Predictor

- ABC (A=scATAC, C=power law) (BMMC5_Myeloid)
- ABC (A=scATAC, C=power law) (PBMC5_Mono)
- Cicero (co-accessibility) (BMMC5_Myeloid)
- Cicero (co-accessibility) (PBMC5_Mono)
- FigR (p-value) (BMMC5_Myeloid)
- FigR (p-value) (PBMC5_Mono)
- In element (ABC) & distance to TSS (BMMC5_Myeloid)
- In element (ABC) & distance to TSS (PBMC5_Mono)
- scE2G (Multiome) (BMMC5_Myeloid)
- scE2G (Multiome) (PBMC5_Mono)
- scE2G (scATAC) (BMMC5_Myeloid)
- scE2G (scATAC) (PBMC5_Mono)
- SCENIC+ (importance) (BMMC5_Myeloid)
- SCENIC+ (importance) (PBMC5_Mono)
- SCENT (FDR) (BMMC5_Myeloid)
- SCENT (FDR) (PBMC5_Mono)
- Signac (p-value) (BMMC5_Myeloid)
- Signac (p-value) (PBMC5_Mono)
- SnapATAC (p-value) (BMMC5_Myeloid)
- SnapATAC (p-value) (PBMC5_Mono)
- STARE (gABC, power law) (BMMC5_Myeloid)
- STARE (gABC, power law) (PBMC5_Mono)
- ArchR (high-confidence) (BMMC5_Myeloid)
- ArchR (high-confidence) (PBMC5_Mono)
- DIRECTNET (high-confidence) (BMMC5_Myeloid)
- DIRECTNET (high-confidence) (PBMC5_Mono)