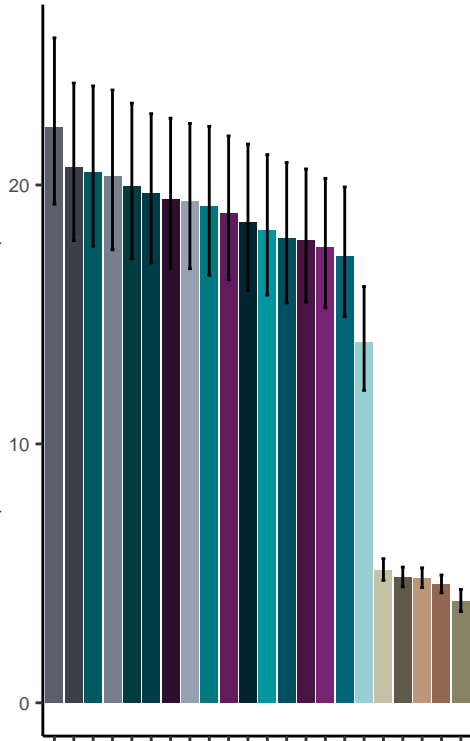Enrichment (eQTLs vs. common variants) at recall 0.05

Predictor (exact recall)

- In element (ABC) & distance to TSS (BMMC22_Normoblast) (0.051)
- In element (ABC) & distance to TSS (BMMC22_Proerythroblast) (0.052)
- ABC (A=scATAC, C=power law) (BMMC22_Normoblast) (0.05)
- In element (ABC) & distance to TSS (BMMC22_Erythroblast) (0.049)
- ABC (A=scATAC, C=power law) (BMMC22_Proerythroblast) (0.05)
- scE2G (scATAC) (BMMC22_Normoblast) (0.05)
- scE2G (Multiome) (BMMC22_Proerythroblast) (0.049)
- In element (ABC) & distance to TSS (K562_Xu) (0.053)
- ABC (A=scATAC, C=power law) (BMMC22_Erythroblast) (0.051)
- scE2G (Multiome) (BMMC22_Erythroblast) (0.051)
- scE2G (scATAC) (BMMC22_Proerythroblast) (0.049)

- ABC (A=scATAC, C=power law) (K562_Xu) (0.051)
- scE2G (scATAC) (BMMC22_Erythroblast) (0.05)
- scE2G (Multiome) (BMMC22_Normoblast) (0.05)
- scE2G (Multiome) (K562_Xu) (0.05)
- scE2G (scATAC) (K562_Xu) (0.052)
- STARE (gABC, power law) (K562_Xu) (0.05)
- SCENIC+ (importance) (K562_Xu) (0.05)
- SnapATAC (p−value) (K562_Xu) (0.049)
- FigR (p−value) (K562_Xu) (0.049)
- Signac (p−value) (K562_Xu) (0.05)
- Cicero (co−accessibility) (K562_Xu) (0.05)