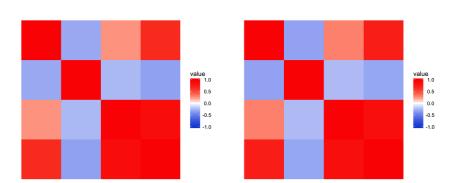
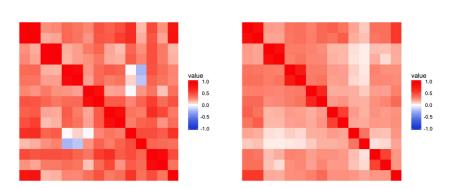
Supplementary Figure S4

Each of the subsequent pages shows the estimated genetic correlation matrix (left) and the raw phenotypic correlation matrix for different subsets of IDPs. The genetic correlation matrices where calculated using the multi-trait mixed model approach described in the main text using the SBAT software. In some cases, phenotypes were pruned to remove pairs of IDPs with raw correlation greater that 0.9. The intention of these plots is to provide an overview of the levels of raw and genetic correlations that exist between the different IDPs.

T1 - global volumes

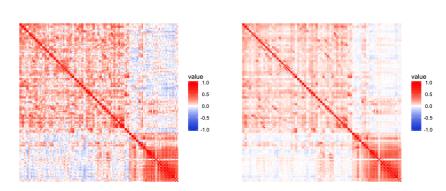


T1 - sub-cortical volumes

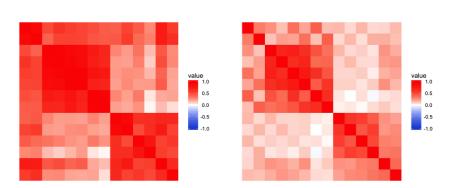


T1 - Cortex ROIs

Genetic correlation

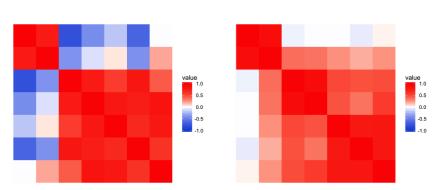


SWI T2* sub-cortical



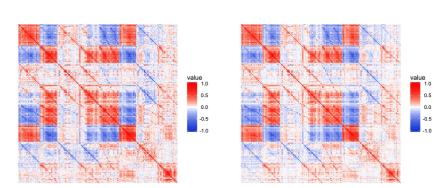
task fMRI





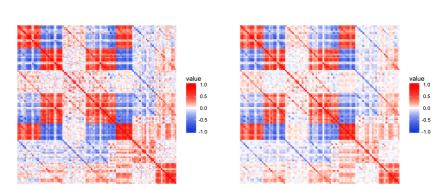
Diffusion MRI - TBSS

Genetic correlation

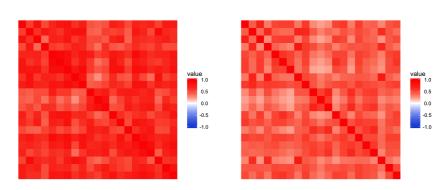


Diffusion MRI - ProbtrackX

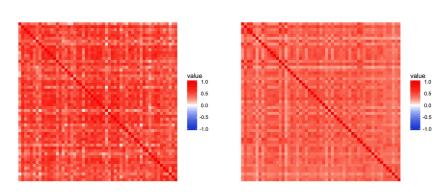
Genetic correlation



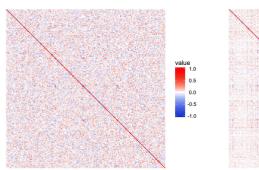
resting fMRI - parcellation25 amplitudes

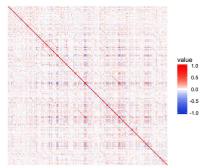


resting fMRI - parcellation100 amplitudes

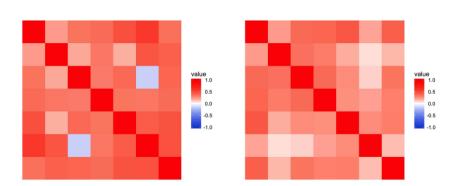


resting fMRI - parcellation25 edges

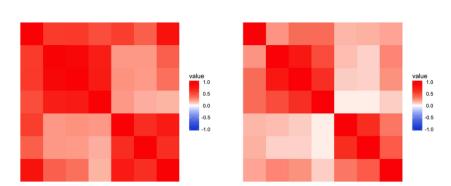




T1 - sub-cortical volumes L+R

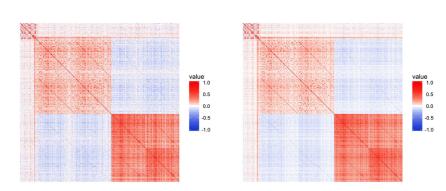


SWI T2* sub-cortical L+R



FreeSurfer

Genetic correlation



resting fMRI - ICA features

