Supplementary Figures

A Genome-Wide Association Study of Frailty Identifies Significant Genetic Correlation with Neuropsychiatric, Cardiovascular and Inflammation Pathways

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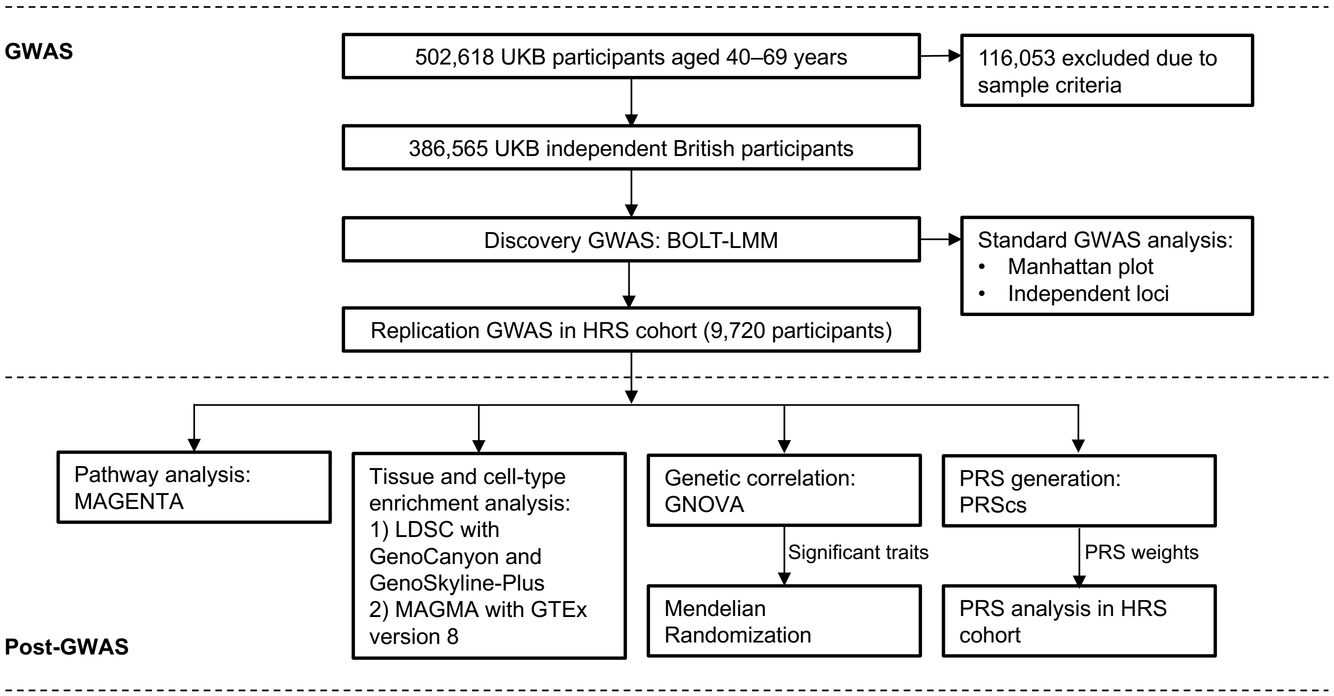
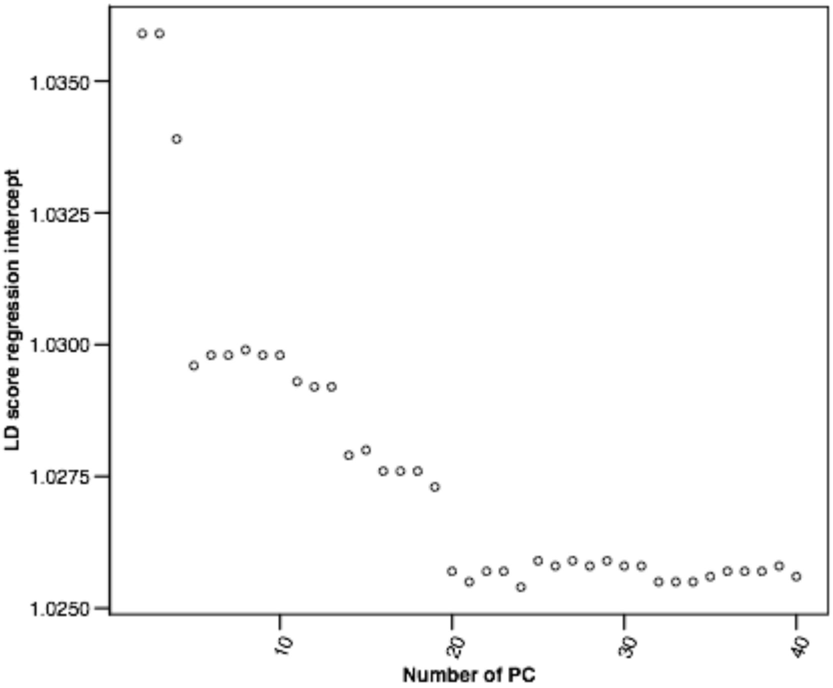
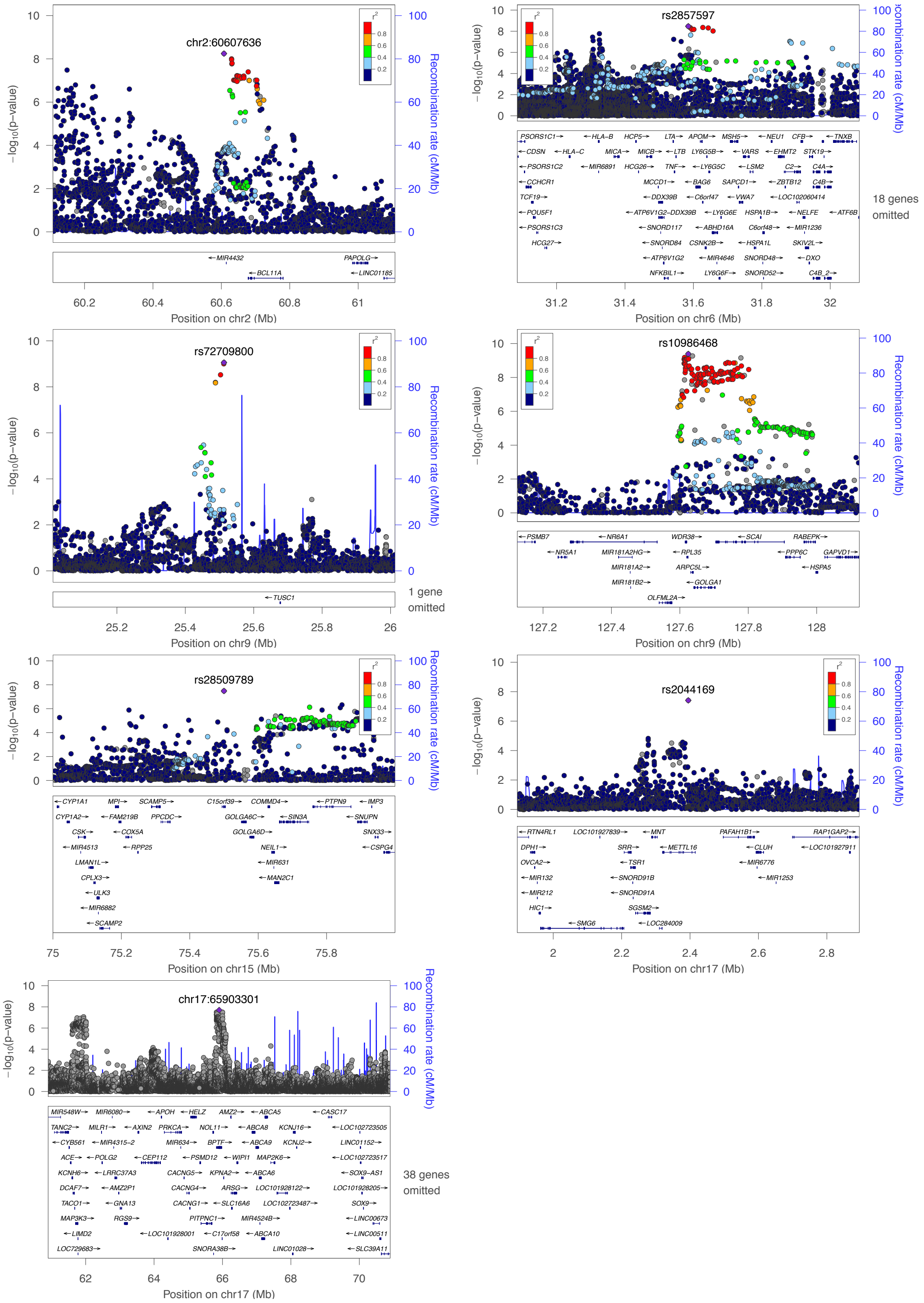
Supp. Figure 1. Workflow



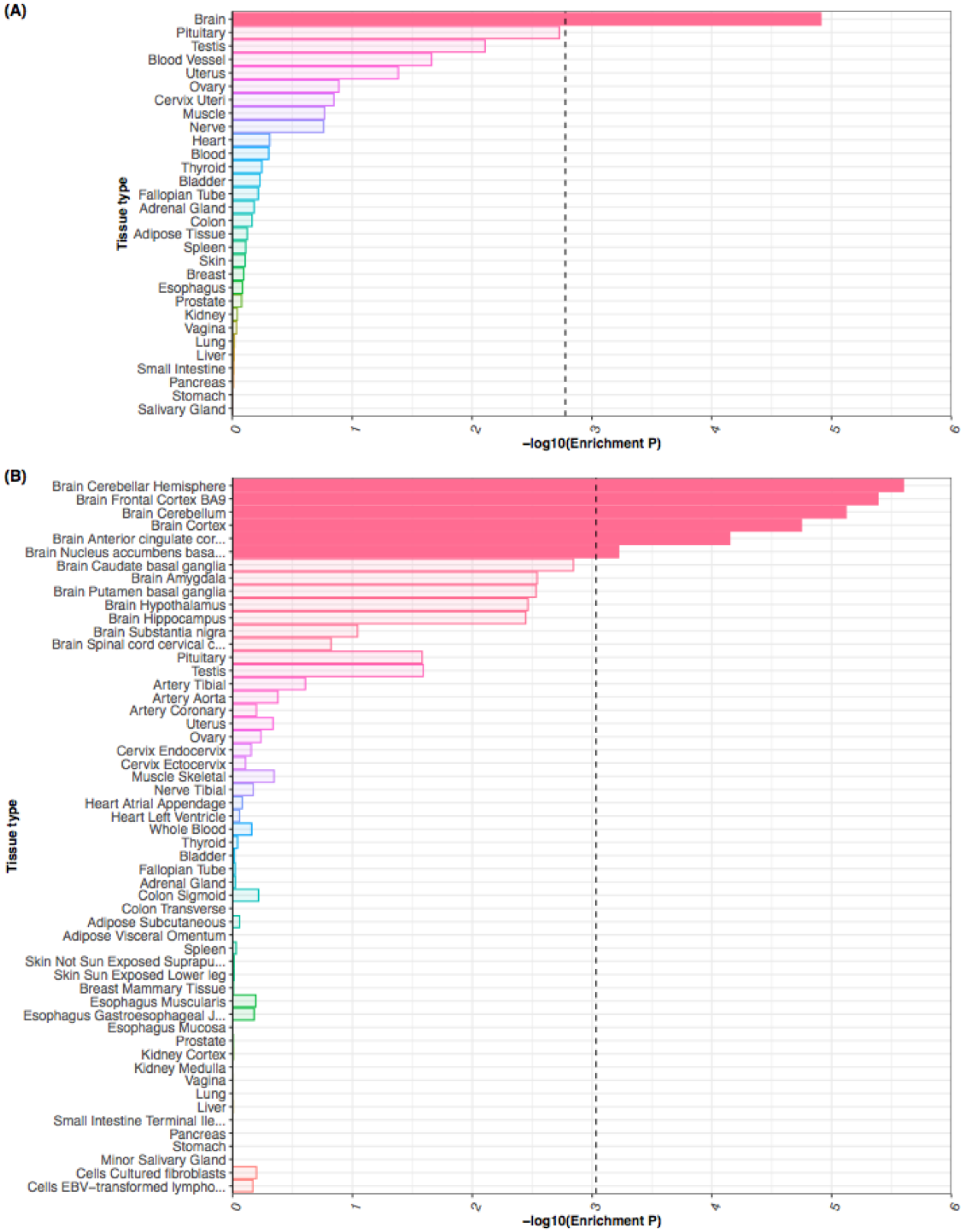
Supp. Figure 2. Selection of number of principal components in GWAS analysis



Supp. Figure 3. LocusZoom plot of novel genetic variants



Supp. Figure 4. MAGMA tissue enrichment results



(A) MAGMA tissue enrichment results based on 30 general tissues from GTEx v8 or (B) MAGMA tissue enrichment results based on 54 specific tissues from GTEx v8. The vertical dotted line indicates the Bonferroni-adjusted P-value for significant enrichment.