-0541-1>



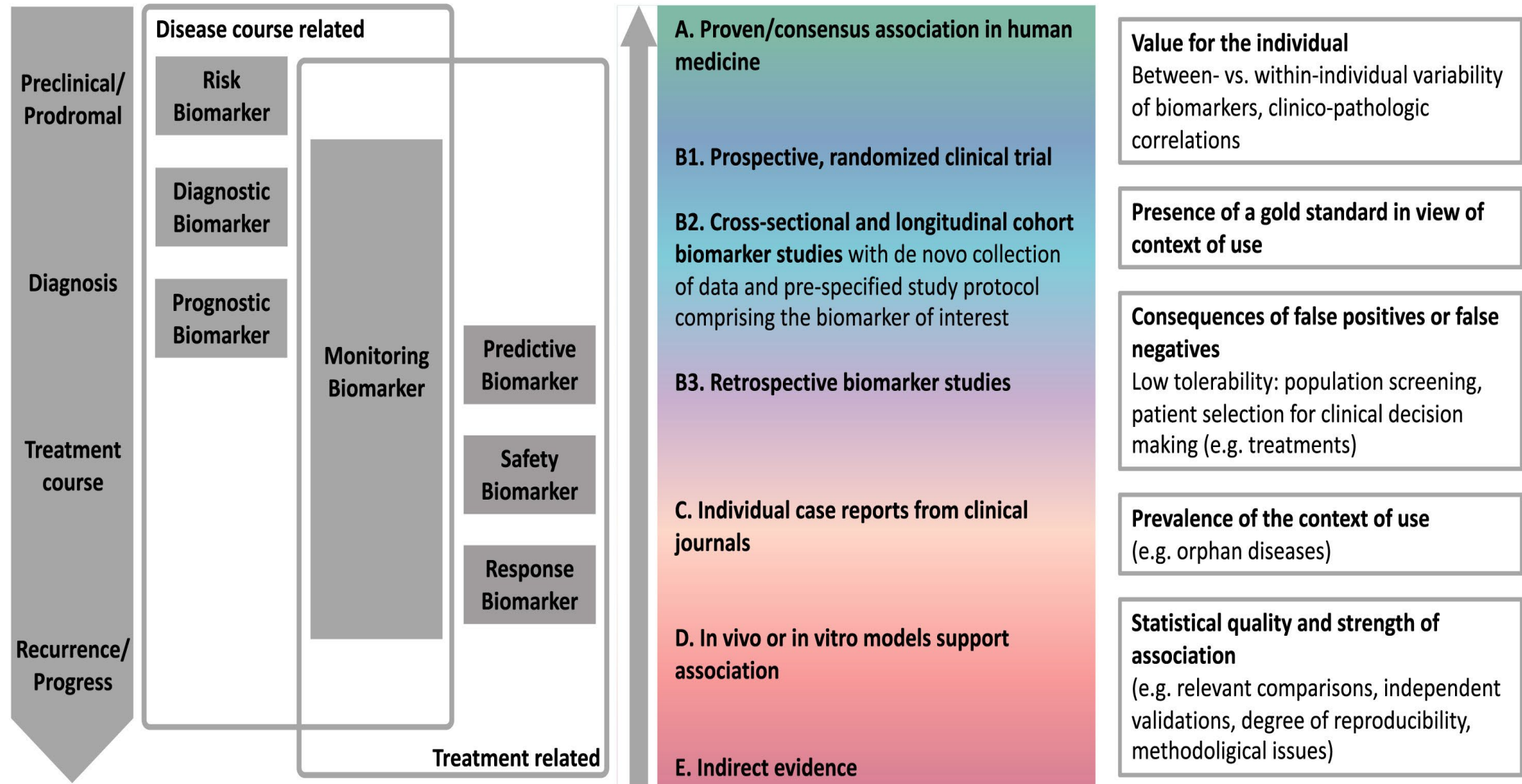
Lopera, F., Marino, C., Chandrabhas, A.S. et al. **Resilience** to autosomal dominant Alzheimer's disease in a Reelin-COLBOS heterozygous man. *Nat Med* 29, 1243–1252 (2023). <https://doi.org/10.1038/s41591-023-02318-3>



I. Defining context of use in relation to clinical endpoint

II. Defining evidence levels

III. Rating clinical utility



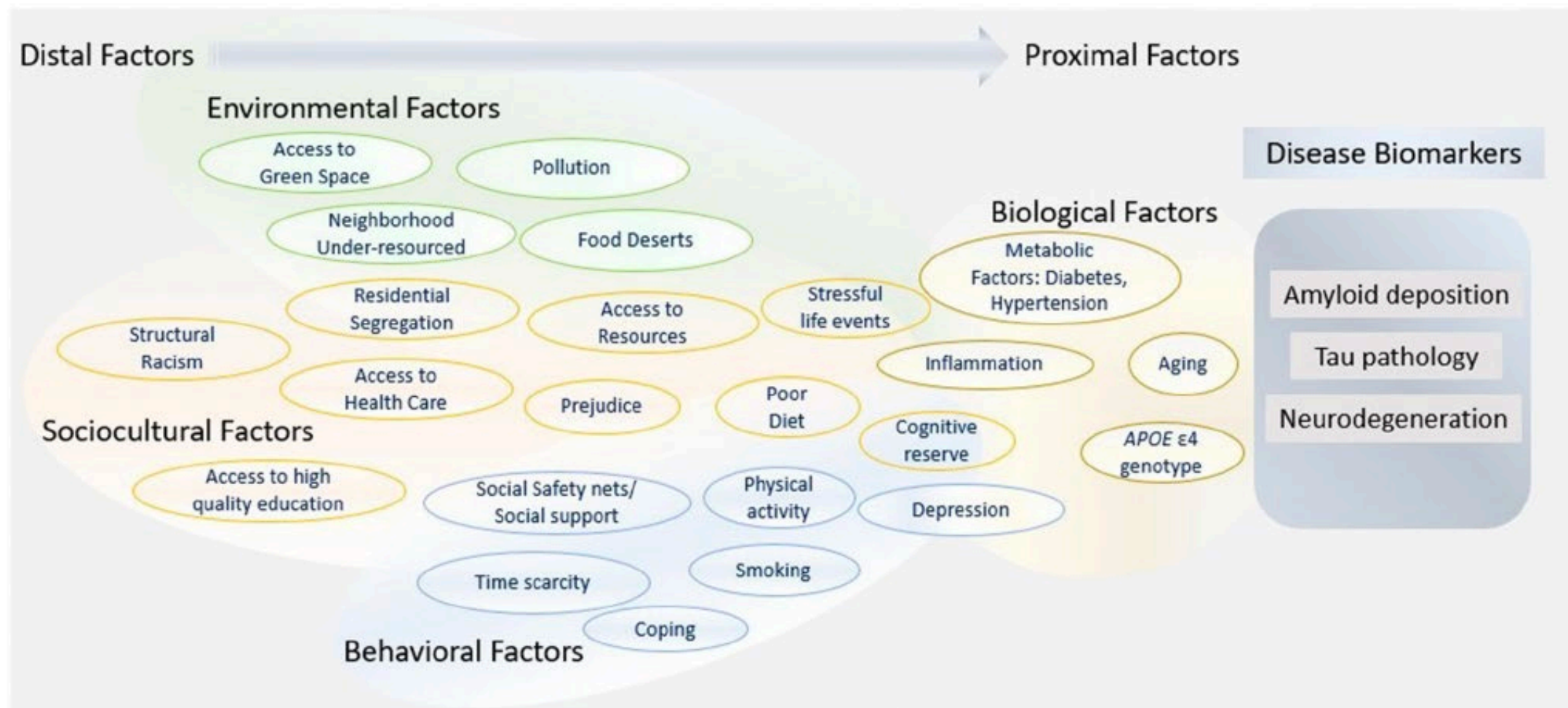
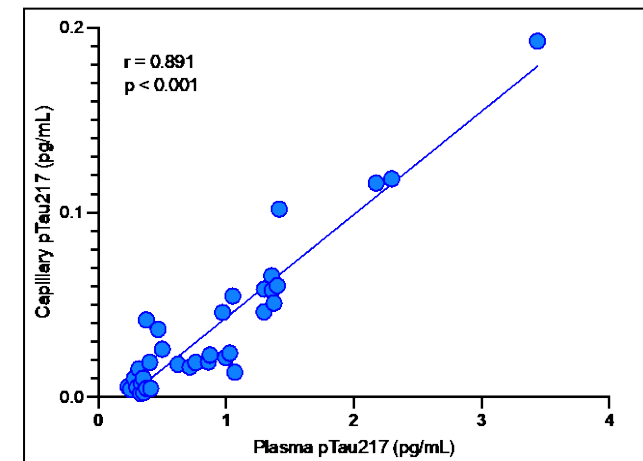
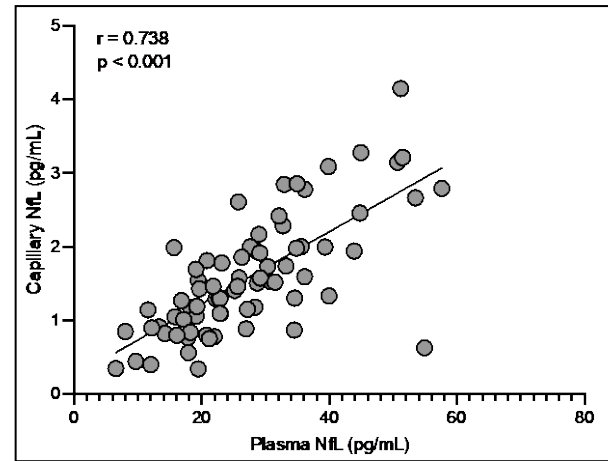
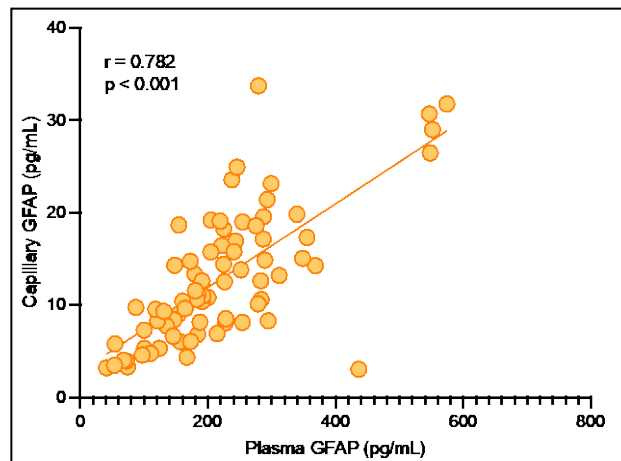
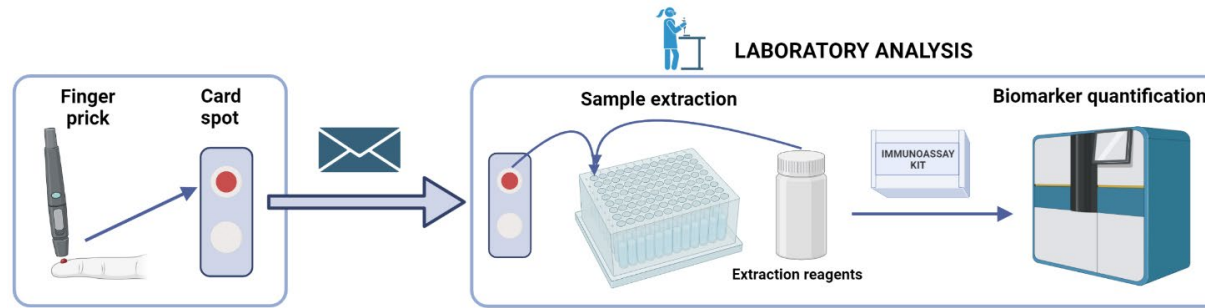


FIGURE 1 Hypothetical model integrating the National Institute on Aging (NIA) Health Disparities Research Framework with the amyloid/tau/neurodegeneration (AT[N]) criteria from the NIA–Alzheimer's Association Research Framework. Evidence supporting individual examples association with Alzheimer's disease biomarkers is hypothetical in some instances. Examples are generally categorized as environmental, sociocultural, behavioral, or biological factors. However, we acknowledge that many examples can belong in multiple categories

DROP-AD: detecting AD blood biomarkers using a finger-prick

- Current blood processing protocols require strict procedures – useful in primary care?
- How do we monitor people overtime (including those on DMT) for personalised management?
- Detecting pre-clinical changes – if/when that it is required?



1.- Align communication to meet the education and support needs that may be unique to each patient and family

2.- Clarify context of use before testing

3.-Pretest-counseling and post-disclosure support

4.-Cautious selection and usage of lay language

5.-Comprehension checks

<https://www.agreedementia.org>

6.-Avoid positive/negative qualifiers



Thank you!

¡Gracias!

