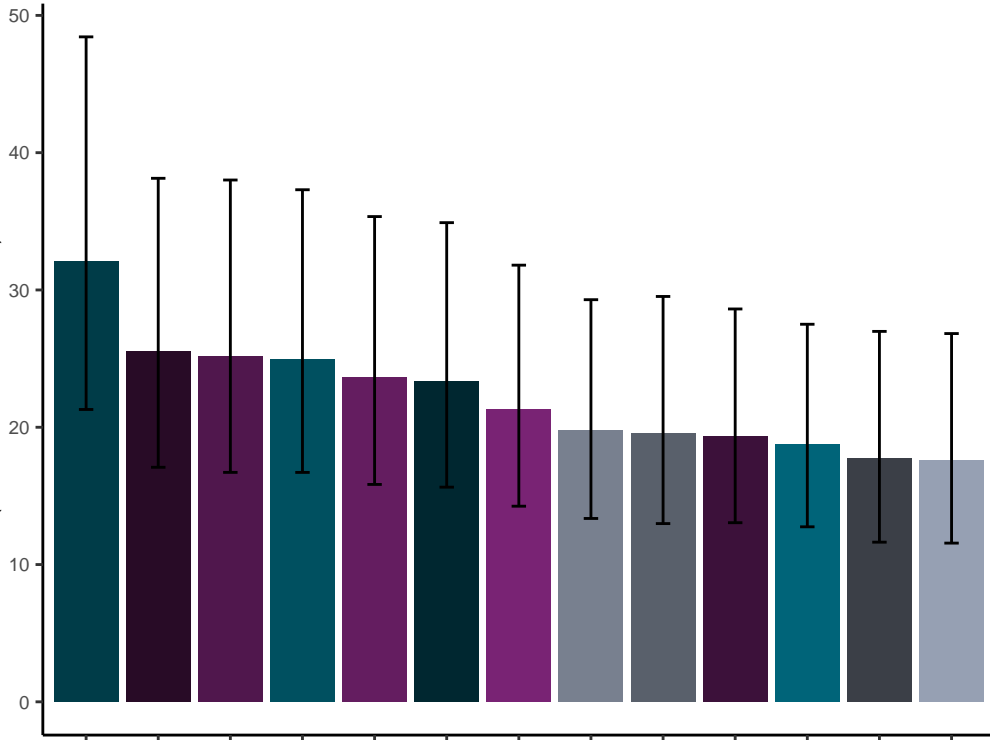


Enrichment (eQTLs vs. common variants) at recall 0.03



Predictor (exact recall)

- scE2G (scATAC) (BMMC22\_ID2\_hi\_myeloid\_prog) (0.03)
- scE2G (Multiome) (BMMC22\_MK\_E\_prog) (0.03)
- scE2G (Multiome) (BMMC22\_ID2\_hi\_myeloid\_prog) (0.03)
- scE2G (scATAC) (BMMC22\_HSC) (0.03)
- scE2G (Multiome) (BMMC22\_HSC) (0.03)
- scE2G (scATAC) (BMMC22\_MK\_E\_prog) (0.03)
- scE2G (Multiome) (BMMC22\_G\_M\_prog) (0.027)
- In element (ABC) & distance to TSS (BMMC22\_HSC) (0.032)
- In element (ABC) & distance to TSS (BMMC22\_Lymph\_prog) (0.03)
- scE2G (Multiome) (BMMC22\_Lymph\_prog) (0.032)
- scE2G (scATAC) (BMMC22\_G\_M\_prog) (0.03)
- In element (ABC) & distance to TSS (BMMC22\_MK\_E\_prog) (0.029)
- In element (ABC) & distance to TSS (BMMC22\_G\_M\_prog) (0.027)