16th MCI Symposium, Special Topic Workshop and Forum

Imaging Genetics of APOE and Non-APOE factors in LOAD

Andrew J. Saykin, PsyD IU Center for Neuroimaging and the Indiana ADC Indiana University School of Medicine

January 20-21, 2018 • Miami Beach, Florida

DISCLOSURES

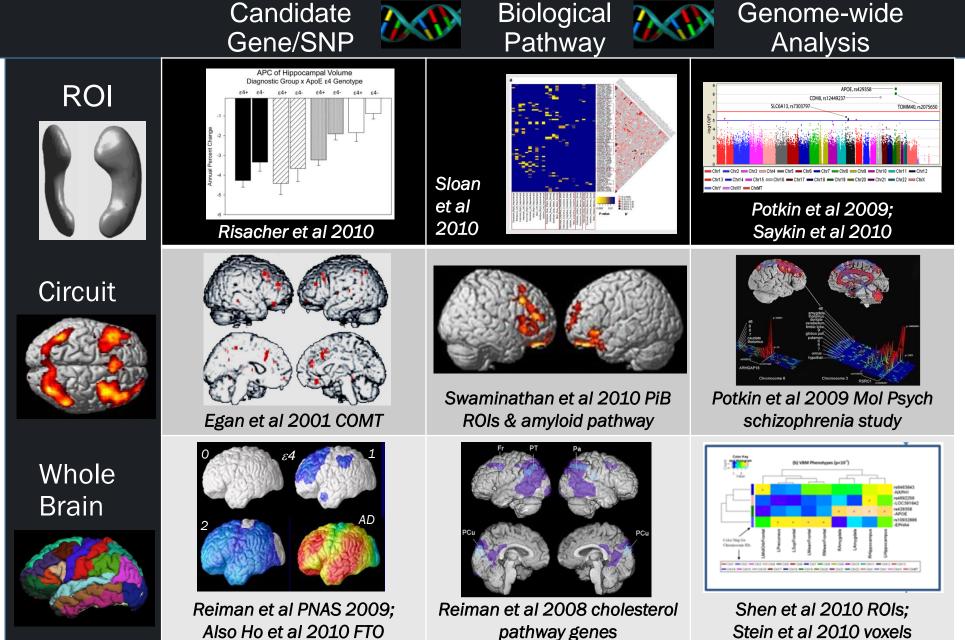
Disclosures

- Eli Lilly (Collaborative Grant), Arkley BioTek (SBIR), Avid Radiopharmaceuticals, Bayer
- Editor-in-Chief, Brain Imaging and Behavior, a Springer Nature journal

Grant Support

- National Institute on Aging P30 AG10133 (IADC)
 - ADNI: U01 AG024904 & RC2 AG036535; Indiana: R01 AG19771
 - U01 AG032984, U24 AG21886, P30 AG010129, K01 AG030514
- National Institute of Biomedical Imaging and Bioengineering
- National Library of Medicine: R01 LM011360 and K99/R00 LM011384
- National Science Foundation: IIS-1117335
- Foundation for the NIH (ADNI-1 GWAS, WGS, AMP-AD)
 - Anonymous Foundation (Challenge Grant), Gene Network Sciences, Merck, Pfizer
 - ADNI methylation project (AbbVie, Biogen and J&J, in-kind support)
- Alzheimer's Association & Brin Wojcicki Foundation Whole Genome Sequencing
- IUSM Strategic Research Initiative (SRI), Indiana Spinal Cord & Brain Injury Research Fund, CTSI

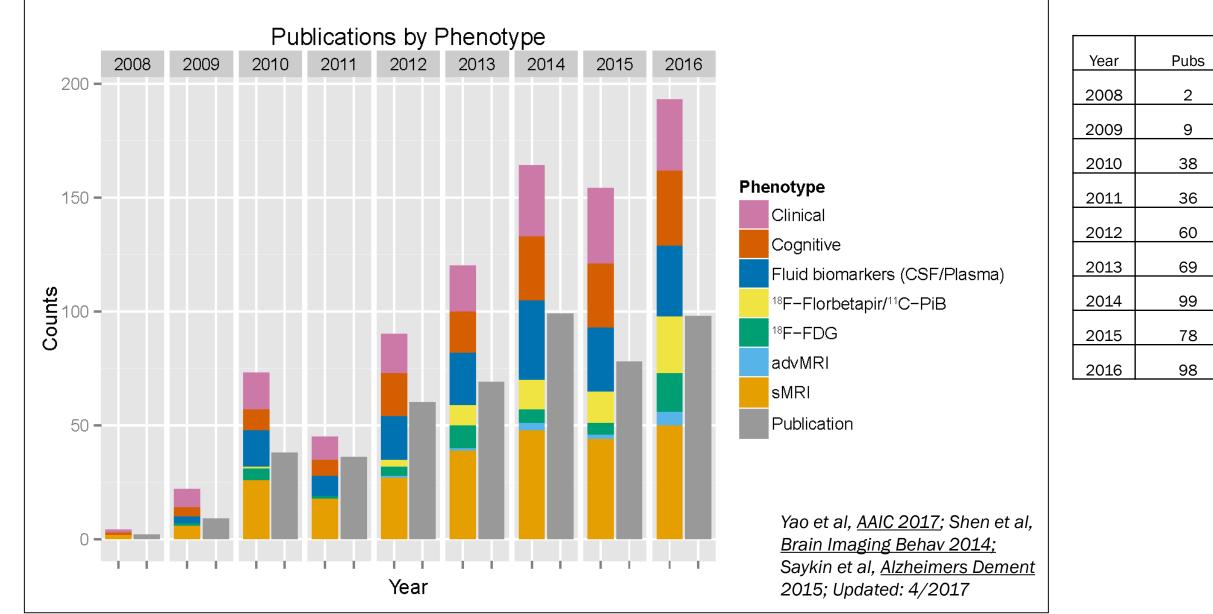
Brain-Genome Association Strategies: First Decade



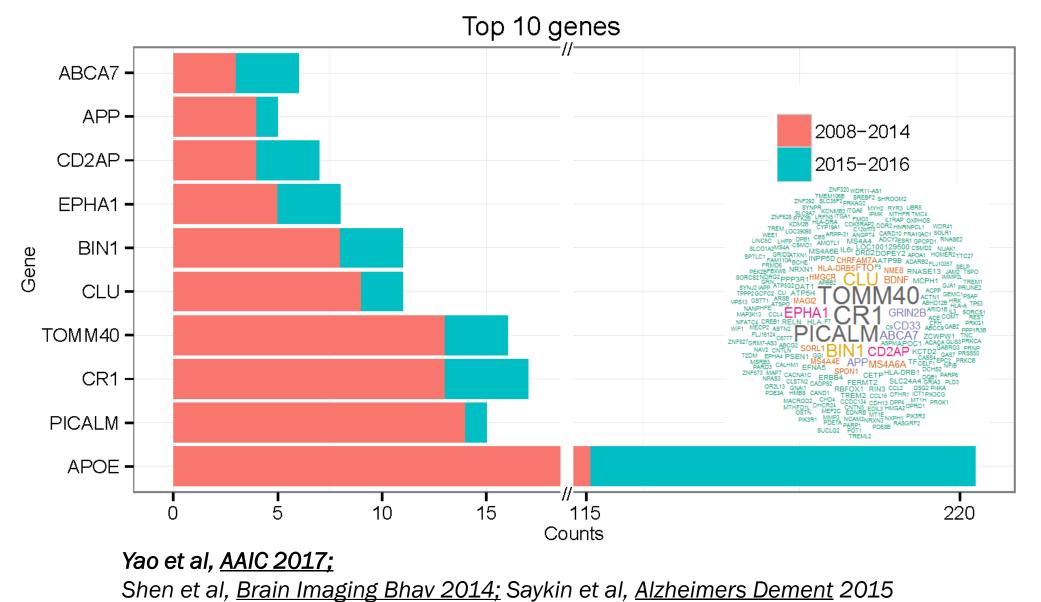
Saykin, 2011

Genetics studies Using ADNI Endophenotypes: Publications Using ADNI Genetic Data (2008–2016)





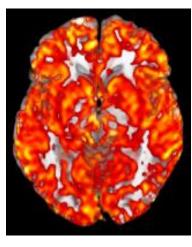
Most frequently reported genes in manuscripts using ADNI genetic data (2008–2016)



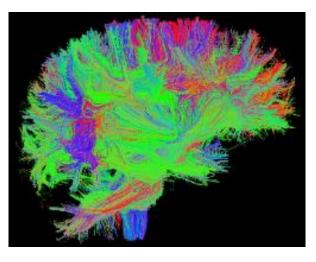
Updated: 4/2017

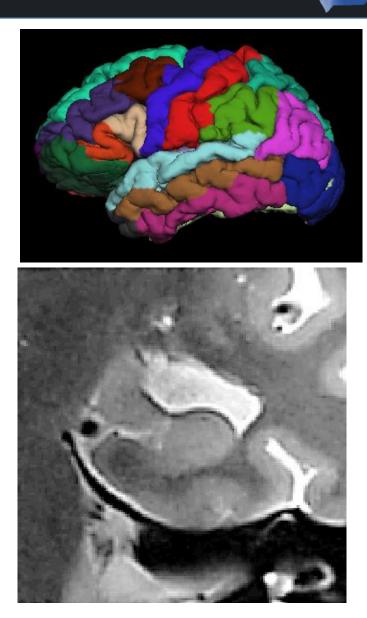
Multimodal MR Phenotypes

- MPRAGE, 3D FLAIR, SWI
- High Res. Hippocampal Sequence
- Diffusion MR (DTI and NODDI)
- fMRI: Resting State & Task-based BOLD
- Arterial Spin Labeled Perfusion









NODDI: neurite orientation dispersion and density

Connectome as Endophenotype

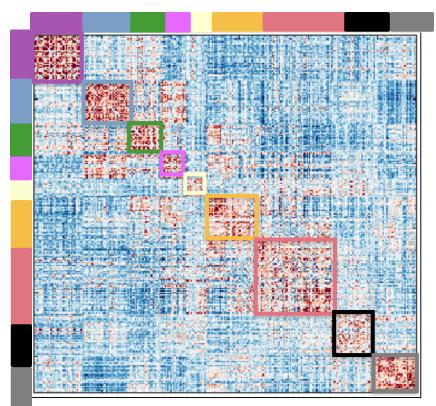
Resting State Networks

Somatomotor Dorsal Attn Ventral Attn Limbic Frontoparietal Default Mode

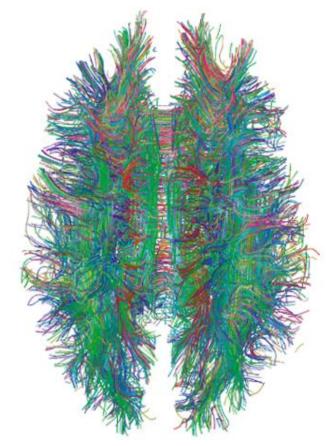
Visual

Subcortical

Cerebellum



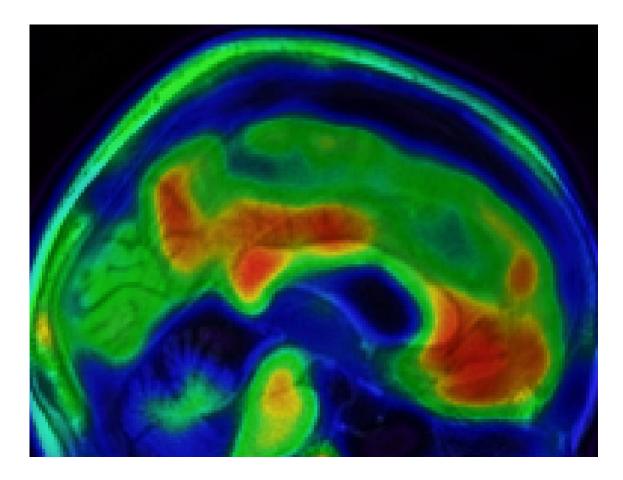
Structural Connectome

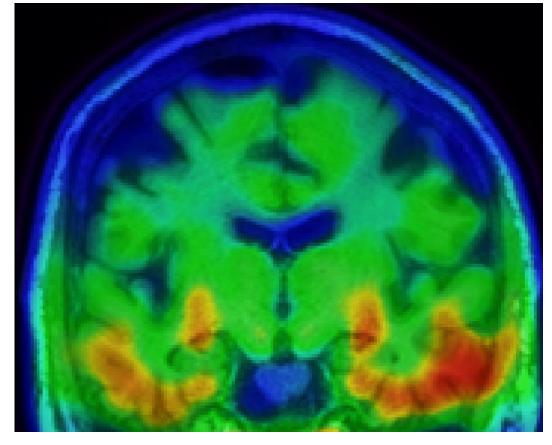


Contreras et al. (2017) Alz & Dementia

Gigandet, Hagmann et al (2008)

Molecular Imaging Phenotypes



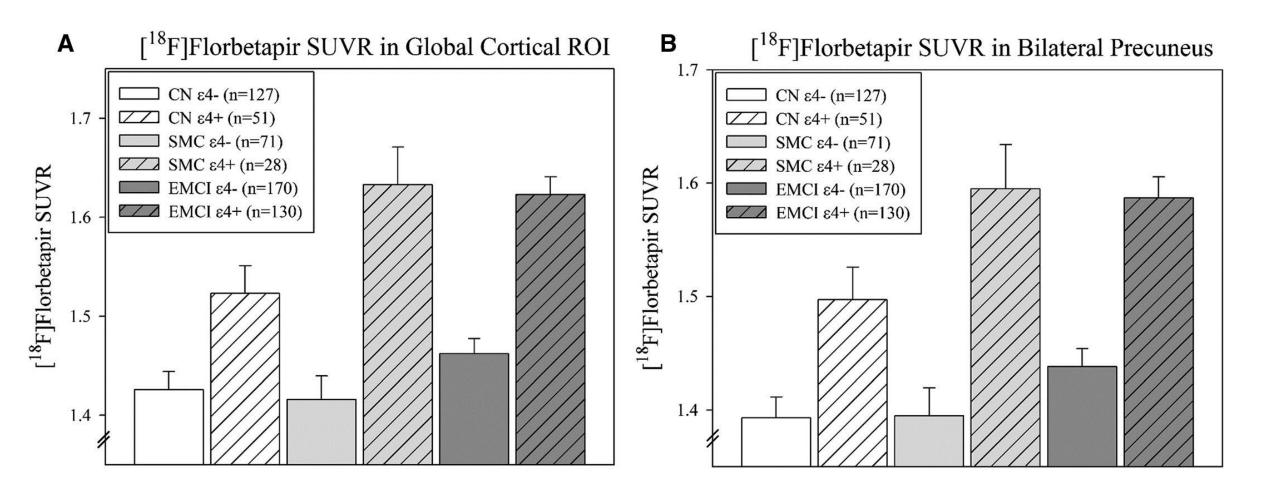


Tau PET [18F]AV-1451

Indiana ADC

Amyloid PET [¹⁸F]Florbetapir

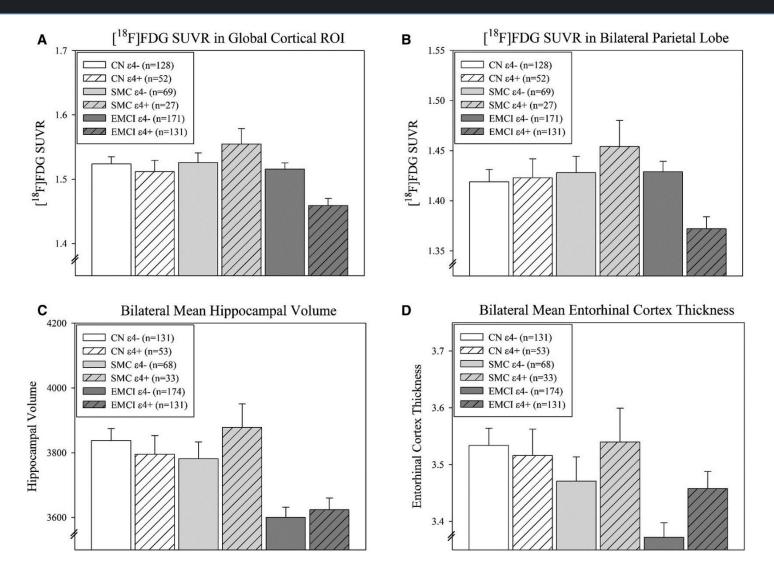
APOE ε4 Status and Early Stage Amyloid Deposition on PET



Risacher et al. Alzheimer's & Dementia (2015): DOI: (10.1016/j.jalz.2015.03.003)



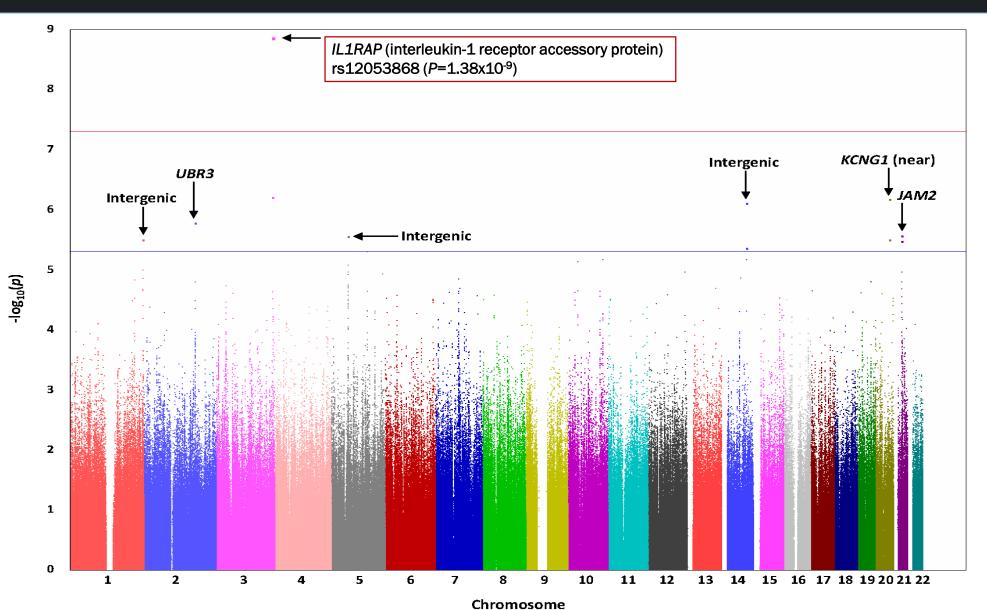
APOE ε4 Status: Early Stage Atrophy and Glucose Metabolism



Risacher et al. Alzheimer's & Dementia (2015): DOI: (10.1016/j.jalz.2015.03.003)

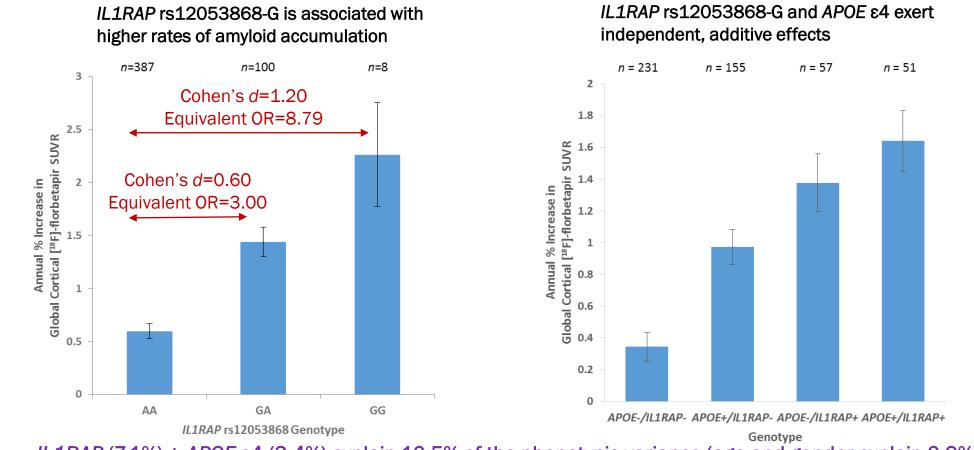


GWAS of Longitudinal Amyloid PET: IL1RAP Example



Ramanan et al, BRAIN, 2015 Oct;138 (Pt 10):3076-3088.

Combined Effects of IL1RAP rs12053868 and APOE



-IL1RAP (7.1%) + APOE ε4 (3.4%) explain 10.5% of the phenotypic variance (age and gender explain 0.9%)

-IL1RAP association remains genome-wide significant (P=5.80x10⁻⁹) with additional covariates of APOE ε4 status, baseline diagnosis, education, baseline amyloid burden and its square, and PCA eigenvectors

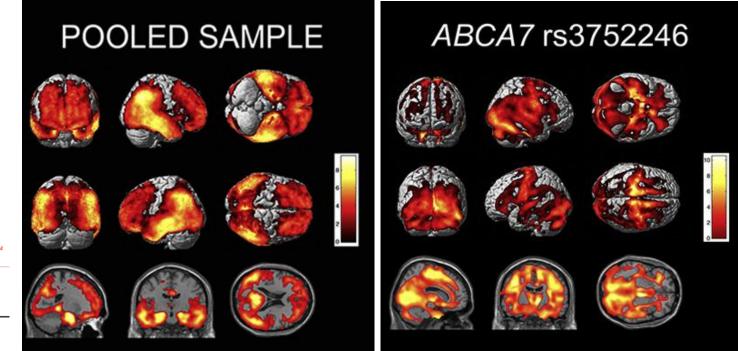
Ramanan et al, BRAIN, 2015 Oct;138(Pt 10):3076-3088.

IGAP – Top AD Candidates: Imaging Associations

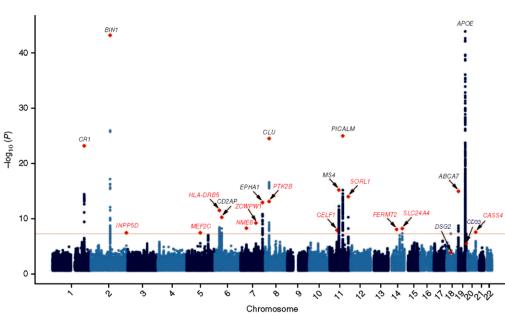
LETTERS

APOE effect: Multivariate analysis of 27 variants in the pooled ADNI sample

Disease stage specific effects in ADNI: *ABCA7* association in AD group only



Stage et al (2016) Alz & Dem: DADM



Meta-analysis of 74,046 individuals identifies 11 new

susceptibility loci for Alzheimer's disease

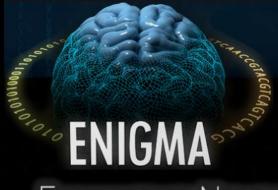
genetics

Lambert et al Nature Genetics (2013)

ENIGMA Consortium

Paul Thompson et al (http://enigma.ini.usc.edu/)

USC University of Southern California



ABOUT ENIG	MA V	Vorking Groups	Publications	Research	Press	EnigmaVis	
Protocols	Events	TRAINING	JOIN ENIGMA	JOIN OUR MAI	LING LIST!		

Enhancing Neuro Imaging Genetics through Meta Analysis

Disease Healthy Variation

Genomics

Algorithm Development

Collaborations

What is **ENIGMA**?

The ENIGMA Network brings together researchers in imaging genomics to understand brain structure, function, and disease, based on brain imaging and genetic data. We welcome brain researchers, imagers, geneticists, methods developers, and others interested in cracking the neuro-genetic code!

The ENIGMA Network has several goals:

- O To create a network of like-minded individuals, interested in pushing forward the field of imaging genetics.
- 2 To ensure promising findings are replicated via member collaborations.
- O To share ideas, algorithms, data, and information on research studies and methods.
- O To facilitate training, including workshops and conferences on key methods and emerging directions in imaging genetics.



Selected studies & data sets:

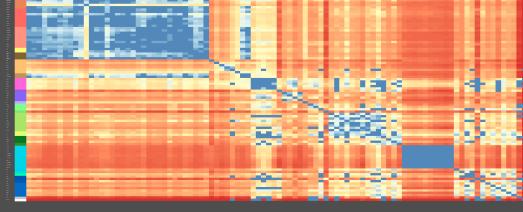
• GWAS of subcortical volumes (ENIGMA2) (<u>Hibar et</u> al., *Nature*, 2015)

• GWAS of intracranial volume (<u>Adams et al., Nature</u> <u>Neuroscience, 2016</u>) from the ENIGMA-CHARGE collaboration

• GWAS of hippocampal volume (<u>Hibar et al., *Nature*</u> <u>*Communications*, 2017</u>) from the ENIGMA-CHARGE collaboration

Allen Atlas: 3D Gene Expression Mapping

Genetic architecture of the human brain



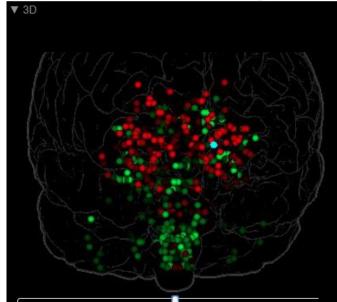
CELL TYPES CELLULAR TAXONOMY OF THE VISUAL CORTEX

http://www.brain-map.org/

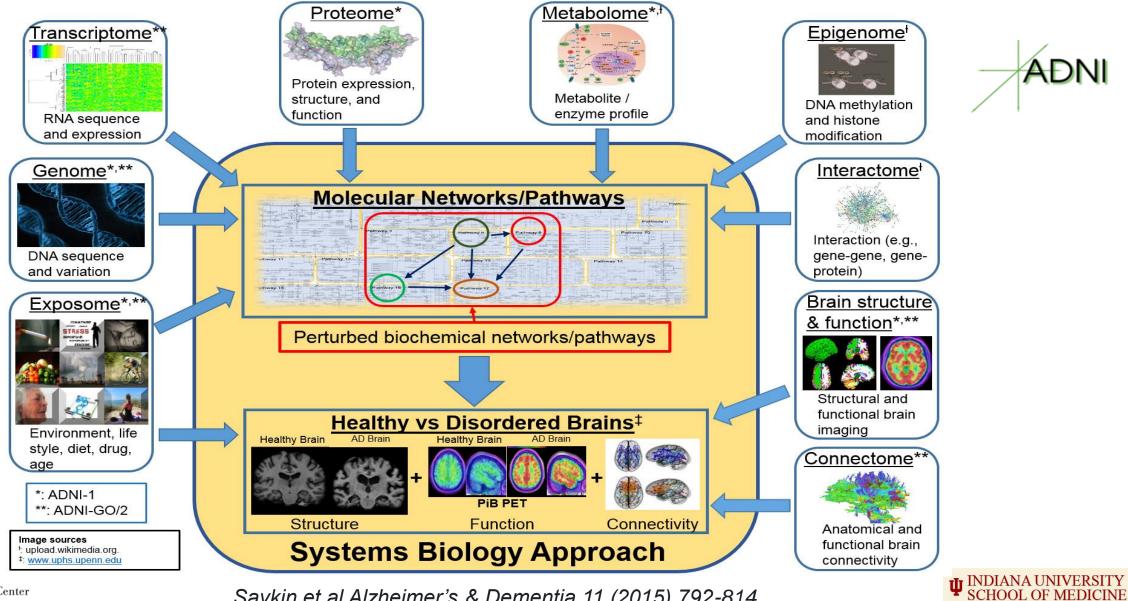
Example of ABCA7 expression --->



Example of APOE expression



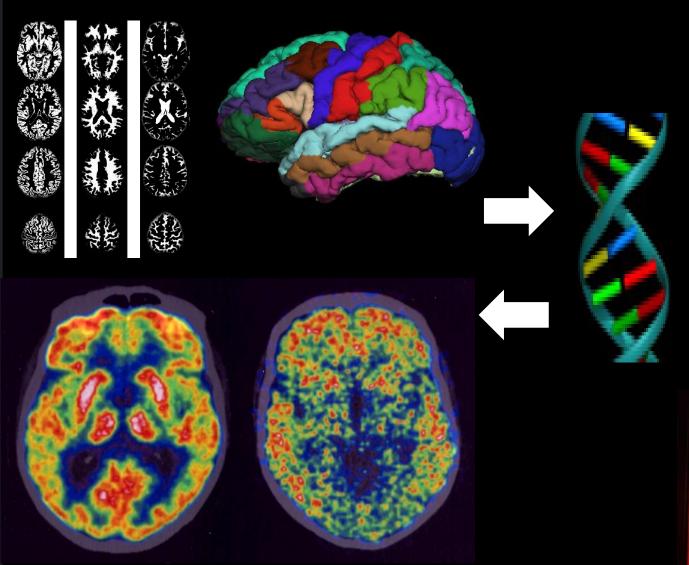
Working Toward a Systems Biology of AD



Neuroscience Center

Saykin et al Alzheimer's & Dementia 11 (2015) 792-814

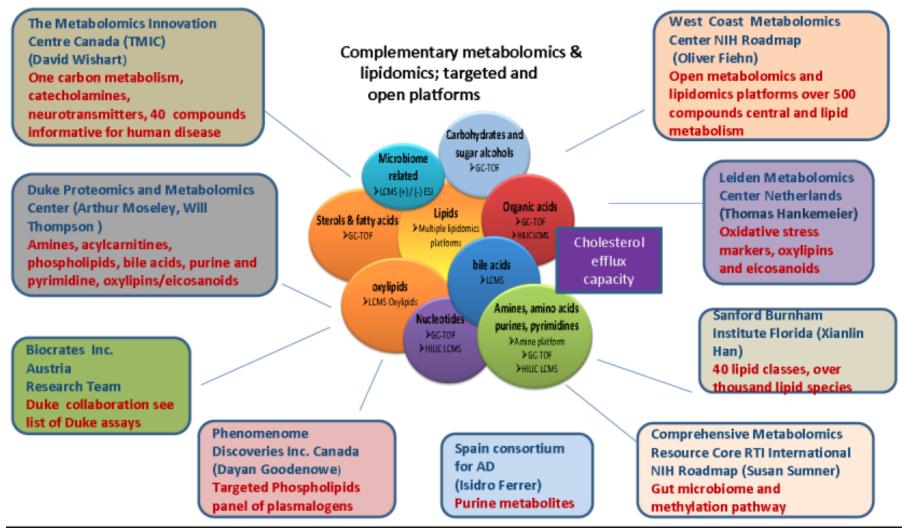
Relating Imaging Phenotypes, Genes & Clinical Information is a <u>BIG DATA</u> Challenge







AD Metabolomics Consortium (ADMC)



PI: Rima Kaddurah-Daouk, Duke University

https://sites.duke.edu/adnimetab/about-us/participating-centers/

Defining Metabolomic Networks in AD

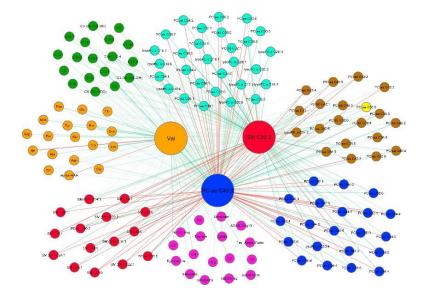


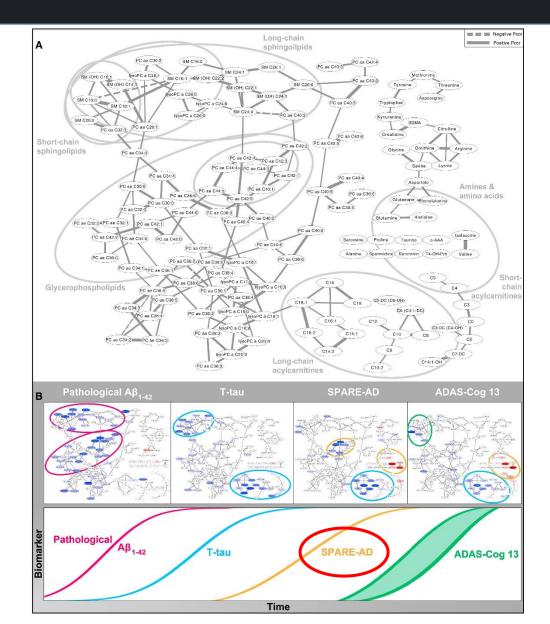
Alzheimer's بئ Dementia

Alzheimer's & Dementia 📕 (2017) 1-20

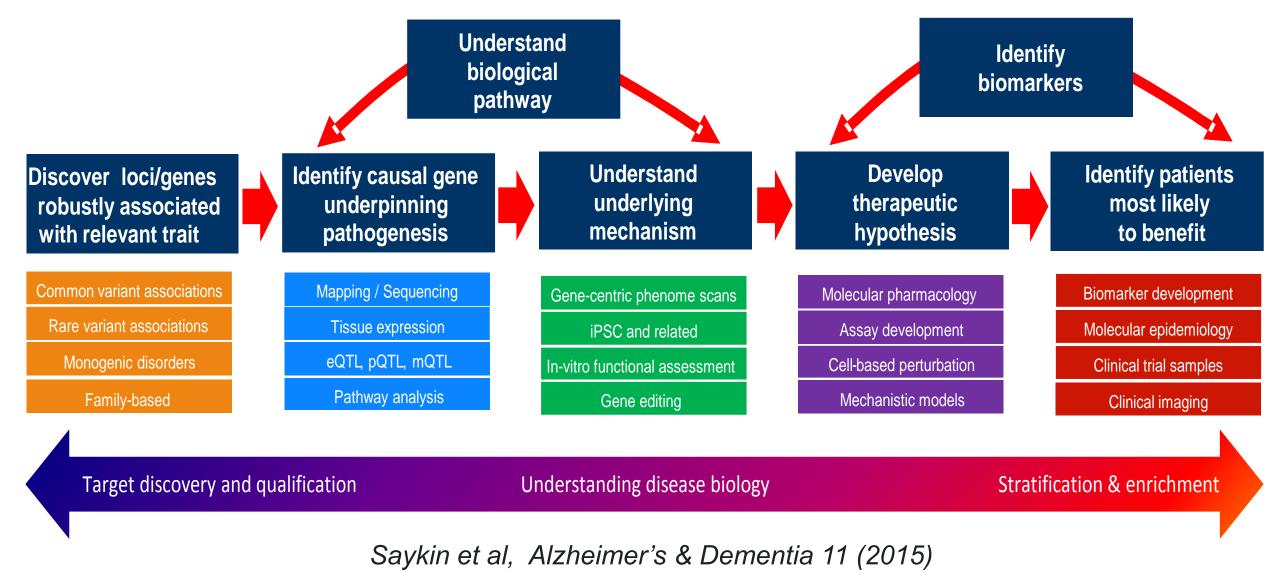
Featured Article Metabolic network failures in Alzheimer's disease—A biochemical road map

Jon B. Toledo^{a,b,***}, Matthias Arnold^c, Gabi Kastenmüller^{c,d}, Rui Chang^e, Rebecca A. Baillie^f, Xianlin Han^g, Madhav Thambisetty^h, Jessica D. Tenenbaum^j, Karsten Suhre^{c,k},
J. Will Thompson¹, Lisa St. John-Williams¹, Siamak MahmoudianDehkordi^m, Daniel M. Rotroff^m, John R. Jack^m, Alison Motsinger-Reif^m, Shannon L. Risacher^{n,o}, Colette Blach^{i,j},
Joseph E. Lucas^p, Tyler Massaro^p, Gregory Louie^{q,r}, Hongjie Zhu^{q,r}, Guido Dallmann^s, Kristaps Klavins^s, Therese Koal^s, Sungeun Kim^{n,o}, Kwangsik Nho^{n,o}, Li Shen^{n,o}, Ramon Casanova^h, Sudhir Varma^h, Cristina Legido-Quigley^t, M. Arthur Moseley¹,
Kuixi Zhu^e, Marc Y. R. Henrion^e, Sven J. van der Lee^u, Amy C. Harms^v, Ayse Demirkan^u, Thomas Hankemeier^{u,v}, Cornelia M. van Duijn^{u,v}, John Q. Trojanowski^a,
Leslie M. Shaw^a, Andrew J. Saykin^{n,o}, Michael W. Weiner^w, P. Murali Doraiswamy^{q,r},
Rima Kaddurah-Daouk^{q,r,x,*}, for the Alzheimer's Disease Neuroimaging Initiative and the Alzheimer Disease Metabolomics Consortium¹





Path from genetic signal to targeted therapeutics



702 011

Molecular Validation & Therapeutics: New Models

Model Organism Development and Evaluation for Late-onset Alzheimer's Disease (MODEL-AD)

Contact PI: Bruce Lamb

ADNI contributes target nominations & characterization MODEL-AD is creating organisms based on ADNI reports



Website: https://Model-AD.org

Contact: ModelAD@iupui.edu

Data: https://www.synapse.org/#!Synapse:syn2580853/wiki/409840

Thanks to IADC, IU Neuroscience, ADNI & AMP-AD





Indiana Alzheimer Disease Center





Directions & Challenges in Imaging Genomics

- Structural, functional & molecular imaging as endophenotypes
 - Connectome as QTL
- Integration of polygenic scoring approaches with endophenotypes
- Epigenetic changes and longitudinal studies
- Network approaches to multi-layered -omics and imaging data
- Connecting peripheral metabolic changes in targeted biochemical pathways to brain structure and function
- Use of imaging genetics and related phenotypes to enrich clinical trials
- Resolving heterogeneity by integrating genetics, other –omics and imaging biomarker profiles → precision medicine of AD & related disorders
- Target discovery and validation, informing model system development and assessment of therapeutic strategies