



Supplemental Figure S16. Correlations of GWAS signal enrichments ($-\log_{10}P$) for 45 complex traits from different methods and cut-offs. **A** Correlation between sum-based method and hypergeometric (count-based) test. **B** Correlations between different cut-offs of gene extension ($\pm 10\text{kb}$, $\pm 20\text{kb}$, $\pm 50\text{kb}$) and top percentages (3%, 5% and 10%) used for defining tissue-specific genes.