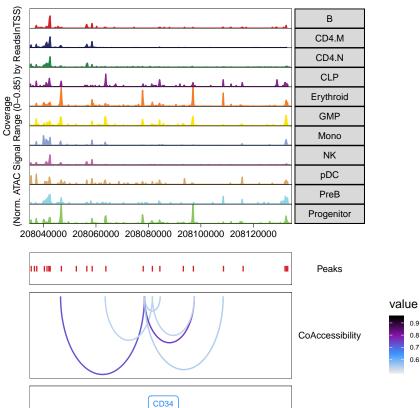
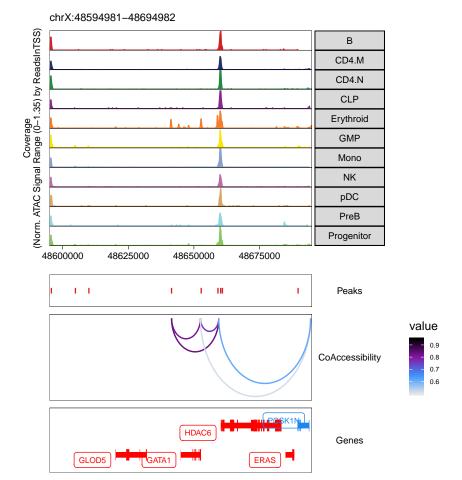
chr1:208034682-208134683

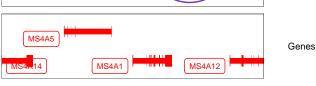


Genes



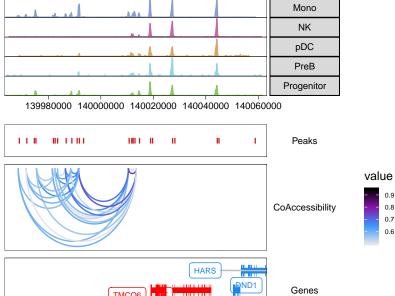
chr11:60173281-60273282 В CD4.M CD4.N CLP Erythroid GMP Mono NK pDC PreB Progenitor 60175000 60200000 60225000 60250000 Peaks ш $\mathbf{I}\mathbf{I}\mathbf{I}$ value 0.9 CoAccessibility 0.8 0.7 0.6

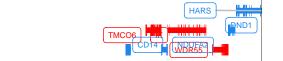
Coverage (Norm. ATAC Signal Range (0–1.03) by ReadsInTSS)



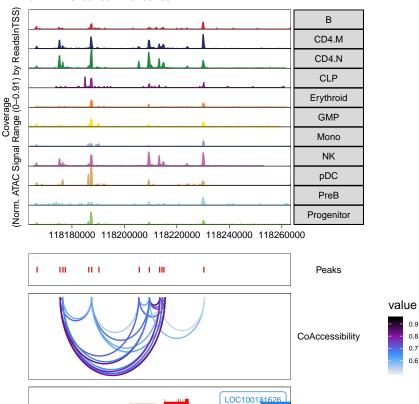
chr5:139963285-140063286 В CD4.M CD4.N CLP Erythroid GMP Mono NK pDC PreB Progenitor 139980000 140000000 140020000 140040000 140060000 1.1.11 III 11 II I IIII I П Peaks ı

Coverage (Norm. ATAC Signal Range (0-1.31) by ReadsInTSS)





chr11:118163458-118263459



Genes

CD3G

CD3E

CD3D

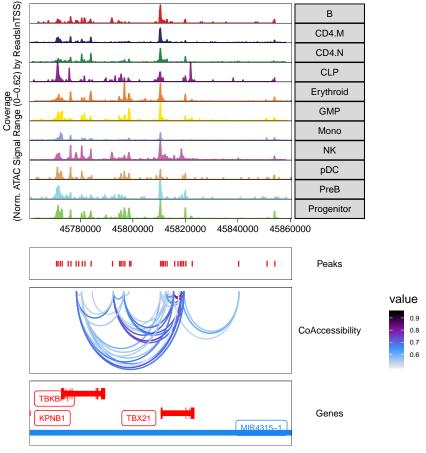
UBE4A

chr2:86985518-87085519 В CD4.M CD4.N CLP Erythroid GMP Mono NK pDC PreB Progenitor 87000000 87020000 87040000 87060000 87080000 Peaks 10 000 0 ш ш ш value 0.9 CoAccessibility 0.8 0.7 0.6

Genes

CD8A

Coverage (Norm. ATAC Signal Range (0-0.81) by ReadsInTSS)



chr5:35806976-35906977 В CD4.M CD4.N CLP Erythroid GMP Mono NK pDC PreB Progenitor 35825000 35850000 35875000 35900000 Peaks 10.11 value 0.9 CoAccessibility 0.8 0.7 0.6 CAPSL Genes

IL7R

Coverage (Norm. ATAC Signal Range (0–0.89) by ReadsInTSS)