

INTRODUCTION

In imaging-genetics, the **heritability** of the quantitative phenotype under study is often considered a prerequisite for future studies, such as genome-wide association studies (GWAS)¹. The current work extends previous work from our group^{5,6} to examine the heritability of the volume and shape of the striatum, thalamus and globus pallidus. It is the first study to examine the **shared heritability** and **genetic correlation** between subcortical structure morphometry (**vertex-wise surface area (SA) and displacement**) and aggregate measures (total brain volume (TBV), subcortical structure volume and total SA) as a means of providing a nuanced description of subtle variation in neuroanatomy.

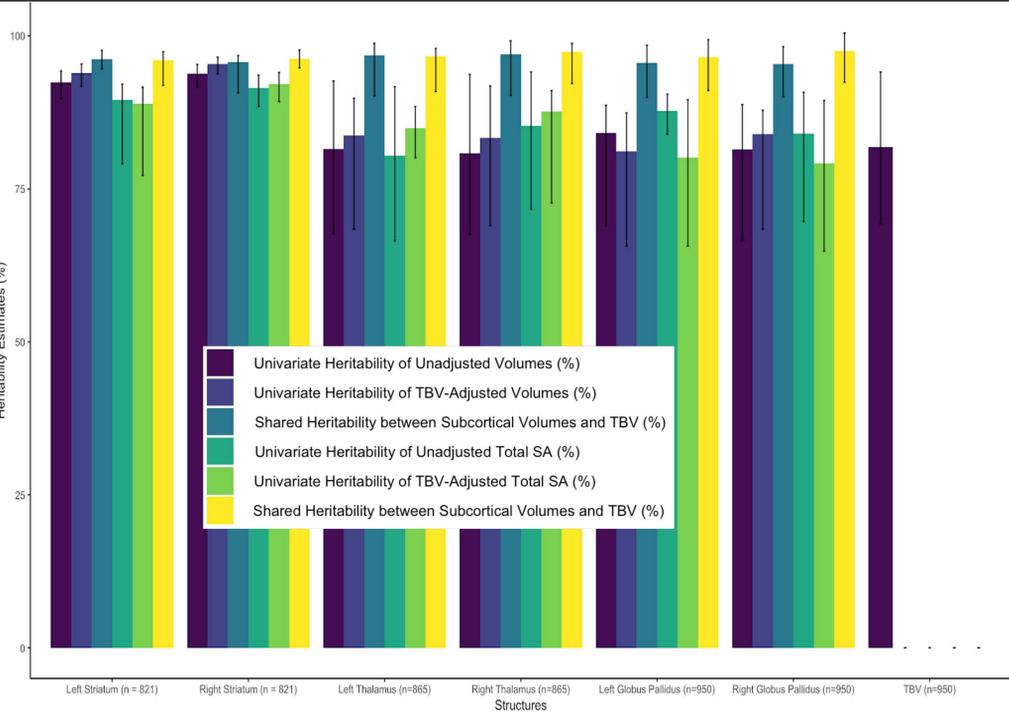


Figure 1. Univariate and bivariate heritability (h^2) estimates of subcortical structure volumes and total SA

GLOSSARY

[Univariate] heritability (h^2): proportion of variance of a phenotype attributable to additive genetic (as opposed to environmental) effects

[Bivariate] shared heritability (h^2): proportion of covariance between two phenotypes attributable to additive genetic (as opposed to environmental) effects

[Bivariate] genetic correlation (r_g): Proportion of the genetic factors that explain the variance of phenotype 1 which also explain the variance of phenotype 2.

METHODS

Data Acquisition: The WU-Minn Human Connectome Project². S1200 Release of structural MRI data on healthy young adult twin and non-twin siblings (N = 1086).

Image Processing: [minc-bpipe library](#) and [MAGeTbrain^{3,4}](#)

Heritability Estimates.

- The OpenMx package (version 2.12.2) in R (version 3.5.1) was used to compute heritability estimates⁵.
- Univariate model: heritability of one phenotype (structure volume, structure total SA, vertex-wise SA, vertex-wise displacement) at a time
- Bivariate model: shared heritability and genetic correlation between two phenotypes at a time: TBV and more local measures (structure volume, vertex-wise SA, vertex-wise displacement)

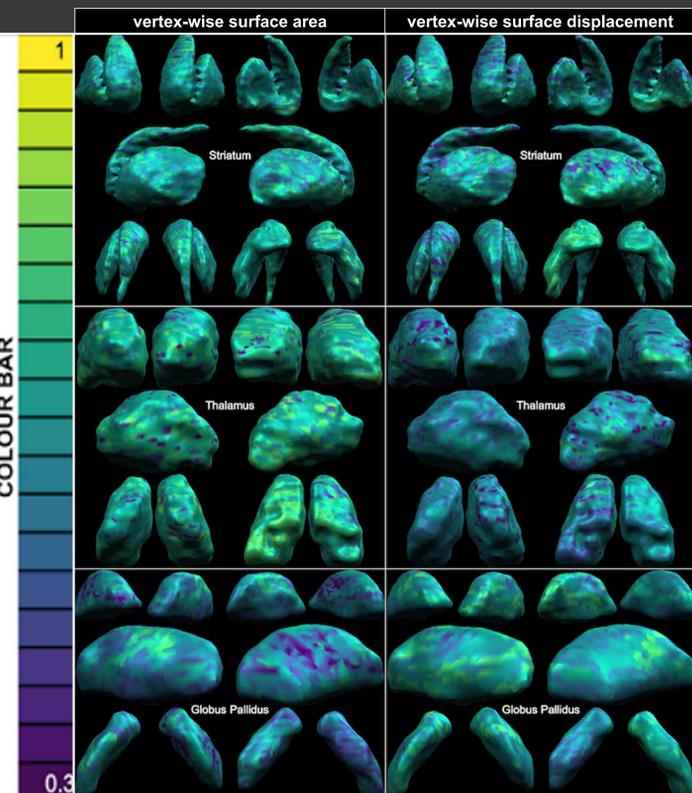


Figure 2. Vertex-wise **univariate heritability**

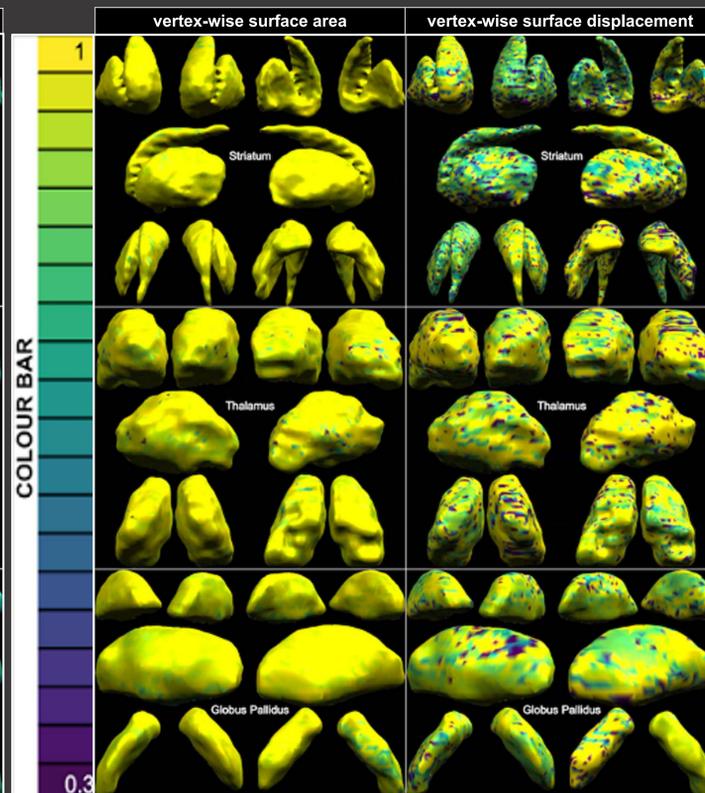


Figure 3. **Shared heritability** between TBV and vertex-wise measures

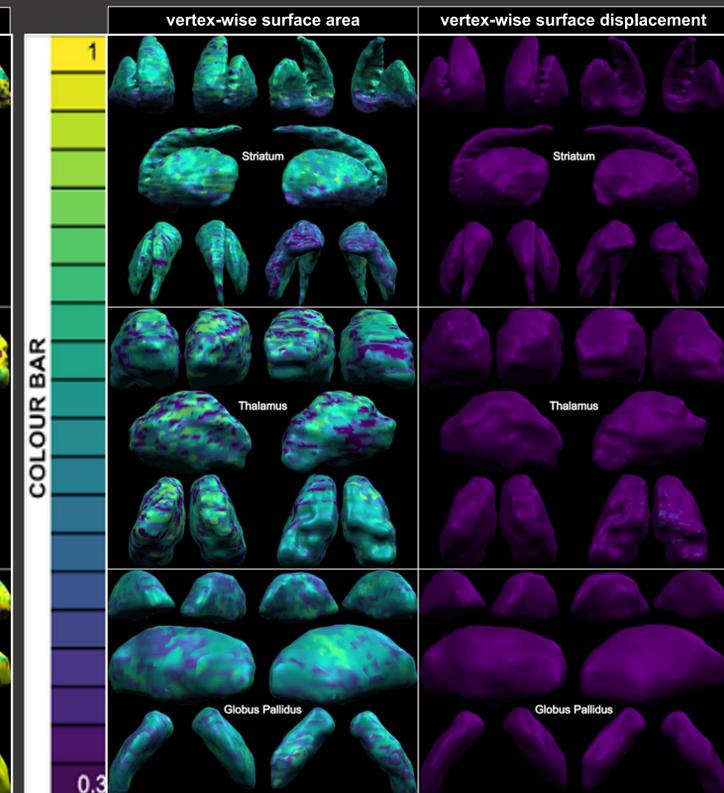


Figure 4. **Genetic correlation** between TBV and vertex-wise measures

MAIN FINDINGS

- Subcortical structure **volumes are highly heritable**. A huge portion of their covariance with TBV is explained by additive genetic effects
- Indirect evidence for the existence of **genetic factors with highly localized effects** on subcortical structures
→ vertex-wise heritability estimates are highly heritable yet only moderately genetically correlated with TBV
- The **ventral striatum** (associated with habit formation and addiction) is a lot less correlated with TBV than the dorsal striatum (associated with motor control).
- [Supplementary results](#)

FUTURE DIRECTIONS

- Intuitively, one would assume an inverse relationship between heritability and brain plasticity. [Our current work](#) therefore seeks to understand the relationship between heritability and evolutionary expansion of subcortical structures across non-human primates.

REFERENCES

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- ³Pipitone J, Park MT, Winterburn J, et al. Multi-atlas segmentation of the whole hippocampus and subfields using multiple automatically generated templates. *Neuroimage*. 2014;
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