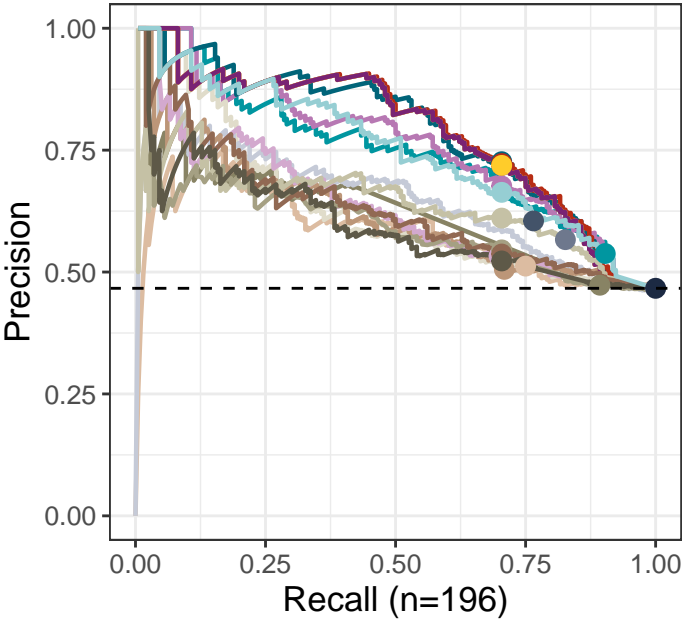
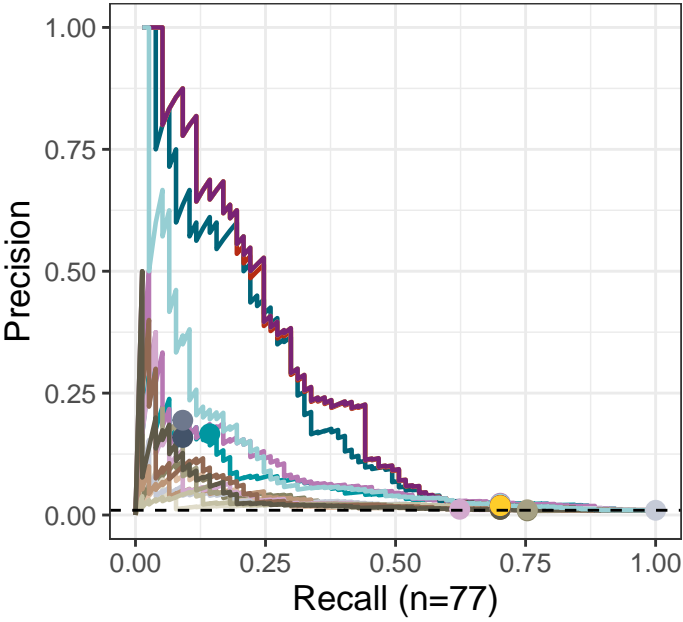


distToTSS (bins)

distToTSS (bins) = [0,20) (420 pairs)



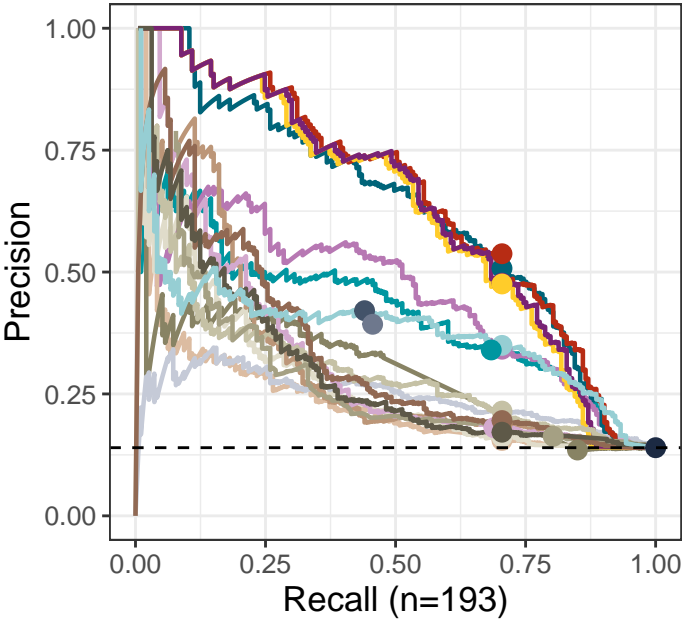
distToTSS (bins) = [100,2.5e+03) (7899 pairs)



- Predictor
- scE2G (Multiome,TPM>1)
 - scE2G (Multiome,qnorm_TPM>1)
 - scE2G (Multiome,ignoreTPM)
 - 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)

- Predictor
- scE2G (Multiome,TPM>1)
 - scE2G (Multiome,qnorm_TPM>1)
 - scE2G (Multiome,ignoreTPM)
 - 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

distToTSS (bins) = [20,100) (1381 pairs)



- Predictor
- scE2G (Multiome,TPM>1)
 - scE2G (Multiome,qnorm_TPM>1)
 - scE2G (Multiome,ignoreTPM)
 - 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