

Genetic Susceptibility of Alzheimer's Disease in African Americans

Risk and Resilience to Alzheimer's Disease in African Americans Conference

Rutgers University

June 20, 2023

Goldie S. Byrd PhD



Presentation

Genetics of AD

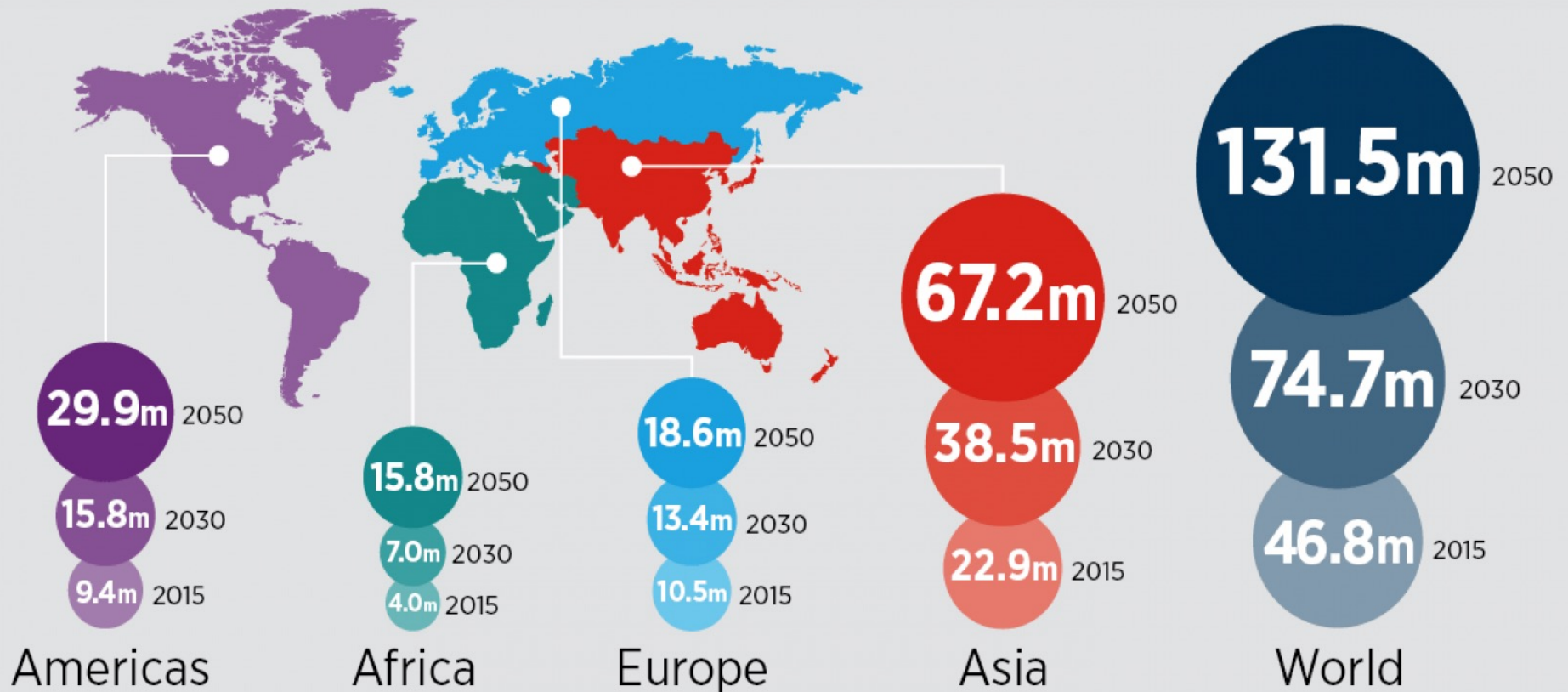
Why Study
AD Genetics
Research in
African Americans

Global and Local
Ancestry

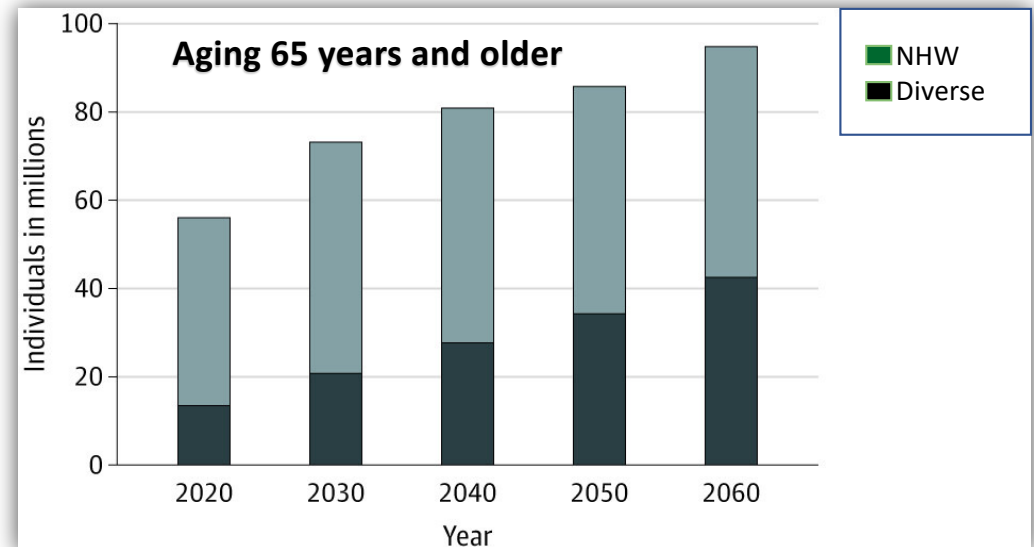
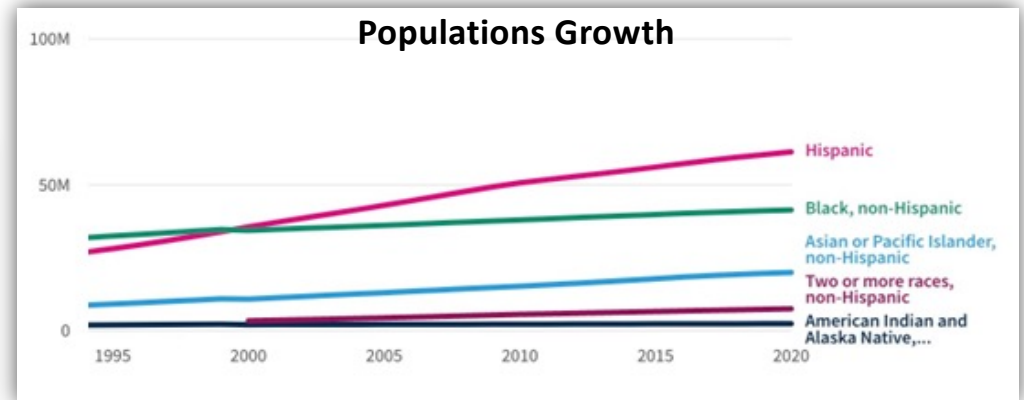
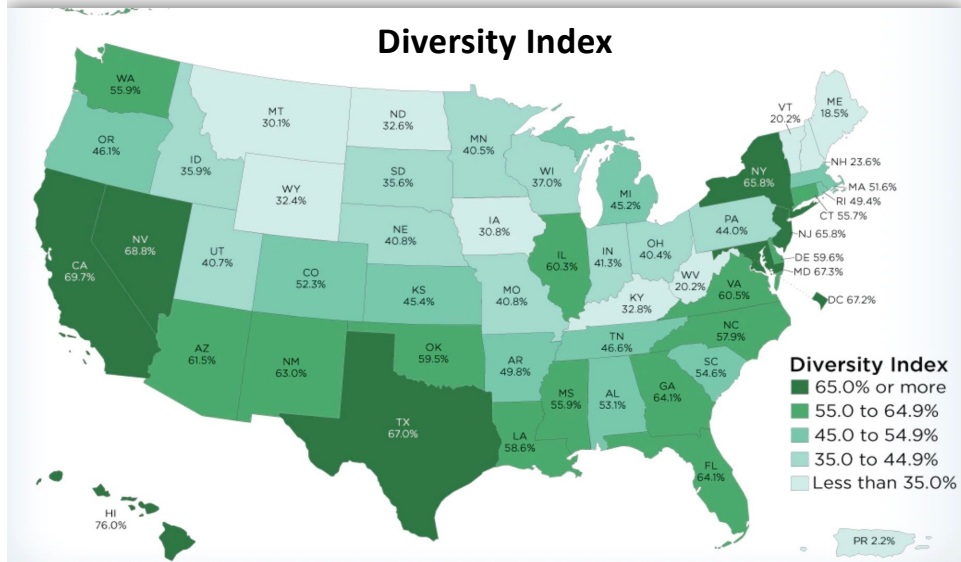
Inclusion
Barriers and Solutions
for Genetics Research
in African Americans

Alzheimer's Disease

People living with dementia around the world



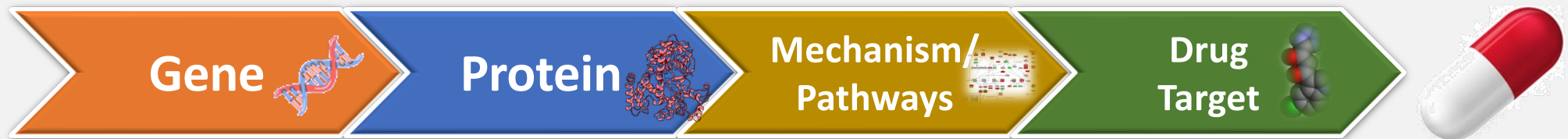
US Populations Diversity, Growth and Aging



2020 US Census
Kawas CH et al. JAMA 2021

Human Genetics Disease Research Goals

- Study human disease mechanism directly in humans
- Prediction
- Mechanism
- Drug targets



Genetic Studies in Alzheimer's Disease

Why study genes?

- To identify new targets for drug discovery
- Genetics targets are 2X as successful in drug trials versus non-genetic targets.

Why study Diverse Groups

- 25 genetic risk factors for Alzheimer's disease discovered-primarily in individuals of European **ancestral** descent
- Different **ancestral** groups have different genetic risk factors
- So that treatments are universally translational

Why Genetic Studies in Alzheimer’s Disease: Drug Discovery: Twice as Successful

ANALYSIS

nature
genetics

The support of human genetic evidence for approved drug indications

Matthew R Nelson¹, Hannah Tipney², Jeffery L Painter¹, Judong Shen¹, Paola Nicoletti³, Yufeng Shen^{3,4}, Aris Floratos^{3,4}, Pak Chung Sham^{5,6}, Mulin Jun Li^{6,7}, Junwen Wang^{6,7}, Lon R Cardon⁸, John C Whittaker² & Philippe Sanssou²

“We estimate that selecting genetically supported targets could double the success rate in clinical development.”

 PLOS | GENETICS

RESEARCH ARTICLE

Are drug targets with genetic support twice as likely to be approved? Revised estimates of the impact of genetic support for drug mechanisms on the probability of drug approval

Emily A. King¹*, J. Wade Davis, Jacob F. Degner

Department of Computational Genomics, AbbVie, North Chicago, Illinois, United States of America

Late Onset Alzheimer's was not thought to have a genetic component until...



Gene hunters. The Duke team (left to right): Warren Strittmatter, Allen Roses, Guy Salvesen, Ann Saunders, John Gilbert, Margaret Pericak-Vance, Mark Alberts, Elizabeth Corder, and Donald Schmechel.

Gene Dose of Apolipoprotein E Type 4 Allele and the Risk of Alzheimer's Disease in Late Onset Families

E. H. Corder, A. M. Saunders, W. J. Strittmatter, D. E. Schmechel, P. C. Gaskell, G. W. Small, A. D. Roses, J. L. Haines, M. A. Pericak-Vance*

SCIENCE • VOL. 261 • 13 AUGUST 1993

The **discovery of *ApoE*** as a genetic risk factor in Alzheimer's Disease **revolutionized the field of complex disease genetics**

Early Findings

The 23 human chromosome pairs contain all of the 30,000 genes that code the biological blueprint for a human being. This interactive illustration highlights the chromosomes containing each of the three genes that cause familial Alzheimer's and the gene with the greatest impact on Alzheimer's risk.

Familial Alzheimer's (Early Onset) Genes

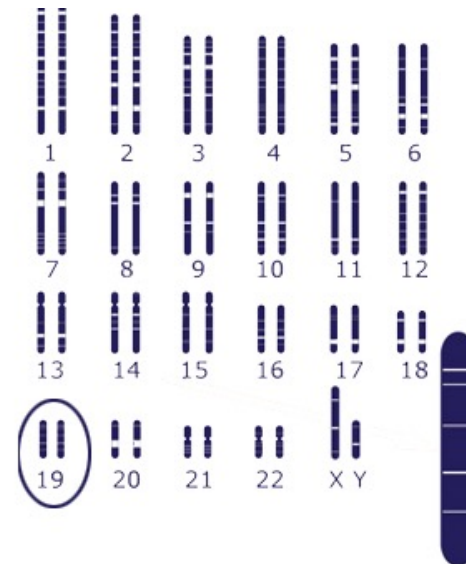
- App Chromosome 21 (1987)
- Ps-1 Chromosome 14 (1991)
- Ps-2 Chromosome 1 (1993)

Sporadic (Late Onset) AD

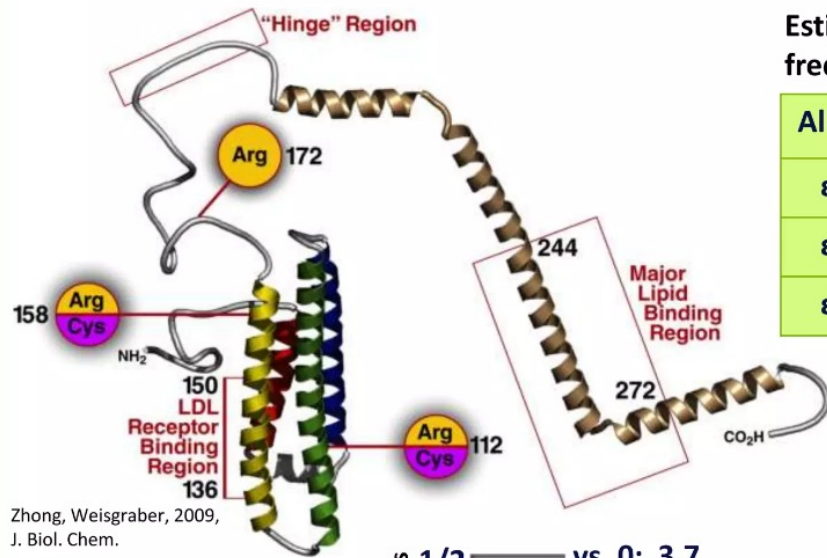
- APOE-4 1993)
- Gene with greatest known impact

23 Chromosome Pairs; 4 Alzheimer's Genes Identified:

APP PS-1 PS-2 APOE4



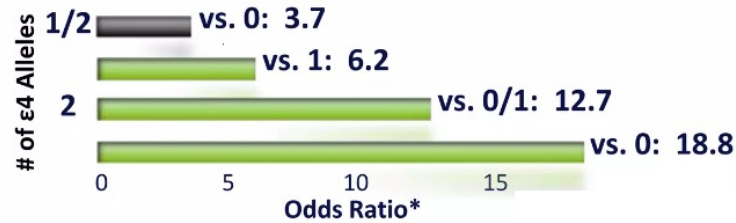
Apolipoprotein E: Genetic Risk



Estimated human genotype frequency of APOE All ethnicities
<http://www.alzgene.org>

Allele	$\epsilon 2$	$\epsilon 3$	$\epsilon 4$
$\epsilon 2$	~1%	~11%	~2%
$\epsilon 3$		~62%	~22%
$\epsilon 4$			~2%

AD: ~43% ~15%



Martins et al., 1995 NeuroReport
 Laws et al., 1999 NeuroReport

Why Study Alzheimer's Disease in African Americans

- **African Americans and Hispanic individuals are more likely to develop AD and dementia compared to non-Hispanic White populations**
 - Greater familial risk for AD
 - Limited health care access
 - AD patients identified at later stages
 - Poorer treatment outcomes
- **NAPA Act: Presidentially mandated during the Obama Administration with continued bi-partisan congressional support to identify new drug targets**



<http://www.socialgradient.org/alzheimers-center-aids-african-americans/>

<http://www.poststat.net/pwp008/pub.49/issue.350/article.528/>

REPORTS

Gene dose of apolipoprotein E type 4
risk of Alzheimer's disease in late on

EH Corder, AM Saunders, WJ Strittmatter, DE Schmechel, PC Gaskell, GW Small, A

**Protective effect of apolipoprotein
E type 2 allele for late onset
Alzheimer disease**

E. H. Corder¹, A. M. Saunders¹, N. J. Risch¹, W. J. Strittmatter^{2,3}, D. E. Schmechel^{1,2,4},
P. C. Gaskell, Jr.¹, J. B. Rimmeler¹, P. A. Locke¹, P. M. Conneally⁴, K. E. Schmechel^{2,3},
G. W. Small⁵, A. D. Roses^{1,2}, J. L. Haines³ & M. A. Pericak-Vance¹

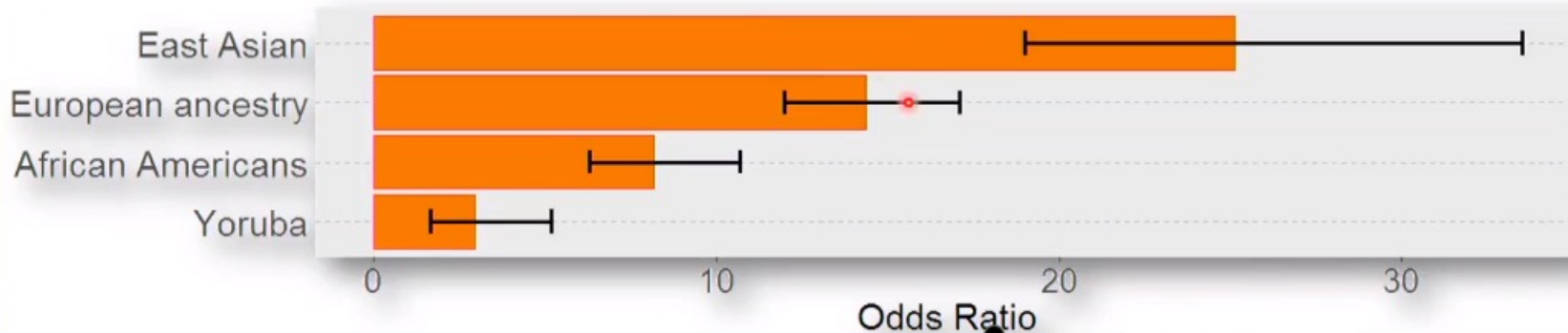
Reviews

**Effects of Age, Sex, and Ethnicity on the
Association Between Apolipoprotein E
Genotype and Alzheimer Disease**

A Meta-analysis

Lindsay A. Farrer, PhD; L. Adrienne Cupples, PhD; Jonathan L. Haines, PhD; Bradley Hyman, MD, PhD;
Walter A. Kukull, PhD; Richard Mayeux, MD; Richard H. Myers, PhD; Margaret A. Pericak-Vance, PhD;
Neil Risch, PhD; Cornelia M. van Duijn, PhD; for the APOE and Alzheimer Disease Meta Analysis Consortium

APOE ε4 risk effect (ε3ε3 vs ε4ε4)



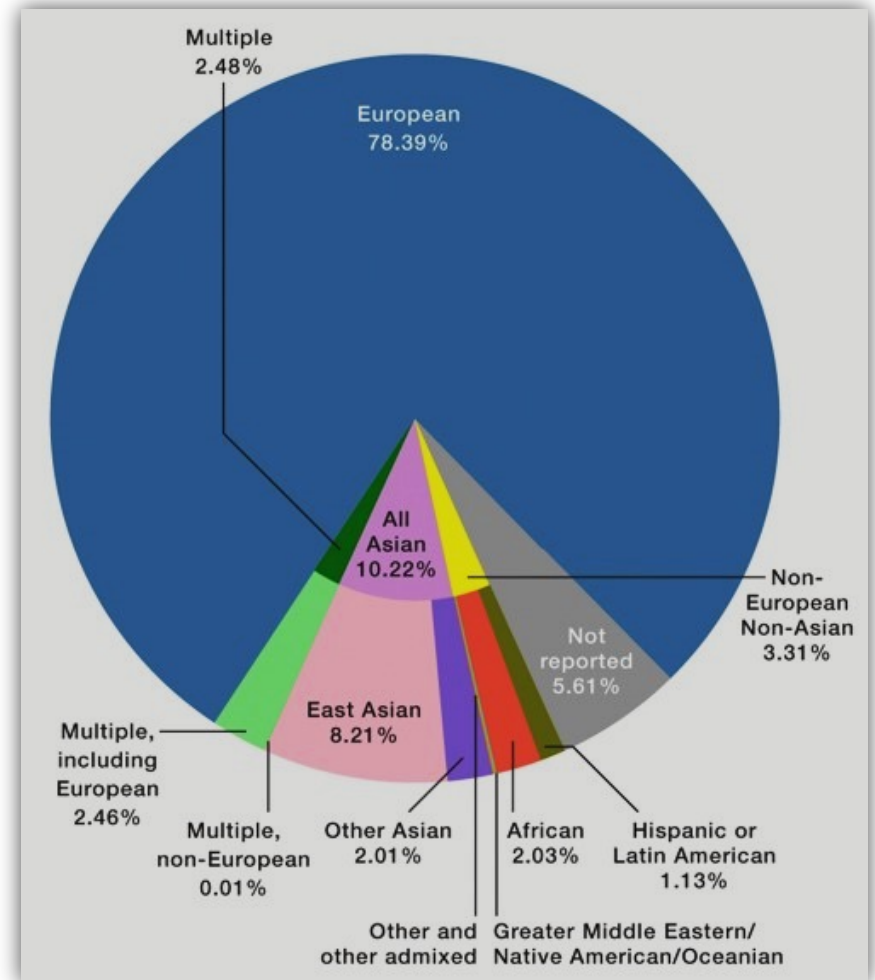
Corder et al. 1993 Science
Corder et al 1994
Farrer et al., 1997 JAMA

Hendrie et al. 2014. Int Psychogeriatr
Choi et al. 2019 J Clin Med
Reitz et al., 2013 JAMA

Why Study African Americans and other Diverse Groups

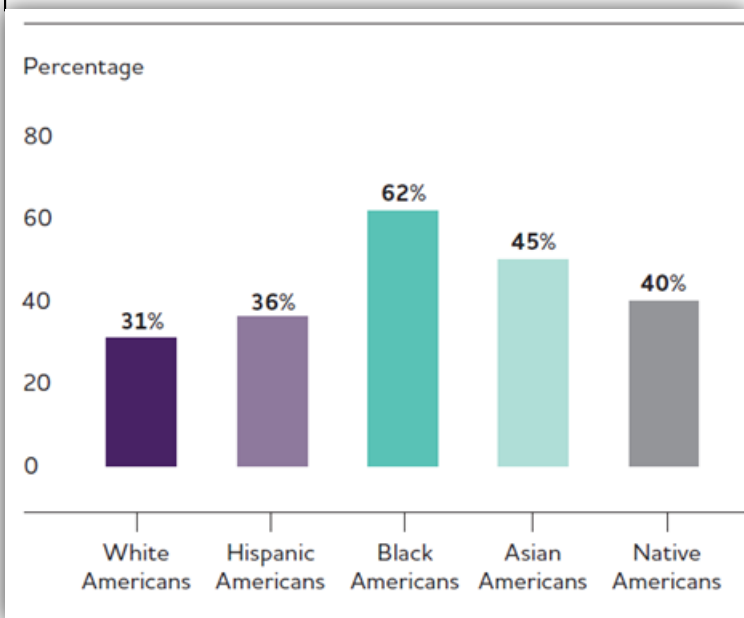
- The underrepresentation of ancestrally diverse populations in genetic studies:
 - hinders our ability to fully understand the genetic architecture of disease, and
 - intensify health inequalities.
 - reduce the power of risk prediction
- The translation of genetic research into clinical practice may be dangerously incomplete or, worse, mistaken

Ancestry category distribution of individuals in study catalog

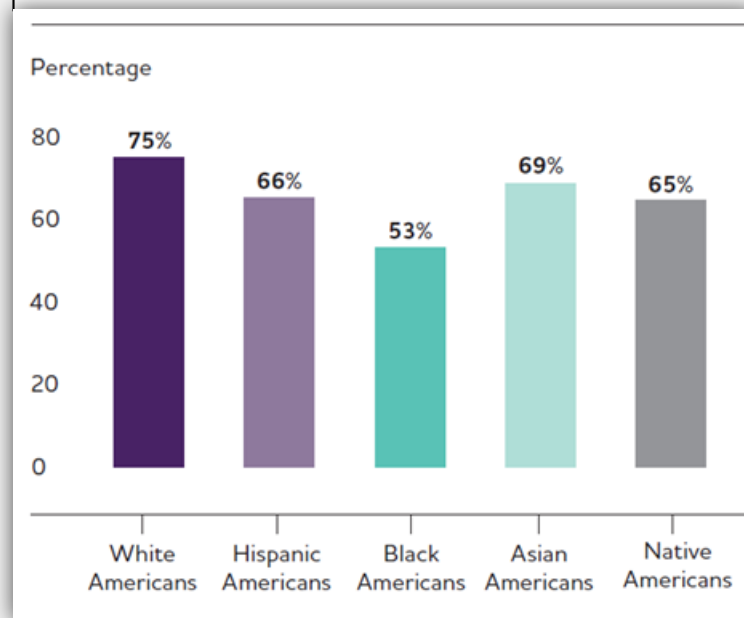


Current Perspectives on Alzheimer's Research

Percentage of US Adults Who Believe Medical Research is Biased Against People of Color



Percentage of US Adults Who Trust an Alzheimer's Cure Will be Shared Equally Regardless of Race, Color or Ethnicity





Welcome to the Alzheimer's Disease Sequencing Project

The overarching goals of the ADSP are to:

1. Identify new genomic variants contributing to increased risk of developing Late-Onset Alzheimer's Disease (LOAD)
2. Identify new genomic variants contributing to protection against developing Alzheimer's Disease (AD)
3. Provide insight as to why individuals with known risk factor variants escape from developing AD
4. Examine these factors in multi-ethnic populations as applicable in order to identify new pathways for disease prevention

Study Design

Learn about study design, sample selection, and data generation procedures

Apply for Data

Instructions on how to apply for ADSP data

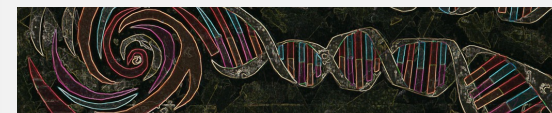
Access Data

Log into NIAGADS DSS

News

- Wednesday, March 3, 2021 - 09:15
[NIAGADS DSS Releases Additional 17K Whole Genomes](#)
- Wednesday, February 19, 2020 - 15:30
[NIAGADS DSS Releases 20K Whole Exomes](#)
- Monday, November 5, 2018 - 20:30
[Additional ADSP Data Released on NIAGADS DSS](#)
- Friday, September 7, 2018 - 19:15
[NIAGADS Data Sharing Service Now Accepting Applications](#)
- Friday, April 13, 2018 - 18:15
[Genetic variation paper published in Dementia and Geriatric Cognitive Disorders](#)

Funded by:
National Institute For Aging



<https://www.niagads.org/adsp/content/home>

Why are Such Studies Possible?

With the completion of the **Human Genome Project in 2003** and the **International HapMap Project in 2005**, researchers now have a set of research tools that make it possible to find the genetic contributions to common diseases. The tools include computerized databases that contain the reference human genome sequence, a map of human genetic variation and a set of new technologies that can quickly and accurately analyze whole-genome samples for genetic variations that contribute to the onset of a disease. **Genome-wide association** studies involve scanning markers across the genomes of many people to find genetic variations associated with a particular disease.

The National Plan has six ambitious goals to:

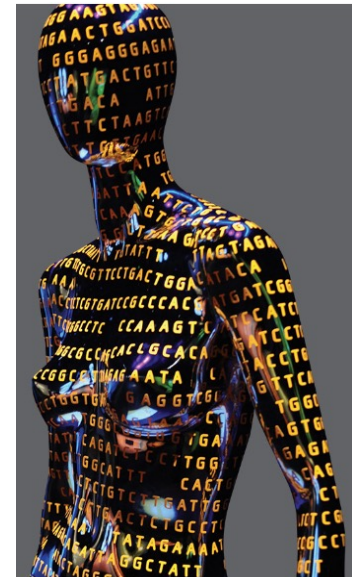
Prevent and Effectively Treat Alzheimer's Disease and Related Dementias by 2025

Enhance Care Quality and Efficiency

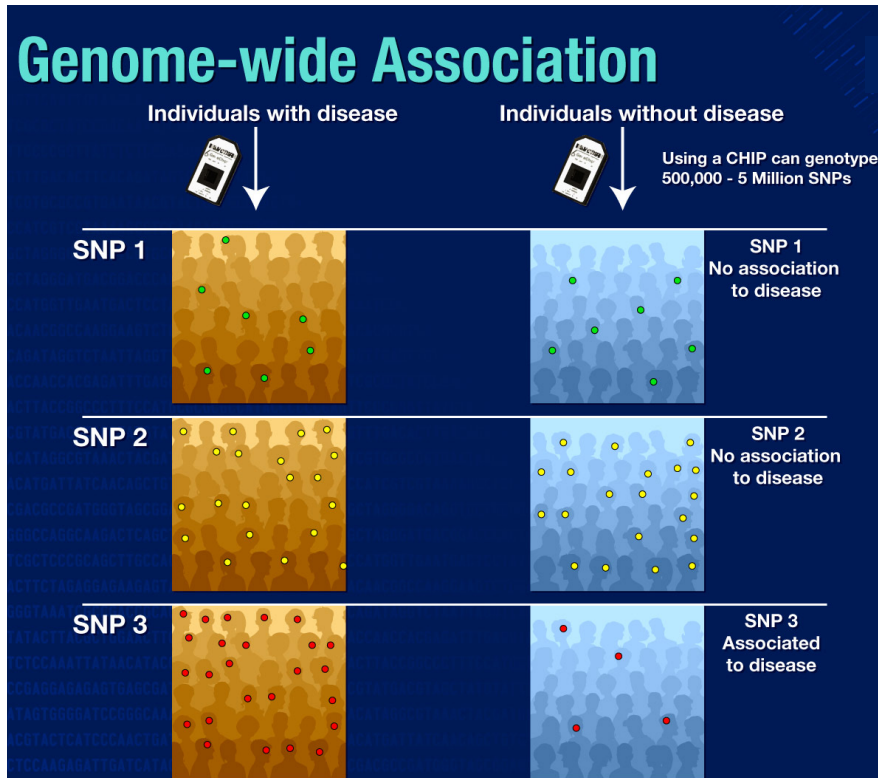
Expand Supports for People with Alzheimer's Disease and Related Dementias and Their Families

Enhance Public Awareness and Engagement
Improve Data to Track Progress

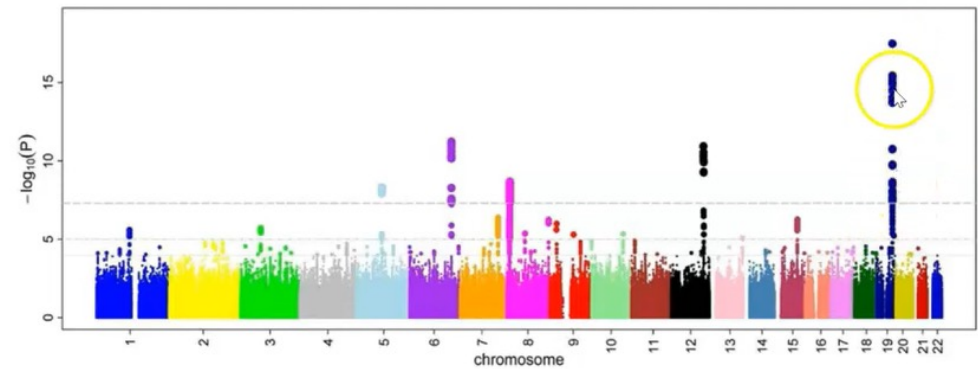
Accelerate Action to Promote Healthy Aging and Reduce Risk Factors for Alzheimer's Disease and Related Dementias



Identifying Risk Loci



GWAS does not identify a specific gene, but points us to Where to begin looking for a gene involved in disease ApoE is on chromosome 19



A Manhattan plot depicting several strongly associated risk loci

Form Strong Collaborations with Academic Partners

- Partnership began 20 years ago
- No formal involvement of African Americans in Genetics Research
- \$50,000 to **start** entire project
- Grass Roots efforts in NC
- Education, Partnerships with pastors, churches and community centers
- Strong foothold in NC, recruited over 1200 individuals in 5 years.
- Largest collection of African American Multiplex AD families



Goldie S. Byrd PhD

Professor and Director of the Maya Angelou Center for Health Equity at Wake Forest School of Medicine

Genetic Distinctions in Alzheimer Disease

Published Data over the past decade supports the need for Diversity in AD Genetics Research

Novel Rare Loci in African American GWAS (Kunkle et al. 2020)

IGFIR: chr15q26

AP15: chr11p12

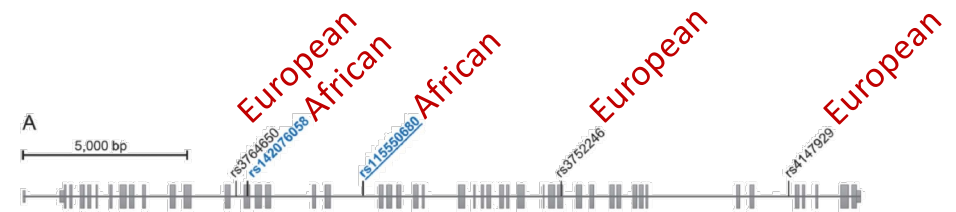
RBFOX1: chr16p13

Rare variants in AKAP9

African Americans (Logue et al. 2014)

Hispanic families in ADSP (Vardarajan et al. 2016)

ABCA7 in European and African Ancestries (Cukier et al. 2016)



- To prevent “genomic” health disparities
- To not miss population specific variants and/or genes

Ancestry and *APOE*

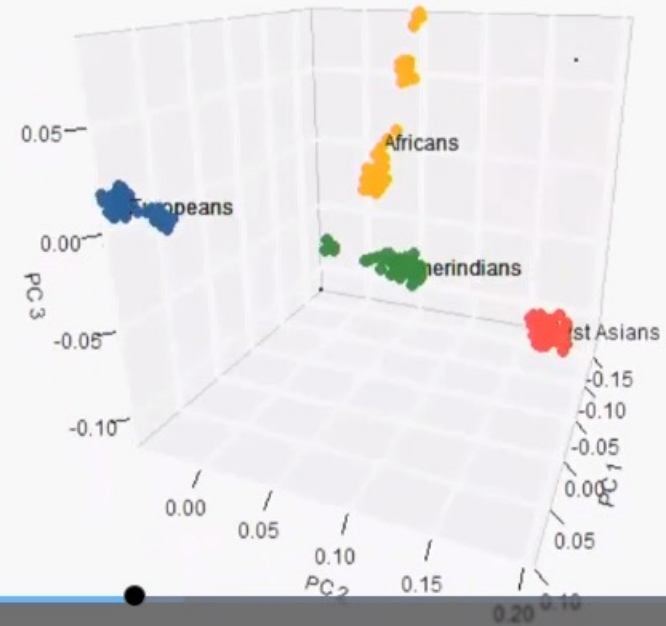
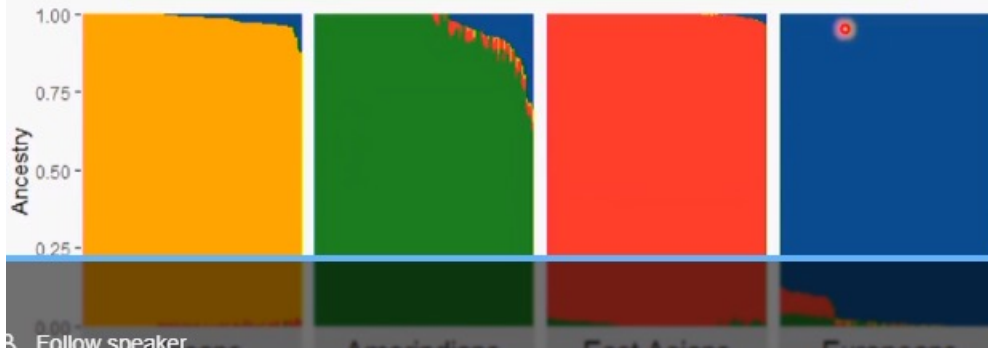
- **Why would *APOE* ϵ 4 have disparate effects across ancestries?**
 - Environmental/exposure effects that vary by population (gene x environment interaction)
 - Population-specific genetic effects that vary by population (gene x gene interaction)
- **Modern genomics methods can explore these hypotheses**
 - Rajabli et al., 2018, *PLoS Genet*
 - Rajabli et al., 2022, *PLoS Genet*

Global Ancestry: Continental Populations

{Algorithmic ancestry estimation: PCA}

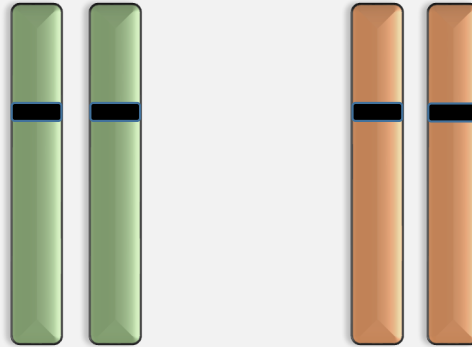


{Model Based Ancestry Estimation}



Admixture – Local Ancestry

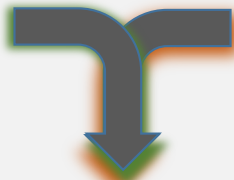
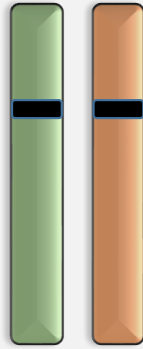
Population A



Population B

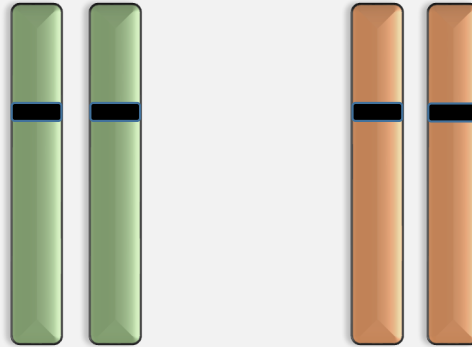


1st Generation

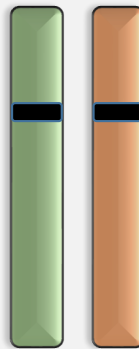


Admixture – Local Ancestry

Population A



1st Generation



Population B

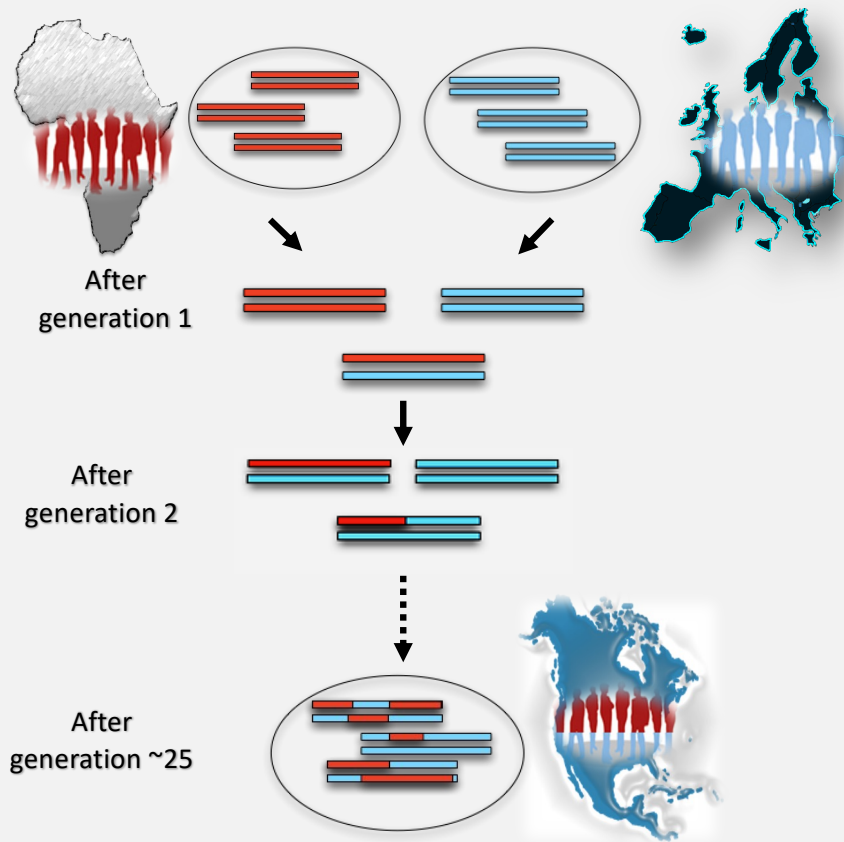


Generation: 1

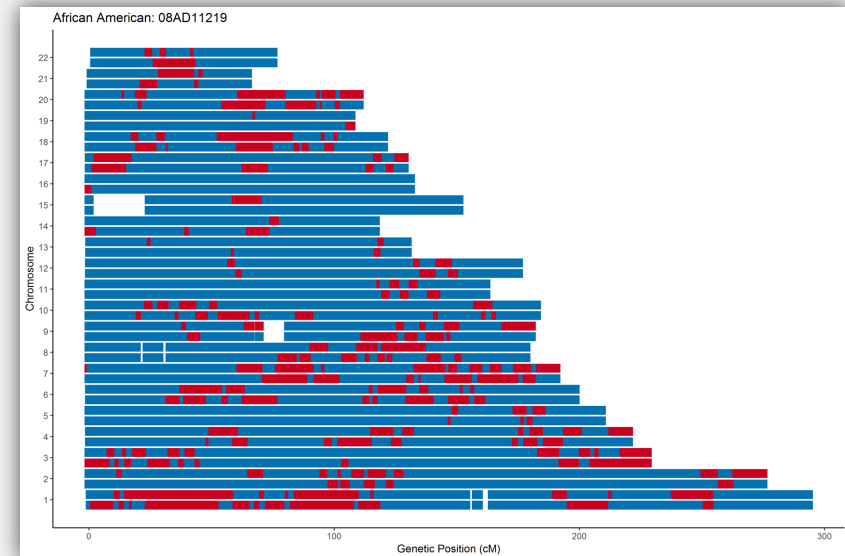


African Americans and Local Ancestry

- AA population is an admixed population

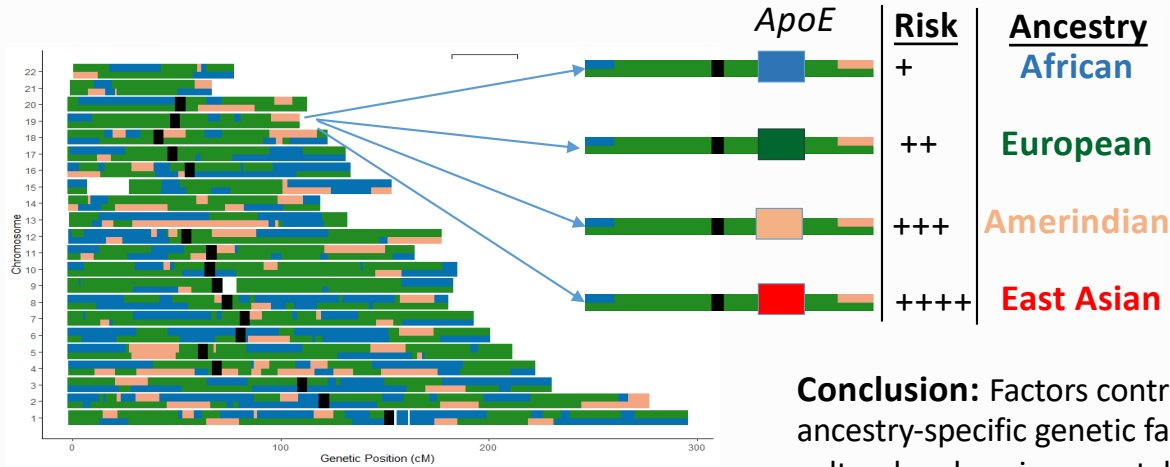


- Local ancestry estimates of a Black American individual



The US Black population is a two-way admixed population with genetic ancestry from African and European ancestors.

Admixture – Local Ancestry – ApoE



PLOS GENETICS

RESEARCH ARTICLE

Ancestral origin of *ApoE* $\epsilon 4$ Alzheimer disease risk in Puerto Rican and African American populations

Farid Rajabli^{1*}, Briseida E. Feliciano², Katrina Cellis¹, Kara L. Hamilton-Nelson¹, Patrice L. Whitehead¹, Larry D. Adams¹, Parker L. Busbies¹, Clara P. Manrique¹, Alejandra Rodriguez², Vanessa Rodriguez², Takiyah Starks³, Grace E. Byfield², Carolina B. Sierra Lopez², Jacob L. McCauley¹, Heriberto Acosta⁴, Angel Chinae², Brian W. Kunkle⁵, Christiane Reitz², Lindsay A. Farrer⁶, Gerard D. Schellenberg⁷, Badri N. Vardarajan⁸, Jeffery M. Vance^{1,8}, Michael L. Cuccaro^{1,8}, Eden R. Martin^{1,8}, Jonathan L. Haines⁹, Goldie S. Byrd², Gary W. Beecham^{1,8*}, Margaret A. Pericak-Vance^{1,8*}

Conclusion: Factors contributing to the lower risk effect in the *ApoE* are due to ancestry-specific genetic factors near *ApoE* rather than non-genetic ethnic, cultural and environmental factors.

ELSEVIER Alzheimer's & Dementia 15 (2019) 1524-1532

Featured Article

Local ancestry at *APOE* modifies Alzheimer's disease risk in Caribbean Hispanics

Elizabeth E. Blue^{a,*}, Andrea R. V. R. Horimoto^b, Shubhabrata Mukherjee^c, Ellen M. Wijsman^{a,b,d}, Timothy A. Thornton^{b,c,e}

Global and local ancestry modulate *APOE* association with Alzheimer's neuropathology and cognitive outcomes in an admixed sample

Michel Satya Naslavsky, Claudia K. Suemoto, Luciano Abreu Brito, Marília Oliveira Scliar, Renata Eloah Ferretti-Rebustini, Roberta Diehl Rodriguez, Renata E. P. Leite, Nathalia Matta Araujo, Victor Borda, Eduardo Tarazona-Santos, Wilson Jacob-Filho, Carlos Pasqualucci, Ricardo Nitrini, Kristine Yaffe, Mayana Zatz, Lea T. Grinberg

doi: <https://doi.org/10.1101/2022.02.02.22270331>

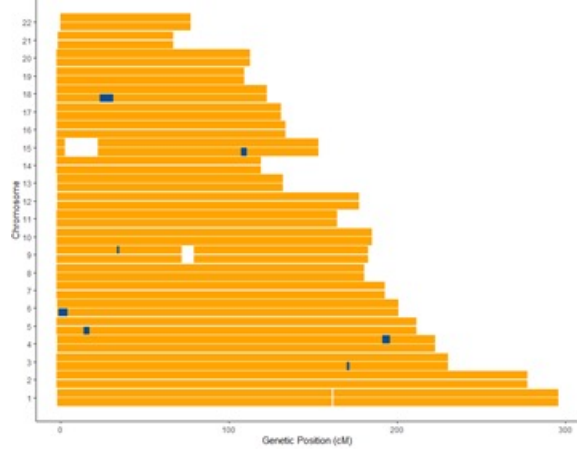


Farid Rajabli

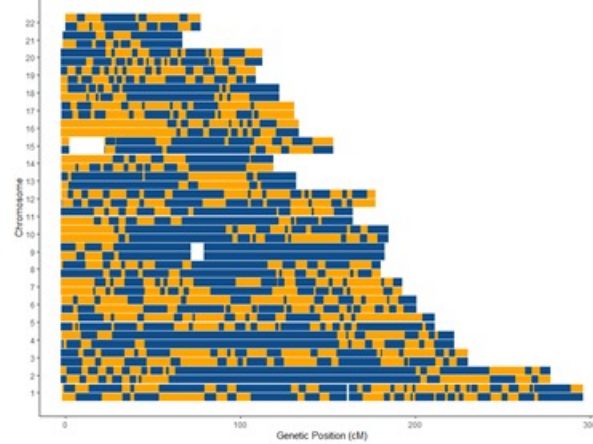


Gary Beecham

Ancestry vs. Race

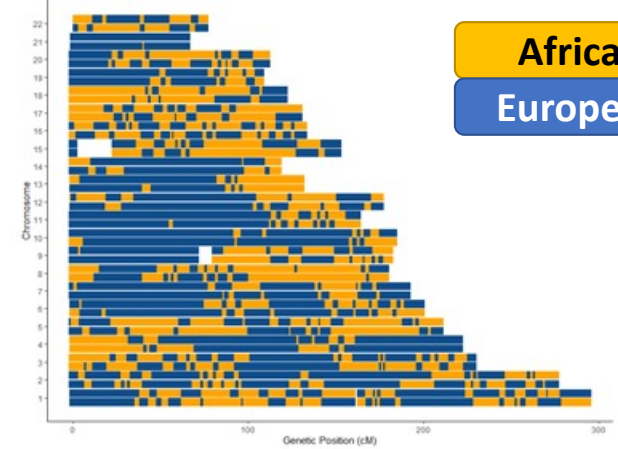


99%



61%

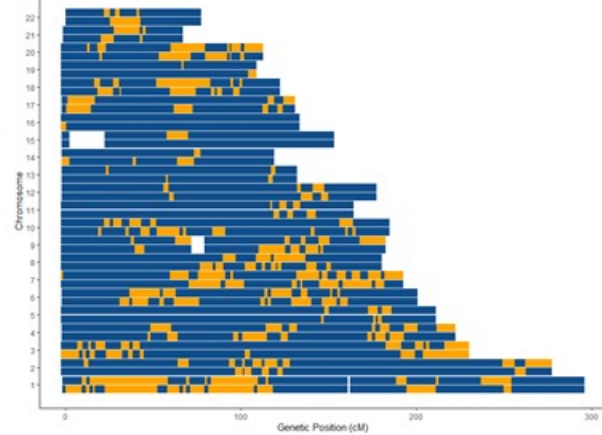
39%



62%

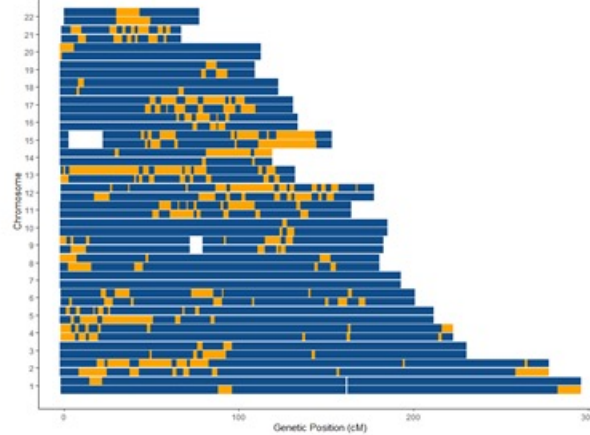
38%

African
European



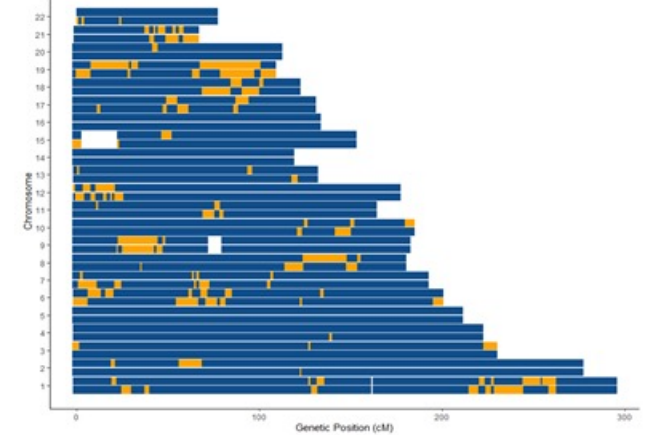
82%

18%



86%

14%



93%

7

Some Recent Studies Applying Ancestry to Genetic Studies

DOI: 10.1002/alz.12865

Alzheimer's & Dementia
THE JOURNAL OF THE ALZHEIMER'S ASSOCIATION

FEATURED ARTICLE

Admixture mapping identifies novel Alzheimer's disease risk regions in African Americans

Farid Rajabli¹ | Giuseppe Tosto² | Kara L. Hamilton-Nelson¹ | Brian W. Kunkle¹ | Badri N. Vardarajan² | Adam Naj³ | Patrice G. Whitehead¹ | Olivia K. Gardner¹ | William S. Bush⁴ | Sanjeev Saryia² | Richard P. Mayeux² | Lindsay A. Farrer⁵ | Michael L. Cuccaro^{1,6} | Jeffrey M. Vance^{1,6} | Anthony J. Griswold^{1,6} | Gerard D. Schellenberg³ | Jonathan L. Haines⁴ | Goldie S. Byrd⁷ | Christiane Reitz² | Gary W. Beecham^{1,6} | Margaret A. Pericak-Vance^{1,6} | Eden R. Martin^{1,6} | for the Alzheimer's Disease Genetics Consortium (ADGC), Collaboration on Alzheimer's Disease Research (CADRe) and Alzheimer's Disease Sequencing Project (ADSP)

RESEARCH ARTICLE

A locus at 19q13.31 significantly reduces the ApoE ε4 risk for Alzheimer's Disease in African Ancestry

Farid Rajabli¹, Gary W. Beecham^{1,2}, Hugh C. Hendrie³, Olusegun Baiyewu⁴, Adesola Ogunniyi⁵, Suljuan Gao¹, Nicholas A. Kuschel¹, Marina Lipkin-Vasquez^{1,2}, Ki L. Hamilton-Nelson¹, Juan I. Young^{1,2}, Derek M. Dykxhoorn^{1,2}, Karen Nuytemans^{1,2}, Brian W. Kunkle^{1,2}, Liyong Wang^{1,2}, Futai Jin⁶, Xiaoxiao Liu⁶, Briseida E. Feliciano Astacio⁷, Alzheimer's Disease Sequencing Project, Alzheimer's Disease Genetic Consortium, Gerard D. Schellenberg⁸, Clifton L. Dalgard⁹, Anthony J. Griswold^{1,2}, Goldie S. Byrd¹⁰, Christiane Reitz¹¹, Michael L. Cuccaro^{1,2}, Jonathan L. Haines¹²

www.nature.com/mp

Check for updates

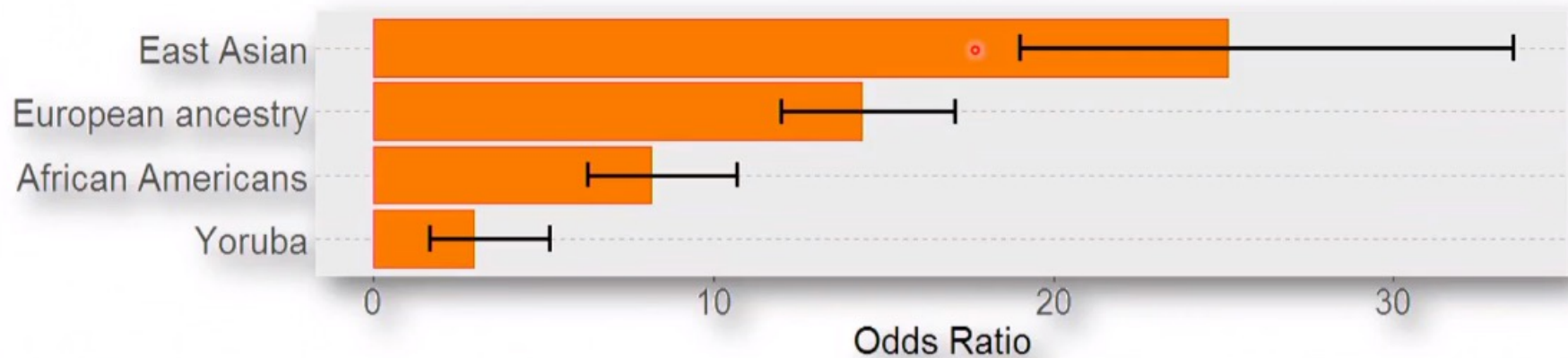
MOLECULAR PSYCHIATRY

Admixture Mapping of Alzheimer's disease in Caribbean Hispanics identifies a new locus on 22q13.1

Zaghan Kozl^{1,2,3,12}, Sanjeev Saryia^{3,2,4,12}, Yoon A. Kim^{1,5}, Farid Rajabli⁶, Eden Martin^{6,7}, Dolly Reyes-Dumeyer^{2,4}, Badri Vardarajan^{2,4}, Uleyda Maldonado⁸, Jonathan L. Haines⁹, Richard Mayeux^{1,2,4,10,11}, Yvonne Z. Jimenez-Velazquez², Ismael Santa-Maria^{1,5} and Giuseppe Tosto^{1,2,4,10}

Back to *APOE*...

APOE $\epsilon 4$ risk effect ($\epsilon 3\epsilon 3$ vs $\epsilon 4\epsilon 4$)



Can genetic ancestry methods bring insight into this differential *APOE* effect?

Hendrie et al., 2014, *Int Psychogeriatr*
Choi et al., 2019, *J Clin Med*
Reitz et al., 2013, *JAMA*

Ancestry vs. Race

- Ancestry is biological and is about the history of genetic variation and the origin of one's population
- Race, itself is not biological, but is often self-ascribed or socially-ascribed by others
- As I have shown populations used to live in isolation with each geographic region having its own genetic map (Continental Populations)
- Populations today particularly in the US are admixed (multiple ancestries) with individuals of European, African, Amerindian and Asian ancestry.

Most GWAS for AD have been conducted in individuals of EU descent

Largest AD Genome-wide Association Studies



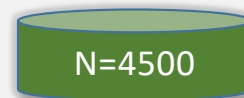
Europeans

(Kunkle et al. 2019)



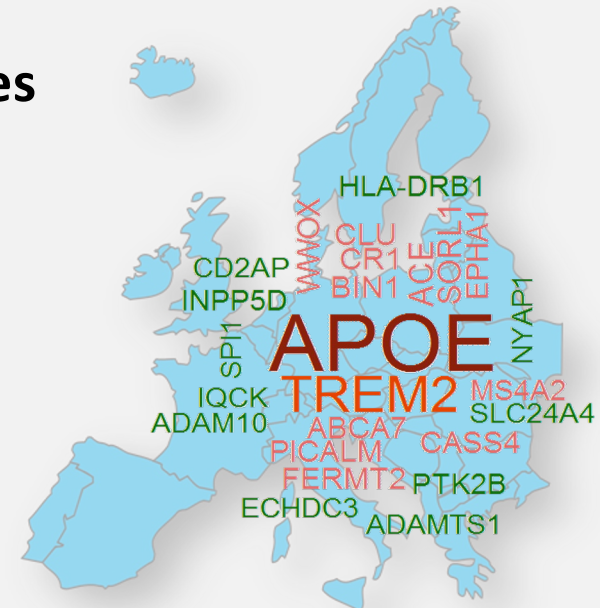
African Americans

(Kunkle et al. 2020)



Hispanics

(Tosto et al. 2015)



Inclusion at all Levels

*Diversity makes for a rich tapestry,
and we must understand
that all the threads of the tapestry are*

EQUAL IN VALUE

no matter what their color.

- Maya Angelou

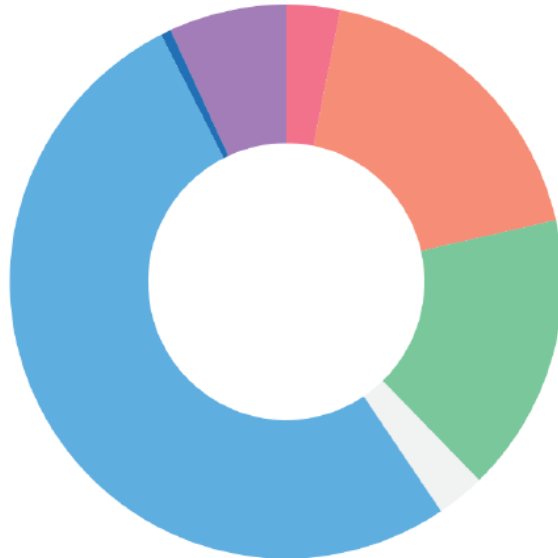
- Researchers
- Workforce
- Participants
- Volunteers



A Transformational Approach to Diversity, Equity and Inclusion in Precision Medicine

All of Us Research Program

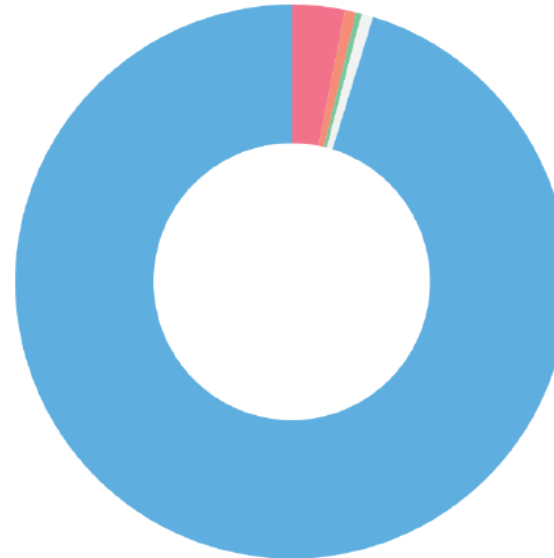
- Asian 3.10%
- BAA 18.10%
- H/L 16.10%
- Other 2.80%
- White 51.40%
- NR 0.60%
- More than one race 6.80%



Source: [All of Us Research Program Data Snapshots, Race and Ethnicity](#) (Updated March 3, 2023)

Other Genomic Studies

- Asian 3.09%
- BAA 0.65%
- H/L 0.33%
- Other/More than one 0.69%
- White/European Descent 95.24%



Source: [Global Genome Wide Association Studies \(GWAS\)](#) (Updated February 3, 2023)

Alzheimer's Disease Sequencing Project (ADSP)*

- ADSP <https://www.niagads.org/adsp/content/home>
 - Sequence over 100,000 individuals from multiple ancestries and ethnic backgrounds
 - 100's of Researchers from around the world are contributing and working together to find new drug targets for therapies
- READD-ADSP part of this larger international ADSP effort
 - Goals:
 - DNA, RNA, Plasma Biomarkers and CVD markers for extensive phenotyping
 - Whole Genome Sequencing

Recruitment and Retention for AD Diversity Genetic Cohorts in the ADSP (REAAD-ADSP)

University of Miami

Margaret Pericak-Vance
Brian Kunkle
Jeffery Vance

Wake Forest University

Goldie Byrd

Columbia University

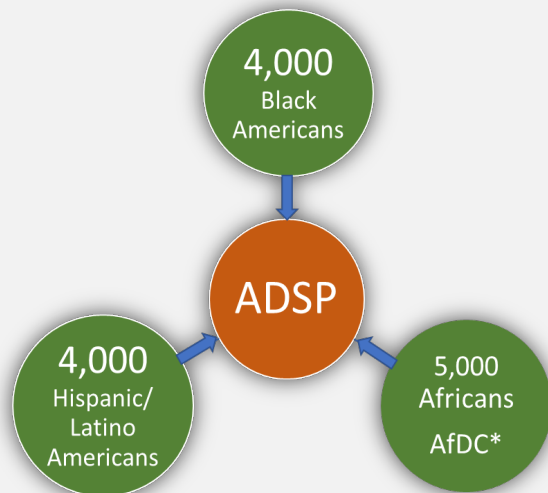
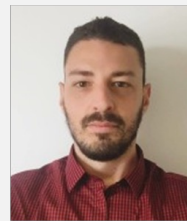
Giuseppe Tosto
Christiane Reitz

University of Ibadan

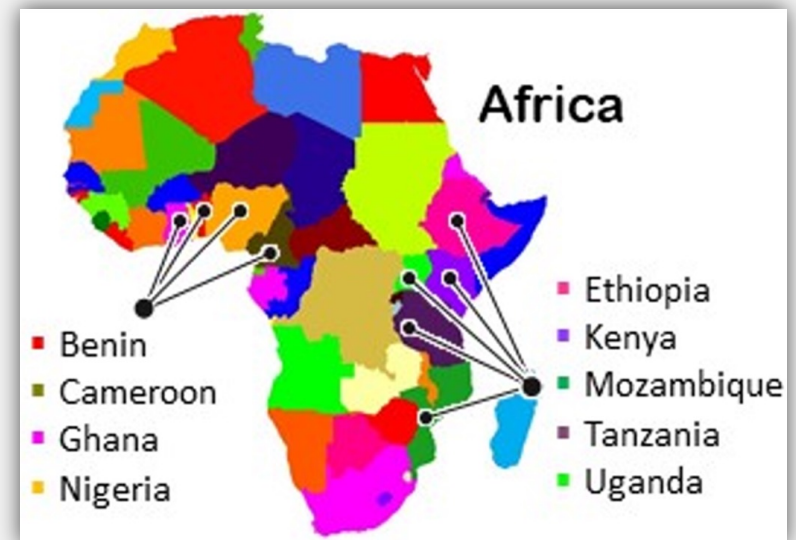
Rufus Akinyemi
Adesola Ogunniyi

Case Western Reserve University

Jonathan Haines
Will Bush

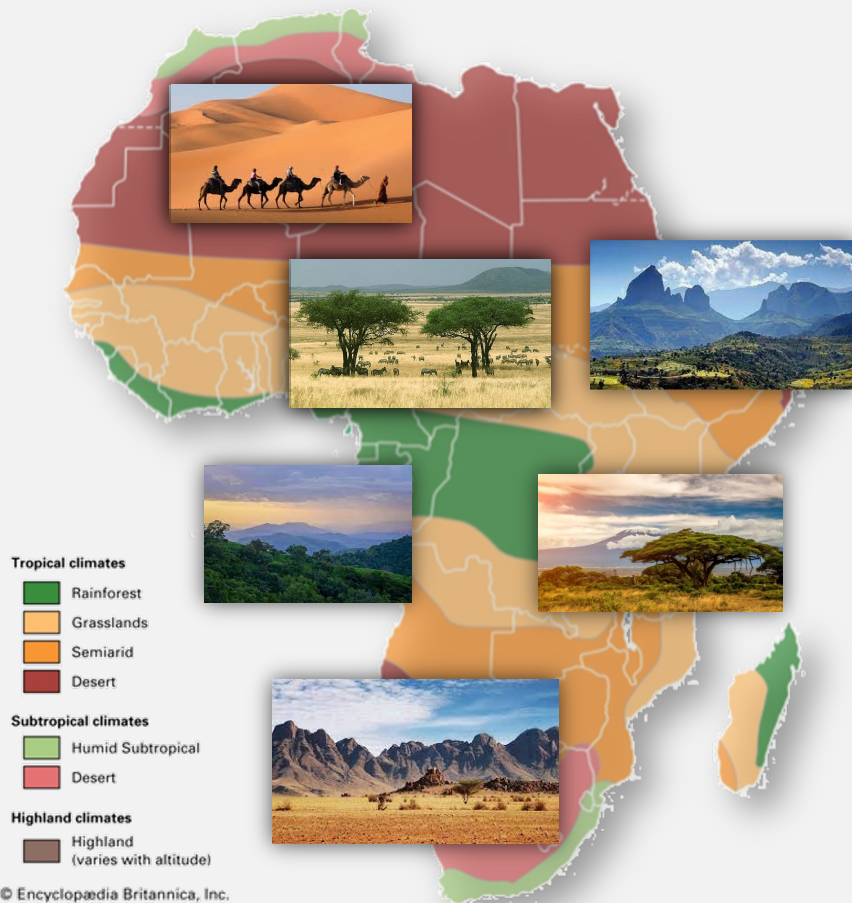


*AfDC; African Dementia Consortium



Diversity: Africa

- Africa is home to ethnic, cultural, linguistic, ancestral, and ecological/environmental diversity
 - 54 Countries
 - ~3,000 different ethnic groups
 - ~2,100 different languages



REAAD-ADSP GOALS

- DNA, RNA, Plasma Biomarkers and Whole Genome Sequencing
- Social Determinants of Health (SDOH) also influence AD risk, but studies collecting both biological and SDOH data are rare. *Collecting SDOH data provides a basis for the integrative studies of biological and social risks of AD.*



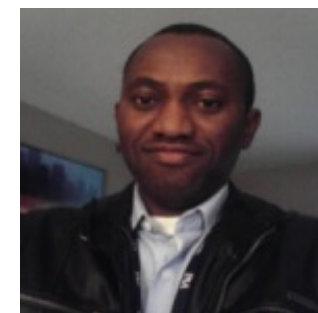
Dr. Farid Rajabli (U of Miami)



Dr. Azizi Seixas. (U of Miami)

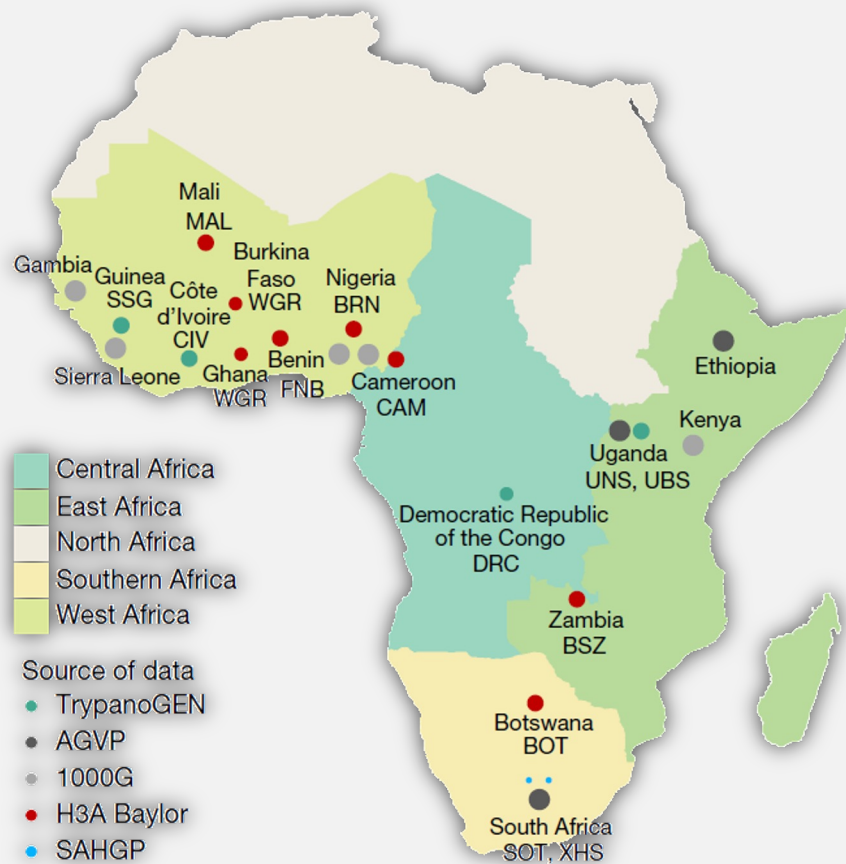


www.cdc.gov



Dr. Joshua Akinyemi
(Ibadan, Nigeria)

Genetic Diversity: sub-Saharan Africa

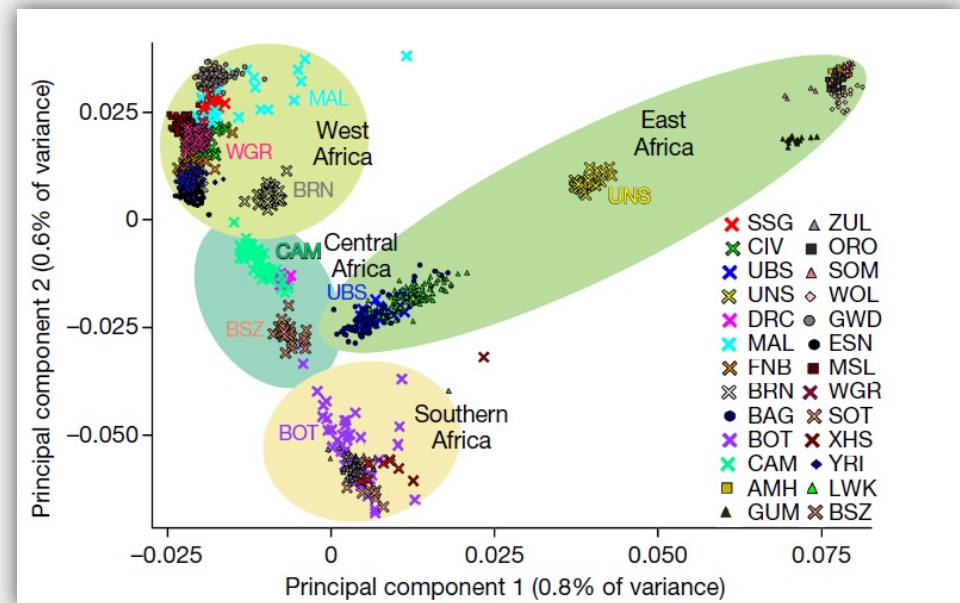


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Africa, the most genetically diverse continent



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Community Partners/Collaborators

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COAACH AAAD Team/Staff and Students

Wake Forest University School of Medicine

North Carolina A&T State University,

Alumni

Endowment Donors

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NHGRI, NIMHD, NIA

Merck, Inc. & Co,

AARP, Inc.

Alzheimer's Association



Collaborators' Meeting at COAACH at NC A&T to Initiate New Genetics of Familial Alzheimer's disease in African Americans