

Enrichment (GWAS variants vs. common variants)

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)
- SCENT (FDR)
- Signac (p-value)
- SnapATAC (p-value)
- STARE (gABC, power law)

Recall (fraction of variants in predicted enhancers)
8641 variants from 15 traits
(15362 variant-biosample pairs) fine-mapped variants

