

Enrichment (eQTLs vs. common variants)

Predictor

- ABC (A=scATAC, C=power law) (PBMC5\_NK)
- Cicero (co-accessibility) (PBMC5\_NK)
- FigR (p-value) (PBMC5\_NK)
- In element (ABC) & distance to TSS (PBMC5\_NK)
- scE2G (Multiome) (PBMC5\_NK)
- scE2G (scATAC) (PBMC5\_NK)
- SCENIC+ (importance) (PBMC5\_NK)
- SCENT (FDR) (PBMC5\_NK)
- Signac (p-value) (PBMC5\_NK)
- SnapATAC (p-value) (PBMC5\_NK)
- STARE (gABC, power law) (PBMC5\_NK)
- ArchR (high-confidence) (PBMC5\_NK)
- DIRECTNET (high-confidence) (PBMC5\_NK)

Recall (variants overlapping prediction linked to eGene)  
701 variants