

# Enrichment of data analytics by whole-brain computational models



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### Introduction

By means of the whole-brain dynamical models we investigated the impact of the neuroimaging data processing on the results of model validation against empirical data. In this study we considered several brain parcellations:

 The functional Schaefer atlas [1] with 100, 200 and 400 cortical parcels (S100, S200 and S400) and anatomical Harvard-Oxford atlas [2] with 96 cortical parcels and several thresholds of the maximal probability (HO96 0% - HO96 45%).

We investigated the impact of the brain atlases and a few data variables on the model fitting and showed that these conditions may strongly influence the modeling results.

**Goal:** To find an optimal parameter setting for data-driven mathematical modeling of the restingstate brain dynamics and data analytics.

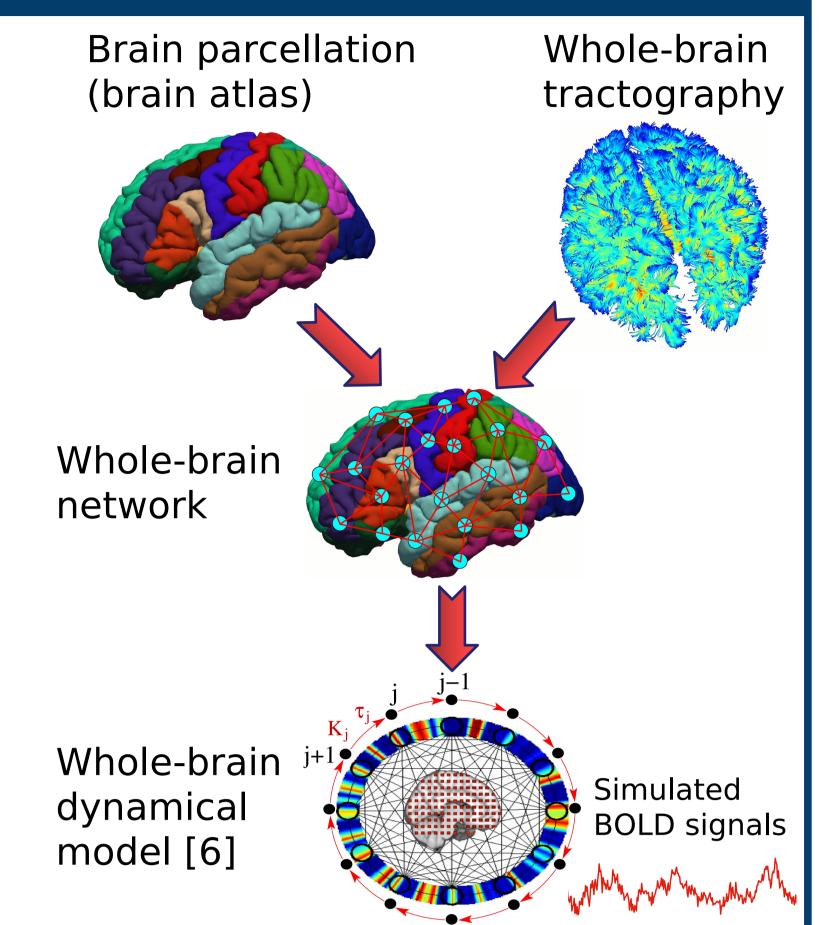
## Methods

#### **Computational model**

Kuramoto model [3] of coupled phase oscillators was used to simulate the dynamics of the phases  $\theta_i(t)$  of network nodes  $i=1,\cdots,N$ , which represent the mean resting-state dynamics of brain regions [4,5]

$$\frac{d\theta_i}{dt} = 2\pi f_i + \frac{C}{N} \sum_{j=1}^{N} k_{ij} \sin\left[\theta_j (t - \tau_{ij}) - \theta_i(t)\right] + \eta_i(t)$$

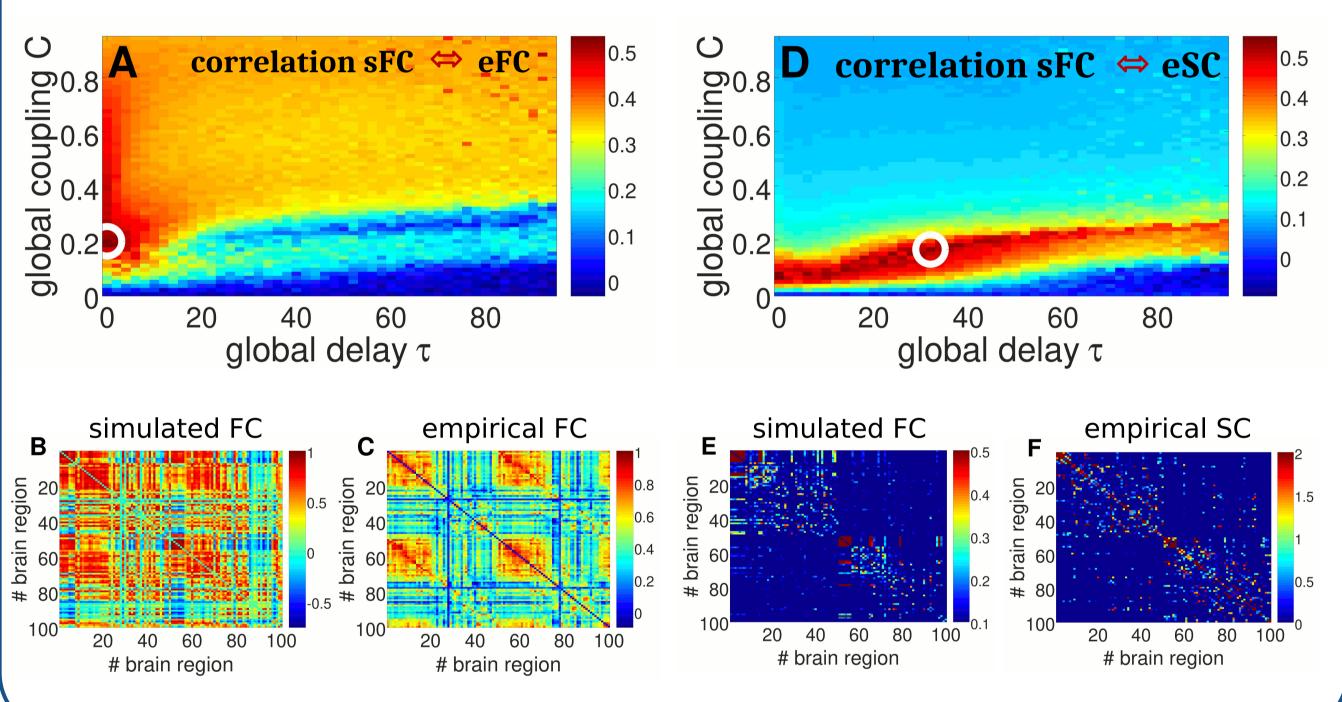
| Model<br>variables  | Description   | Model<br>variables                   | Description  |
|---|---|--------------------------------------|--|
| $\theta_i(t)$   | phase of node $i$ at time $t$   | $L_{ij}$                             | average fiber path length of eSC from node $j$ to node $i$ |
| $f_i$   | natural frequency of node i   | V                                    | mean conduction velocity                                   |
| C   | parameter of global coupling strength   | $\tau = \langle L_{ij} \rangle / V$  | parameter of global delay                                  |
| $k_{ij} = n_{ij}/\langle n_{ij} \rangle$                            | relative number of streamlines in the empirical structural connectivity (eSC) from node $j$ to node $i$ | $\eta_i(t)$                          | independent noise  |
| $\tau_{ij} = L_{ij}/V$ $= \tau \cdot L_{ij}/\langle L_{ij} \rangle$ | coupling delay (signal conduction time) from node $j$ to node $i$                                       | $S_i = \sin\left[\theta_i(t)\right]$ | simulated BOLD signals                                     |



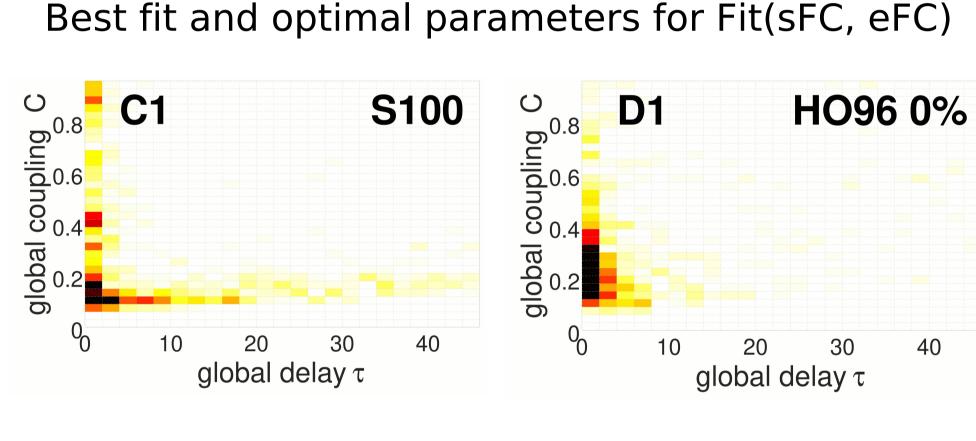
#### Results

# Model validation (personalized simulations)

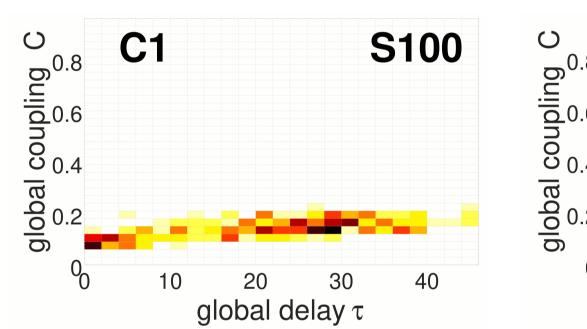
Simulated functional connectivity (sFC) is compared (by Pearson's correlation) with empirical FC (eFC) and structural connectivity (eSC) for the best fitting Fit(sFC, eFC) and Fit(sFC, eSC).

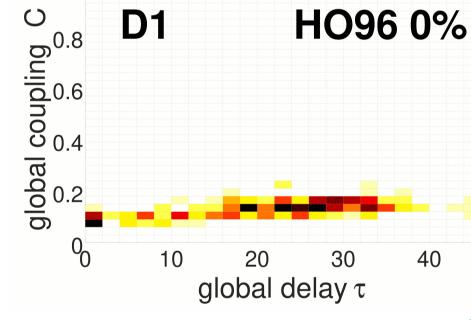


# Model validation (group-level analysis, n=272 HCP) eSC)

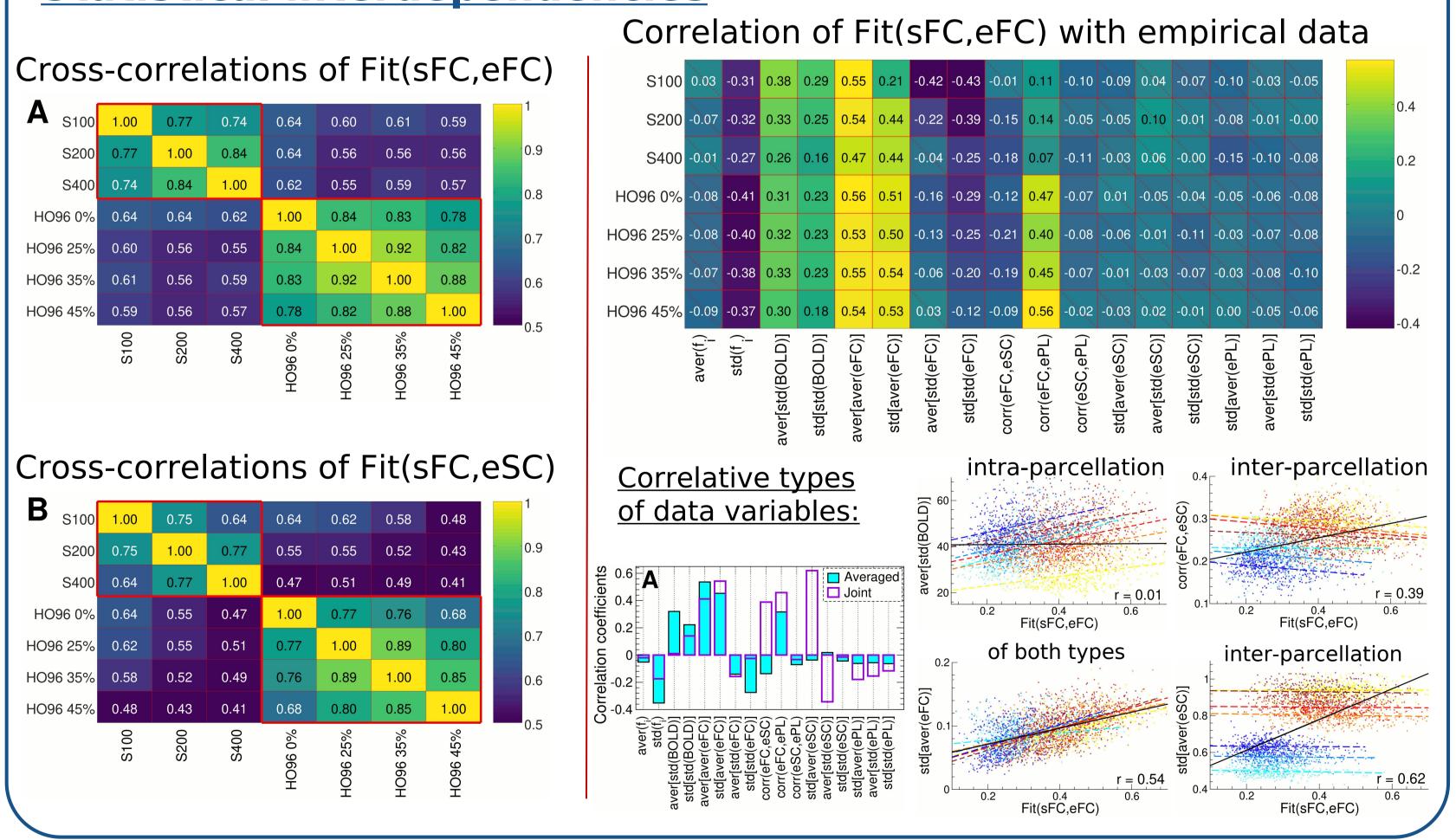




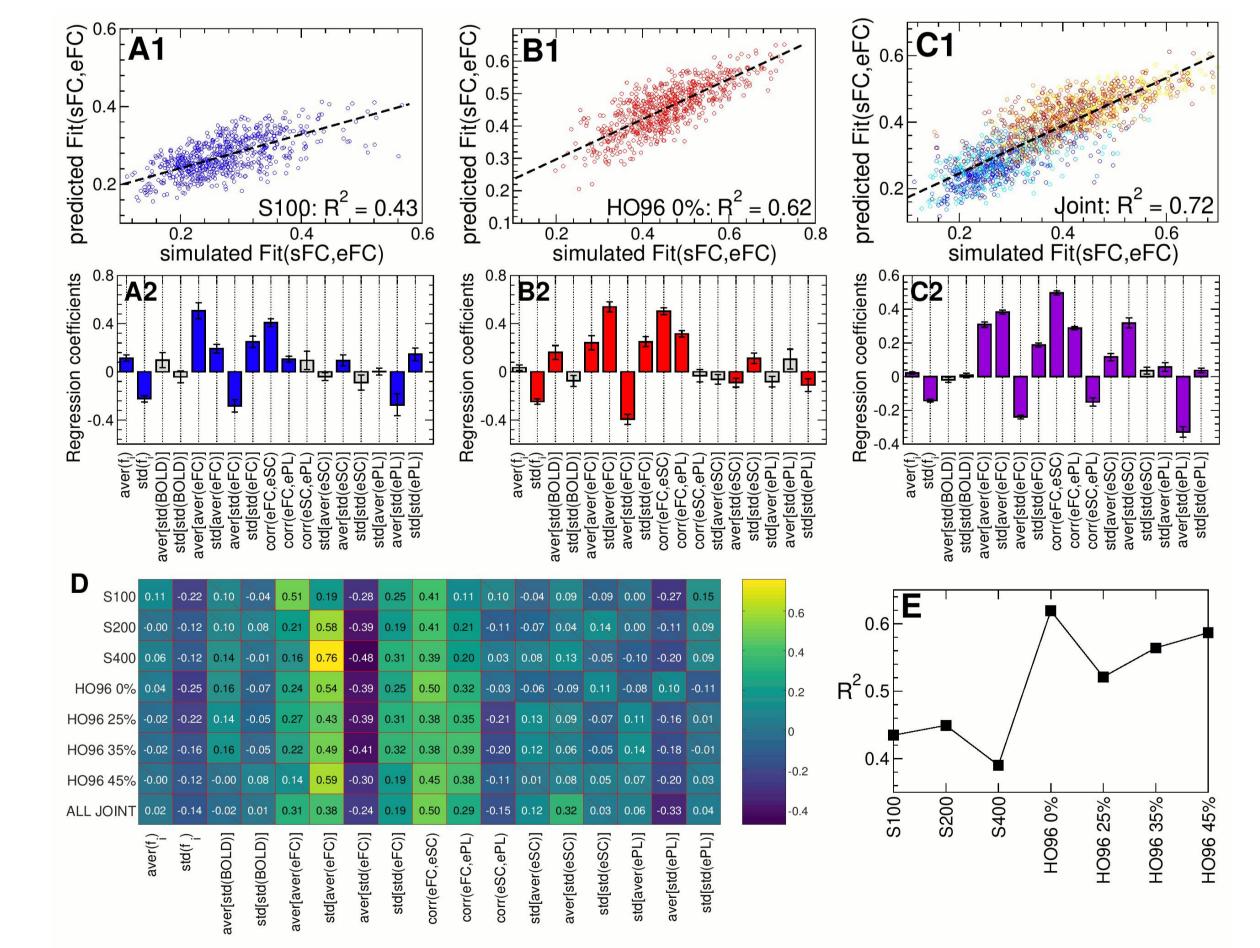




# Statistical interdependencies



# Multiple linear regression model (MLR)



#### Conclusions

- A choice of a particular brain parcellation can cause a strong impact on the quality of the model validation and structure of the model parameter space, which may deviate for different fitting modalities.
- The variation of Fit-values across subjects and parcellations can be accounted for by data variables of different correlative types: intra-parcellation variables, interparcellation variables and of both types.
- The results of MLR modeling showed that up to 70% of the variance of modeling results (Fit-values) can be explained by the properties of empirical data used for model derivation and validation.

**References**: [1] A. Schaefer et al., Cereb. Cortex 28, 3095 (2018). [2] R.S. Desikan et al., Neuroimage **31**, 968 (2006).

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