Predictor

- ABC (A=scATAC, C=power law) (BMMC22_Erythroblast)
- ABC (A=scATAC, C=power law) (BMMC22_Normoblast)
- ABC (A=scATAC, C=power law) (BMMC22_Proerythroblast)
- ABC (A=scATAC, C=power law) (K562_Xu)
- Cicero (co-accessibility) (K562_Xu)
- FigR (p-value) (BMMC22_Erythroblast)
- FigR (p-value) (BMMC22_Normoblast)
- FigR (p-value) (BMMC22_Proerythroblast)
- FigR (p-value) (K562_Xu)
- In element (ABC) & distance to TSS (BMMC22_Erythroblast)
- In element (ABC) & distance to TSS (BMMC22_Normoblast)
- In element (ABC) & distance to TSS (BMMC22_Proerythroblast)

- In element (ABC) & distance to TSS (K562_Xu)
- scE2G (Multiome) (BMMC22_Erythroblast)
- scE2G (Multiome) (BMMC22_Normoblast)
- scE2G (Multiome) (BMMC22_Proerythroblast)
- scE2G (Multiome) (K562_Xu)
- scE2G (scATAC) (BMMC22_Erythroblast)
- scE2G (scATAC) (BMMC22_Normoblast)
- scE2G (scATAC) (BMMC22_Proerythroblast)
- scE2G (scATAC) (K562_Xu)
- SCENIC+ (importance) (BMMC22_Erythroblast)
- SCENIC+ (importance) (BMMC22_Normoblast)
- SCENIC+ (importance) (BMMC22_Proerythroblast)

- SCENIC+ (importance) (K562_Xu)
- SCENT (FDR) (K562_Xu)
- Signac (p-value) (BMMC22_Erythroblast)
- Signac (p-value) (BMMC22_Normoblast)
- Signac (p-value) (BMMC22_Proerythroblast)
- Signac (p-value) (K562_Xu)
- SnapATAC (p-value) (BMMC22_Erythroblast)
- SnapATAC (p-value) (BMMC22_Normoblast)
- SnapATAC (p-value) (BMMC22_Proerythroblast)
- SnapATAC (p-value) (K562_Xu)
- STARE (gABC, power law) (K562_Xu)
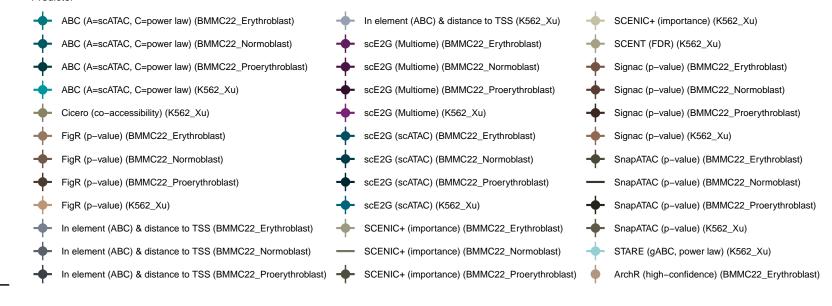- ArchR (high-confidence) (BMMC22_Erythroblast)

- ArchR (high-confidence) (BMMC22_Proerythroblast)
- ArchR (high-confidence) (K562_Xu)
- DIRECTNET (high-confidence) (BMMC22_Erythroblast)
- DIRECTNET (high-confidence) (BMMC22_Normoblast)
- DIRECTNET (high-confidence) (BMMC22_Proerythroblast)
- DIRECTNET (high-confidence) (K562_Xu)