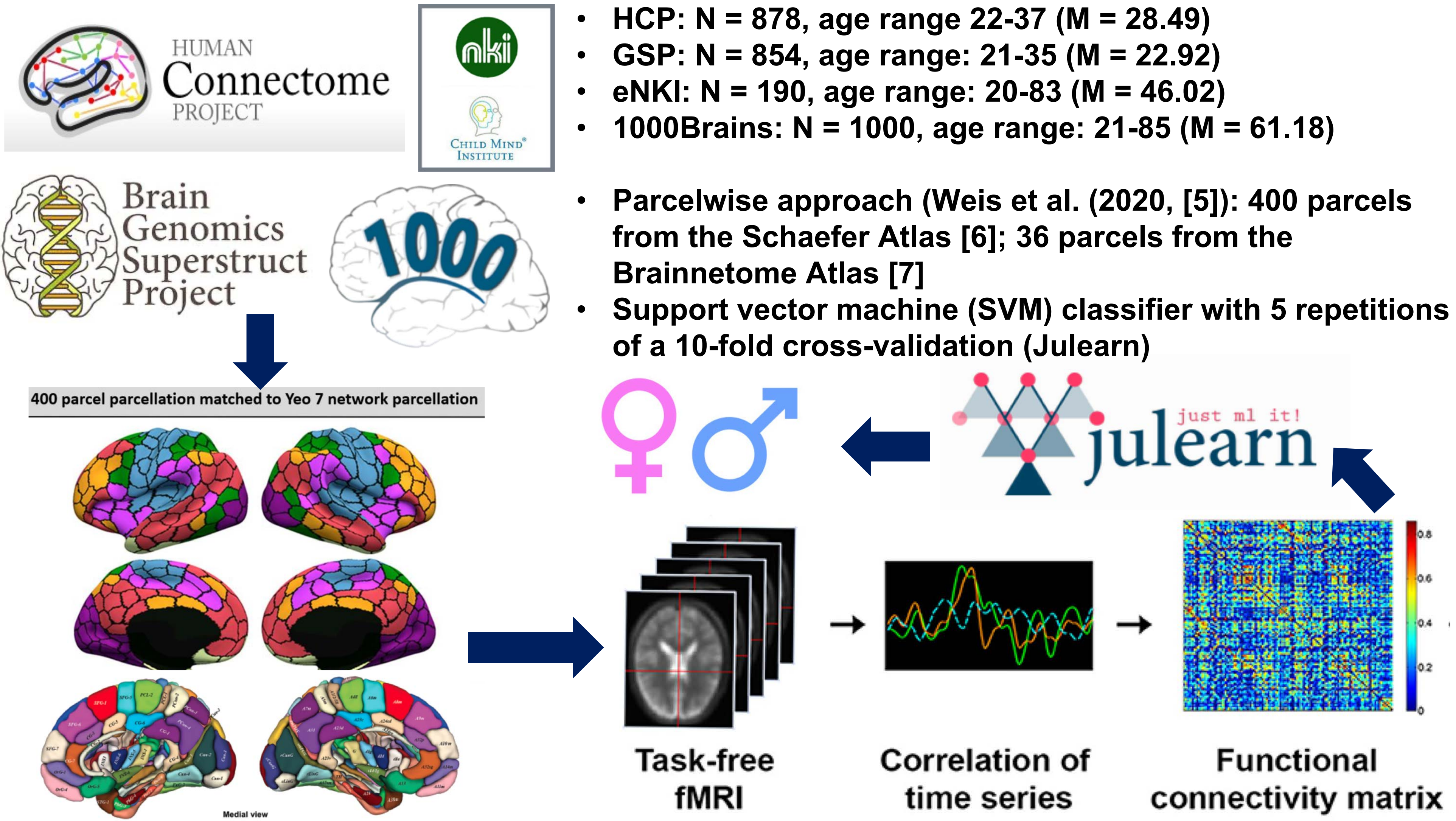


Introduction

- Machine-learning analyses allow for the prediction of phenotypes from neuroimaging data (e.g. sex)
- Which sample characteristics provide highest model performance for within- and between-sample predictions?
- The present study addresses this question for sex classification analyses based on the resting-state functional connectivity (RSFC)
- RSFC of four cohorts differing in sample size, age range and image quality (HCP [1], GSP [2], eNKI [3] and 1000Brains [4])
- Data of the four cohorts were combined in various ways to examine which model provides highest classification accuracies and high generalizability

Methods



Results

Table 1. Sex classification accuracies for within- and between sample predictions for model 1-5

|         | Model trained on                                | Within – predictions | Model applications                                                                                                                       |
|---------|-------------------------------------------------|----------------------|------------------------------------------------------------------------------------------------------------------------------------------|
| Model 1 | 75% of 1000Brains, eNKI, GSP and HCP (N = 2190) | 62.9 (53.7 - 70.1)   | HCP (25%): 65.3 (45.9 - 74.6)<br>GSP (25%): 60.6 (49.1 - 70.1)<br>eNKI (25%): 64.8 (47.9 - 83.3)<br>1000Brains (25%): 62.9 (52.4 - 75.2) |
| Model 2 | eNKI, GSP, HCP (N = 1922)                       | 64.9 (53.0 - 71.3)   | 1000Brains: 58.0 (49.5 - 68.0)                                                                                                           |
| Model 3 | 1000Brains, GSP, HCP (N = 2732)                 | 63.8 (53.0 - 70.5)   | eNKI: 61.4 (50.0 - 72.6)                                                                                                                 |
| Model 4 | 1000Brains, eNKI, HCP (N = 2068)                | 63.9 (52.8 - 71.7)   | GSP: 57.3 (48.7 - 68.6)                                                                                                                  |
| Model 5 | 1000Brains, eNKI, GSP (N = 2044)                | 61.5 (54.1 - 68.4)   | HCP: 59.4 (48.8 - 69.9)                                                                                                                  |

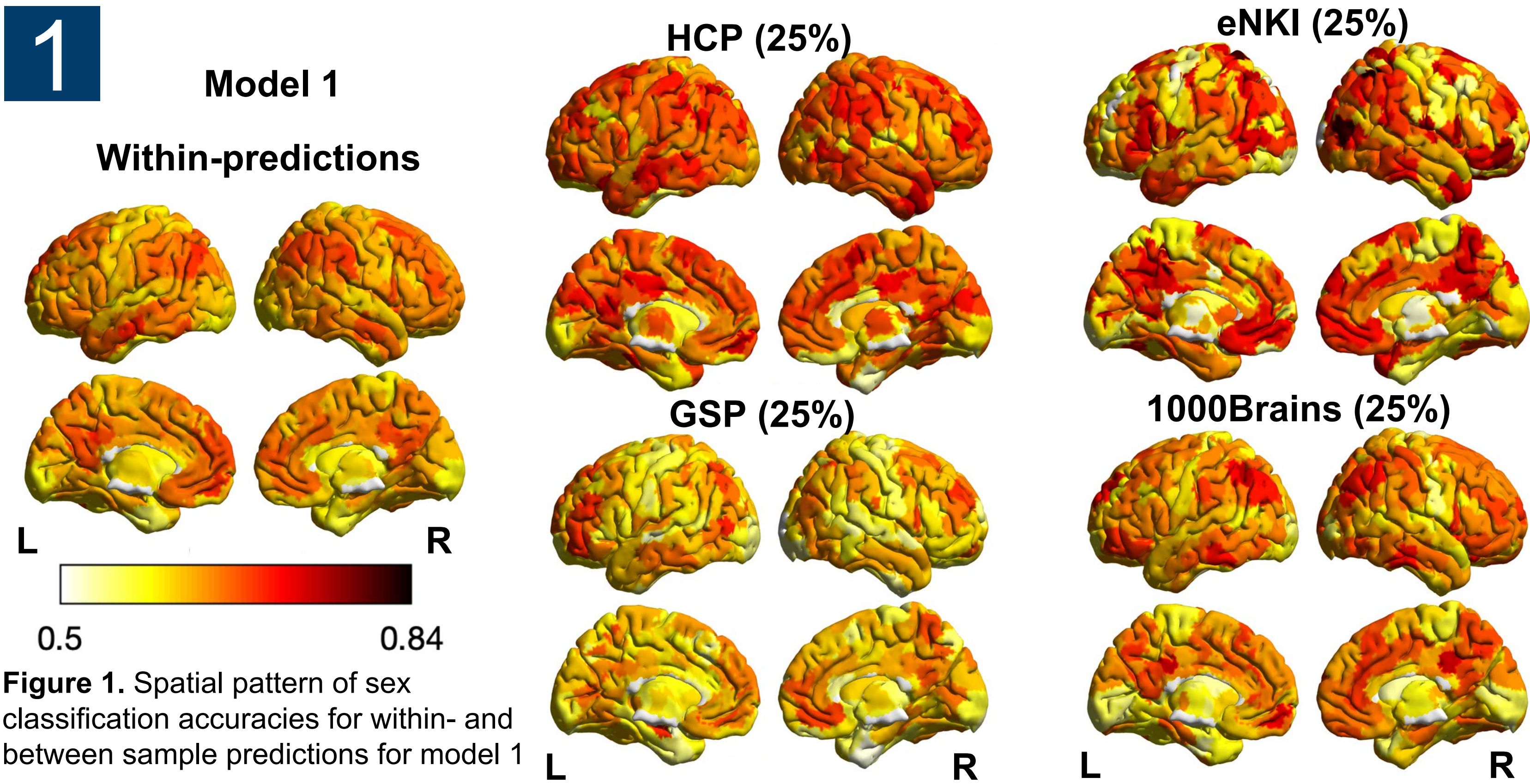


Figure 1. Spatial pattern of sex classification accuracies for within- and between sample predictions for model 1

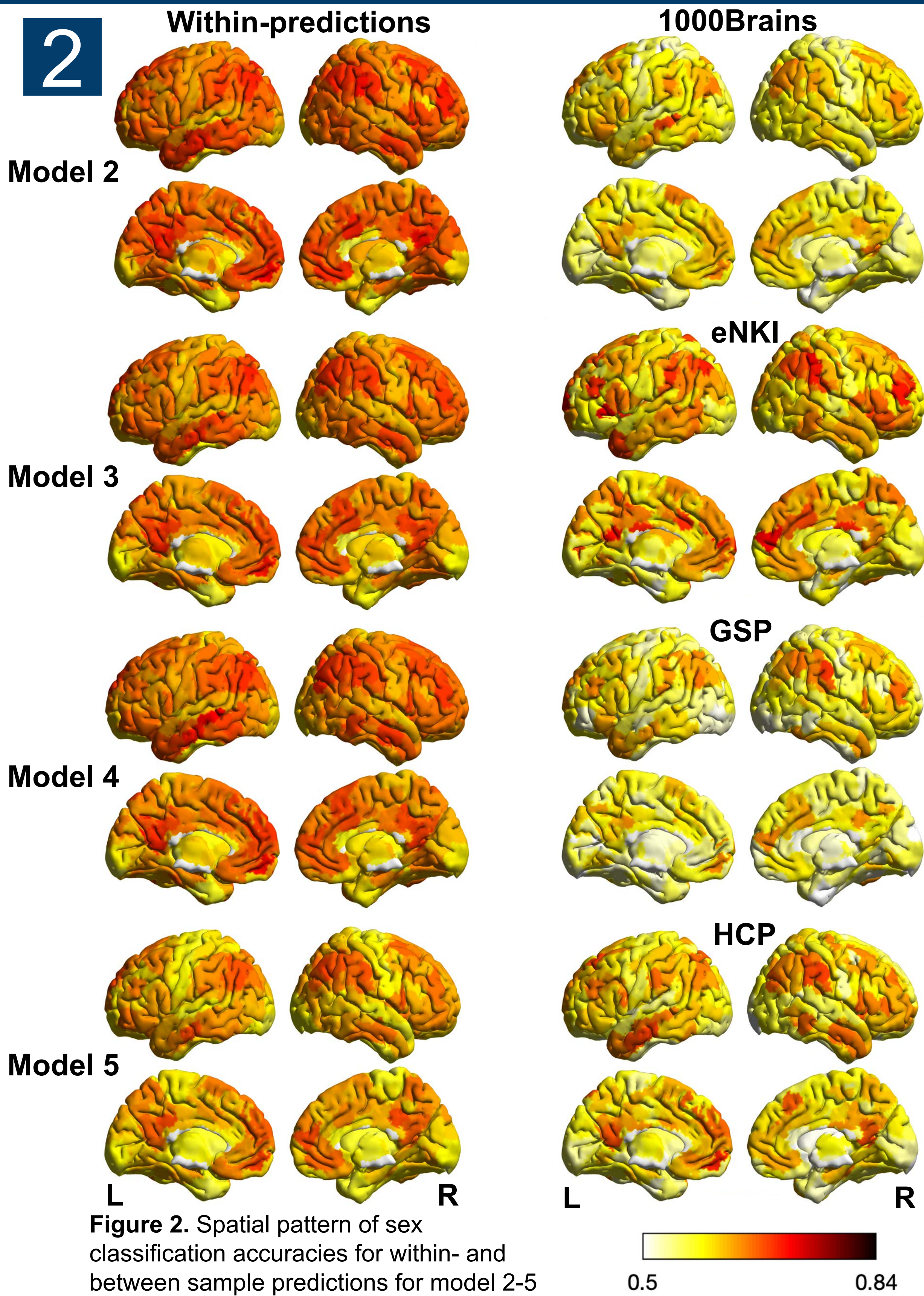


Figure 2. Spatial pattern of sex classification accuracies for within- and between sample predictions for model 2-5

Discussion

- Model 2 was trained on smallest sample size but outperformed all other models the in within-predictions
- Model 3 displayed higher between-prediction accuracy compared to model 2, 4 and 5
- Despite similar sample sizes, model 4 and 5 differed in classification accuracies:  
→ model 4 achieved higher within-predictions, model 5 achieved higher between-predictions
- Model 2,3 and 4 were trained on samples of different size but exhibited similar ranges of classification accuracies for within-predictions  
→ including HCP in training sample results in similar accuracies for within-predictions (63-64% mean prediction accuracy)
- Model 1 achieved highest accuracies for the eNKI-dataset with up to 83%, exceeding also the within-predictions for that model  
→ Model 1 generalized best

- Consistent spatial pattern of highly classifying parcels despite differently trained models
- Higher sample size in training does not necessarily lead to higher accuracies
- Best generalization performance for model 1 with heterogenous training sample, including parts of the test-dataset to adapt the model accordingly

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