Genetic Drivers of Resilience to Alzheimer's Disease

Timothy Hohman, PhD Associate Professor of Neurology Vanderbilt University Medical Center

Disclosures

• Scientific Advisory Board for Vivid Genomics

Explosion of Big Data in Alzheimer's Disease



Precision Medicine in AD



Ferretti et al., Nature Reviews Neurology, 2018

Integrating AI into Genomic Discovery at NIA





Drug Discovery Framework for the Neuroresilience Program

Outline

- Defining Resilience
- Genetic Drivers of Resilience
- Future Directions

Amyloid Cascade Hypothesis



Resistance and Resilience



Heterogeneity in Cognitive Performance



Resilience as a Pathway to New Targets



All Participants:

- Amyloid⁺
- Tau⁺
- *ΑΡΟΕ* ε4⁺

Harmonizing Data to Increase Sample Size



Fluid Biomarker Harmonization



Carlos Cruchaga, PhD Washington University

Cognitive Harmonization



Paul Crane, MD, MPH University of Washington

Jesse Mez. MD. MS Boston University

Coordinating Centers



Timothy Hohman, PhD Vanderbilt University Medical Center MPI

Storage & Informatics



Arthur Toga, PhD University of Southern California MPI

CHARGE Coordination

Michael Cuccaro, PhD

University of Miami MPI



Mohamad Habes, PhD University of Tex as Health San Antonio

Diffusion MRI Harmonization



Bennett Landman, PhD Vanderbilt University

Structural MRI Harmonization





Shannon Risacher, PhD Indiana University

Christos Davatzikos, PhD University of Pennsylvania

Vascular Harmonization



Adam Brickman, PhD Columbia University



Richard Mayeux, MD, MSc Columbia University

Integration & Analytics



Paul Thompson, PhD Andrew Saykin, PsyD University of Southern Indiana University California



Thomas Montine, MD. PhD Stanford University



Garv Beecham, PhD University of Miami

Elizabeth Mormino, PhD Stanford University

Duygu Tosun, PhD University of California San Francisco



PET Harmonization





Example Domain: Cognition

| | NACC | ACT | ADNI | ROSMAP & MARS | TOTAL |
|--------------------------|-------|------|-------|------------------|--------|
| Total Cog N | 41459 | 5546 | 3189* | 4386 | 54,580 |
| Total ADSP N | 10486 | 1392 | 1574 | 1575 | 15,027 |
| Total Cog & ADSP N | 8458 | 1340 | 1574 | 1560 | 12,932 |



Paul Crane, MD, MPH University of Washington Anchor Items

 Tests administered consistently across studies serve as anchor items



Logical Memory Immediate

Harmonization Approach

Applied Psychometrics

- An expert panel assigns items to one of 4 domains
 - Memory
 - Language
 - Executive functioning
 - Visuospatial ability



Harmonization Approach

All Diagnoses

Cognitively Normal







Data Integration to Define Resilience



Genetic Architecture of Resilience

Analytical Plan



Logan Dumitrescu, PhD

Resilience GWAS Workflow

Post-GWAS Analytical Workflow



Stratified Analyses:

- *ΑΡΟΕ-ε4*
- Cognitively Normal

Dumitrescu et al., Brain, 2020

Genetic Correlation and Pathway Results



Dumitrescu et al., Brain, 2020

Cognitively Normal Resilience Results



Gene Mapping and Functional Annotation

| | Methylation targets for rs2571244 (18:55473651) | | | |
|---|---|--------------------------|--------------|------------------------|
| _ | target | target start position | Spearman's p | Ρ |
| | cg19596477 | 18:55472454 | 0.33 | 2.24x10 ⁻¹³ |
| | cg16310513 | 18:55471075 | 0.17 | 1.79x10-4 |
| | cg16141316 | 18:55469758 | -0.12 | 8.14x10-3 |
| | | | | |



Prefrontal Cortex Methylation

Dumitrescu et al., Brain, 2020

| | | | | | 1 |
|-----|---|------------|--------------------------|-----------|----------|
| | Adipose- | 1.000 | 1.000 | 1.000 | adjusted |
| | Blood- | 1.000 | 1.000 | 1.000 | aujusteu |
| | Blood Vessel- | 1.000 | 1.000 | 1.000 | p-value |
| | Brain- | 1.000 | | 1.000 | p value |
| | Circulating Cell | 1.000 | 1.000 | 1.000 | 10 |
| | Connective Tissue- | 1.000 | | 1.000 | 1.0 |
| | Digestive- | 1.000 | 1.000 | 1.000 | |
| | Embryo- | 1.000 | 1.000 | 1.000 | |
| | Endocrine | 1.000 | 0.001 | 1.000 | |
| | Epithelial- | 1.000 | 0.001 | 1.000 | |
| | ESC- | 1.000 | 1.000 | 1.000 | |
| Lob | Eye- | 1.000 | 1.000 | 1.000 | |
| | emale Reproductive- | 1.000 | 1.000 | 1.000 | |
| | Hair Follicle- | 1.000 | 1.000 | 1.000 | |
| Ť. | Heart- | 1.000 | | 1.000 | |
| 8 | Immune Organ- | 1.000 | | 1.000 | |
| neo | iPSC- | 1.000 | 1.000 | 1.000 | |
| | Liver- | 1.000 | | 1.000 | |
| SS | Lung | 1.000 | | 1.000 | |
| | Male Reproductive- | 1.000 | 1 000 | 1.000 | |
| - | Nervous- | 1.000 | 1 000 | 1.000 | |
| | Olfactory- | 1.000 | 1.000 | 1.000 | |
| | Placenta- | 1.000 | 1,000 | 1.000 | |
| | Skeletal Muscles | 1,000 | | 1.000 | |
| | Smooth Muscle | 1,000 | 1,000 | 1.000 | |
| | Stron Coll- | 1.000 | 1.000 | 1.000 | 0 1 |
| | Throat | 1.000 | 1.000 | 1.000 | 0.1 |
| | Tongua | 1.000 | 1.000 | 1.000 | 0.0 |
| | Tanaila | 1.000 | 1.000 | 1.000 | 0.0 |
| | | 1.000 | 1.000 | 1.000 | |
| | Umbilical Cord- | 1.000 | 1.000 | 1.000 | |
| | Urinary- | 1.000 | 1.000 | 1.000 | 1 |
| | and the second se | 5 Enhancer | A Ethancel | PHMM Entr | |
| | FENIO, | Poadmaphia | ON ⁵ EnhRoadm | | |
| | | 4P | 200 | | |
| | | | Annotation | | |

ATP8B1 Function and Posthoc Analyses

- ATPase phospholipid transporting 8B1
 - Codes an aminophospholid translocase protein
 - Operates in the Liver to maintain bile acid homeostasis

| | SNP Association with Bile Acids | | | |
|-----------|---------------------------------|------|------|--|
| Bile Acid | β | DF | Ρ | |
| ТСА | 0.40 | 1019 | 0.01 | |
| GLCA | 0.33 | 1019 | 0.02 | |
| GCA | 0.31 | 1019 | 0.02 | |
| TDCA | 0.33 | 1019 | 0.04 | |
| TCDCA | 0.30 | 1019 | 0.04 | |

• Exome sequencing identified risk variants in ATP8B4 (Holstege, 2020)

Summary of Genetic Resilience

- Genetic architecture of resilience is distinct from clinical AD
 - Small contribution of APOE
- Observed a shared architecture with cognition and education
- Modest genetic correlation with <u>vascular</u> and <u>psychiatric</u> phenotypes
- *ATP8B1* is a novel resilience gene along the <u>bile acid metabolism</u> and phospholipid transport pathway

Gene Modifiers of Neuropathology

Data Integration to Identify Gene Modifiers



Amyloid Burden

Identify genetic variants that confer neuroprotection from AD



Genome Wide Association Study

Study Design



Mabel Seto, PhD

Study Design

Annual Change in Hippocampal Volume (slopes)



Results



 β = -0.005; p = 0.004

 β = -0.24, p = 0.0112

Semaphorin 5b

- Semaphorin 5B is a member of the semaphorin family
 - Development of nervous system
 - Regulation of neuronal proliferation and migration
- Overexpression of *SEMA5B* is associated with reduction in hippocampal synapse number (O'Connor et al. 2011)
- SEMA5B knock-out causes aberrant branching of neurons (Jung et al. 2019)





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- *ATP8B1* is a novel resilience gene along the <u>bile acid metabolism</u> and phospholipid transport pathway
- SEMA5B is a novel susceptibility gene that may be <u>beneficial</u> in absence of amyloid, but <u>detrimental</u> with onset of pathology

Incorporating Sex and Gender into Resilience Models

Females Have More AD Pathology at Autopsy



Neuropathology at Autopsy

Oveisgharan et al., Acta Neuropathologica, 2018



Carroll et al., Brain Research, 2010



Females with Pathology Decline More Rapidly



CSF Biomarker Association with Atrophy



APOE Association with AD is Stronger in Females



Farrer et al., JAMA Neurology, 1997



Neu et al., JAMA Neurology, 2017

Summary of Sex Differences



Leveraging Genome-Wide Data to Explore Sex Differences in AD

GWAS of CSF A β -42

Deming et al., Acta Neuropathologica, 2017

GWAS of CSF A β -42

Deming, ... Hohman, Acta Neuropathologica, 2018

Locus Zoom

rs316341 is eQTL for SERPINB1, SERPINB6, and SERPINB9 in Braineac and GTex

Deming, ... Hohman, Acta Neuropathologica, 2018

SERPINB1 Functional Evidence

 Female-specific association between prefrontal cortex expression of SERPINB1 (p=0.02) and SERPINB6 (p=0.00007) and amyloid levels in brain tissue

Serpin Signaling and Amyloidosis

- Serpins are Protease Inhibitors
 - Serpin-B1 Regulates Neutrophil Infiltration
- Serpins have been shown to inhibit Aβ toxicity

- Likely through regulation of neutrophils

- Some evidence of sex difference in neutrophil infiltration and clearance
 - Female mice show more activated neutrophils than male mice following stroke
 - Estradiol modulates neutrophil infiltration and clearance

SERPINB1 in Brain Tissue

RNA AND PROTEIN EXPRESSION SUMMARY

Testis

SERPINB1 Staining in AD Cortex

Hot off the Presses (or Computer...)

Sex-Stratified Memory GWAS

rs2590395 is eQTL for SERPINB2 and SERPINB10 in Blood

Precision Medicine through Collaboration

Future Directions

APOE Modifiers

Protective Proteomic Effects

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reas, BS <u>Columbia</u> • Richar