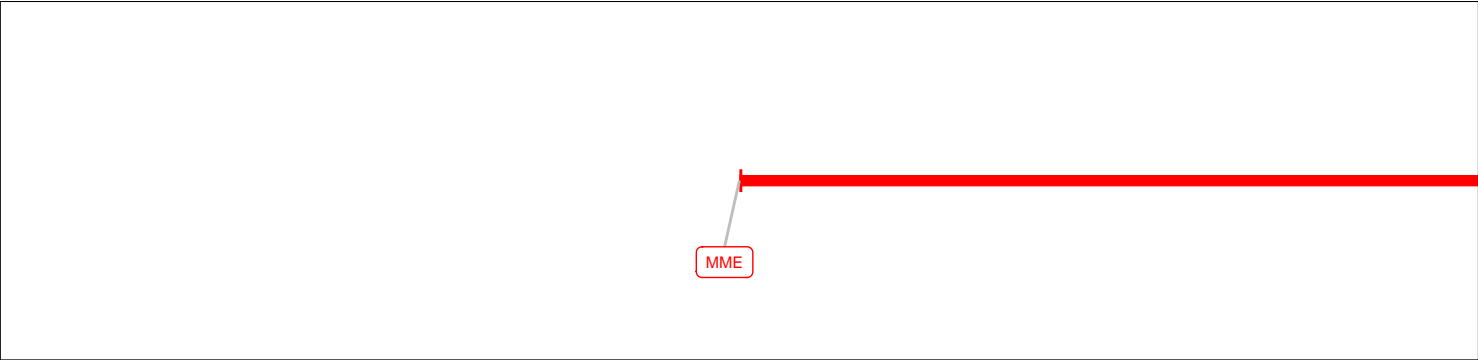
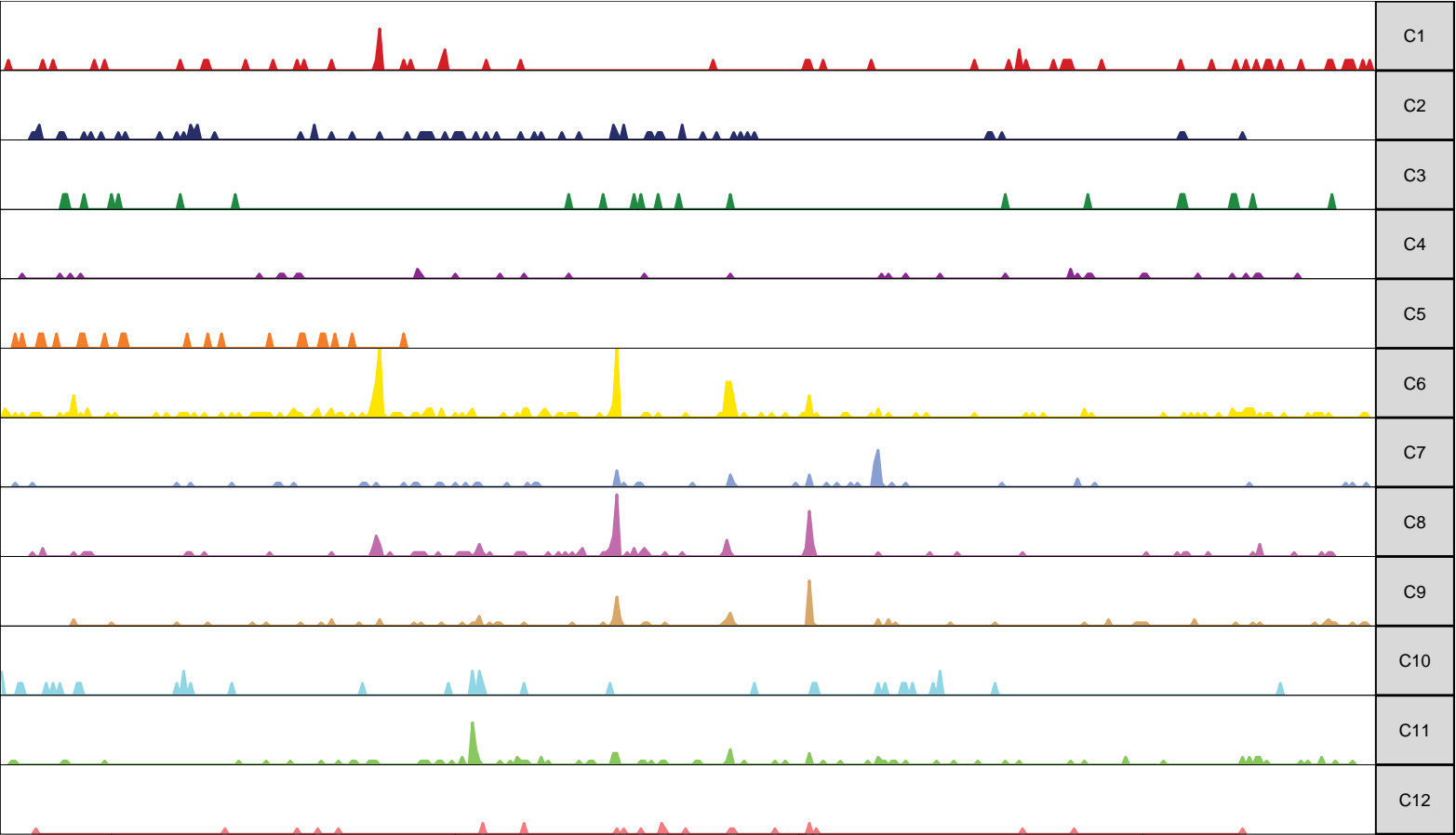


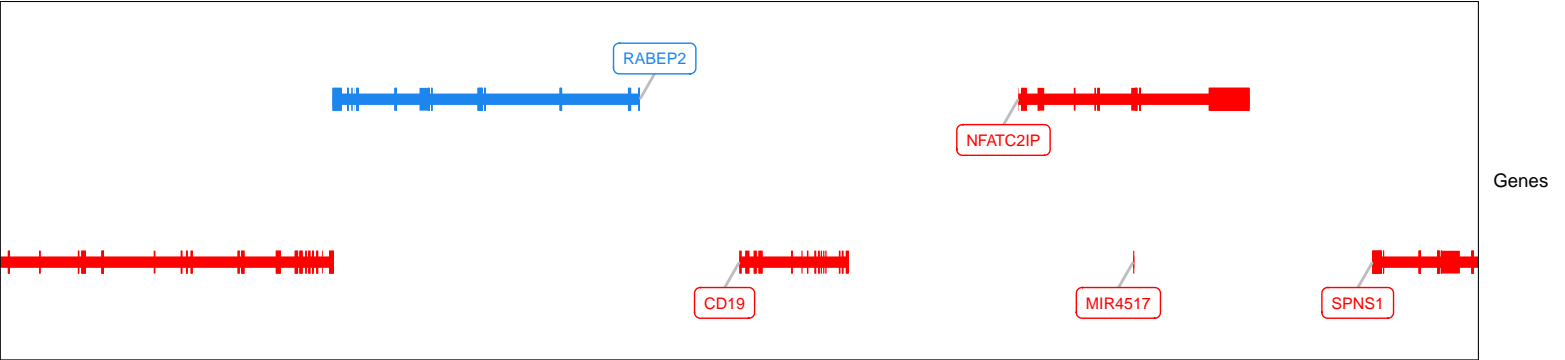
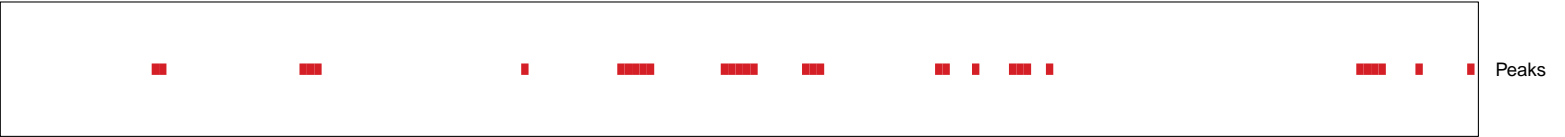
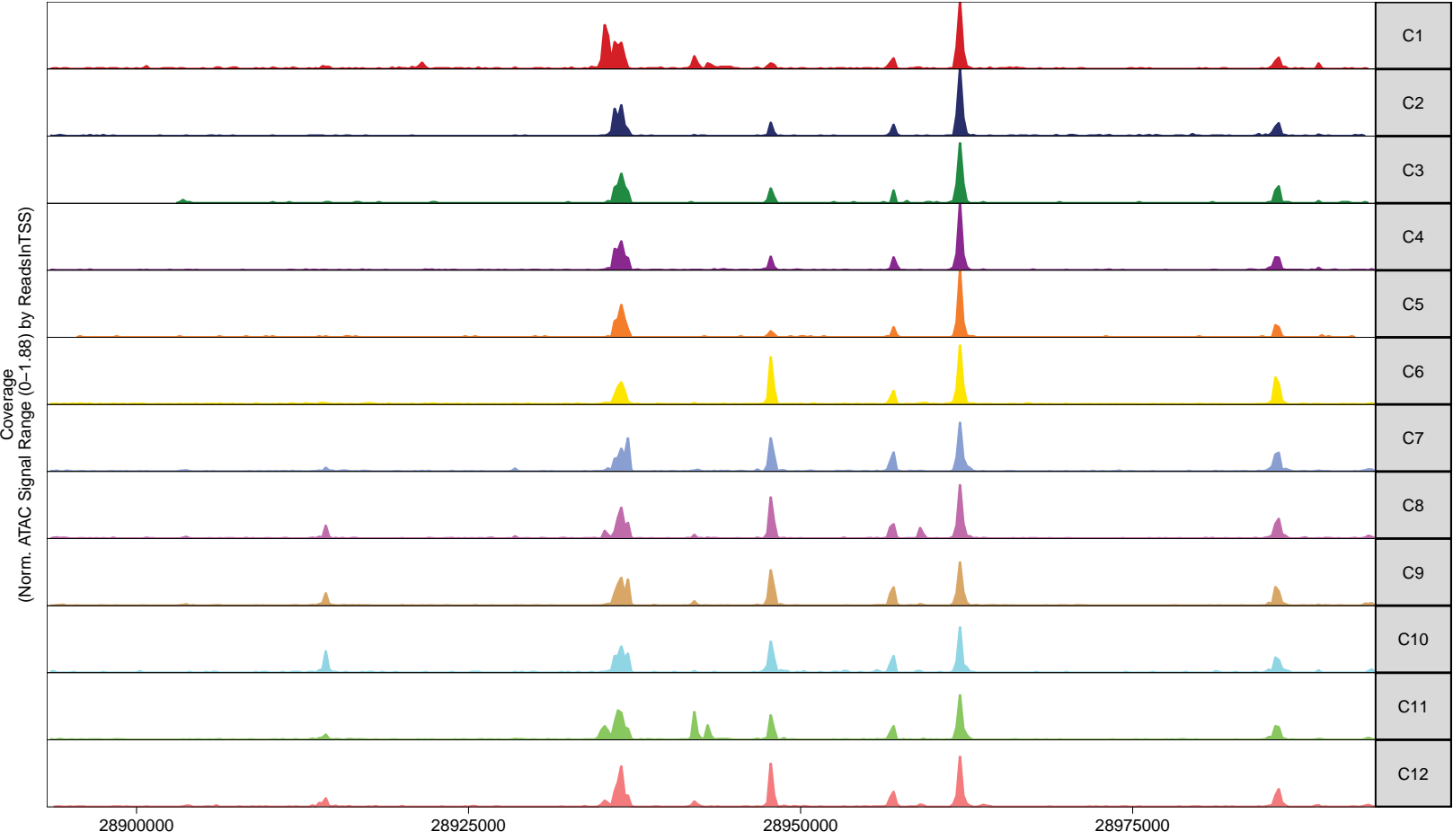
Coverage
(Norm. ATAC Signal Range (0-0.14) by ReadsInTSS)



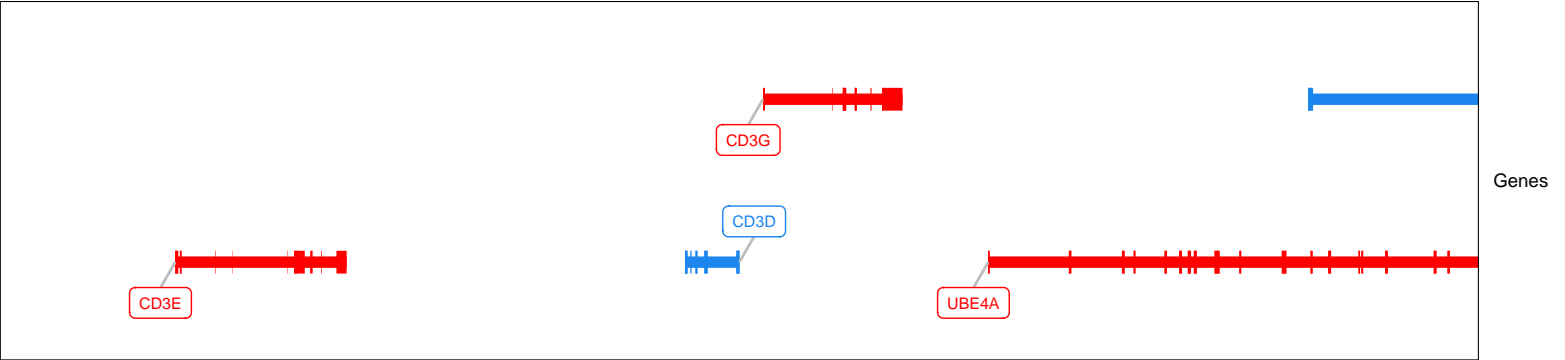
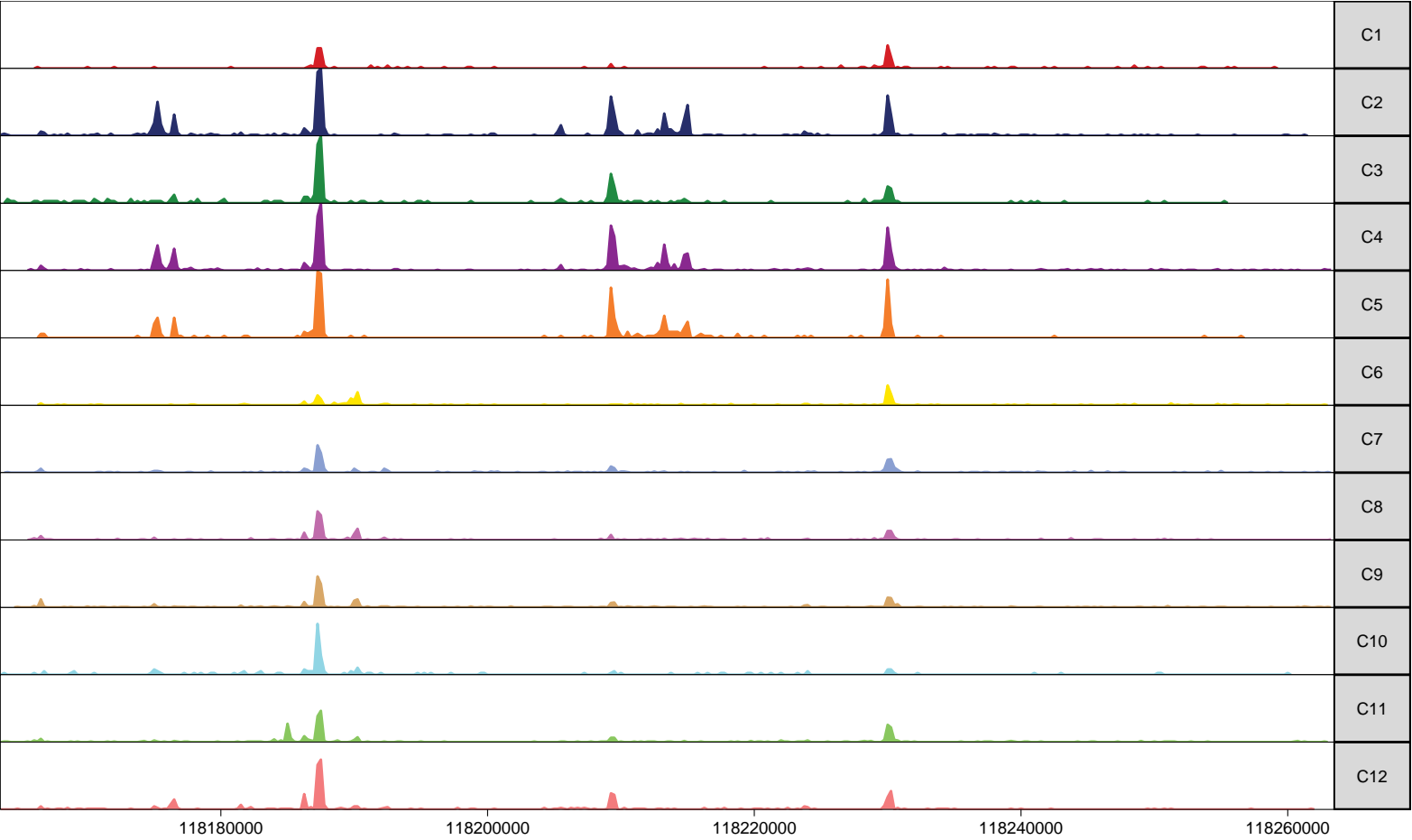
C1
C2
C3
C4
C5
C6
C7
C8
C9
C10
C11
C12

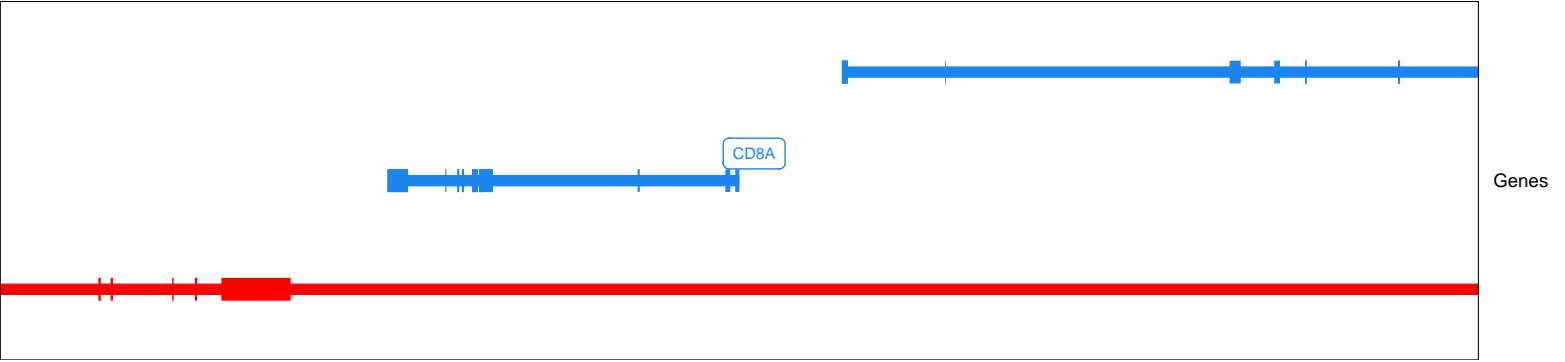
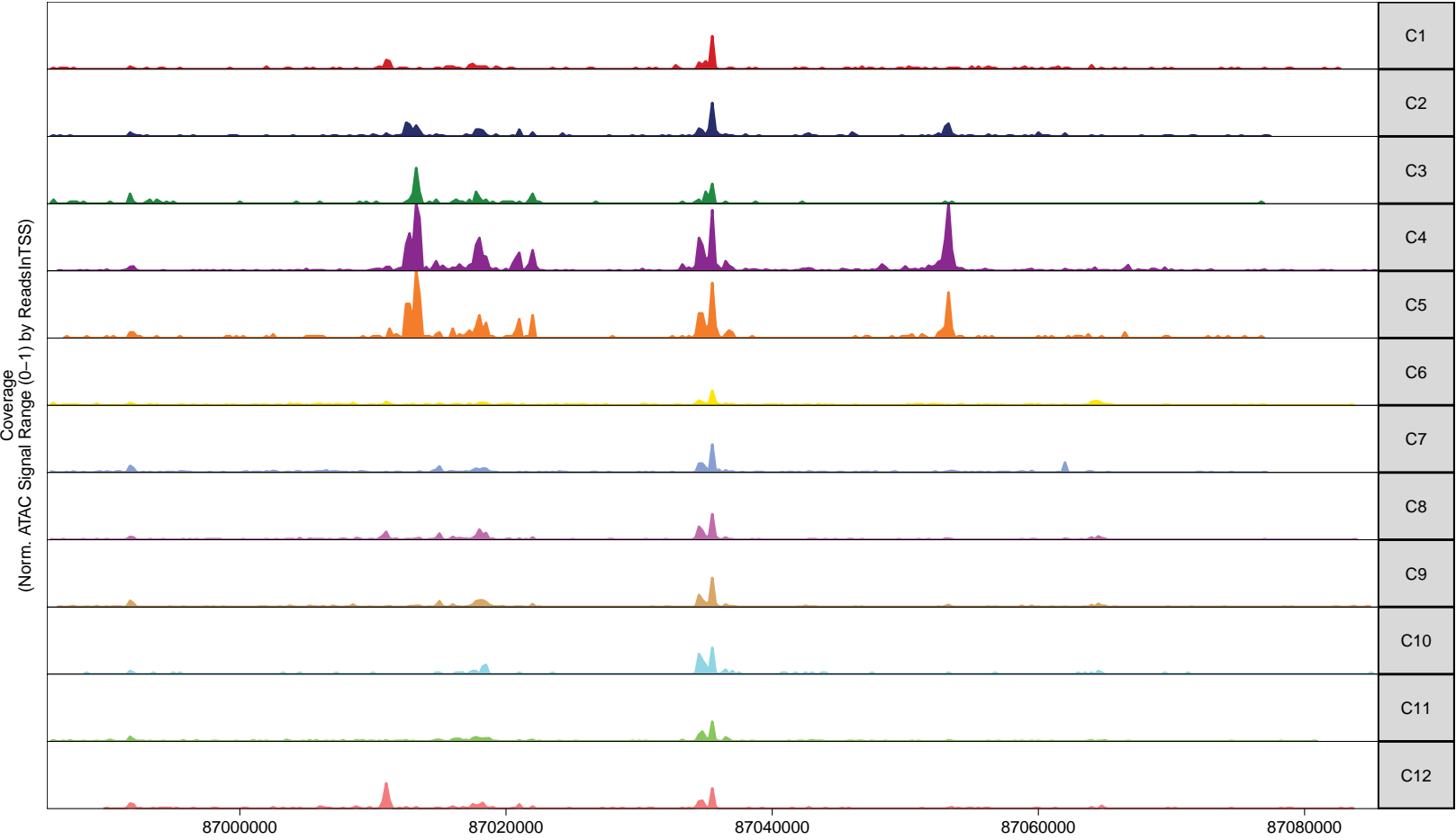
Peaks

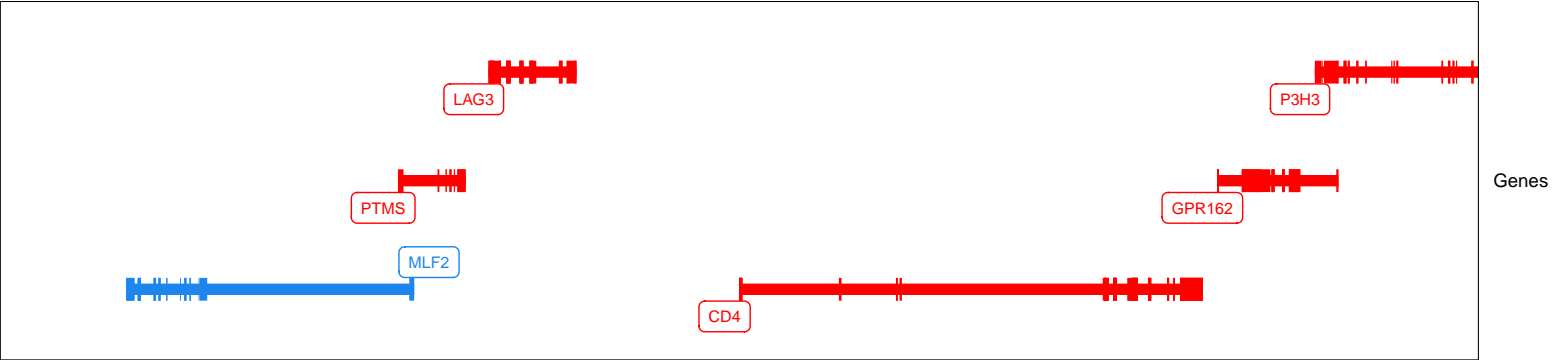
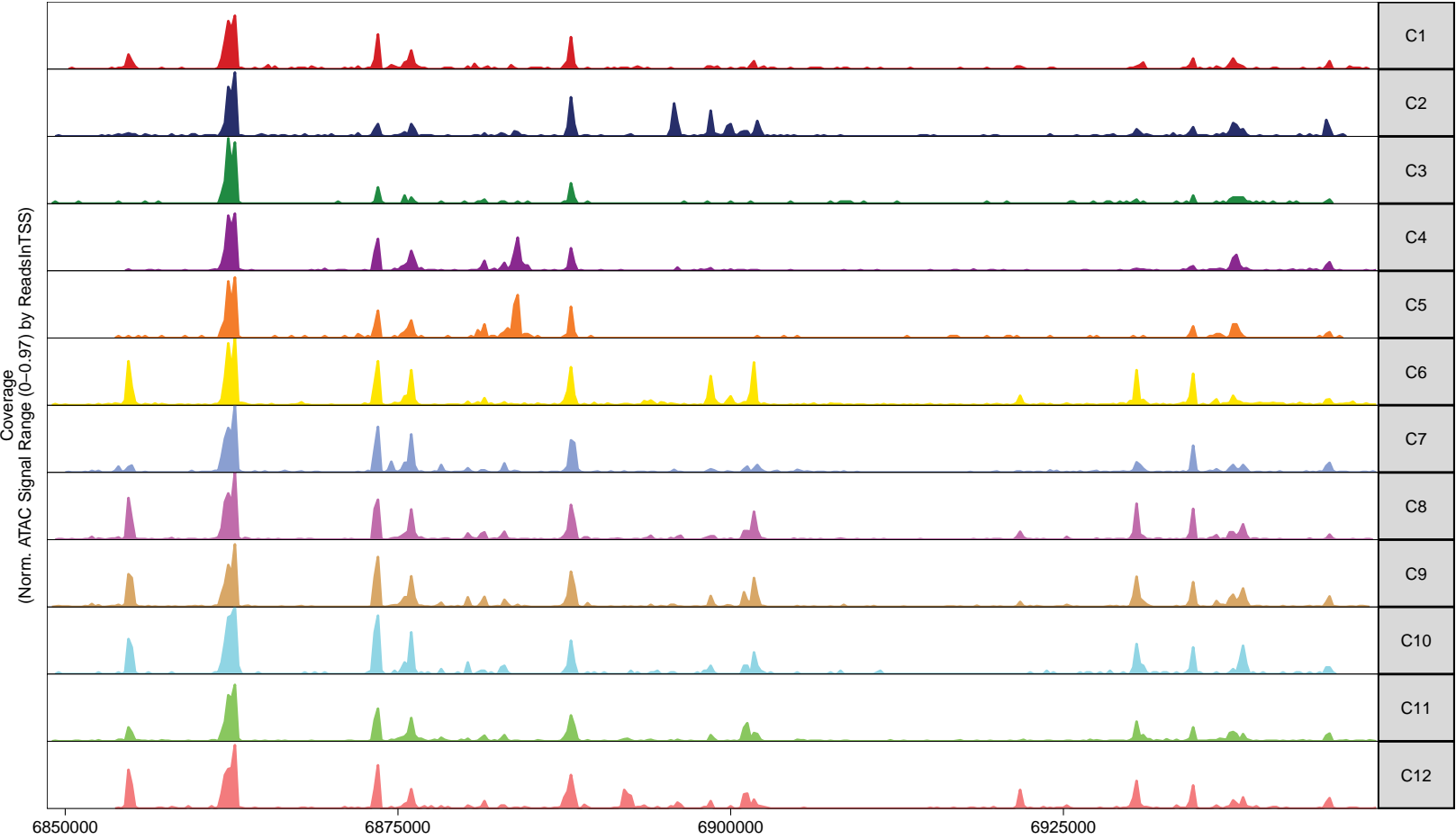
Genes

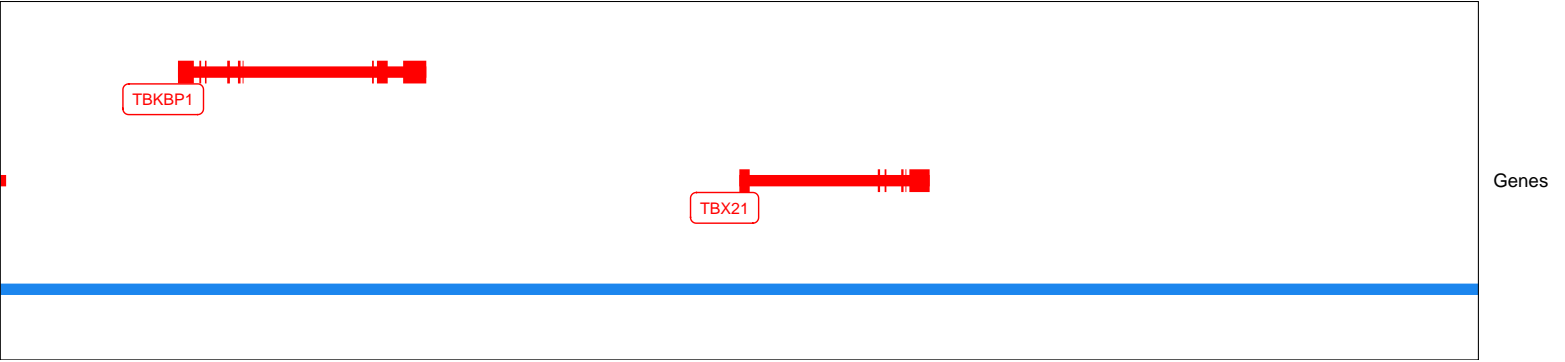
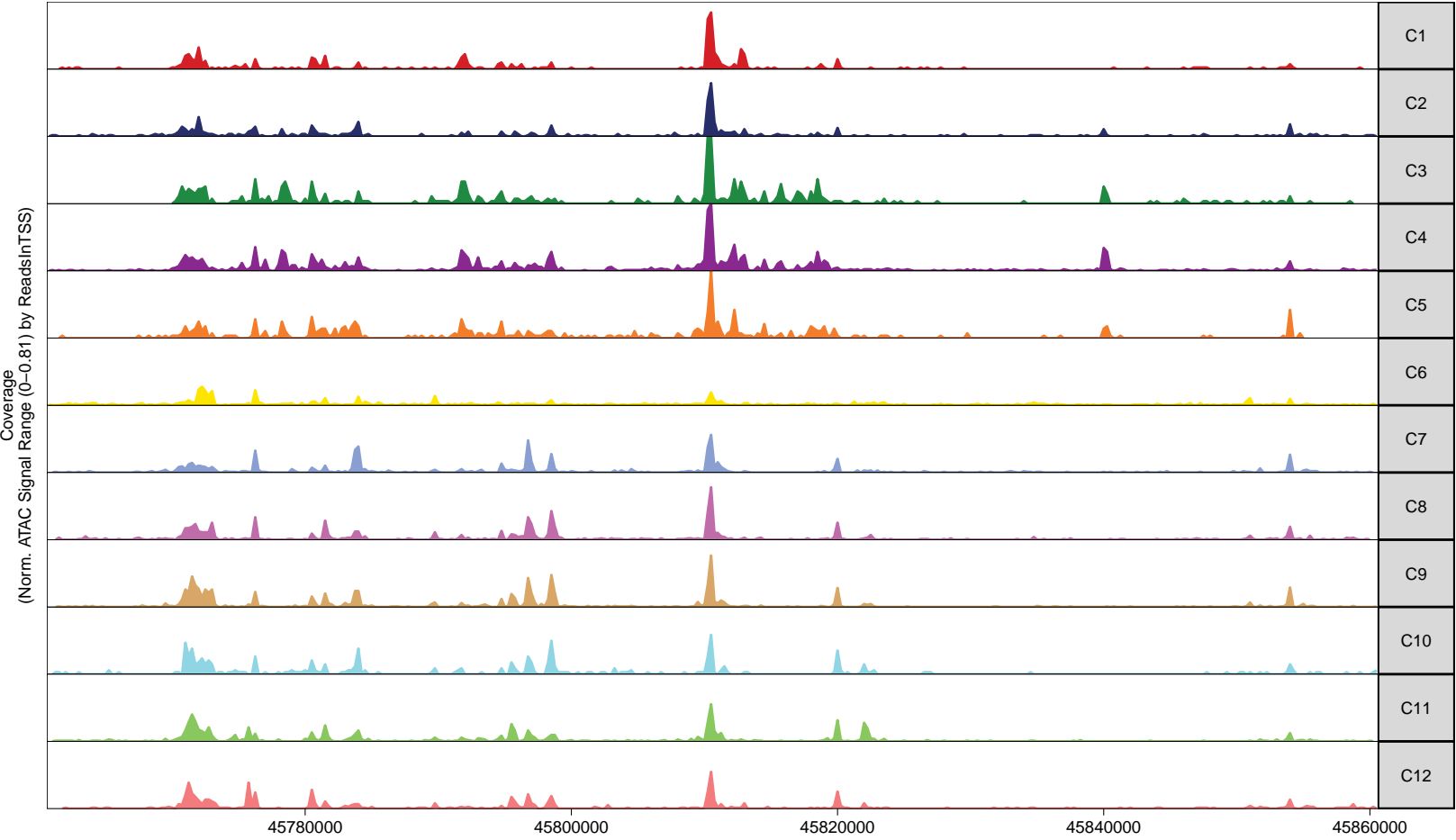


Coverage
(Norm. ATAC Signal Range (0-0.95) by ReadslnTSS)

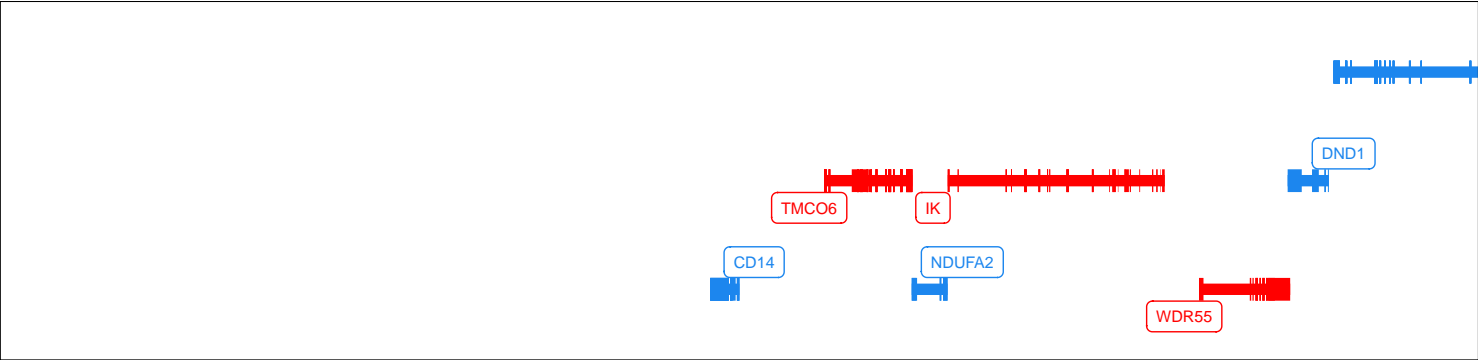
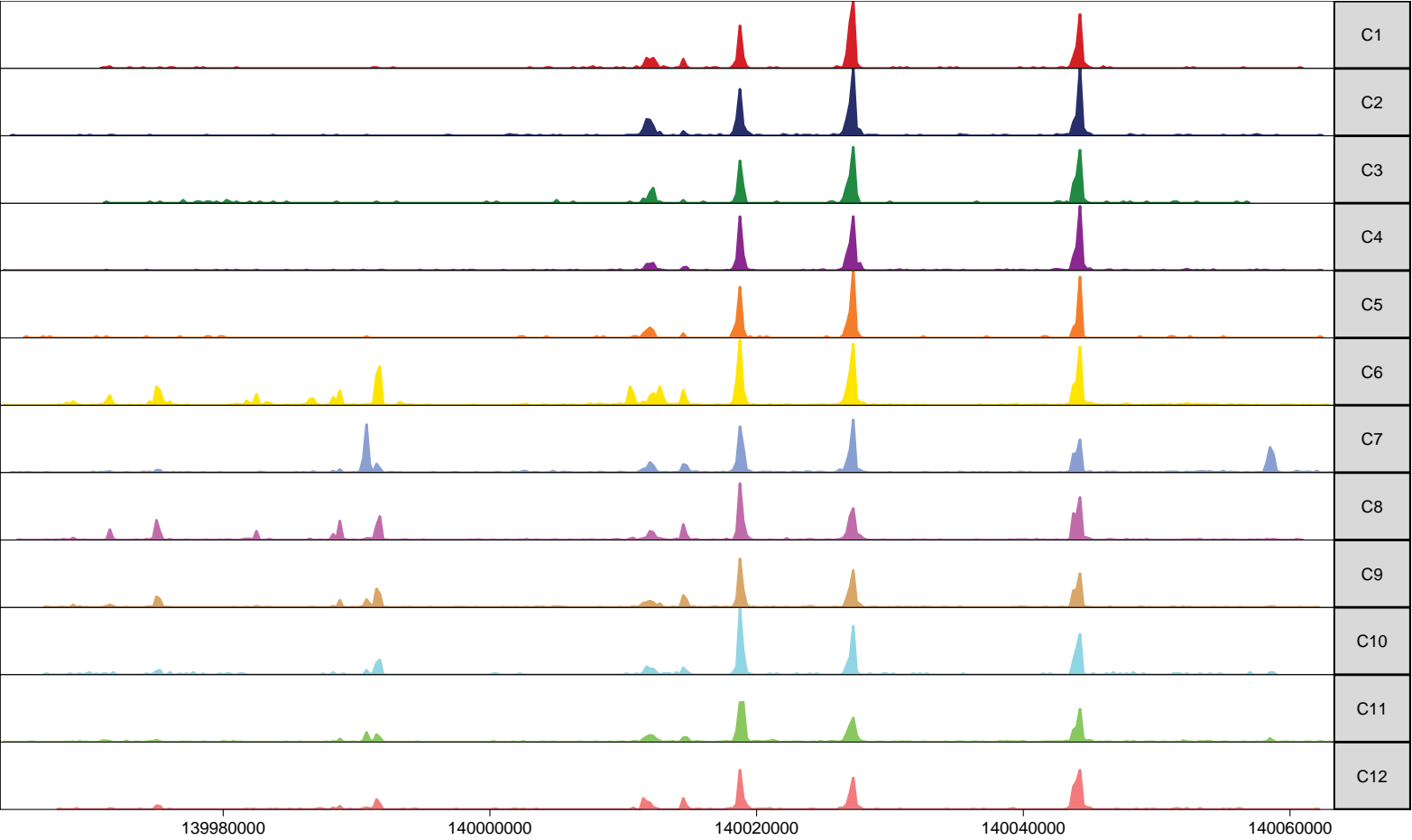


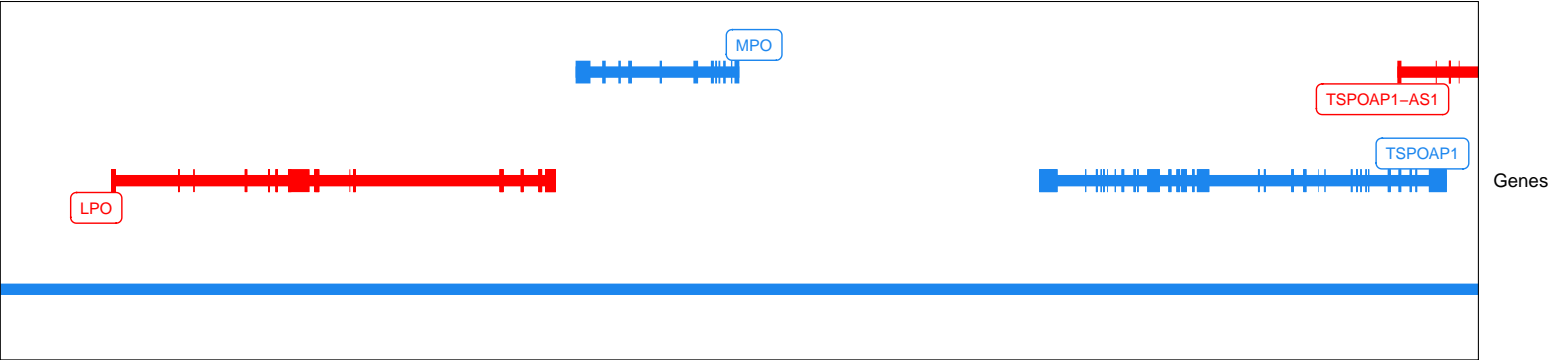
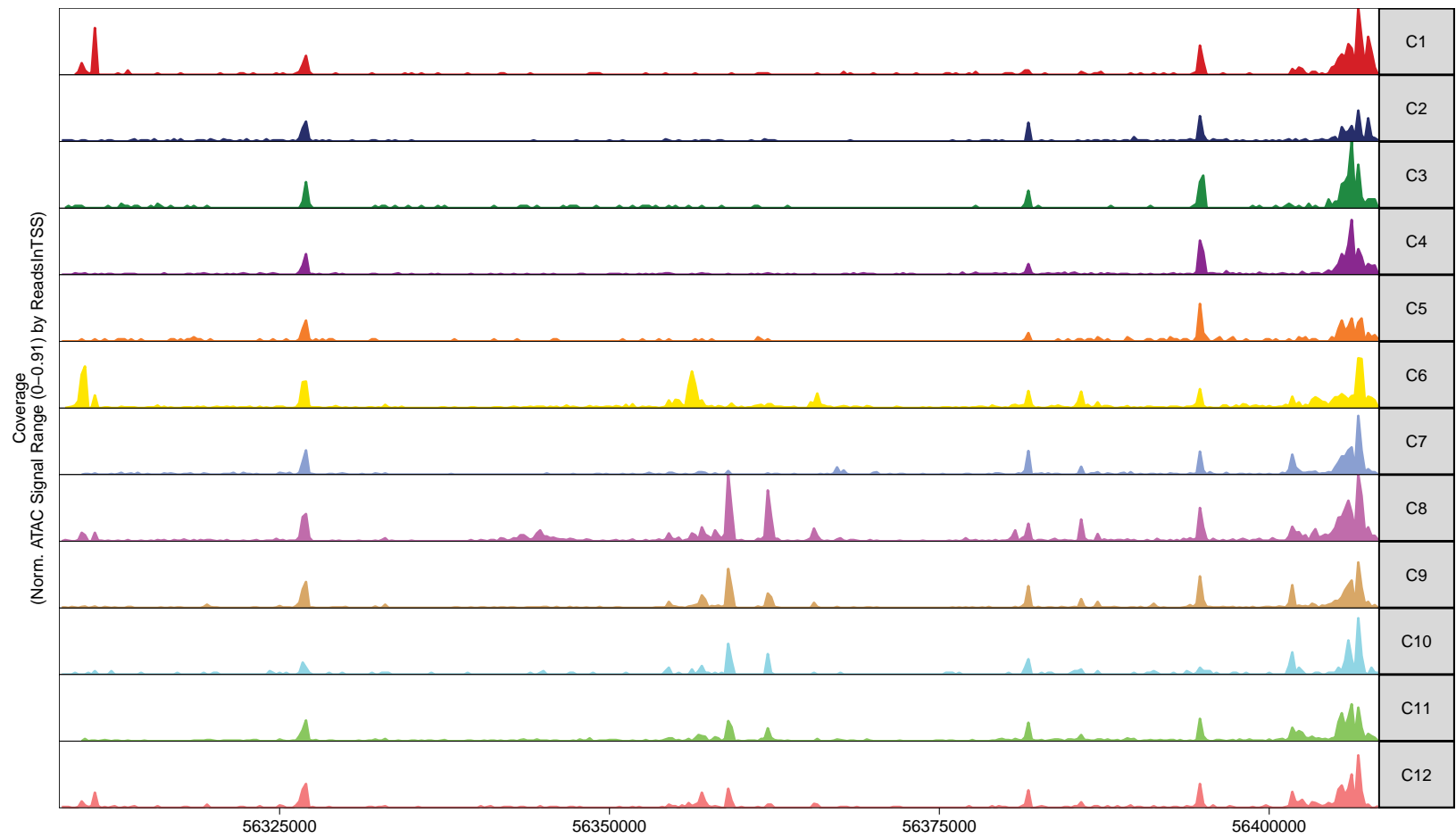


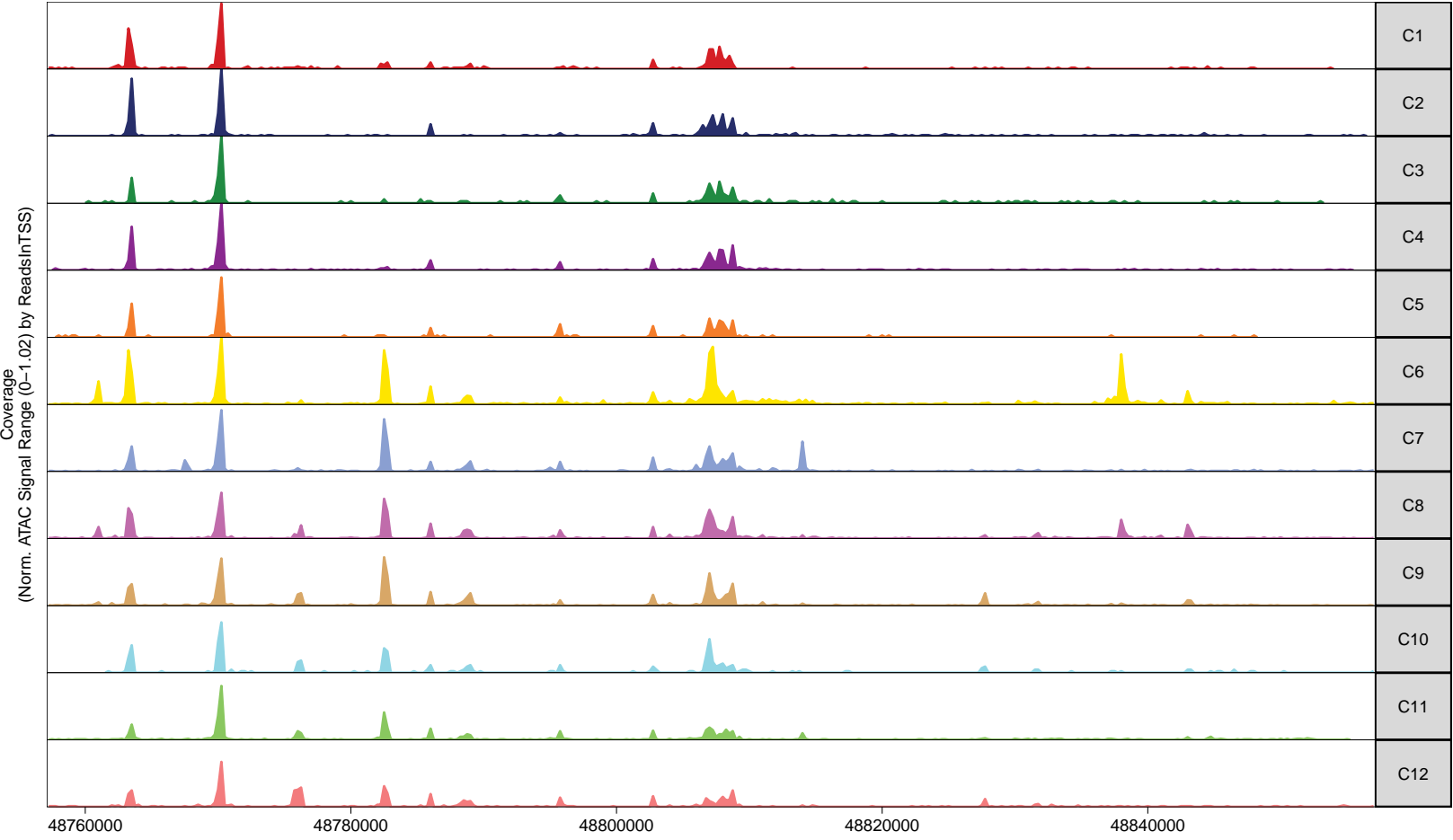


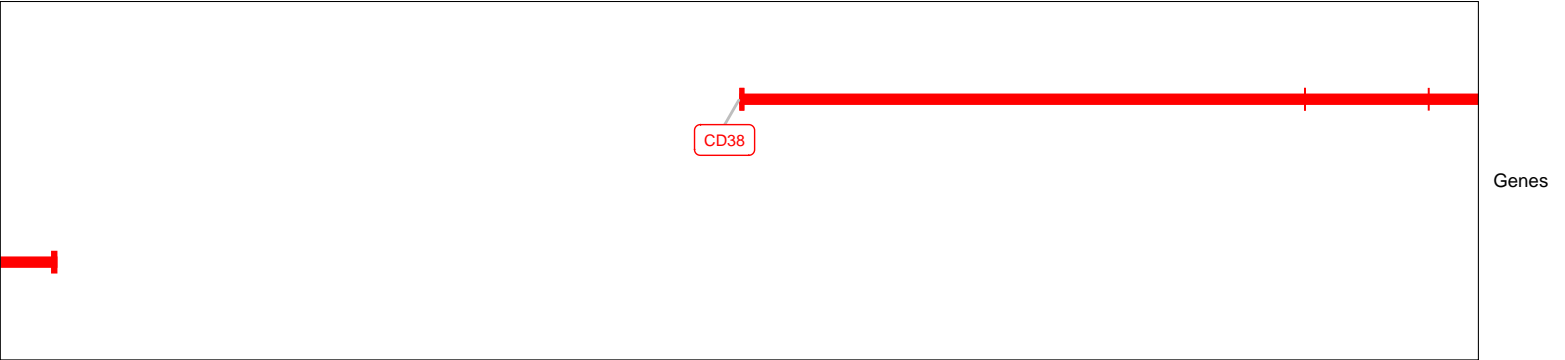
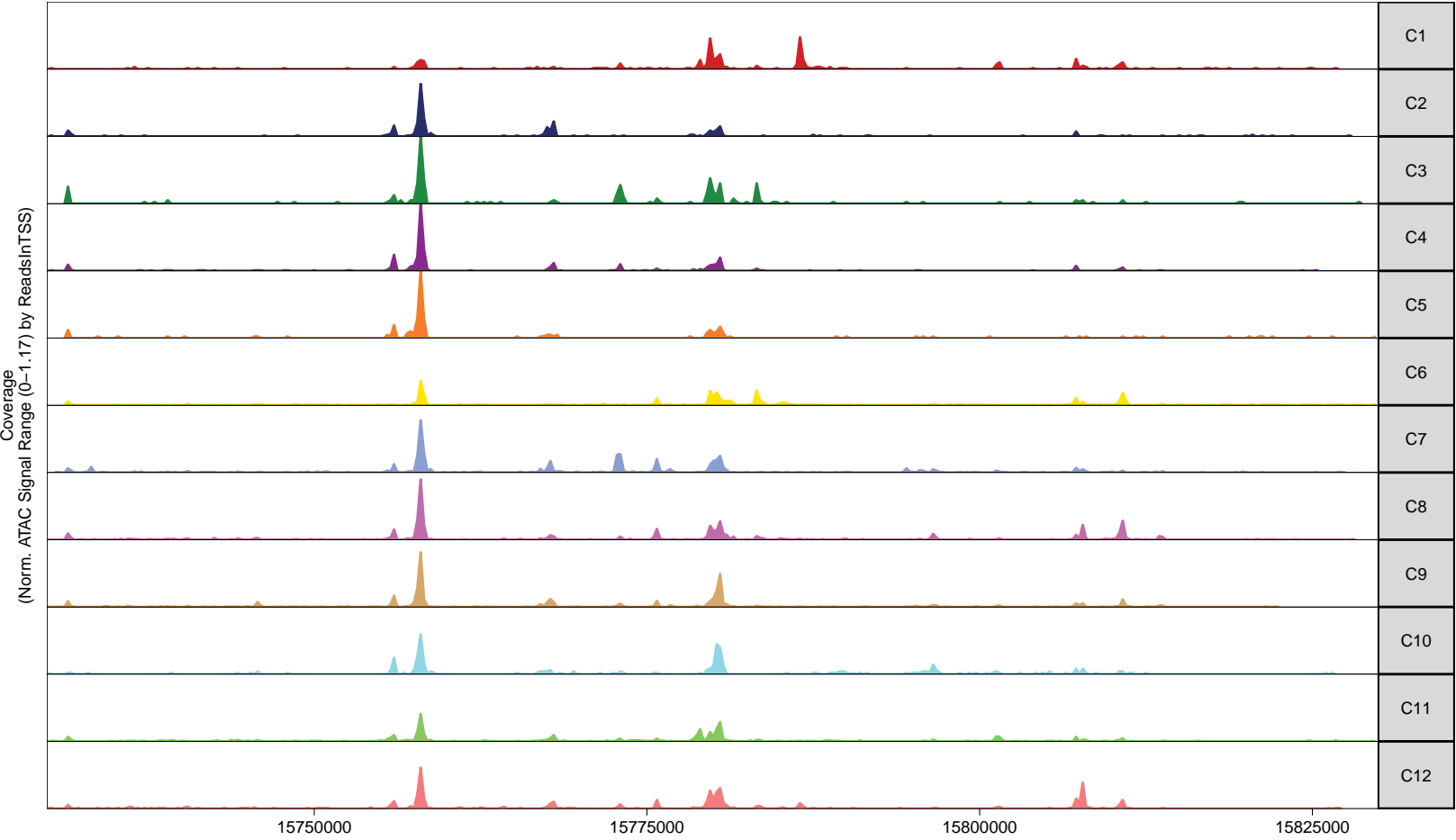


Coverage
(Norm. ATAC Signal Range (0–1.31) by ReadslnTSS)

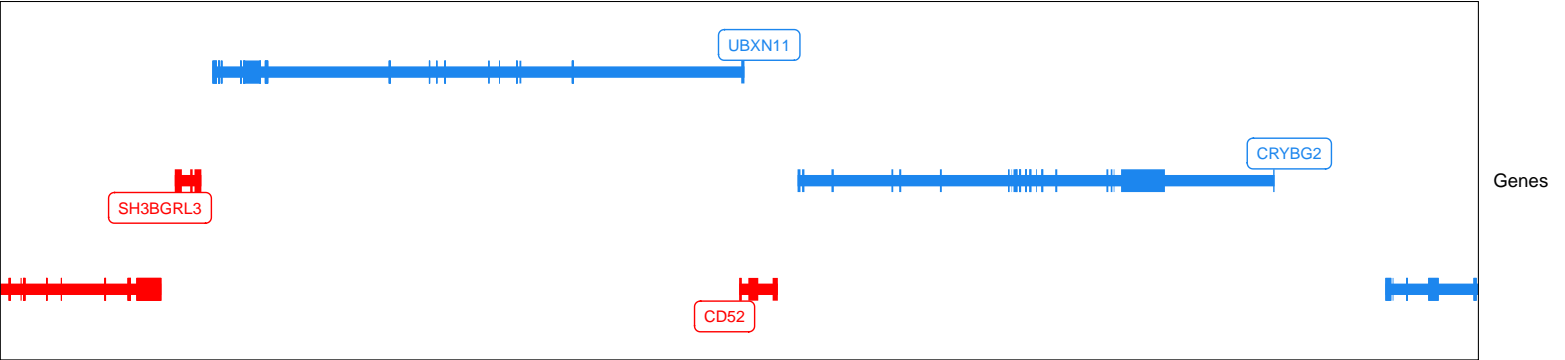


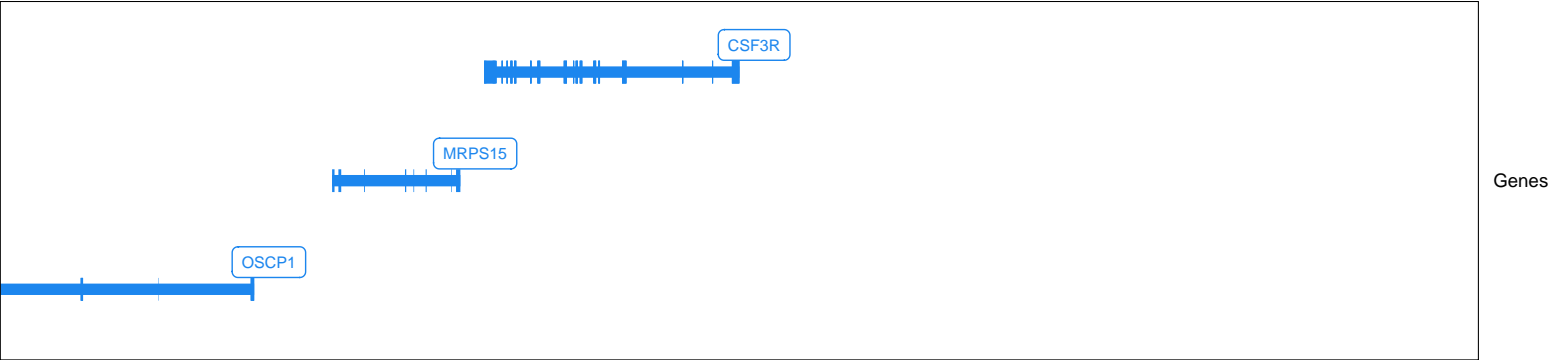
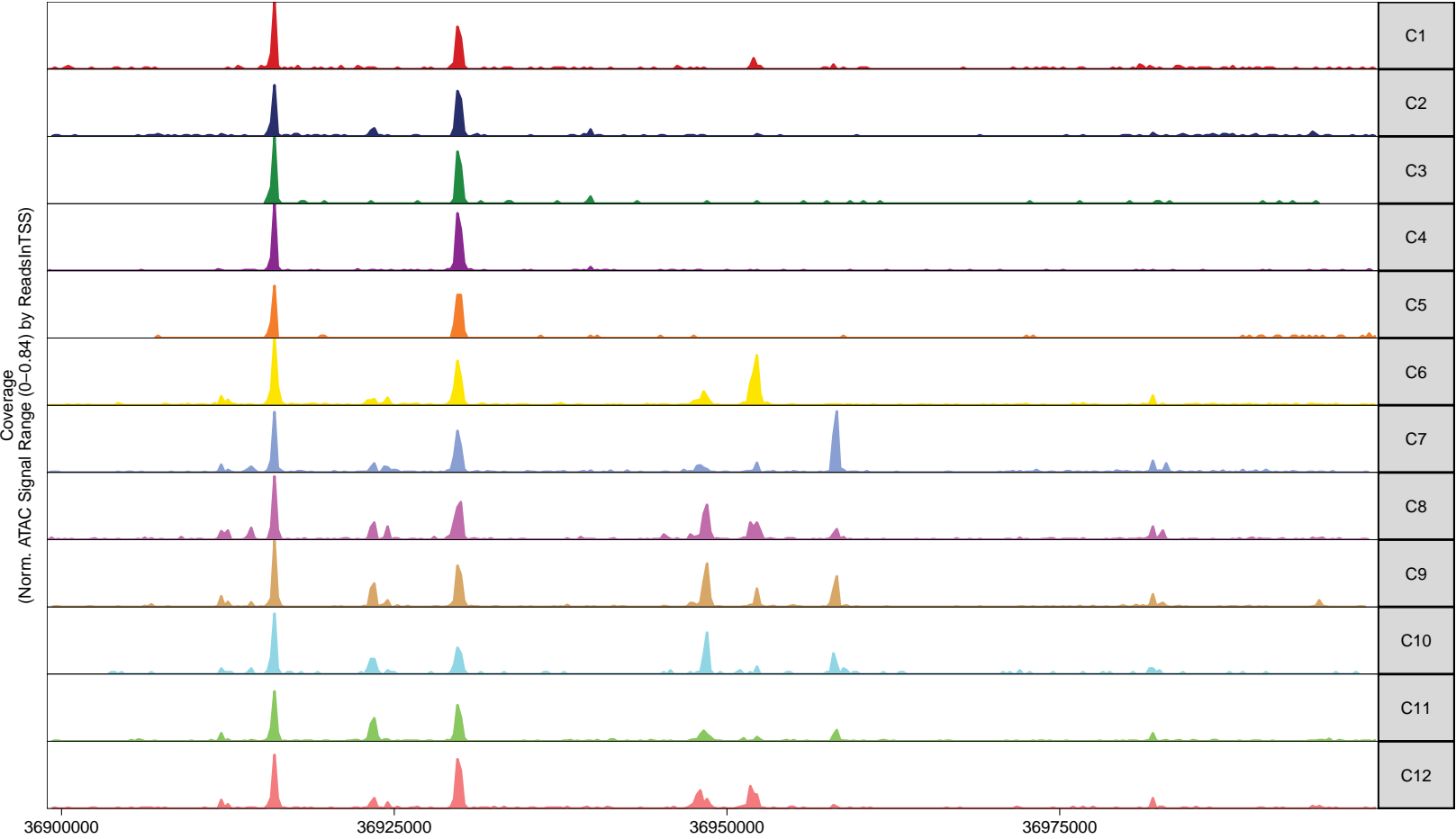




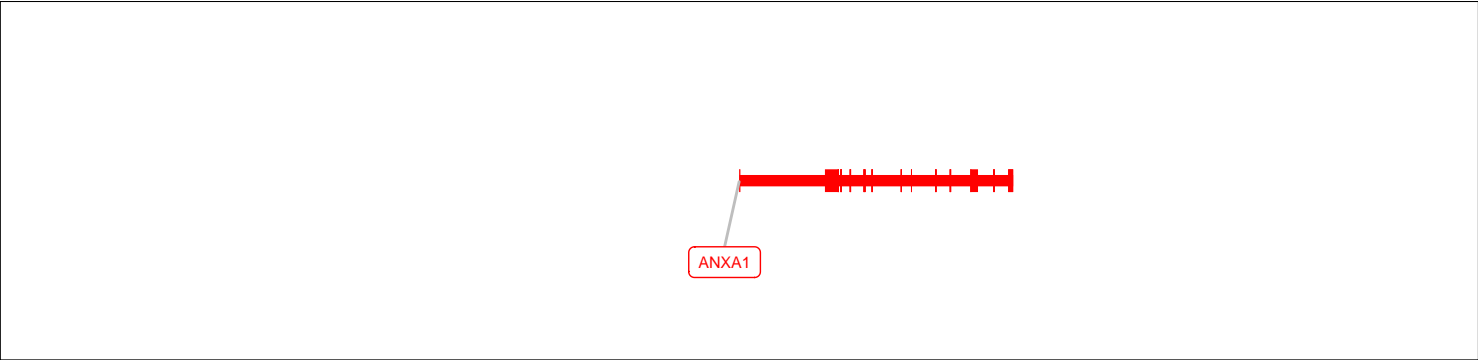
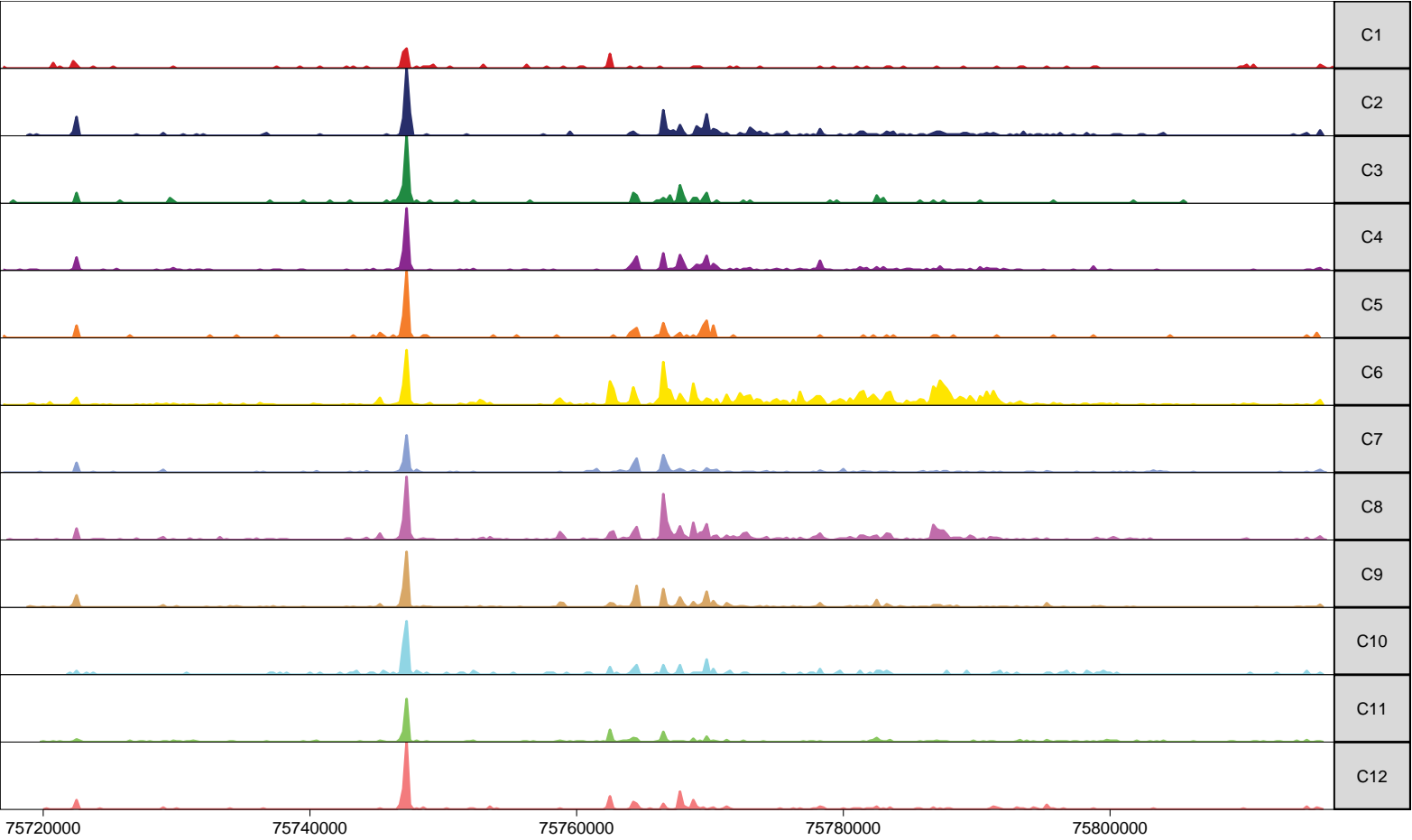


chr1:26594410–26694411





Coverage
(Norm. ATAC Signal Range (0-0.78) by ReadslnTSS)



C1

C2

C3

C4

C5

C6

C7

C8

C9

C10

C11

C12

Peaks

Genes

