



# Stability of resting state functional connectivity within intensive longitudinal designs



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## Background

Traditional group-based neuroimaging studies often lack the power to detect brain-behavior associations<sup>1</sup>. Recent perspectives<sup>2,3</sup> emphasized two complementary strategies to address these limitations:

- 1. Intensive longitudinal designs (Fig. 1)** - Derives statistical power from repeated measurements within individuals, enabling detection of within-subject effects over time
- 2. Precision neuroimaging** - Improves measurement reliability through high-quality, individualised data acquisition.

Together, these approaches make the investigation of within-subject brain-behaviour associations possible.

To inform these goals, here we investigate naturally occurring between-person and within-person (i.e. session-to-session) variance of functional connectivity to evaluate how powered precision intensive longitudinal designs actually are

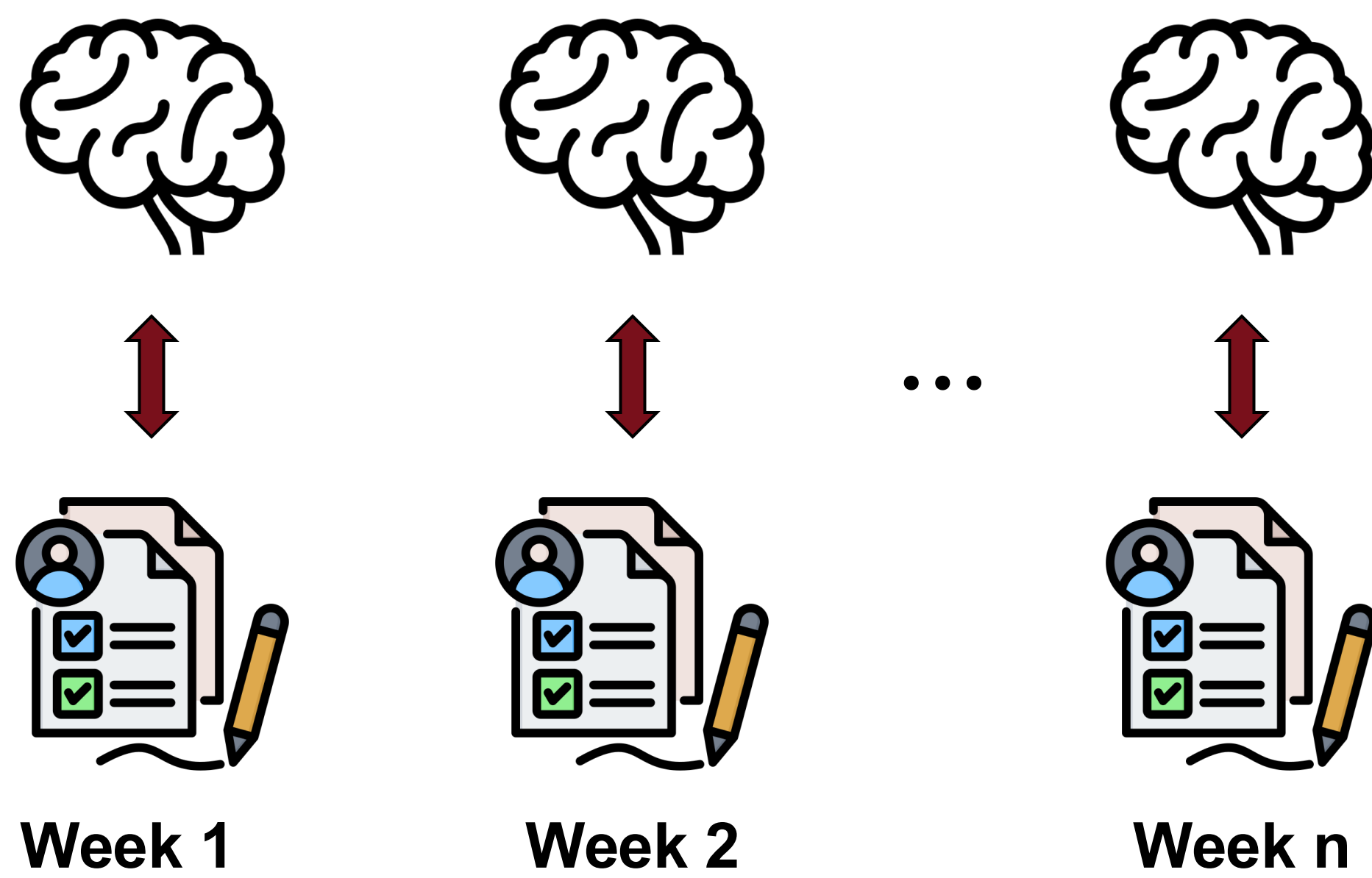


Figure 1. Dense longitudinal neuroimaging design

## Methods

### Dataset

- N = 14 participants (5 males, ages = 9-11), each scanned 3-4 times in 1-to-8-week intervals
  - Minimum of **30 minutes** of **low motion** resting-state fMRI acquired using **multi-echo** (ME) multi-band sequence (echo = 4, TR = 1761ms, 2.0 mm isotropic voxels) per session
- All preprocessing was performed using fMRIPrep<sup>4</sup> and XCPd<sup>5</sup>. Data were parcellated with Glasser<sup>6</sup> atlas

### Analysis

- Linear mixed effects model with random intercepts for each subject fitted for every edge
- Edgewise interclass correlation coefficient (ICC) reliability calculated as the ratio of between and total variance
- Disaggregated within-subject power was calculated using simulated data with variance components from empirical data.
  - Effect sizes of brain-behaviour associations set to:  $r = 0.1$  and  $r = 0.2$
  - Behavioural variance components were taken from previous work<sup>7</sup>

$$Y_{ij} = \beta_0 + \beta_B \bar{X}_i + \beta_W (X_{ij} - \bar{X}_i) + u_i + \varepsilon_{ij}$$

## Results

### 1. Natural occurring within-subject variance in functional connectivity is very low (Fig. 2)

Between-subject variance (i.e., stable individual differences between participants) in higher-order networks was up to an order of magnitude higher than within-subject variance (i.e., session-to-session changes).

### 2. Functional connectivity is most stable (highest ICC) within and between higher order and attention networks (Fig 3). Sensory networks should most variability

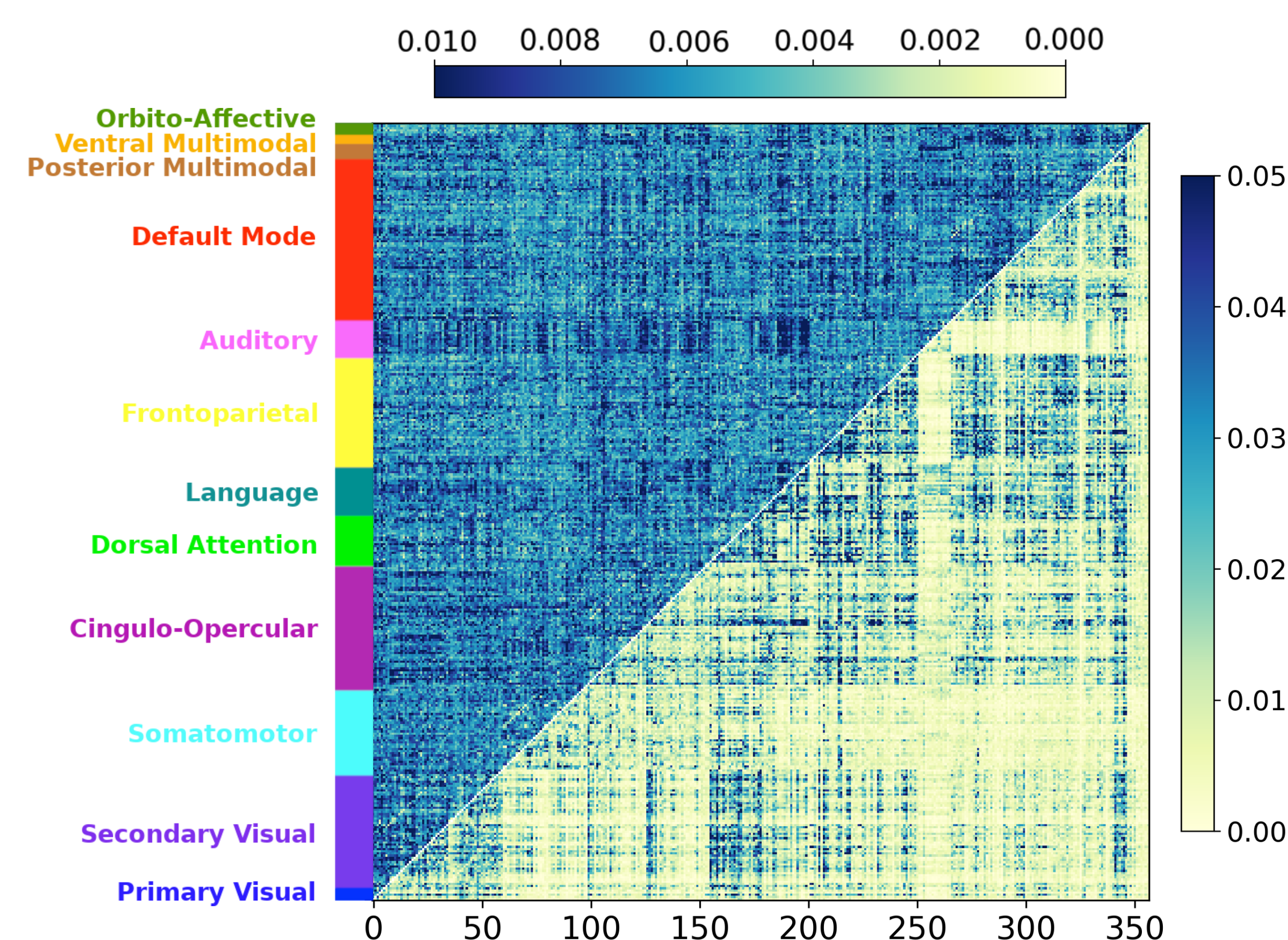


Figure 2. Edgewise between and within subject variance

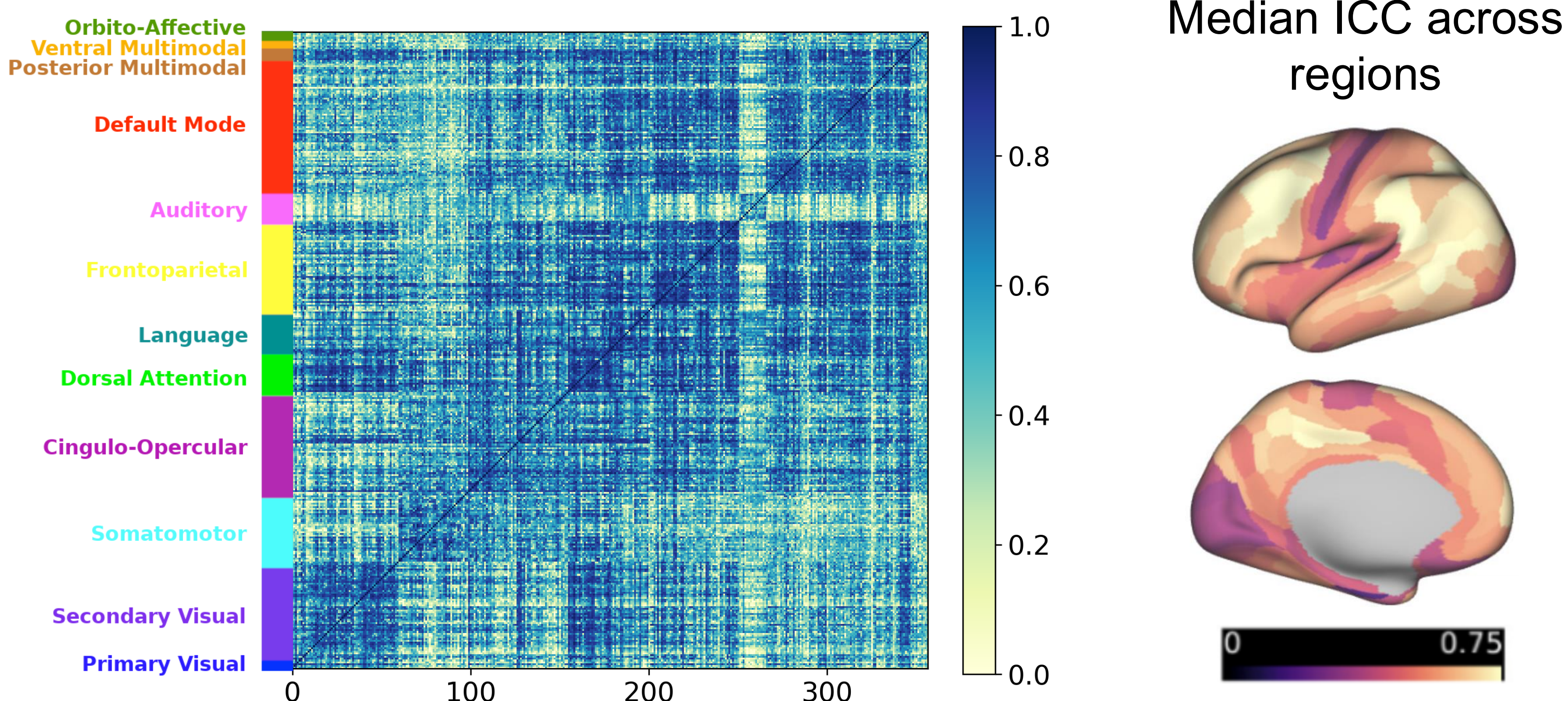


Figure 3. Edgewise and nodewise stability (ICC) of functional connectivity

### 3. Within-subject brain-behaviour associations likely require modest sample sizes and large number of repeat sessions (Fig. 3)

Figure 3 displays statistical power for a range of sample sizes and within-subject sessions to identify a weak association between the connectivity of frontoparietal and dorsal attention networks and a reliable phenotype (ICC = 0.75)

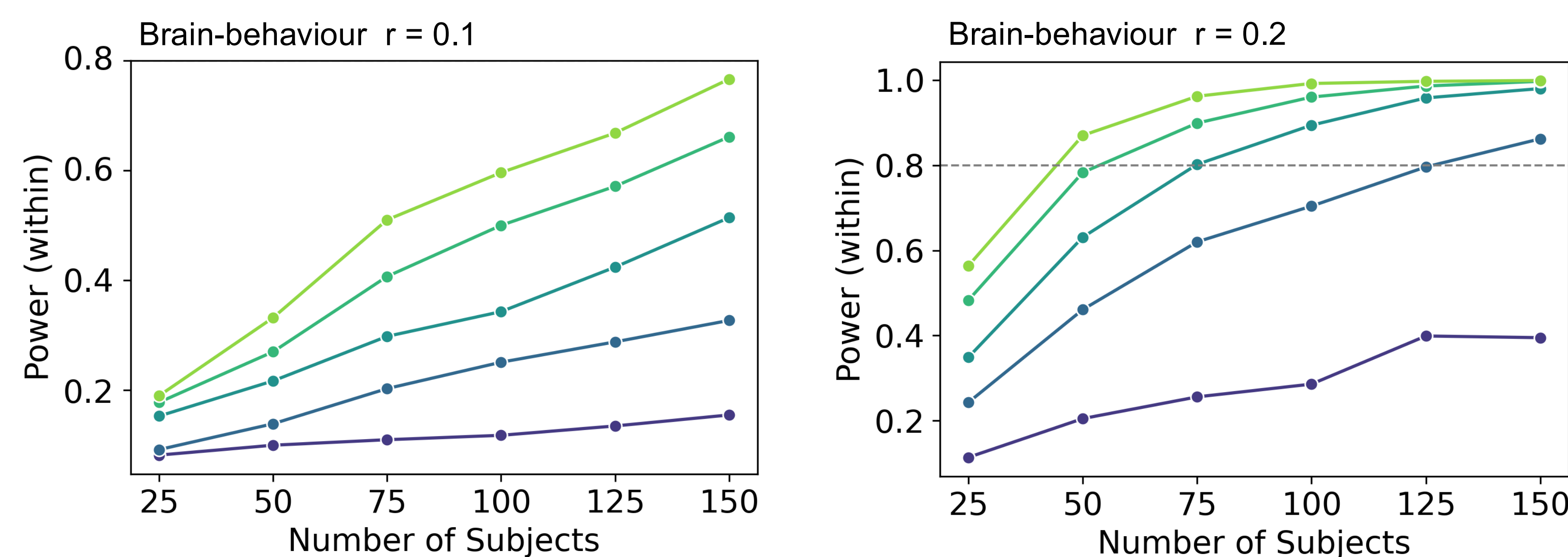


Figure 3. Statistical power to identify small effect size within-subject brain-behaviour associations

## Conclusions

These findings strongly support “trait-like” stability of the functional connectome across scanning sessions  
Unstructured, naturally occurring within-person changes in functional connectivity over weeks to months are minimal

**Tracking brain-behavior associations in densely sampled individuals will likely require large associations, sizable samples, or perturbations (e.g., task paradigm) to amplify within-subject variability.**

**References:** 1. Marek, S., Tervo-Clemmens, B., et al., (2022). Reproducible brain-wide association studies require thousands of individuals. *Nature*; 2. Gratton, C., et al., (2022). Brain-behavior correlations: Two paths toward reliability. *Neuron*; 3. Kraus, B., et al., (2023). Insights from personalized models of brain and behavior for identifying biomarkers in psychiatry. *Neuroscience & Biobehavioral Reviews*; 4. Esteban, O., et al., (2019). fMRIPrep: A robust preprocessing pipeline for functional MRI. *Nature Methods*; 5. Mehta, K., et al., (2023). XCP-D: A Robust Pipeline for the post-processing of fMRI data. *bioRxiv*; 6. Glasser, M. F., et al., (2016). A multi-modal parcellation of human cerebral cortex. *Nature*; 7. Voelkle, M. C., et al., (2014). Toward a Unified Framework for the Study of Between-Person and Within-Person Structures: Building a Bridge Between Two Research Paradigms. *Multivariate Behavioral Research*.