distToTSS (bins)

1.00 -

0.75

0.25

0.00

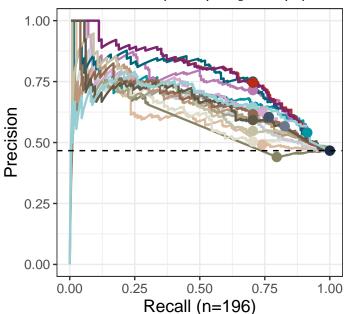
0.00

Precision

Predictor

- scE2G (Multiome)
- scE2G (Multiome,ignoreTPM)

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

Predictor

scE2G (Multiome)

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

0.50

Recall (n=77)

0.25

0.75

1.00

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

Predictor

- scE2G (Multiome)
- scE2G (Multiome,ignoreTPM)

distToTSS (bins) = [20,100) (1381-paies) (ATAC)

0.75

1.00

0.50

Recall (n=193)

0.25

1.00 -

0.75

Precision 0.50

0.25

0.00

0.00

- 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