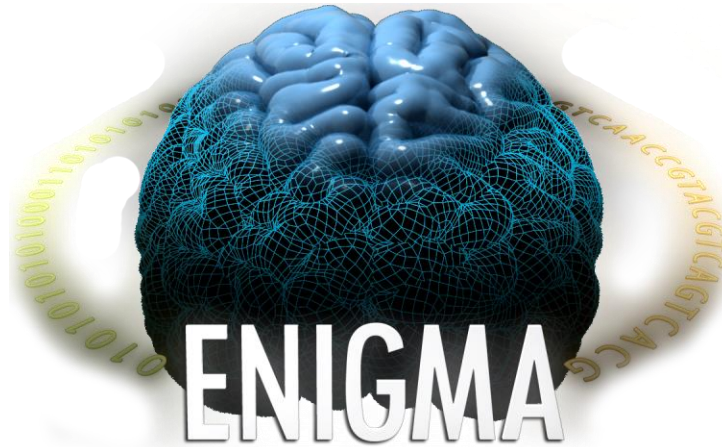

The ENIGMA project



Martine Hoogman, PhD
jPI, Human Genetics, Radboudumc

Enhancing Neuro Imaging Genetics through Meta-Analysis



PI: Paul Thompson (USC)

The ENIGMA Network brings together researchers in imaging genomics to understand brain structure, function, and disease, based on brain imaging and genetic data.

ENIGMA

Enhancing Neuro Imaging Genetics through Meta-Analysis

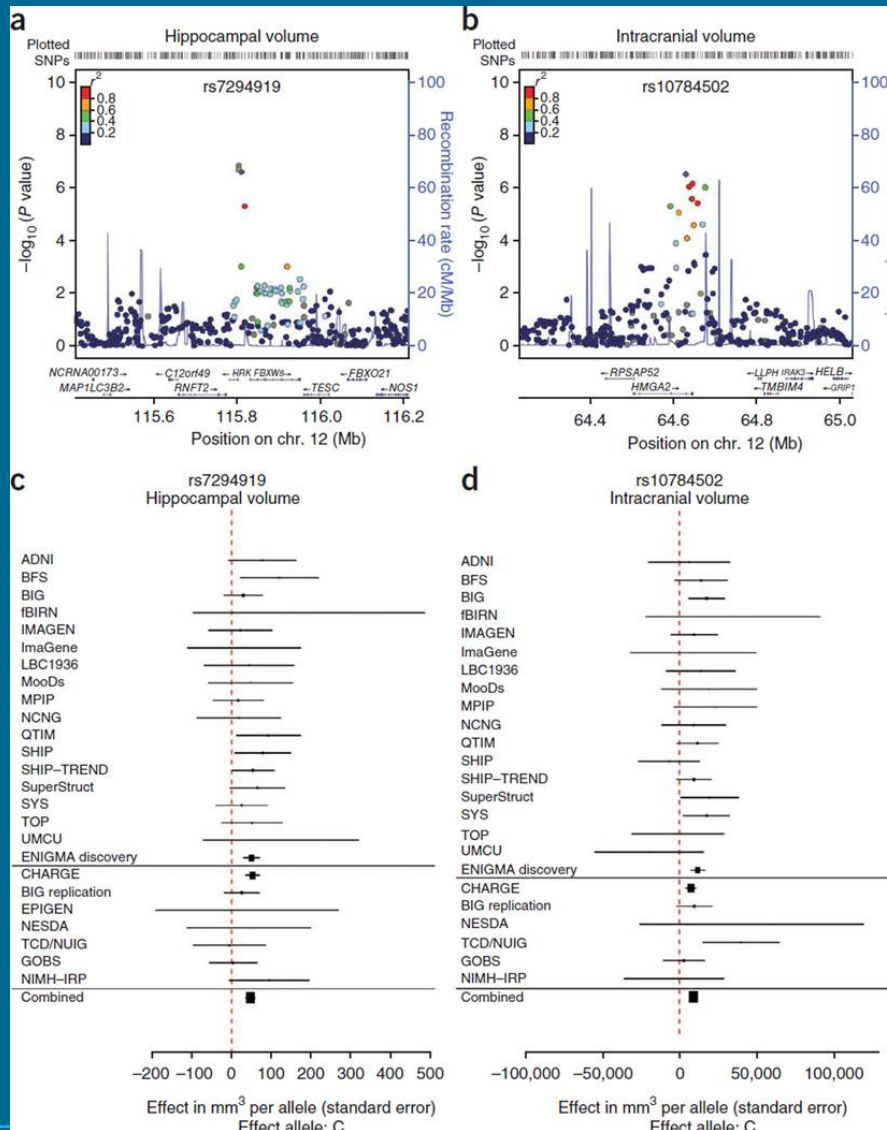


How it works:

- Making use of already collected brain and genetic data
- Using harmonized methods to analyze all raw data again
- Share summary stats or individual level data
- Collaborating to increase power
- Share expertise

ENIGMA

First success: Stein et al, 2012, Nature Genetics

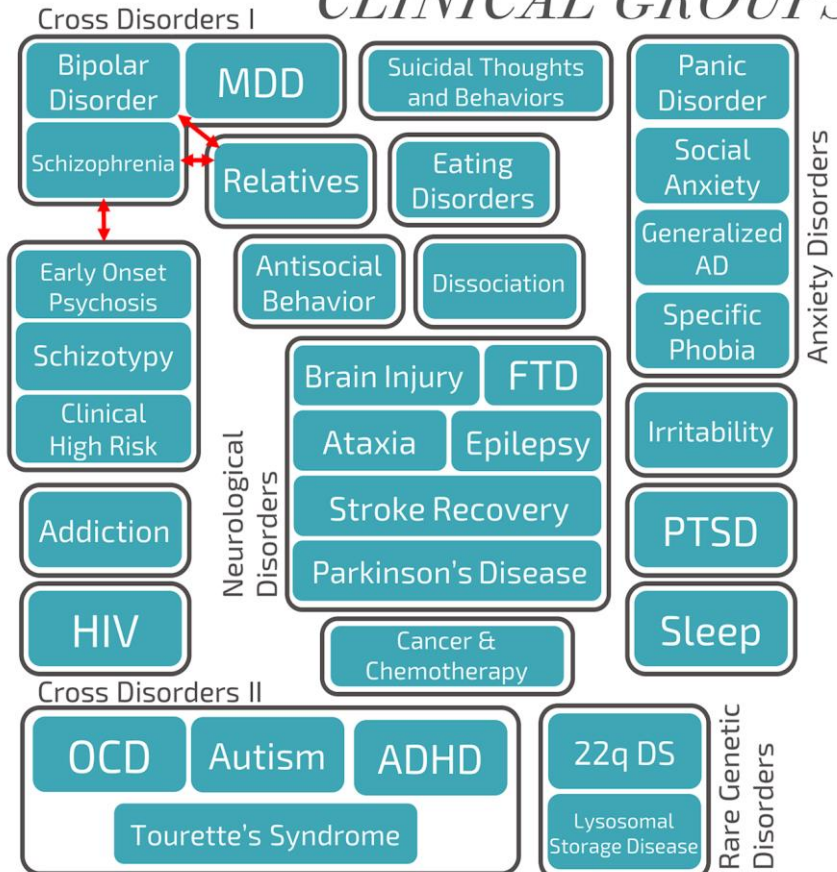


N = 21,151

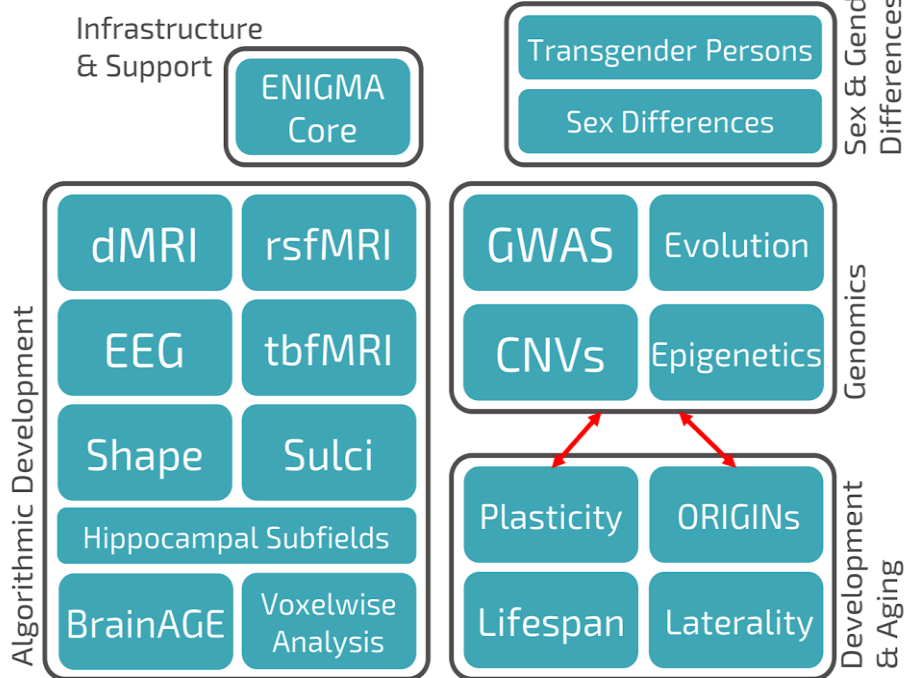
Working Groups: Chart Overview

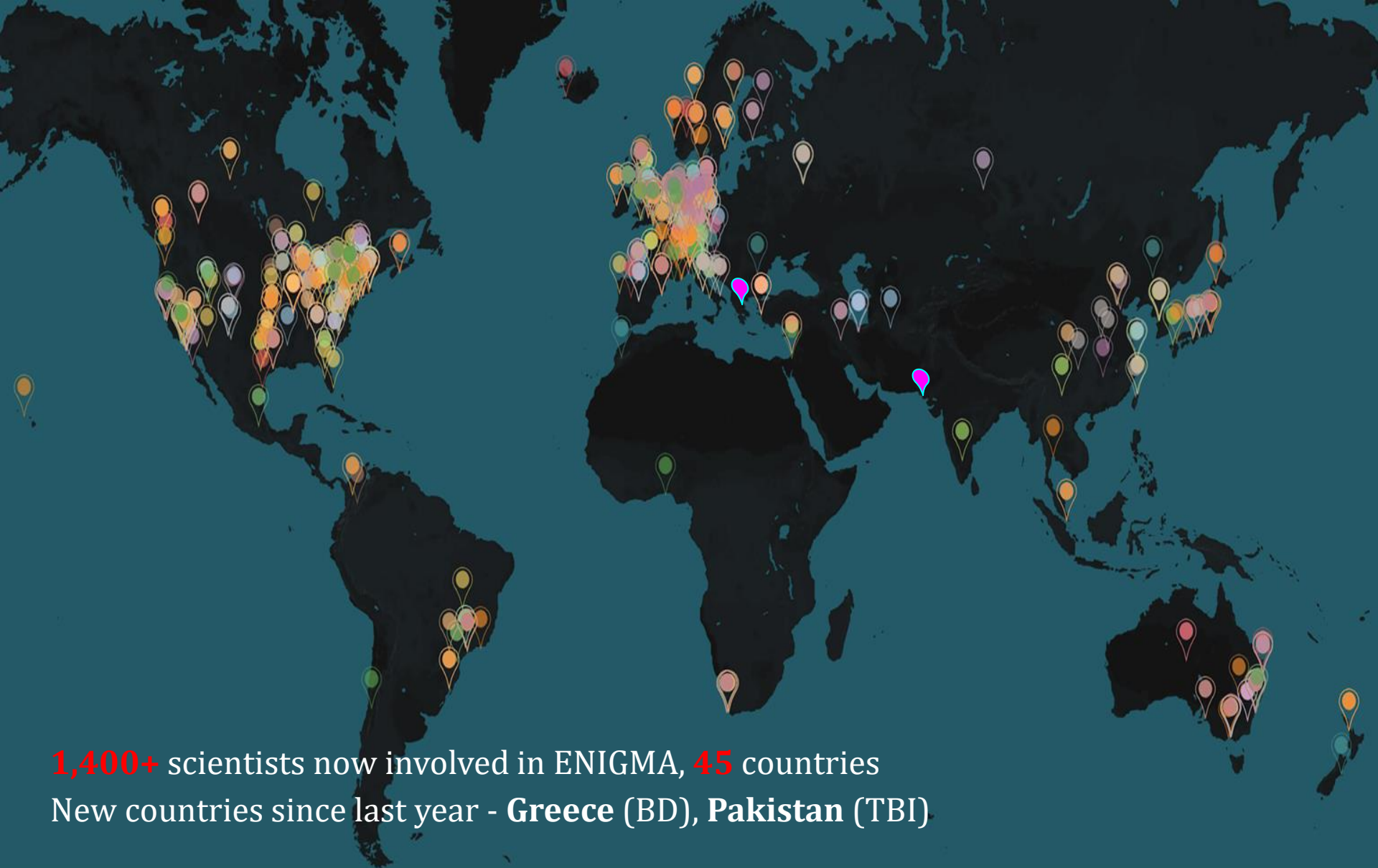
50 total working groups →
30 clinical, **20** non-clinical

CLINICAL GROUPS



NON-CLINICAL GROUPS

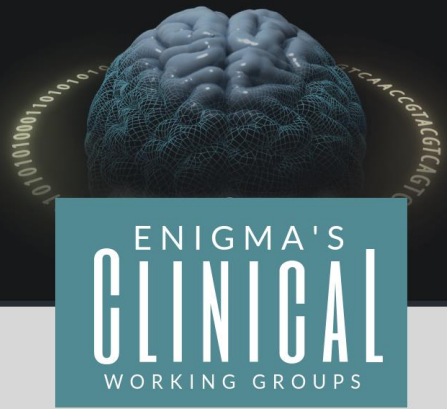




1,400+ scientists now involved in ENIGMA, **45** countries
 New countries since last year - **Greece (BD)**, **Pakistan (TBI)**

Primary Working Group	
SCZ	Ataxia
Addiction	Brain Injury
22q Deletion Syndrome	BrainAGE
MDD	Clinical High Risk
BD	ASD
ADHD	Anxiety
Core	CNVs
DTI	Eating Disorders
Early Onset Psychosis	EEG
Evolution	Frontotemporal Dementia
Epigenetics	GWAS
Epilepsy	Hippocampal subfields
Evolution	HIV
Frontotemporal Dementia	Irritability
GWAS	Lateralization
Hippocampal subfields	Lifespan
HIV	OCD
Irritability	ORIGInS
Lateralization	Parkinson's
Lifespan	Plasticity
OCD	PTSD
ORIGInS	Relatives
Parkinson's	rsfMRI
Plasticity	Schizotypy
PTSD	Sleep Disorders
Relatives	Storage Disease
rsfMRI	Suicidal Thoughts & Behaviors
Schizotypy	Sulci
Sleep Disorders	Tourette's Syndrome
Storage Disease	Transgender Persons
Suicidal Thoughts & Behaviors	
Sulci	
Tourette's Syndrome	
Transgender Persons	

Nijmegen researchers



9 LARGEST
NEUROIMAGING STUDIES
IN THE FOLLOWING CLINICAL POPULATIONS

10,105

Major Depression
Schmaal et al 2016

6,503

Bipolar Disorder
Hibar et al 2016

3,665

OCD
Boedhoe et al 2018

1,868

PTSD
Logue et al 2018

789

22q DS
Sun et al 2018

9,572

Schizophrenia
van Erp et al 2018

3,876

Epilepsy
Whelan et al 2018

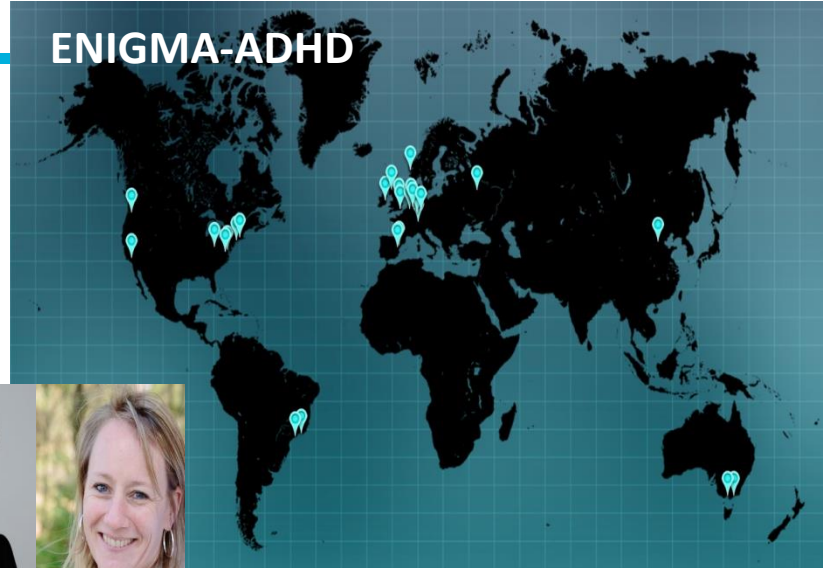
3,242

ADHD
Hoogman et al 2017

3,222

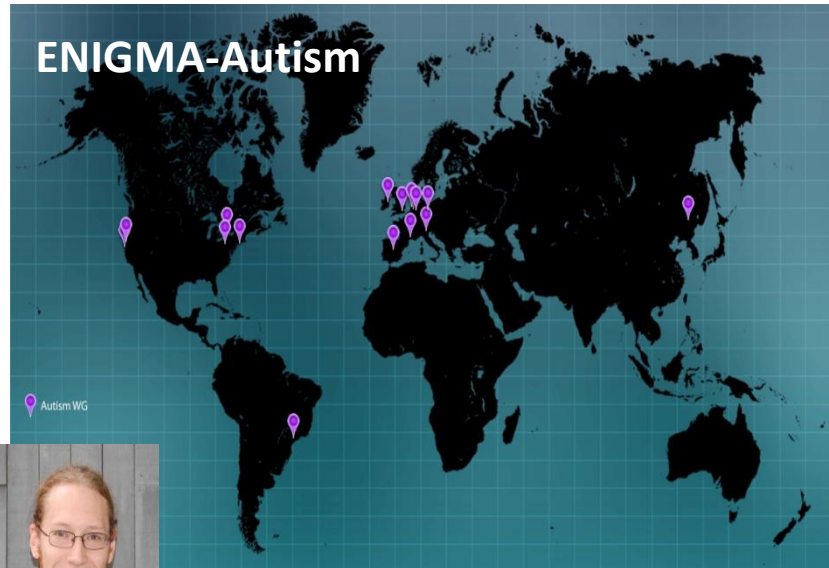
Autism Spectrum
Disorder
van Rooij et al 2018

ENIGMA-ADHD



Barbara Franke & Martine Hoogman

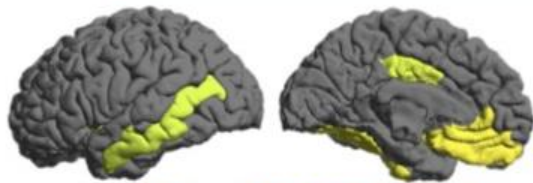
ENIGMA-Autism



Jan Buitelaar
& Daan van Rooij

Radboudumc

Comparing results becomes easy!

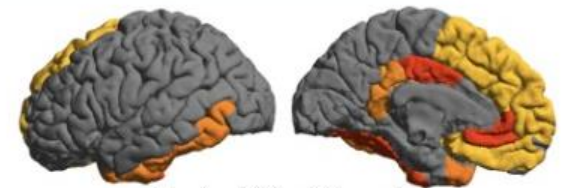


MDD: [Schmaal 2017 Mol Psych](#)



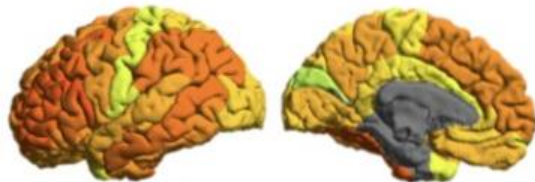
OCD

[Boedhoe 2017 Am J Psych](#)



Alcohol Use Disorder:

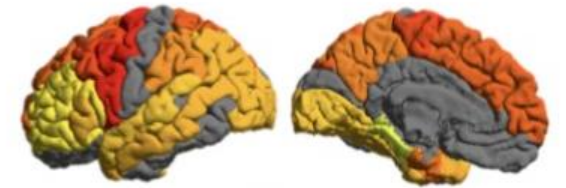
[Mackey 2018 Am J Psych](#)



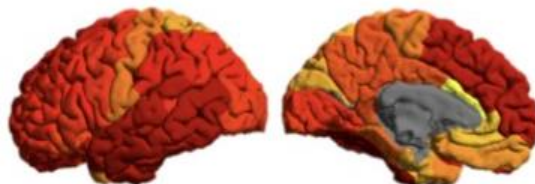
BD: [Hibar 2018 Mol Psych](#)



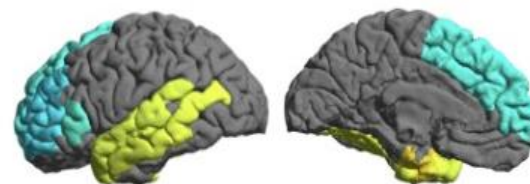
ADHD: [Hoogman 2019 Am J Psych](#)



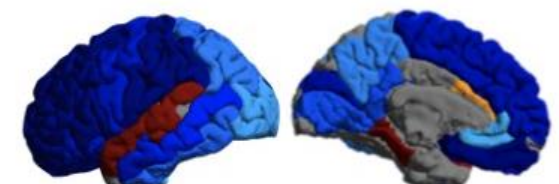
Epilepsy: [Whelan 2018 Brain](#)



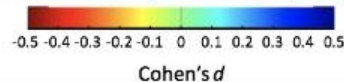
SCZ: [van Erp 2018 Biol Psych](#)



ASD: [van Rooij 2017 Am J Psych](#)



22q: [Sun 2019 Mol Psych](#)



Advantages

- Sample size and power
- Collaborating is fun and easy
- Life-span perspective (broad age range)
- Harmonized segmentation procedures and protocols for QC and analysis

Disadvantages

- Old data
- Collected with various aims (inclusion criteria etc)
- Heterogeneity of other phenotypes of interest (example: ADHD severity measures collected with different instruments)
- Cross-sectional data
- (Submission systems/journals still don't understand consortia papers...)

Access to the data via proposal form to PI's of WG, more information: <http://enigma.ini.usc.edu/>