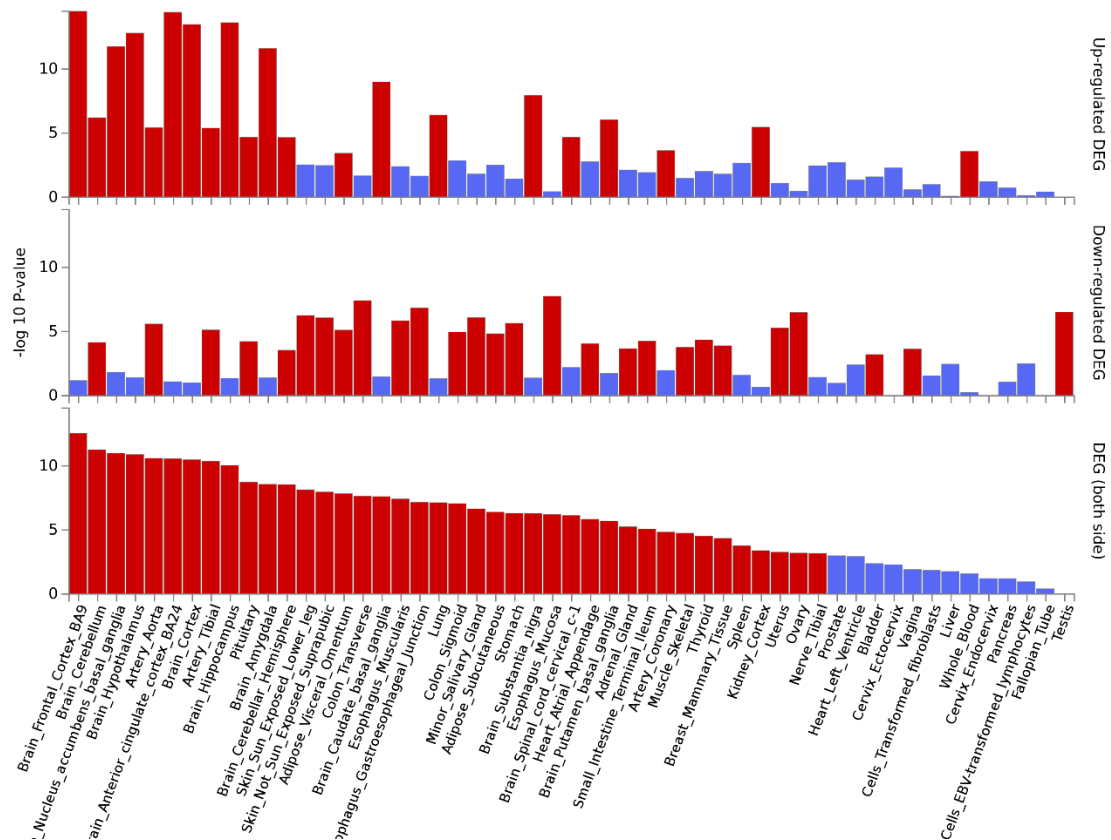
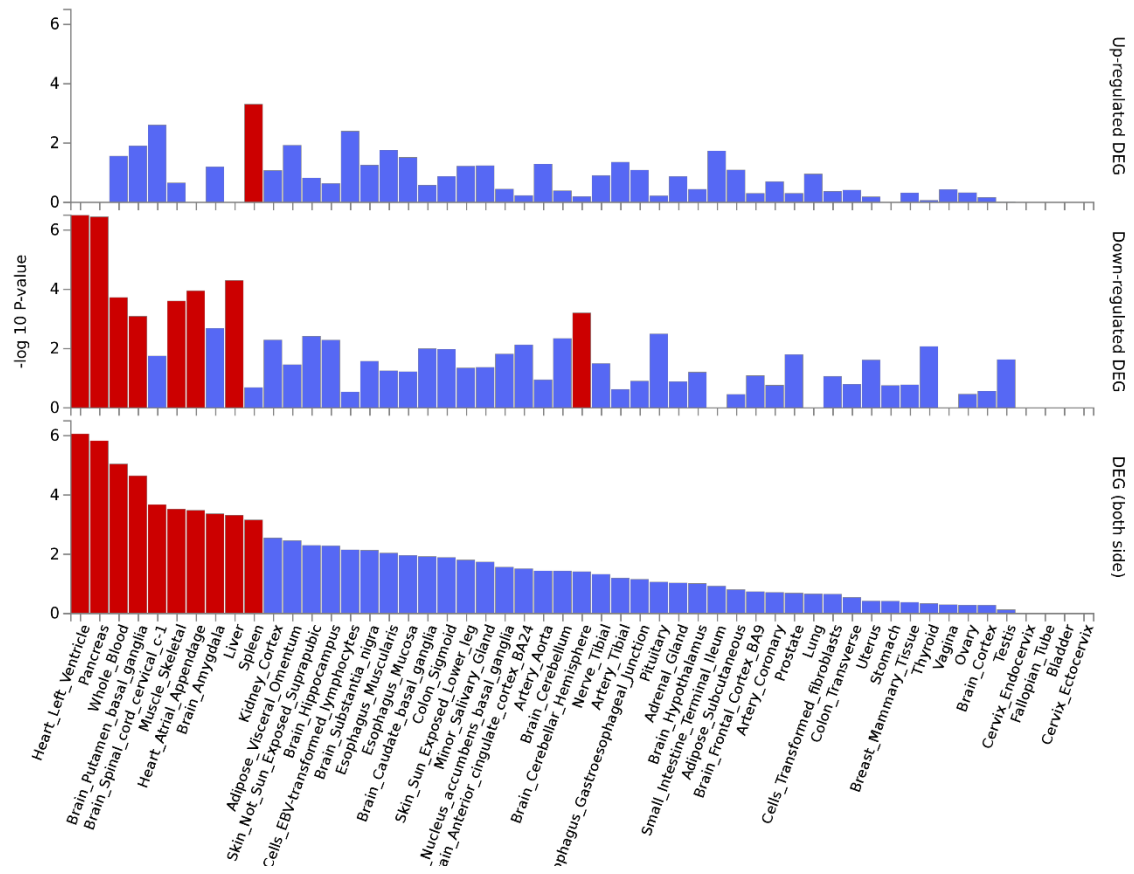


Supplementary Data 4. Tissue-specific patterns of expression for genes associated with Olink neurology proteins exhibiting genome-wide significant CpG sites. Differentially expressed gene sets in 53 tissue types (GTEx v7) with Bonferroni-corrected P values of < 0.05 and an absolute log fold change ≥ 0.58 are shown in red (default settings). Differentially expressed genes (DEG).

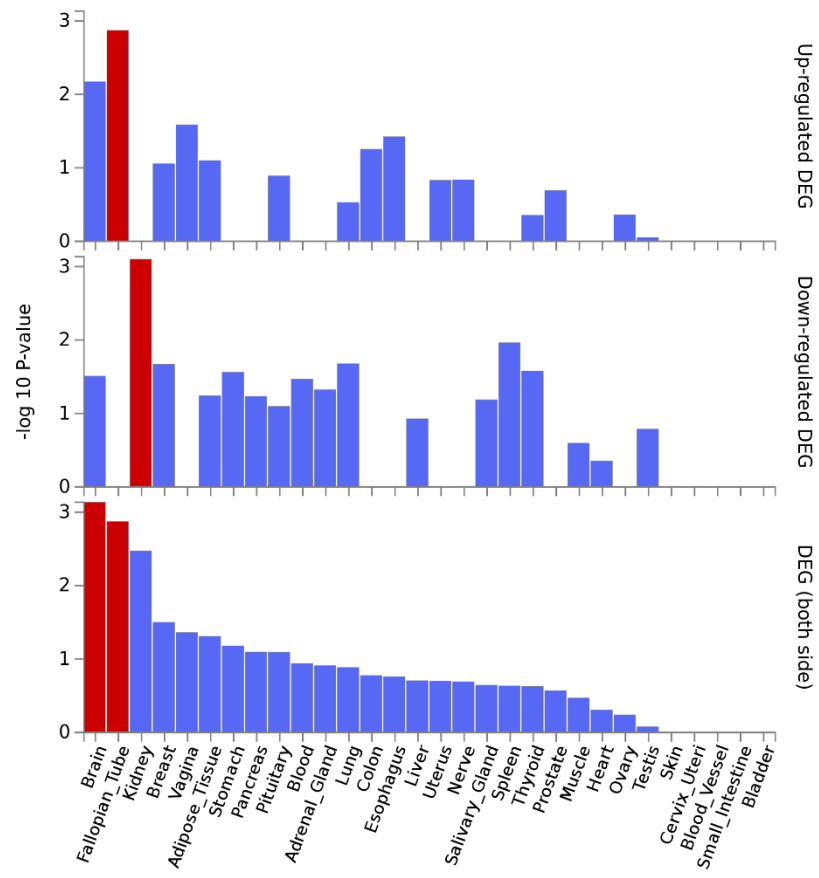
CRTAM



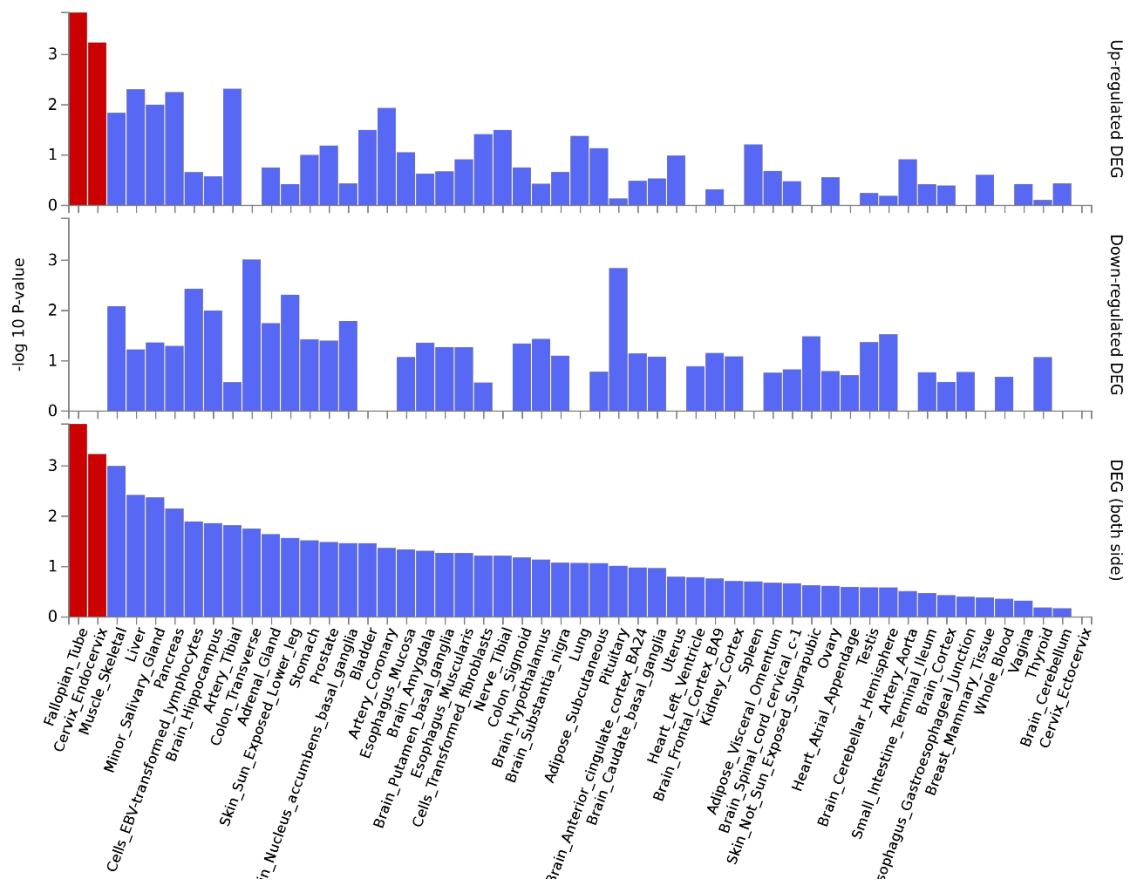
G-CSF



MDGA1



NEP



SIGLEC1

