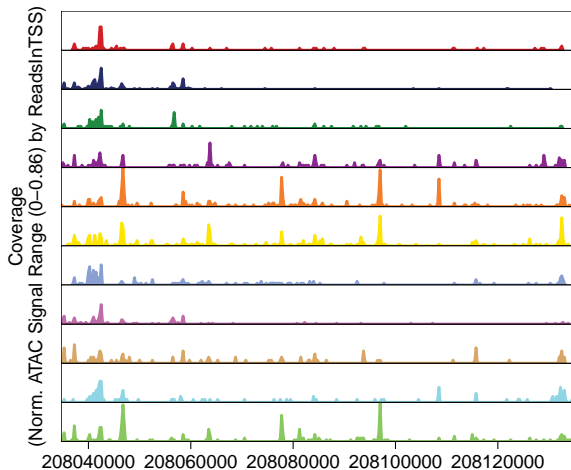


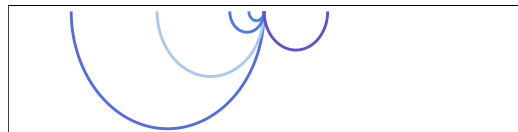
chr1:208034682–208134683



B
CD4.M
CD4.N
CLP
Erythroid
GMP
Mono
NK
pDC
PreB
Progenitor

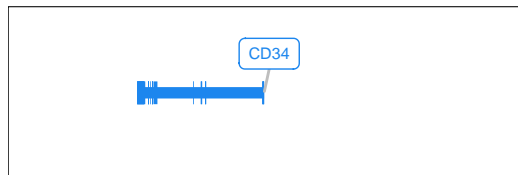
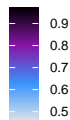


Peaks



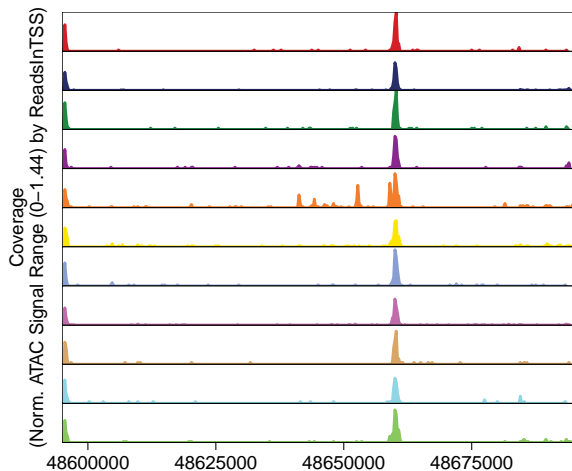
Peak2GeneLinks

value



Genes

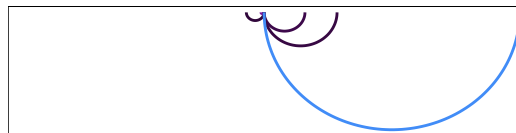
chrX:48594981-48694982



B
CD4.M
CD4.N
CLP
Erythroid
GMP
Mono
NK
pDC
PreB
Progenitor

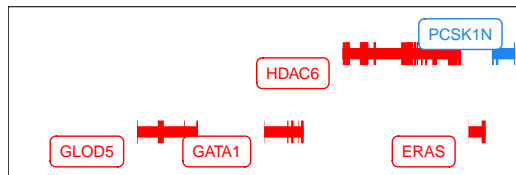
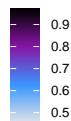


Peaks



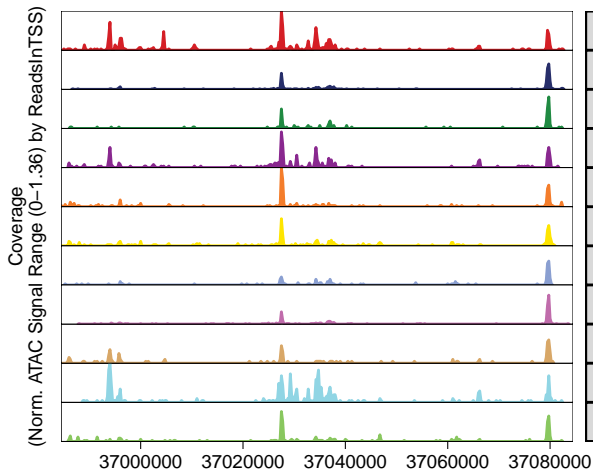
Peak2GeneLinks

value



Genes

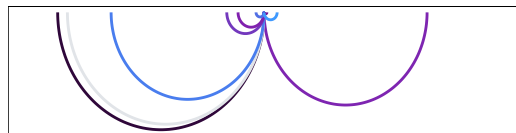
chr9:36984475–37084476



B
CD4.M
CD4.N
CLP
Erythroid
GMP
Mono
NK
pDC
PreB
Progenitor

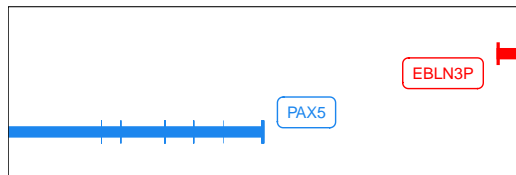
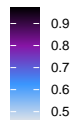


Peaks



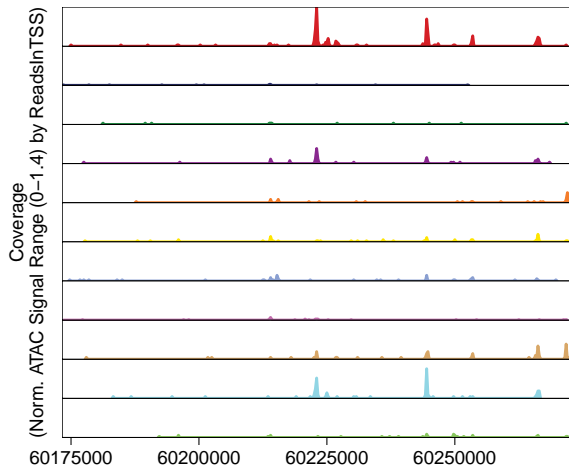
Peak2GeneLinks

value



Genes

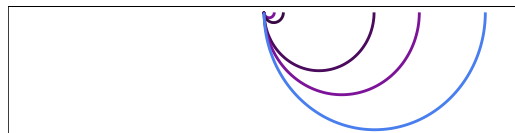
chr11:60173281-60273282



B
CD4.M
CD4.N
CLP
Erythroid
GMP
Mono
NK
pDC
PreB
Progenitor

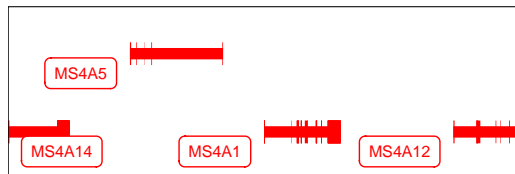
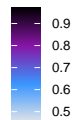


Peaks



Peak2GeneLinks

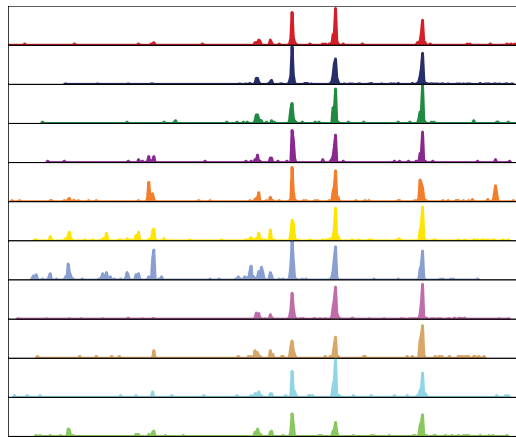
value



Genes

chr5:139963285–140063286

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



139980000 140000000 140020000 140040000 140060000

B
CD4.M
CD4.N
CLP
Erythroid
GMP
Mono
NK
pDC
PreB
Progenitor

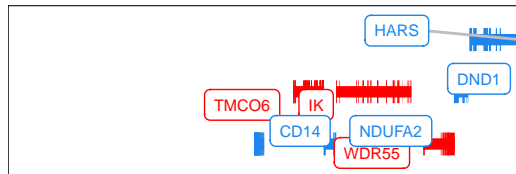
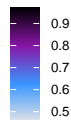


Peaks



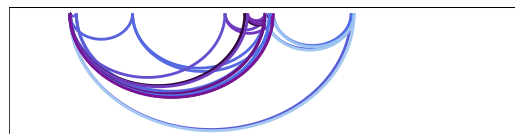
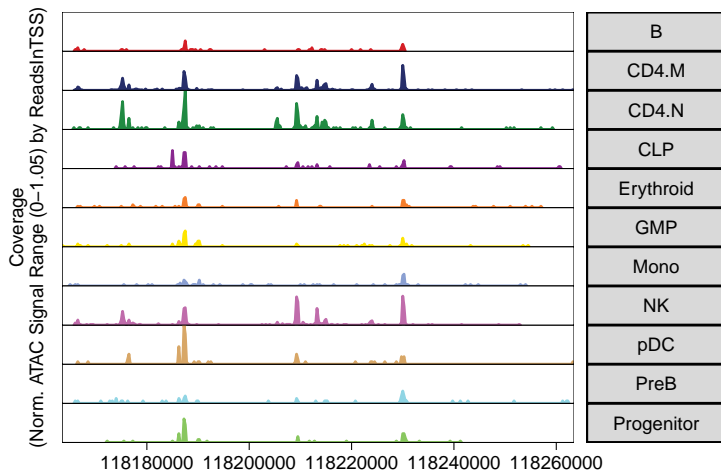
Peak2GeneLinks

value

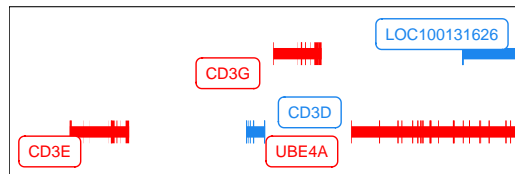
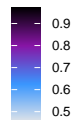


Genes

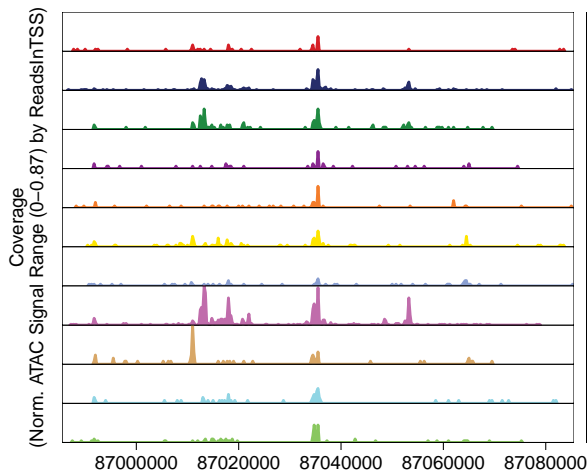
chr11:118163458–118263459



value



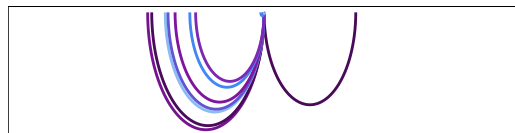
chr2:86985518–87085519



B
CD4.M
CD4.N
CLP
Erythroid
GMP
Mono
NK
pDC
PreB
Progenitor

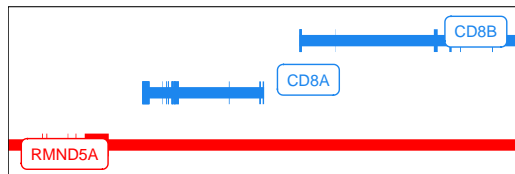
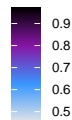


Peaks



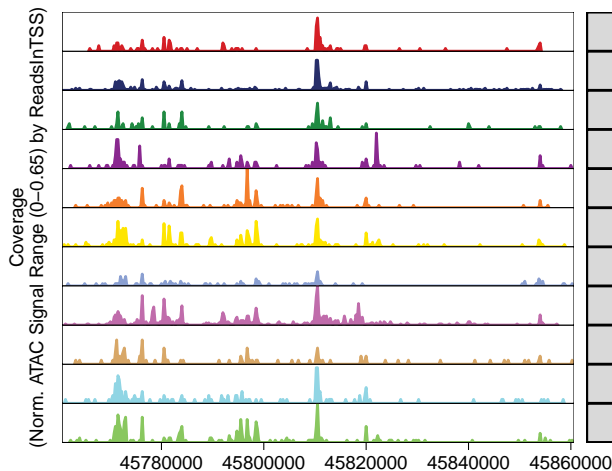
Peak2GeneLinks

value



Genes

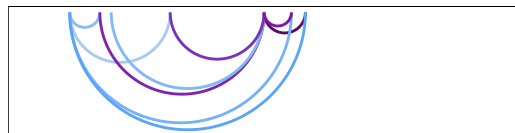
chr17:45760609–45860610



B
CD4.M
CD4.N
CLP
Erythroid
GMP
Mono
NK
pDC
PreB
Progenitor

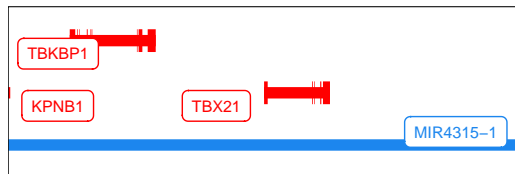
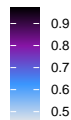


Peaks



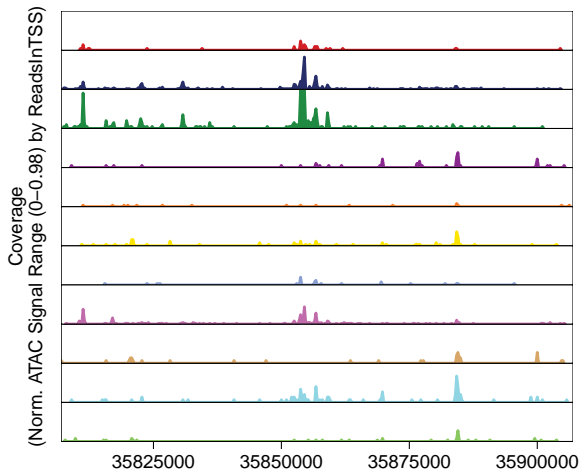
Peak2GeneLinks

value



Genes

chr5:35806976–35906977



B
CD4.M
CD4.N
CLP
Erythroid
GMP
Mono
NK
pDC
PreB
Progenitor

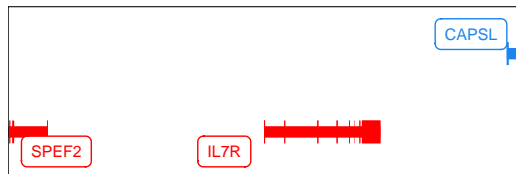
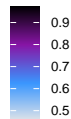


Peaks



Peak2GeneLinks

value



Genes