

Molecular drivers of Alzheimer's disease resilience through genomic and proteomic analyses

Timothy Hohman, PhD Professor of Neurology Vanderbilt University Medical Center

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Computational Neurogenomics Team (CNT)





Abel Belachew



Aditi Sathe MS



MS



Julia Libby

MS

Michelle Clifton MS



Skylar Walters, MS

Yiyang Wu MD, PhD

Alaina Durant





Melisa Lara Gomez



Alexis Smith

Vaibhav Janve PhD





Emily Mahoney

Dimitrios Zaras PhD

Hui Shi, MS



Edward Miller



MIT





PhD

Jackie Eissman PhD

Annah Sclafani PhD



Rebecca Winfree PhD



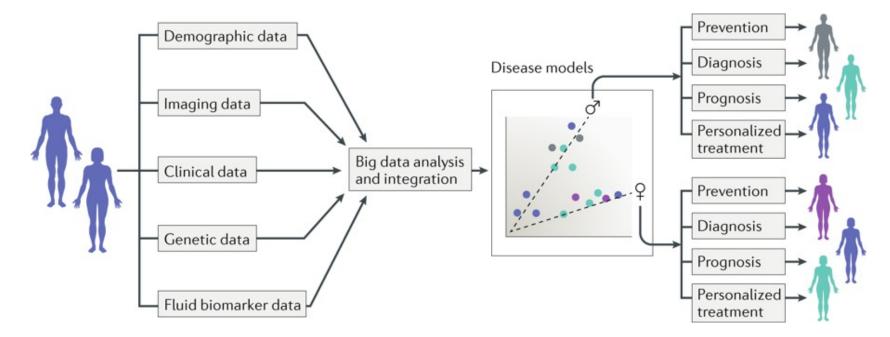




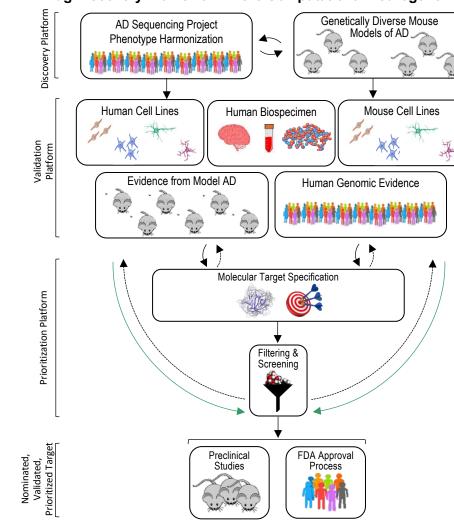




Precision Medicine in AD

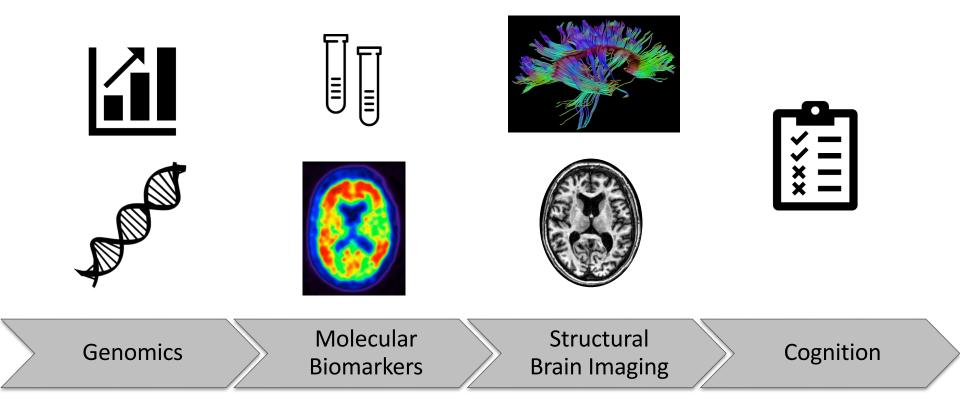


Ferretti et al., Nature Reviews Neurology, 2018

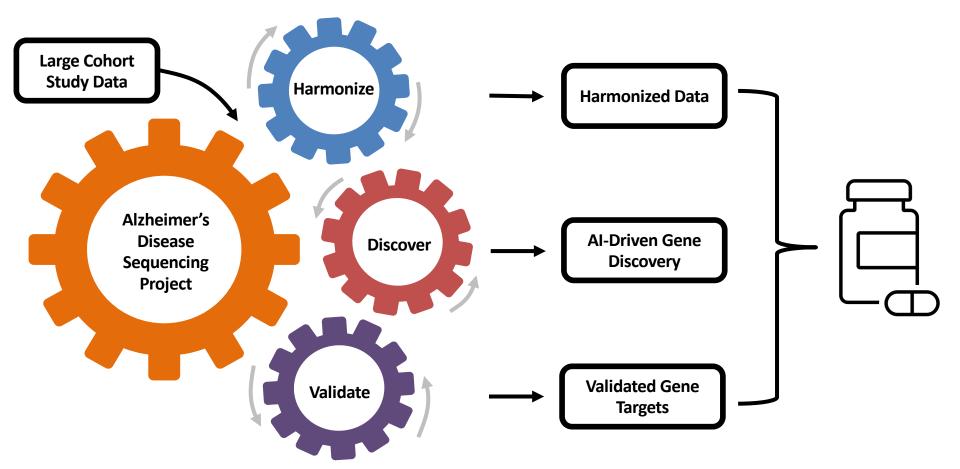


Drug Discovery Framework in the Computational Neurogenomics Team

Explosion of Big Data in Alzheimer's Disease

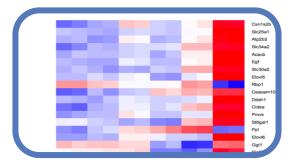


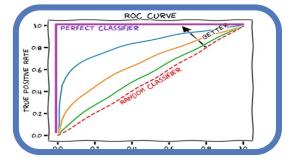
Integrating AI into Genomic Discovery at NIA



New ADSP Programs

Maximum Score	Patient's Score	Questions
5		"What is the year? Season? Date? Day? Month?"
5		"Where are we now? State? County? Town/city? Hospital? Floor?"
3		The examiner names three unrelated objects clearly and slowly, then the instructor asks the patient to name all three of them. The patient's response is used for scoring. The examiner repeats them until patient learns all of them, if possible.
5		"I would like you to count backward from 100 by sevens." (93, 86, 79, 72, 65,) Alternative: "Spell WORLD backwards." (D-L-R-O-W)
3		"Earlier I told you the names of three things. Can you tell me what those were?"
2		Show the patient two simple objects, such as a wristwatch and a pencil and ask the patient to name them.
1		"Repeat the phrase: 'No ifs, ands, or buts."
3		"Take the paper in your right hand, fold it in half, and put it on the floor. (The examiner gives the patient a piece of blank paper.)
1		"Please read this and do what it says." (Written instruction is "Close your eyes.")
1		"Make up and write a sentence about anything." (This sentence must contain a noun and a verb.)
		"Please copy this picture." (The examiner gives the patient a blank piece of paper and asks him/her to draw the symbol below. All 10 angles must be present and two must intersect.)
1		





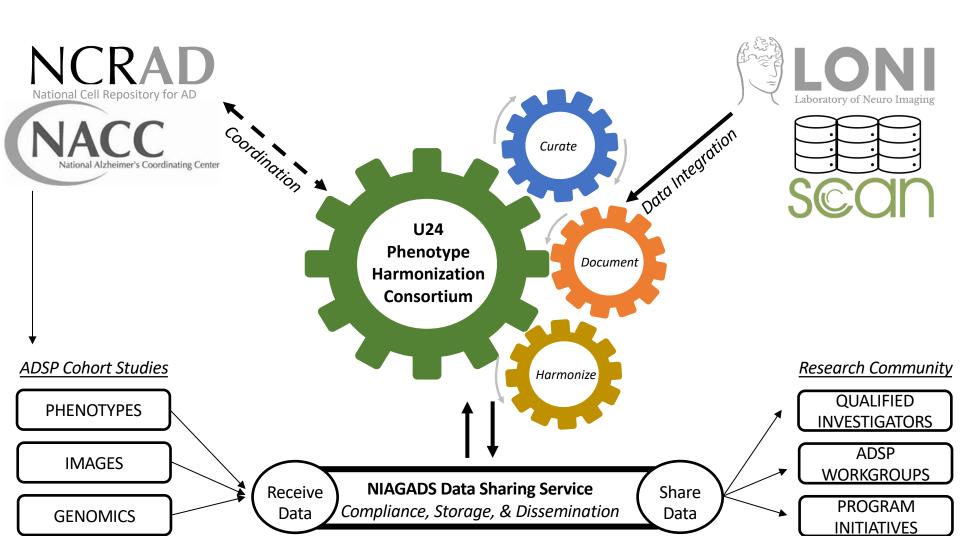












ADSP-PHC Teams

Fluid Biomarker Harmonization



Carlos Cruchaga, PhD Washington University

Cognitive Harmonization





Paul Crane, MD, MPH University of Washington

Jesse Mez, MD, MS Boston University

Coordinating Center



Timothy Hohman, PhD Vanderbilt University Medical Center Contact-PI

Storage & Informatics



Arthur Toga, PhD University of Southern California MPI

Michael Cuccaro, PhD University of Miami MPI

CHARGE Coordination



Mohamad Habes, PhD University of Texas Health Sán Antonio

Diffusion MRI Harmonization



Bennett Landman, PhD Vanderbilt University

Structural MRI Harmonization





Shannon Risacher, PhD Indiana University

Christos Davatzikos. PhD University of Pennsvlvania

Vascular Harmonization



Adam Brickman, PhD Columbia University



Integration & Analytics



Andrew Saykin, PsyD Indiana University

Neuropathology Harmonization



Thomas Montine, MD. PhD Stanford University



Gary Beecham, PhD University of Miami

Elizabeth Mormino, PhD Stanford University

Duygu Tosun, PhD University of California San Francisco

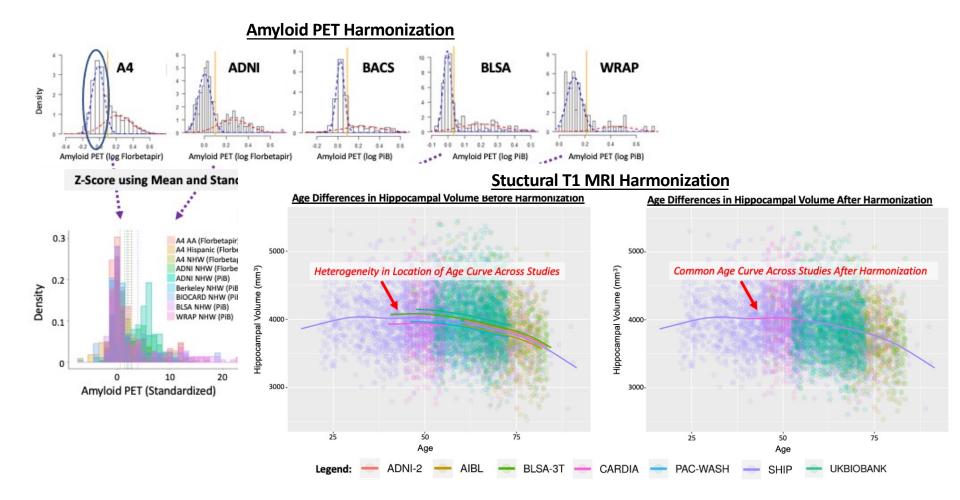








Paul Thompson, PhD University of Southern California



Data Harmonized in Year 2

Cohort	Autopsy	Vascular Risk Factors	Cognition	Fluid Biomarker	Diffusion Tensor Imaging (DTI)	Fluid Attenuated Inversion Recovery (FLAIR)	Magnetic Resonance Imaging (MRI) Freesurfer	Magnetic Resonance Imaging (MRI) MUSE	Positron Emission Tomography (PET) Amyloid	Positron Emission Tomography (PET) Tau
TOTAL	23,020	57,083	83,548	3,047	2,830	10,020	7,679	10,590	7,602	1,836
A4	0	0	6,595	0	0	1,198	0	0	4,486	447
ACT	532	0	5,546	0	0	0	0	0	0	0
ADNI	0	1,577	3,539	1,249	1,198	7,054	2,560	2,592	1,803	944
EFIGA	0	5,965	6,903	0	0	0	0	0	0	0
Knight ADRC	0	0	2,477	64	0	0	0	0	0	0
NACC	19,899	41,172	41,538	2,094	646	0	3,690	5,004	920	102
NIA-AD FBS	570	405	4,564	0	0	0	0	0	0	0
ROS/MAP/ MARS	2,019	4,500	4,550	0	643	0	0	1,183	0	0
WHICAP	0	1,830	6,257	0	0	1,768	929	1,197	0	0
WRAP	0	1,634	1,579	0	343	0	500	614	393	343



Freeze 2: Harmonized Data with Sequence Data

Cohort	Autopsy	Vascular Risk Factors	Cognition	Fluid Biomarker	Diffusion Tensor Imaging (DTI)	Fluid Attenuated Inversion Recovery (FLAIR)	Magnetic Resonance Imaging (MRI) Freesurfer	Magnetic Resonance Imaging (MRI) MUSE	Positron Emission Tomography (PET) Amyloid	Positron Emission Tomography (PET) Tau
TOTAL	6,795	15,565	25,278	2,034	967	3,278	3,630	4,329	4,955	912
A4	0	0	3,373	0	0	963	0	0	3,378	345
ACT	532	0	1,337	0	0	0	0	0	0	0
ADNI	0	876	1,546	1,165	502	1,009	1,558	1,564	1,128	321
EFIGA	0	1,183	1,248	0	0	0	0	0	0	0
Knight ADRC	0	0	837	64	0	0	0	0	0	0
NACC	4,908	10,872	10,615	805	203	0	1,056	1,462	213	26
NIA-AD FBS	262	28	553	0	0	0	0	0	0	0
ROS/MAP/ MARS	1,093	1,254	1,271	0	77	0	0	165	0	0
WHICAP	0	604	3,786	0	0	1,306	712	770	0	0
WRAP	0	748	712	0	185	0	304	368	236	220
https://dss.n	iagads org/	1	1	1	1	1	1	Sample size	s reflect individu	als with ADSP

https://dss.niagads.org/

December 2023



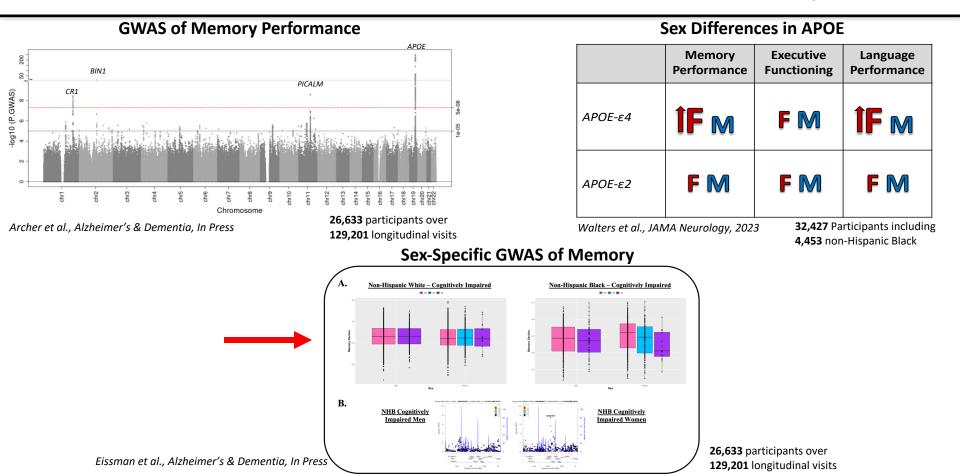


sequencing data, with exception to A4, which

will have sequenced data available in R5.

Exploring the Genetic Architecture of AD Leveraging Endophenotypes

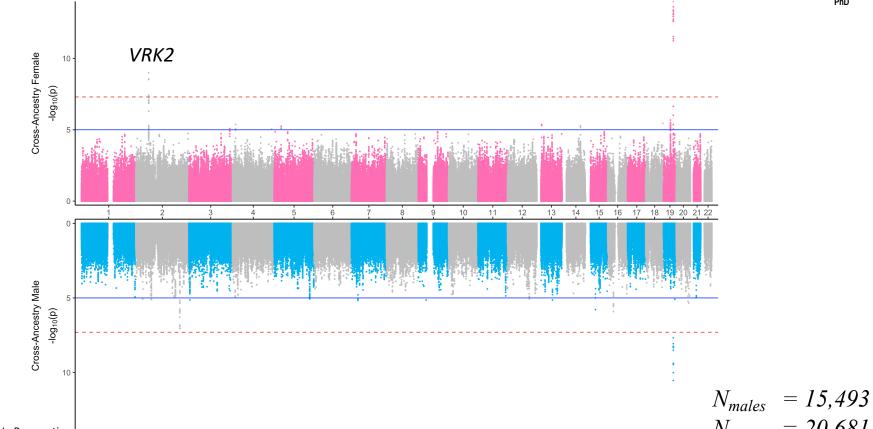
Explorations into the Genetic Architecture of Memory



Sex-Specific Genetic Drivers of Language



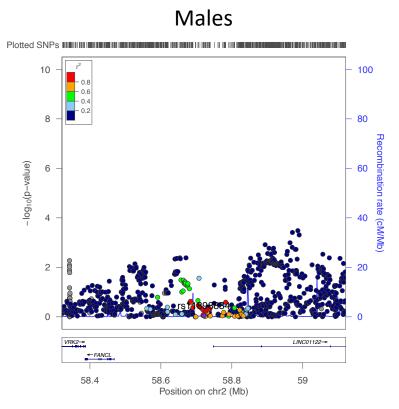




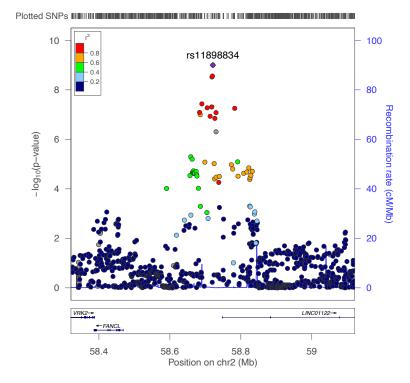
Eissman et al., In Preparation

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N_{females} = 20,681
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Sex Differences in VRK2 Association with Language

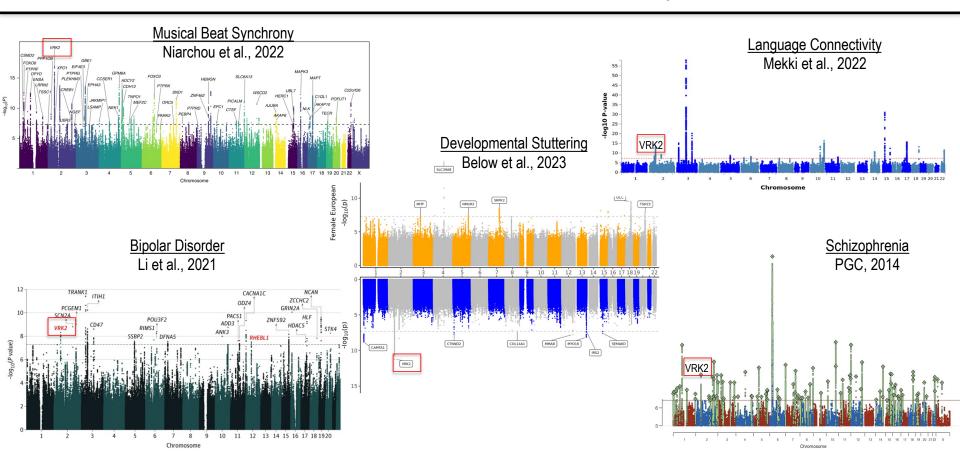


Females



Eissman et al., In Preparation

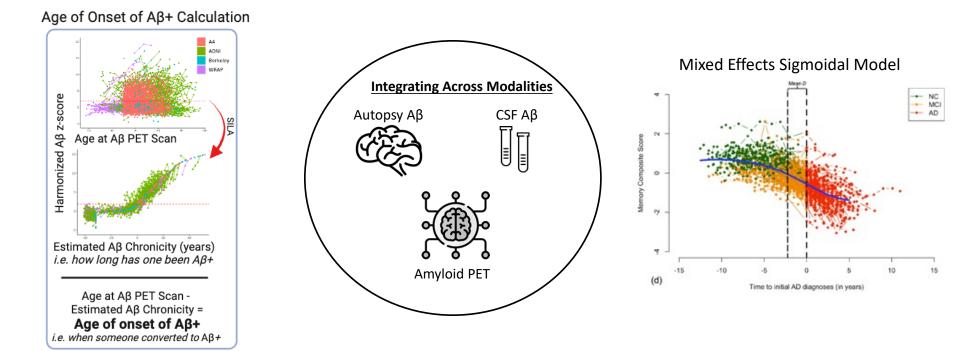
VRK2 Association with other Neuropsychiatric Traits



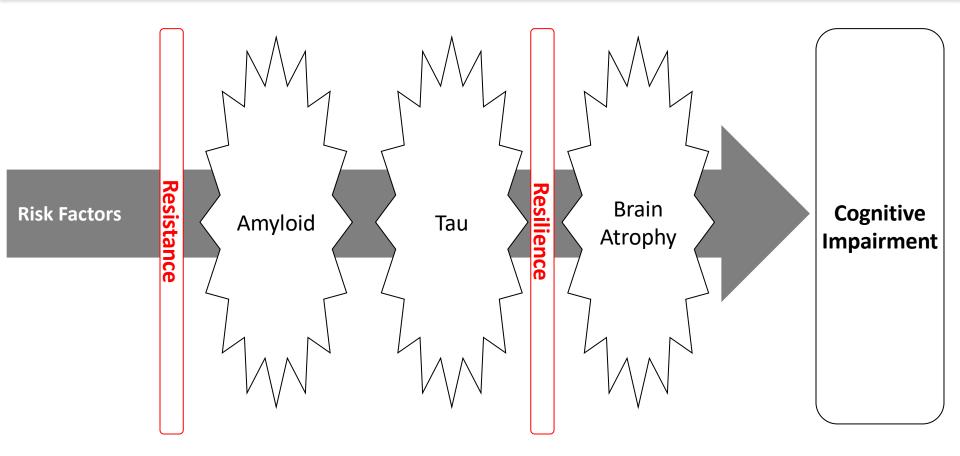
Integrative Analysis for More Precise Phenotypes

Integrative Analytics

Applying Advanced Algorithms to Increase Statistical Power



Resistance and Resilience



Approximating Resilience w/o Path Measures



Jared Phillips Graduate Student

Gold Model

Mixed Effects Model Predictors: Age, Sex, Race, APOE, Baseline Performance, Time and 3-way Interactions

Variance explained = 26%

Sample Size (3+ Visits) = 20,339

Age, Sex, Race, APOE, Pathology, Baseline

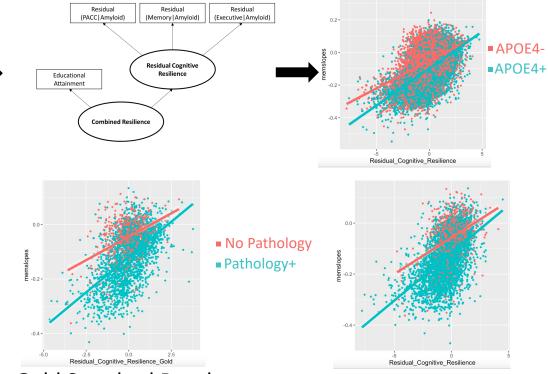
Variance explained = 55%

Performance, Time

and 3-way Interactions

Sample Size: 5,107

Mixed Effects Model Predictors:

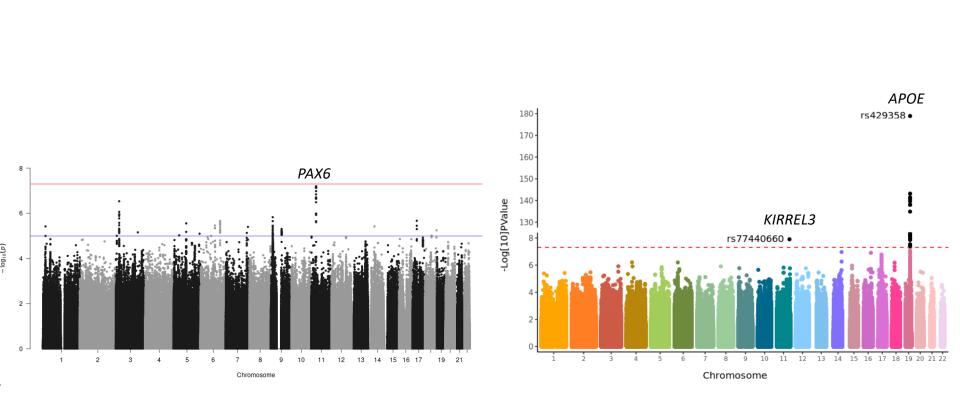


0.4 -

Gold Standard Results

Silver Standard Results

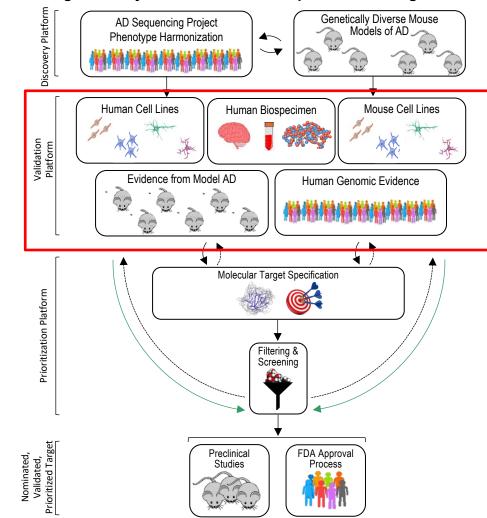
New Resilience and Resistance Loci



Phillips et al, In Preparation

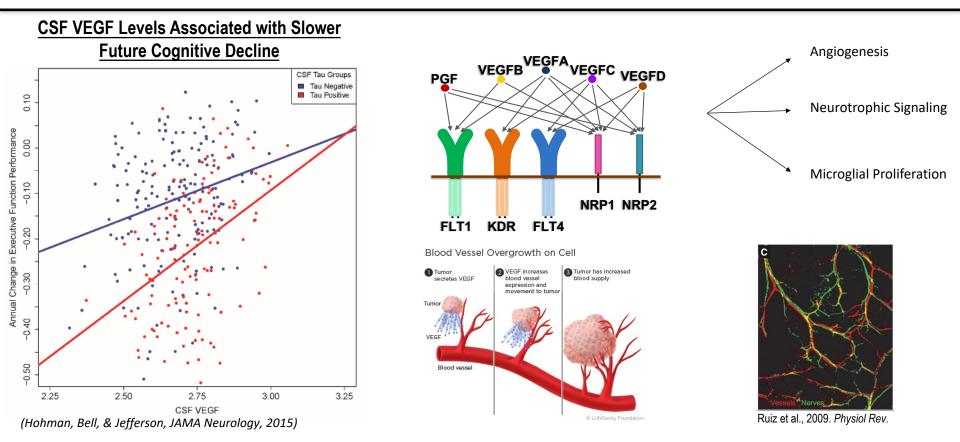
Castellano, ... Koran, In Preparation

Deep Molecular Characterization of High Quality Candidates



Drug Discovery Framework in the Computational Neurogenomics Team

Vascular Endothelial Growth Factor

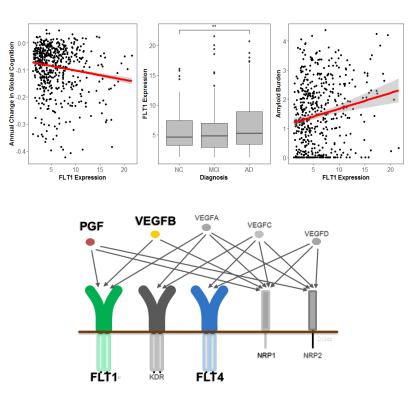


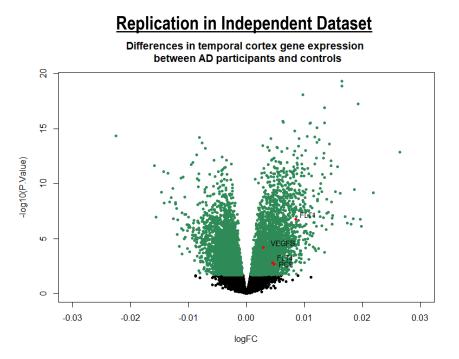
VEGF Transcript Association in the Brain



Emily Mahoney

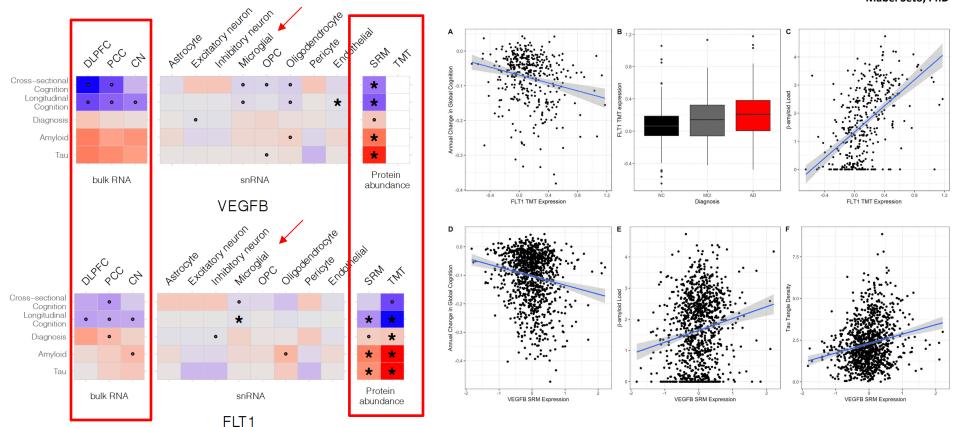
Brain Expression of Vascular Endothelial Growth Factor





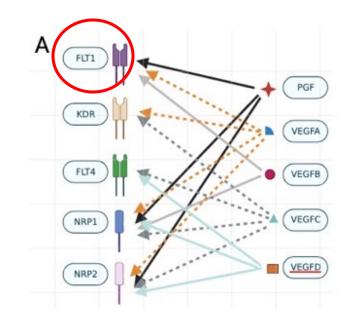
FLT1 and VEGFB Robustly Relate to AD outcomes

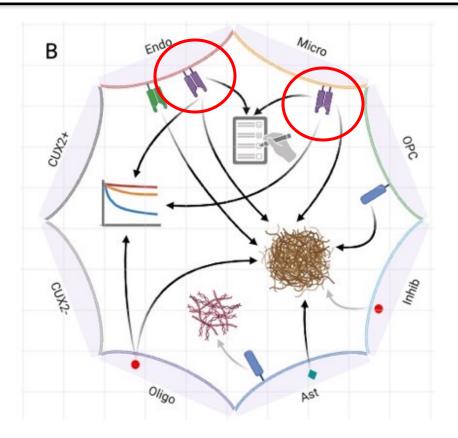
Mabel Seto, PhD



Seto et al., Neurobiology of Aging, 2023

Confirming snRNA Associations in Prefrontal Cortex





Sample size: 424

Post-translational Modifications

The FASEB Journal express article 10.1096/fj.03-0767fje. Published online March 4, 2004.

The c-Cbl/CD2AP complex regulates VEGF-induced endocytosis and degradation of Flt-1 (VEGFR-1)

Satsuki Kobayashi, *
. † Asako Sawano,* Yoshihisa Nojima, † Masabumi Shibuya,
* and Yoshiro Maru *,‡

PLOS ONE

🔓 OPEN ACCESS 🖻 PEER-REVIEWED

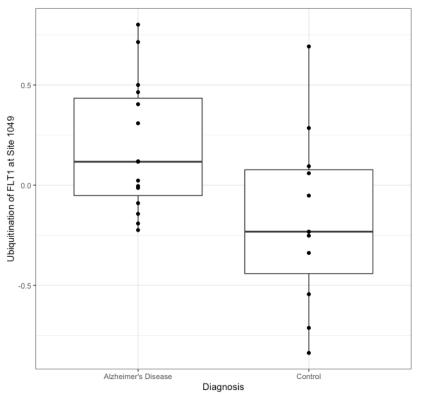
RESEARCH ARTICLE

N-Terminal Cleavage and Release of the Ectodomain of Flt1 Is Mediated via ADAM10 and ADAM 17 and Regulated by VEGFR2 and the Flt1 Intracellular Domain

Nandita S. Raikwar, Kang Z. Liu, Christie P. Thomas 🔤

Published: November 11, 2014 • https://doi.org/10.1371/journal.pone.0112794

FLT1 Ubiquitination is Upregulated in AD

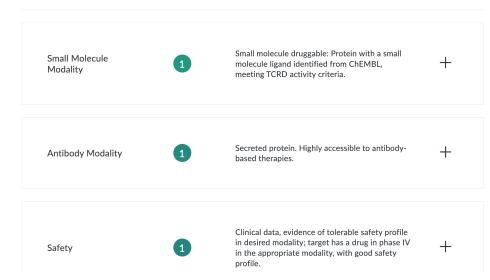


FLT1 is a Druggable Target

Druggability of FLT1

AMP-AD pharmaceutical partners scored the tractability of targets using publicly-available resources and have generated ratings of small-molecule druggability, therapeutic antibody feasibility, and safety.

i) Learn more about the analysis



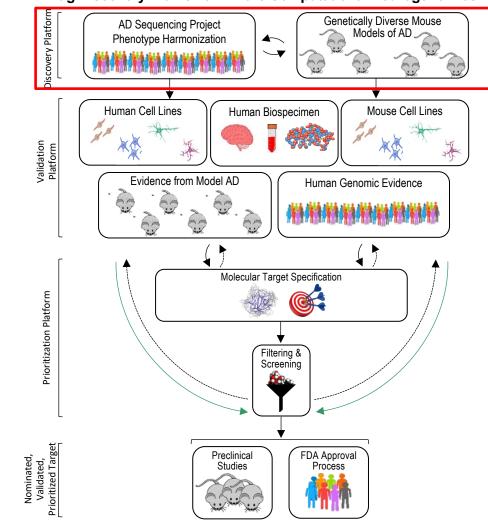


TREAT-AD

TaRget Enablement to Accelerate Therapy Development for AD

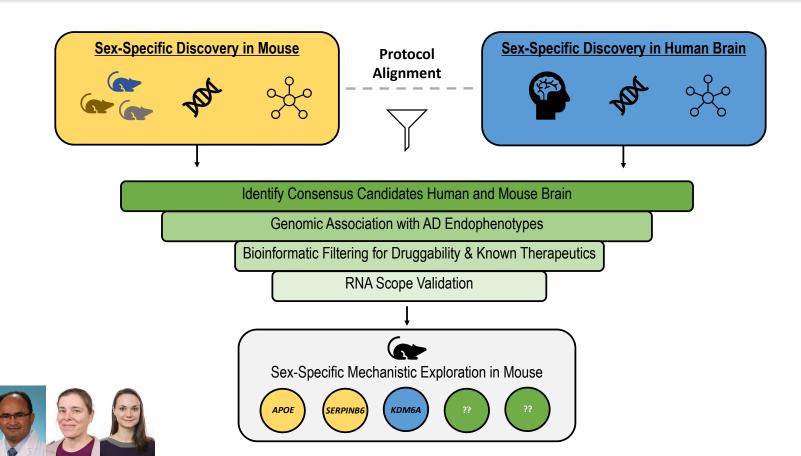


https://agora.adknowledgeportal.org/

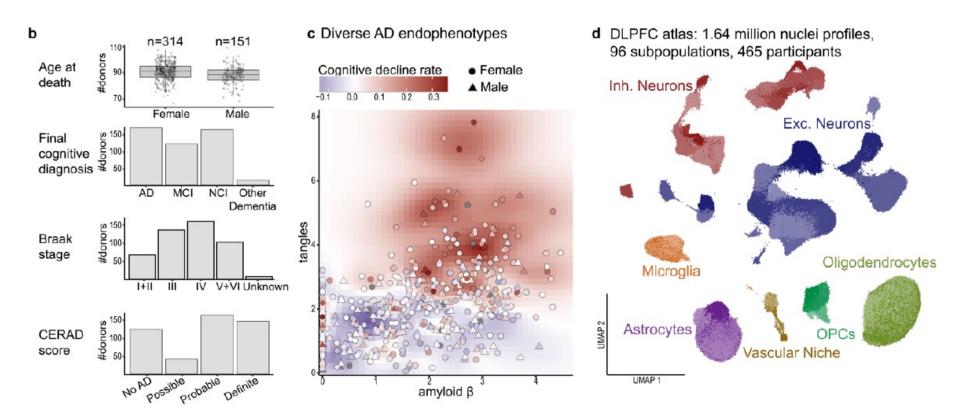


Drug Discovery Framework in the Computational Neurogenomics Team

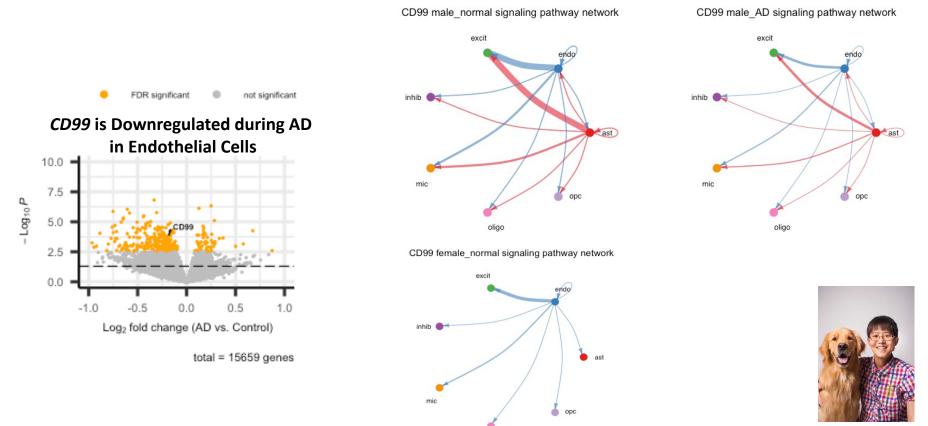
Collaborations to Move Targets Forward



Exploring Sex Differences with Single Cell Resolution



Striking Sex Differences in Immune Cells in AD



Yiyang Wu, MD, PhD

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