



Eliana Nicolaisen-Sobesky¹, Shahrzad Kharabian-Masouleh^{1,2}, Agoston Mihalik^{3,4,5}, Fabio Ferreira^{3,4}, Felix Hoffstaedter^{1,2}, Holger Schwender⁶, Somayeh Maleki Balajoo^{1,2}, Sofie L. Valk^{1,2,7}, Simon B. Eickhoff^{1,2}, B.T. Thomas Yeo^{8,9}, Janaina Mourao-Miranda^{3,4}, Sarah Genon^{1,2}

¹Institute of Neuroscience and Medicine (INM-7: Brain and Behaviour), Research Centre Jülich, Jülich, Germany; ²Institute of Systems Neuroscience, Heinrich Heine University Düsseldorf, Düsseldorf, Germany; ³Centre for Medical Image Computing, Department of Computer Science, University College London, London, United Kingdom; ⁴Max Planck University College London Centre for Computational Psychiatry and Ageing Research, University College London, London, United Kingdom; ⁵Department of Psychiatry, University of Cambridge, Cambridge, United Kingdom; ⁶Mathematical Institute, Heinrich Heine University Düsseldorf, Düsseldorf, Germany; ⁷Otto Hahn Research Group "Cognitive Neurogenetics", Max Planck Institute for Human Cognitive and Brain Sciences, Leipzig, Germany; ⁸Department of Electrical and Computer Engineering, ASTAR-NUS Clinical Imaging Research Centre, Singapore Institute for Neurotechnology and Memory Networks Program, National University of Singapore, Singapore; ⁹Centre for Cognitive Neuroscience, Duke-NUS Medical School, Singapore, Singapore.

e.nicolaisen@fz-juelich.de

Introduction

- Brain-behaviour associations are often not replicable^{1,2,3}.
- Our aim is to identify robust and replicable associations between interindividual variability in **brain structure and behaviour**.
- We use a multivariate method to link brain and behaviour: **Regularized Canonical Correlation Analysis (RCCA)**^{4,5}.
- To test the generalisability of the model, we use a novel machine learning framework that embeds RCCA in **multiple holdouts** of the data^{4,6}.
- Crucially, we tested the **cross-cohort replicability** of the brain-behaviour associations.
- In addition, the nature vs nurture influence on associations was studied with **heritability**.

Methods

1) Datasets:

- HCP Young Adult (HCP-YA):** (S1200), n=1047 (560 females), age = 28.8, 22-37 years (mean, range)
- HCP-aging (HCP-A):** n=601 (353 females), age= 58.5, 36-100 years (mean, range)

Brain data (639 measures):

- Cortical Thickness (CT)
- Surface Area (SA)
- Grey Matter Volume (GMV)

Behavioural data (32 measures):

- Alertness
- Cognition
- Emotion

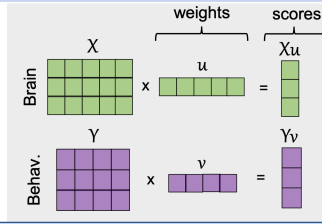
Age and gender were regressed out from brain and behaviour.

Processing of brain data:

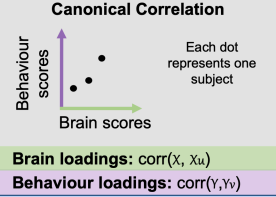
- CT and SA: FreeSurfer v5.3.0-HCP in HCP-YA; v6.0 in HCP-A
- GMV: CAT12.5 (both cohorts)
- Averaged by regions (200 cortical⁷, 32 subcortical⁸, 7 cerebellar⁹)
- Normalised within subjects by brain size

2) Regularized Canonical Correlation Analysis:

- X and Y: subjects are in rows, measures are in columns
- RCCA finds the weights which maximize the canonical correlation
- Regularization: L2-norm constraints on weights
- RCCA was applied independently in each cohort.



Latent Dimension and Canonical Correlation



3) Machine Learning Framework: Multiple holdouts^{4,6}:

- Outer split (x5): 1) Train set (64%): Train several RCCA models with different regularization parameters
- Inner split (x5): 2) Test set (16%): Test models and select model with best generalizability
- 3) Optimisation set (80%): Train best RCCA model
- 4) Hold-out set (20%): Test best model

Code available at: https://github.com/anaston/cca_pls_toolkit

4) Statistical analyses:

- 1000 permutations, shuffling rows of Y, respecting family structure
- Omnibus hypothesis⁶: states that there is no association in any of the outer splits. Hence, if a significant association is found in at least one split, the null hypothesis can be rejected.

5) Replicability:

- Cross-cohort similarity of loadings:
- Pearson's correlation for behaviour
- Spin test for CT and SA

6) Heritability:

- Heritability (h^2) and genetic correlation (ρ_g) of brain and behavioural scores were studied using a twin-based analysis in SOLAR-Eclipse¹⁰ (HCP-YA).

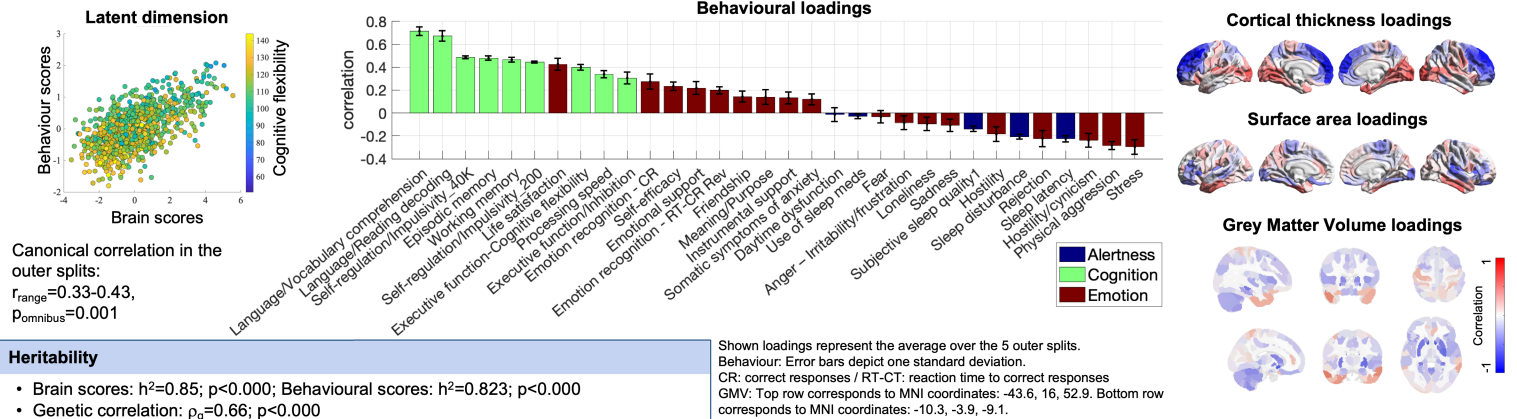
Results

The RCCA model in the HCP-YA yielded 3 significant latent dimensions. The RCCA model in the HCP-A yielded 2 significant latent dimensions. Of these, only **one latent dimension replicated across cohorts**, with significant correlations on the behavioural ($r=0.73$, $p<0.001$), CT ($r=0.81$, $p<0.001$) and SA ($r=0.56$, $p<0.001$) loadings.

This latent dimension can be understood as an axis capturing:

- Behaviour: one pole of **good cognitive-control/executive-functions and positive affect**, and another pole of **cognitive dysfunction/impulsivity and negative affect**.
- Brain: one pole associated to areas related to **higher cognitive functions**, and another pole associated with **sensorimotor regions**.

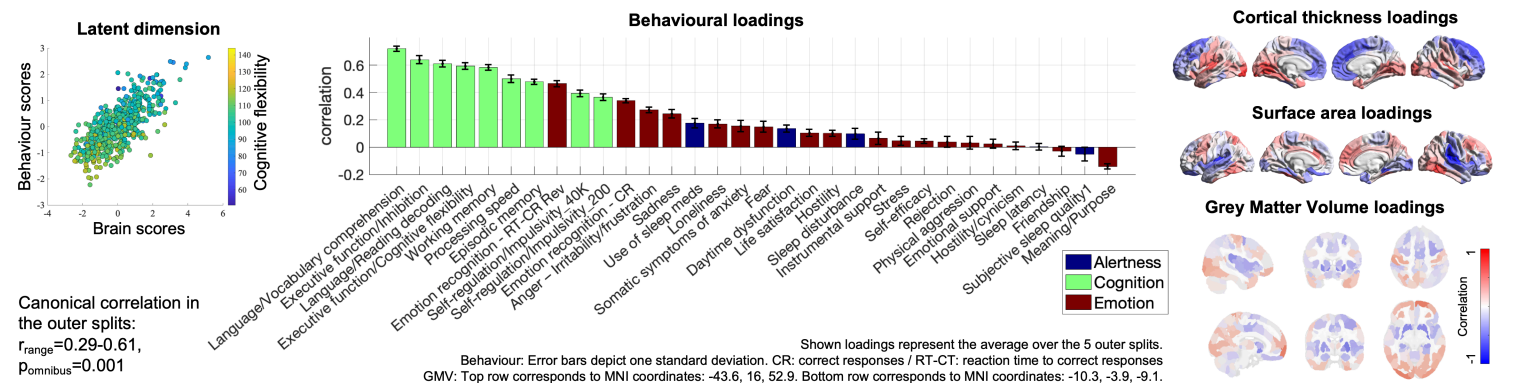
Latent dimension in HCP-YA



Heritability

- Brain scores: $h^2=0.85$; $p<0.000$; Behavioural scores: $h^2=0.823$; $p<0.000$
- Genetic correlation: $\rho_g=0.66$; $p<0.000$

Latent dimension in HCP-A



Discussion

- Our results show one heritable and replicable latent dimension linking interindividual variability in brain structure with interindividual variability in behaviour.
- The behavioural profile of this latent dimension was previously captured in association with brain function in the HCP-YA cohort^{11,12}. We extend such findings by analysing the brain structural profile of this dimension.
- This latent dimension can be interpreted as an axis in which

subjects are spread based on their covariance between brain structure and behaviour.

- Crucially, we test the replicability of this latent dimension and extend it to a sample with a wider age range (HCP-A).
- We also show that the brain and behavioural scores of this latent dimension are at least partly influenced by overlapping genetic mechanisms.

References

- Kharabian-Masouleh 2019. *Elife*.
- Kharabian-Masouleh 2020. *bioRxiv*.
- Marek 2022. *Nature*.
- Mihalik 2020. *Biol Psychiatry*
- Hardoon 2004. *Neural Computation*
- Monteiro 2016. *J Neuroscience Methods*
- Schaefer 2018. *Cereb Cortex*
- Tian 2020. *Nature Neuroscience*
- Yeo 2011. *J Neurophysiol*
- <http://www.solar-eclipse-genetics.org>
- Smith 2015. *Nature Neuroscience*
- Han 2020. *Neuroimage*