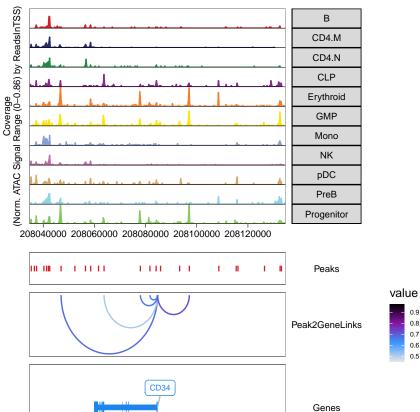
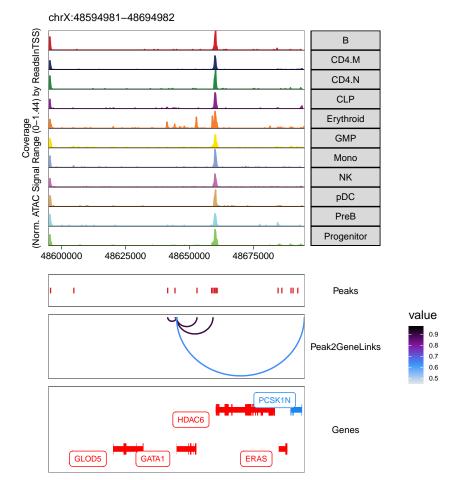
chr1:208034682-208134683





chr11:60173281-60273282 В CD4.M CD4.N CLP Erythroid **GMP** Mono NK pDC PreB Progenitor 60175000 60200000 60225000 60250000 111 Peaks ш 1.1 value 0.9 Peak2GeneLinks 8.0 0.7 0.6 0.5 MS4A5

MS4A12

Genes

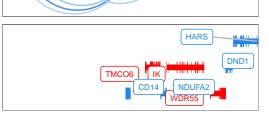
Coverage (Norm. ATAC Signal Range (0-1.4) by ReadsInTSS)

MS4A14

MS4A1

chr5:139963285-140063286 В CD4.M CD4.N CLP Erythroid GMP Mono NK pDC PreB Progenitor 139980000 140000000 140020000 140040000 140060000 Peaks III 1111 101 0 ī 11 1 value 0.9 Peak2GeneLinks 8.0 0.7 0.6 0.5

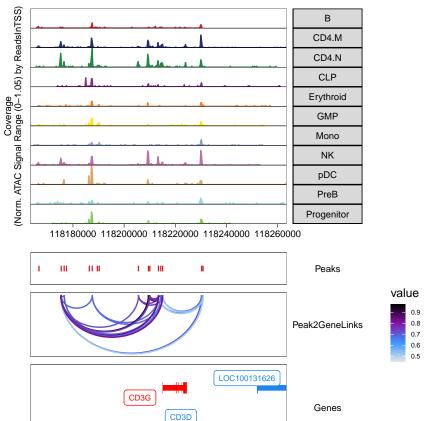
Genes



Coverage (Norm. ATAC Signal Range (0-1.38) by ReadsInTSS)

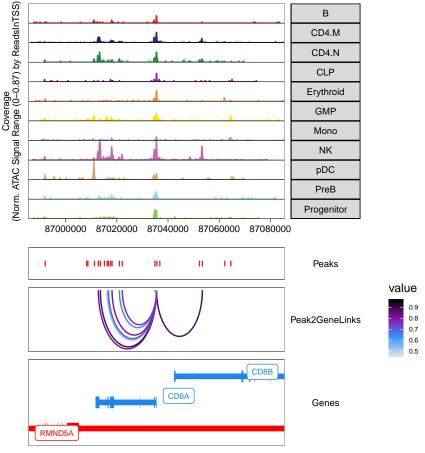
chr11:118163458-118263459

CD3E



UBE4A

chr2:86985518-87085519 В CD4.M CD4.N CLP Erythroid GMP Mono NK pDC PreB Progenitor 87000000 87020000 87040000 87060000 87080000 0.10100.0 11 Peaks Ш П value 0.9 Peak2GeneLinks 8.0 0.7 0.6



MIR4315-1

Coverage (Norm. ATAC Signal Range (0-0.65) by ReadsInTSS)

chr5:35806976-35906977 Coverage (Norm. ATAC Signal Range (0-0.98) by ReadsInTSS) В CD4.M CD4.N CLP Erythroid GMP Mono NK pDC PreB Progenitor 35825000 35850000 35875000 35900000 Peaks 11 11 10.01 value 0.9 Peak2GeneLinks 8.0 0.7 0.6 0.5 CAPSL Genes IL7R