

Enrichment (eQTLs vs. common variants) at recall 0.01

30  
20  
10  
0

Predictor (exact recall)

- scE2G (Multiome) (PBMC5\_Mono) (0.01)
- scE2G (scATAC) (PBMC5\_Mono) (0.01)
- STARE (gABC, power law) (BMMC5\_Myeloid) (0.01)
- STARE (gABC, power law) (PBMC5\_Mono) (0.01)
- SnapATAC (p-value) (PBMC5\_Mono) (0.009)
- SCENIC+ (importance) (PBMC5\_Mono) (0.01)
- Signac (p-value) (PBMC5\_Mono) (0.009)
- Signac (p-value) (BMMC5\_Myeloid) (0.01)
- scE2G (scATAC) (BMMC5\_Myeloid) (0.01)
- SCENIC+ (importance) (BMMC5\_Myeloid) (0.011)
- Cicero (co-accessibility) (PBMC5\_Mono) (0.01)
- FigR (p-value) (BMMC5\_Myeloid) (0.01)
- Cicero (co-accessibility) (BMMC5\_Myeloid) (0.01)

