Biosample: Progenitors Trait: Blood_counts Predictor ABC (A=scATAC, C=power law) In element (ABC) & distance to TSS scE2G (Multiome) scE2G (scATAC) 0 0.0 0.1 0.2 Recall (fraction of variants in predicted enhancers) 6017 variants from 9 traits

(6017 variant-biosample pairs)

Enrichment (GWAS variants vs. common variants)