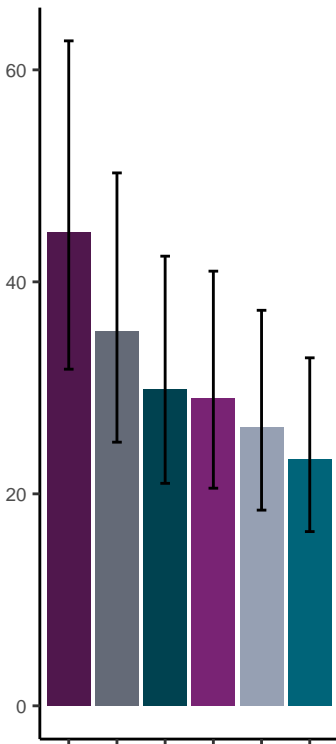


Enrichment (eQTLs vs. common variants) at recall 0.03



Predictor (exact recall)

- scE2G (Multiome) (BMMC22_ID2_hi_myeloid_prog) (0.031)
- In element (ABC) & distance to TSS (BMMC22_ID2_hi_myeloid_prog) (0.031)
- scE2G (scATAC) (BMMC22_ID2_hi_myeloid_prog) (0.03)
- scE2G (Multiome) (BMMC22_G_M_prog) (0.031)
- In element (ABC) & distance to TSS (BMMC22_G_M_prog) (0.029)
- scE2G (scATAC) (BMMC22_G_M_prog) (0.031)