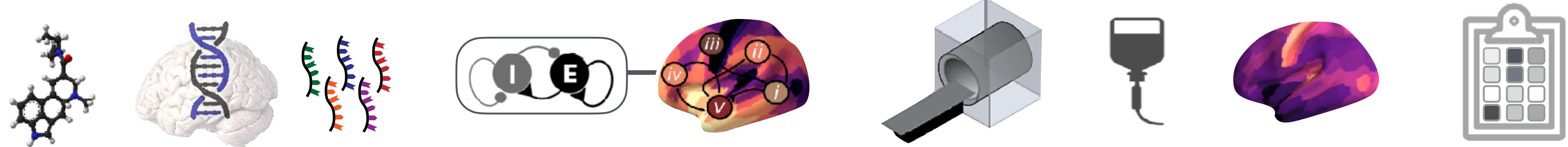


# Modeling Connectivity: Biophysical Network Modeling



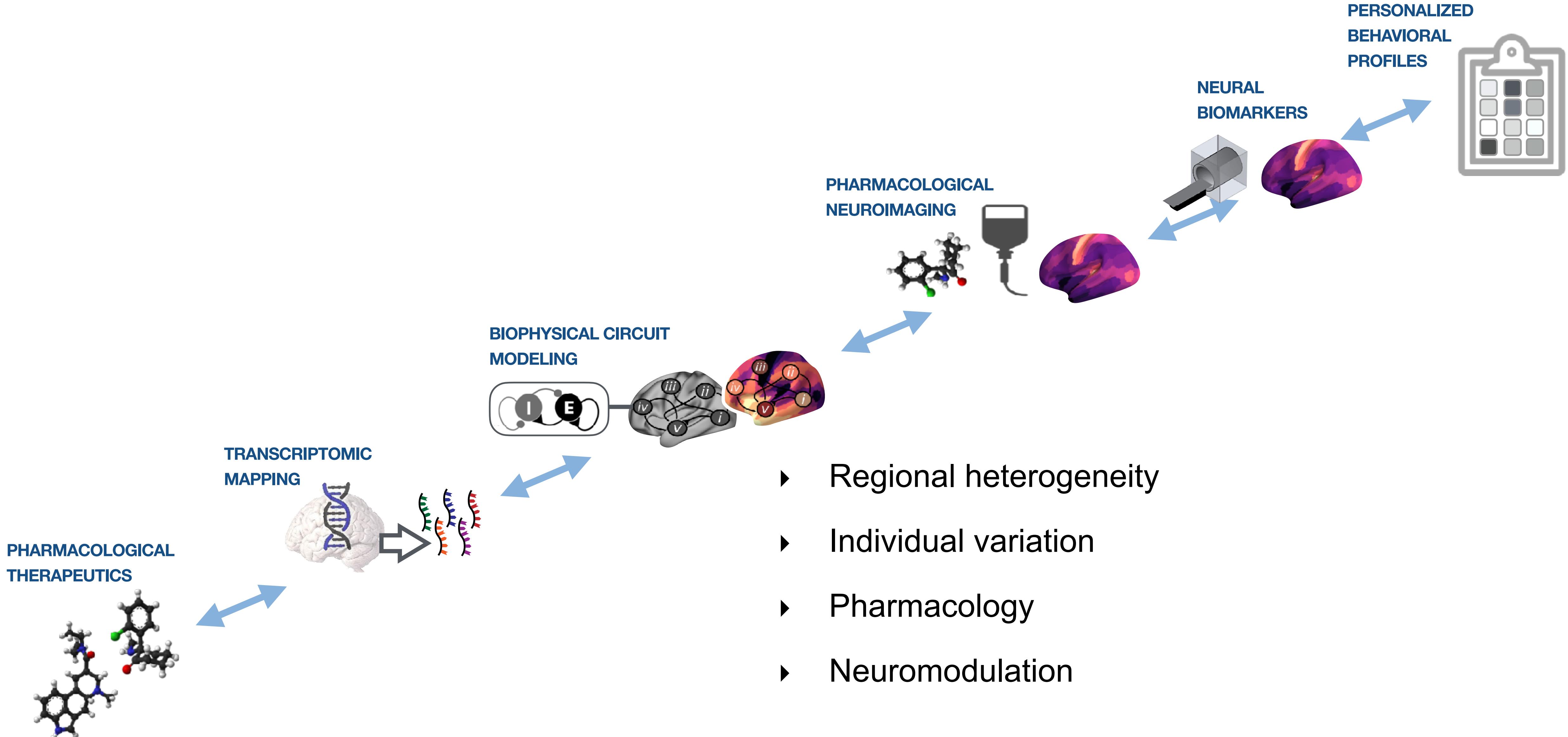
John D. Murray



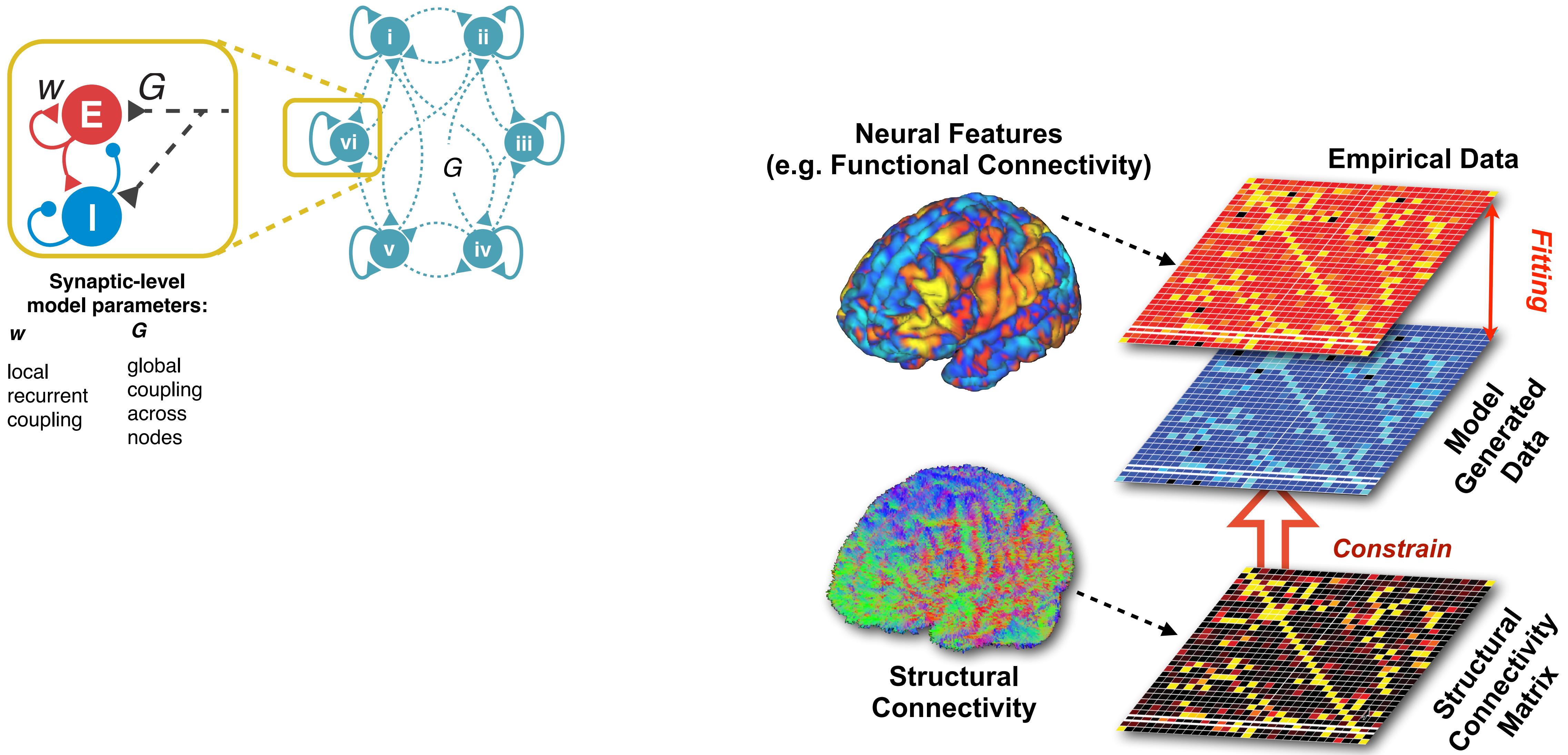
Division of Neurocognition,  
Neurocomputation, and  
Neurogenetics (N3)

Department of Psychiatry  
Yale School of Medicine

# Computational Psychiatry via bridging levels of analysis with neuroimaging and neural circuit modeling

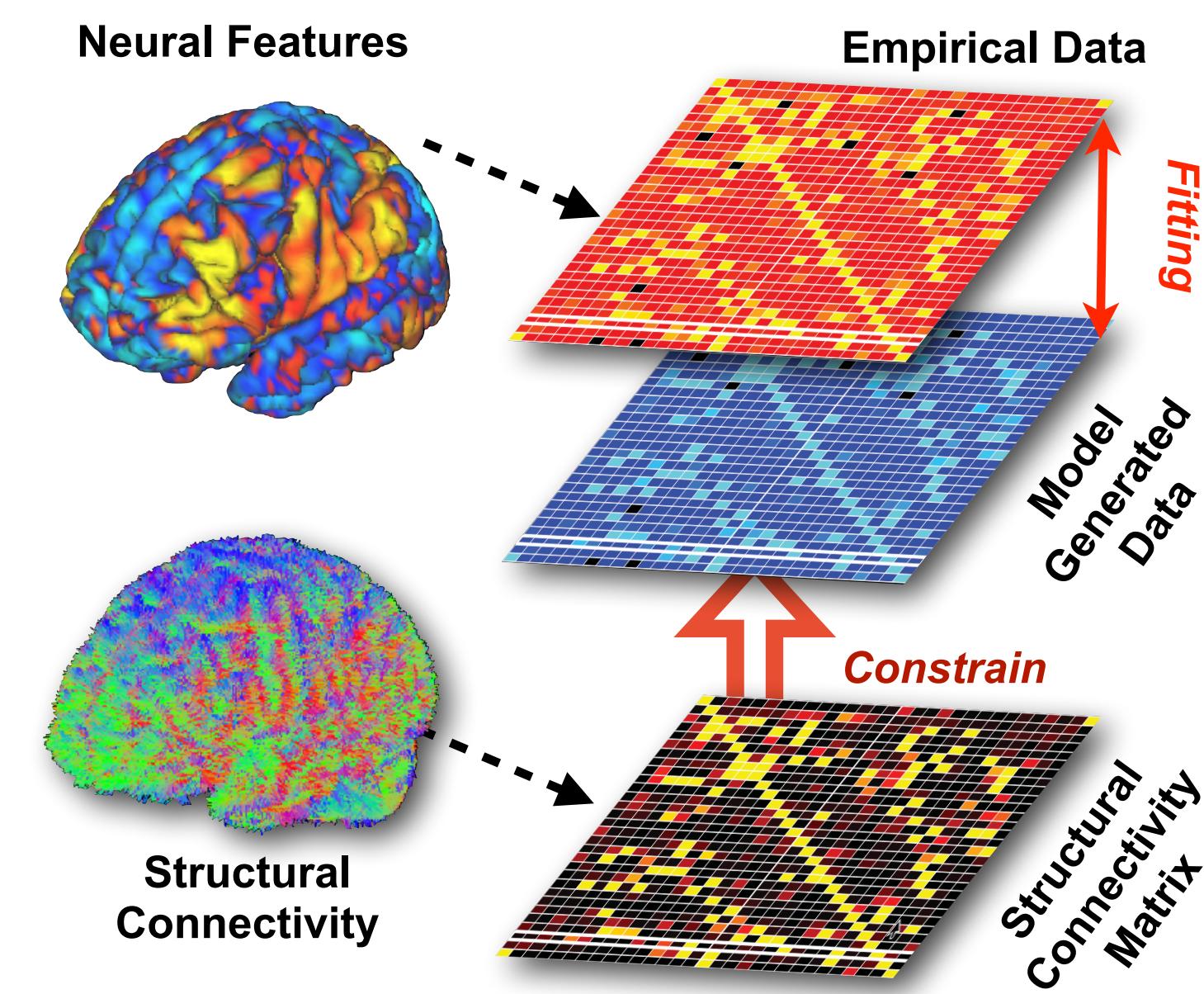
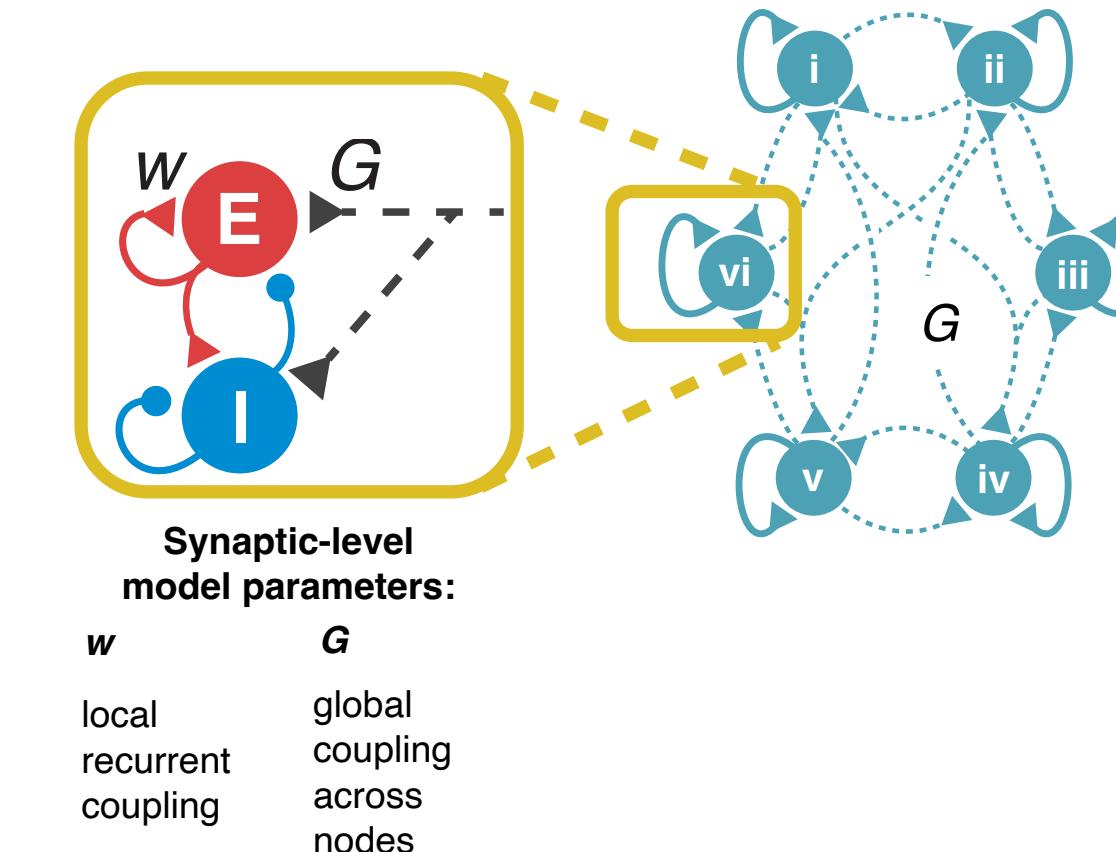


# Biophysical modeling of large-scale resting-state networks

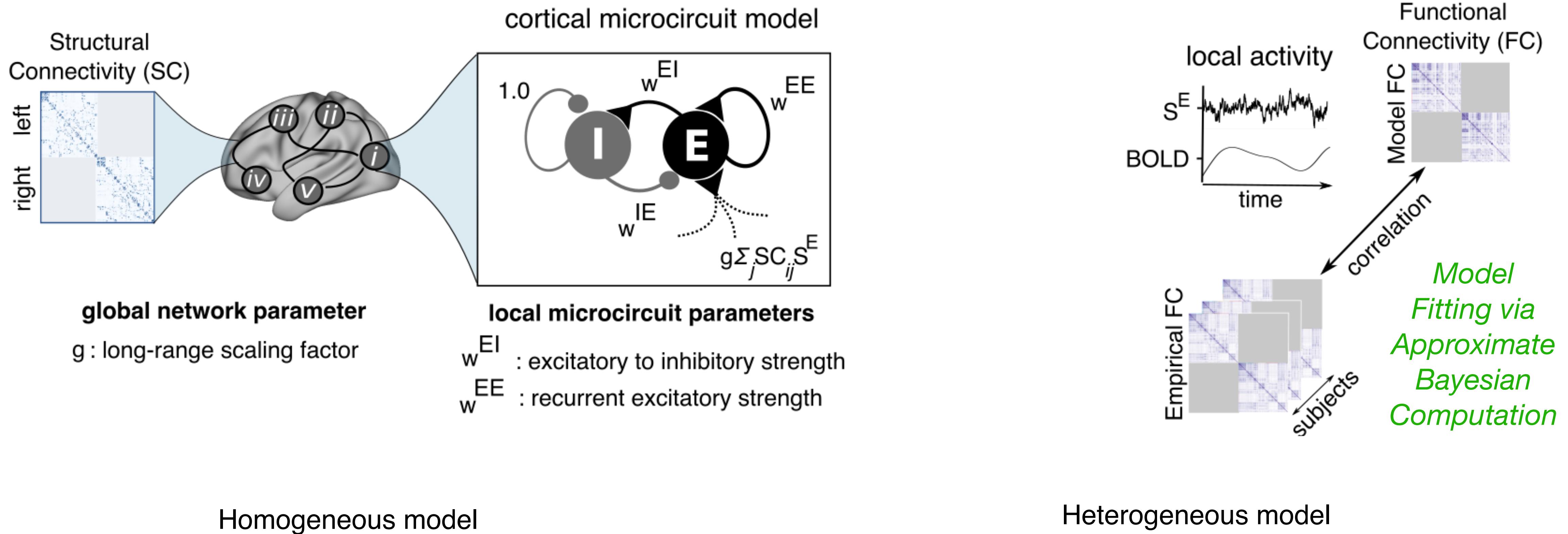


# Biophysical modeling of large-scale resting-state networks

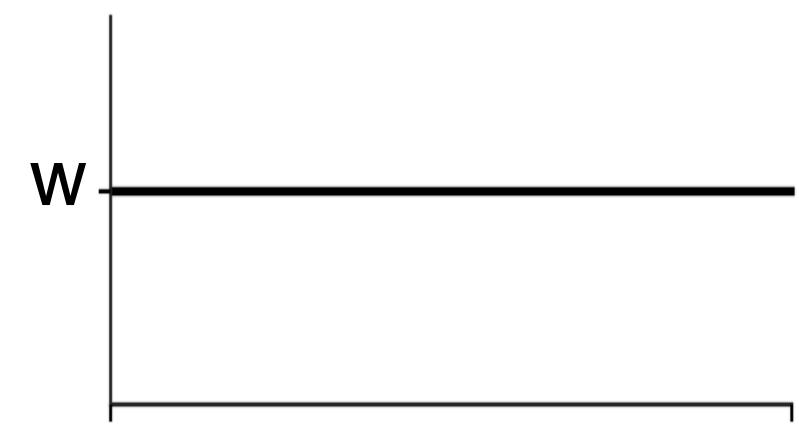
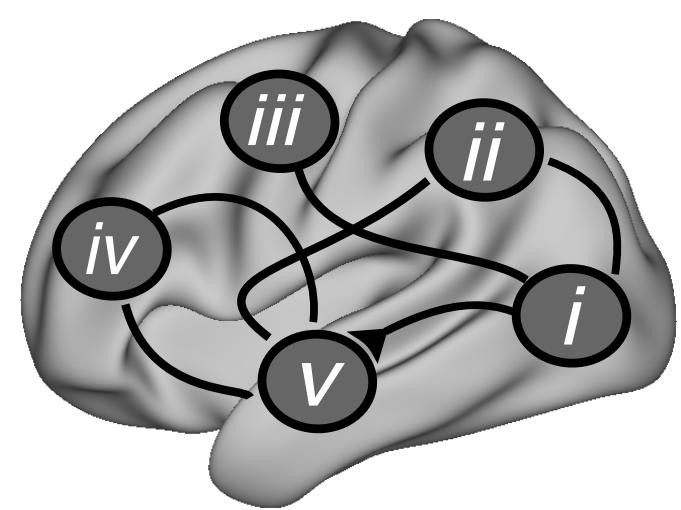
- ▶ Need to select local node model
- ▶ Depends on phenomena and parameters of interest
- ▶ Typically long-range connectivity pattern is fixed
- ▶ Fewer fit parameters, goal is not to get a great fit to data
- ▶ Key aspect of models is that parameters are neurobiologically interpretable
- ▶ Examples of types of questions one can pursue:
  - ▶ How does variation in local circuit properties across brain areas produce interesting features of empirical data?
  - ▶ How does a hypothesized synaptic-level perturbation produce systems-level effects observed in empirical data?



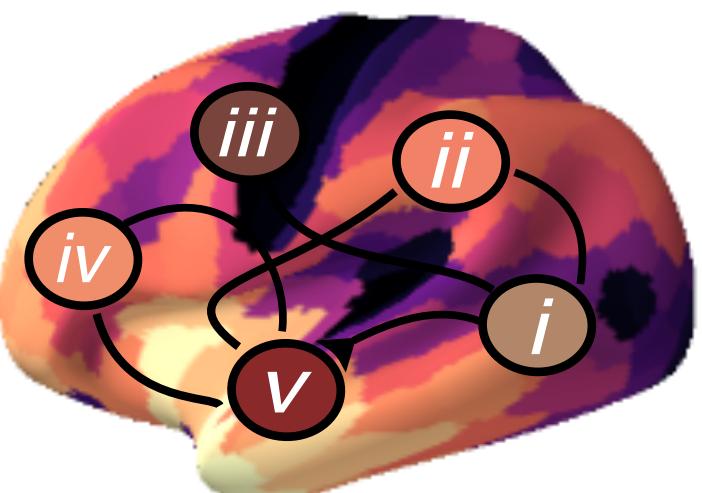
# Model with heterogeneous local circuit properties



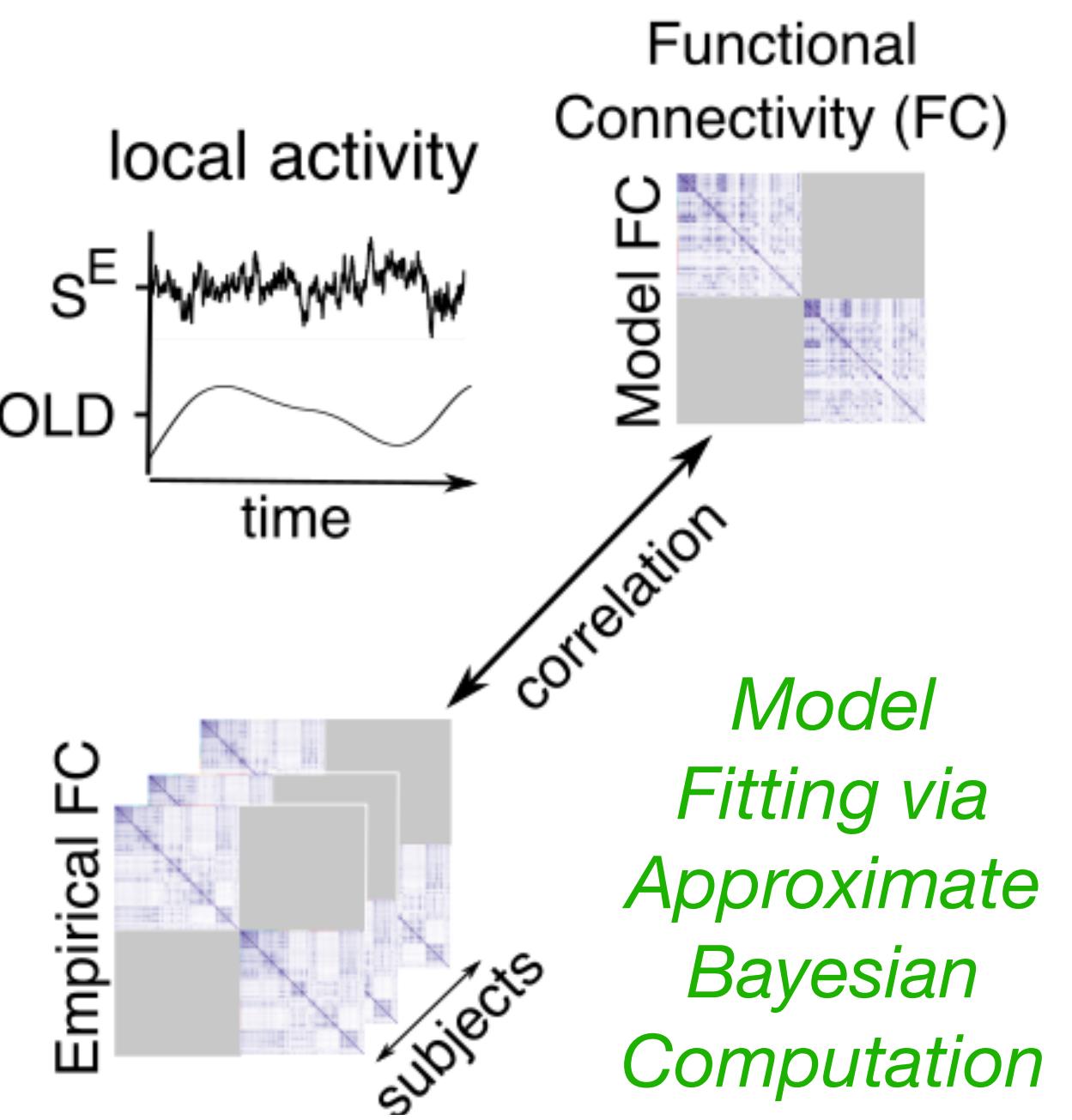
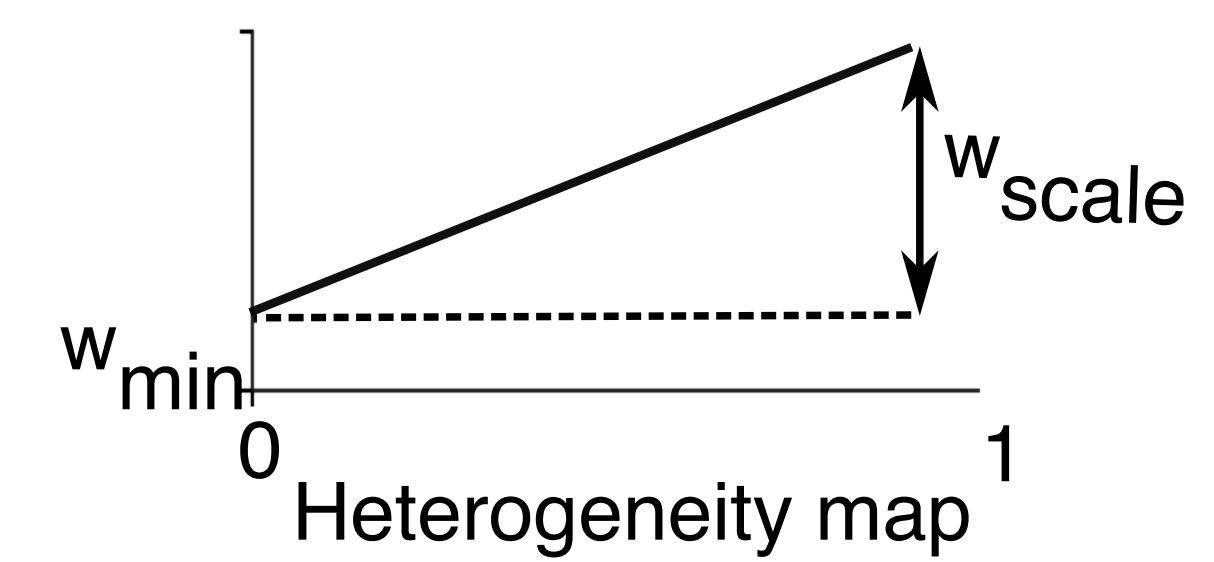
Homogeneous model



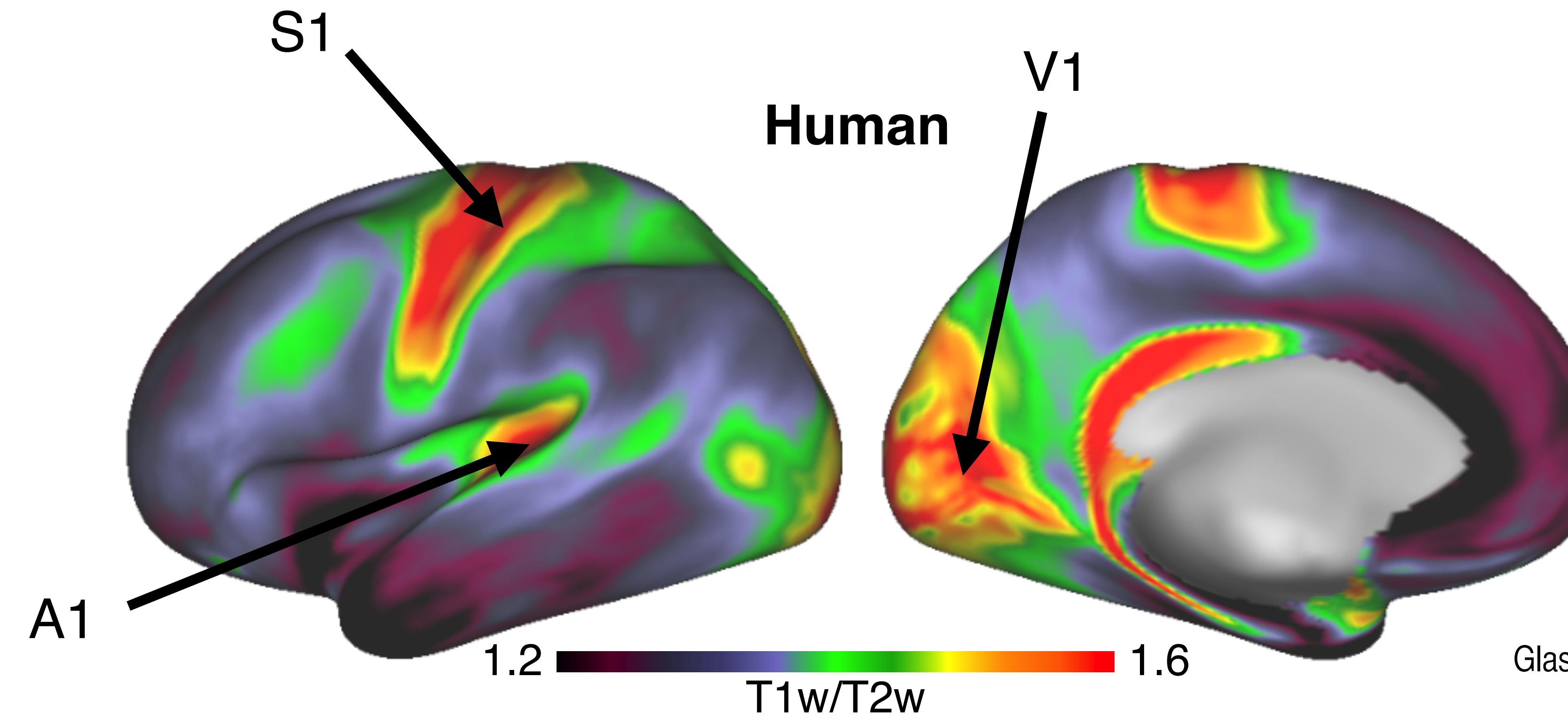
Heterogeneous model



Reverse  $T1w/T2w$   
myelin map



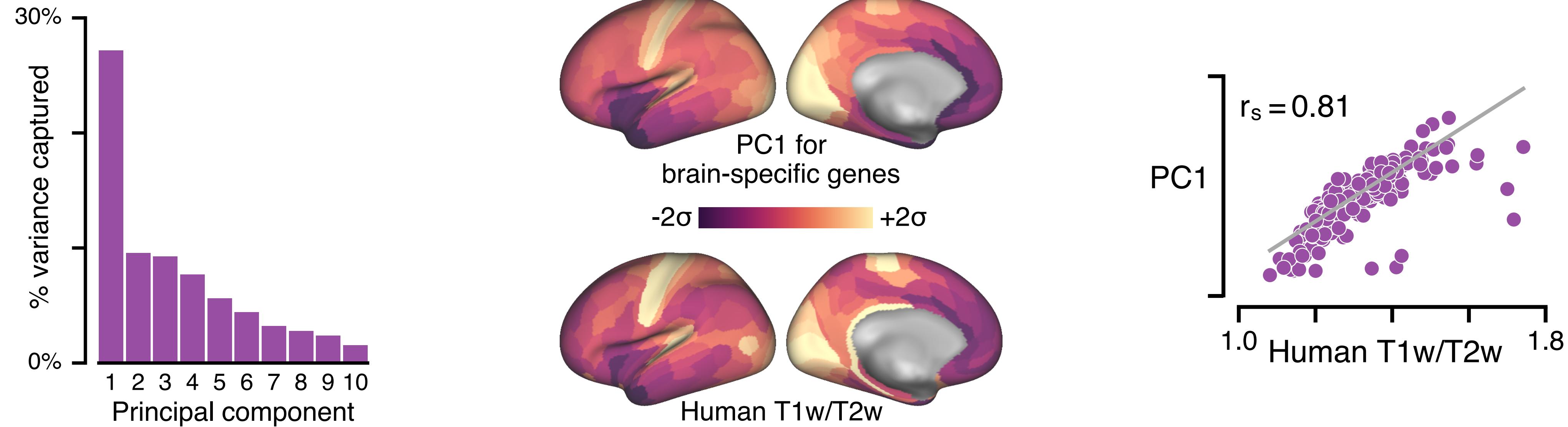
# Cortical T1w/T2w map as a proxy measure for cortical hierarchy



Glasser et al. (2011)  
*J Neurosci*

Burt et al. (2018) *Nature Neurosci*

# T1w/T2w topography captures the dominant axis of transcriptional variation across human cortex



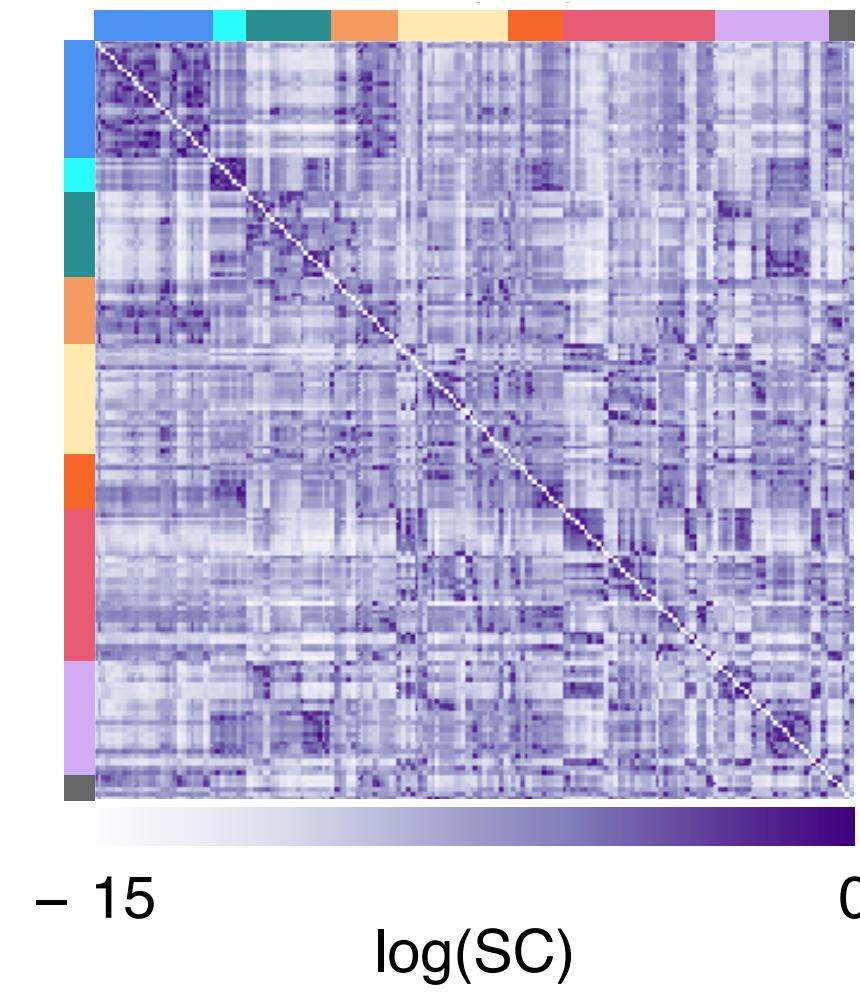
- ▶ Cortical gene expression variation is dominated by a single principal axis which is highly correlated with the T1w/T2w map
- ▶ Hierarchical gradients of microscale properties may contribute to sensory–association specialization of cortical function.



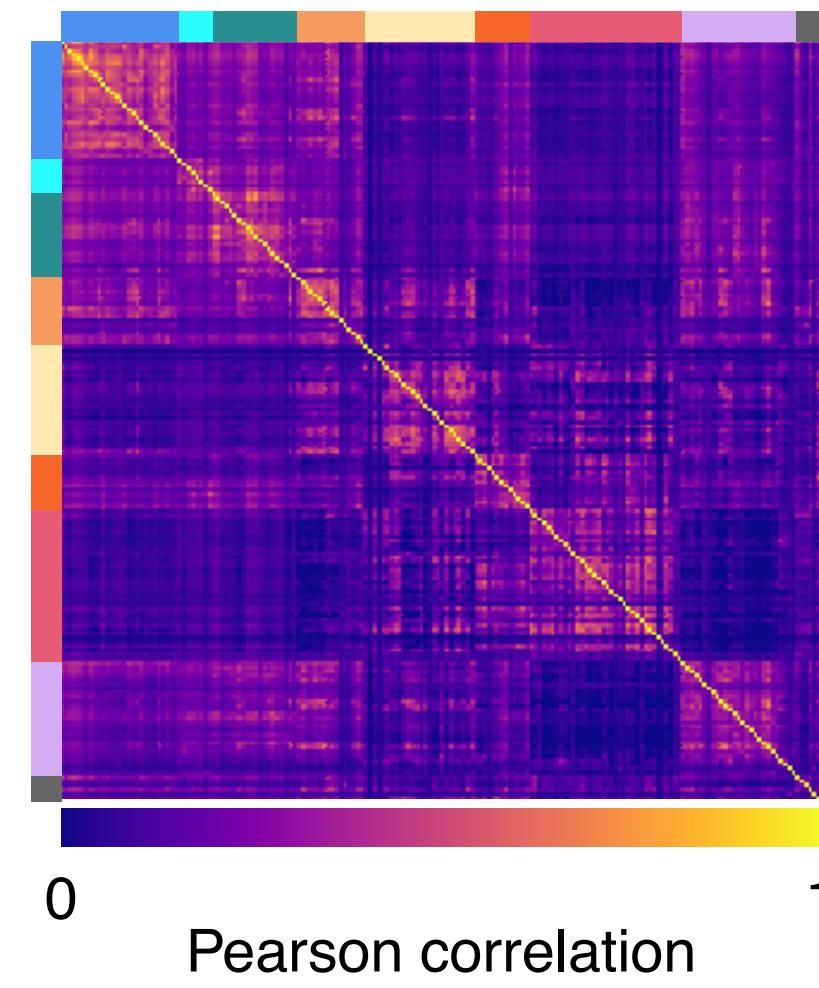
# Simulated functional connectivity in large-scale models

Murat Demirtaş

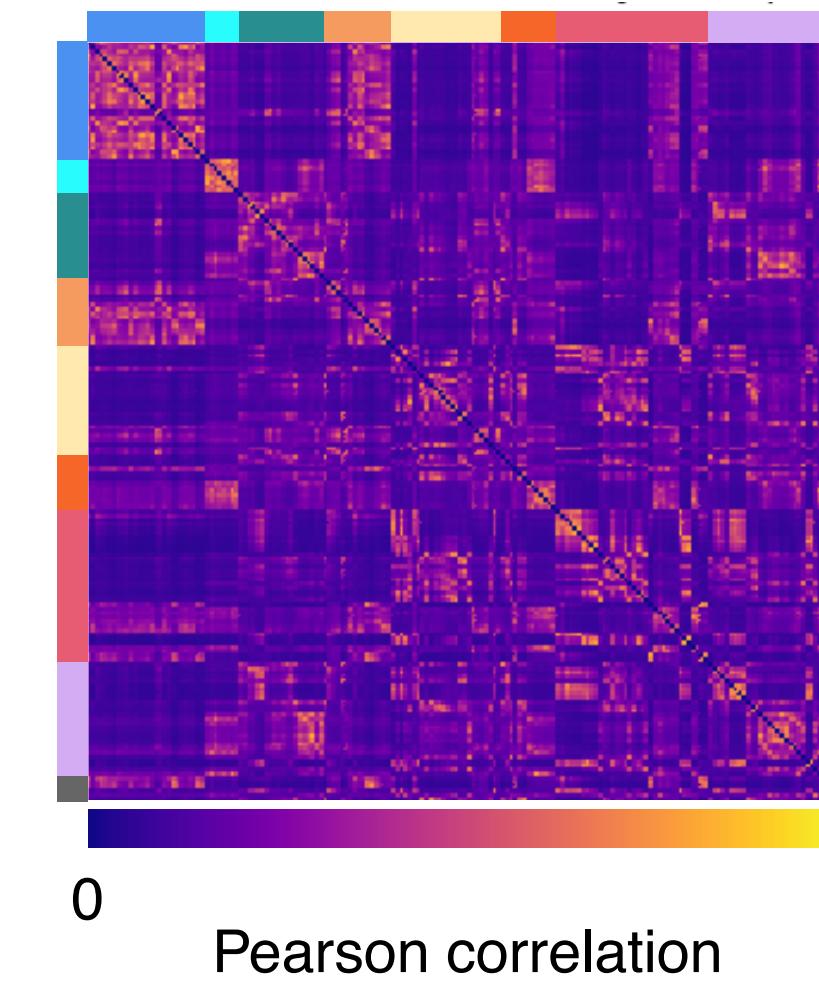
**Empirical structural connectivity (SC)**



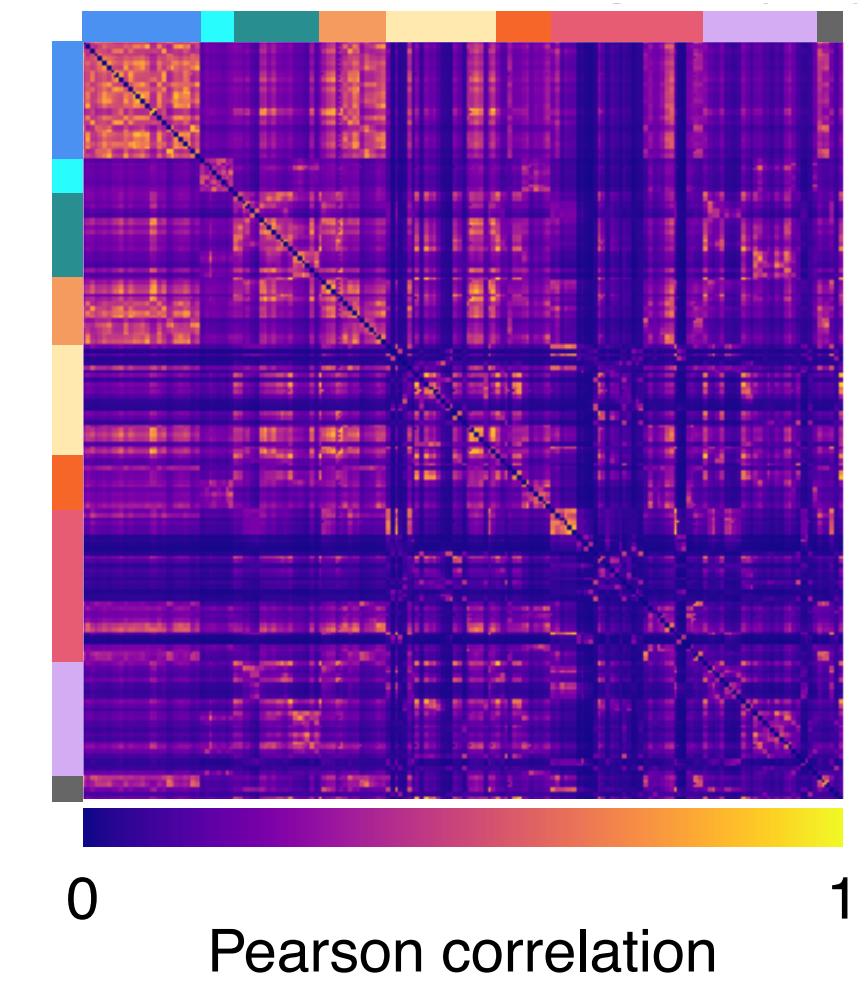
**Empirical functional connectivity (FC)**



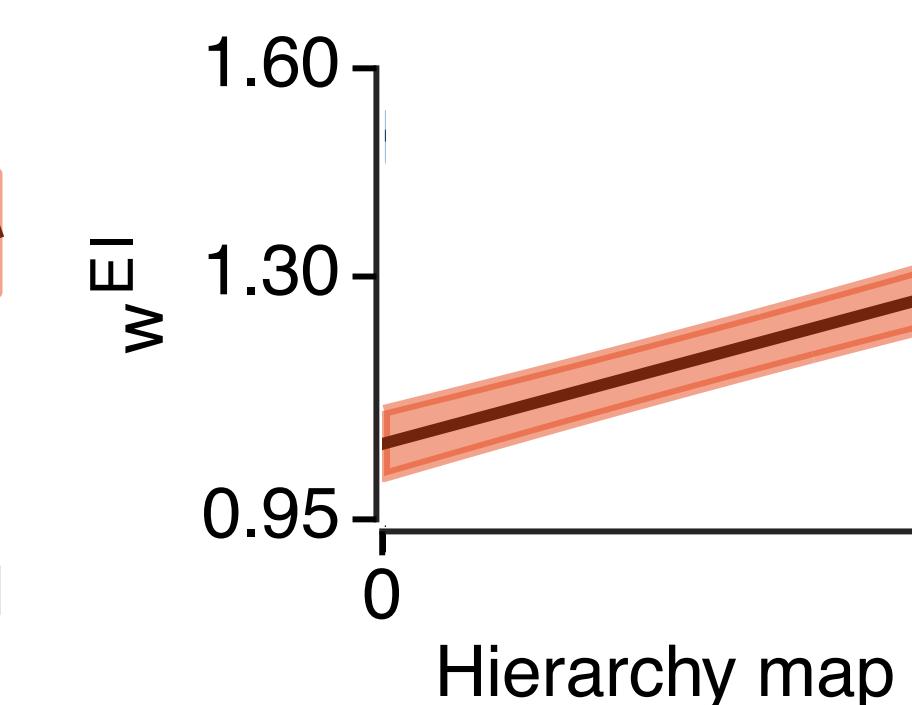
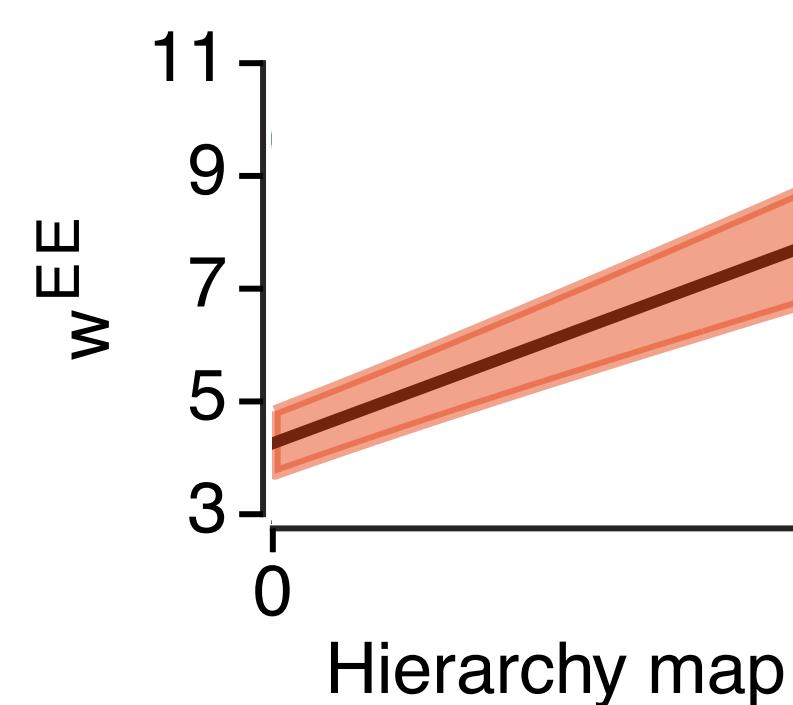
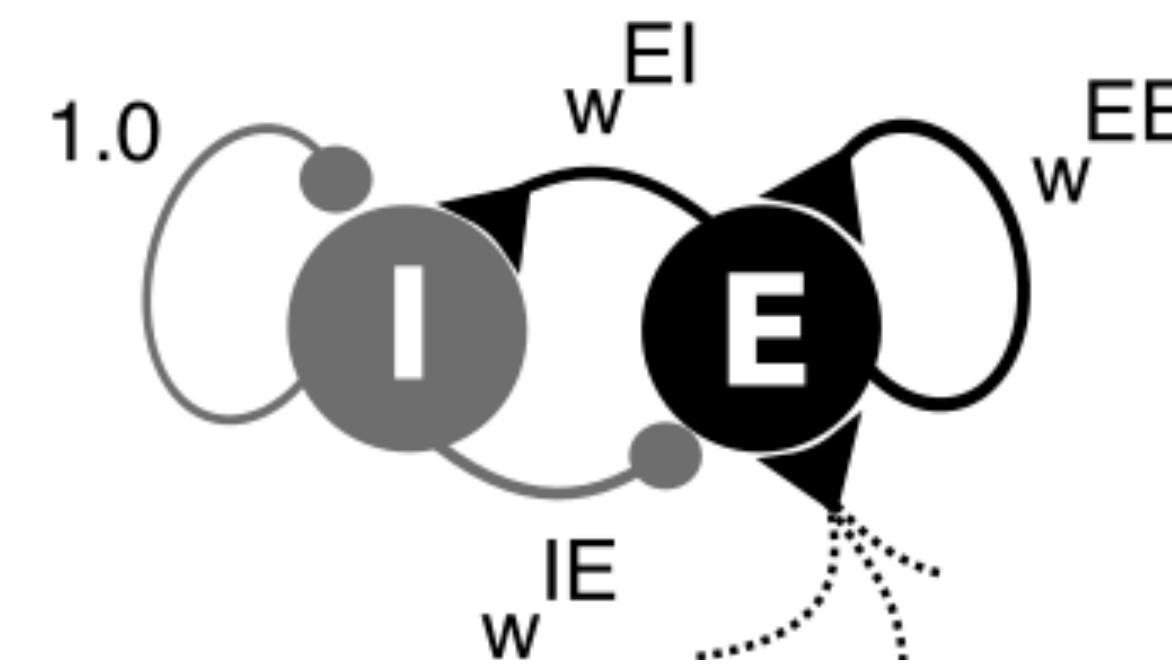
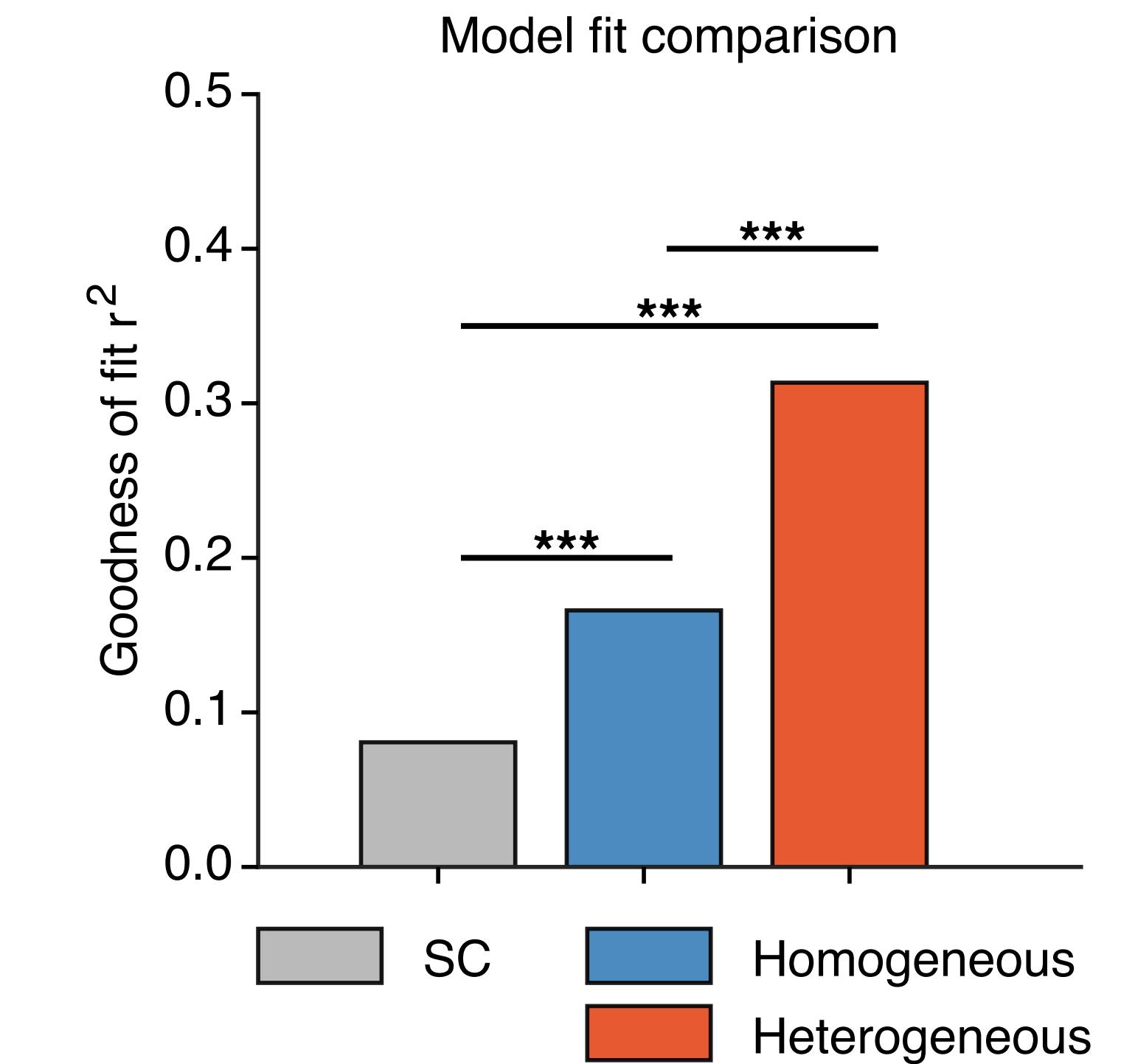
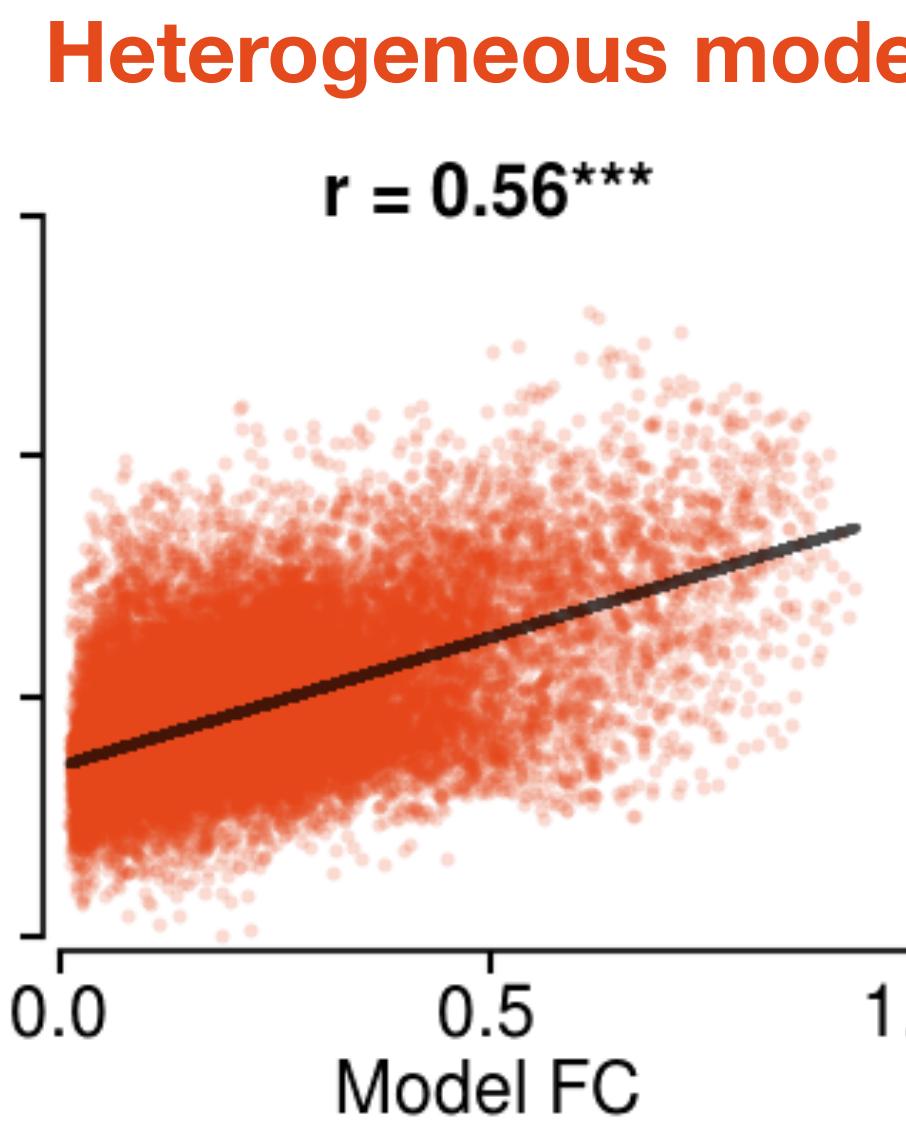
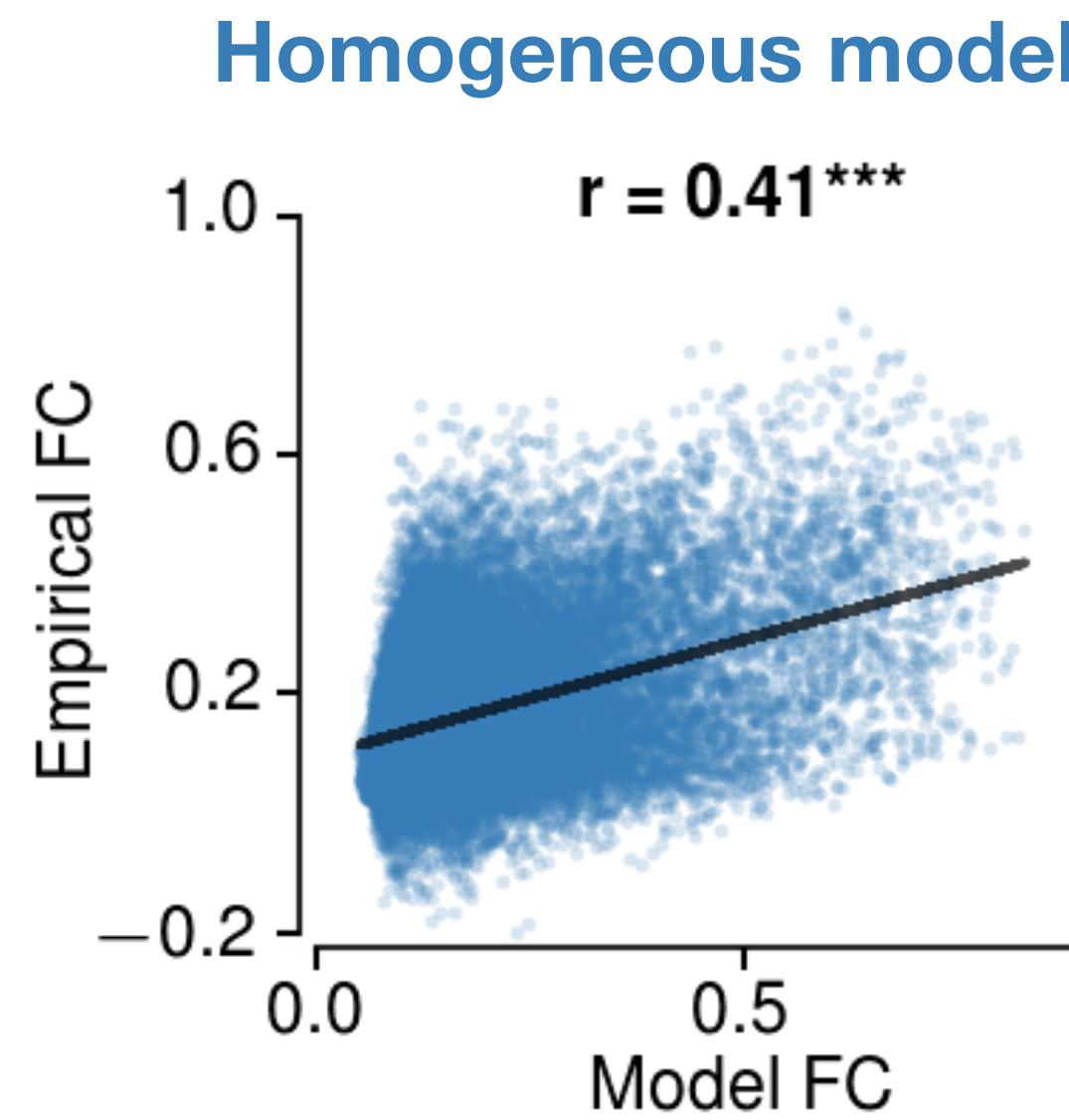
**Homogeneous model FC**



**Heterogeneous model FC**



# Hierarchical heterogeneity improves model fit to functional connectivity

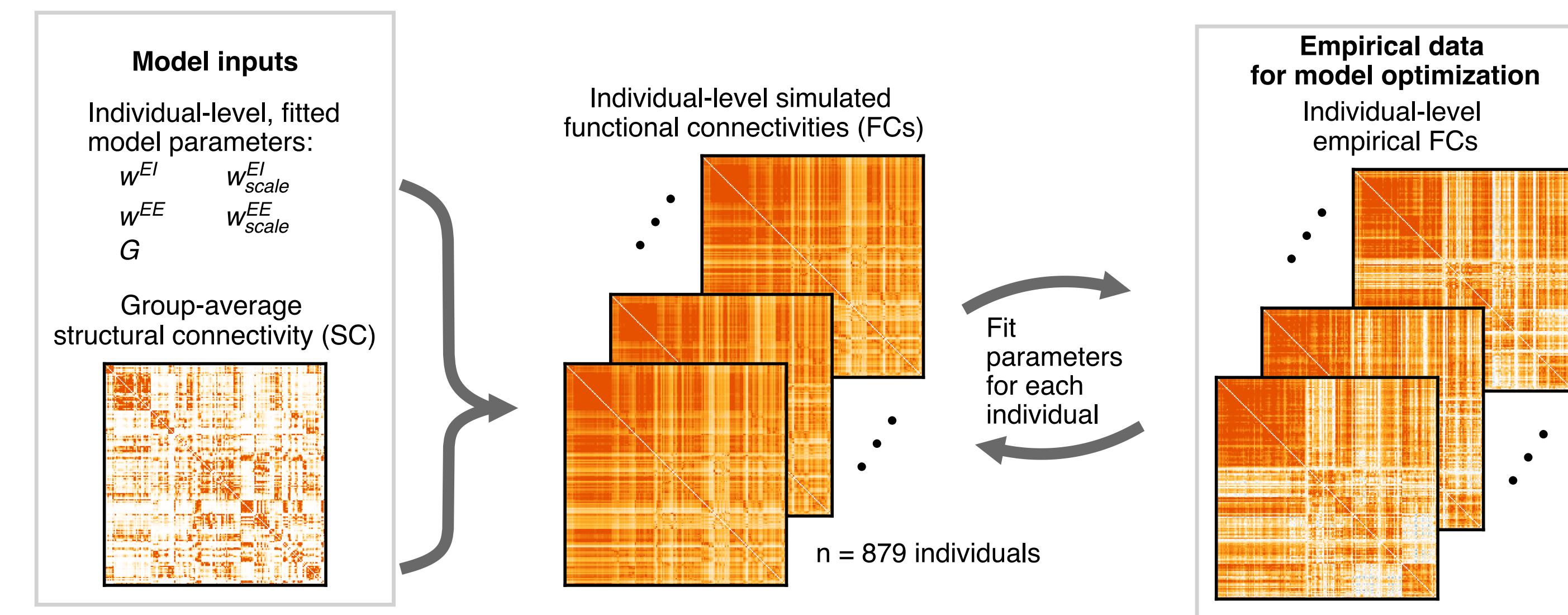
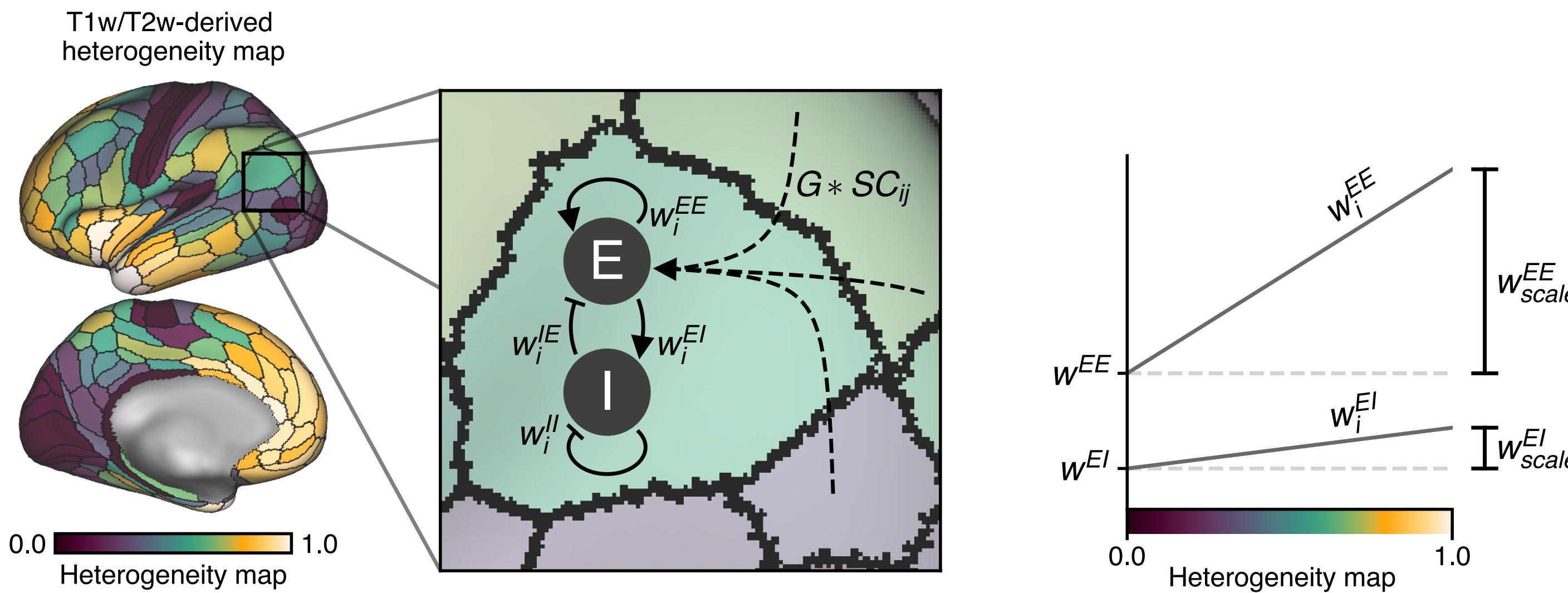


Increasing local  
recurrent strength  
along cortical  
hierarchy

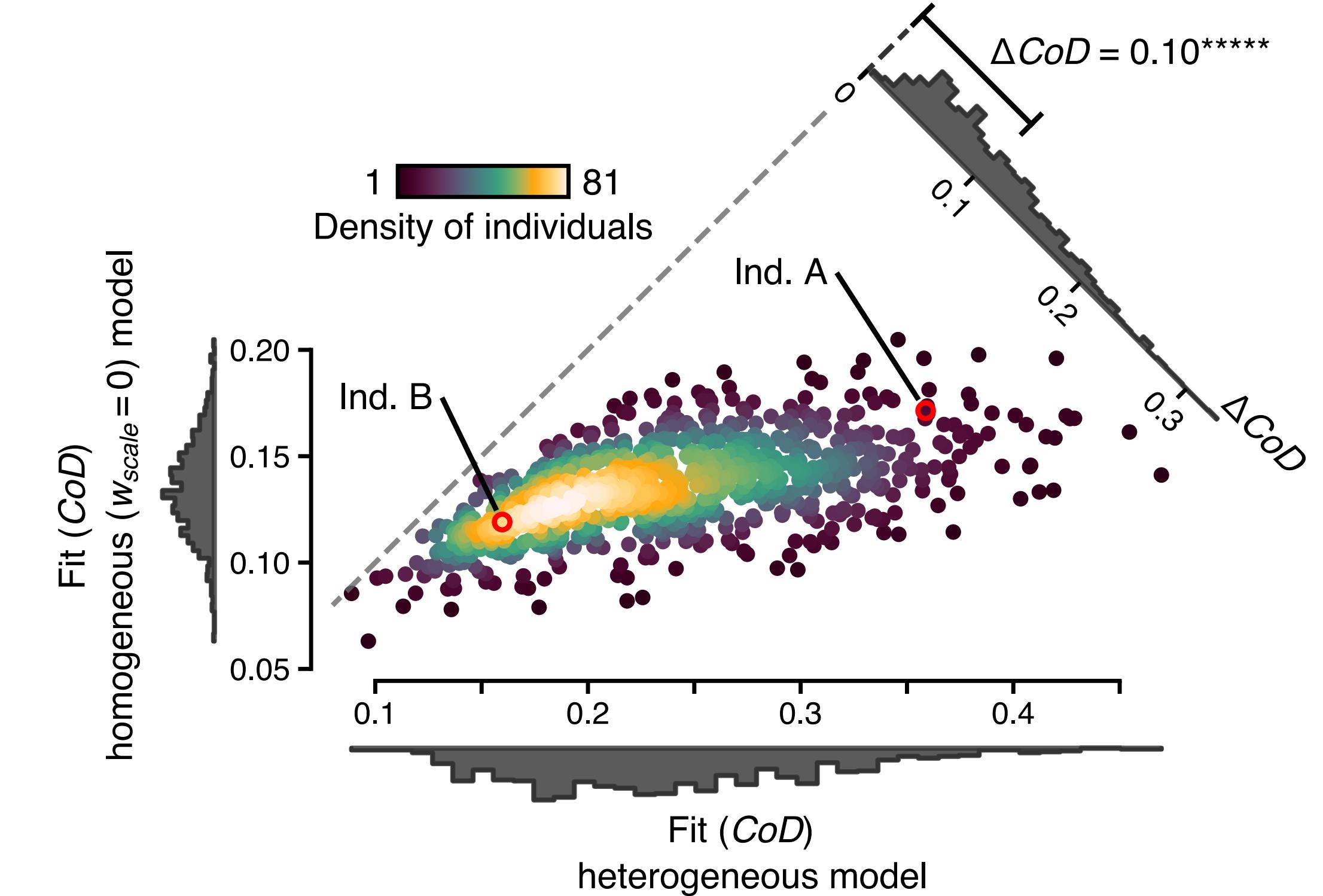
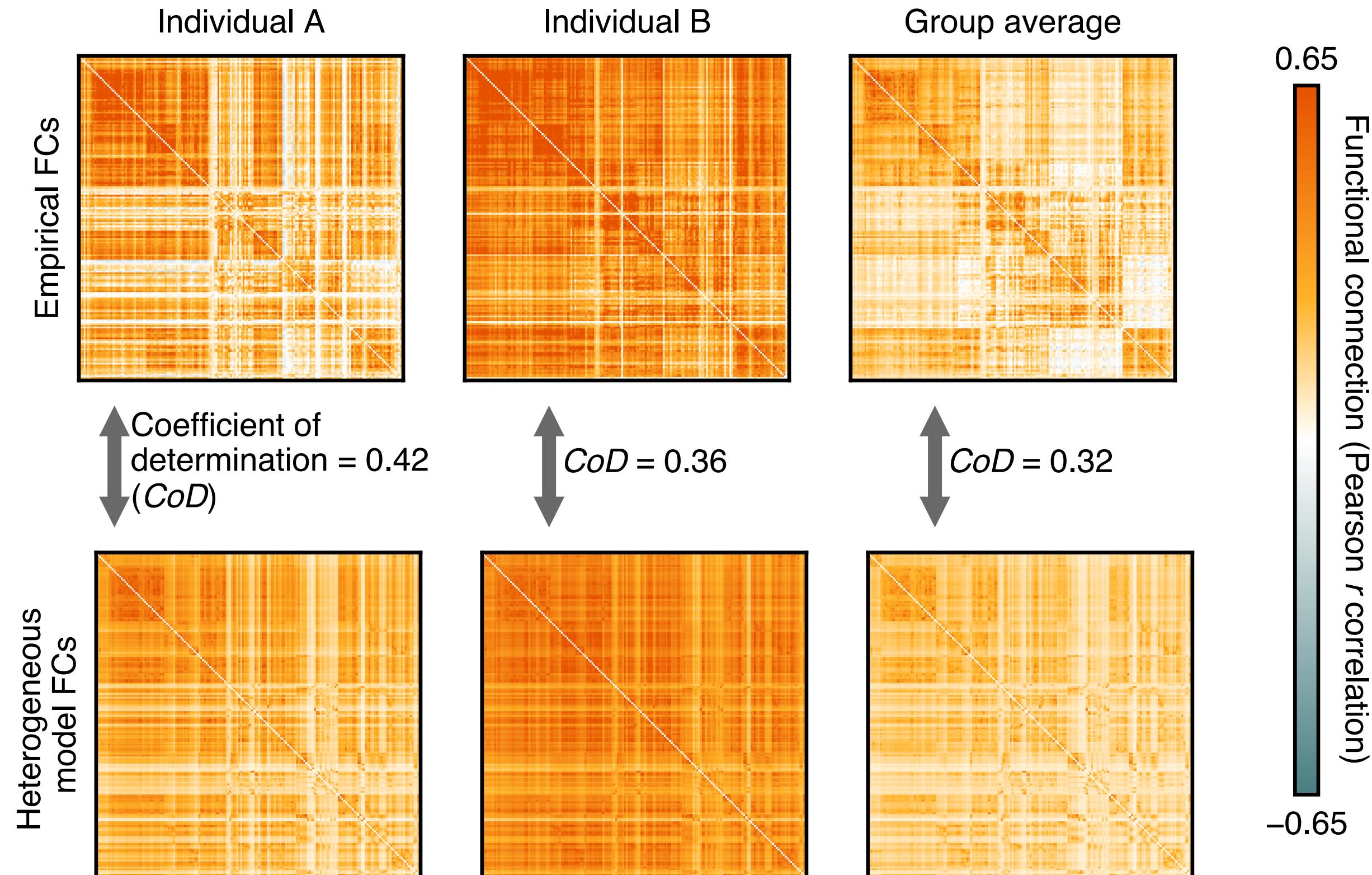


# Individual-subject modeling of functional connectivity

Rachel Cooper

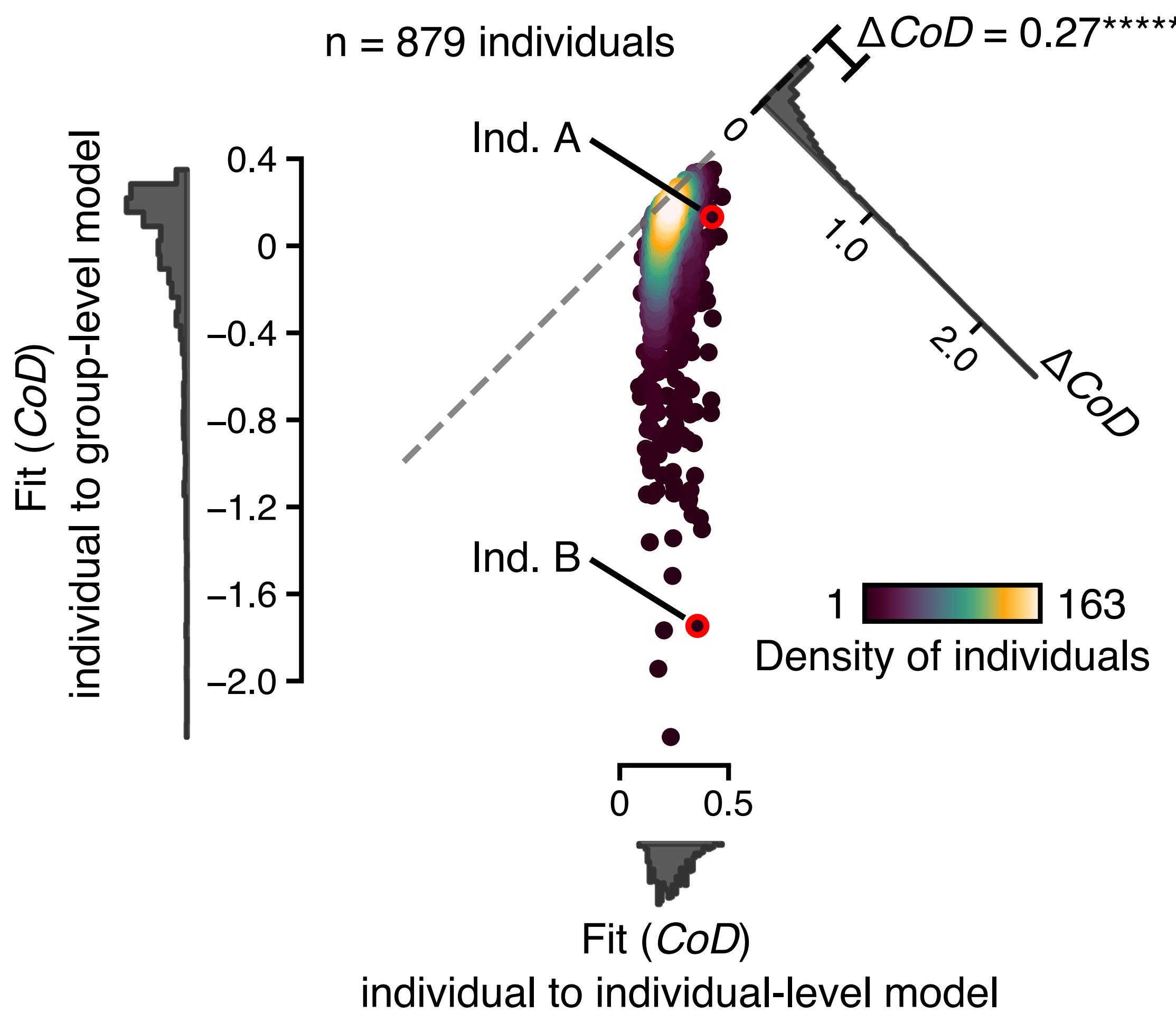


# Individual-subject modeling of functional connectivity

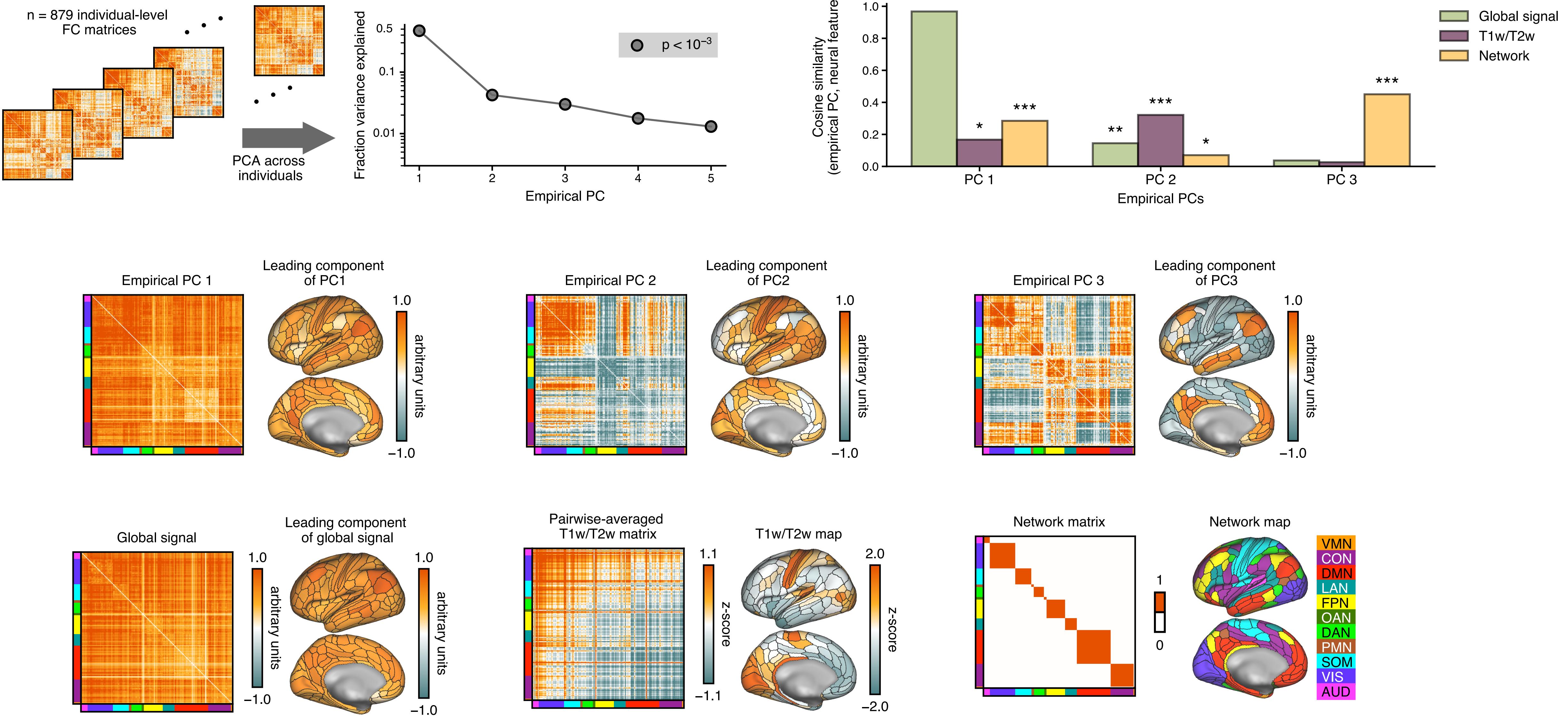


*Fit measure:* Coefficient of Determination (COD) =  $1 - \frac{MSE(\text{predicted, observed})}{MSE(\text{predicted}, \hat{\mu})}$

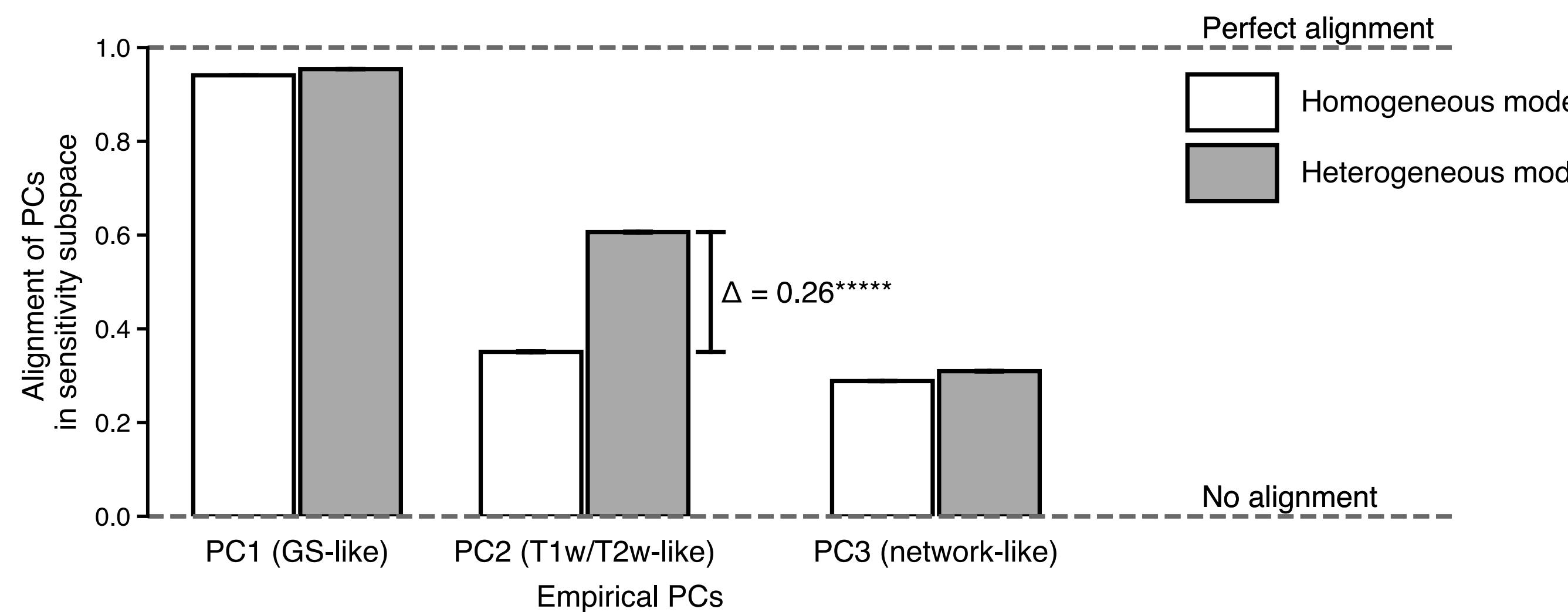
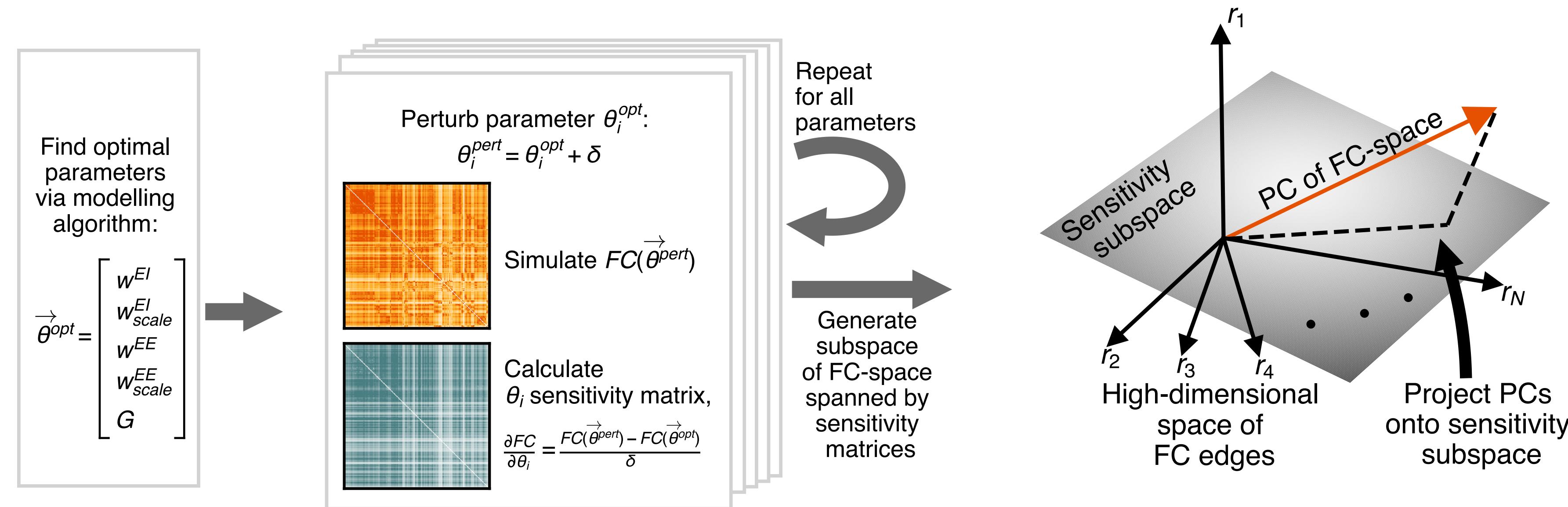
# Individual-subject modeling of functional connectivity



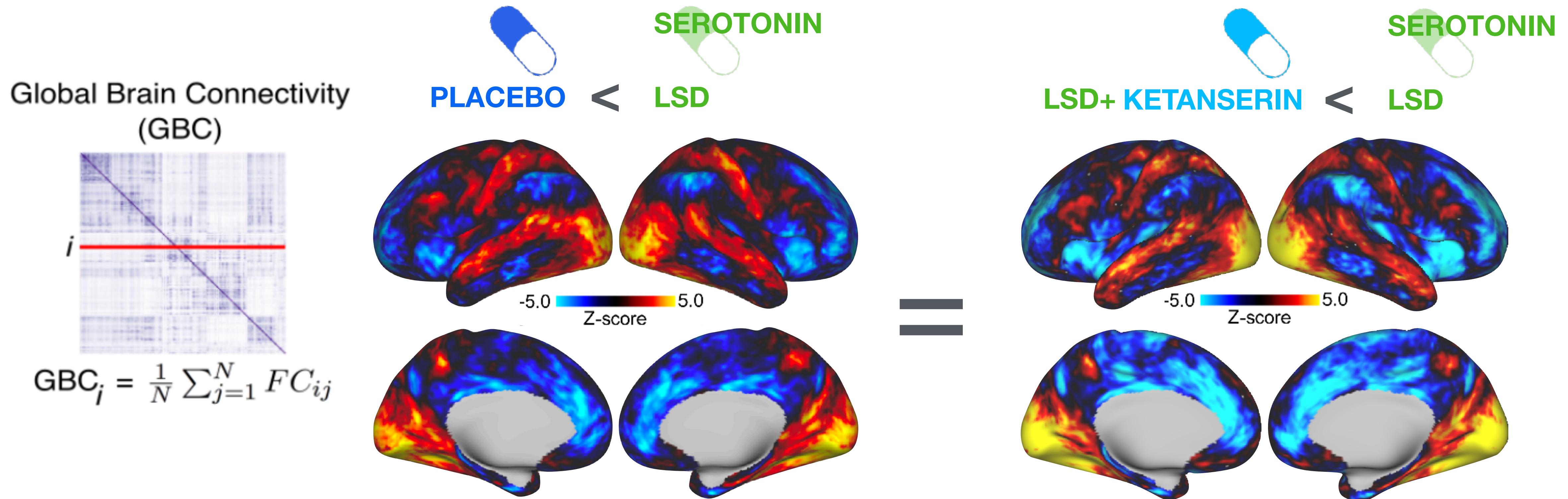
# Modes of population variation



# Model expressivity to capture modes of population variation



# LSD effects on resting-state functional connectivity

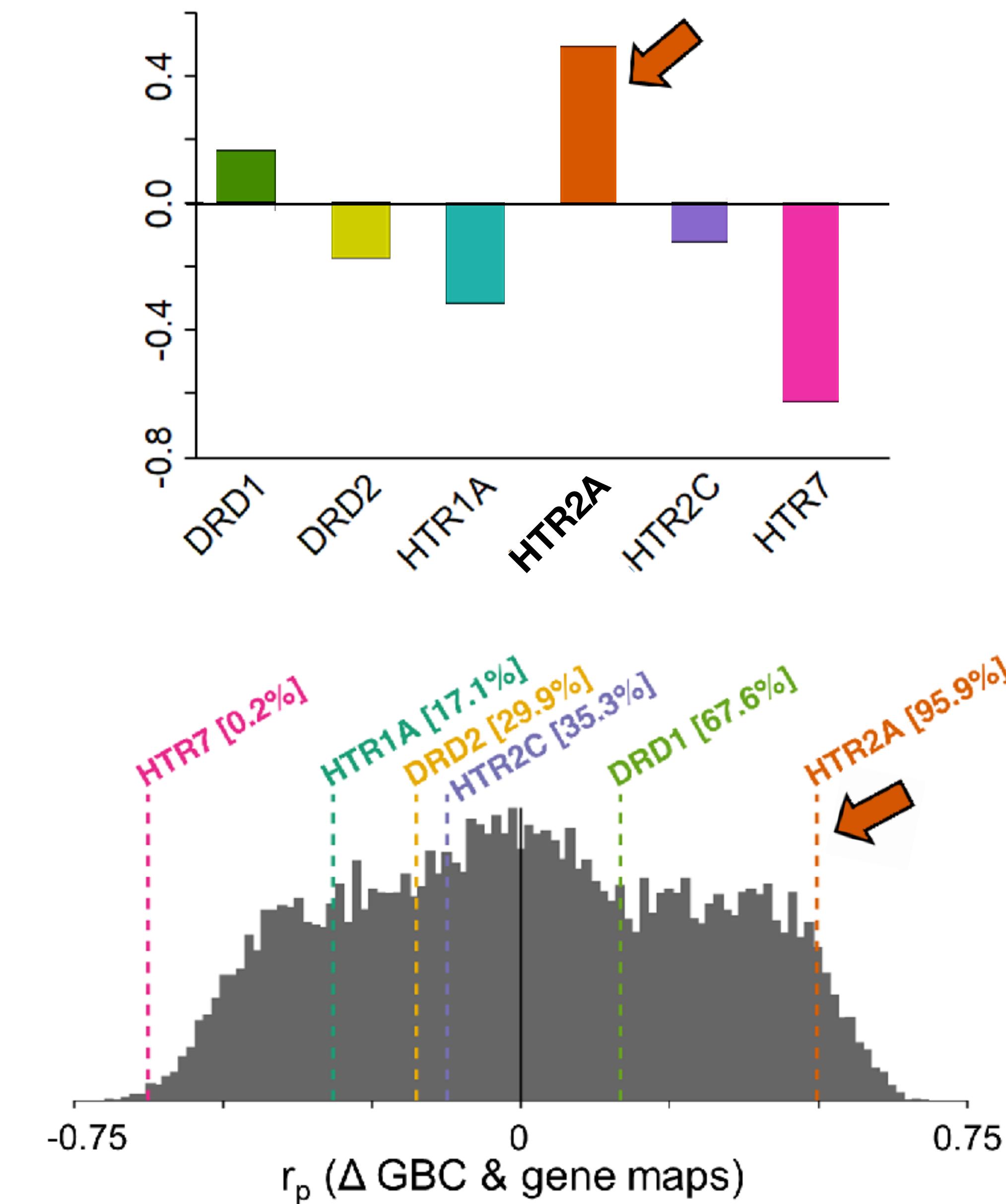
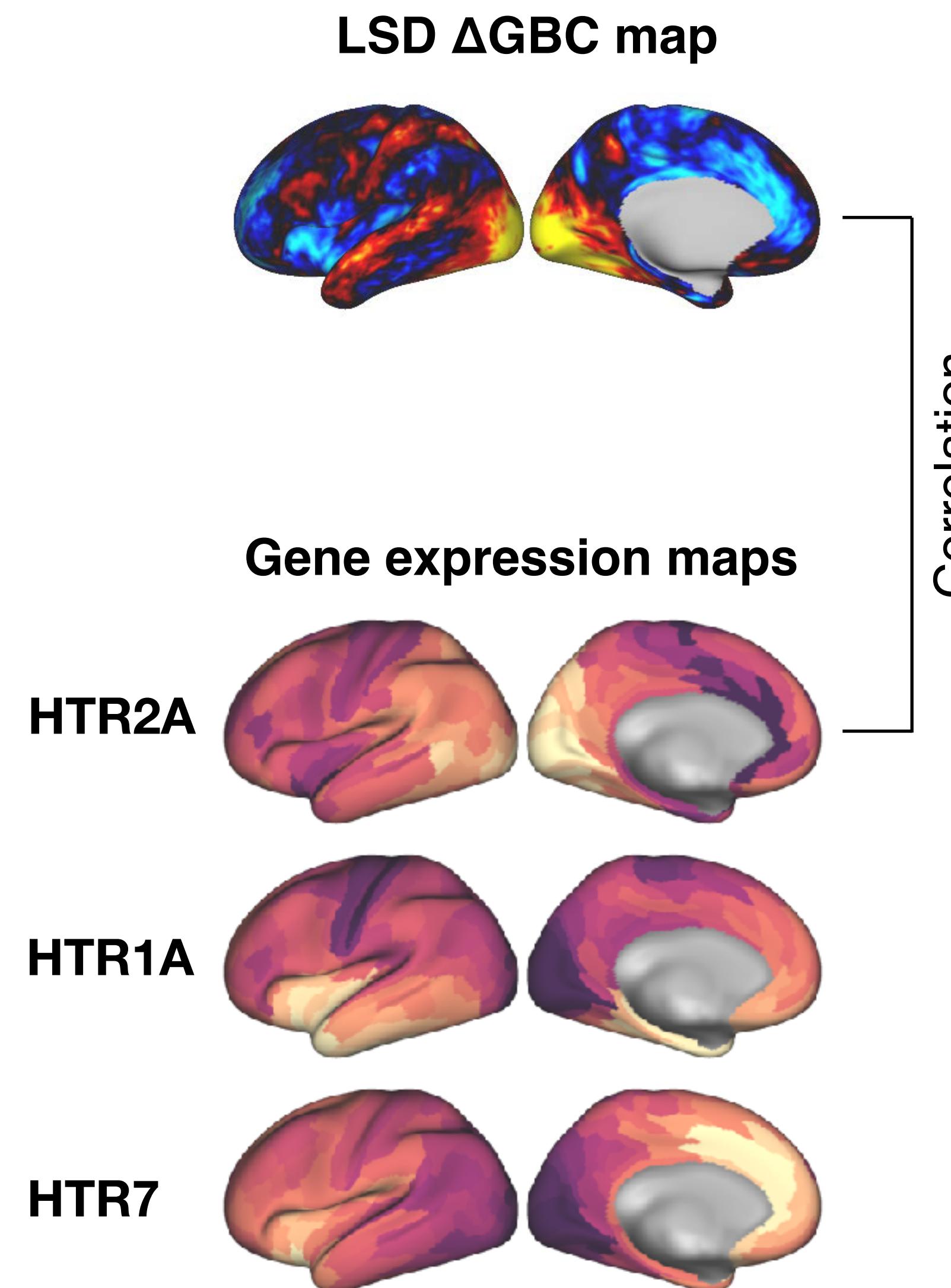


- ▶ Neural and behavioral effects of LSD blocked by ketanserin, a selective antagonist of the serotonergic **5-HT2A** receptor
- ▶ Hypothesis: map aligns with expression of gene **HTR2A**

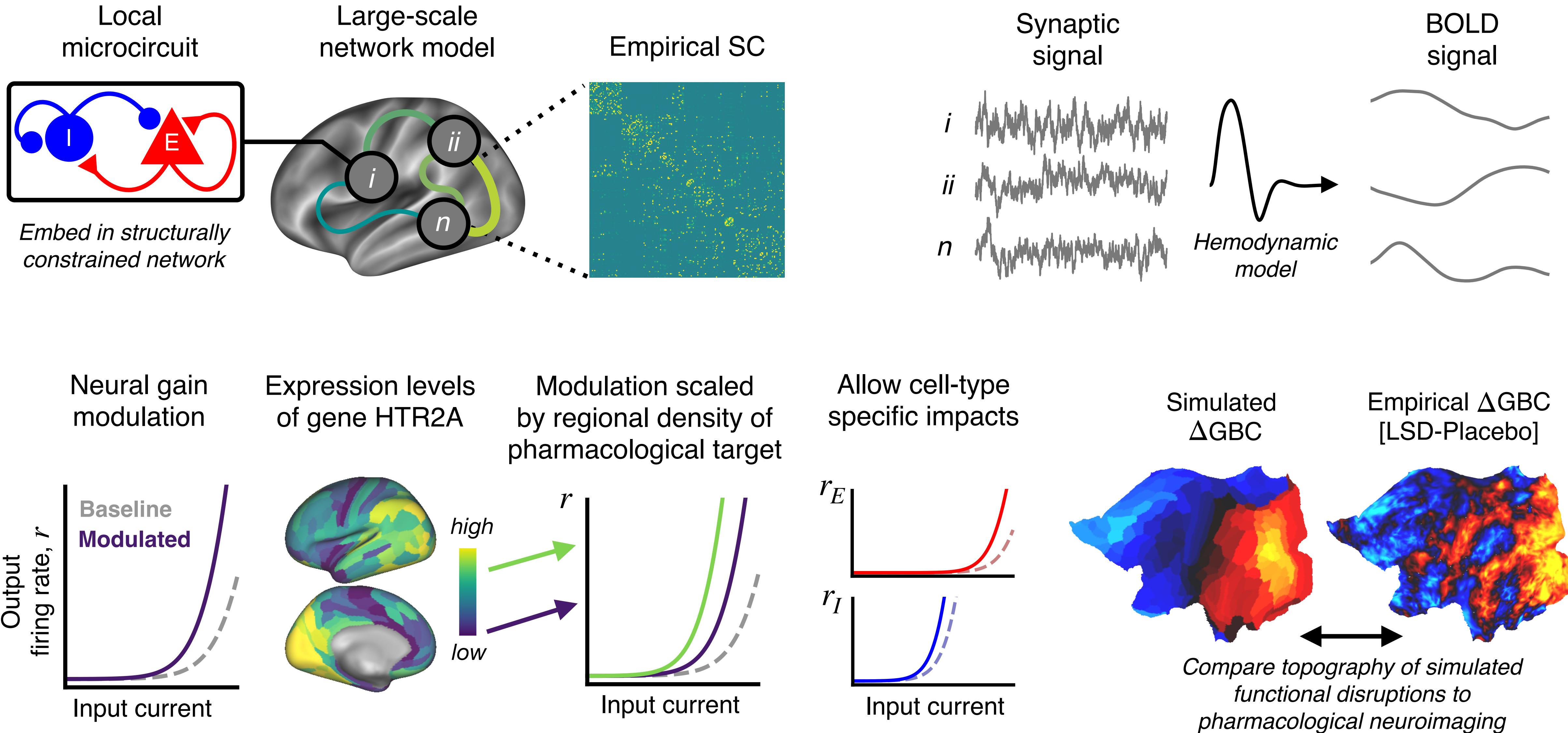
# Identifying molecular candidates of neuroimaging effects



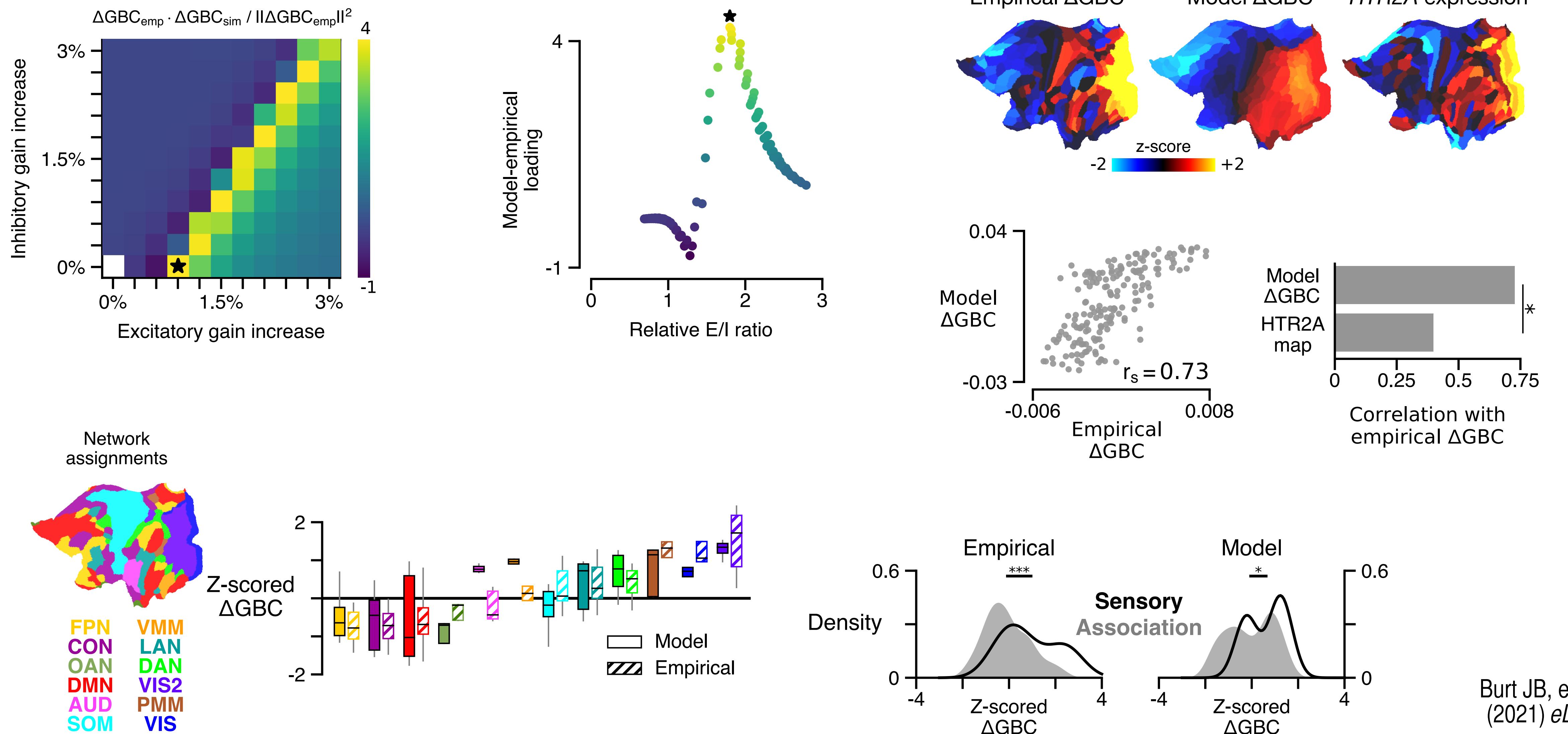
**Josh Burt**



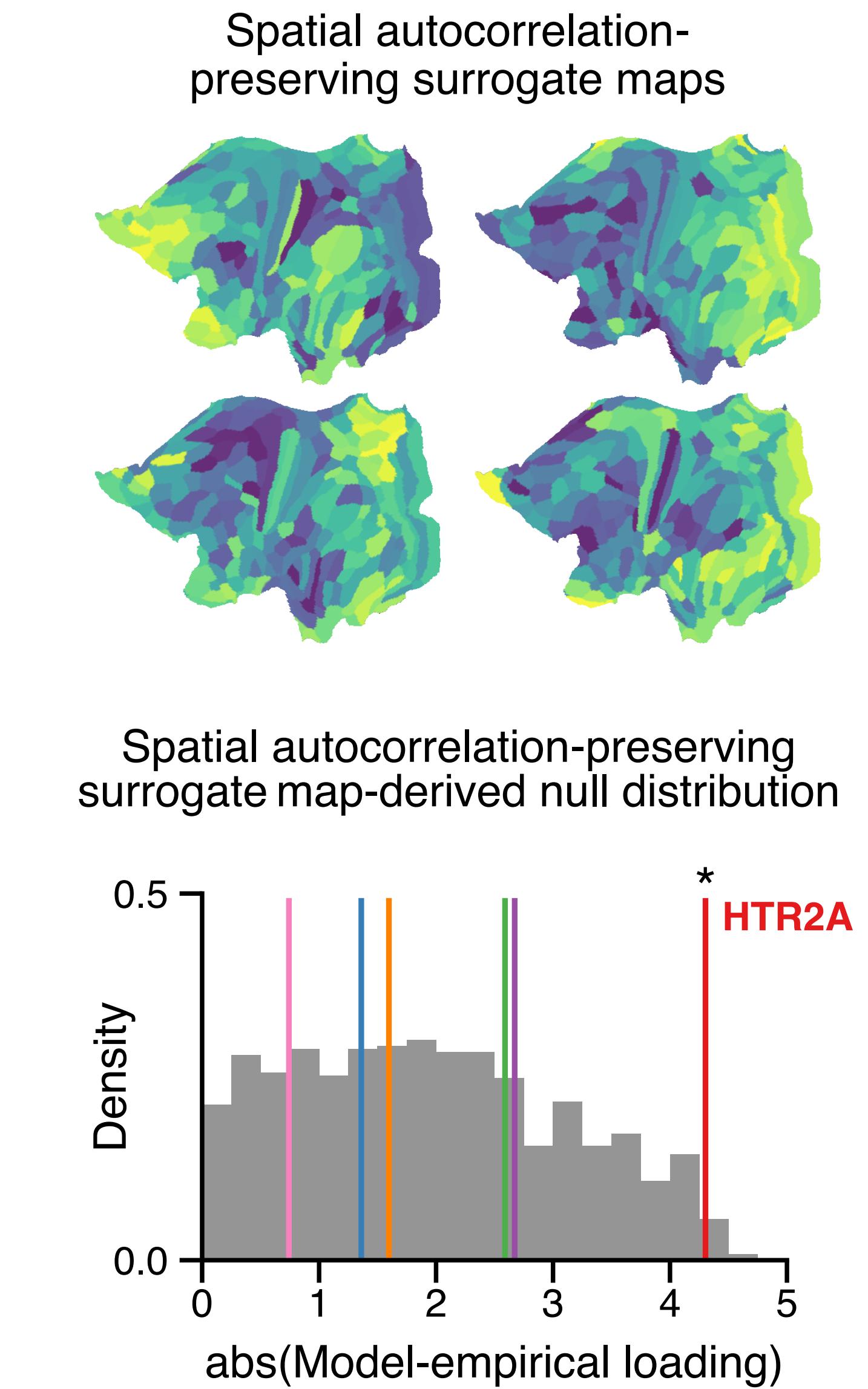
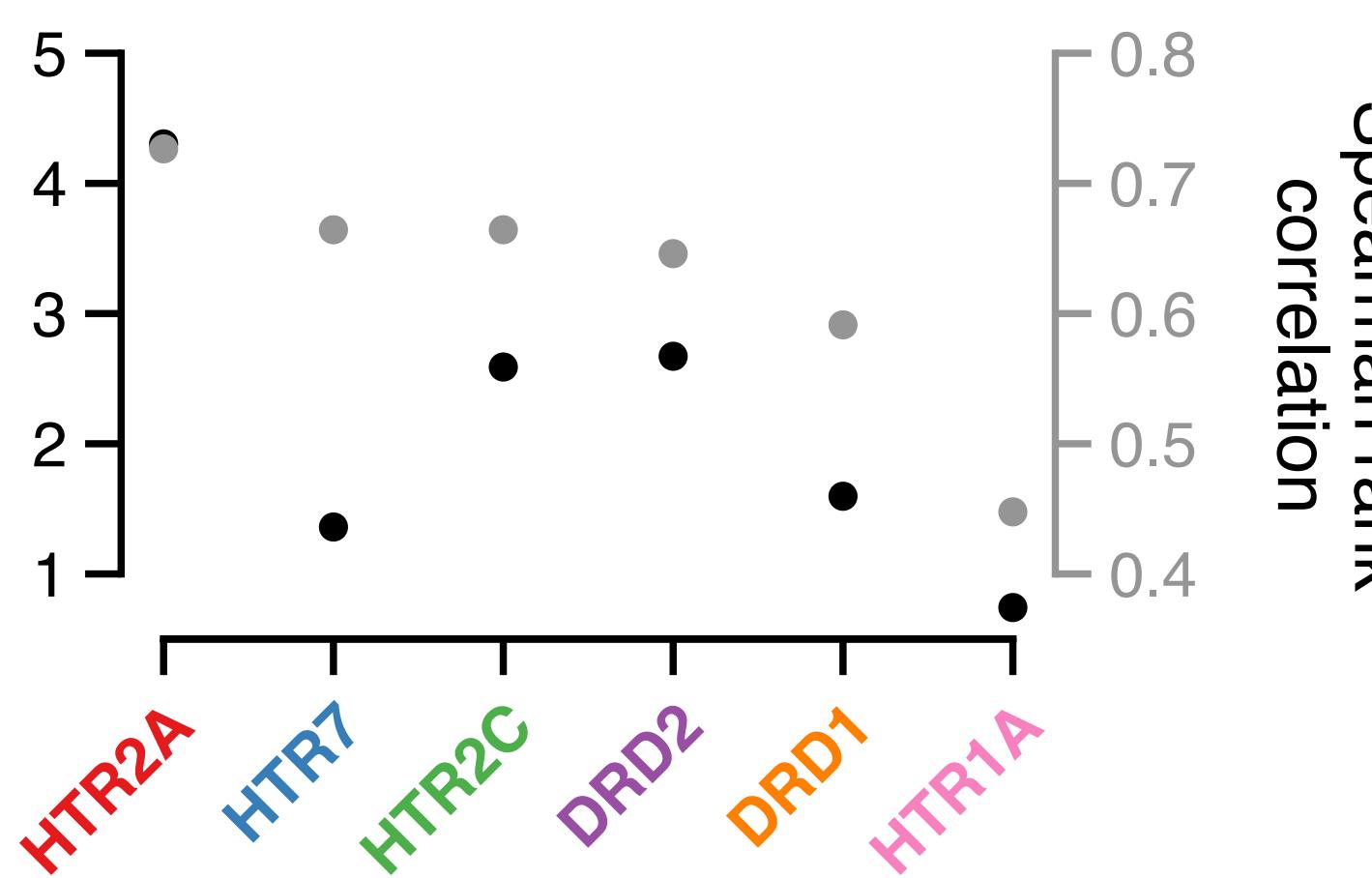
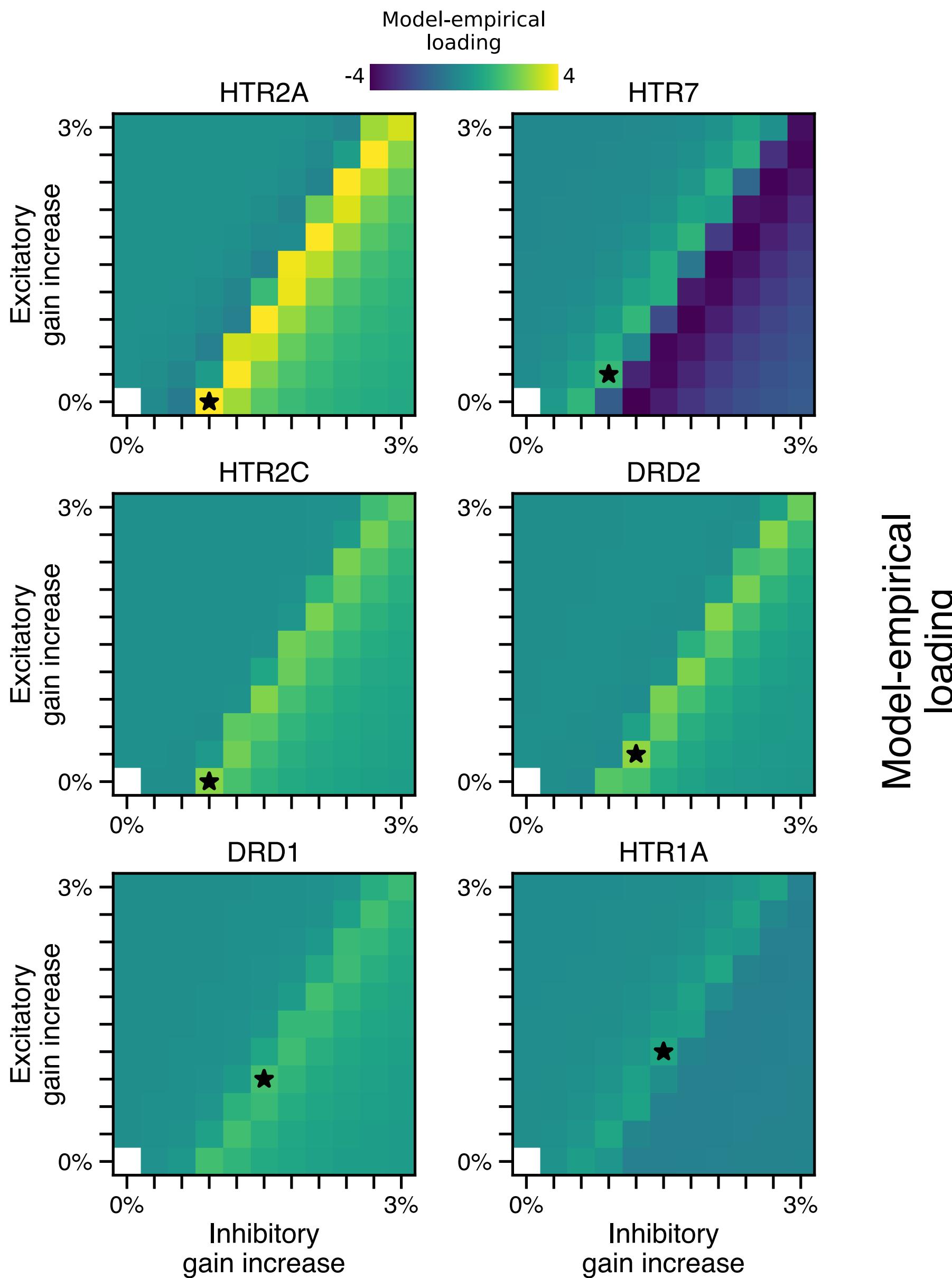
# Modeling pharmacological neuroimaging



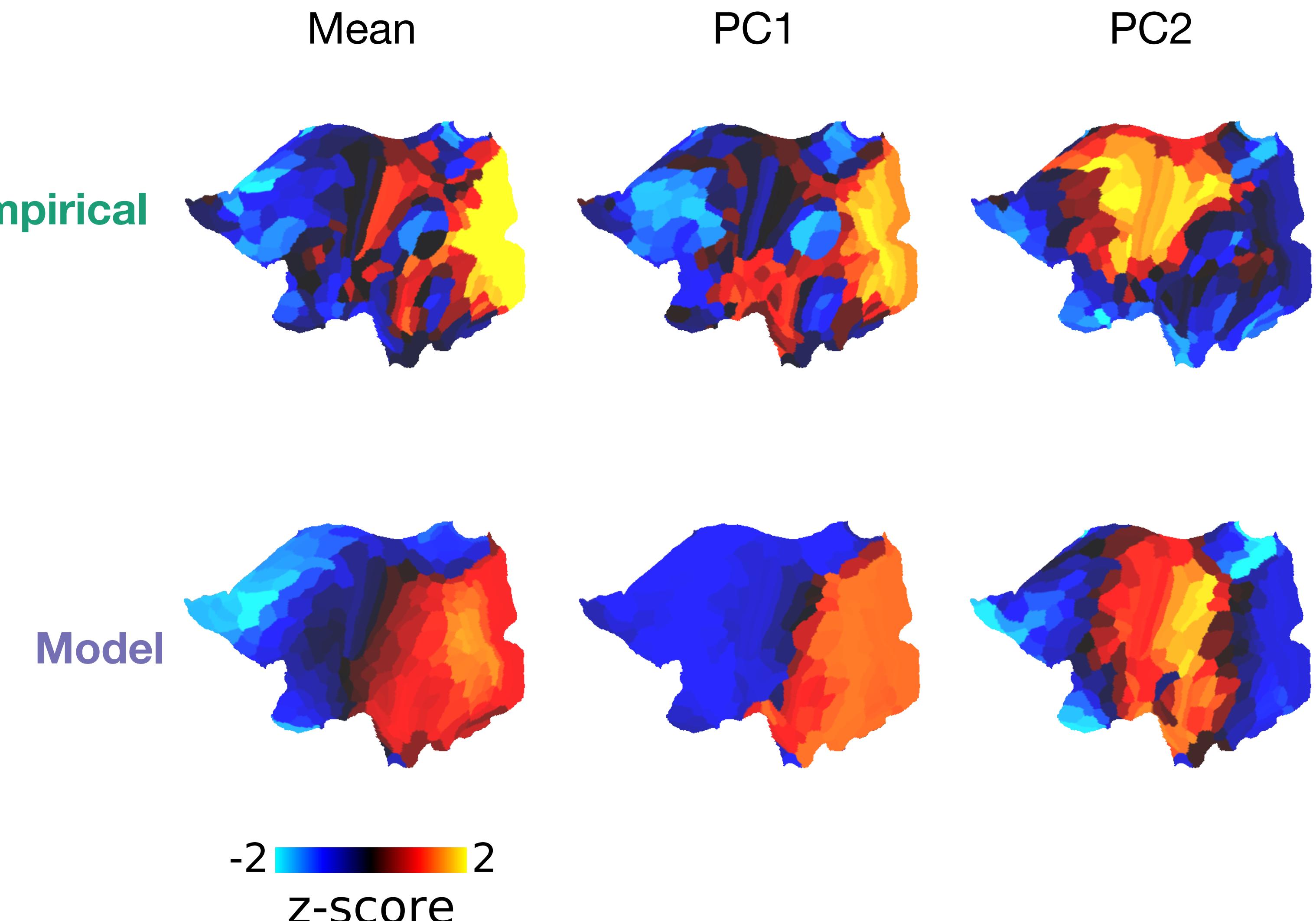
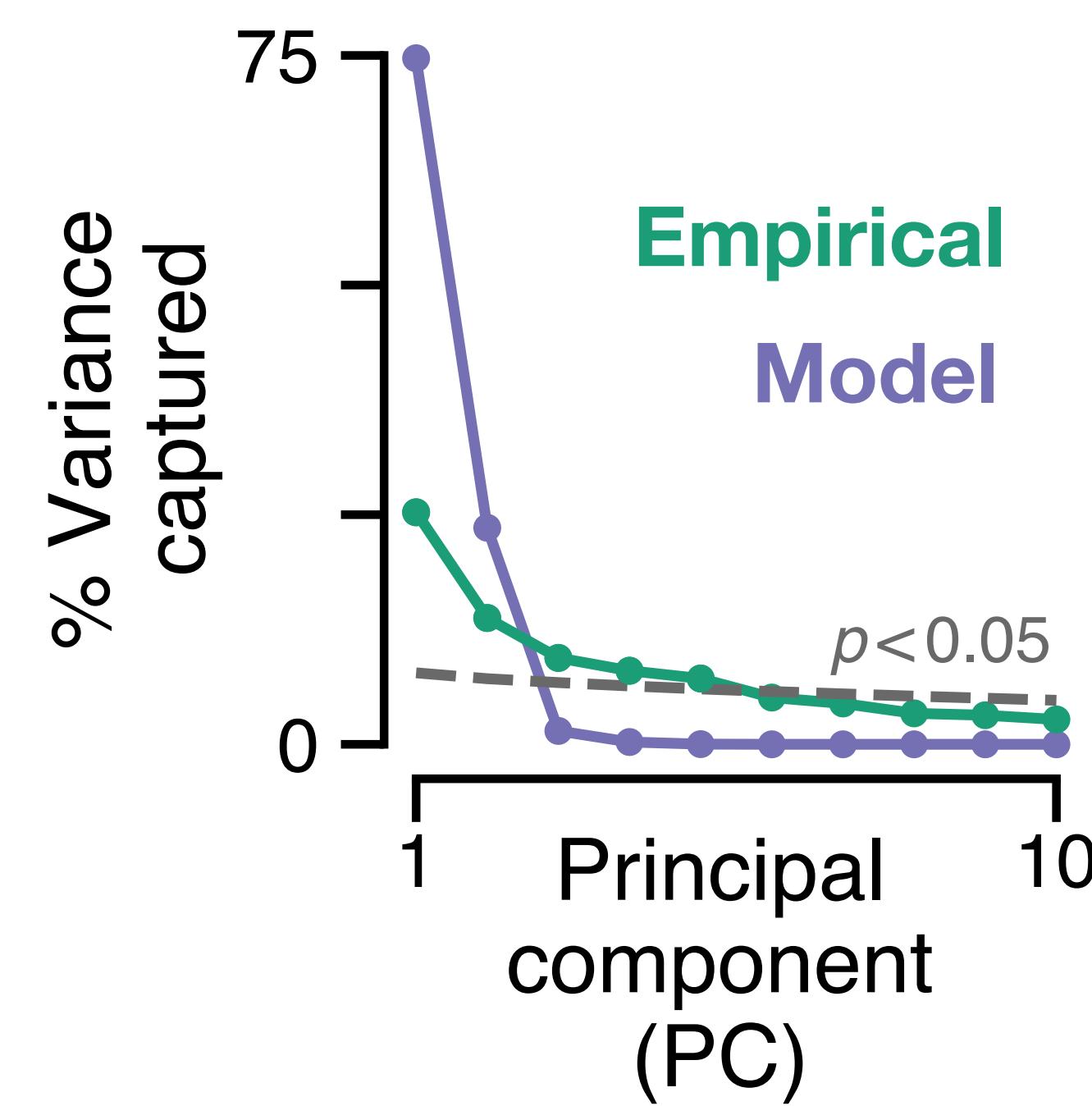
# Model captures topography of LSD-induced connectivity changes via HTR2A-mediated gain modulation



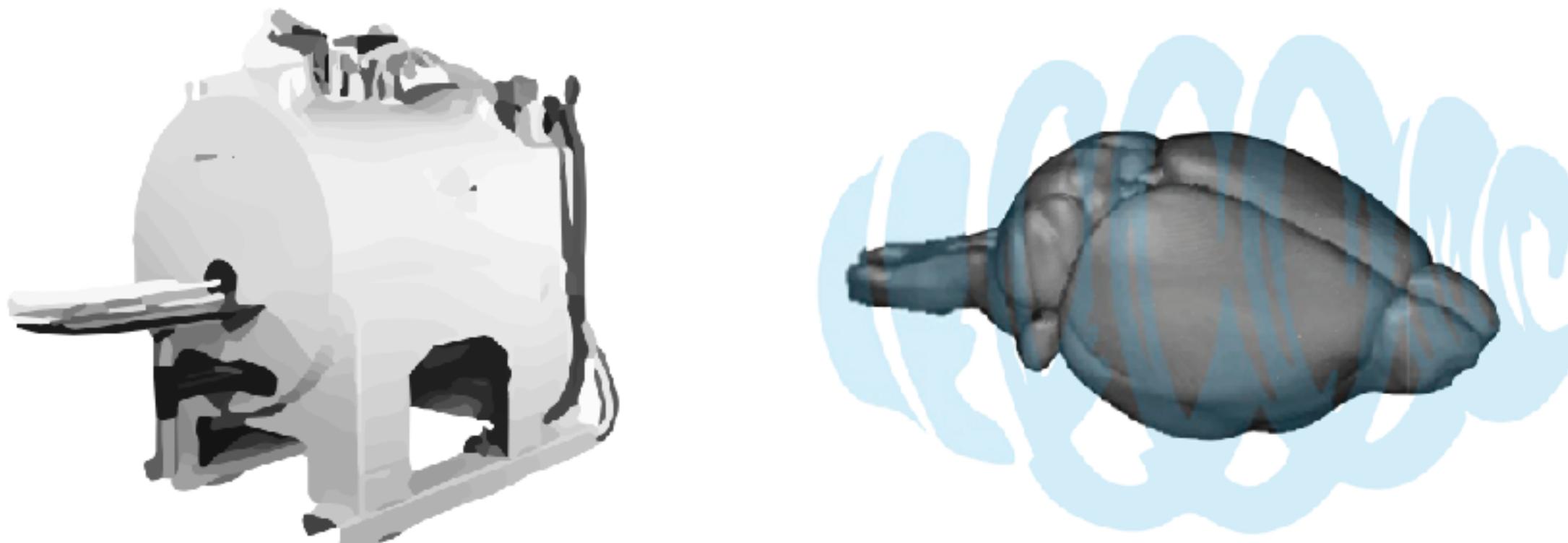
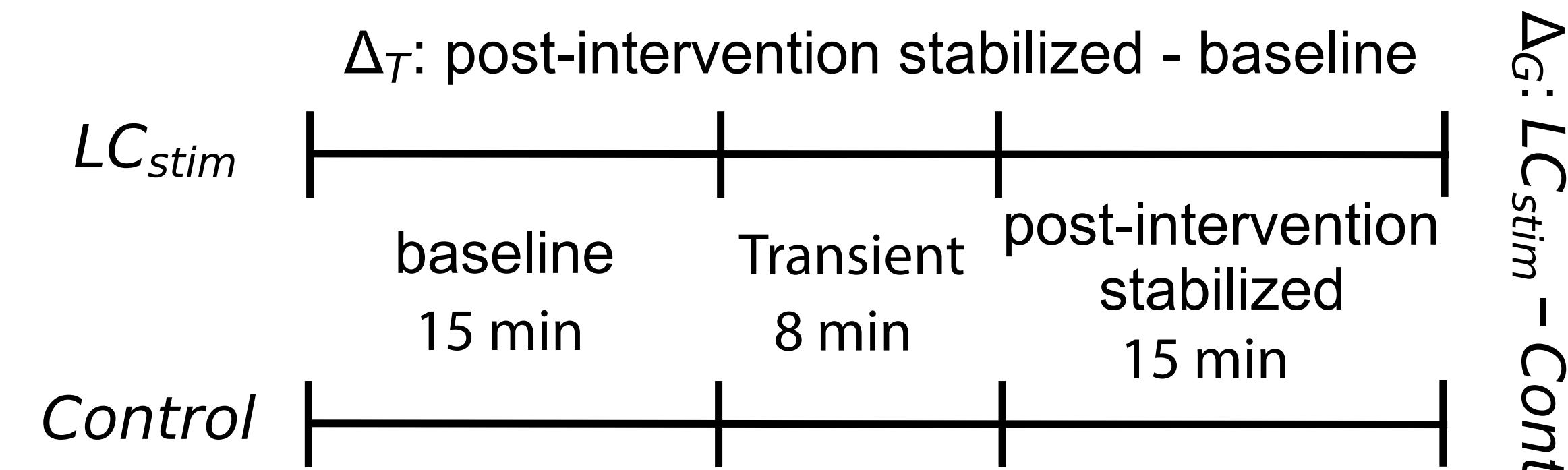
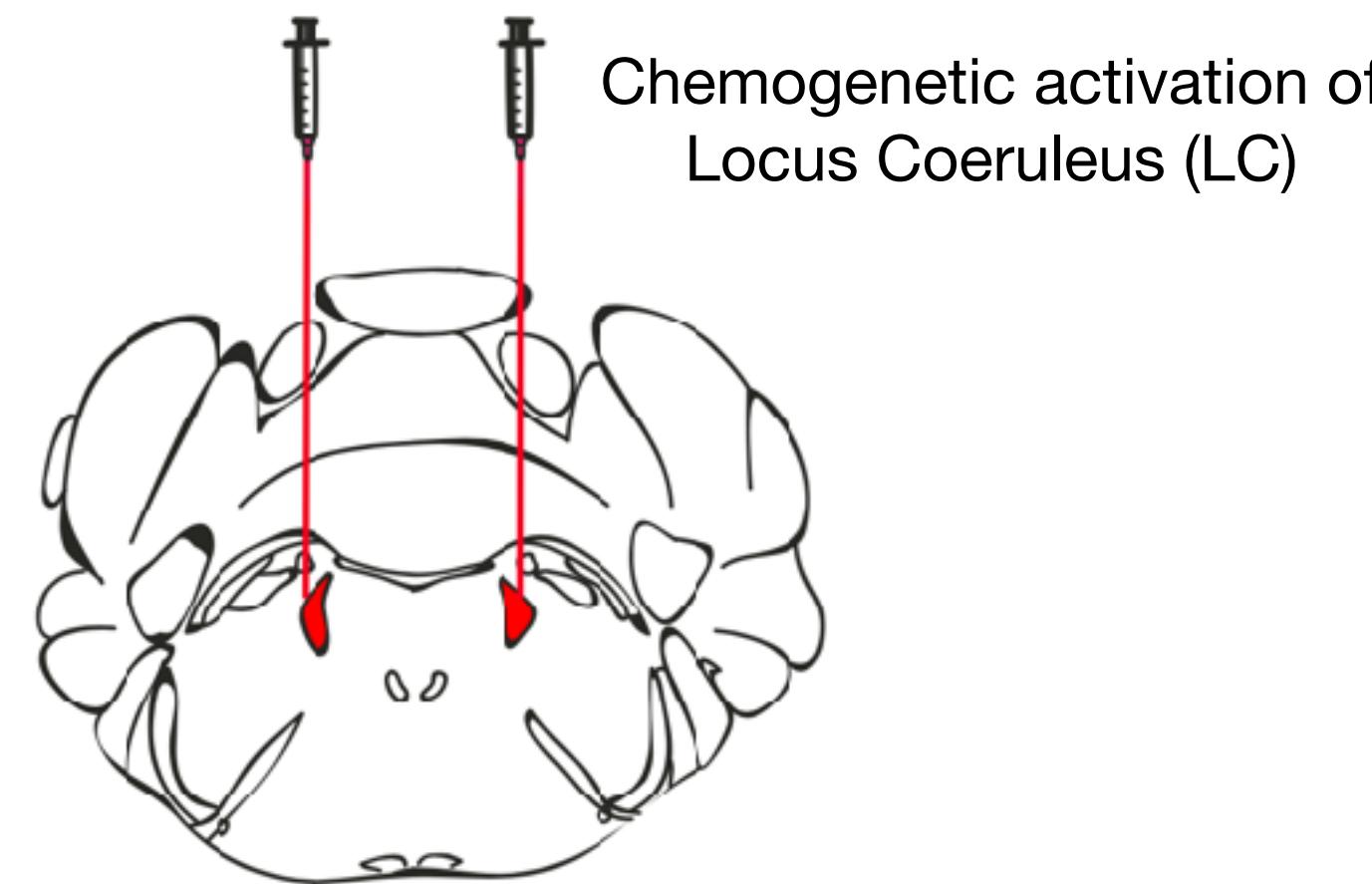
# Comparison of HTR2A to alternative gain modulation maps



# Individual variation in pharmacological fMRI response



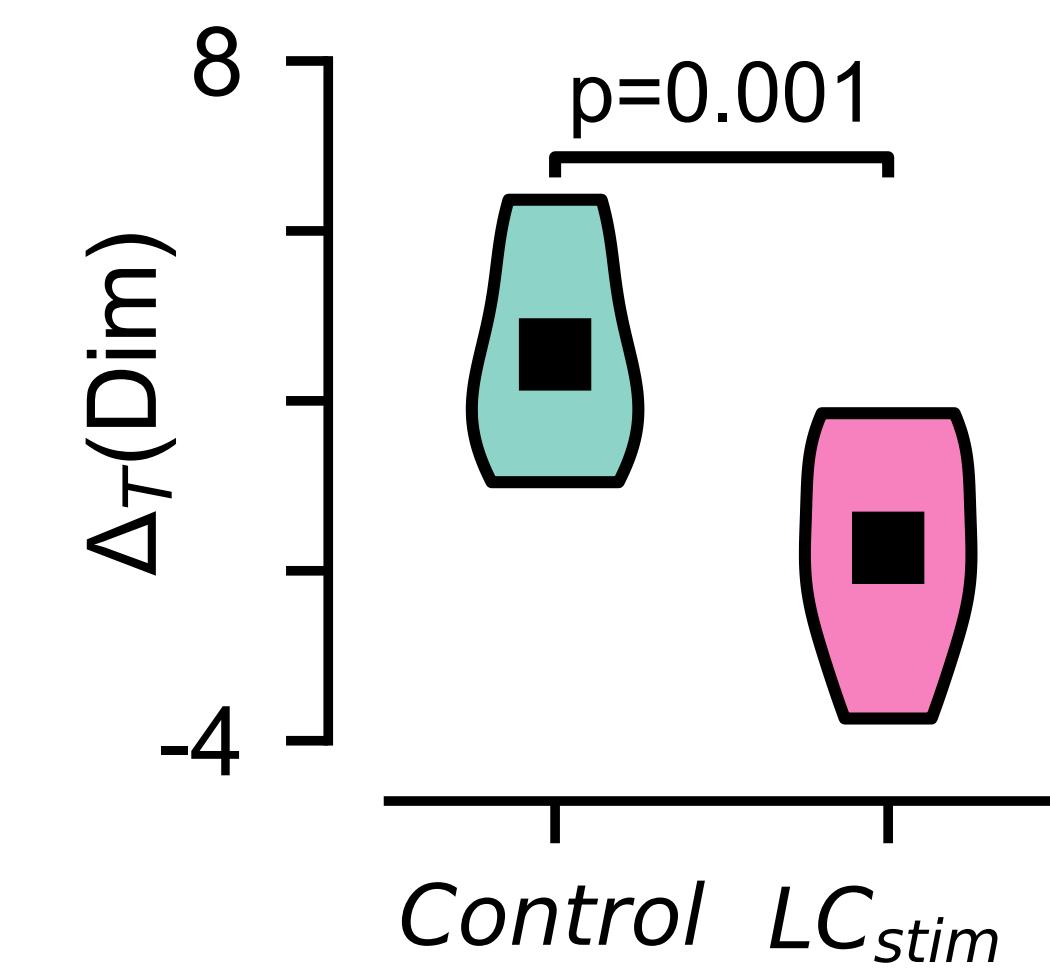
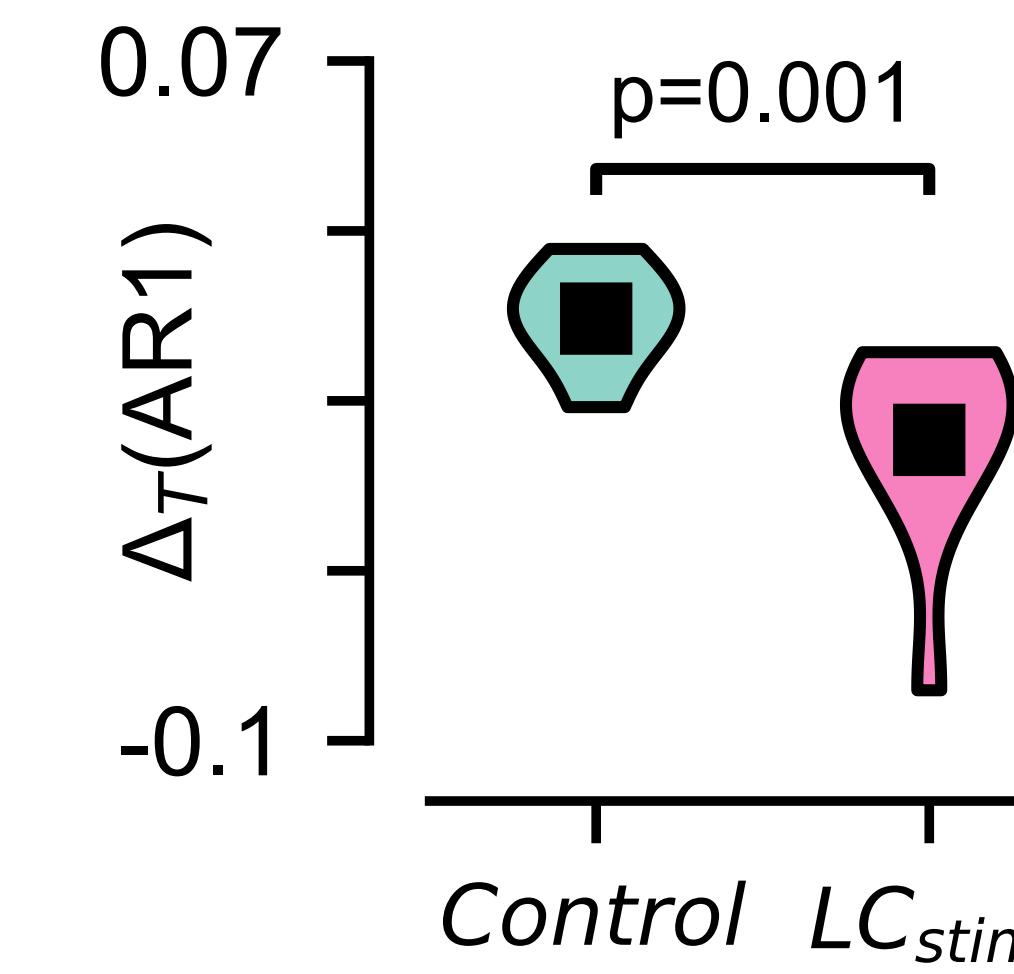
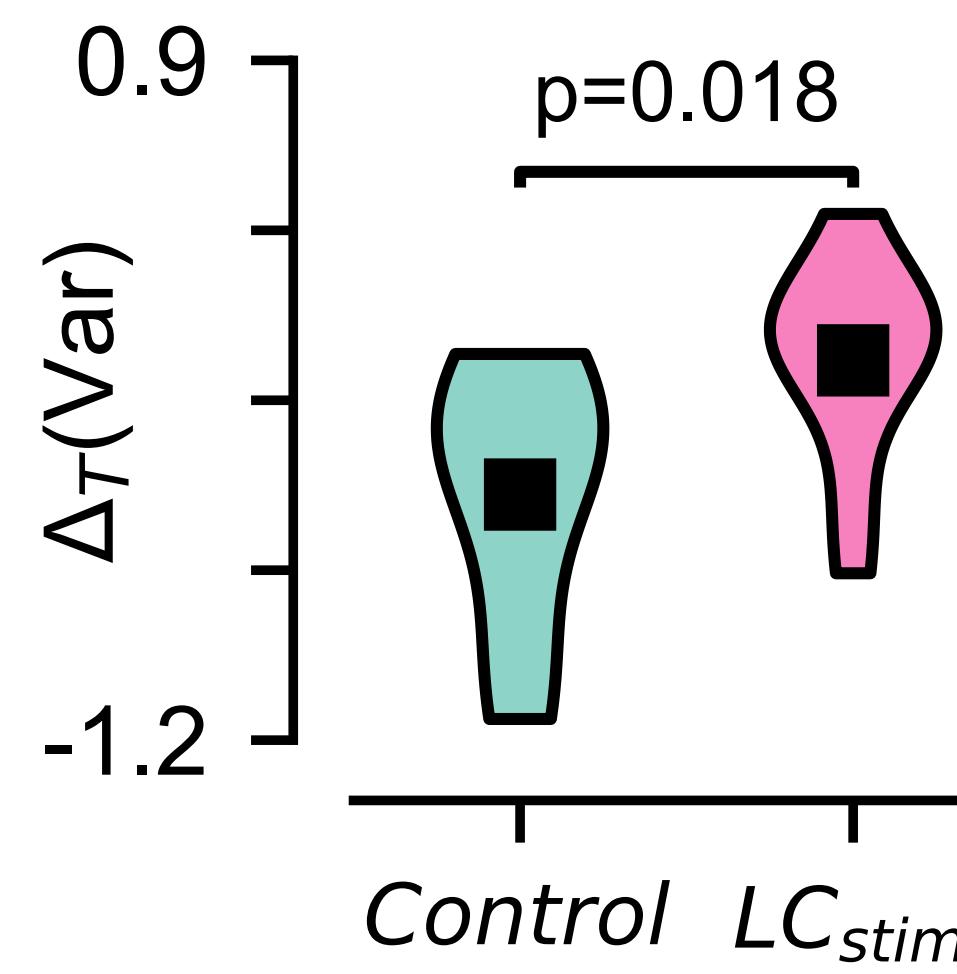
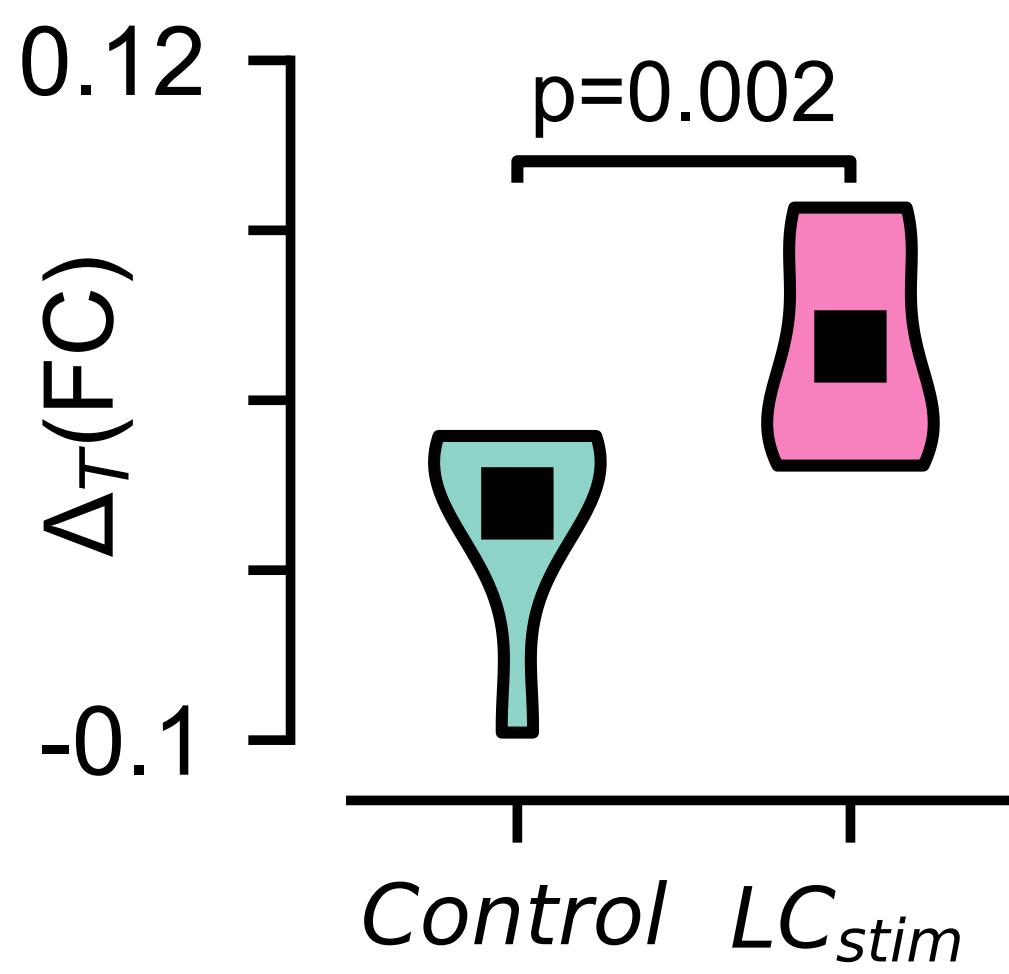
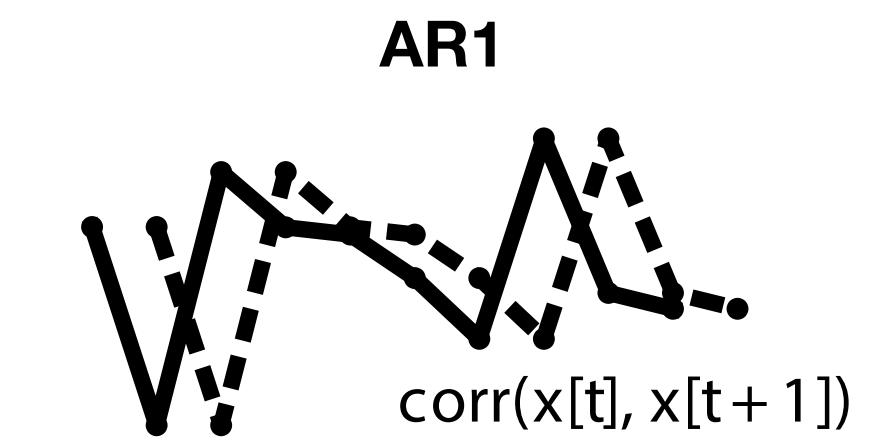
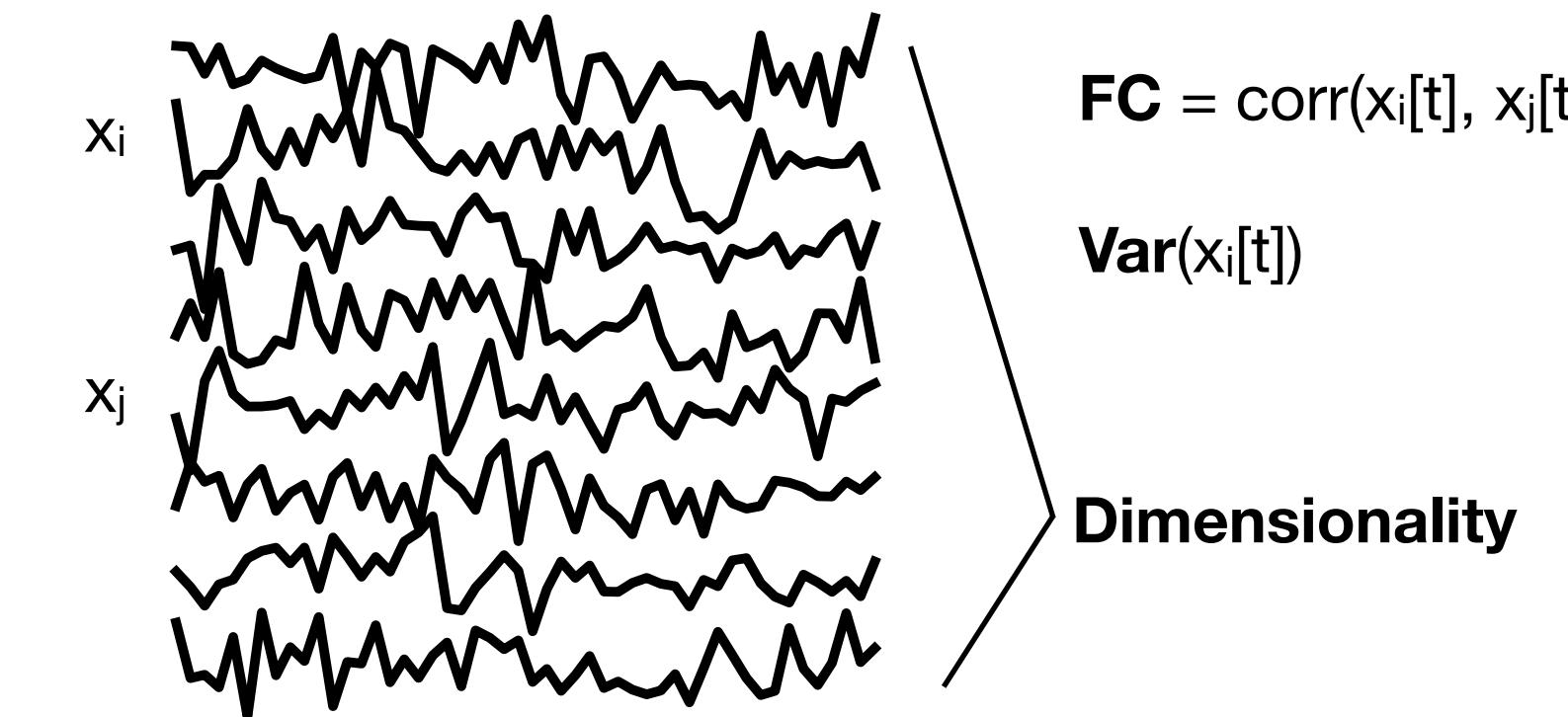
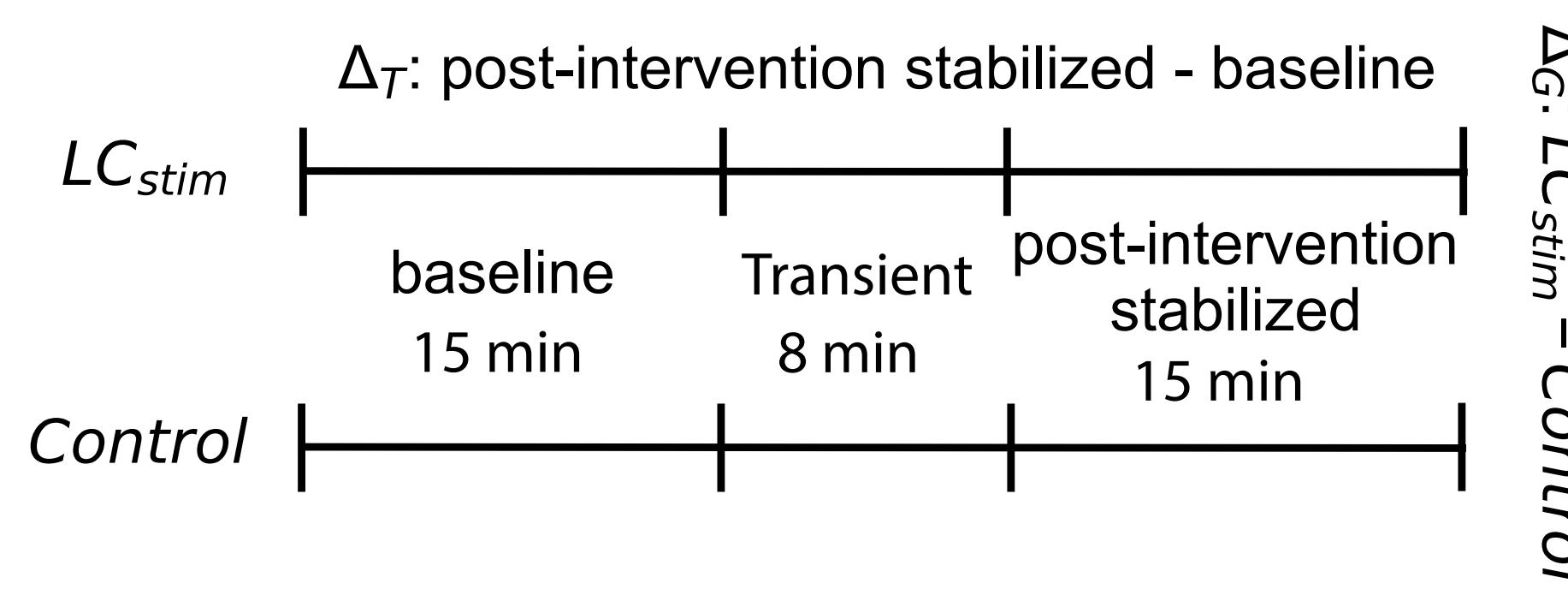
# Chemogenetic activation of locus coeruleus (LC) in mouse to study brainwide neuromodulation by norepinephrine



Collaboration with Valerio Zerbi (ETH)

Zerbi et al. (2019) *Neuron*

# LC stimulation effects on BOLD dynamics



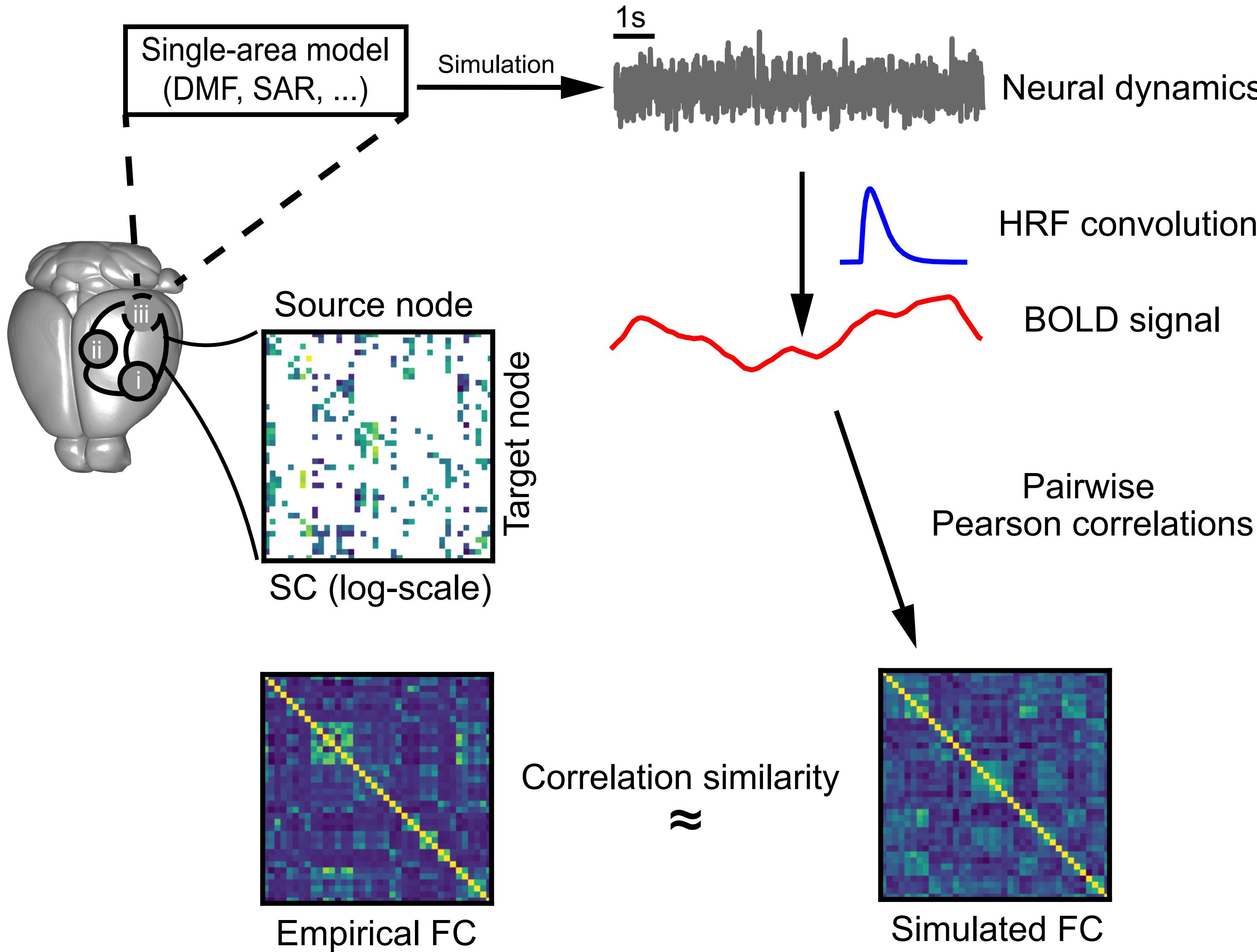
# Large-scale model of mouse cortical BOLD dynamics



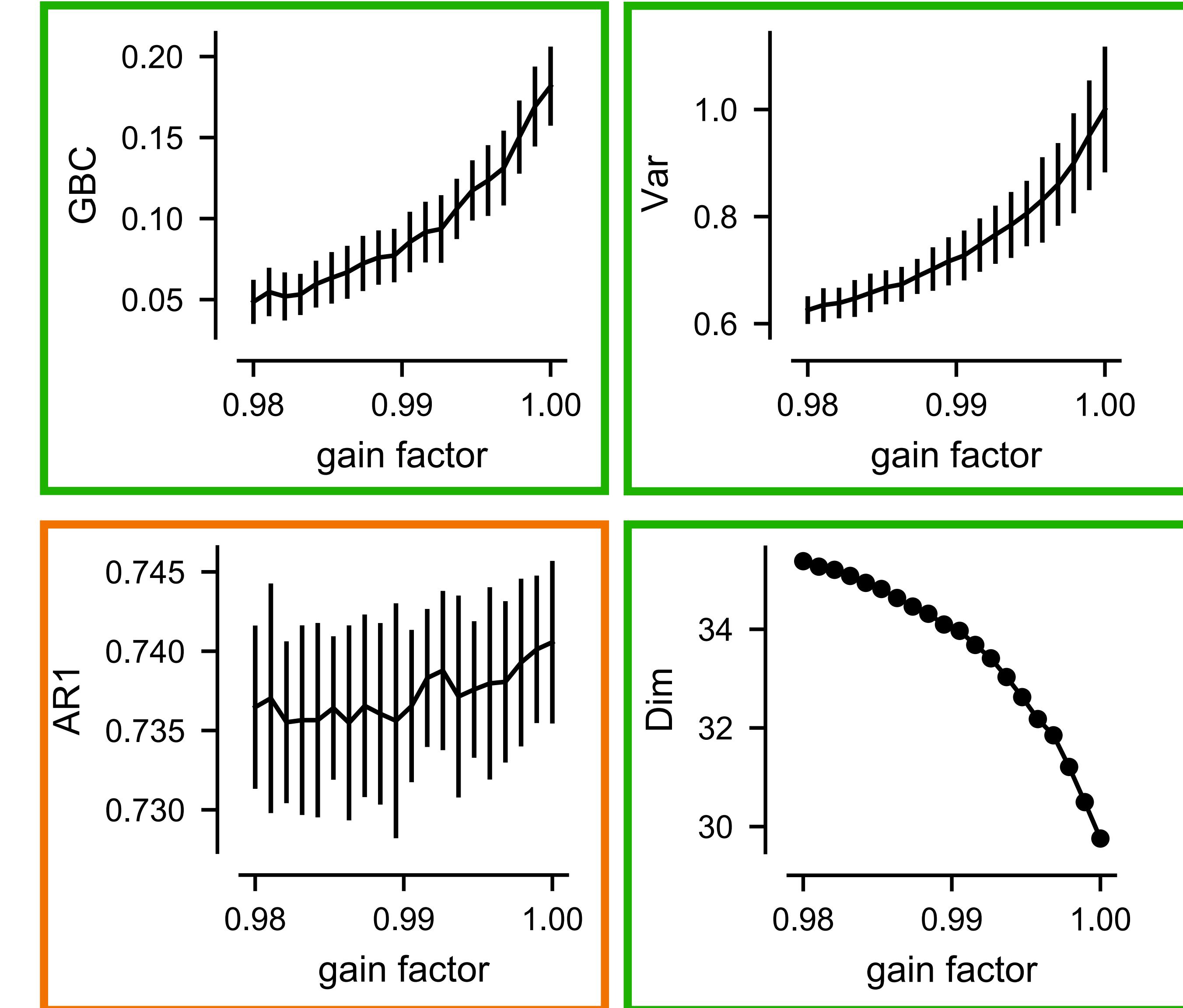
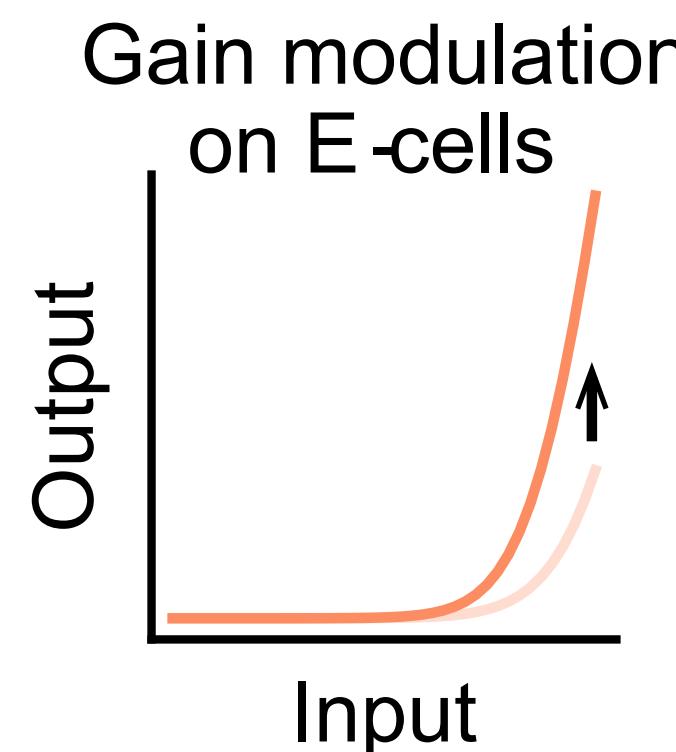
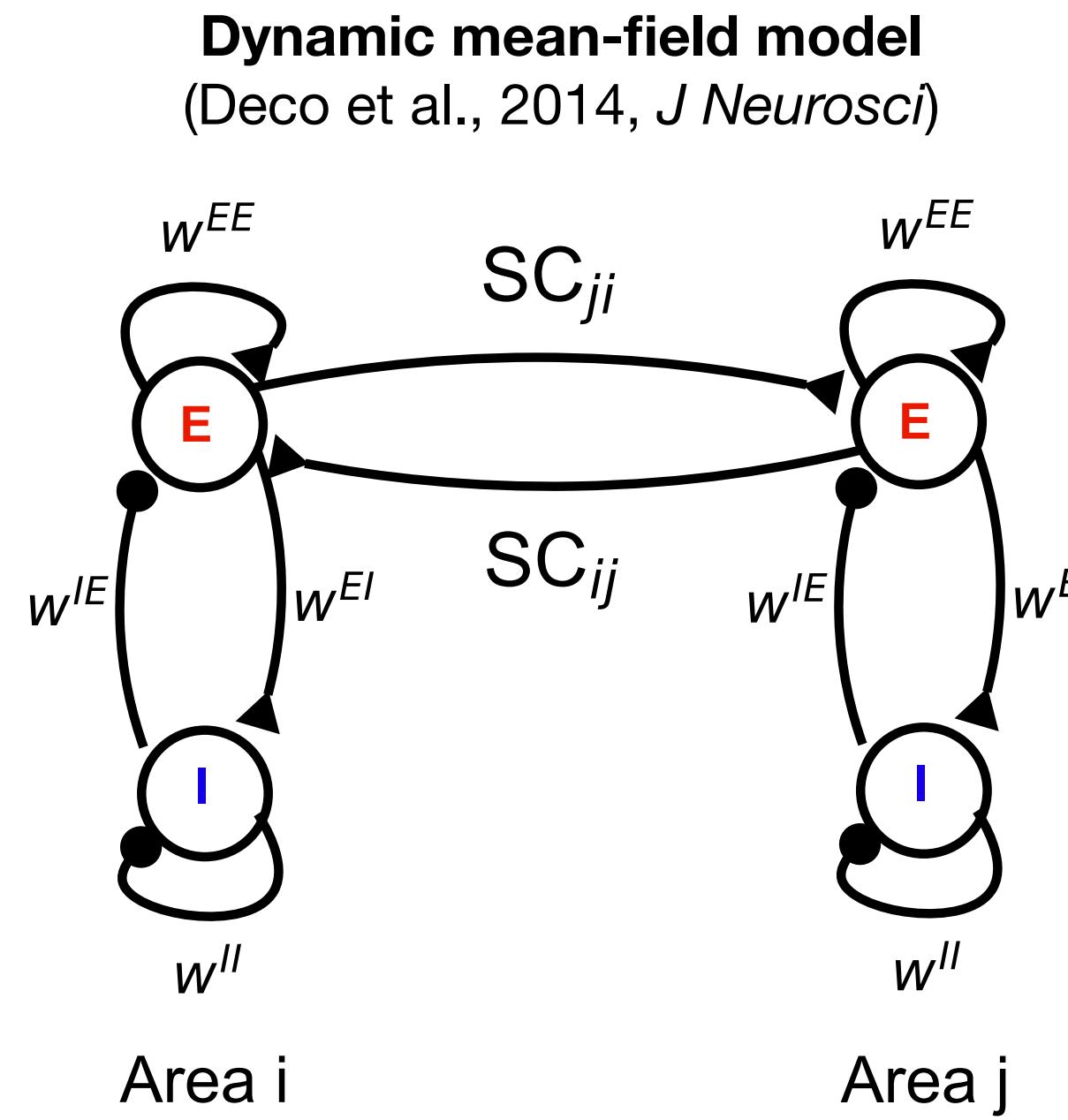
Daming Li



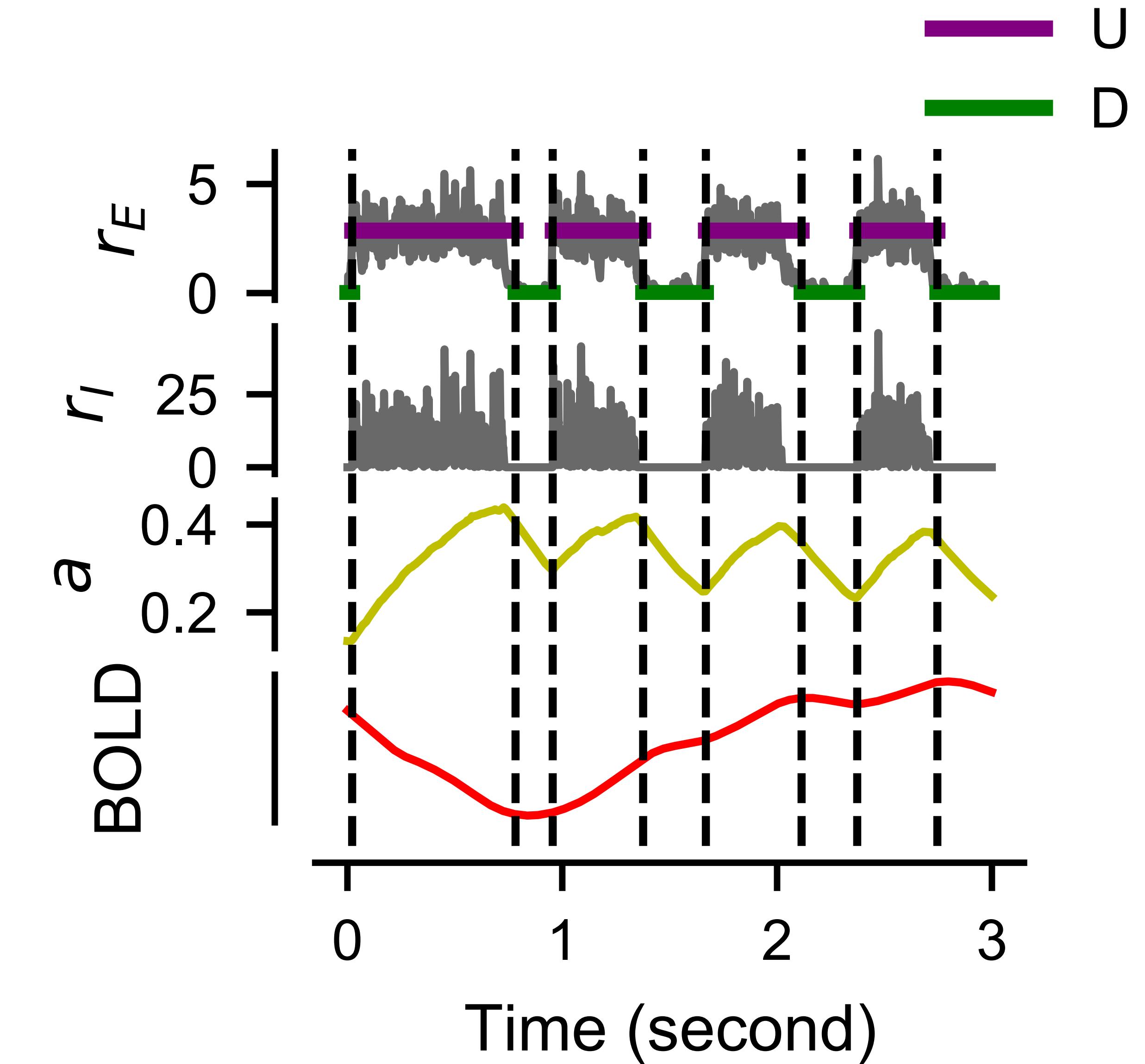
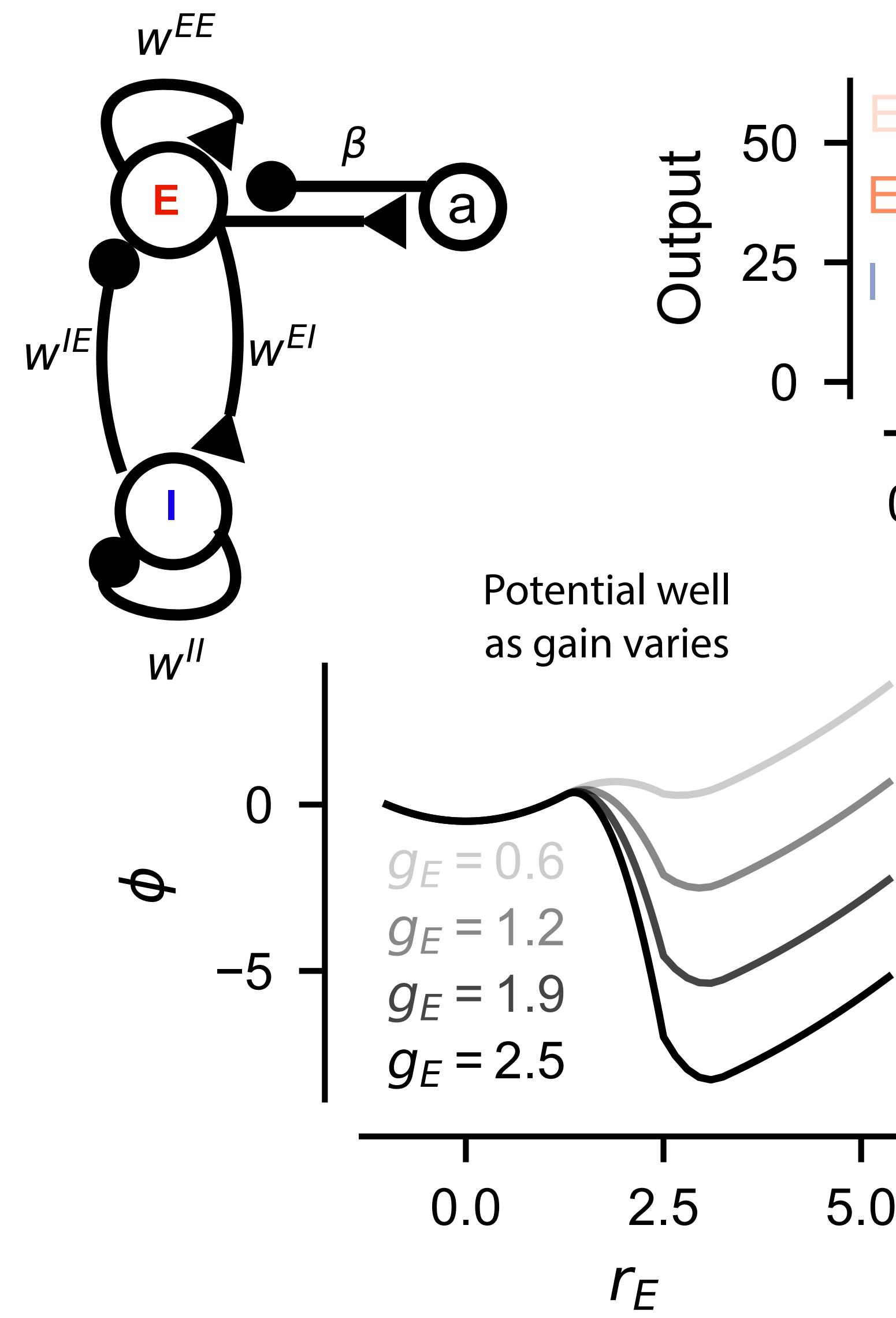
Qinglong Gu



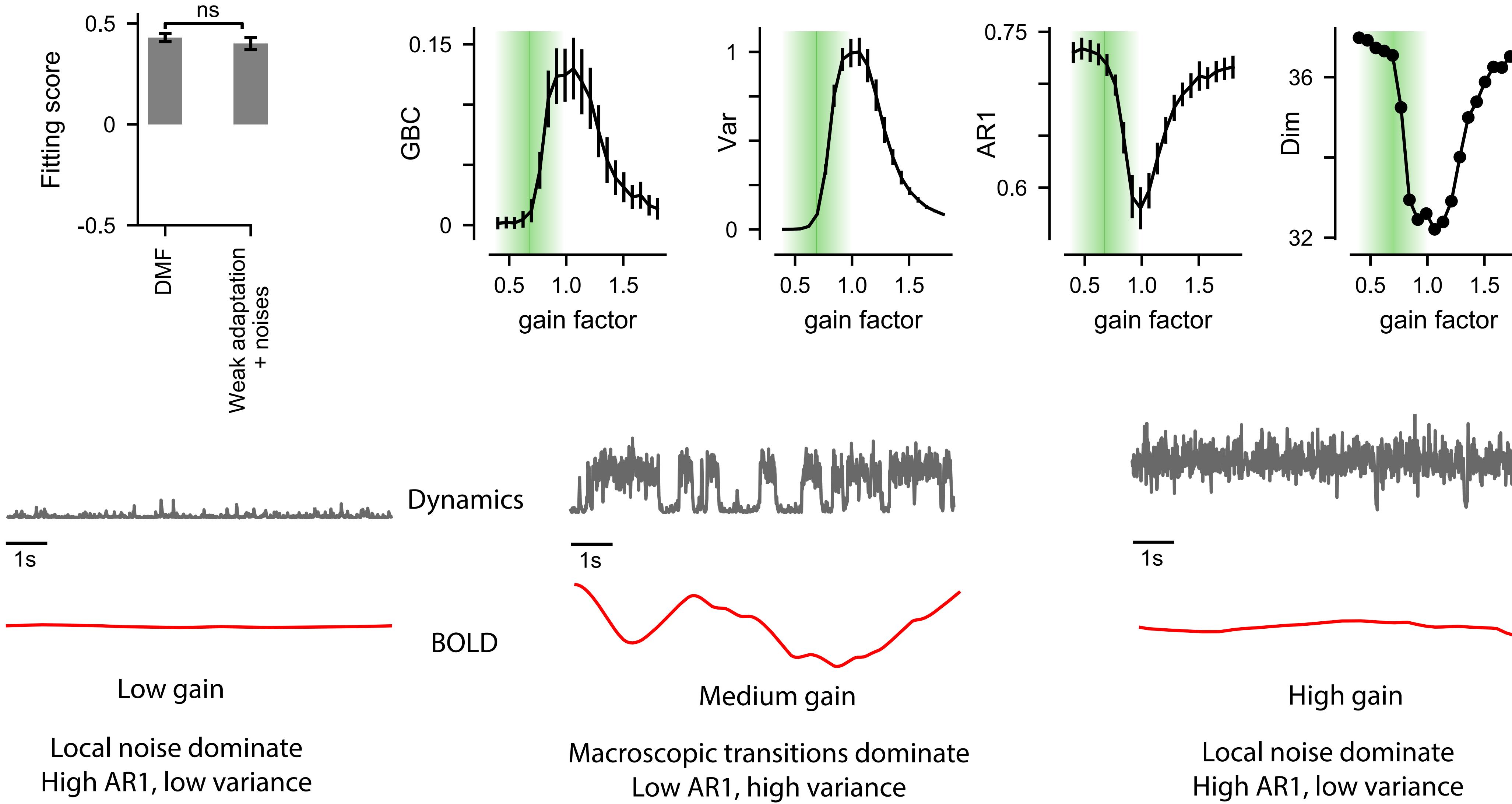
# Dynamic mean-field model of mouse cortex with gain modulation



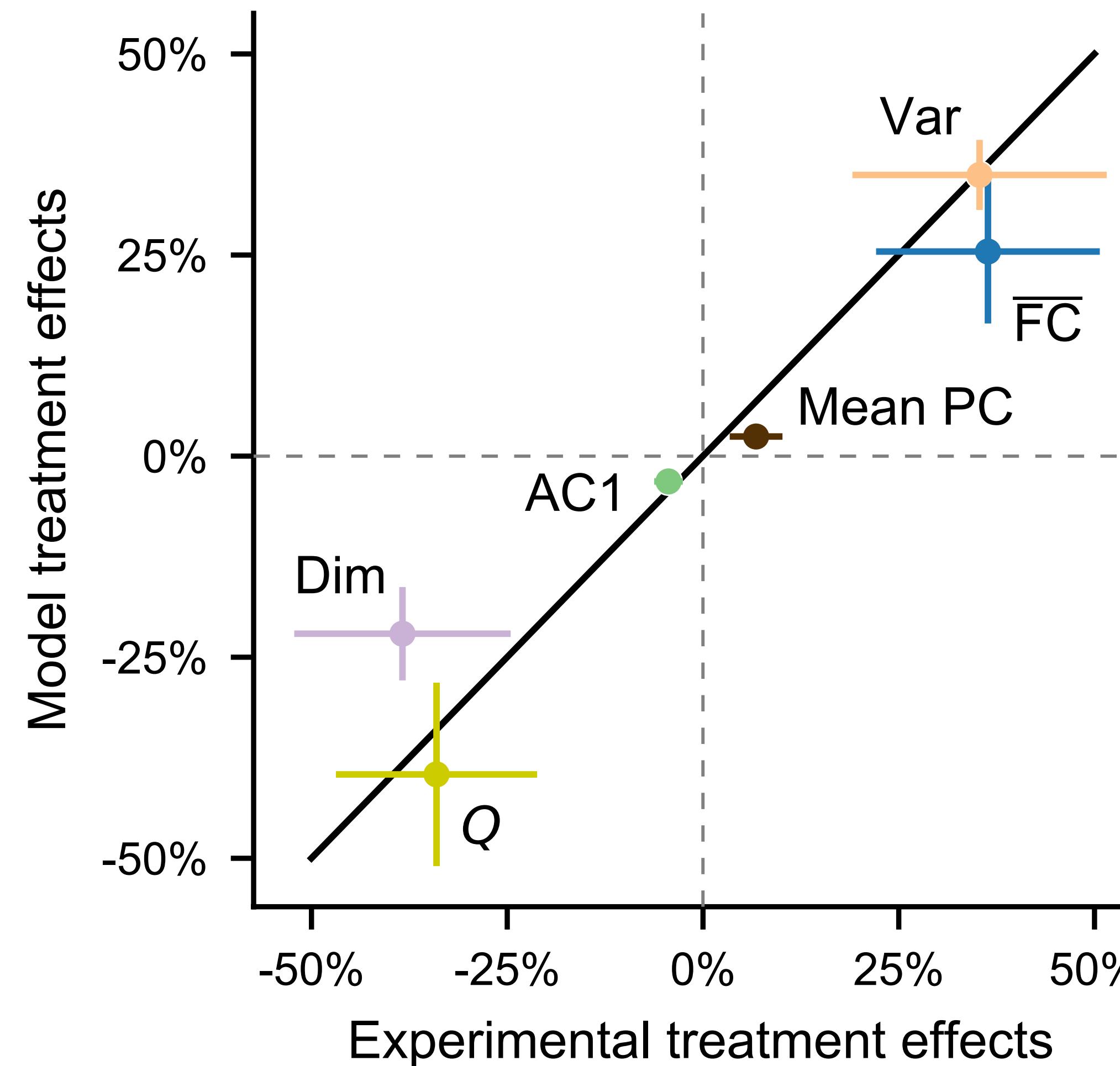
# Local E-I circuit with adaptation with bistable dynamics



# Gain modulation in bistable model captures LC stimulation effects



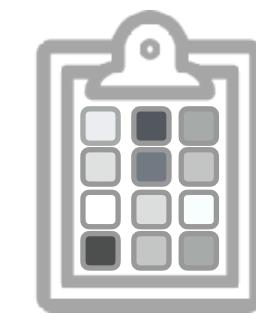
# Gain modulation in bistable model captures LC stimulation effects



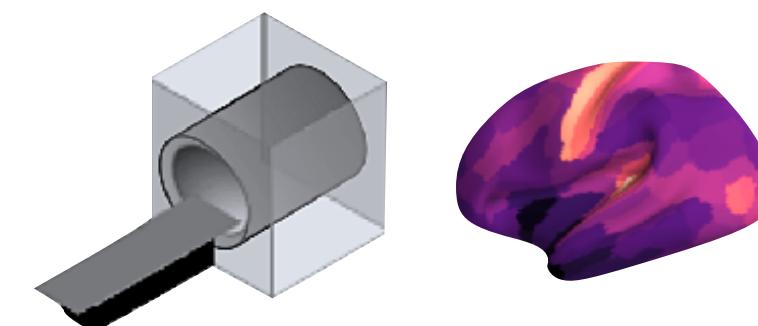
- When the local node is in a suitable regime of stochastic bistable dynamics, then multiple neural features of LC stimulation are produced under gain modulation of excitatory neurons

# Conclusions

PERSONALIZED  
BEHAVIORAL  
PROFILES



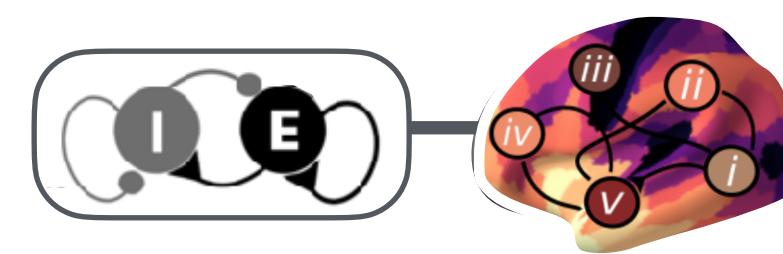
NEURAL  
BIOMARKERS



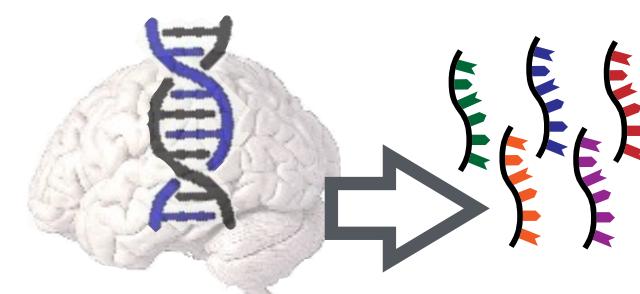
PHARMACOLOGICAL  
NEUROIMAGING



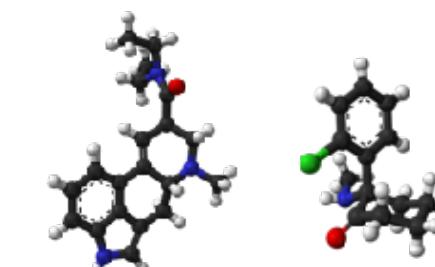
BIOPHYSICAL  
CIRCUIT  
MODELING



TRANSCRIPTOMIC  
MAPPING



PHARMACOLOGICAL  
THERAPEUTICS



- ▶ **Bridging across levels of analysis, from molecular to systems**
- ▶ Platforms to iteratively expand and constrain computational models with biophysical detail needed for capture important empirical phenomena
- ▶ Mapping transcriptomic expression
- ▶ Modeling biophysical circuit dynamics at individual level
- ▶ Modeling brainwide response to pharmacology and neuromodulation

# Acknowledgements



Murat Demirtaş



Joshua Burt



Rachel Cooper



Daming Li



Qinglong Gu

## Main collaborators on this project

Alan Anticevic  
John Krystal  
Katrín Preller

Stam Sotiropoulos  
Franz Vollenweider  
Valerio Zerbi



R01 MH112746

