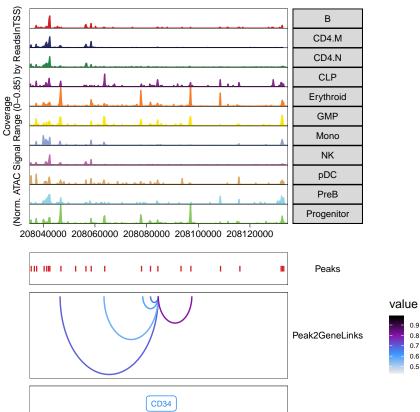
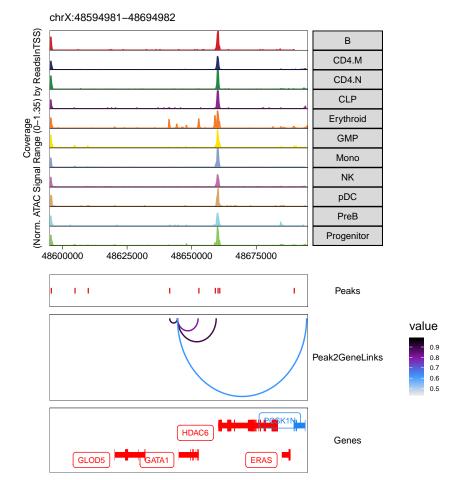
### chr1:208034682-208134683



Genes



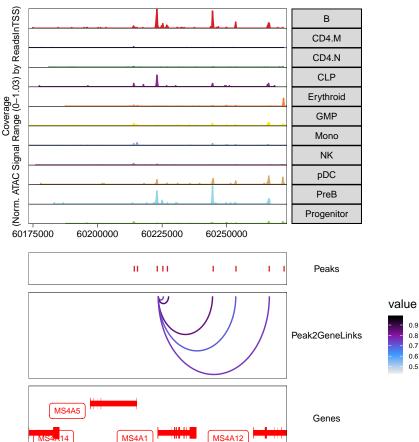
chr9:36984475-37084476 В CD4.M CD4.N CLP Erythroid GMP Mono NK pDC PreB Progenitor 37000000 37020000 37040000 37060000 37080000 Peaks 100111 1 0001 II = IП value 0.9 Peak2GeneLinks 0.8 0.7 0.6 0.5 EBLN3P

PAX5

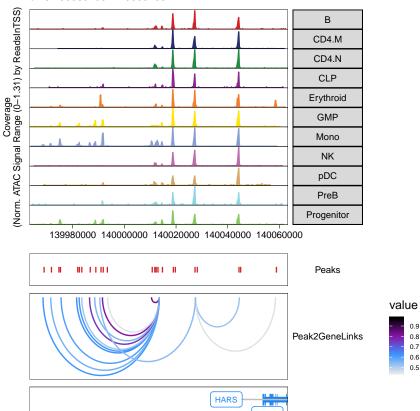
Genes

Coverage (Norm. ATAC Signal Range (0–1.29) by ReadsInTSS)

### chr11:60173281-60273282

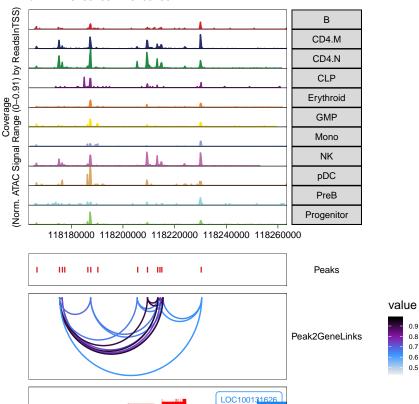


#### chr5:139963285-140063286



Genes

#### chr11:118163458-118263459



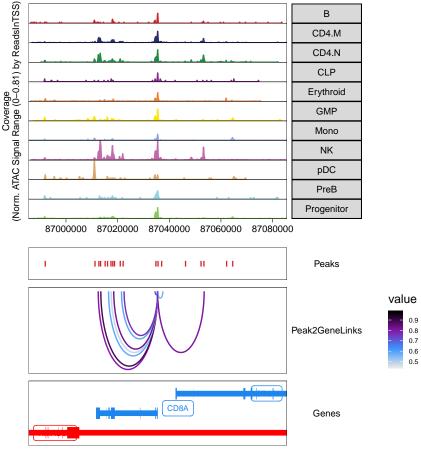
Genes

CD3G

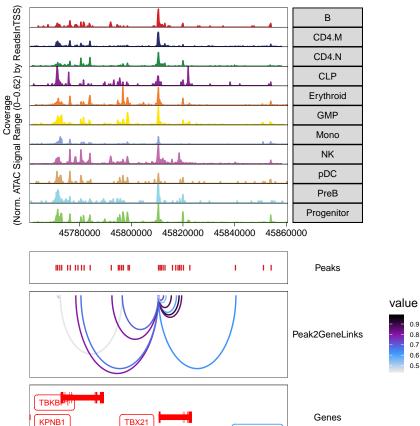
CD3E

CD3D

## chr2:86985518-87085519 В CD4.M CD4.N CLP Erythroid GMP Mono NK pDC PreB Progenitor 87000000 87020000 87040000 87060000 87080000 Peaks 10 00 0 ш П П value 0.9 Peak2GeneLinks 0.8 0.7



# chr17:45760609-45860610



MIR4315-1

chr5:35806976-35906977 Coverage (Norm. ATAC Signal Range (0–0.89) by ReadsInTSS) В CD4.M CD4.N CLP Erythroid GMP Mono NK pDC PreB Progenitor 35825000 35850000 35875000 35900000 Peaks ш 10.1.1 1 1 value 0.9 Peak2GeneLinks 0.8 0.7 0.6 0.5 CAPSL Genes

IL7R