

Enrichment (GWAS variants vs. common variants)

0.0

0.1

0.2

Recall (fraction of variants in predicted enhancers)  
6017 variants from 9 traits  
(6017 variant-biosample pairs)

Predictor

- ABC (A=scATAC, C=power law)
- In element (ABC) & distance to TSS
- scE2G (Multiome)
- scE2G (scATAC)

0

2

4

6