



Assessing individual differences with fMRI

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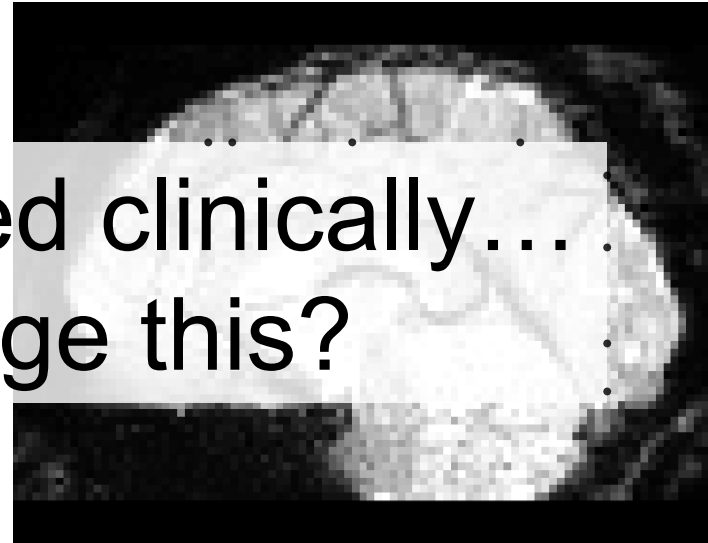
NIH Summer Neuroimaging Course

June 29, 2018

Structural MRI



Functional MRI)



fMRI is barely used clinically...
can we change this?

Outline

1. **What** do we mean by individual differences?
2. **Why** should we care about individual differences?
3. **How** can we study individual differences?
(some practical guidelines)
4. **Where** are we going next?

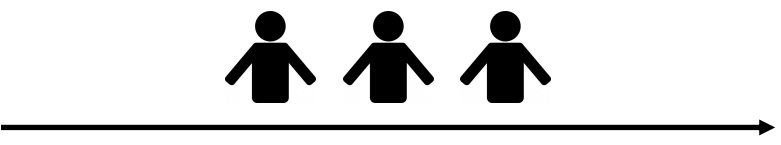
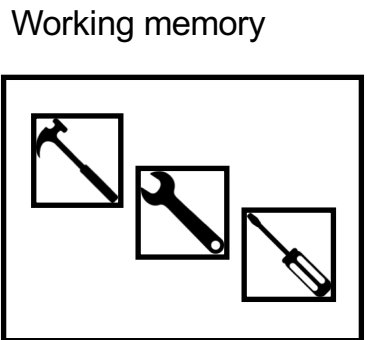
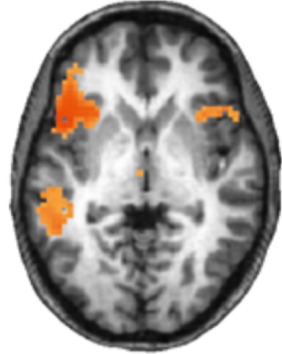
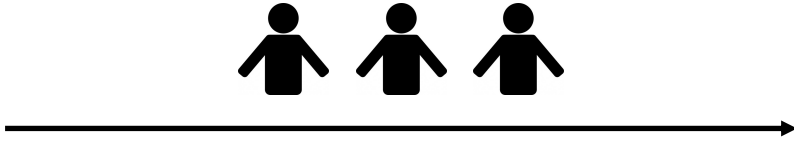
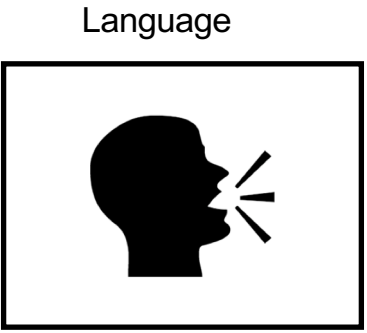
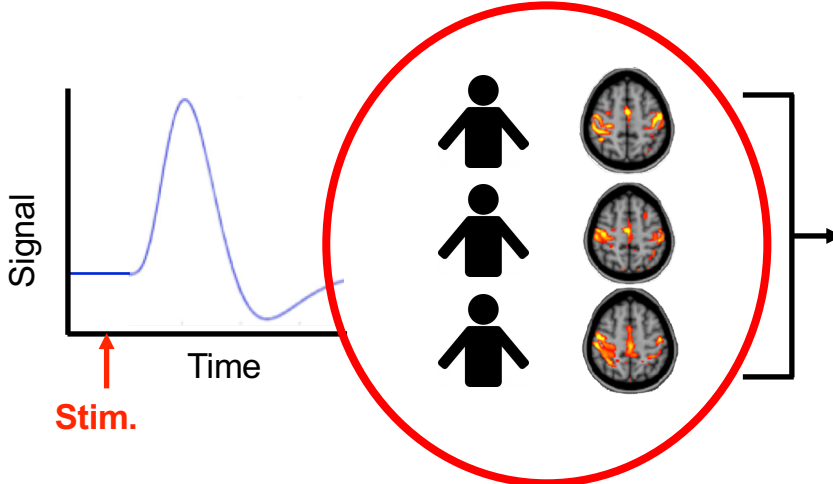
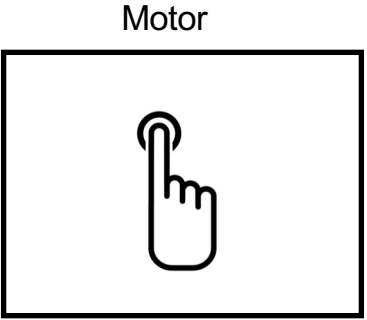


Outline

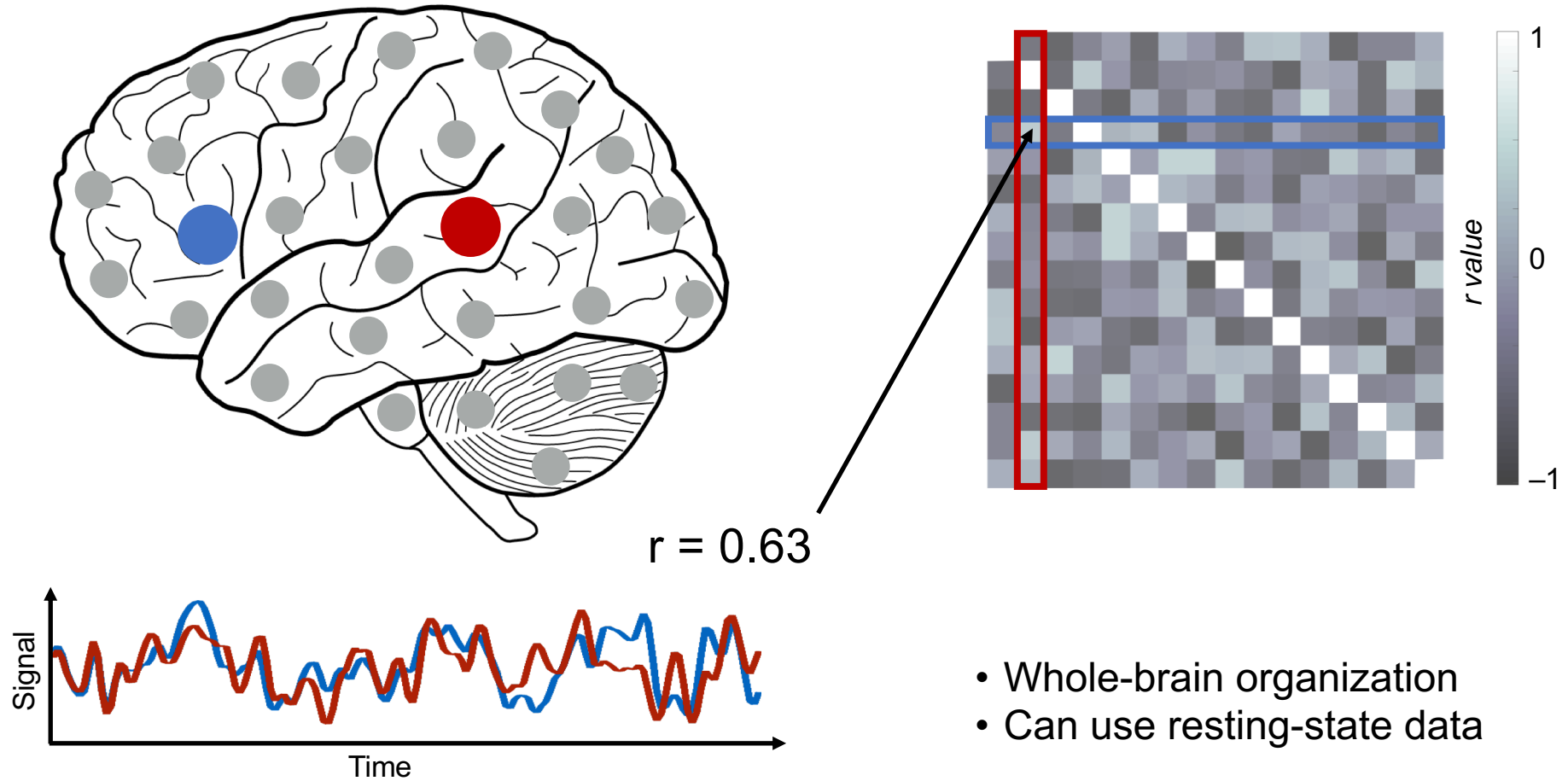
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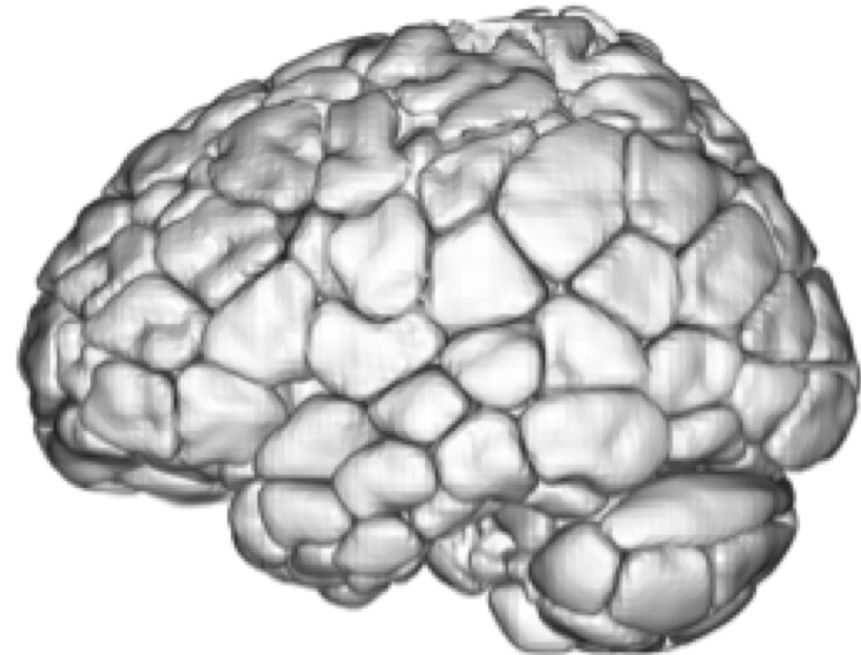
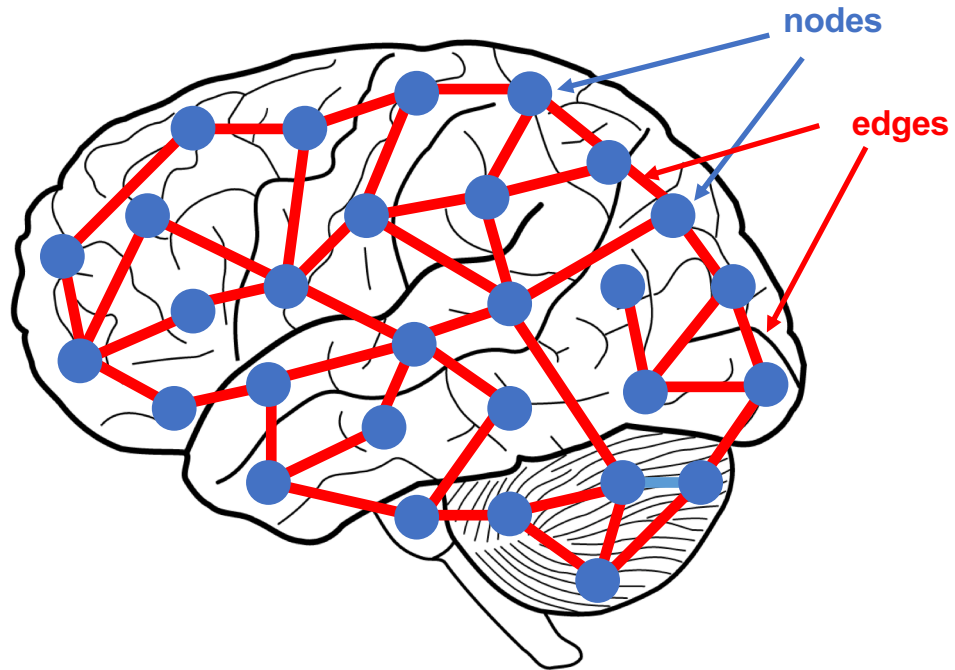
Traditional fMRI analyses



Functional connectivity



Brain networks

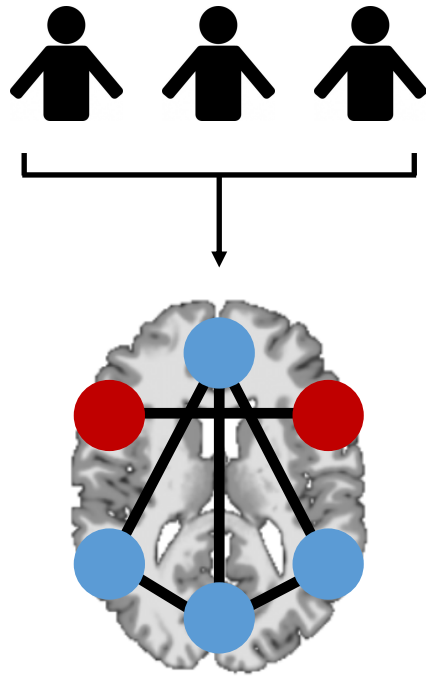


Shen et al., *NeuroImage* (2013)

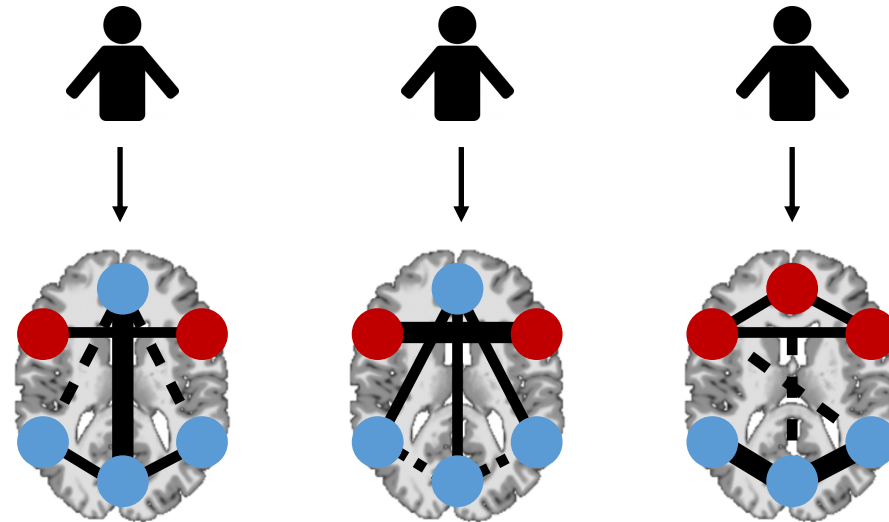
268 nodes = 35,778 edges

Individual differences

Group analyses



Individual differences



Identification experiments

Human Connectome Project

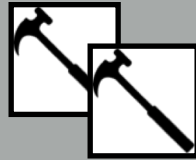
- 126 healthy subjects (50 sets of twins)
- Age 22-35 years old

Day 1

Resting
R1



Working memory
WM



Motor
Mt



Day 2

Resting
R2



Language
Lg



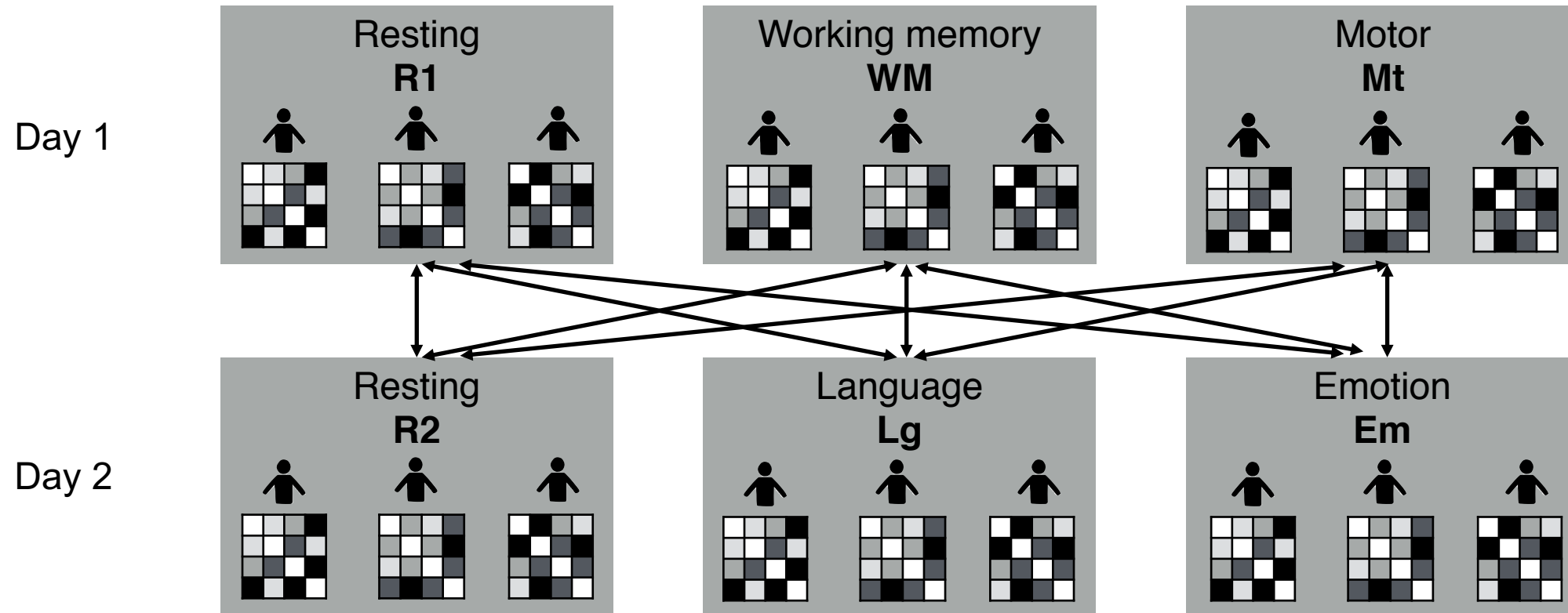
Emotion
Em



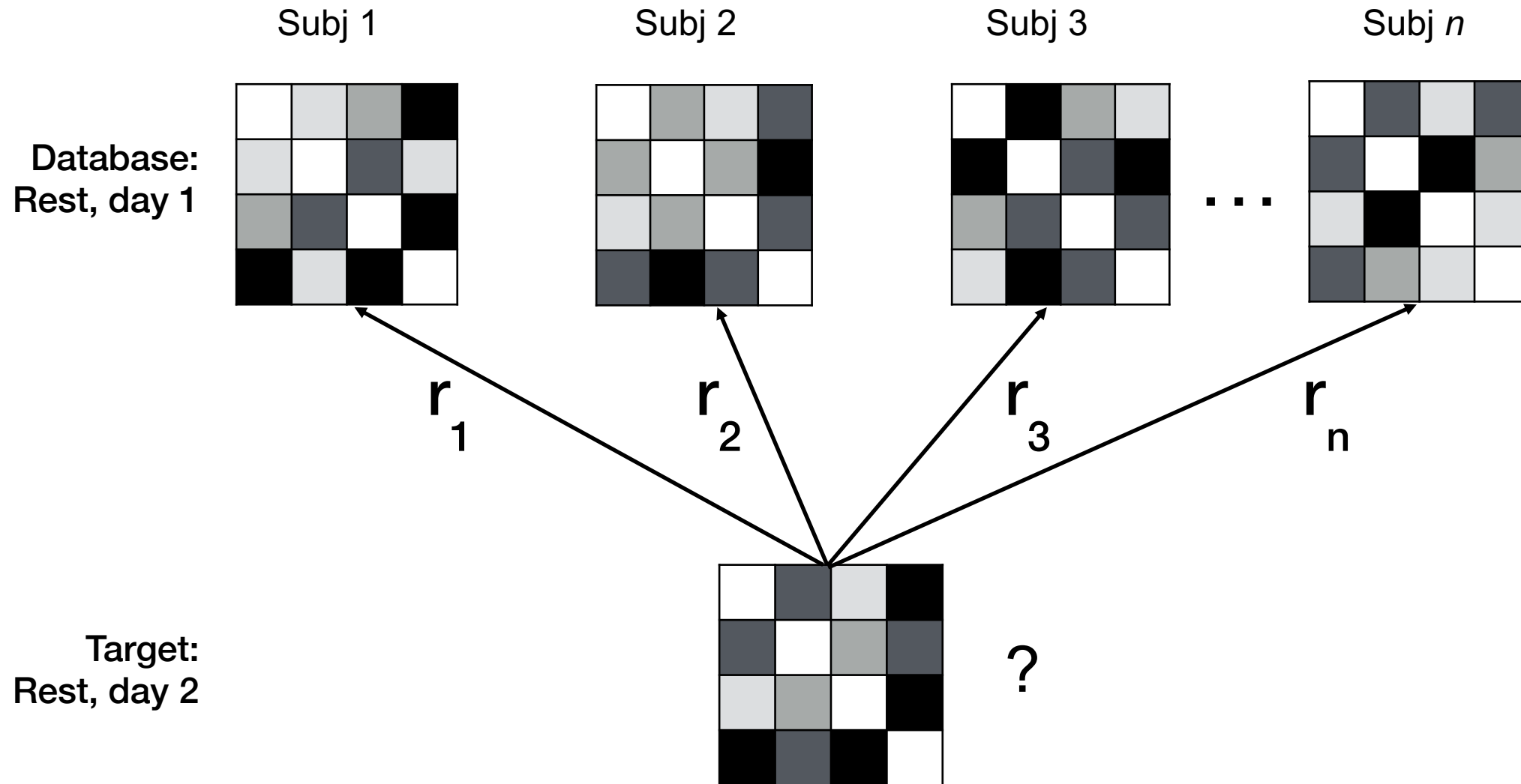
Identification experiments

Human Connectome Project

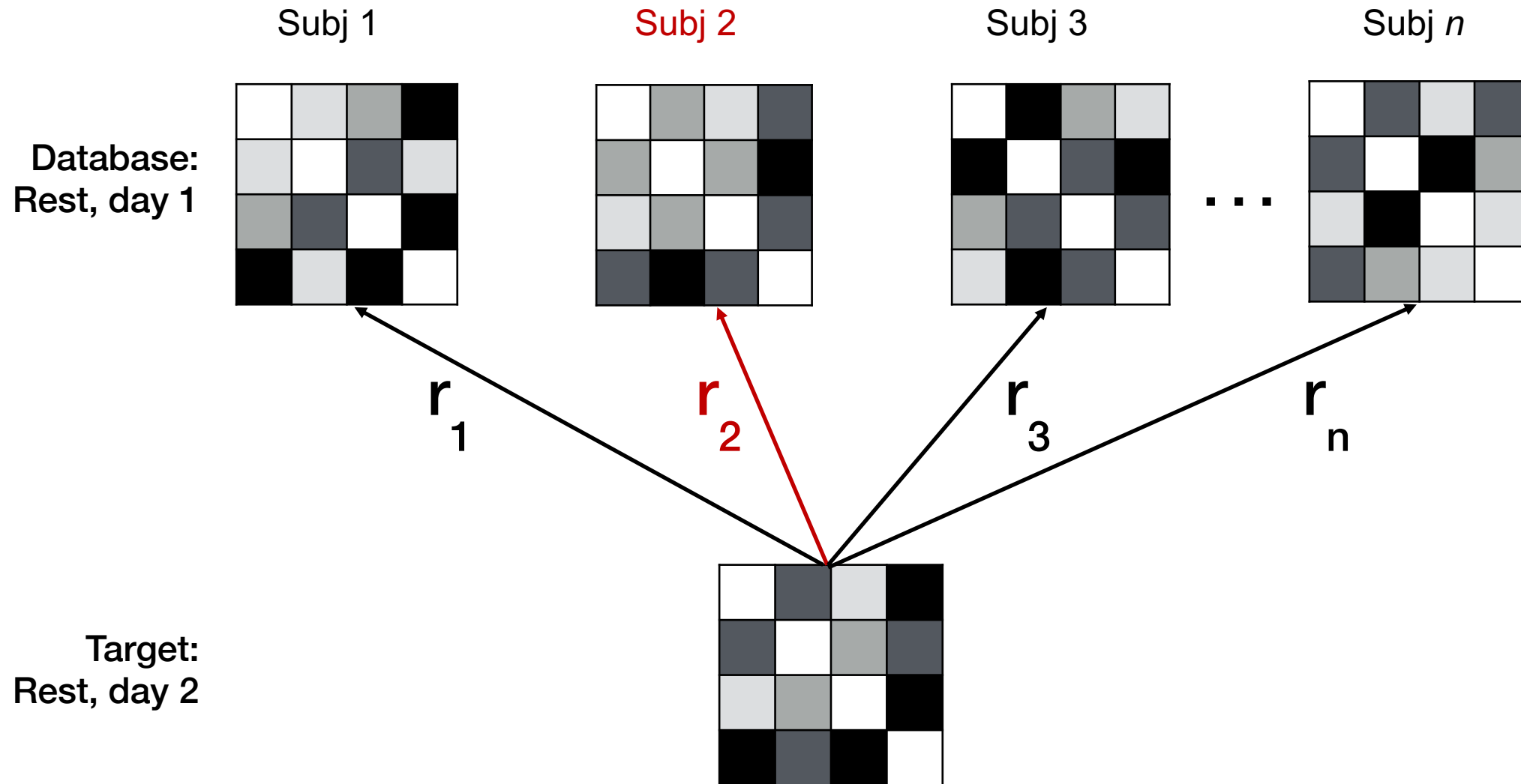
- 126 healthy subjects (50 sets of twins)
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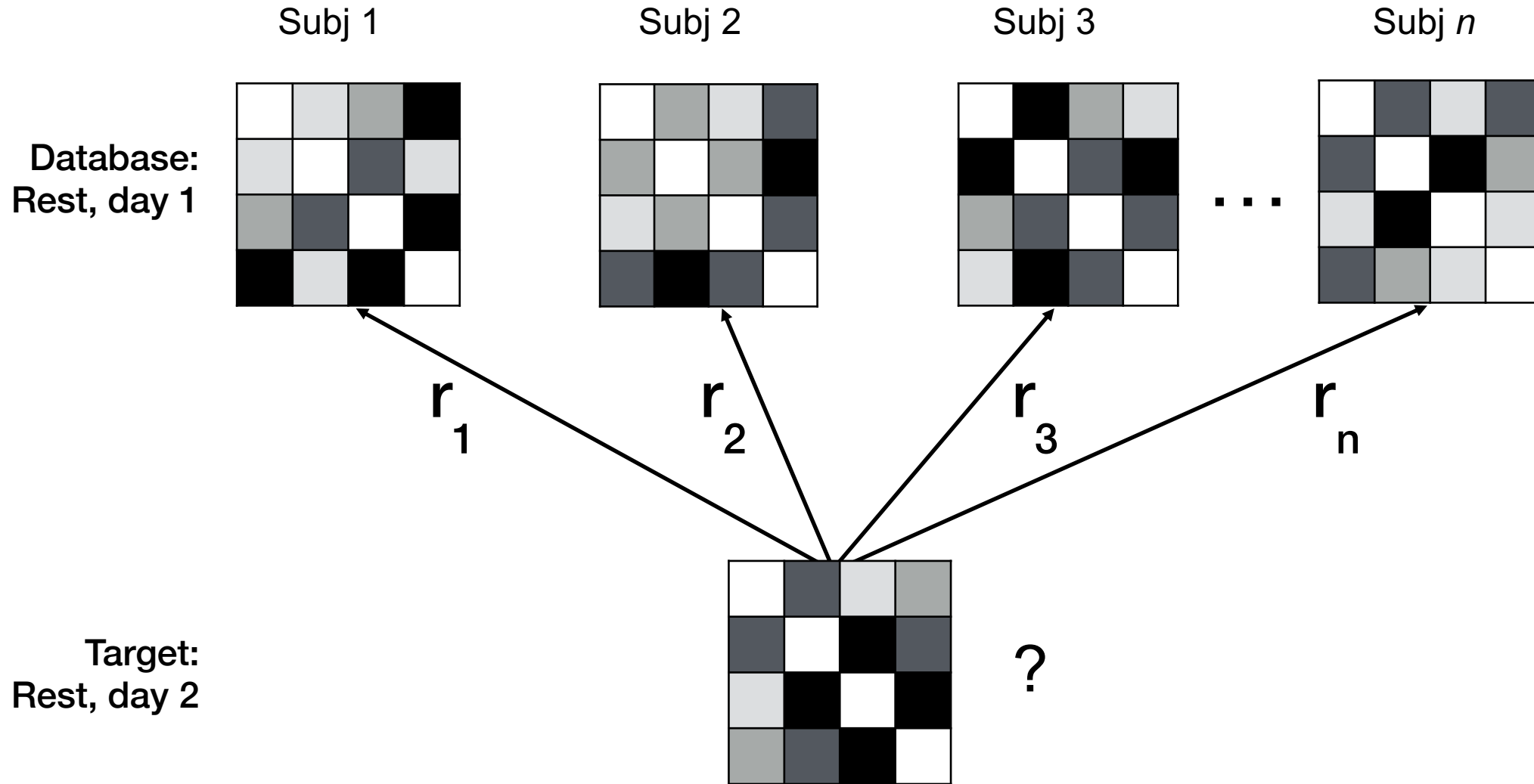
Identification experiments



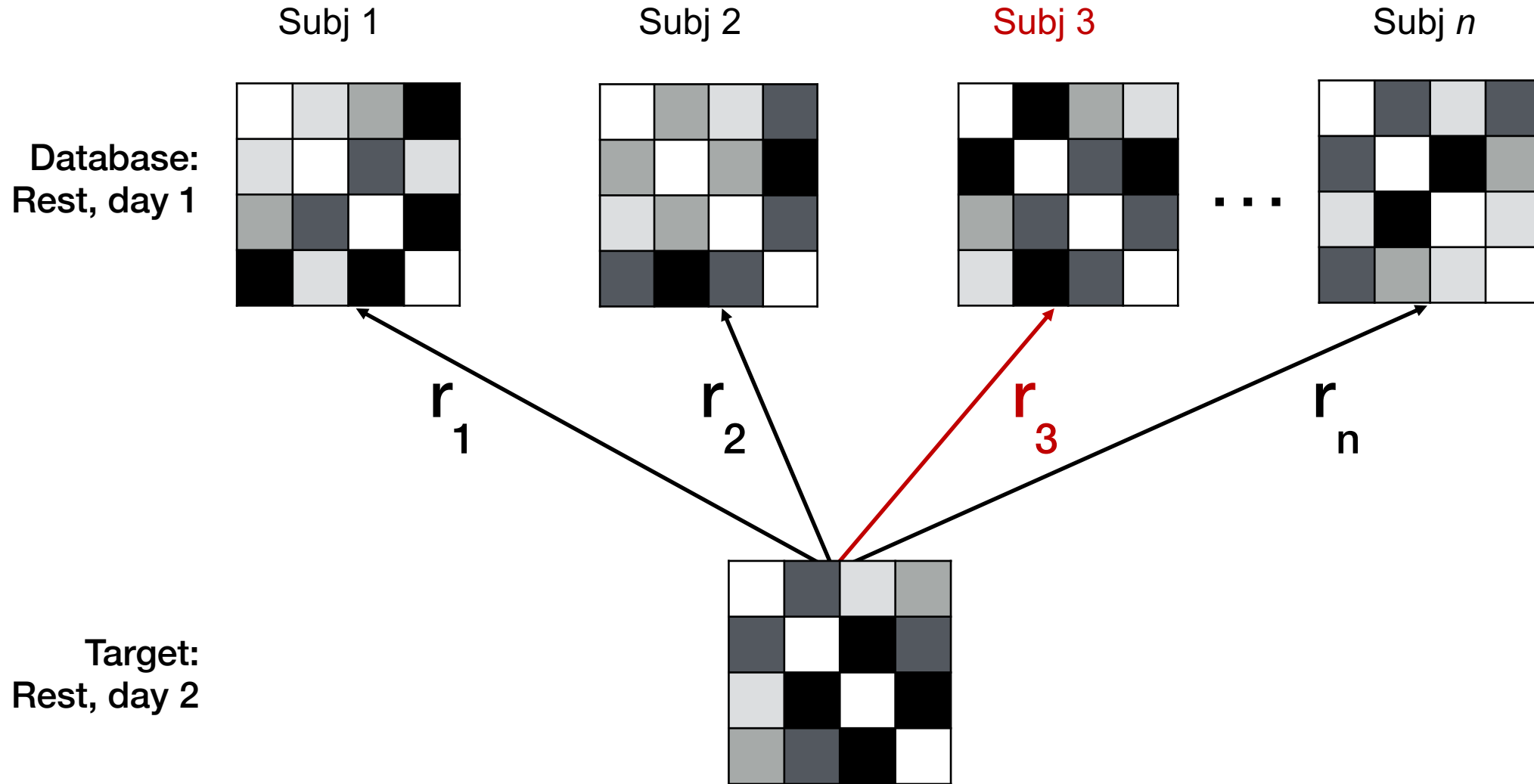
Identification experiments



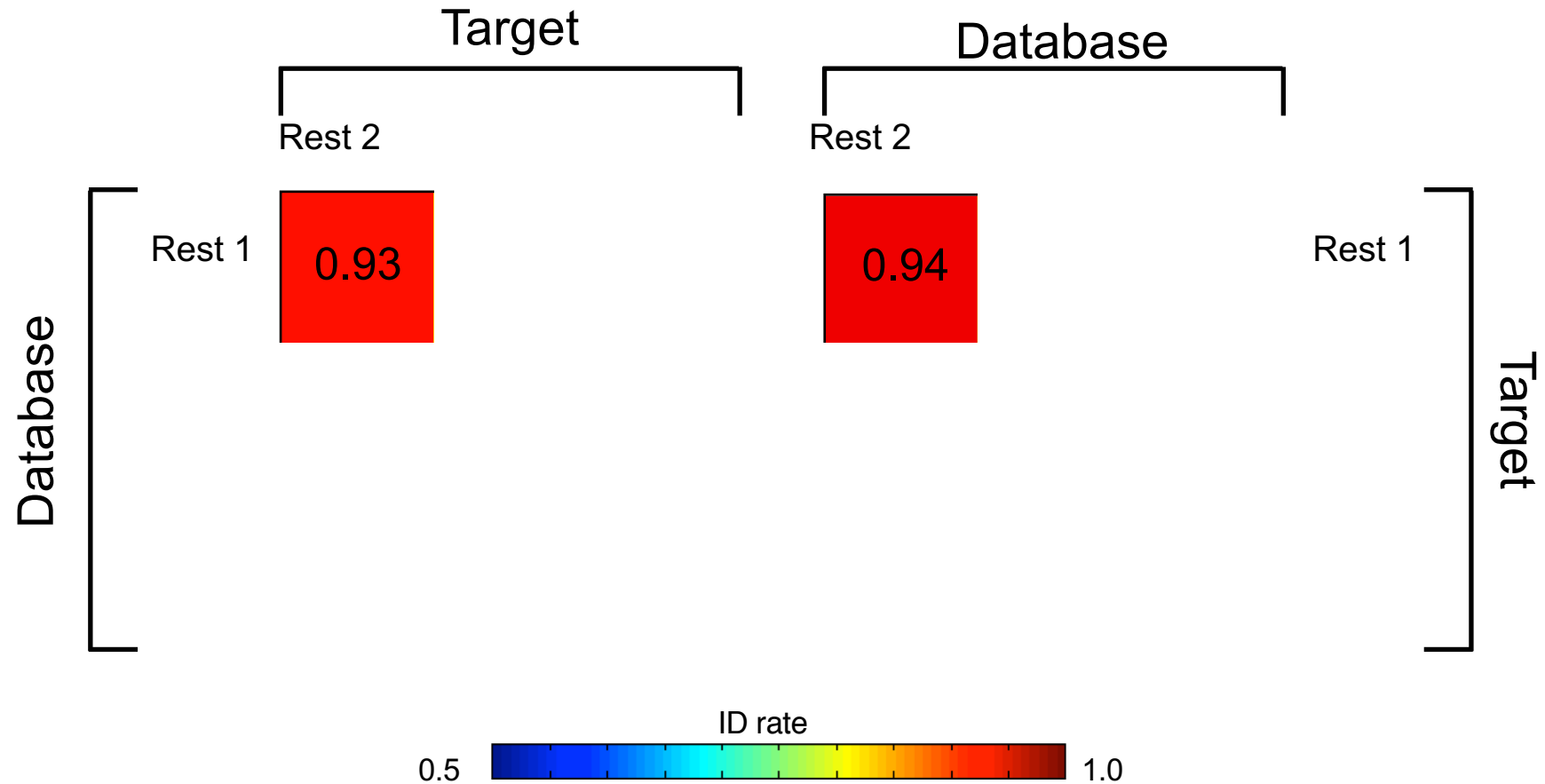
Identification experiments



Identification experiments



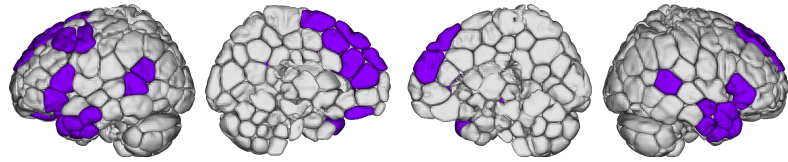
Identification results



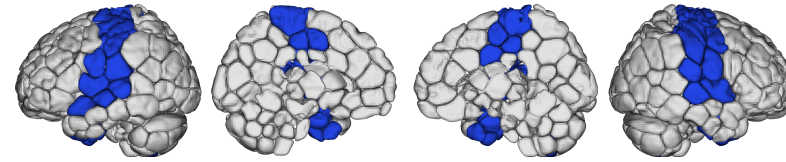
Chance: ~0.008

Network-based identification

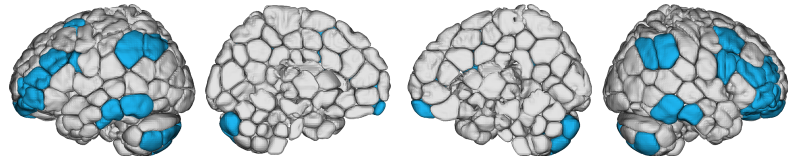
1. Medial frontal



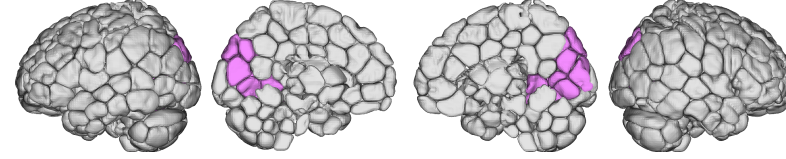
5. Somato-motor



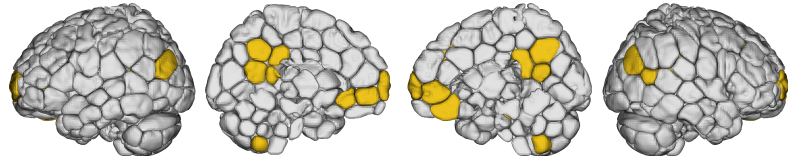
2. Frontoparietal



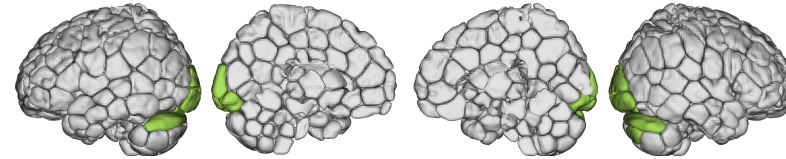
6. Visual I



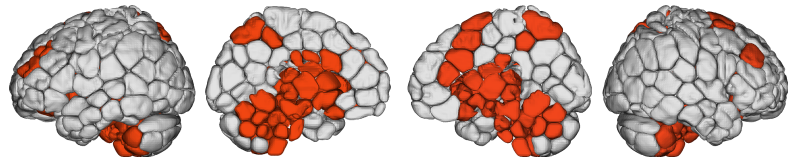
3. Default mode



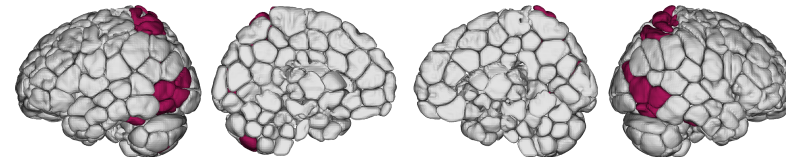
7. Visual II



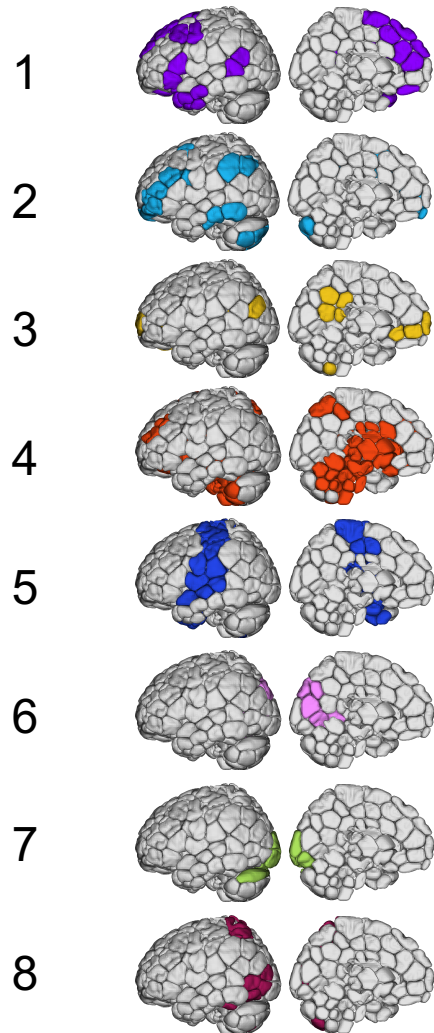
4. Subcortical/cerebellum



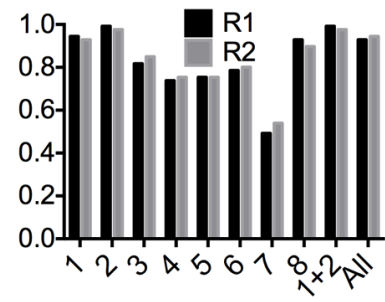
8. Visual association



Network-based identification



Identification rate

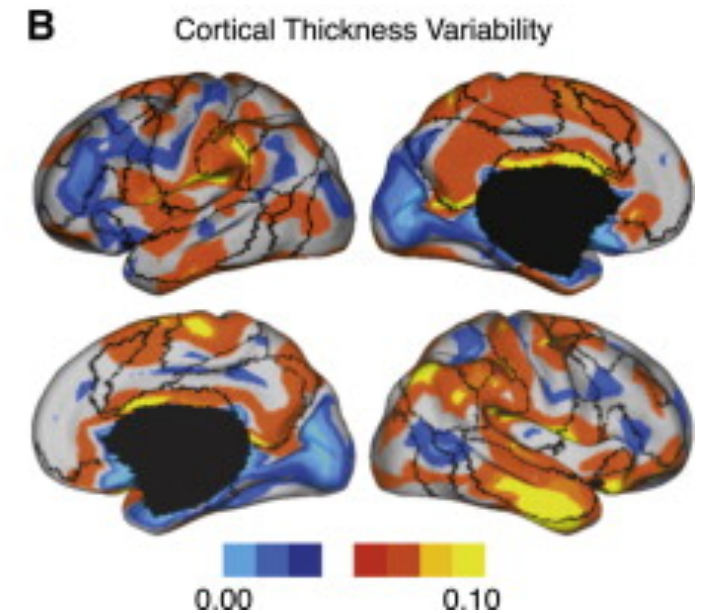
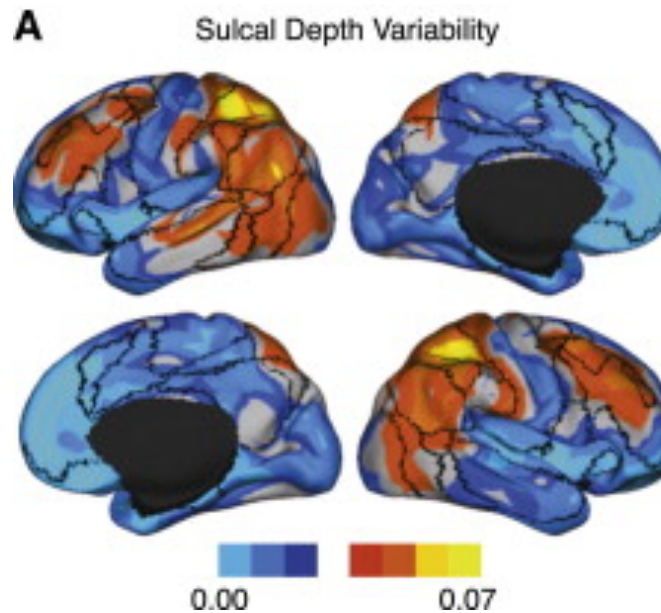
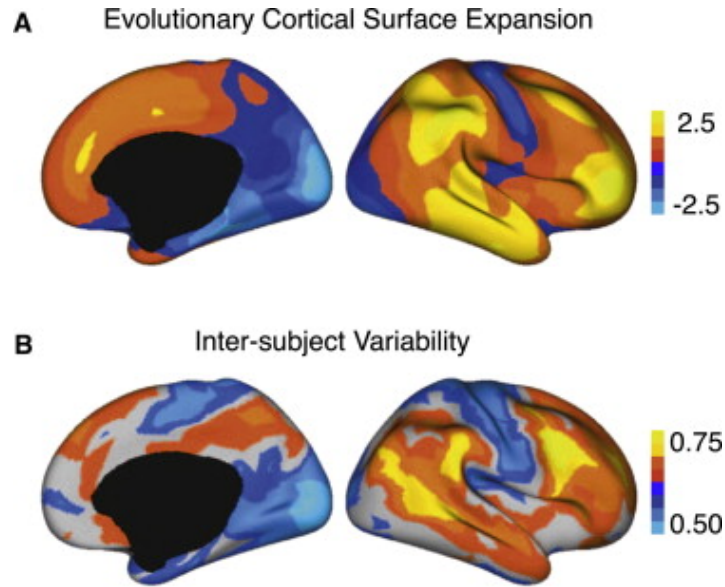


Network

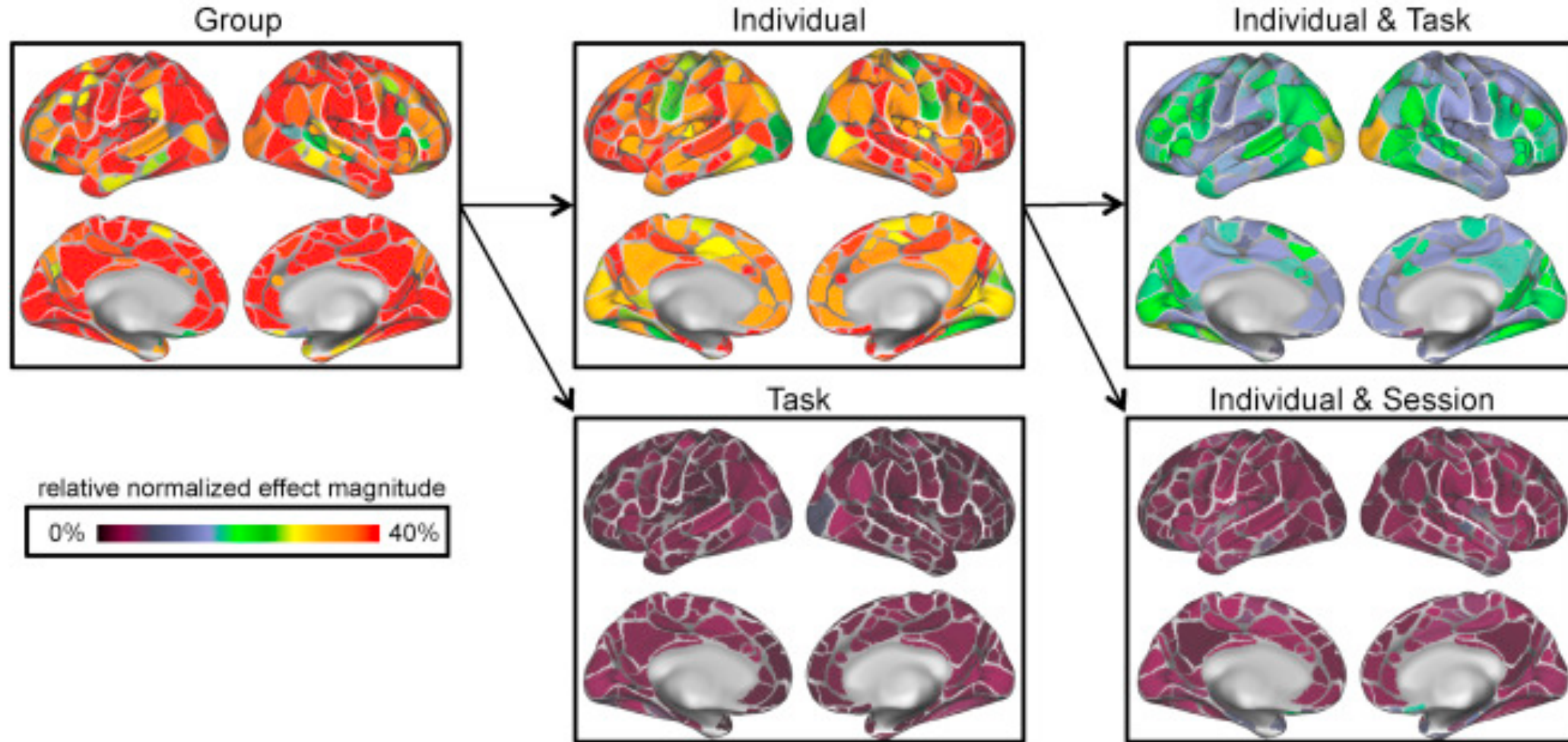
Localizing individual differences

Biggest differences found in most evolutionarily recent regions:

Anatomical differences also play a (large) role:



Individuals account for the most variance!



Gratton et al., *Neuron* (2018)

Outline

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(some practical guidelines)
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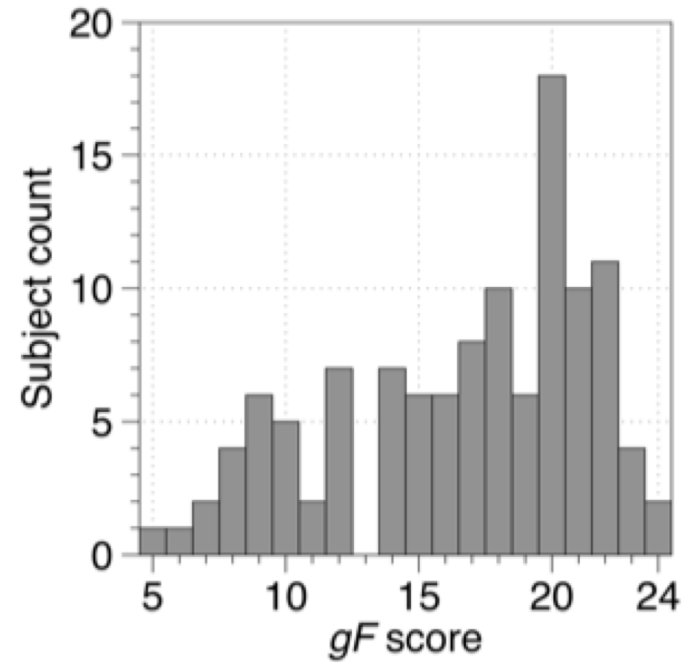
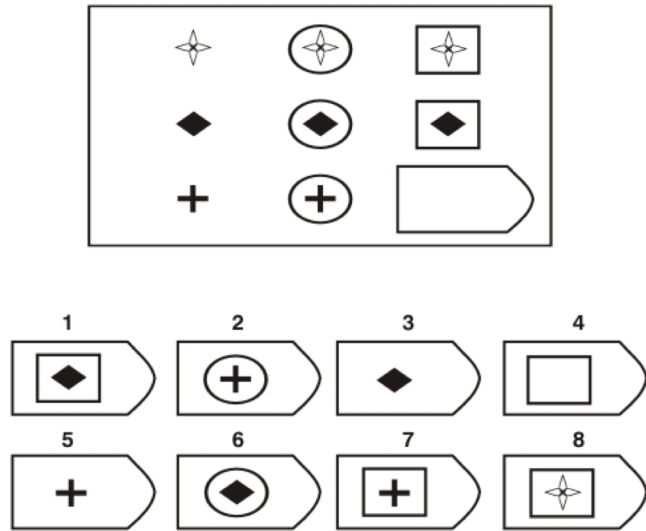
Why is this important?

- It's trivial to ID someone based on a structural scan
- Could just be anatomy, other confounds
- How can we prove these differences are meaningful?


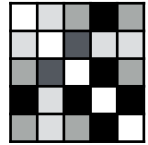

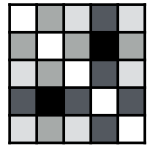

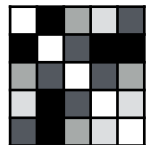

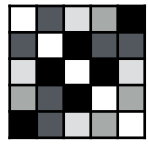

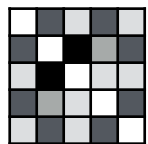
BEHAVIOR!

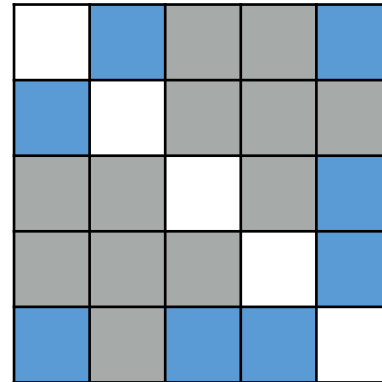
Predicting fluid intelligence

- ability to discern patterns
- independent of learned knowledge

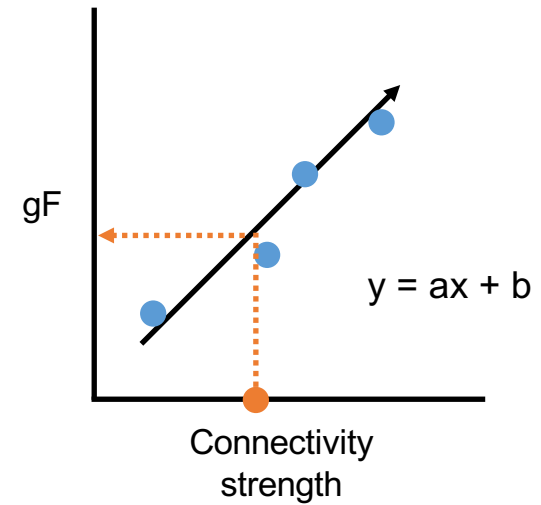


Predicting fluid intelligence

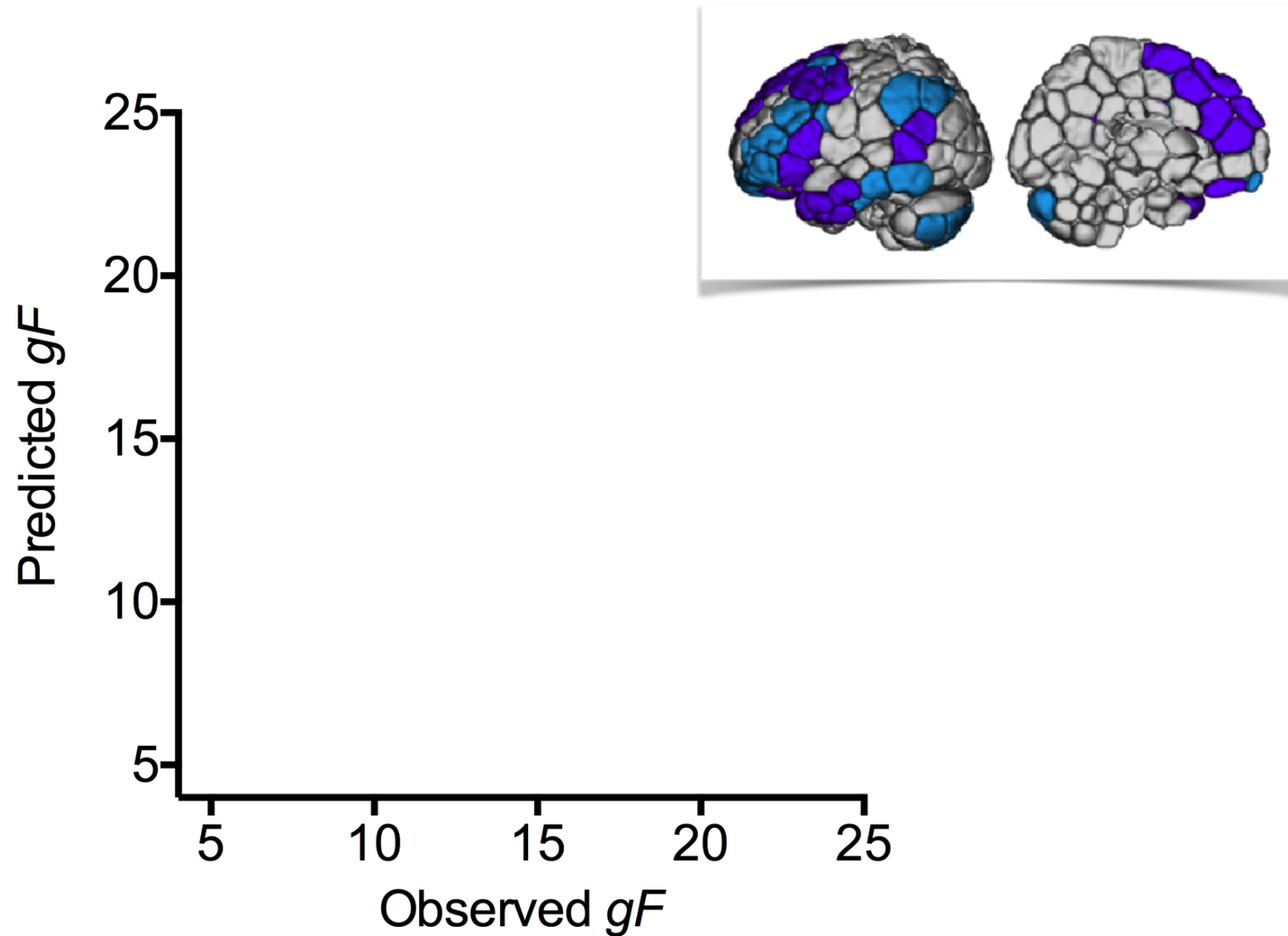
Subj	Matrix	Score
		17
		9
		24
		15
		



Connectome-based Predictive Model (CPM)

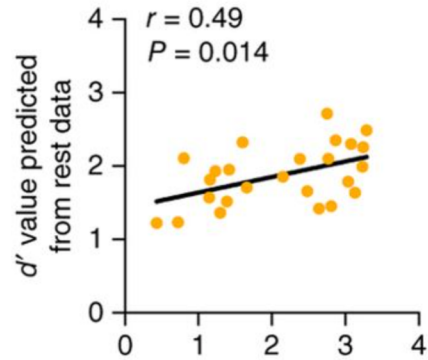


Predicting fluid intelligence



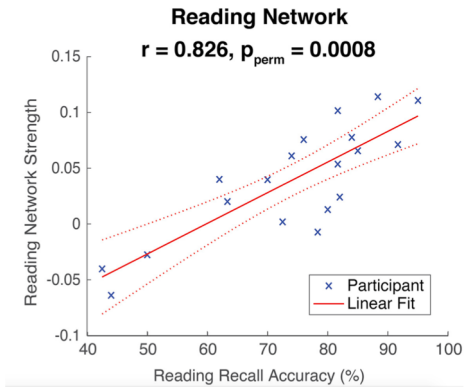
Predicting other behaviors

Sustained attention



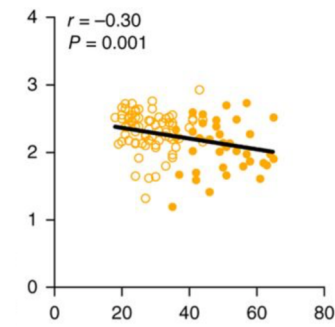
Rosenberg, Finn et al., *Nat Neurosci* (2016)

Reading ability

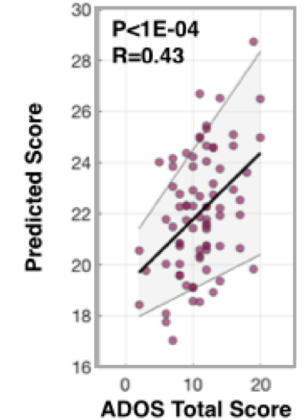


Jangraw et al., *NeuroImage* (2017)

ADHD, autism symptom severity

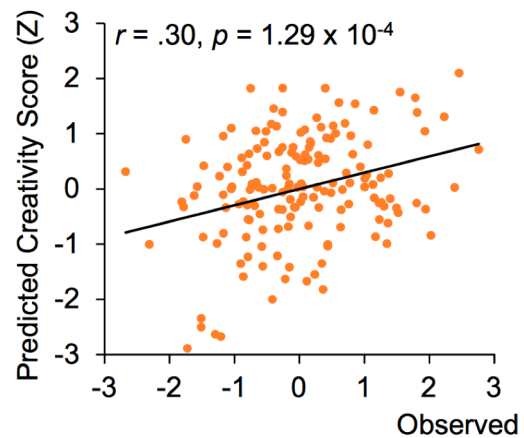


Rosenberg, Finn et al., *Nat Neurosci* (2016)



Lake et al., submitted

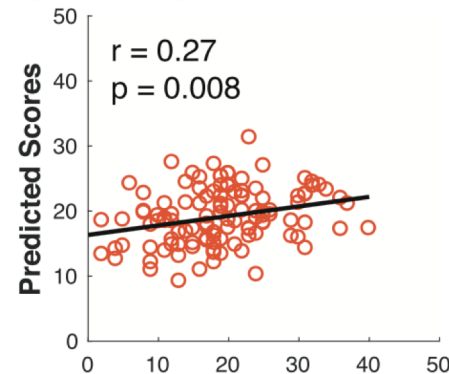
Creativity



Beaty et al., *PNAS* (2018)

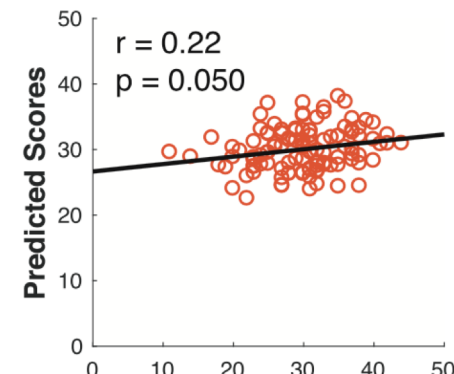
Personality traits

Neuroticism

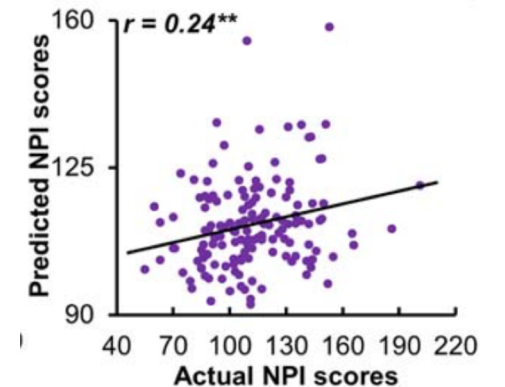


Hsu et al., *Soc Cogn Aff Neurosci* (2018)

Extraversion



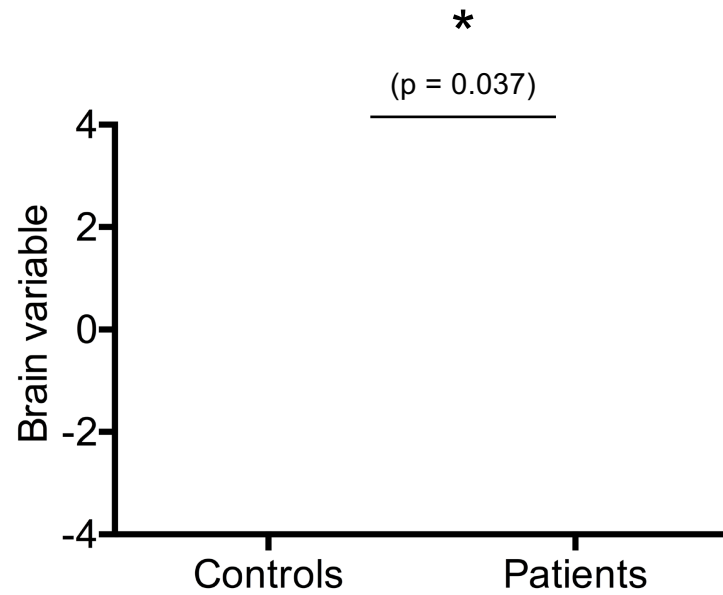
Narcissism



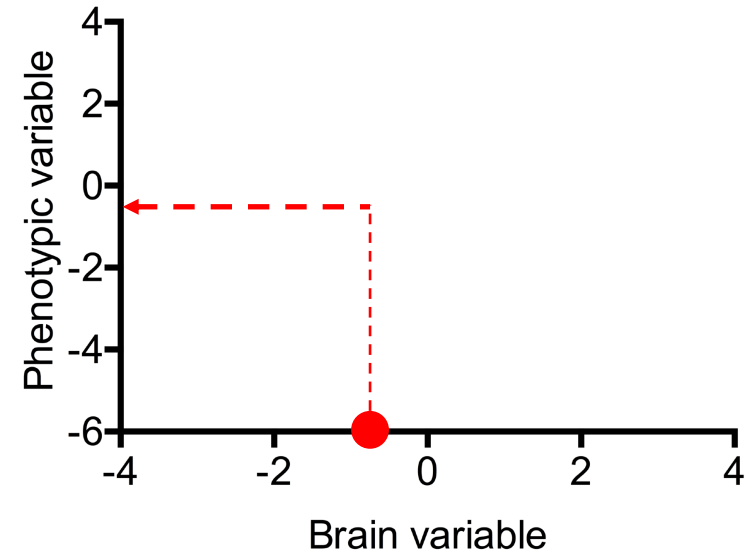
Feng et al., *Hum Brain Mapp* (2018)

Translational applications

Categorical approach



Dimensional approach



Outline

1. **What** do we mean by individual differences?
2. **Why** should we care about individual differences?
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Outline

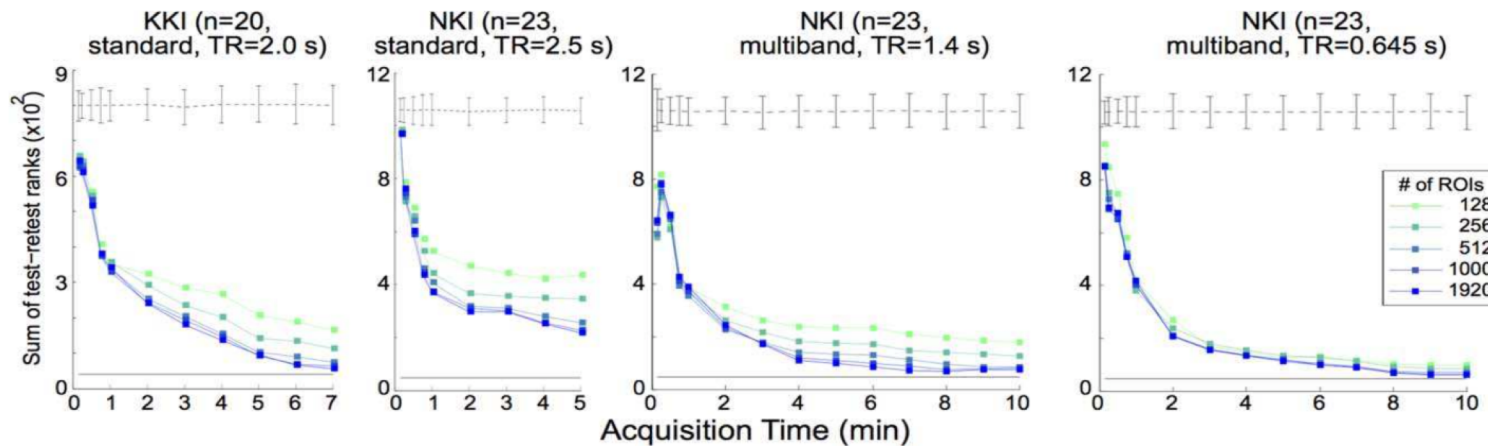
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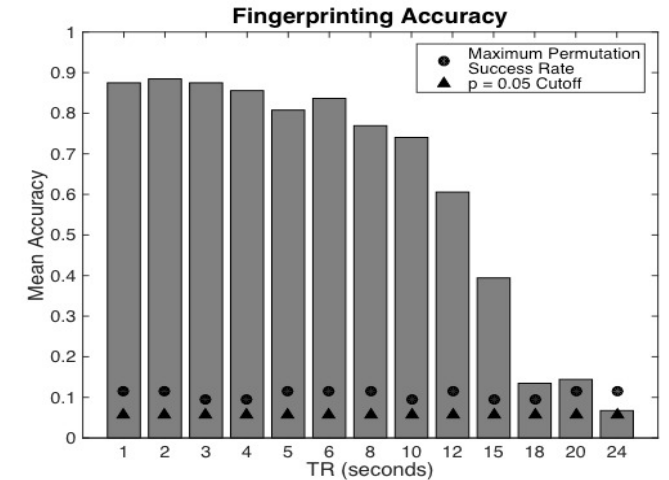
Q. Do you need HCP-quality data?

A. Not really

ID is fairly robust even at more standard spatial & temporal resolutions:



Airan et al., Hum Brain Mapp (2016)



Courtesy of Jason Druzgal

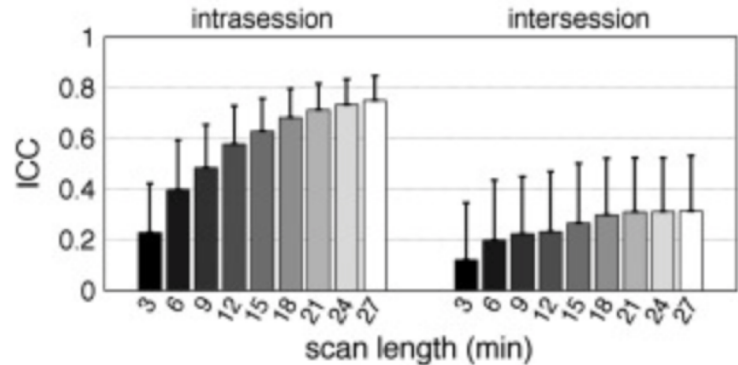
- More nodes → higher identification rate
 - Parcellation method (random vs. functional) did not matter
 - *Caution:* Higher resolution may amplify effects of anatomical diffs/registration error
 - Parcellations in the 200-300 node range seem like a good compromise

Q. What about amount of data?

A. Scan duration matters!

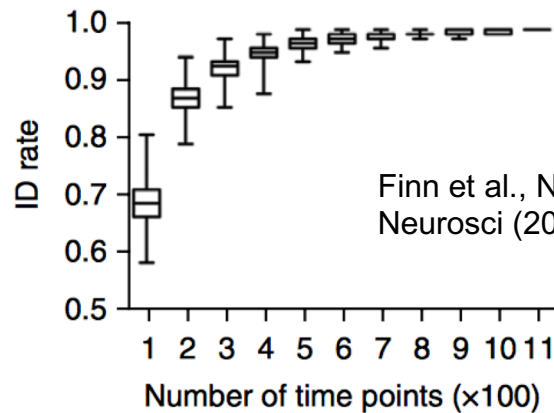
Longer acquisitions are better:

- higher reliability within subjects



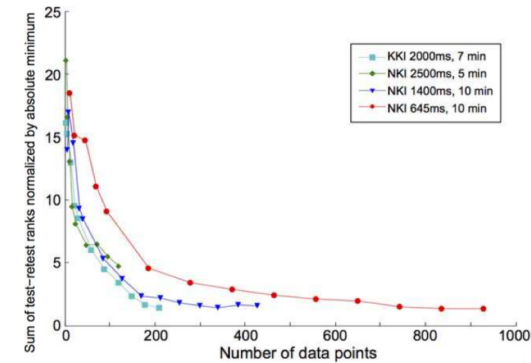
Birn et al., NeuroImage (2013)

- higher identifiability across subjects

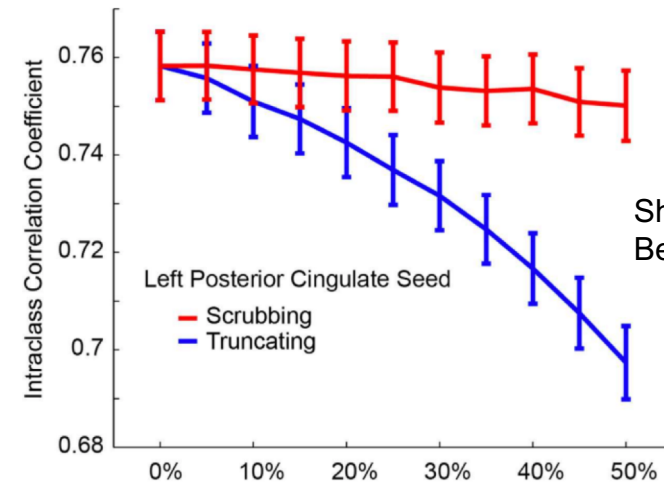


Finn et al., Nat Neurosci (2015)

- ▶ higher sampling rate (shorter TR) cannot make up for shorter scan duration



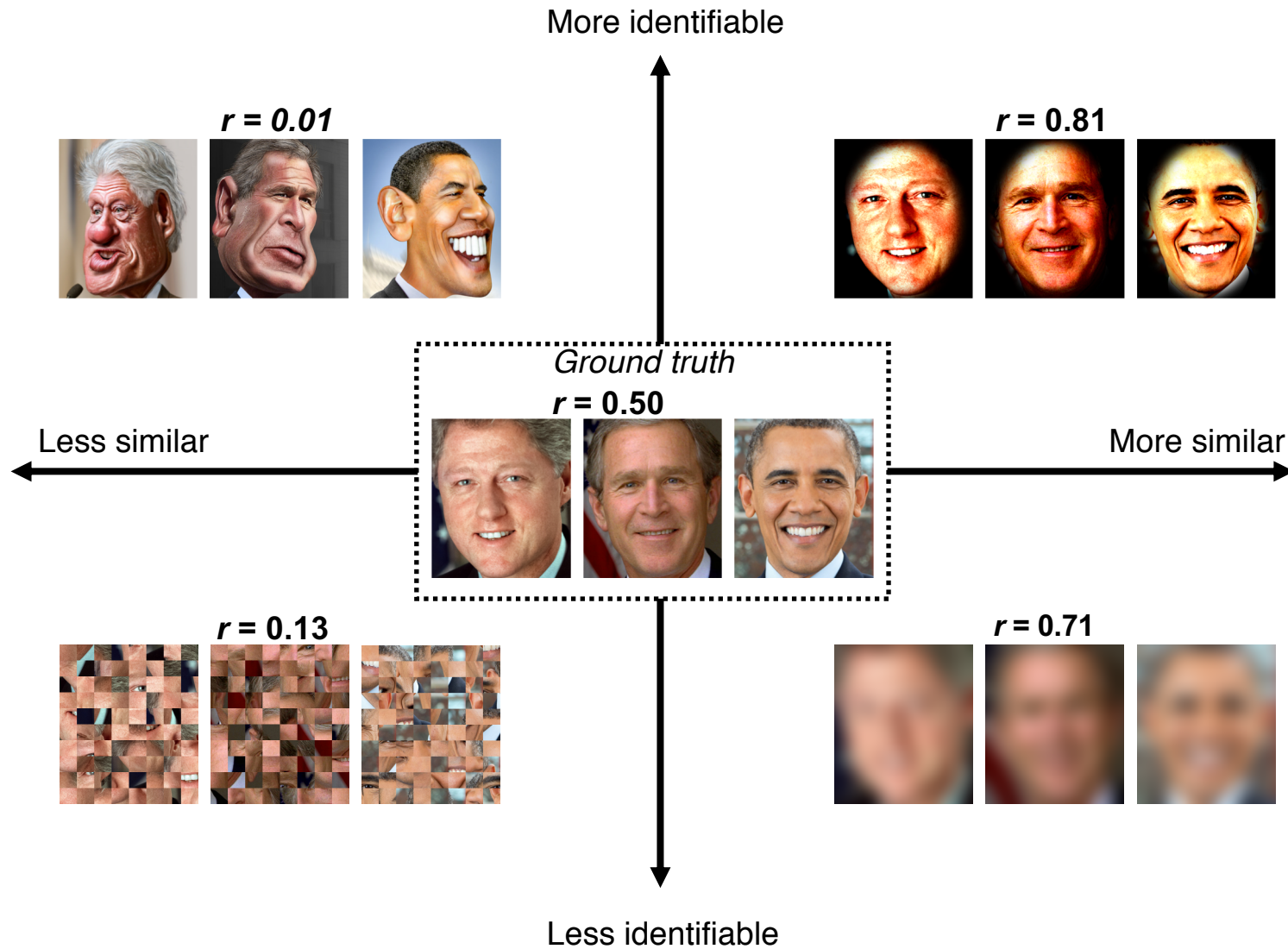
Airan et al., Hum Brain Mapp (2016)



Shah et al., Brain & Behav (2016)

Q. Does scan condition matter?

A. Yes!

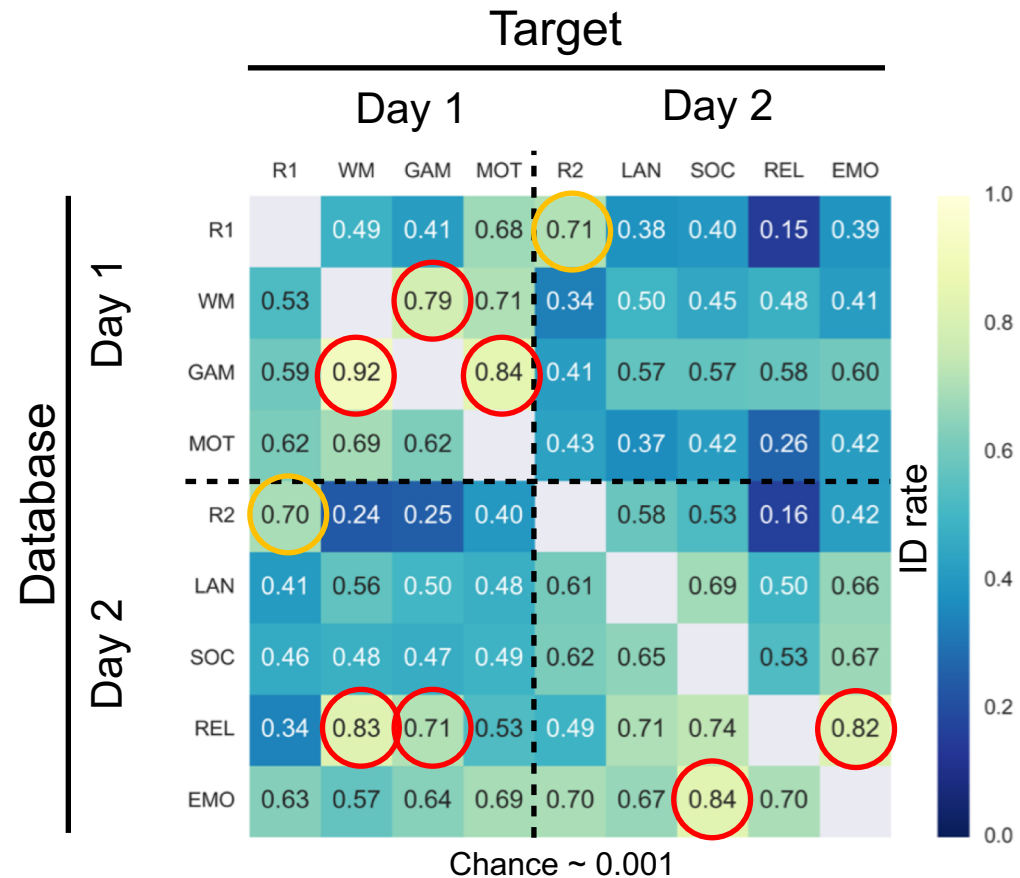


Q. Does scan condition matter?

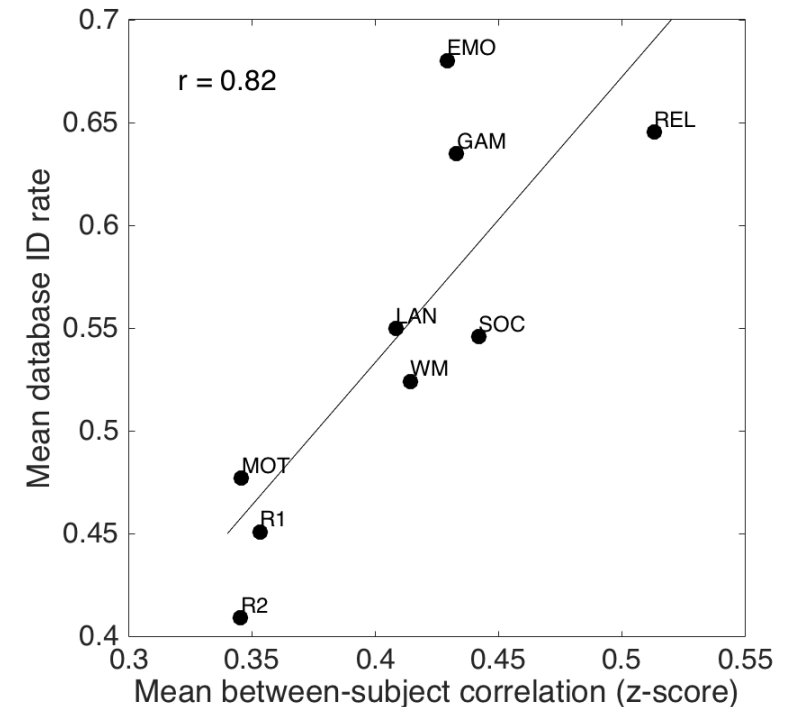
A. Yes!

Rest has become the default condition for FC & individual differences, but tasks may increase signal-to-noise

Replicating identification experiments:



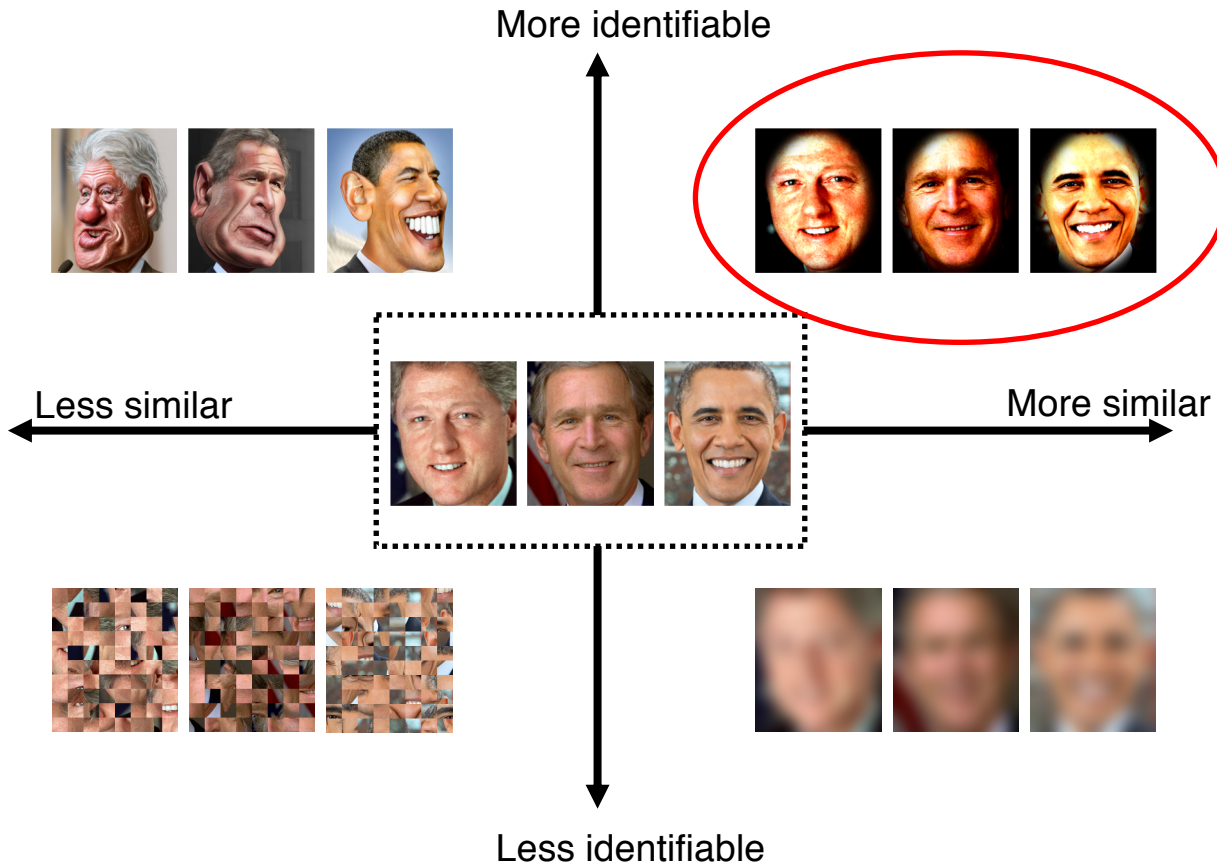
Conditions that make subjects look **more similar** to one another actually make **better databases** for identification:



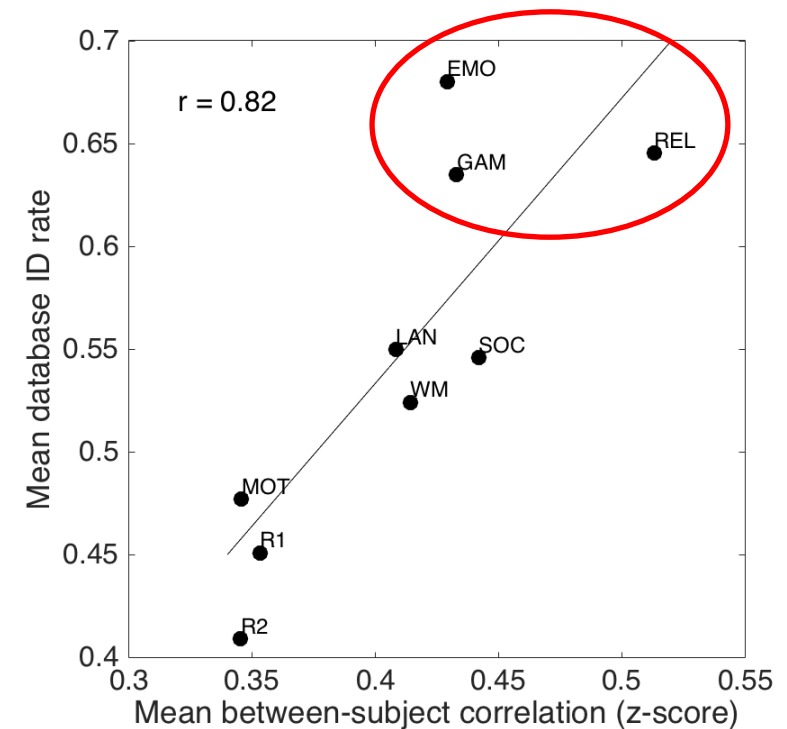
Q. *Does scan condition matter?*

A. Yes!

Rest has become the default condition for FC & individual differences, but tasks may increase signal-to-noise

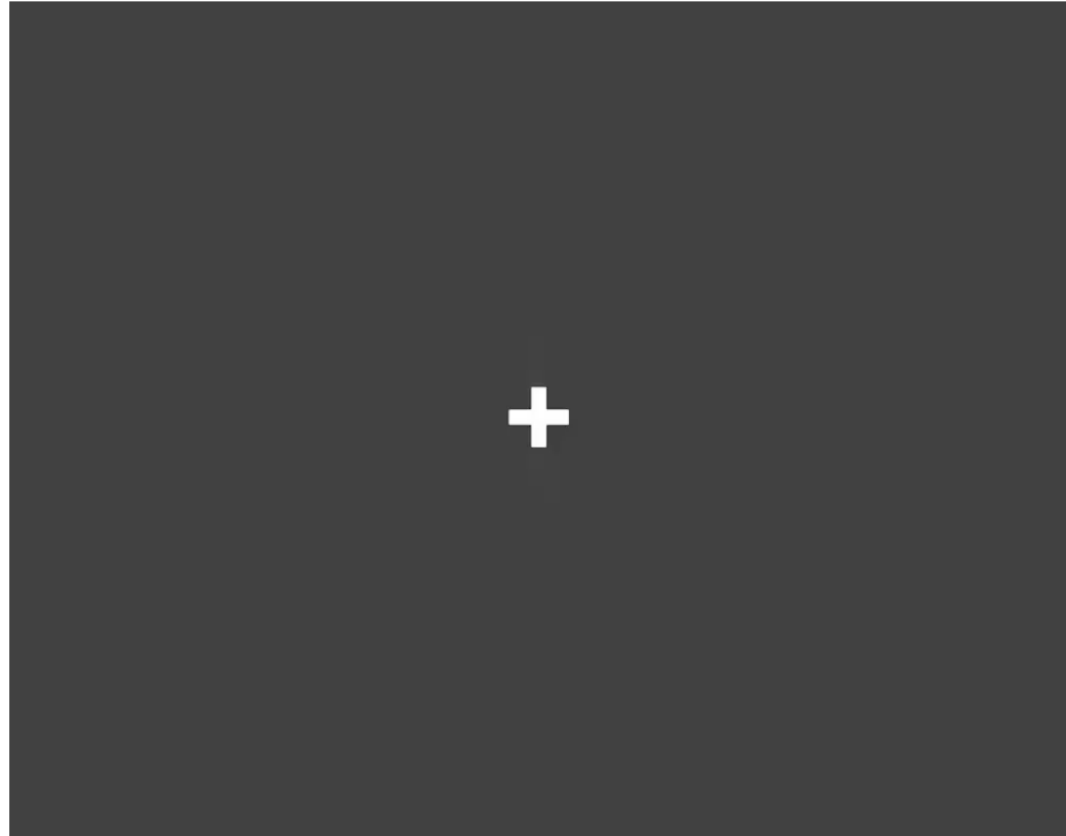


Conditions that make subjects look **more similar** to one another actually make **better databases** for identification:



Q. *Is rest best?*
A. Probably not

Consider naturalistic tasks:

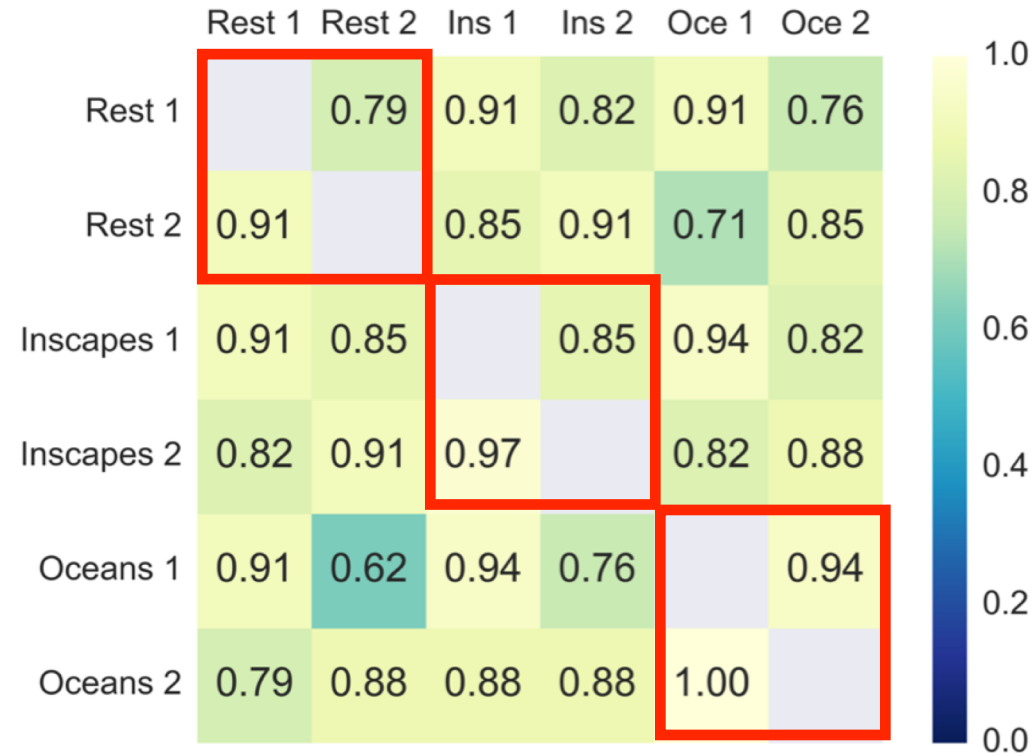
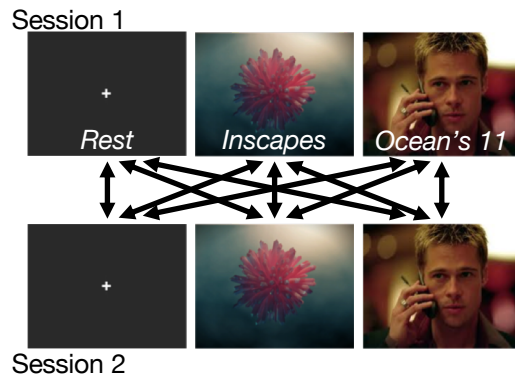


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Consider naturalistic tasks:

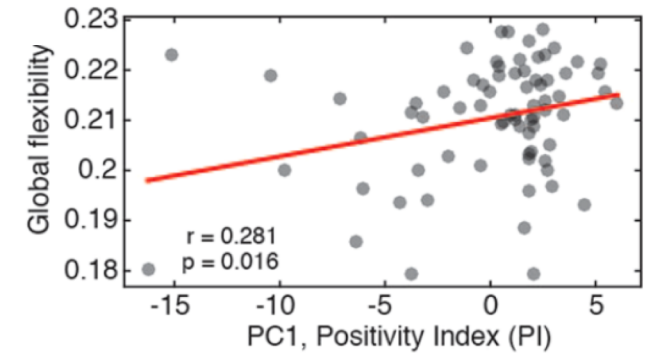
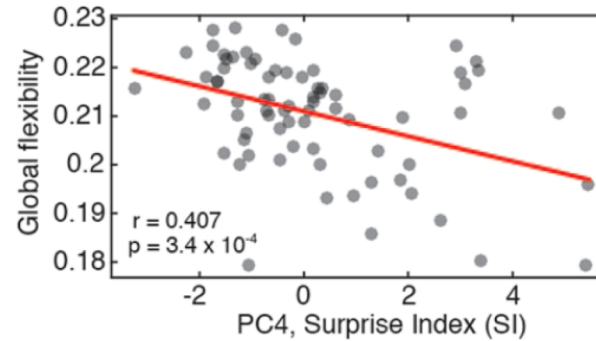
- ID rate is just as good as (if not better than) rest



How to choose behavior

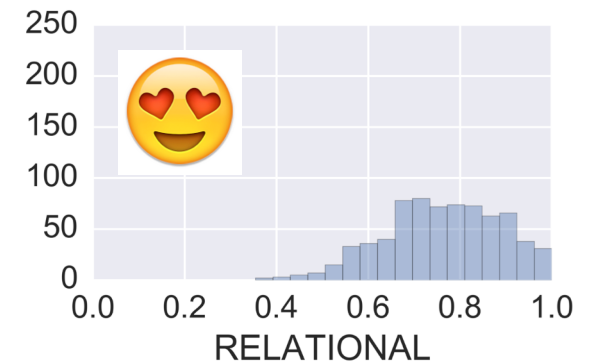
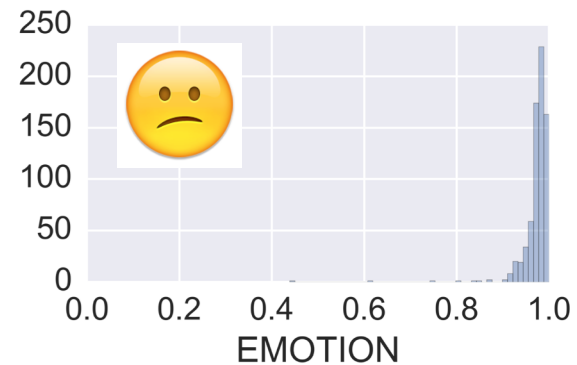
Is it stable?

- Trait vs. state
- State variables may be better suited to within-subject analysis



Betzel et al., Sci Rep (2017)

Does it show a good distribution in your population?



Behavior: Mitigating confounds

Many behaviors/phenotypes are correlated with head motion!

Negatively:

Positively:

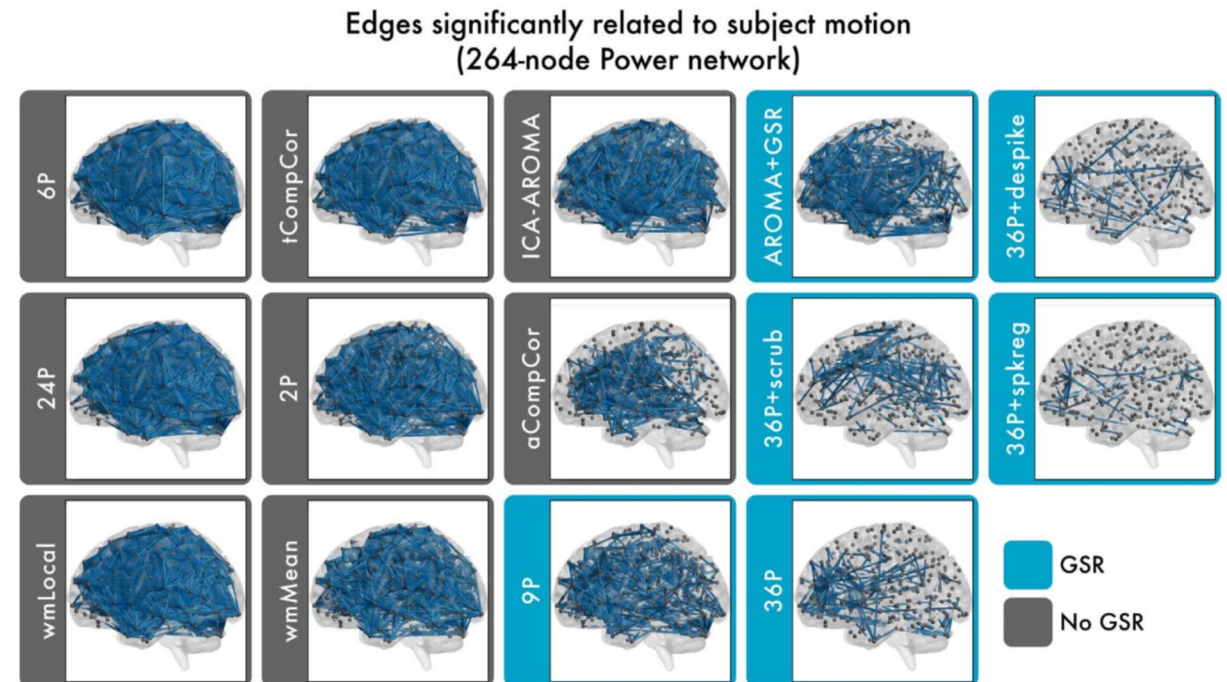
Subject measures	Pearson <i>r</i>		
ReadEng (AgeAdj)	-0.23	DSM somatic problems (pct)	0.16
ReadEng (Unadj)	-0.23	DSM antisocial (raw)	0.16
Vocabulary (AgeAdj)	-0.19	ASR externalizing (raw)	0.16
Dexterity (Unadj)	-0.18	DSM somatic problems (raw)	0.16
CardSort (Unadj)	-0.18	Tobacco use 7 day	0.18
Dexterity (AgeAdj)	-0.18	Diastolic blood pressure	0.18
CardSort (AgeAdj)	-0.18	ASR externalizing	0.18
Education	-0.17	Tobacco use today	0.2
Fluid intelligence	-0.17	Systolic blood pressure	0.23
Spatial orientation	-0.17	Weight	0.52
Vocabulary (unadjj)	-0.17	Body mass index (BMI)	0.66
Emotion recognition	-0.16		

- Patients of any kind move more
- Children move more
- Older adults move more

Behavior: Mitigating confounds

Many behaviors/phenotypes are correlated with head motion!

- ▶ Check correlation in your sample
- ▶ Consider excluding particularly high-motion subjects
- ▶ Choose appropriate preprocessing techniques
- ▶ Use motion as an explicit covariate

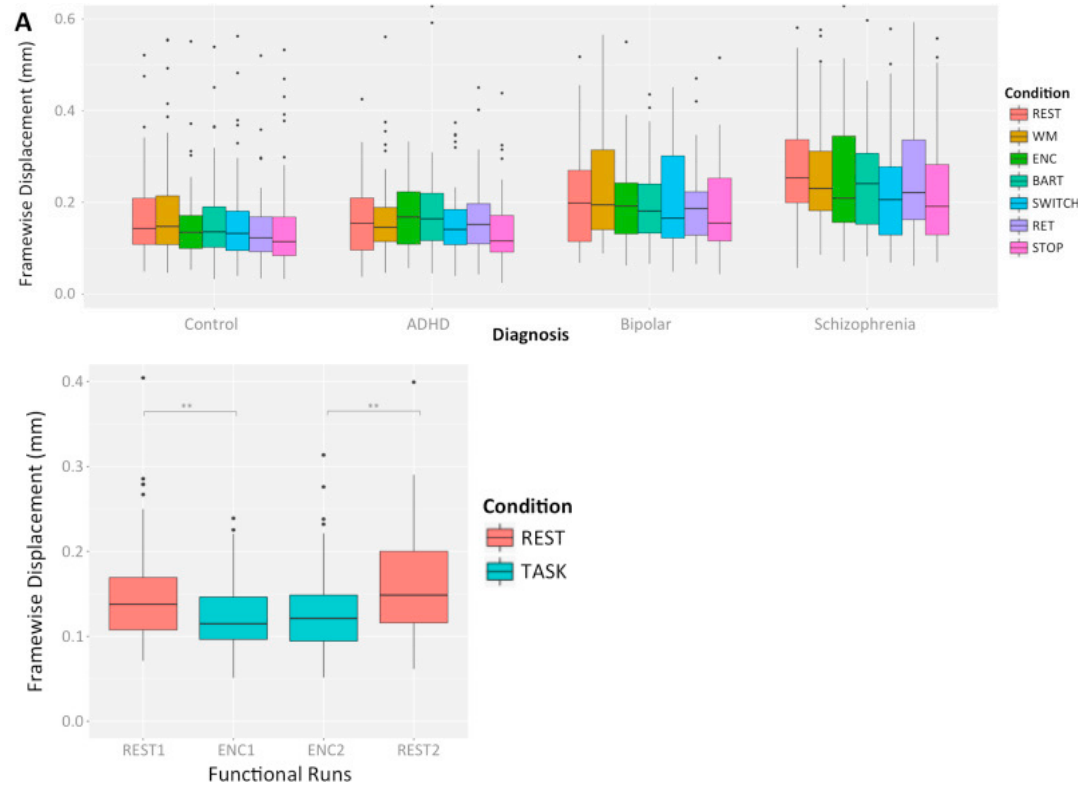


Ciric et al., NeuroImage (2017)

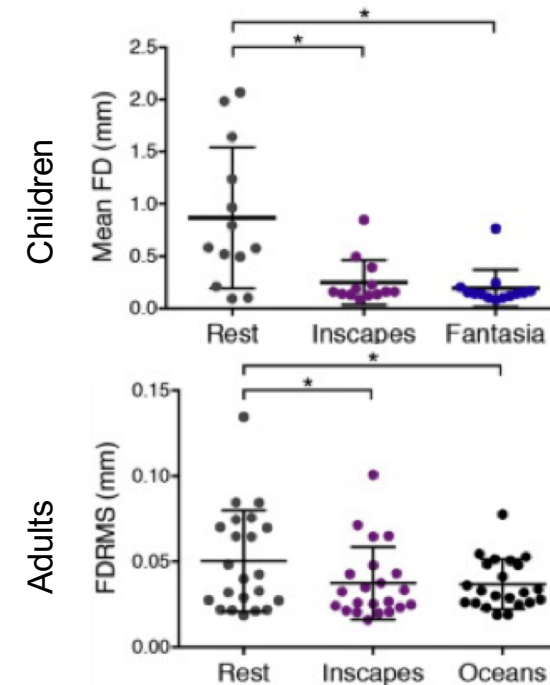
Q. Is rest best? A. Probably not

Tasks also have purely practical advantages:

- ▶ increase subject compliance (i.e., decrease head motion), especially in certain populations



Huijbers et al., NeuroImage (2017)

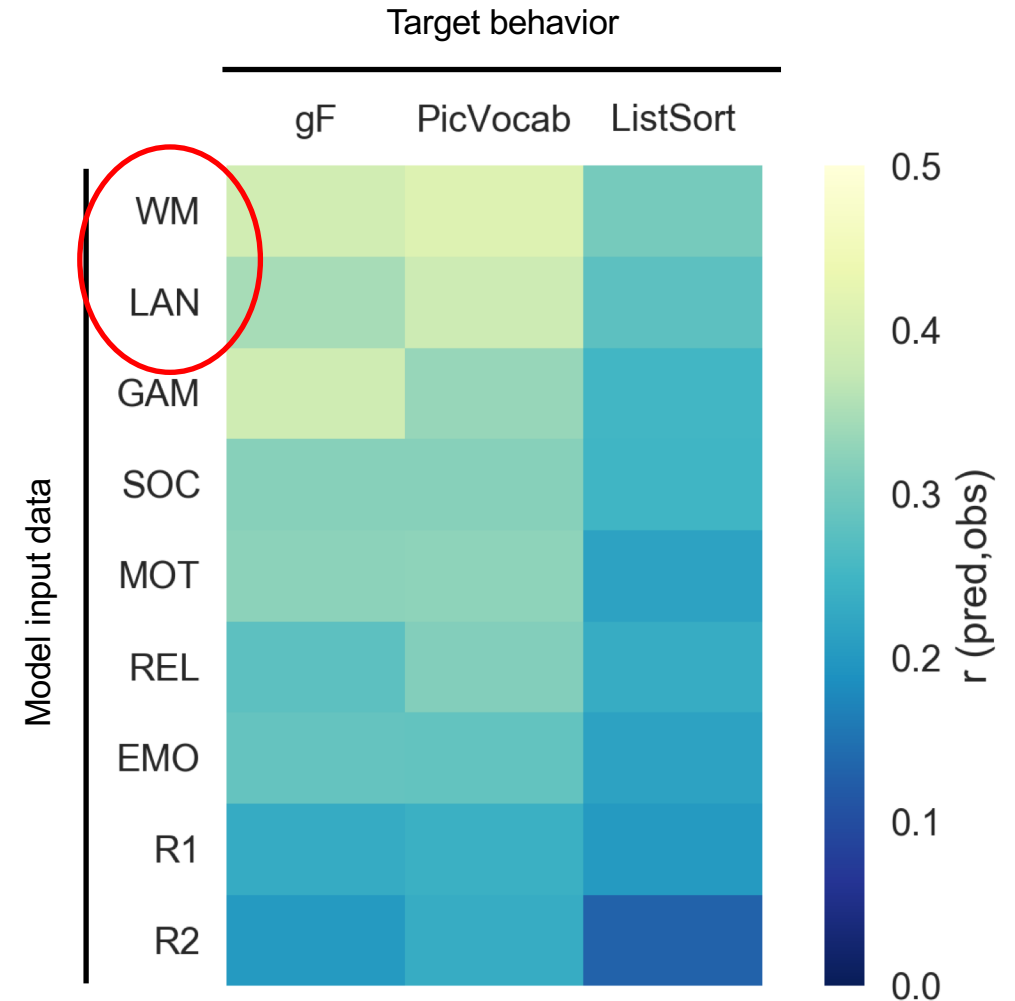
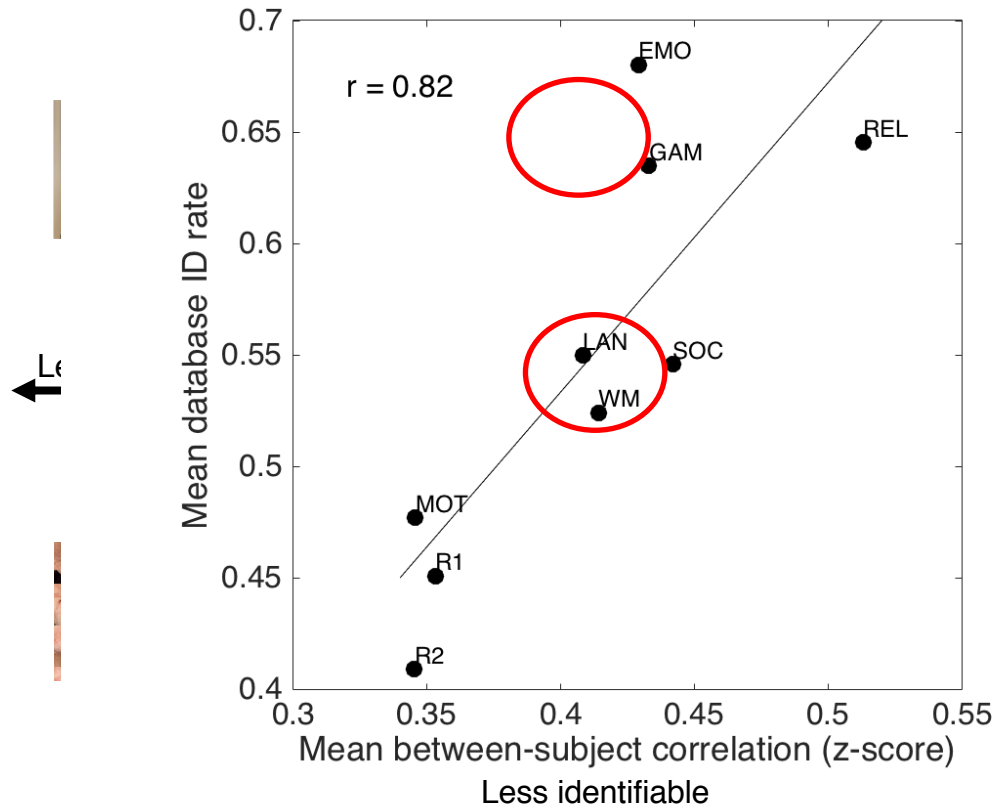


Vanderwal et al., NeuroImage (2015)

Q. What is the best brain state?

A. Maybe it depends on your behavior

Certain task conditions generate better predictions of behavior:



n = 716, 10-fold cross-validation
Connectome-based Predictive Modeling (CPM; Shen et al., *Nat Protocols* 2017)

Outline

1. **What** do we mean by individual differences?
2. **Why** should we care about individual differences?
3. **How** can we study individual differences?
(some practical guidelines)
4. **Where** are we going next?



Outline

1. **What** do we mean by individual differences?
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Outstanding questions

Data acquisition

Populations

- Many individuals lightly sampled, or few individuals densely sampled?
- Patients, controls?
- Longitudinal studies?

Imaging

- Scan condition? “Stress test”?
- Function, anatomy, both?

Behavior

- Robust measures?
- *Inter- vs intra-subject* variability?

Data analysis

Strategy

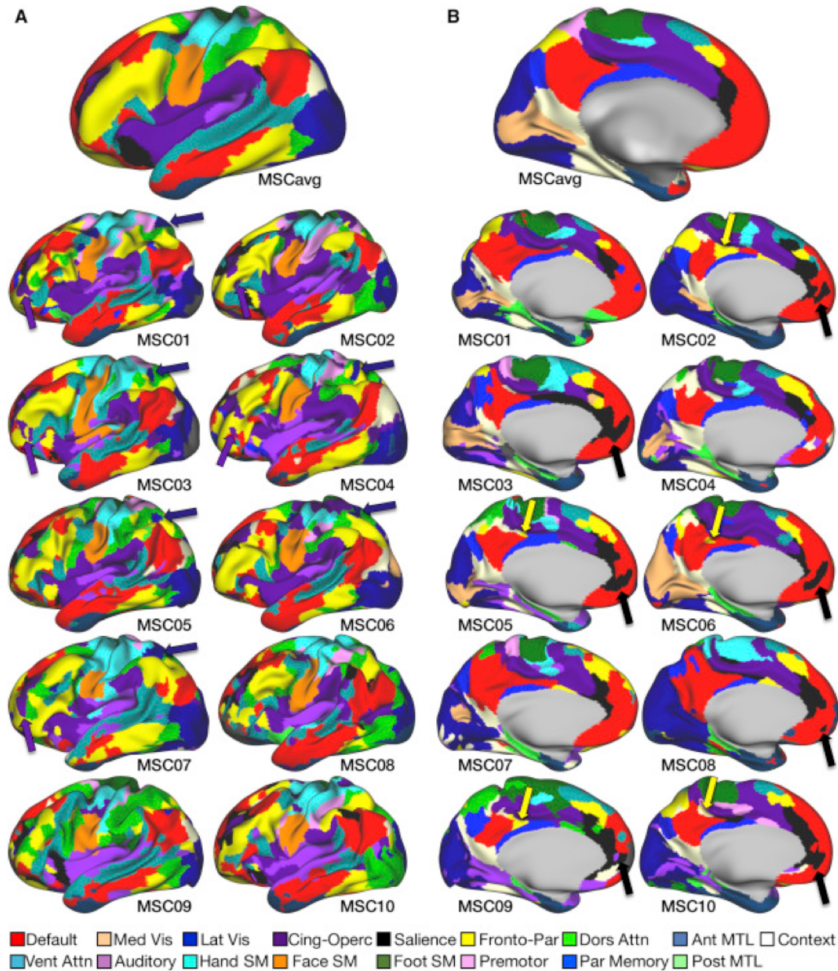
- Functional connectivity?
- Activation?
- Combination?

Specifics

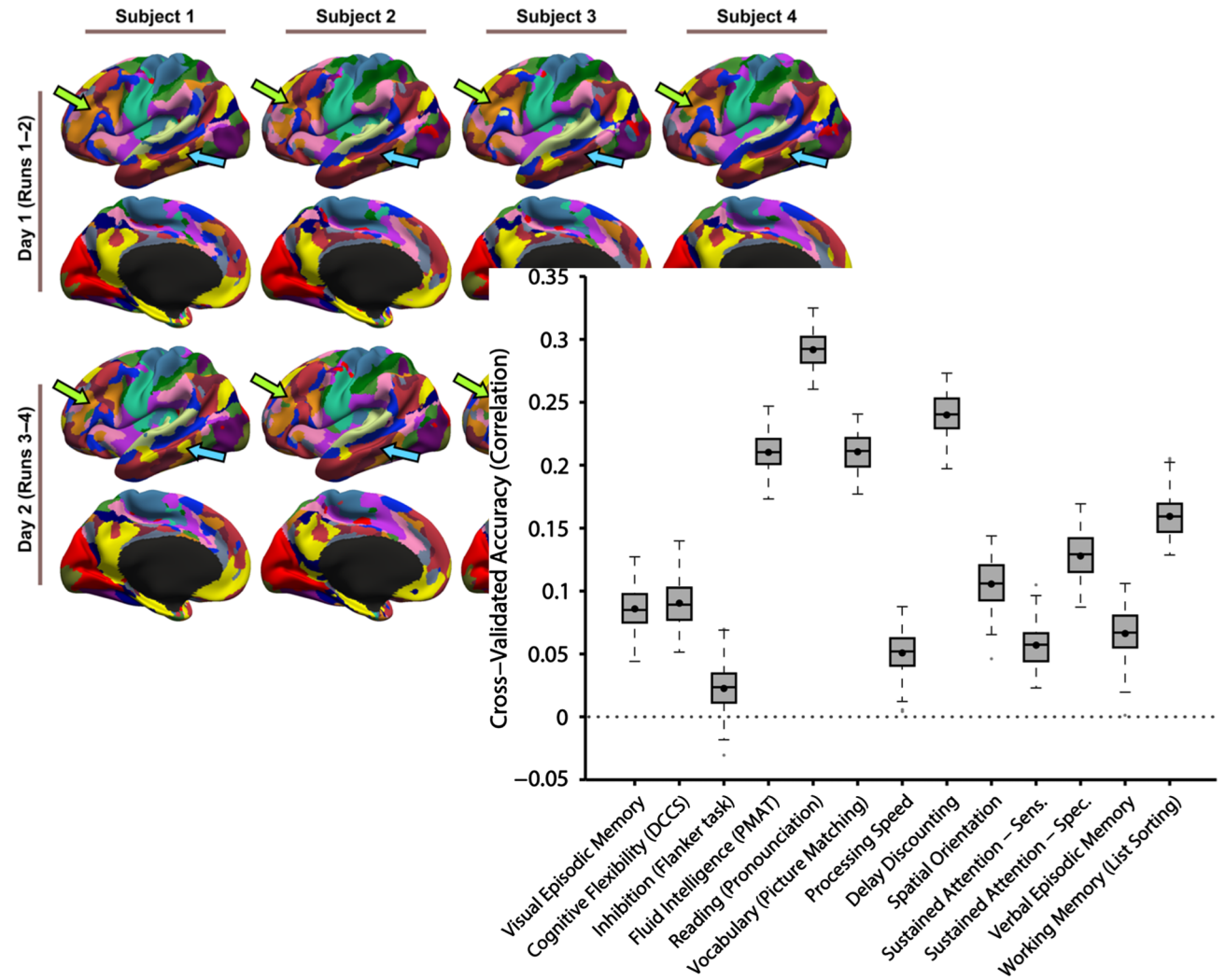
- Parcel boundaries?
Connections between parcels? Both?

Interpretations & applications

Individual-specific parcellations



Gordon et al., *Neuron* (2017)



Kong et al., *Cereb Cortex* (2018)

Outstanding questions

Data acquisition

Populations

- Many individuals lightly sampled, or few individuals densely sampled?
- Patients, controls?
- Longitudinal studies?

Imaging

- Multisite studies?
- Scan condition? “Stress test”?
- Function, anatomy, both?

Behavior

- Robust measures?
- *Inter-* vs *intra*-subject variability?

Data analysis

Strategy

- Functional connectivity?
- Activation?
- Combination?

Specifics

- Parcel boundaries?
Connections between parcels? Both?

Applications & interpretations

Mutability

- Development?
- Disease progression?
- Plasticity/training?

Applications

- Translational utility?
- Ethics?

Further reading & open data sets

Selected reviews:

Prediction as a humanitarian and pragmatic contribution from human cognitive neuroscience

Gabrieli, Ghosh & Gabrieli, *Neuron* (2015)

Building a science of individual differences from fMRI

Dubois & Adolphs, *Trends in Cognitive Sciences* (2016)

From regions to connections and networks: new bridges between brain and behavior

Misic & Sporns, *Current Opinion in Neurobiology* (2016)

Can brain state be manipulated to emphasize individual differences in functional connectivity?

Finn et al., *NeuroImage* (2017)

Open data sets with brain and behavior:



Use these on their own or in combination with your own data to generate or test hypotheses, see if a finding generalizes, etc

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