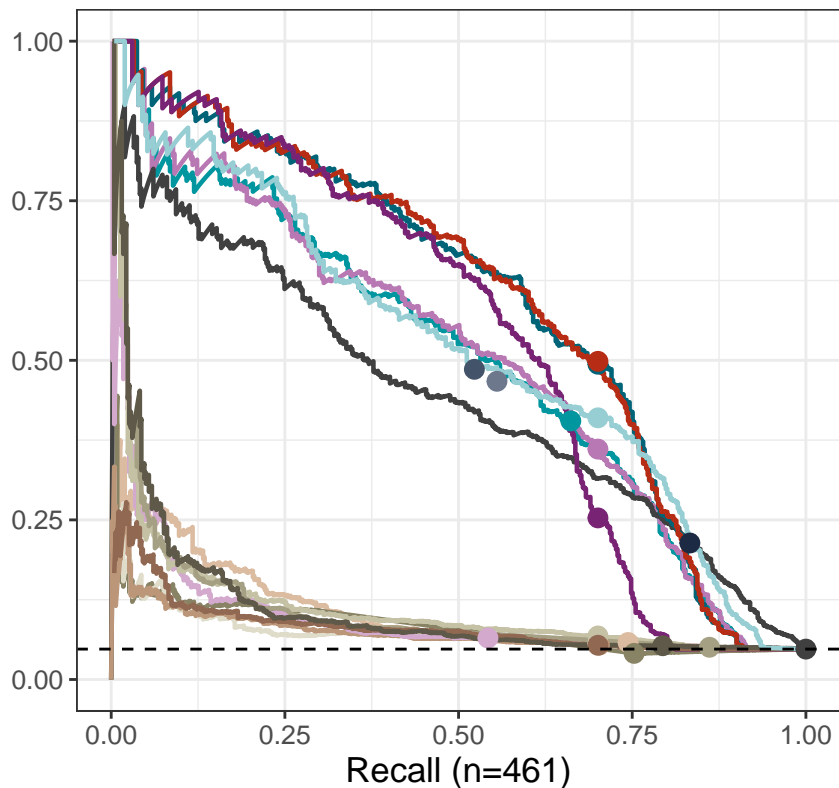sc.250425.K562.IGVF.ignore_TPM (9686 pairs)

Legend:
- scE2G (Multiome)
- scE2G (Multiome, ignore_TPM)
- scE2G (ATAC)
- ARC–E2G score
- Kendall correlation
- ABC (A=scATAC, C=power law)
- STARE (gABC, power law)
- SnapATAC (z–score)
- Signac (p–value)
- Cicero (co–accessibility)
- FigR (p–value)
- SCENT (bootstrap p–value, –log10, signed)
- SCENIC+ (importance)
- DIRECTNET (importance)
- ArchR (Pearson correlation)
- Distance to TSS
- Closest TSS
- Closest gene
- Within 100kb of TSS