

Effective workflow from multi-modal MRI data to model-based prediction

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Introduction

- Comparing structural connectivity (SC) and functional connectivity (FC) led to the **structure-function relationship** as a possible methodological approach to explore the interdependence between structure and function of the human brain. However, this relationship between **empirical SC (eSC)** and **empirical FC (eFC)** is **relatively low**, might depend on many factors, and its mechanism is still unclear [1].
- Integration of model-based approaches into whole-brain connectome research** can expand the scope of investigation to understand the brain [2,3]. The models can be used to generate **simulated FC (sFC)** as **an additional data modality**. Accordingly, it can be suggested as a possible mediator between brain structure and function.
- We suggest a framework that advances the applicability of the model-based approach by applying simulated data to machine-learning analysis.

Methods: A workflow for model-based machine-learning research using multi-modal MRI data

Step 1: Multi-modal MRI data

HCP young adults (n=270, 142 females, 28.5±3.5 years old)

- Structural and functional MRI data: T1-weighted MRI, Diffusion-weighted MRI, Resting-state functional MRI

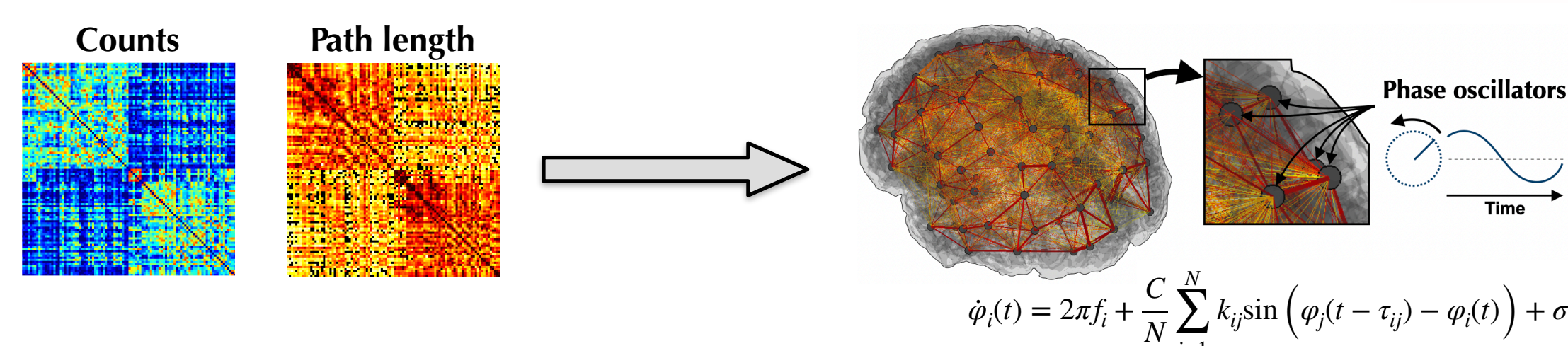
Step 2: MRI processing

- Correction: bias field, head motion, eddy
- Denoising, tissue segmentation
- Image registration to MNI space

Step 3: Whole-brain connectome

- Functional atlas: **Schaefer 100 cortical regions**
- Structural atlas: **Harvard-Oxford 96 cortical regions**

✓ Tracking streamlines → whole-brain tractography



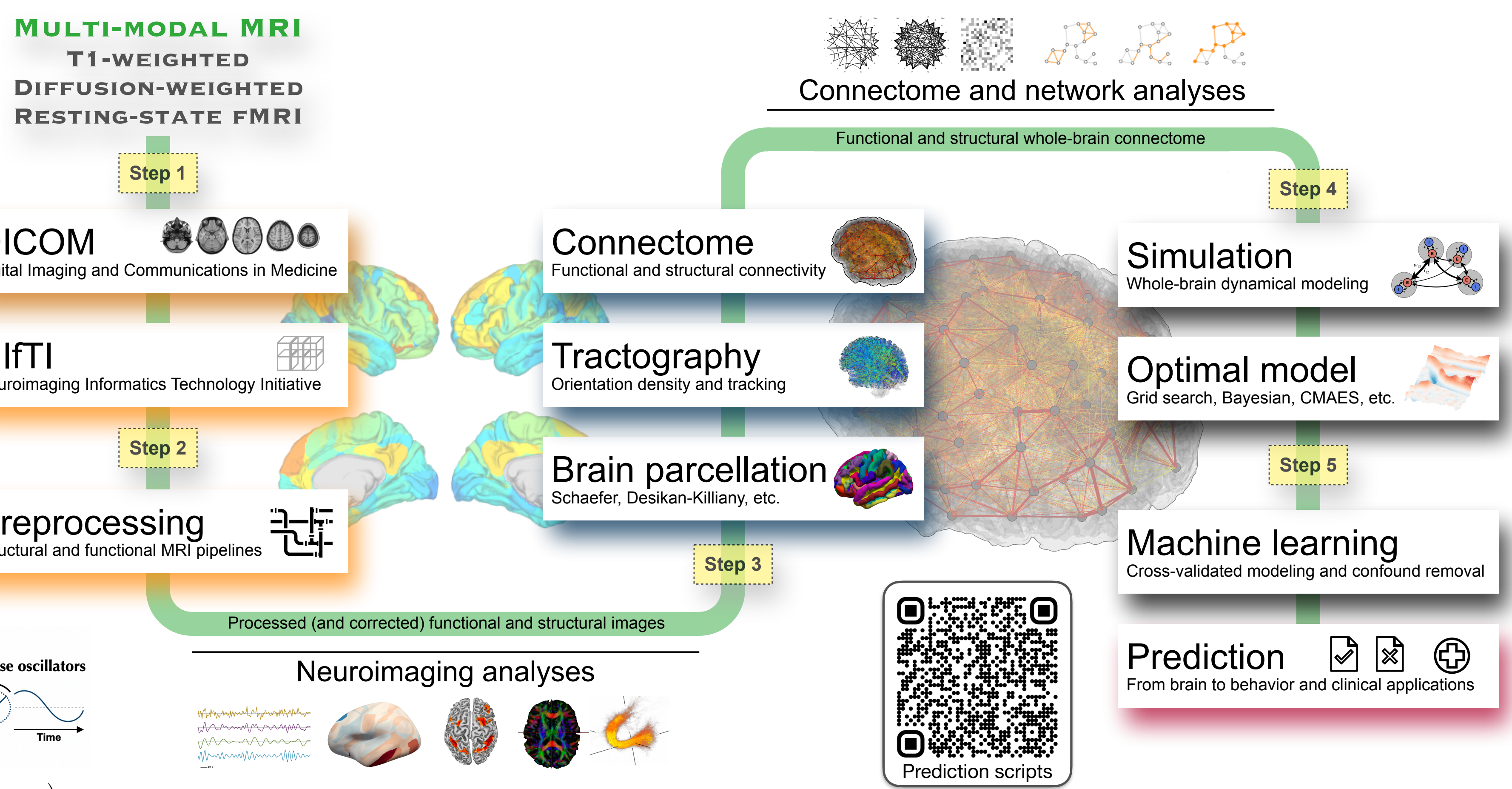
Step 4: Whole-brain dynamical modeling

- A whole-brain model** (coupled phase oscillators with delay)
- Simulated Blood-Oxygen-Level Dependent signals ([0.01,0.1] Hz)

✓ **Correlation between eFC and sFC** (goodness-of-fit)

✓ Parameter optimization (Covariance Matrix Adaptation Evolution Strategy) [4,5]

Low dimensional (2 parameters), **High dimensional** (around 100 parameters)



Step 5: Machine learning for model-based prediction

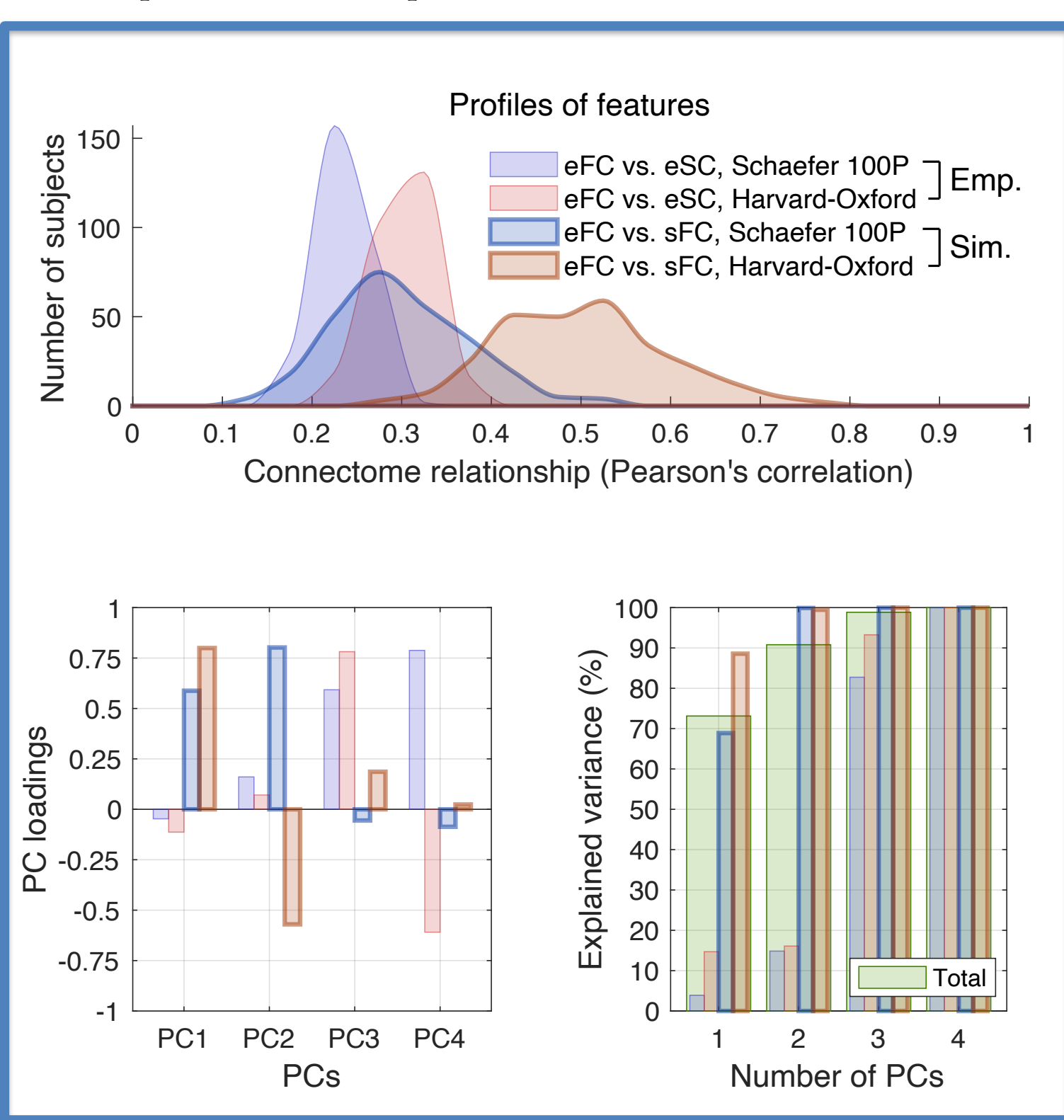
- Empirical feature**: corr(eFC, eSC) and **Simulated feature**: corr(eFC, sFC)
- Cross-validated confound removal scheme (5-fold nested cross validation; n=100)

✓ Classification of **females** and **males** (confound: brain volume)

✓ Prediction of **cognitive composite score** and the **Big-Five personality traits** (confound: brain volume and age)

Results: Simulated features outperforming empirical features in machine-learning analysis

1. Empirical (Emp.) and simulated (Sim.) features



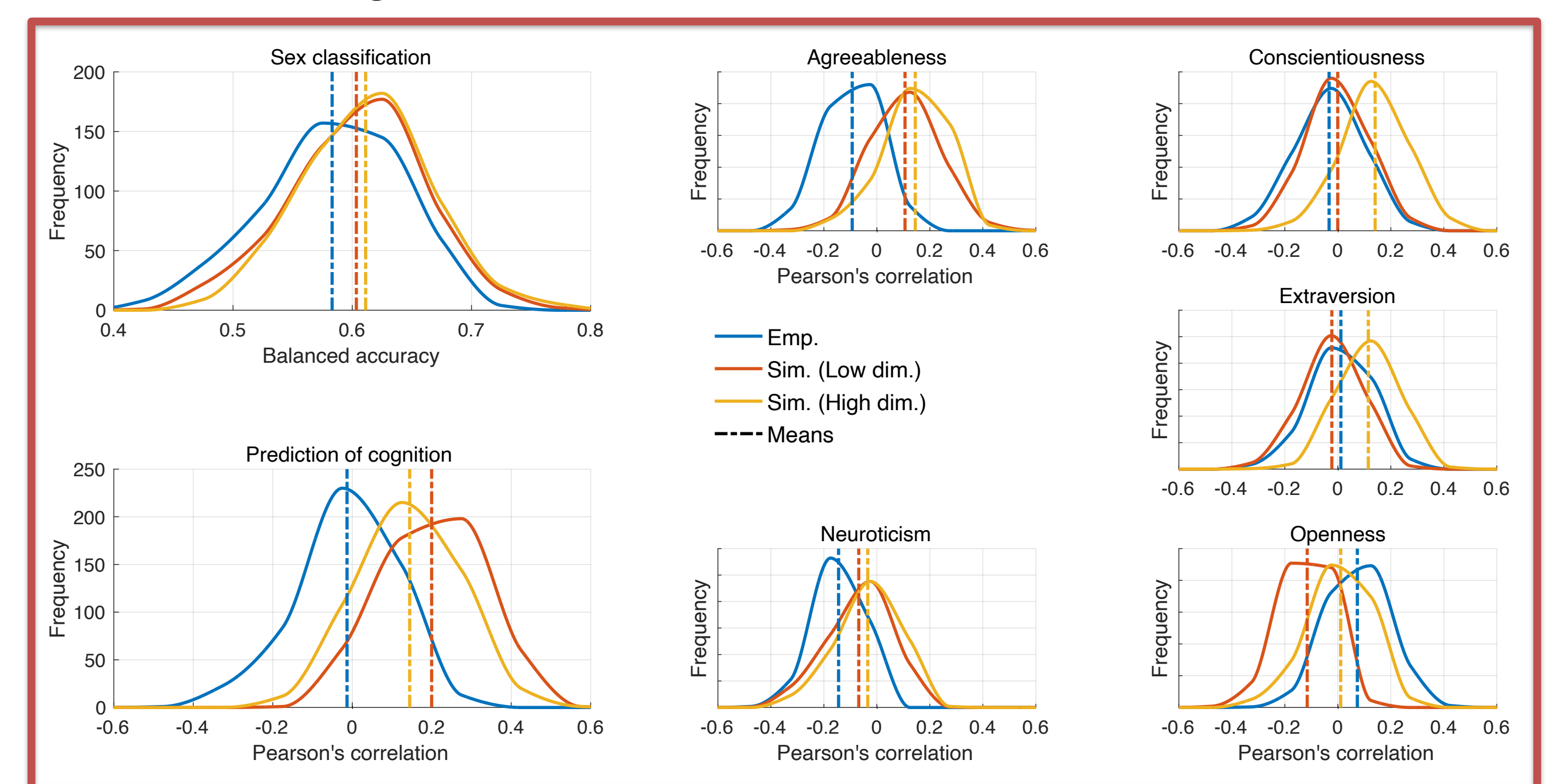
- Feature distributions across individual subjects for two brain parcellations.
- PC1 and PC2 are related to the simulated features and cumulatively explain 90% of the variance of features.
- PC2 and PC4 distinguish the two parcellation schemes.

	*ES (p value)
Sex classification: Sim. (High dim.)	1.16 (0.00)
Cognition: Sim. (Low dim.)	0.84 (0.00)
Agreeableness: Sim. (High dim.)	0.76 (0.00)
Conscientiousness: Sim. (High dim.)	0.76 (0.00)
Extraversion: Sim. (High dim.)	0.63 (0.00)
Neuroticism: Sim. (High dim.)	-0.21 (0.00)
Openness: Emp.	0.46 (0.00)

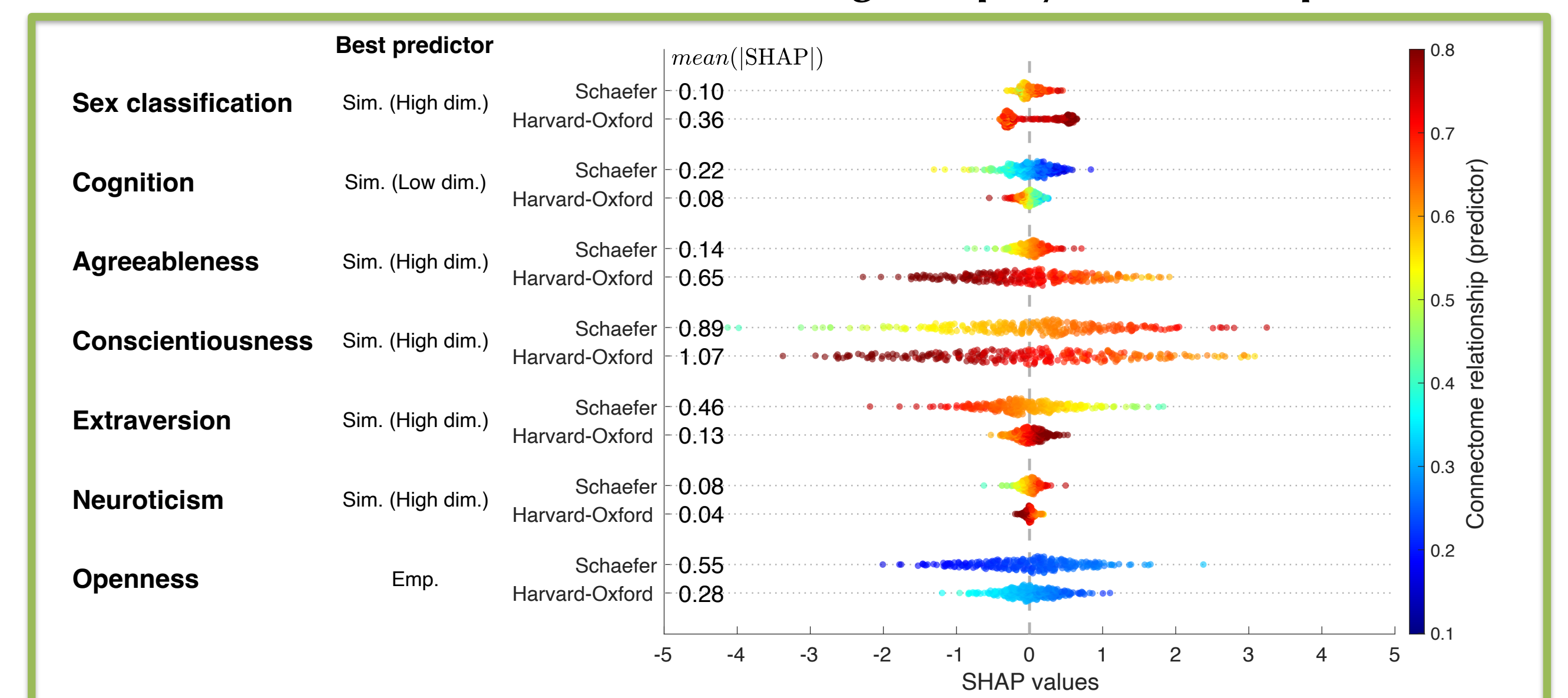
- Distributions of Shapley additive explanation* (SHAP) values [6] (data points = individual subjects).
- SHAP value itself does not indicate the magnitude of performance.
- Parcellation schemes can contribute to the prediction in different ways.

*A large |SHAP| means a strong contribution.

2. Machine-learning results



3. Feature contribution in machine learning (Shapley additive explanation)



Conclusion

- By incorporating **model-based features** alongside empirical data, we can explore brain connectomes and their interrelationships, thereby **enhancing performance** and bringing additional benefits in neuroimaging analysis.
- We propose to consider the simulated data as **an additional neuroimaging data modality** that captures distinct properties barely present in empirical data and can be integrated into machine-learning applications.

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