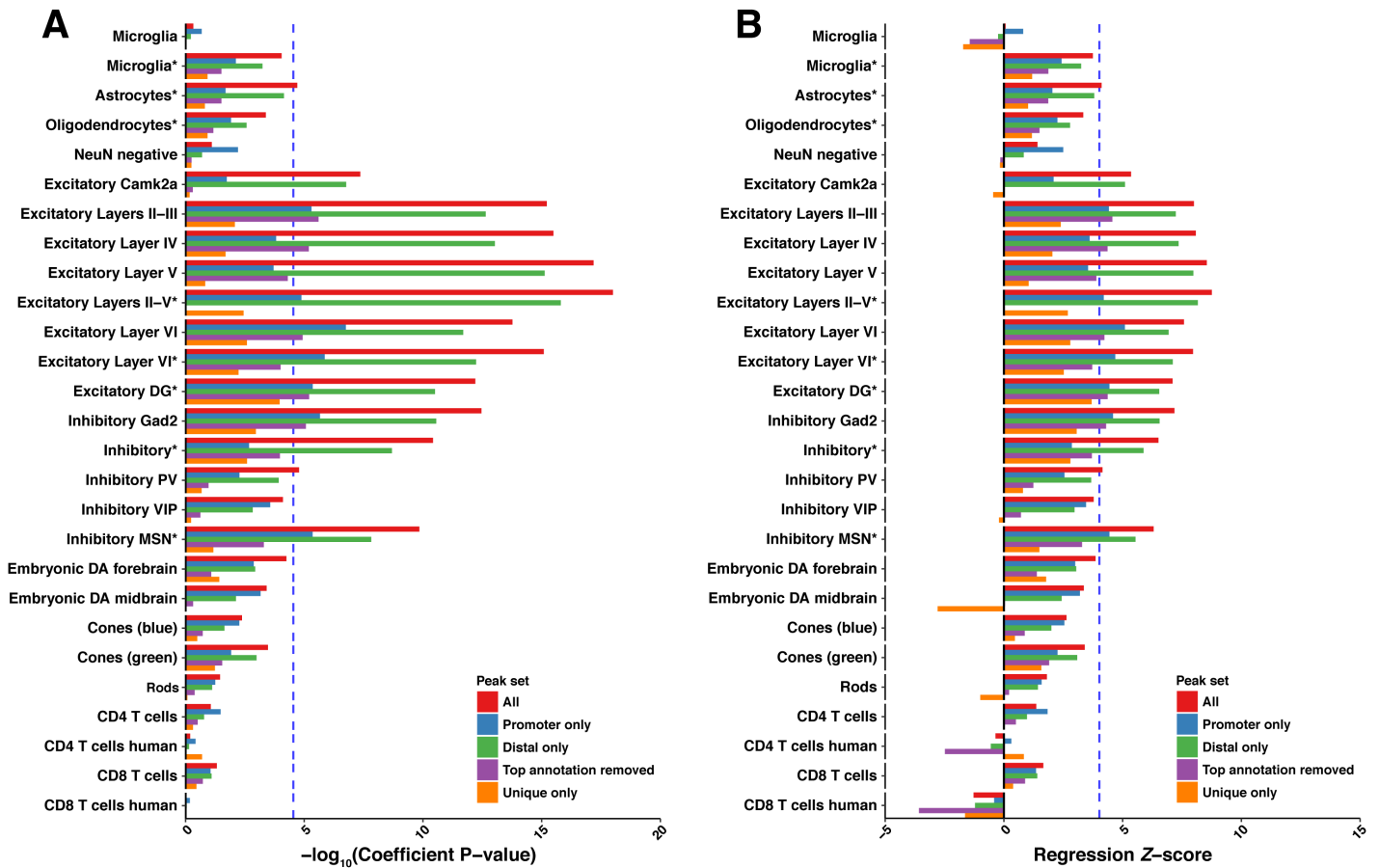


# Supplemental Figure S8



*Supplemental Figure S8.* CLOZUK SCZ S-LDSC results for all peaks, promoter peak sequences, distal peak sequences, peaks not in the top CLOZUK SCZ cell population, and peaks unique to each cell population. A) Barplots displaying  $-\log_{10}$  (heritability coefficient P-values). B) Barplots displaying regression coefficient Z-scores. Across trait significance level is shown ( $-\log_{10}(\text{coefficient p-value}) = 4.53857$  and equivalent Z-score = 4.02133; blue dashed line). Asterisks in the cell population name indicate single-nuclei ATAC-seq data. All results can be found in Supplemental Table 7.