

Enrichment (GWAS variants vs. common variants)

0.0

0.1

0.2

0.3

0.4

0.5

Recall (fraction of variants in predicted enhancers)

6017 variants from 9 traits

(6017 variant-biosample pairs)

Predictor

- ABC (A=scATAC, C=power law)
- ArchR (high-confidence)
- Cicero (co-accessibility)
- DIRECTNET (high-confidence)
- FigR (p-value)
- In element (ABC) & distance to TSS
- scE2G (Multiome)
- scE2G (scATAC)
- SCENIC+ (importance)
- Signac (p-value)
- SnapATAC (p-value)