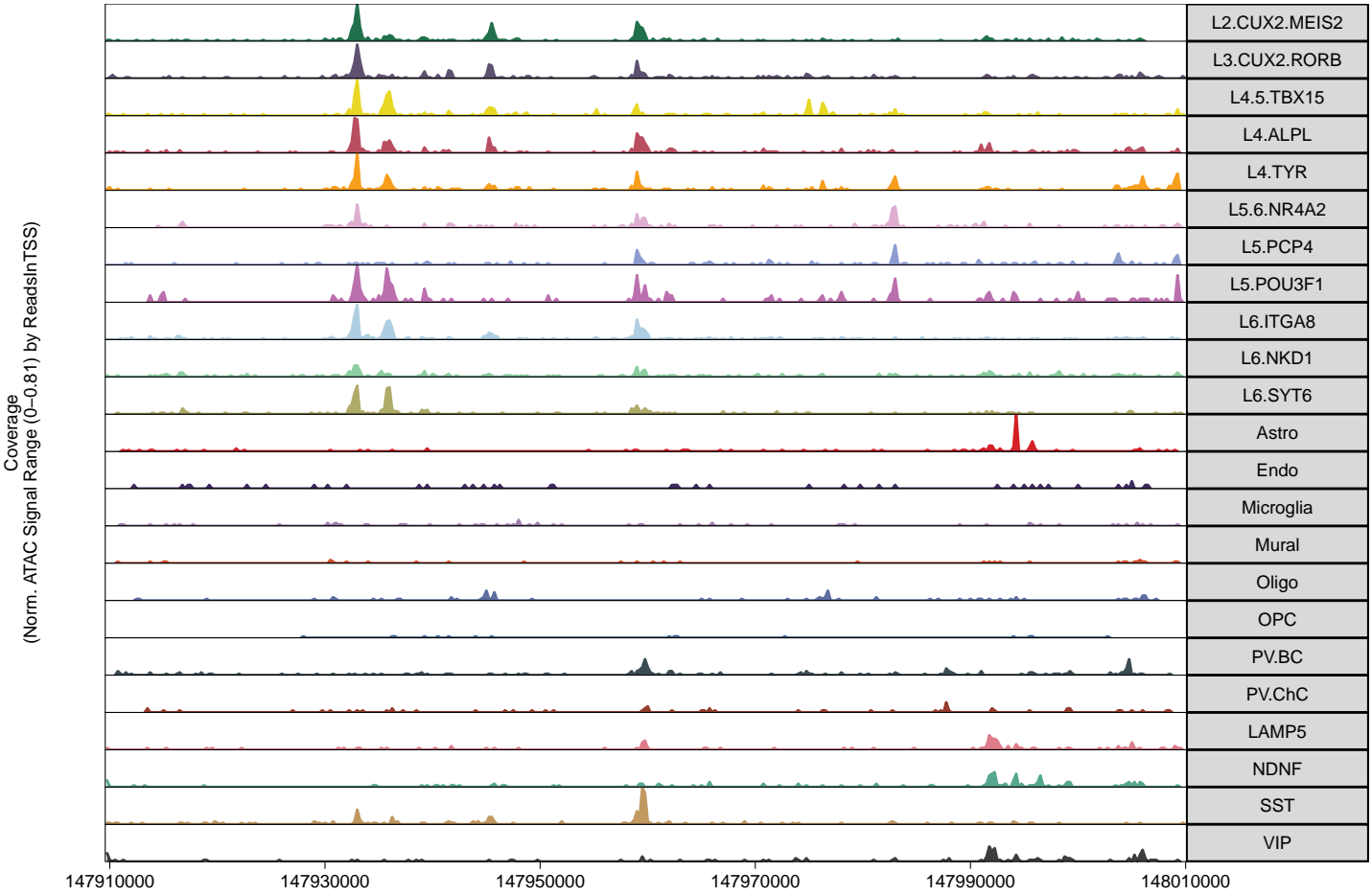
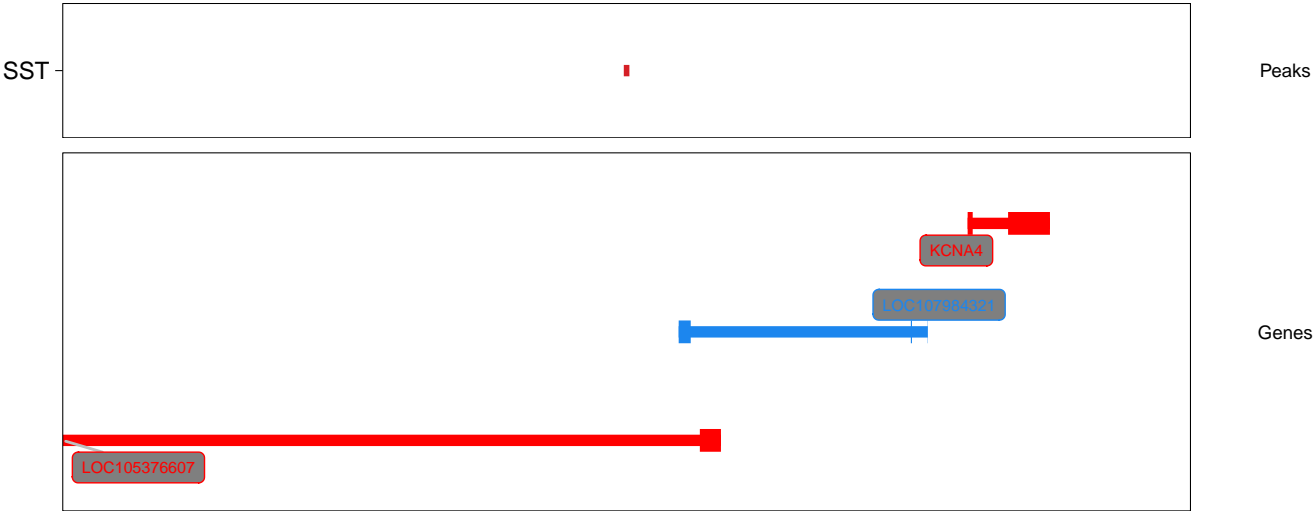
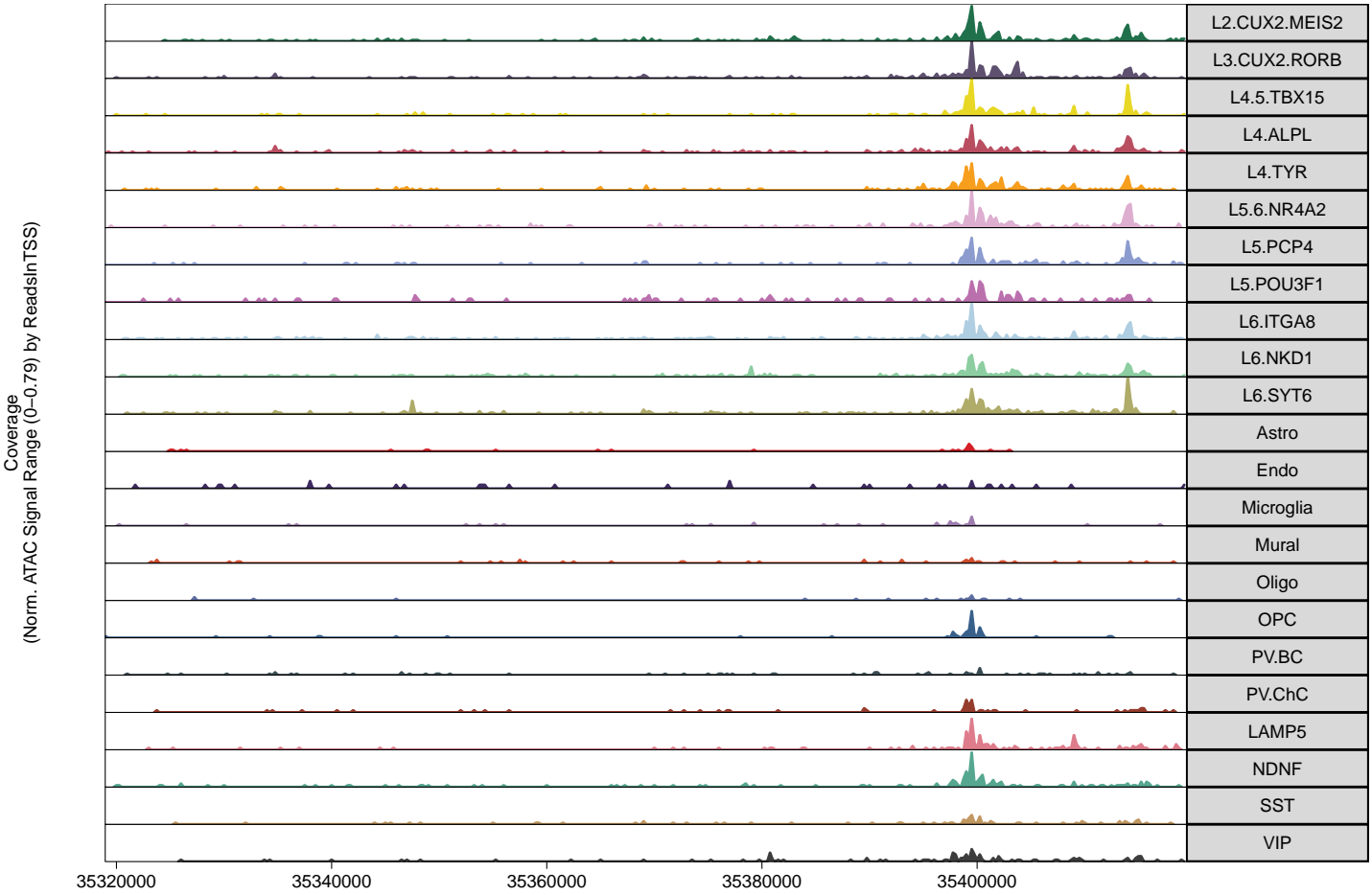


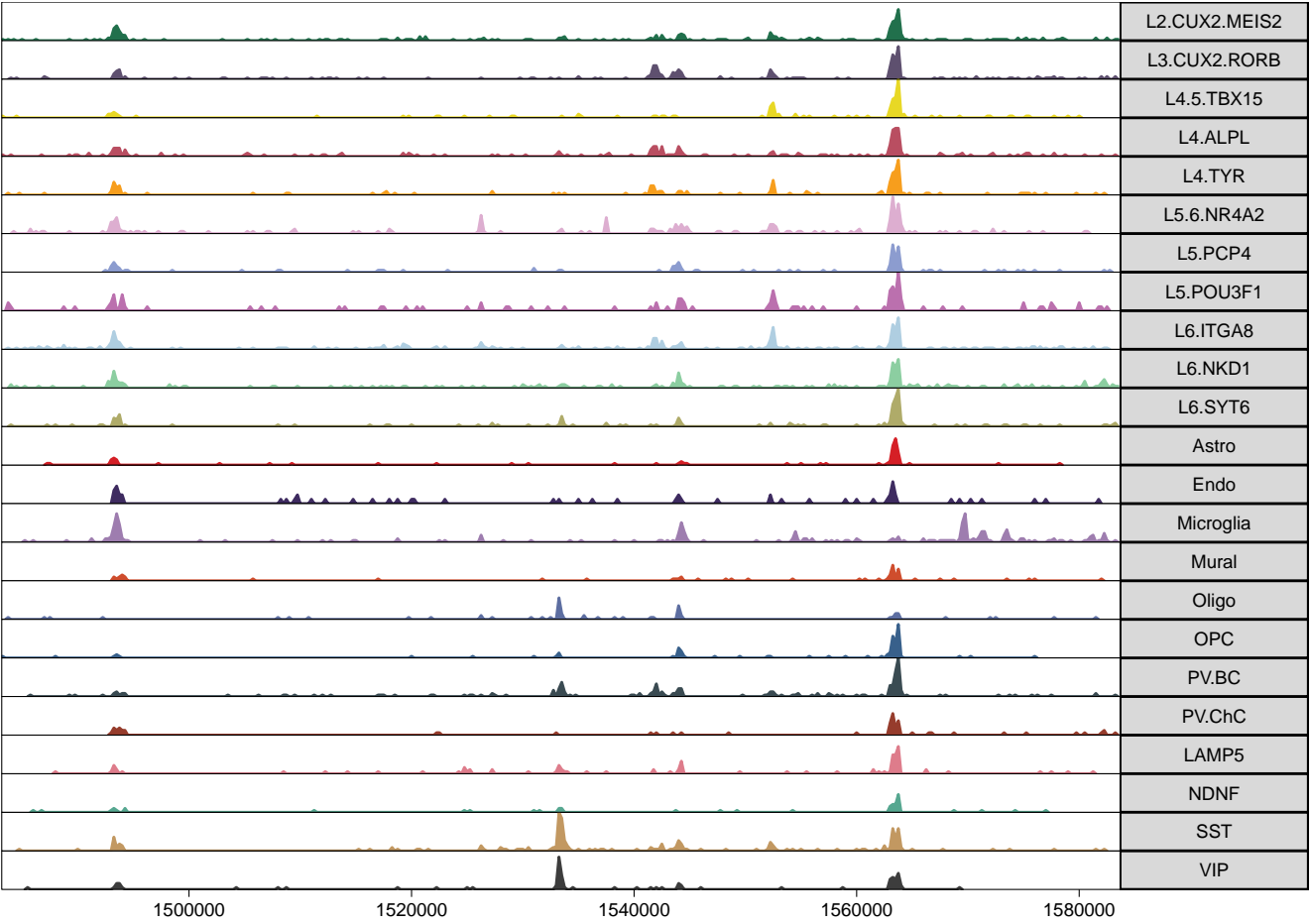
chr1:147909572–148010073 Marker Peak Tracks for: SST



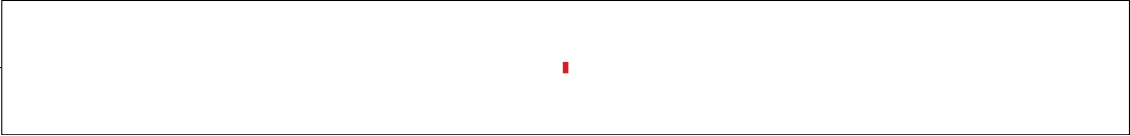
chr14:35318955–35419456 Marker Peak Tracks for: SST



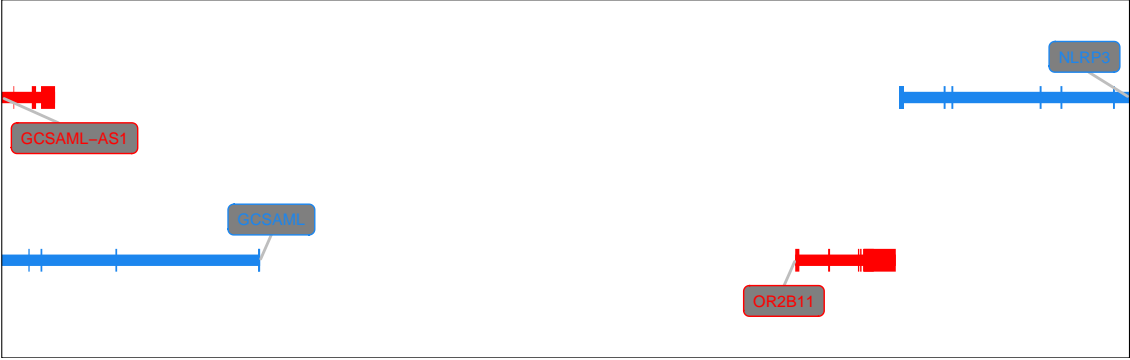
Coverage  
(Norm. ATAC Signal Range (0–0.71) by ReadsInTSS)



SST



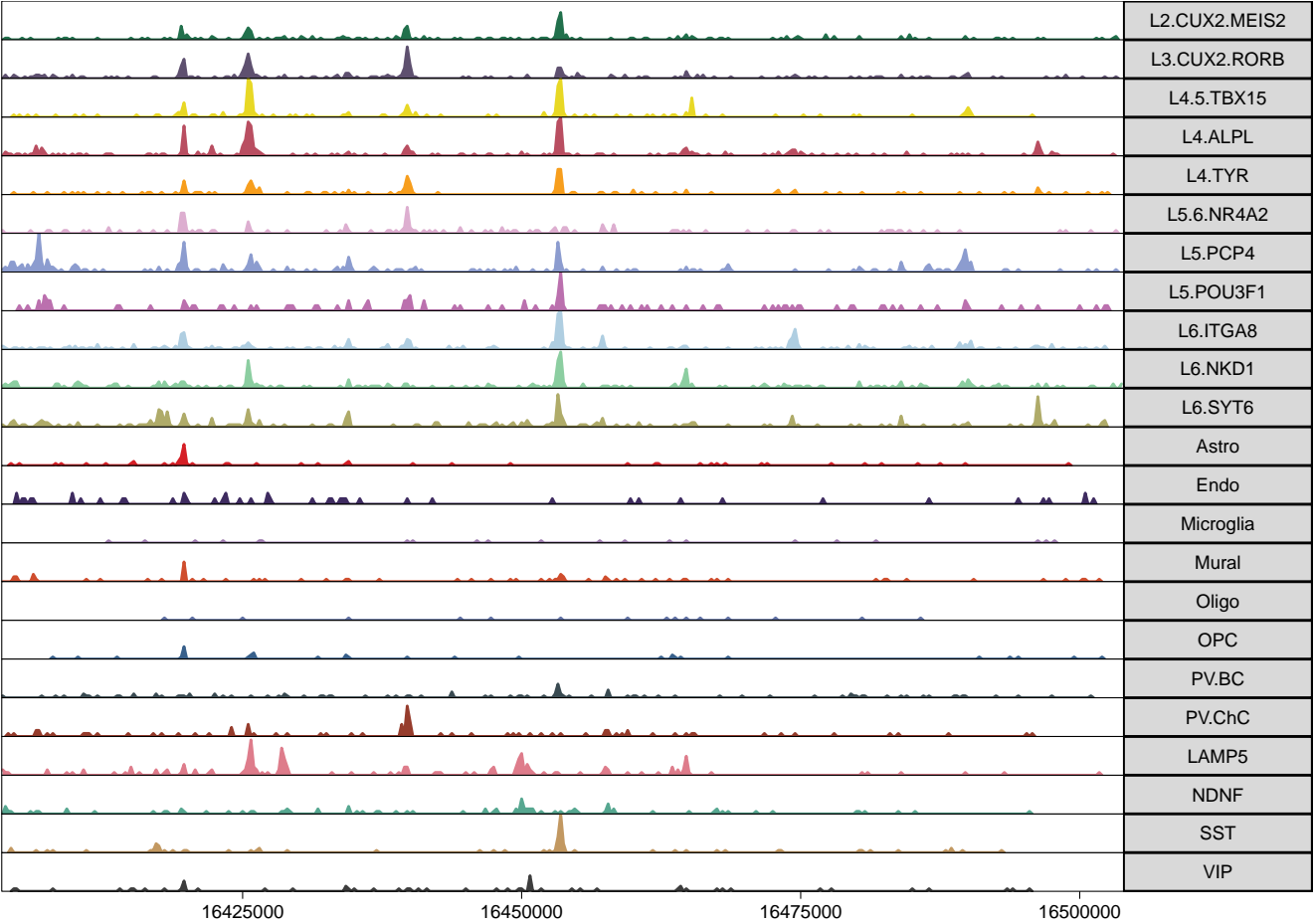
Peaks



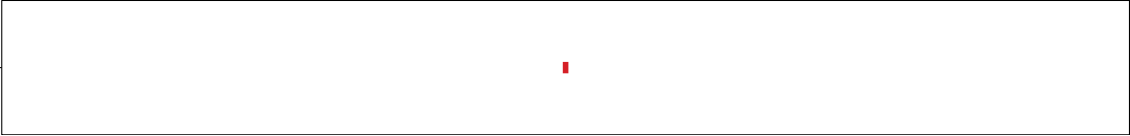
Genes

chr14:16403418–16503919 Marker Peak Tracks for: SST

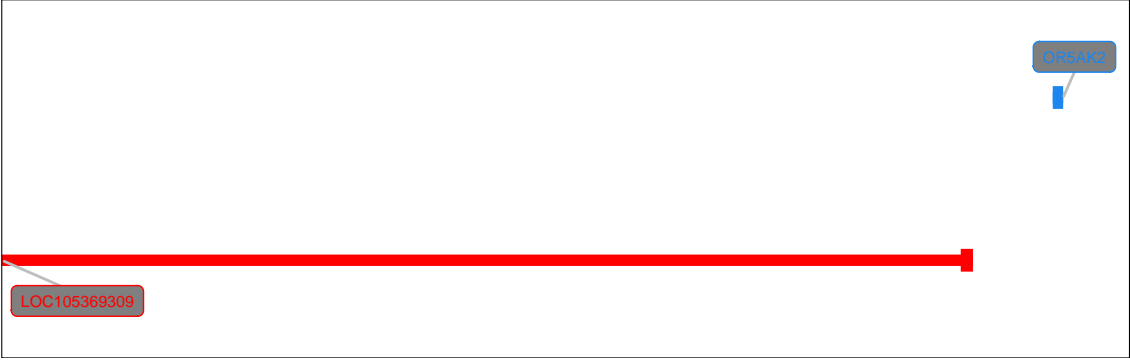
Coverage  
(Norm. ATAC Signal Range (0–0.56) by ReadsInTSS)



SST

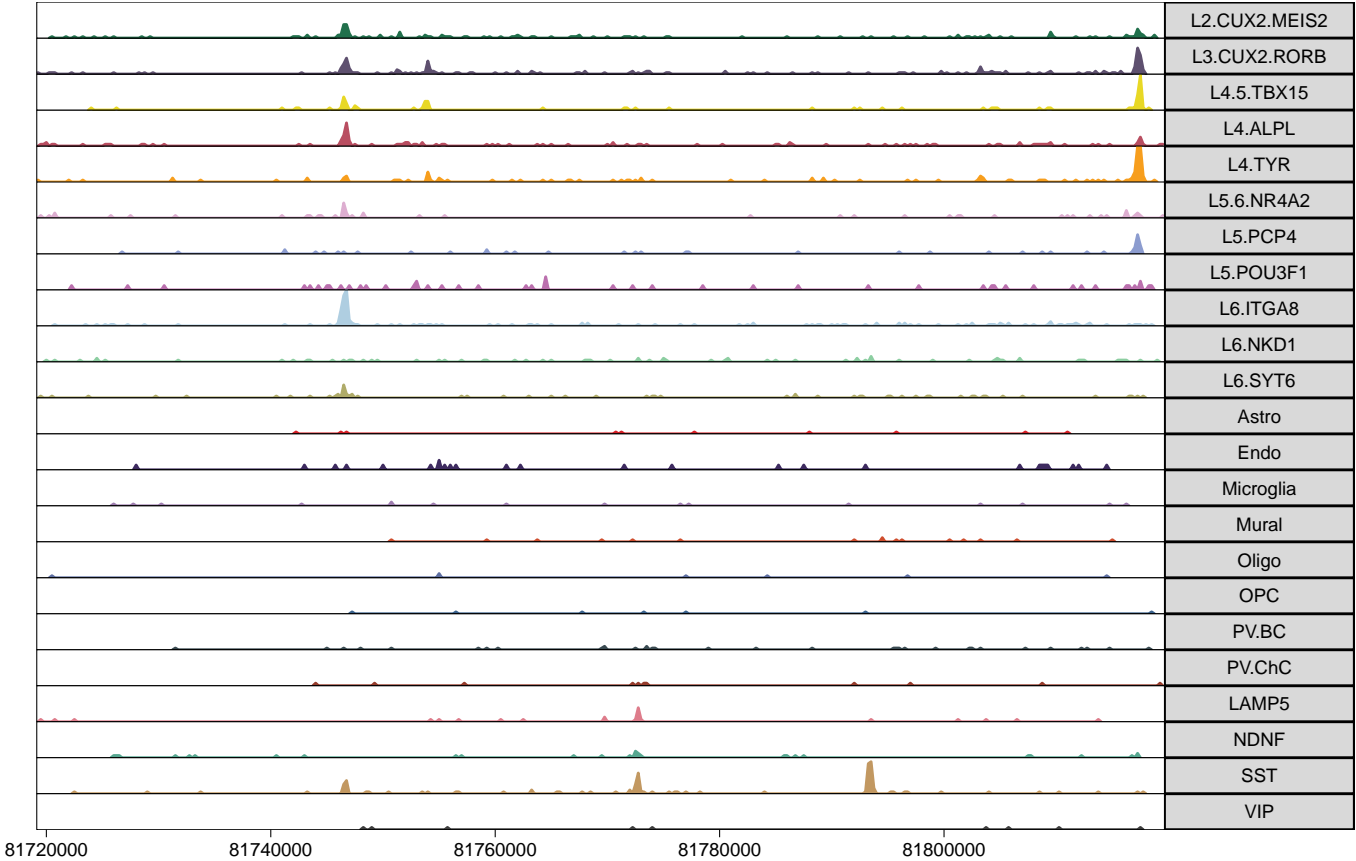


Peaks



Genes

Coverage  
(Norm. ATAC Signal Range (0–0.6) by ReadsInTSS)



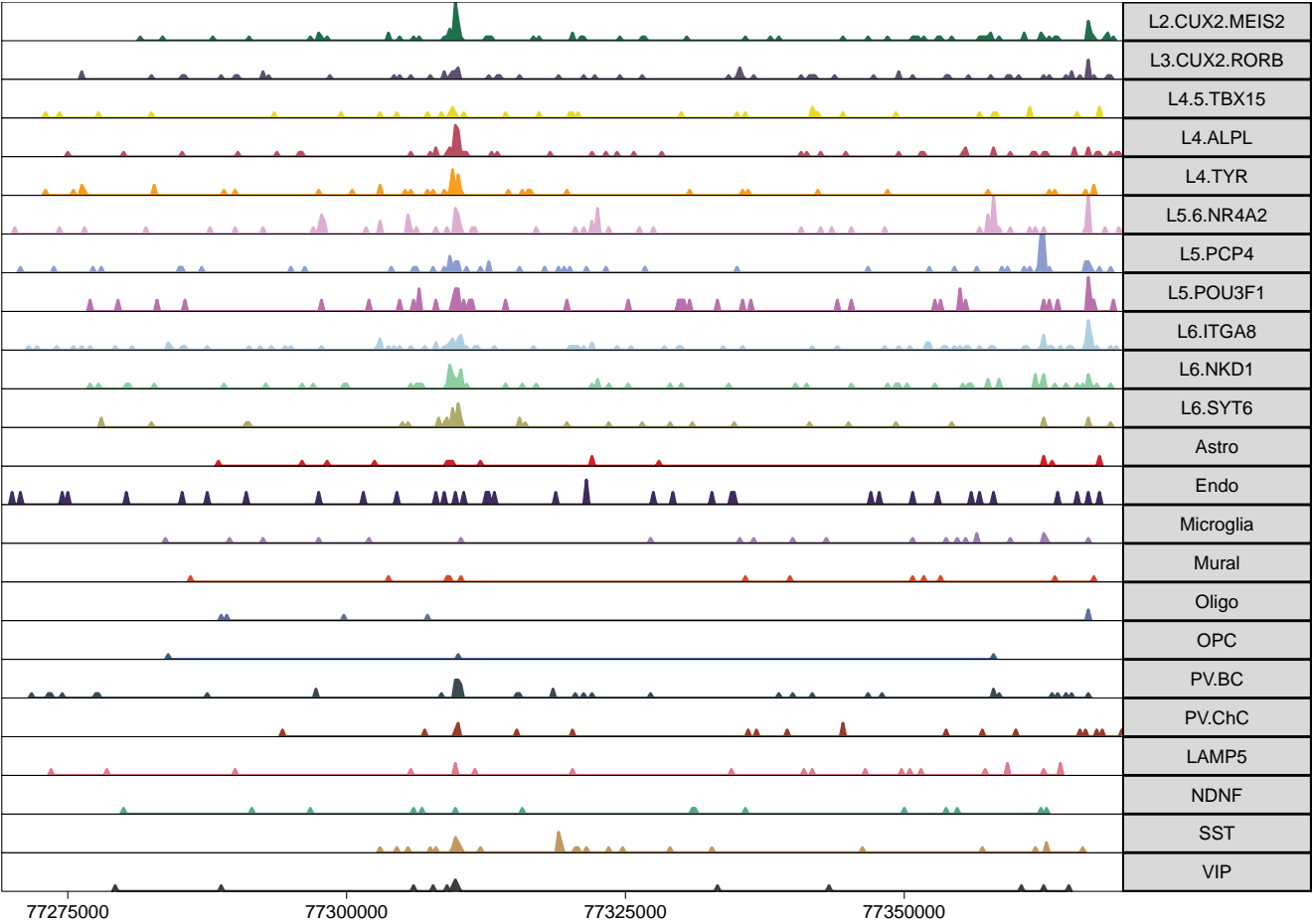
SST

Peaks

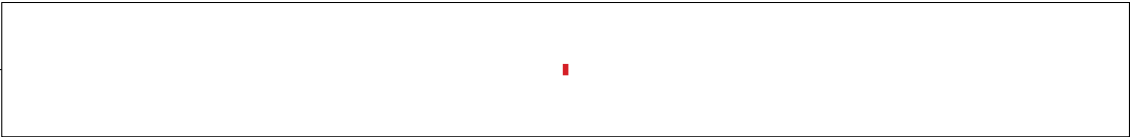
GeneTrack

chr13:77269075–77369576 Marker Peak Tracks for: SST

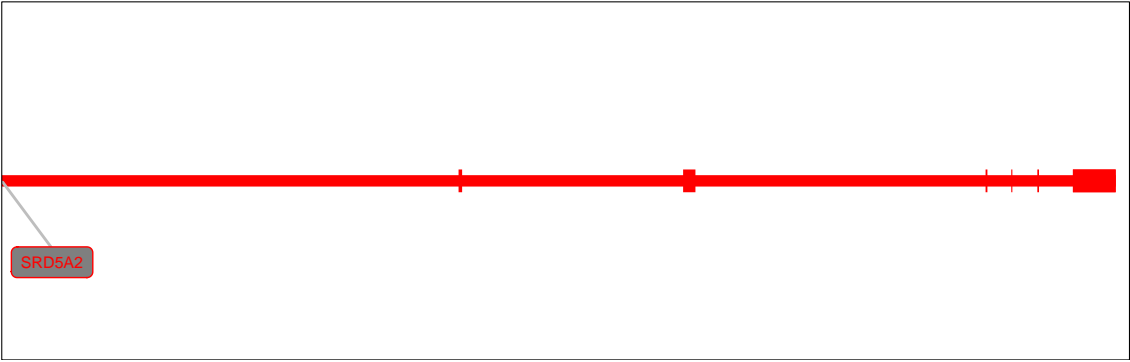
Coverage  
(Norm. ATAC Signal Range (0–0.25) by ReadsInTSS)



SST

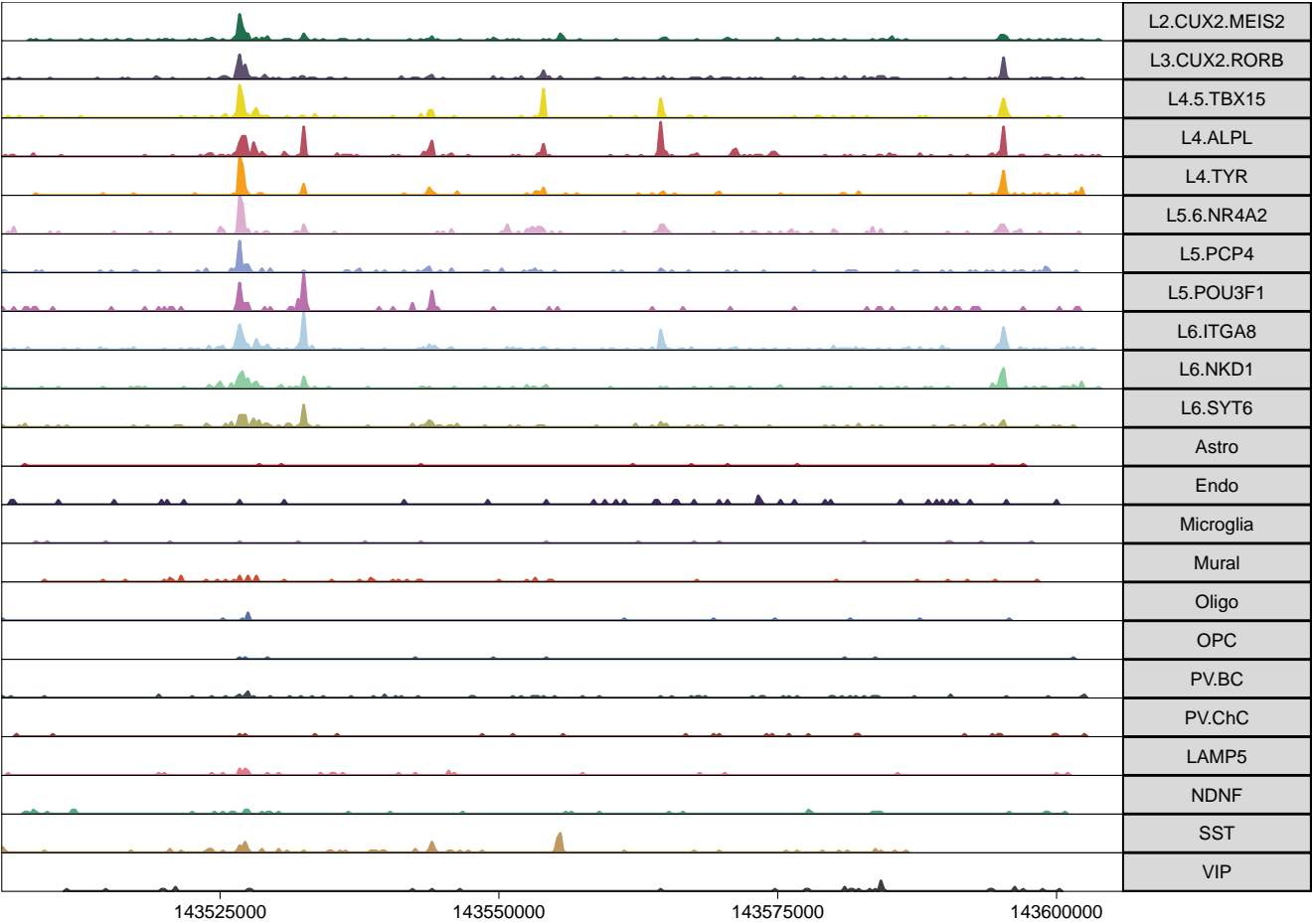


Peaks

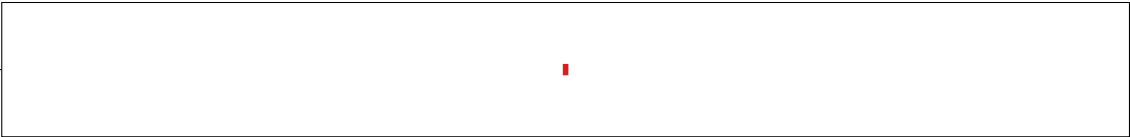


Genes

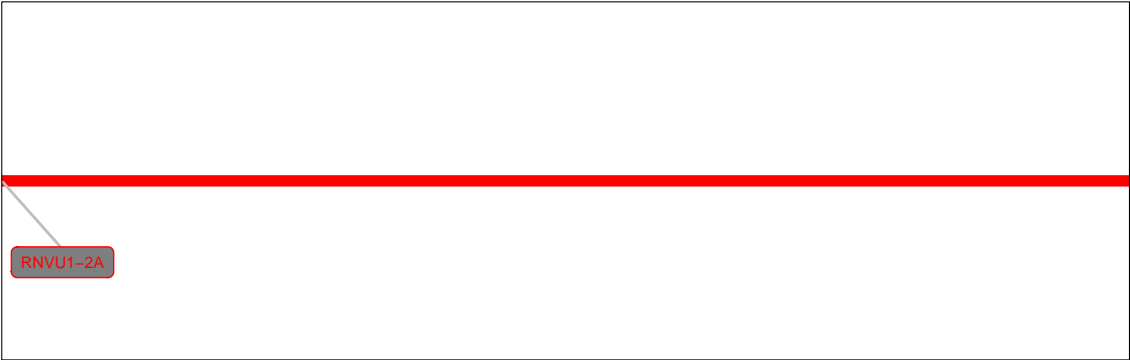
Coverage  
(Norm. ATAC Signal Range (0-0.7) by ReadsInTSS)



SST

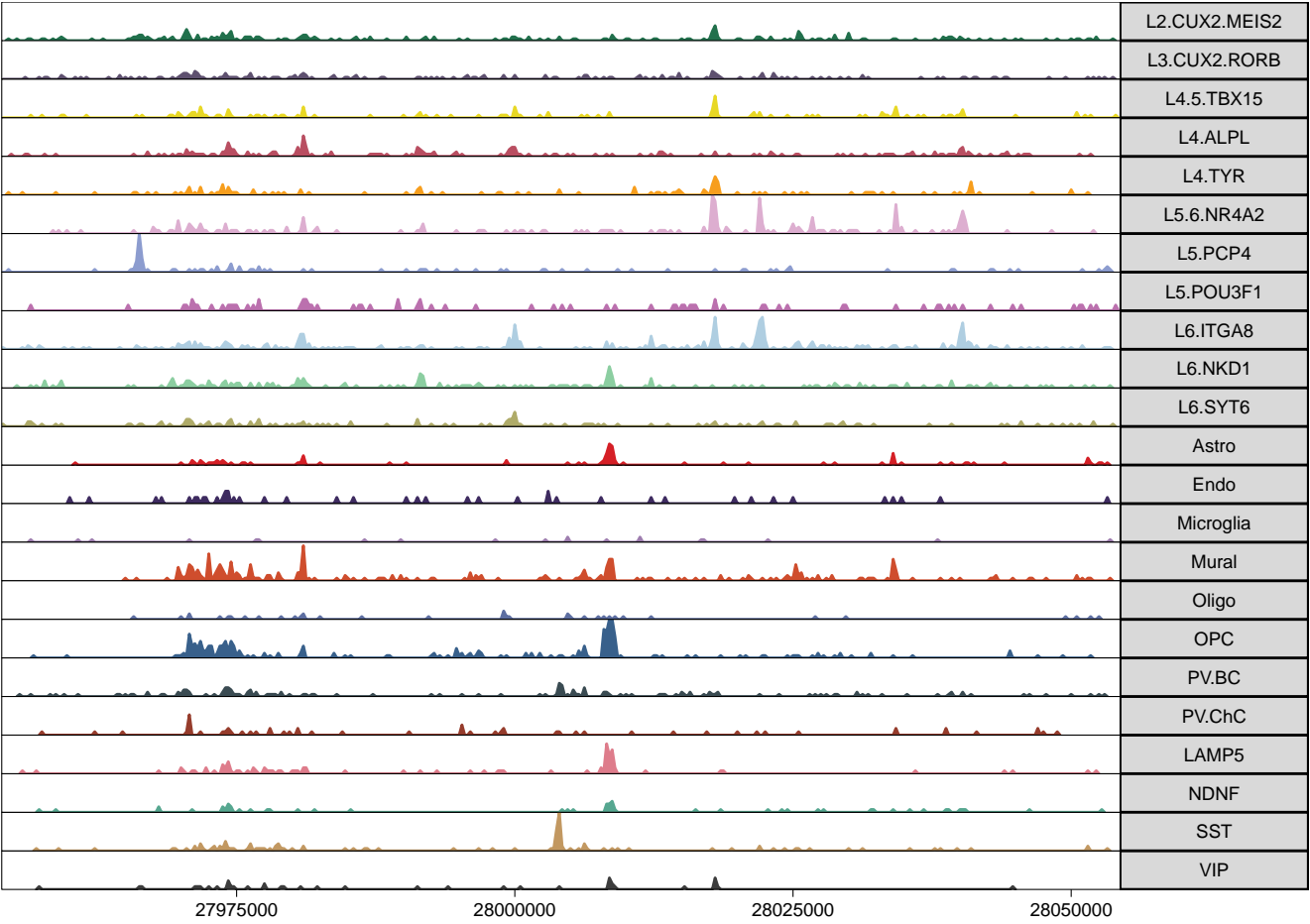


Peaks

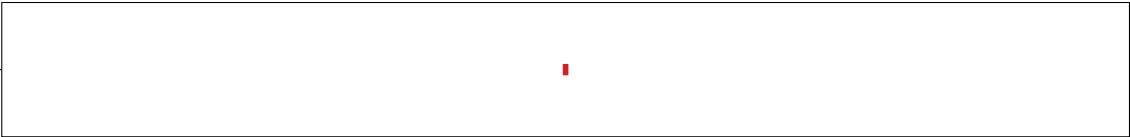


Genes

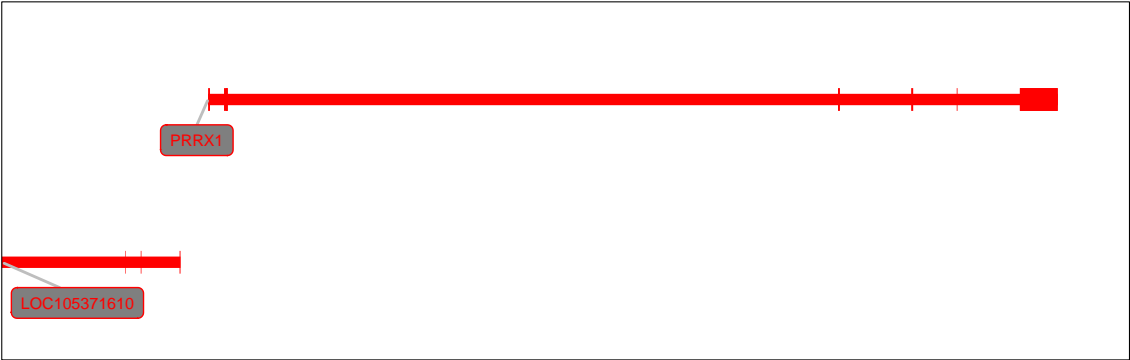
Coverage  
(Norm. ATAC Signal Range (0-0.5) by ReadsInTSS)



SST



Peaks

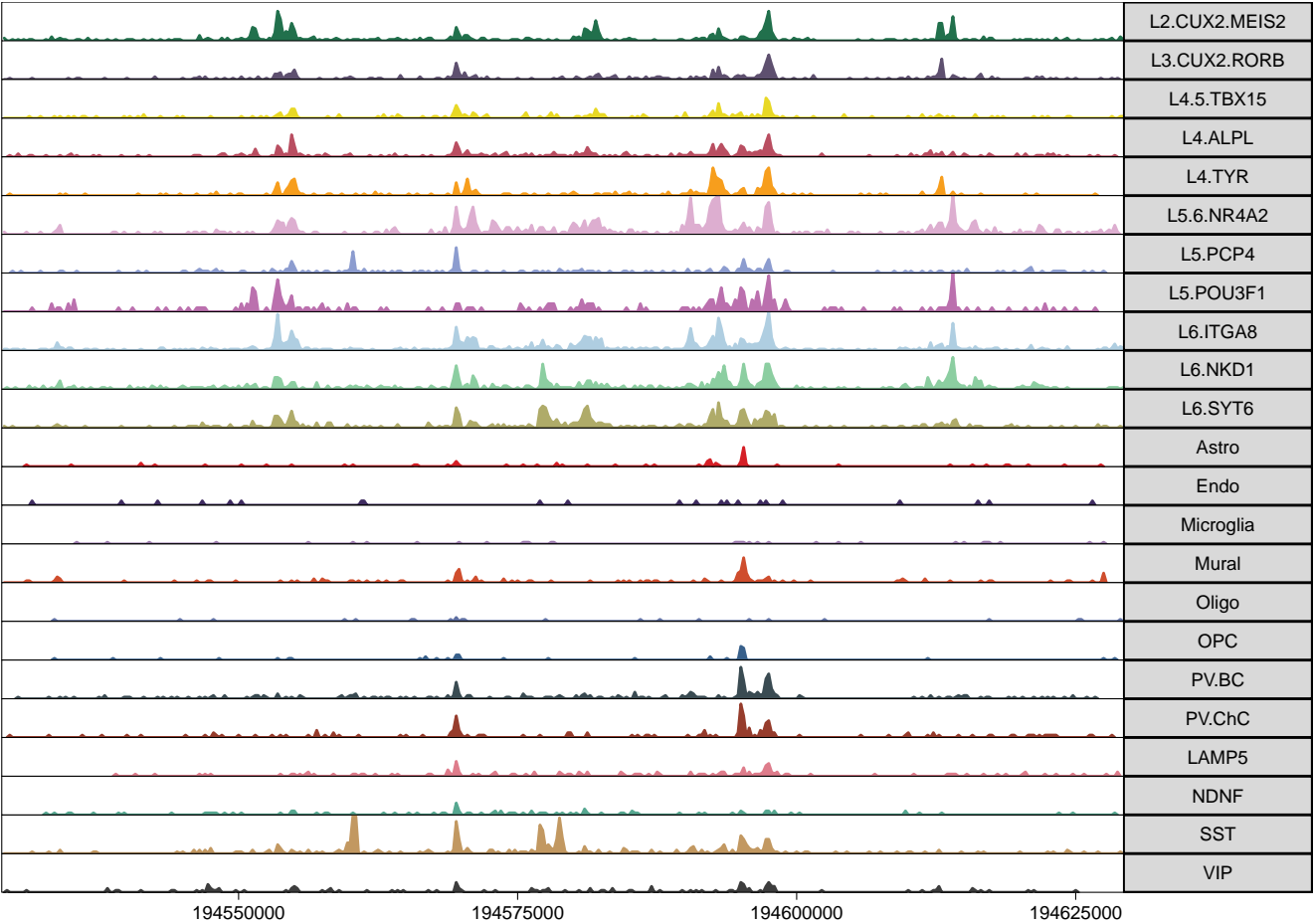


Genes

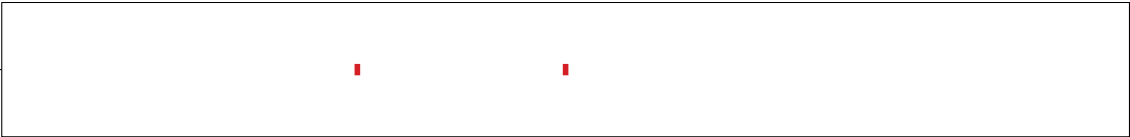


chr1:194528778–194629279 Marker Peak Tracks for: SST

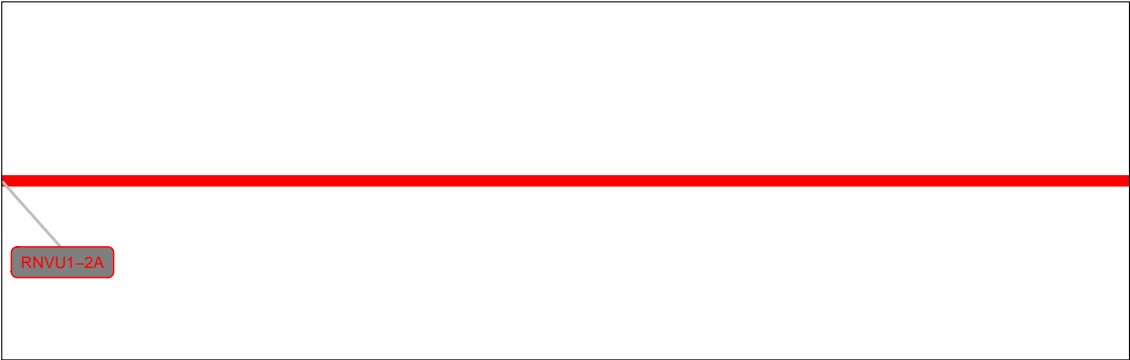
Coverage  
(Norm. ATAC Signal Range (0–0.71) by ReadsInTSS)



SST

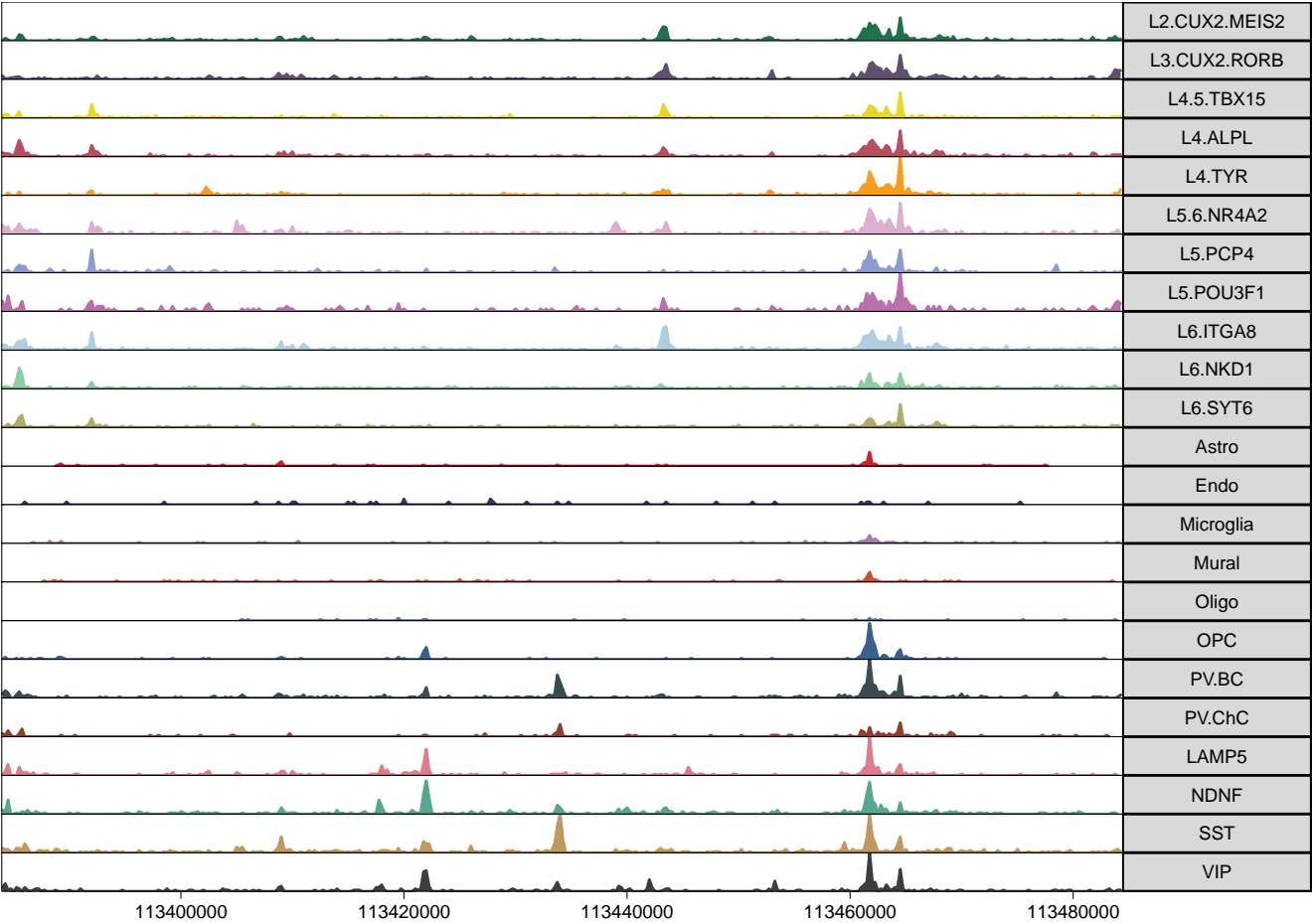


Peaks

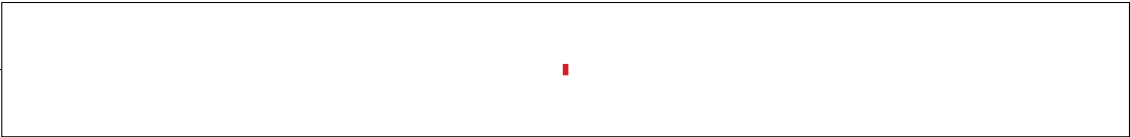


Genes

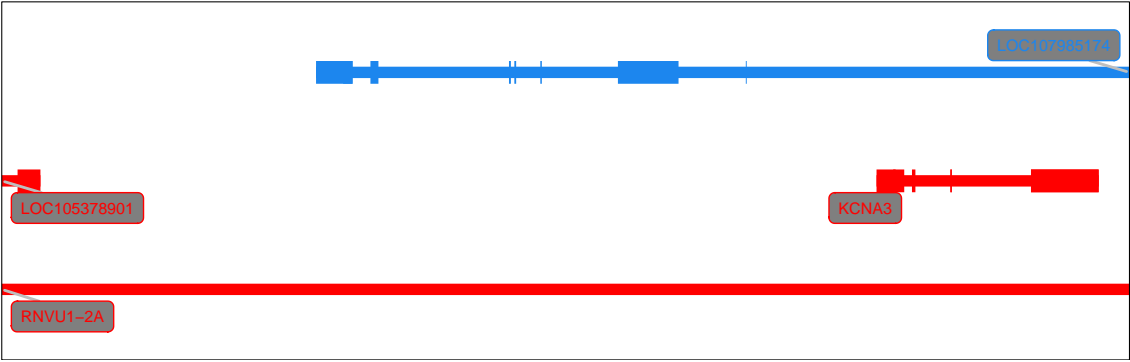
Coverage  
(Norm. ATAC Signal Range (0-1.09) by ReadsInTSS)



SST

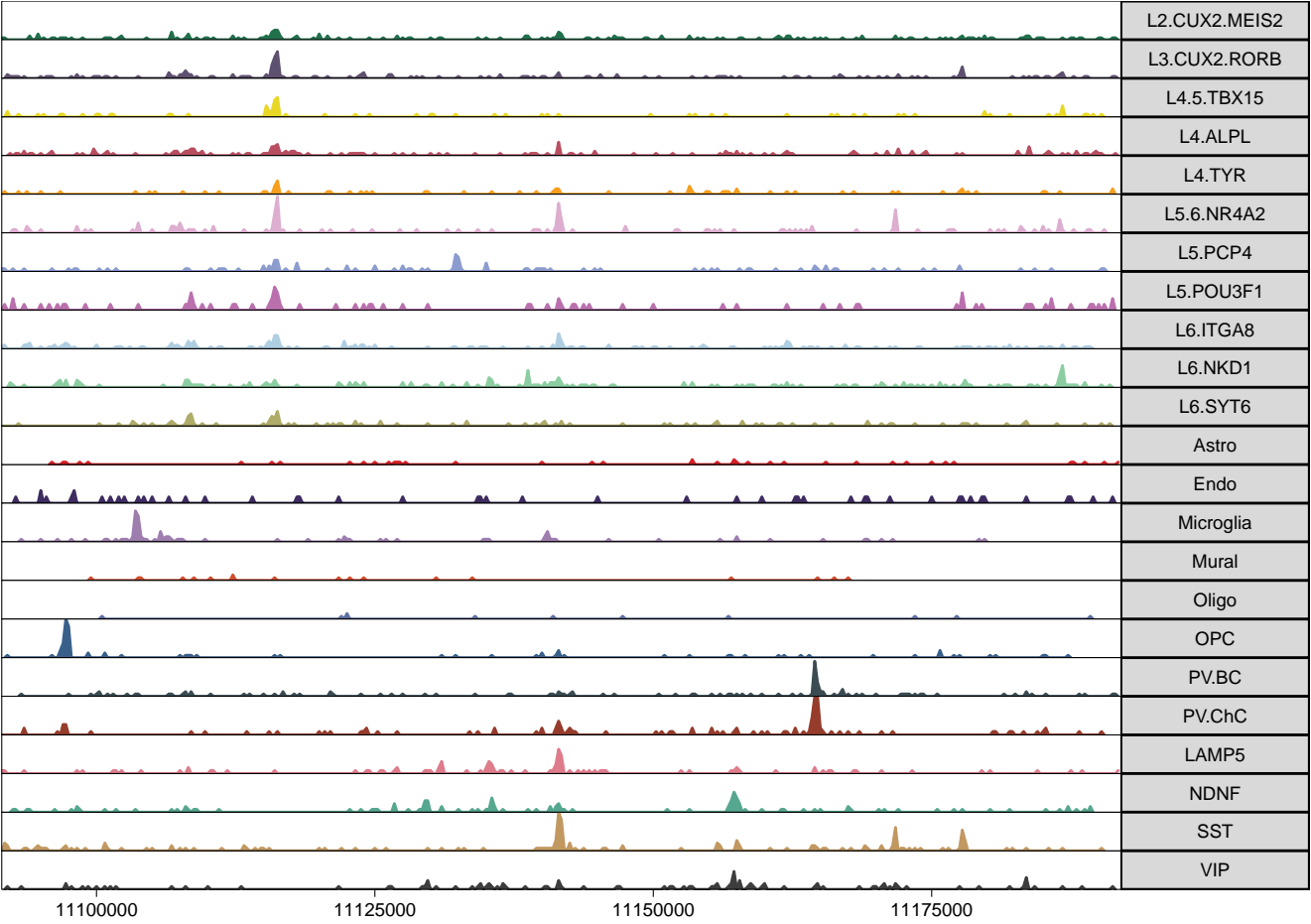


Peaks

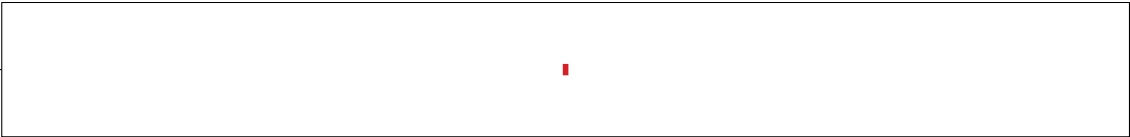


Genes

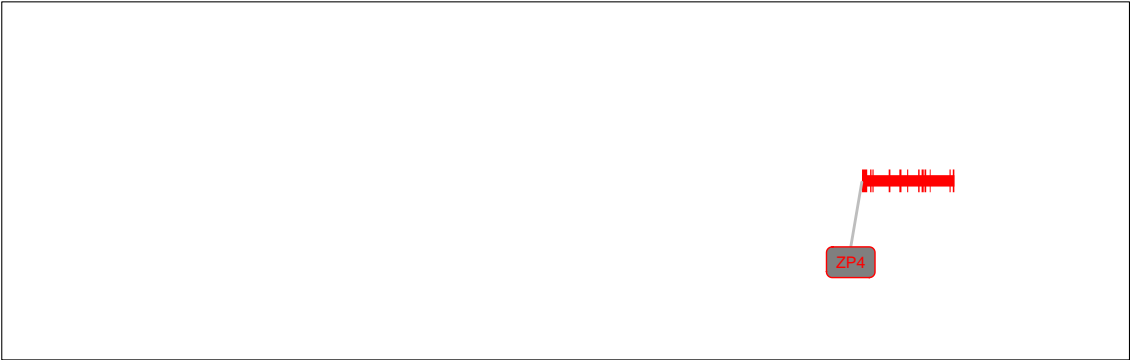
Coverage  
(Norm. ATAC Signal Range (0–0.49) by ReadsInTSS)



SST

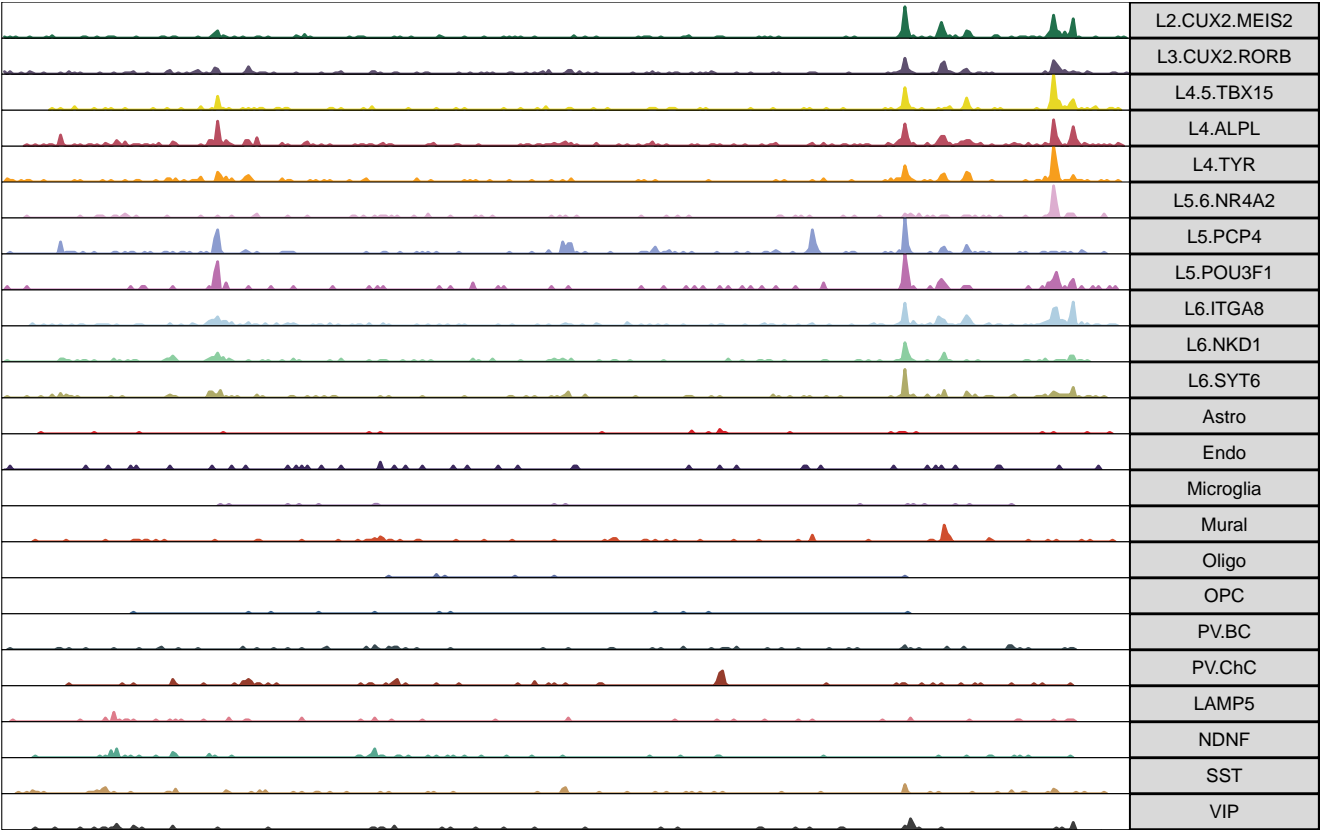


Peaks



Genes

Coverage  
(Norm. ATAC Signal Range (0–0.75) by ReadsInTSS)



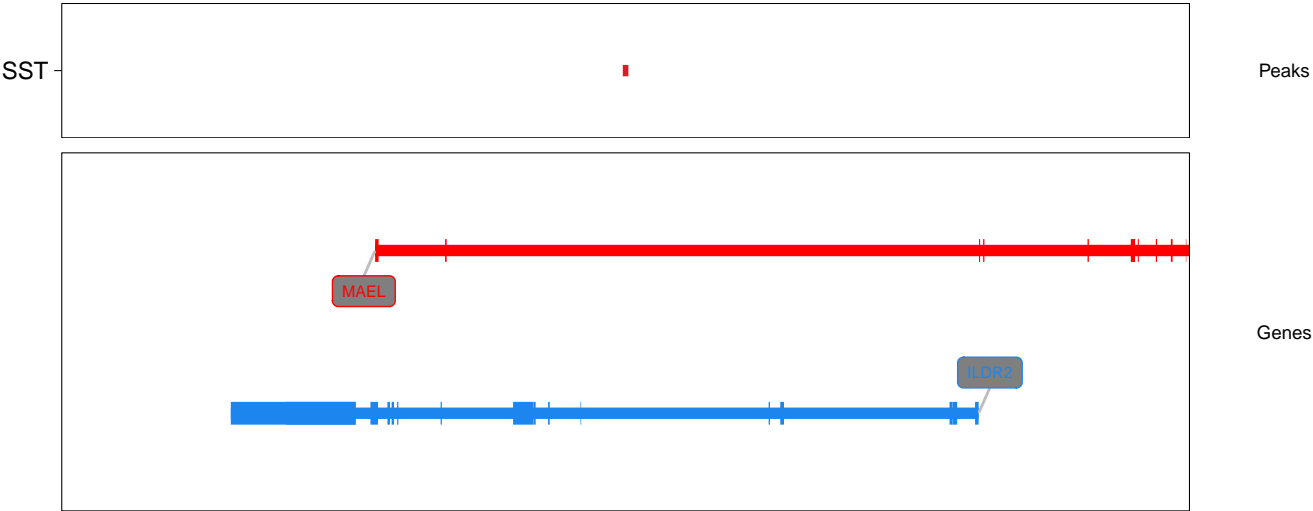
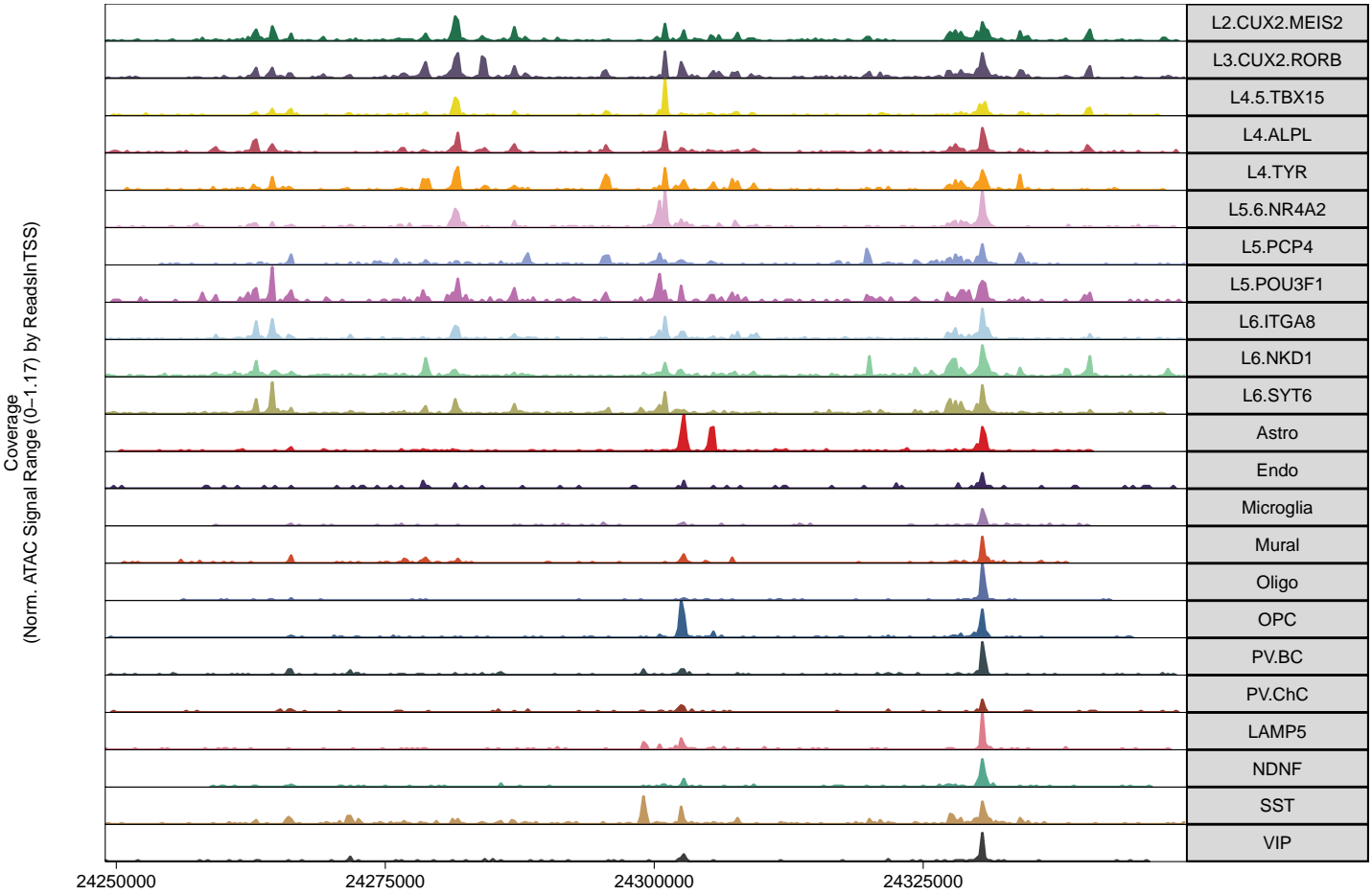
SST



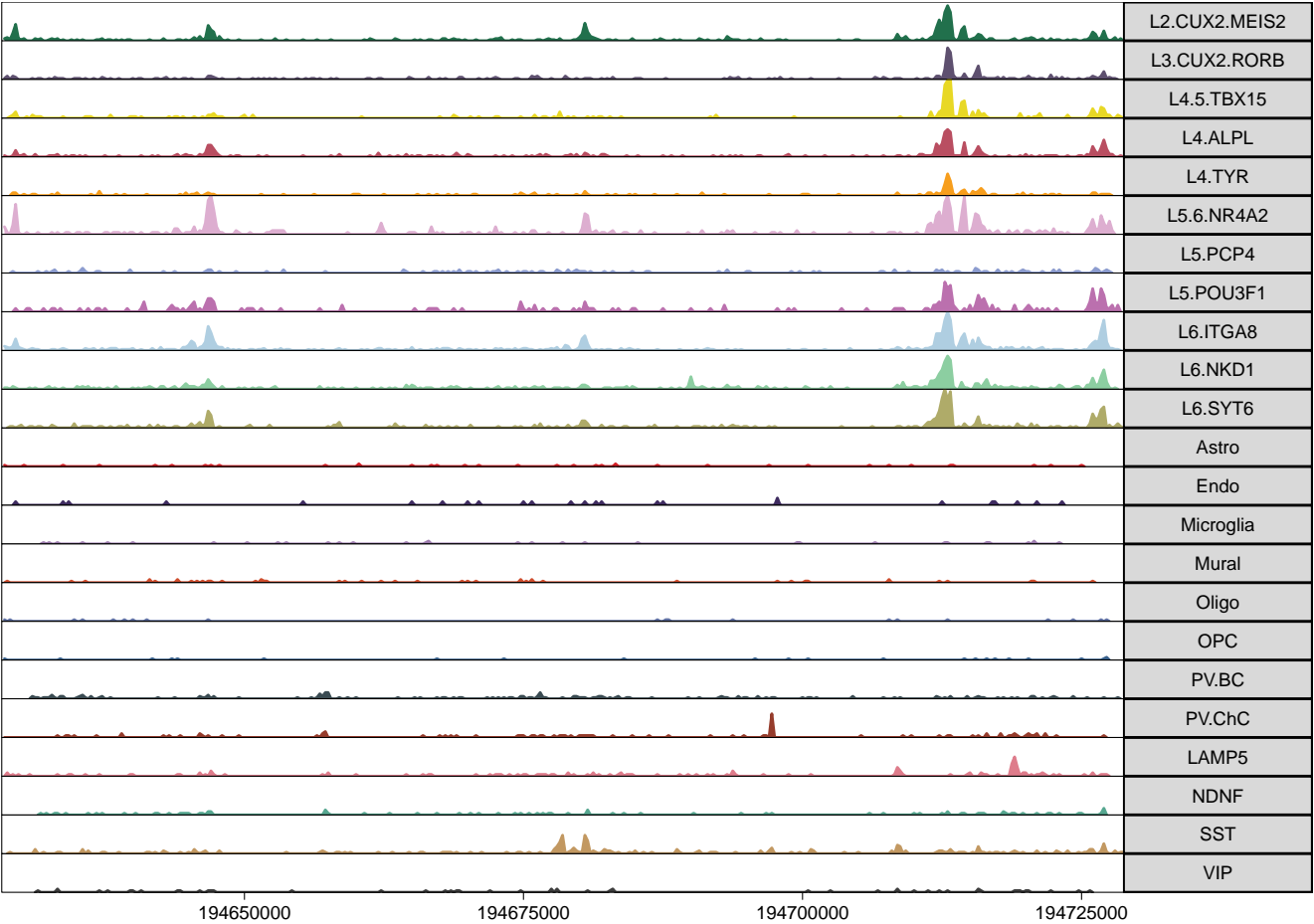
Peaks

GeneTrack

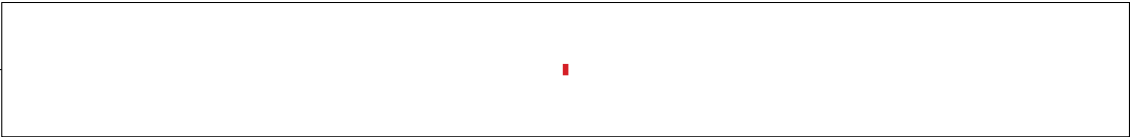
strand ● ·



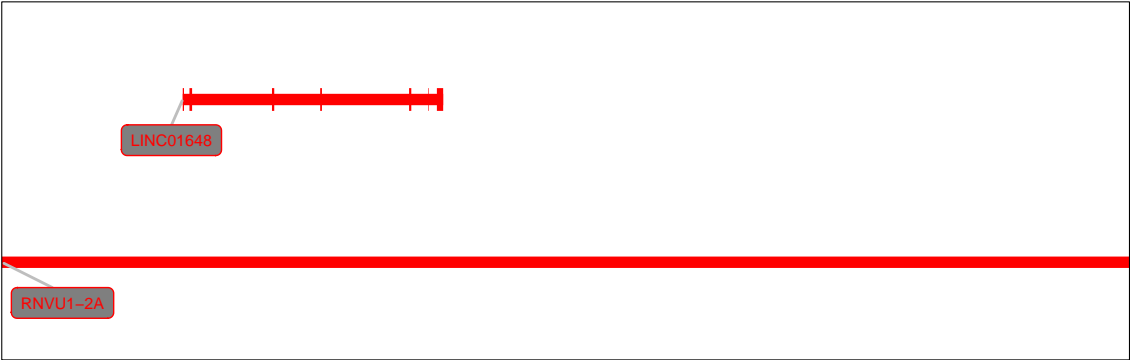
Coverage  
(Norm. ATAC Signal Range (0–0.86) by ReadsInTSS)



SST

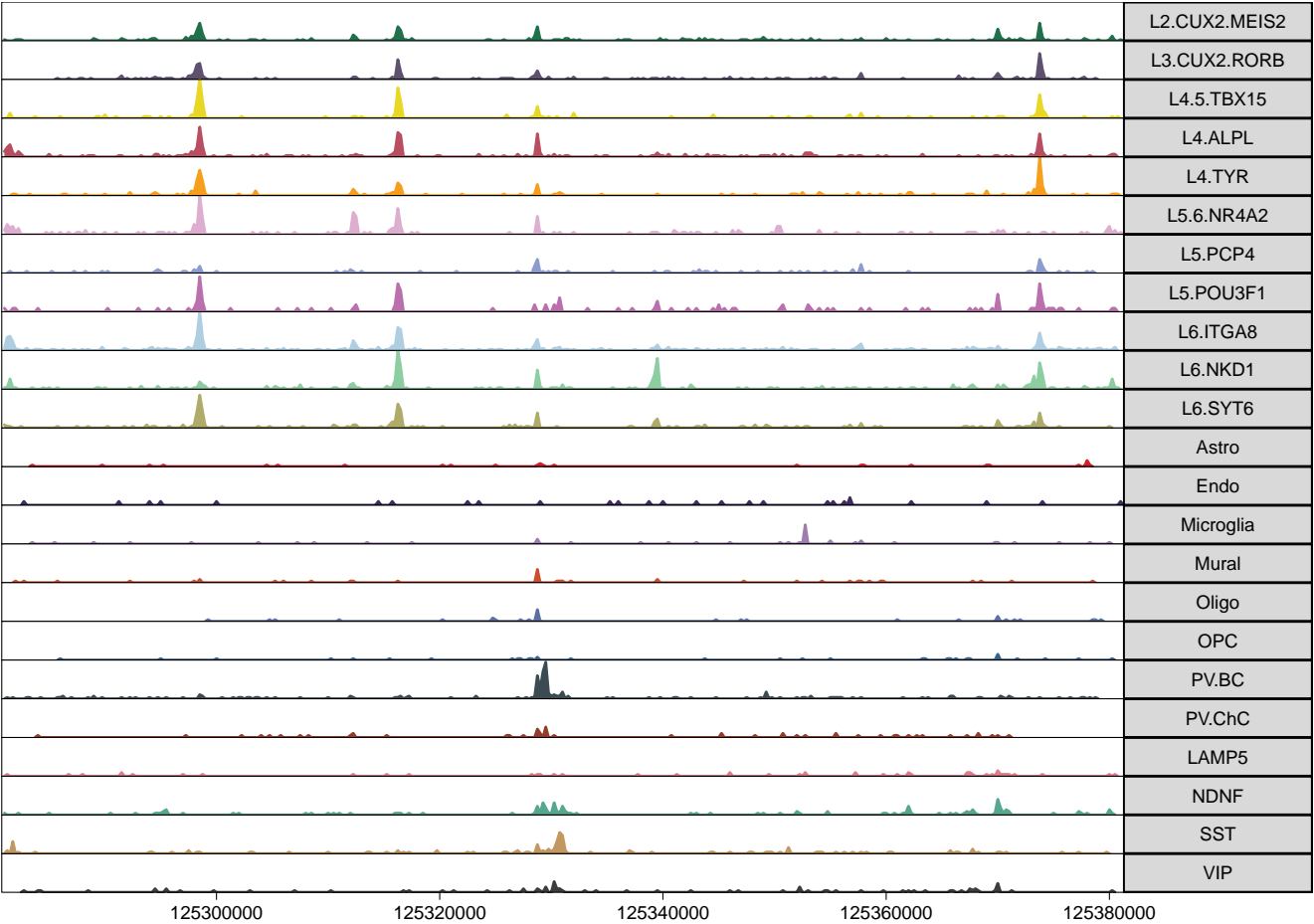


Peaks

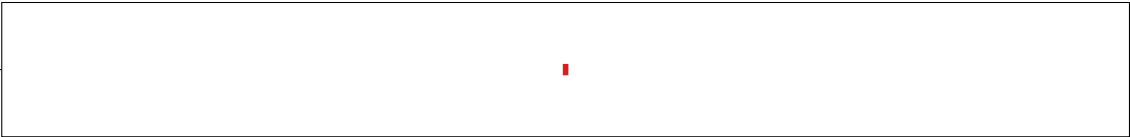


Genes

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



SST

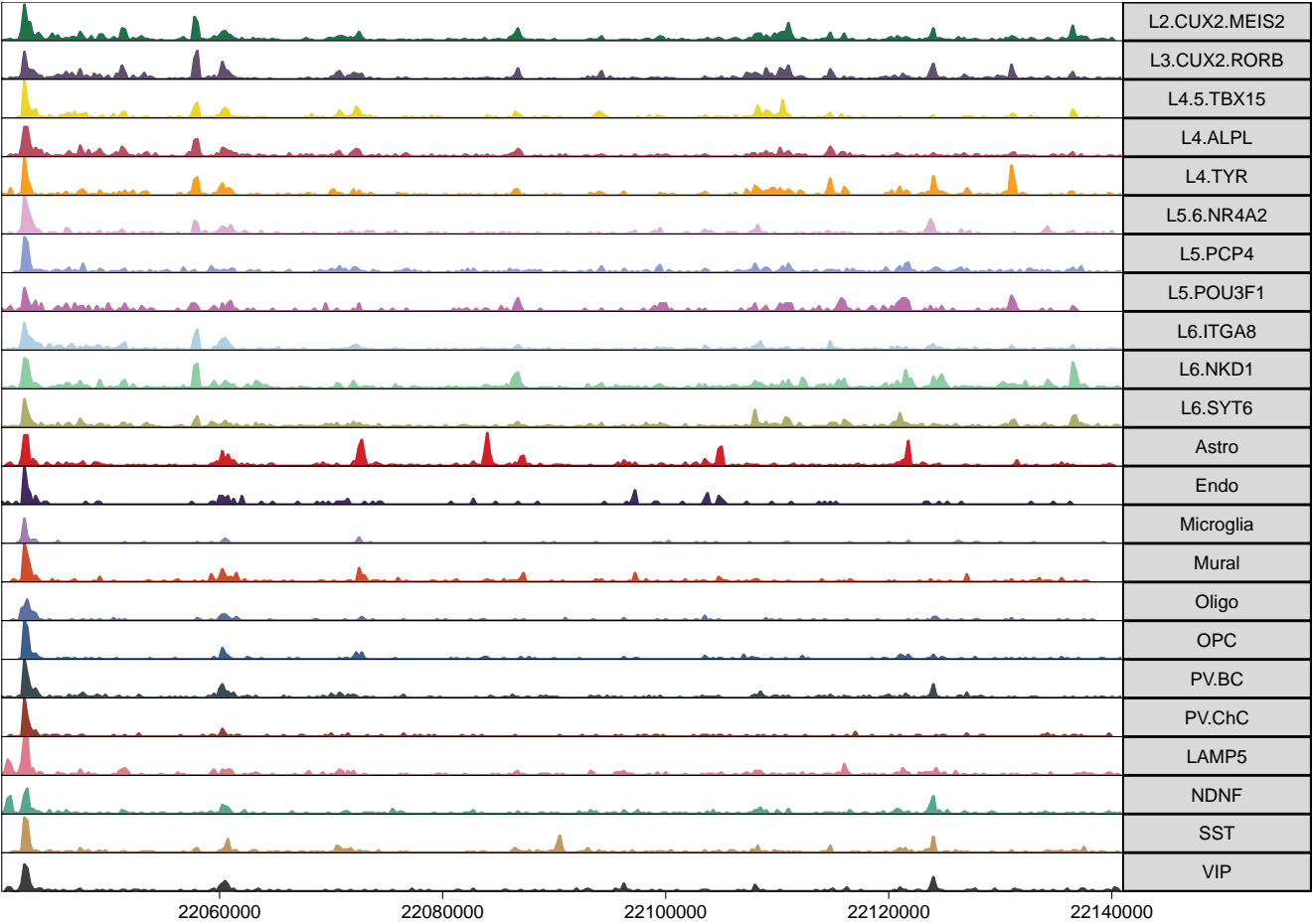


Peaks

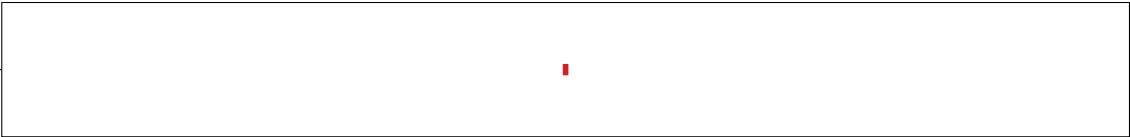


Genes

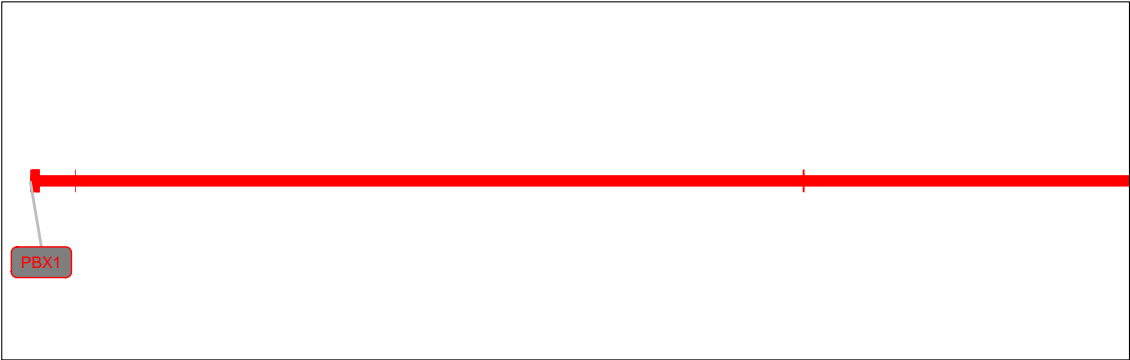
Coverage  
(Norm. ATAC Signal Range (0–1.09) by ReadsInTSS)



SST

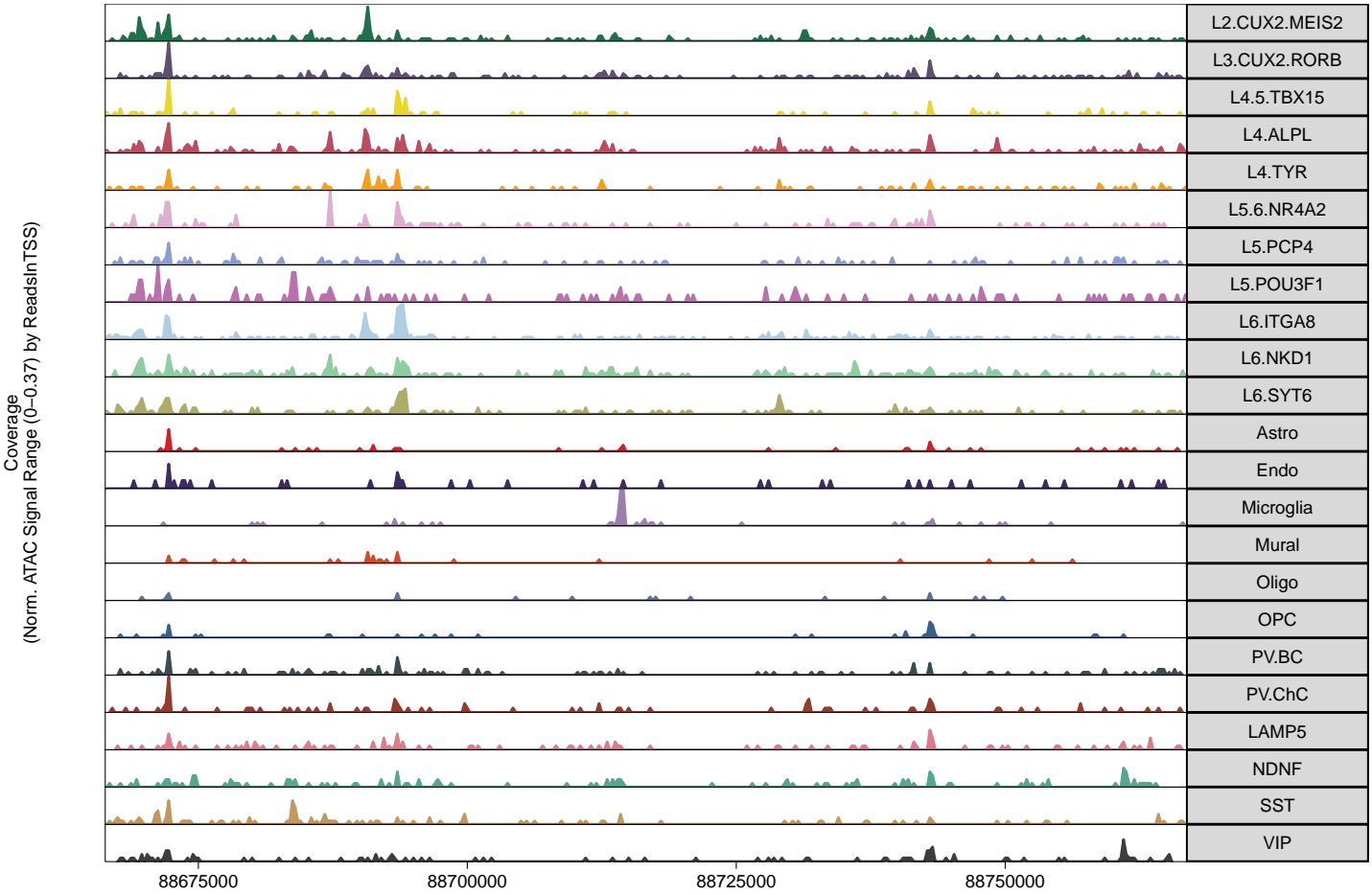


Peaks

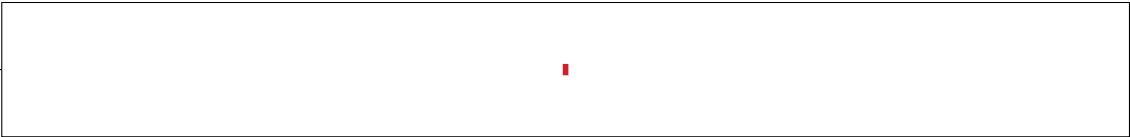


Genes

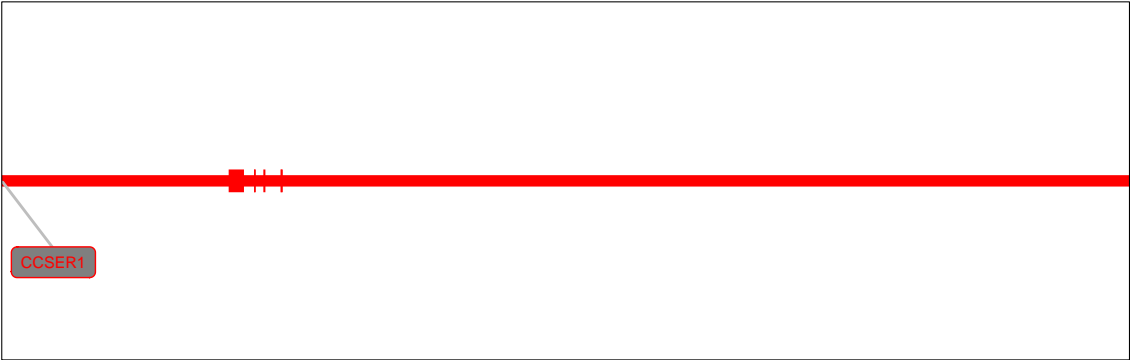




SST

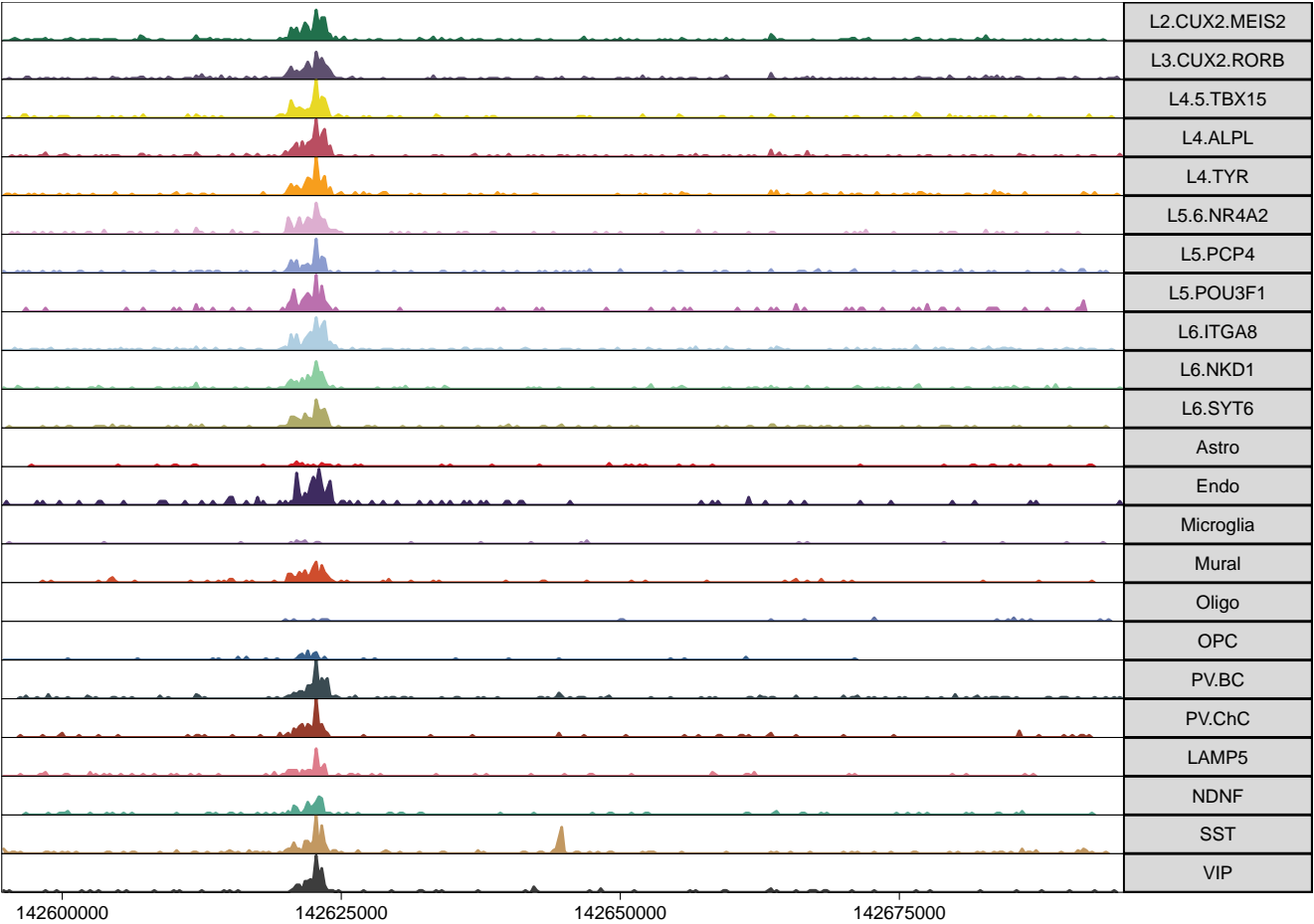


Peaks

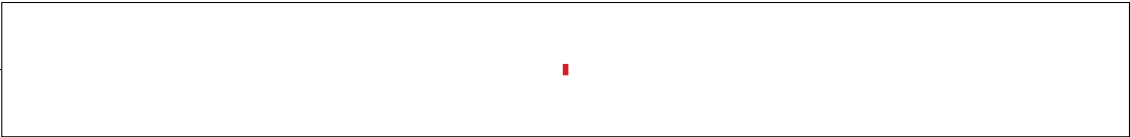


Genes

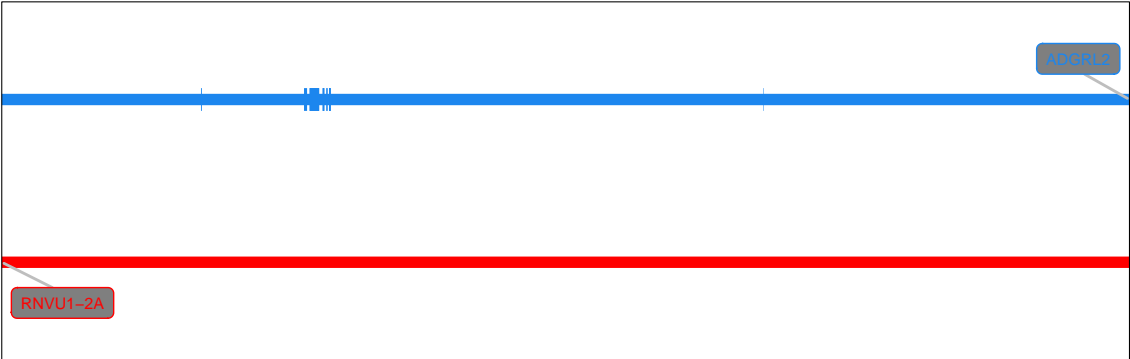
Coverage  
(Norm. ATAC Signal Range (0–0.77) by ReadsInTSS)



SST



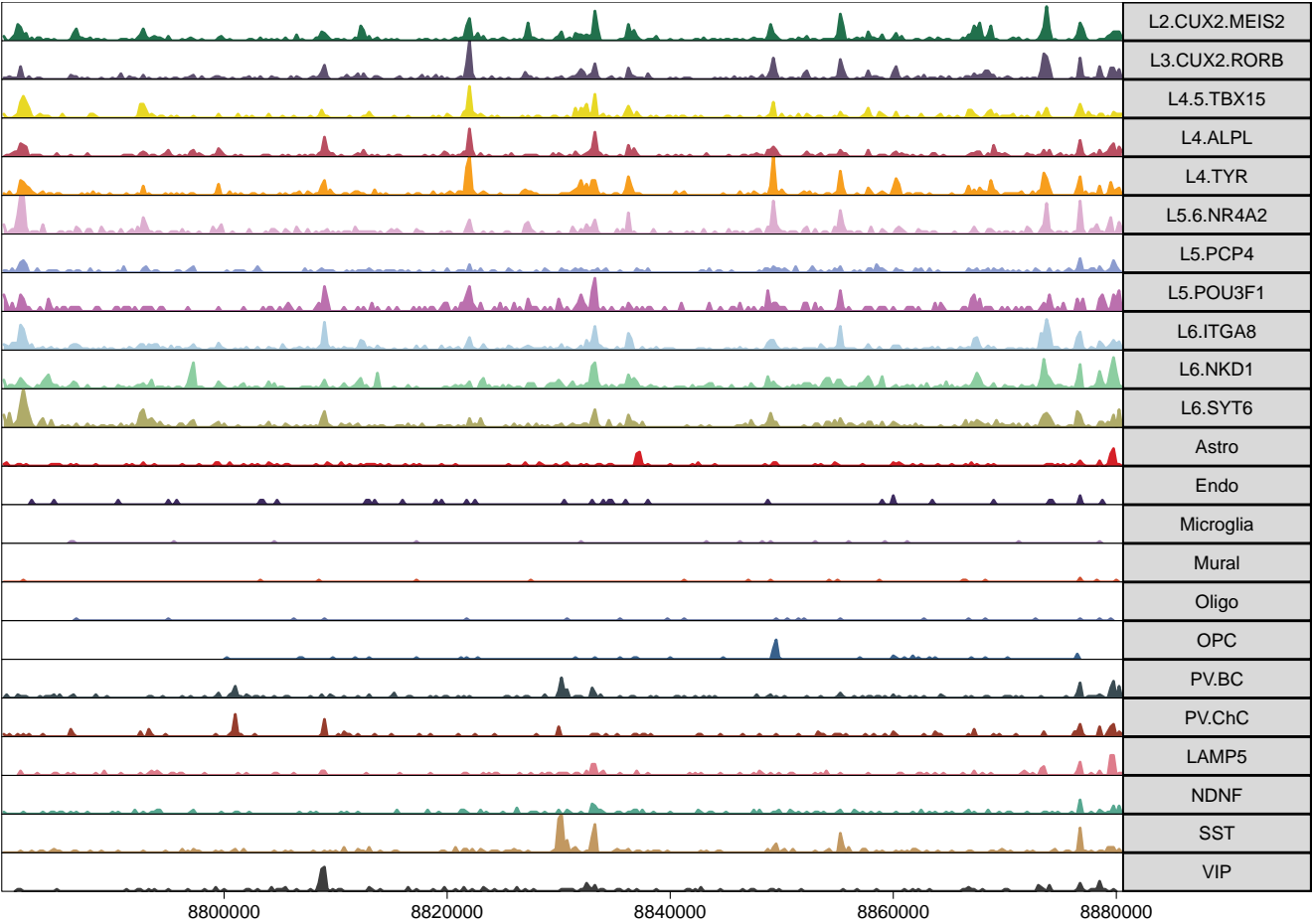
Peaks



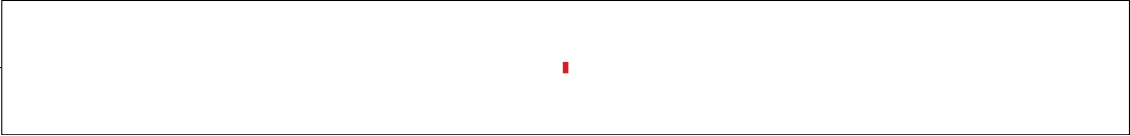
Genes

chr1:8780061–8880562 Marker Peak Tracks for: SST

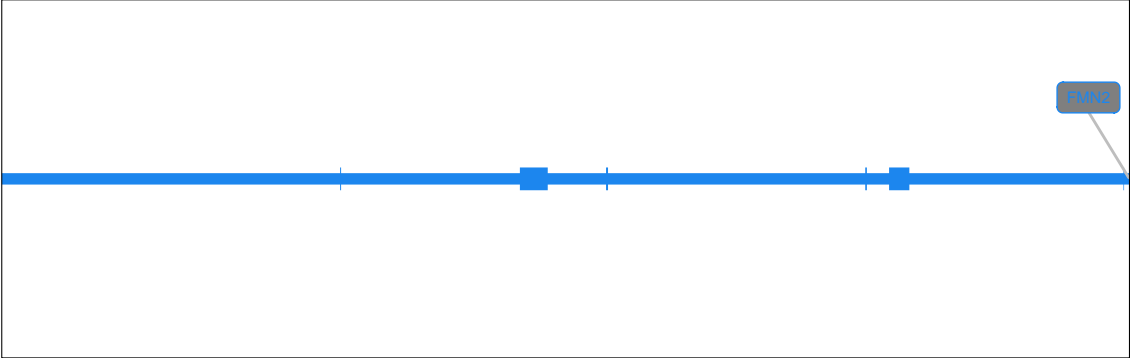
Coverage  
(Norm. ATAC Signal Range (0–0.68) by ReadsInTSS)



SST

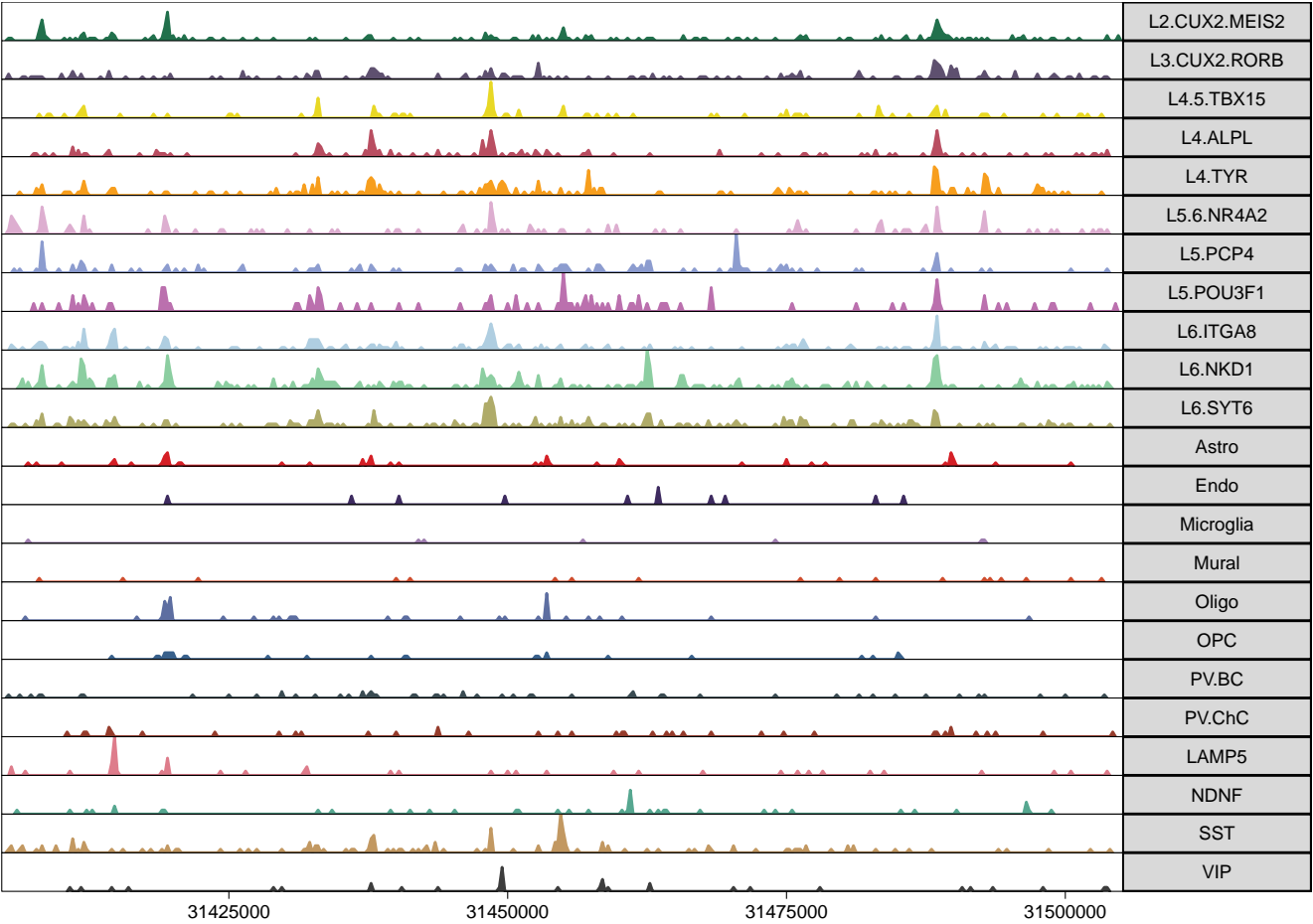


Peaks

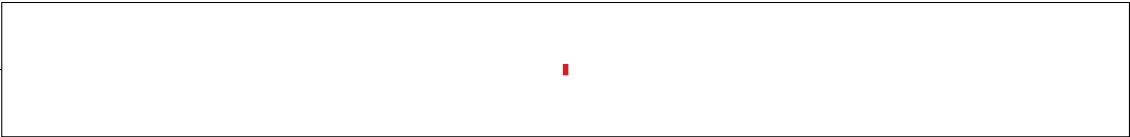


Genes

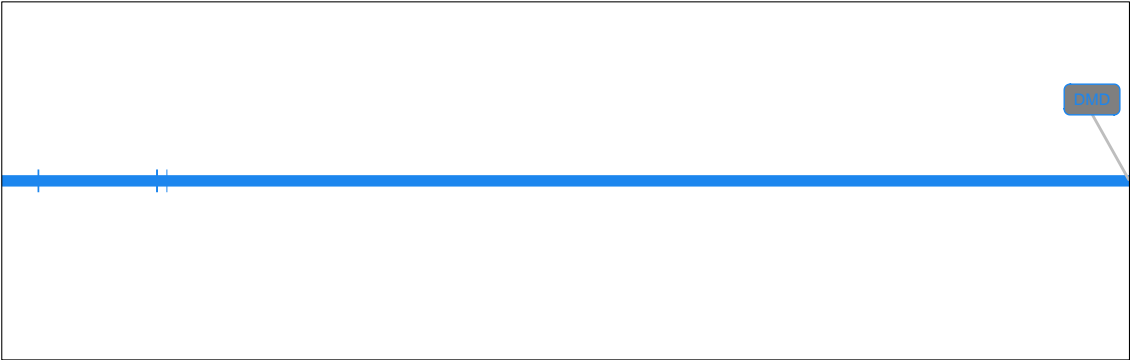
Coverage  
(Norm. ATAC Signal Range (0–0.36) by ReadsInTSS)



SST

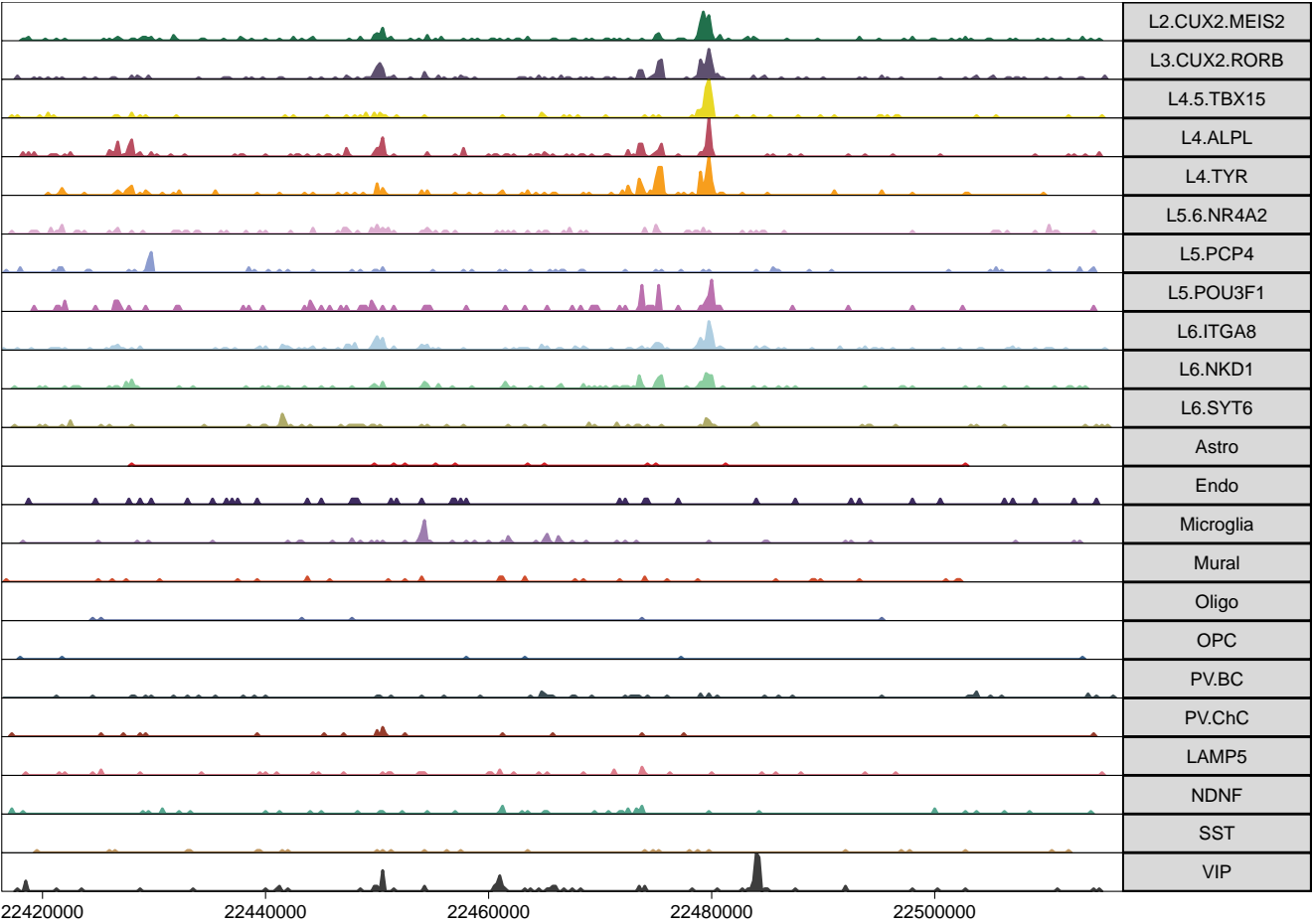


Peaks

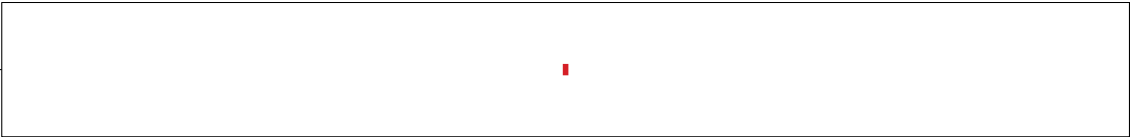


Genes

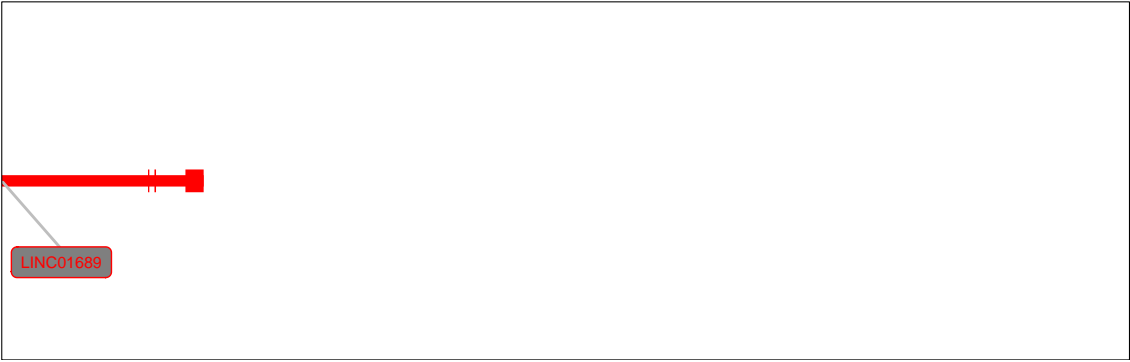
Coverage  
(Norm. ATAC Signal Range (0–0.54) by ReadsInTSS)



SST

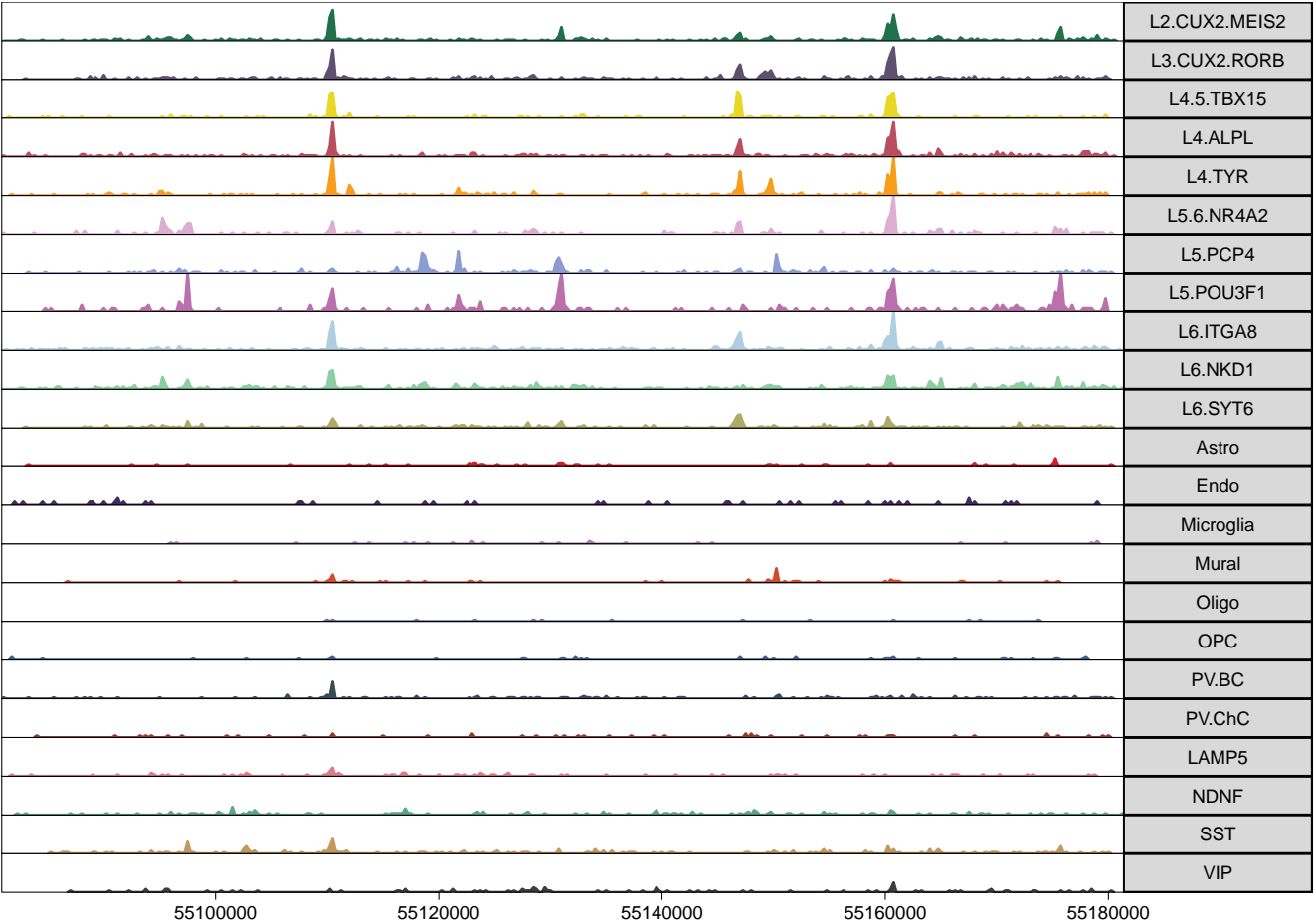


Peaks

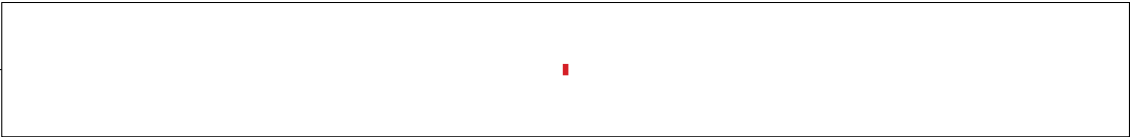


Genes

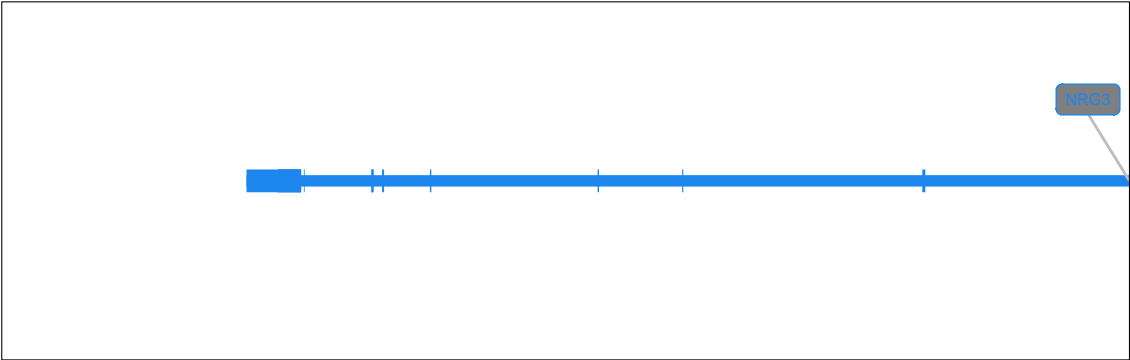
Coverage  
(Norm. ATAC Signal Range (0–0.88) by ReadsInTSS)



SST

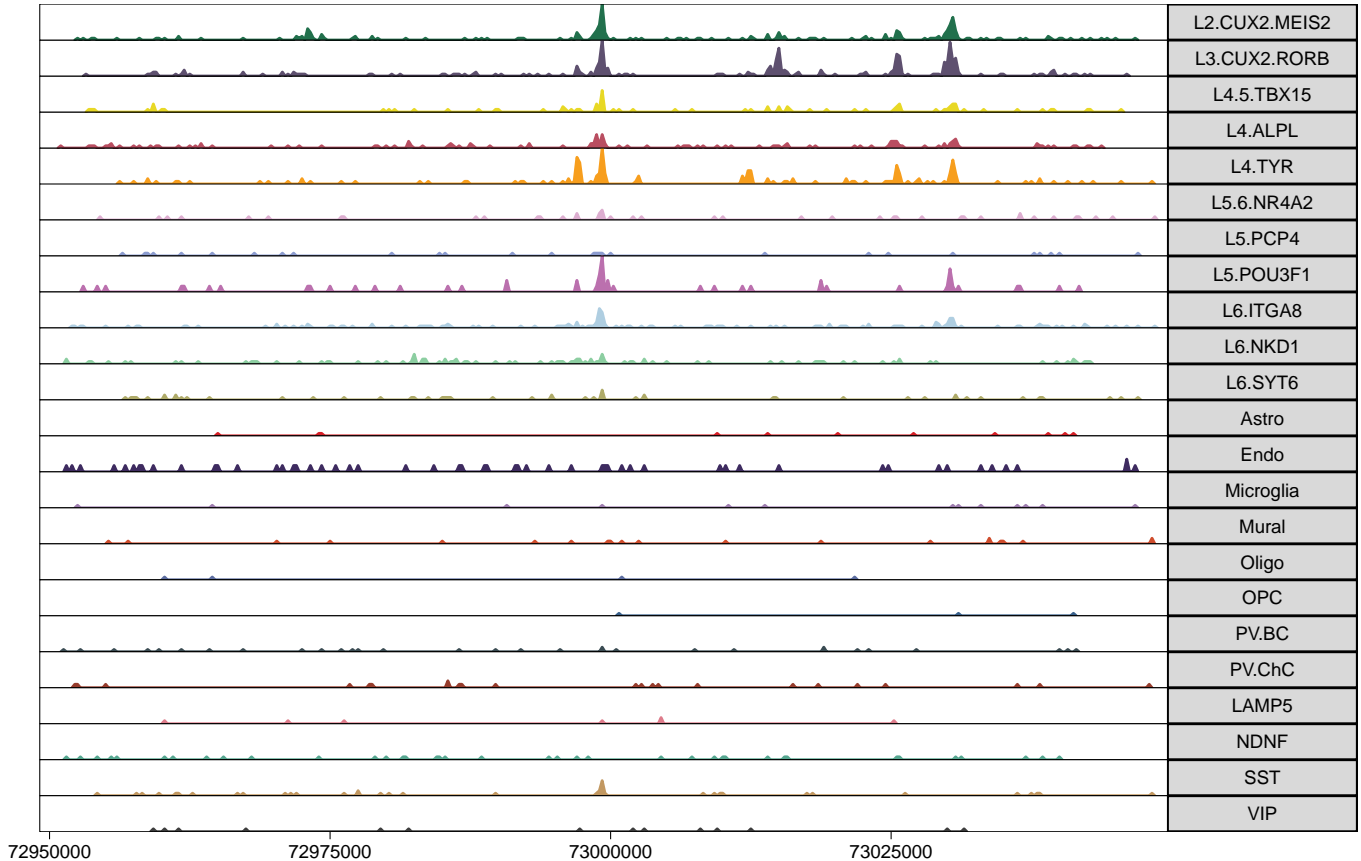


Peaks

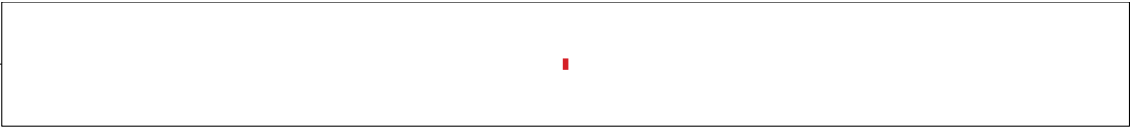


Genes

Coverage  
(Norm. ATAC Signal Range (0-0.46) by ReadsInTSS)



SST

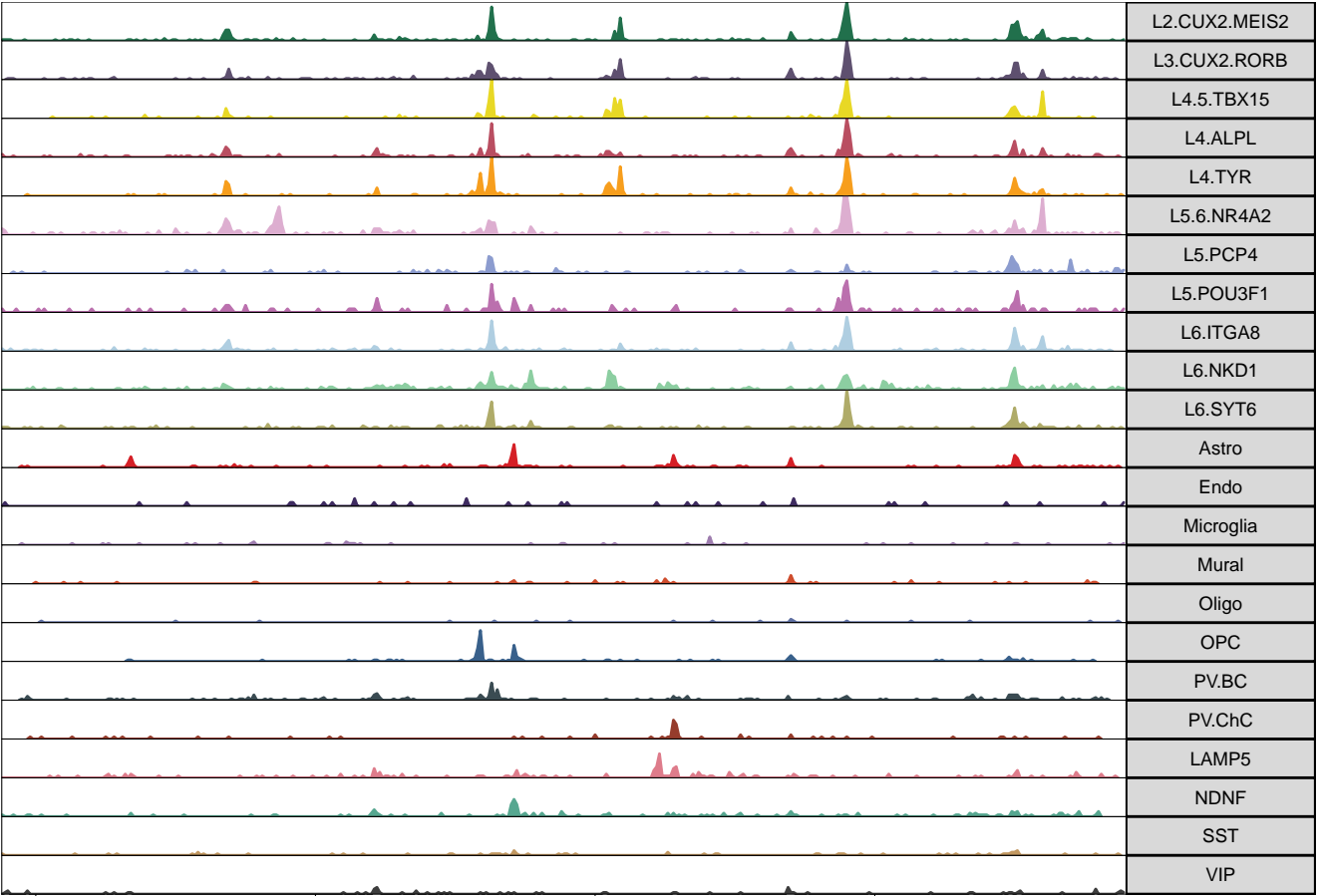


Peaks

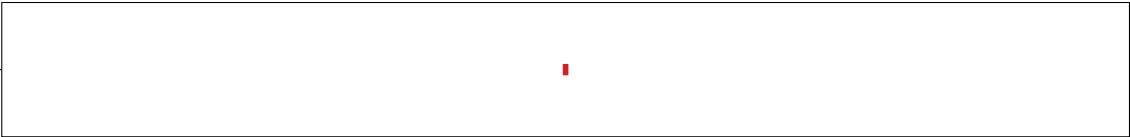
GeneTrack

strand    ●    ·

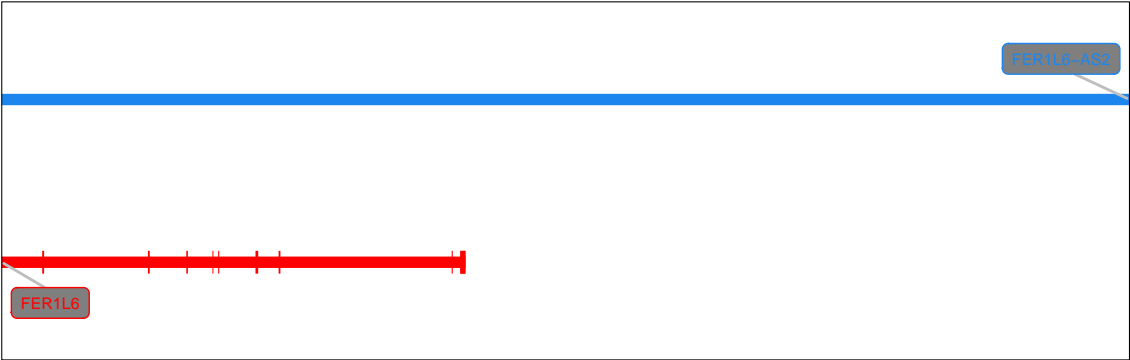
Coverage  
(Norm. ATAC Signal Range (0–0.81) by ReadsInTSS)



SST



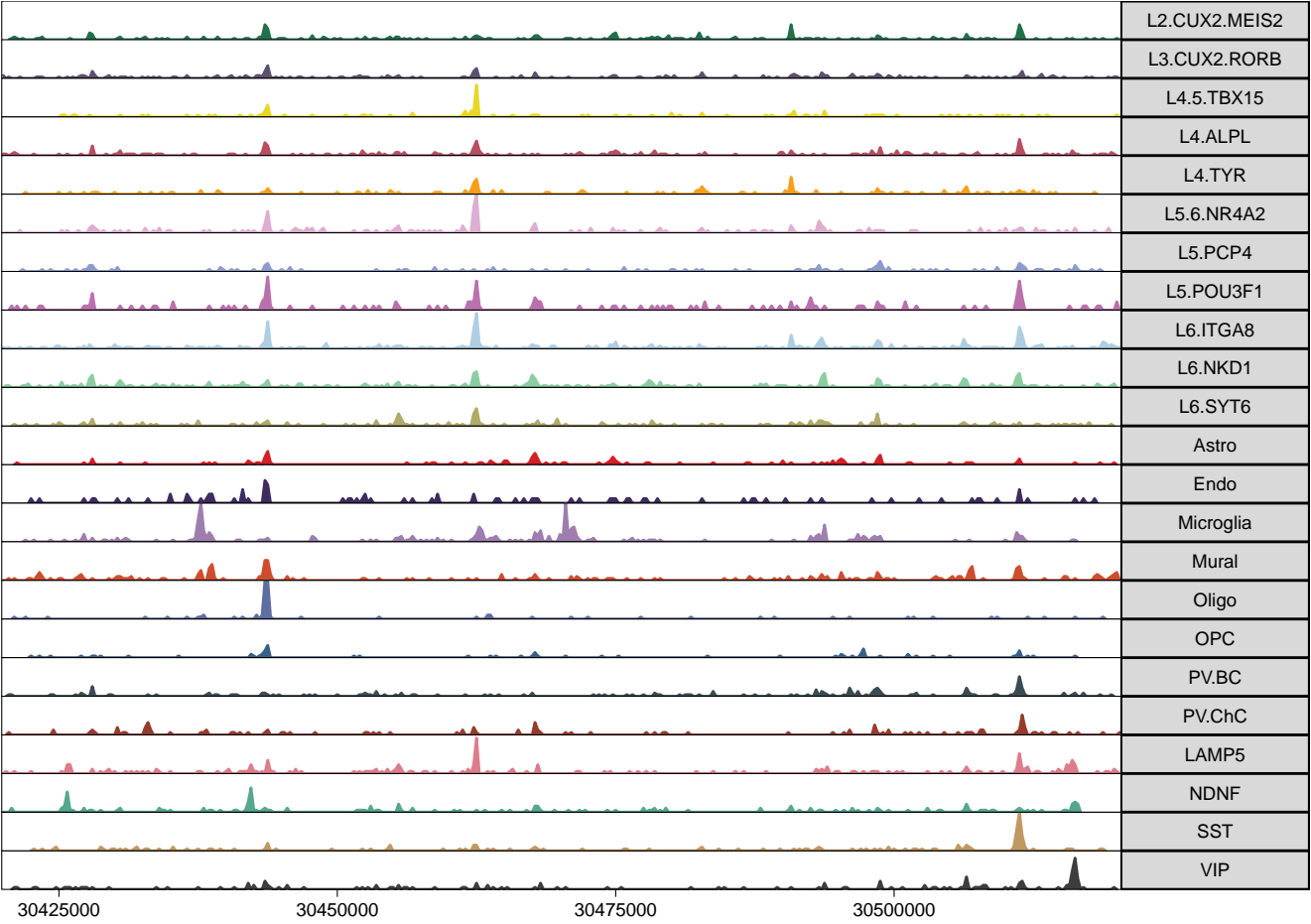
Peaks



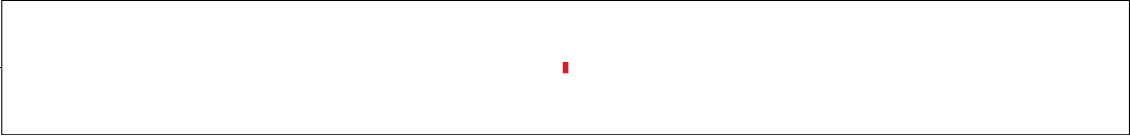
Genes



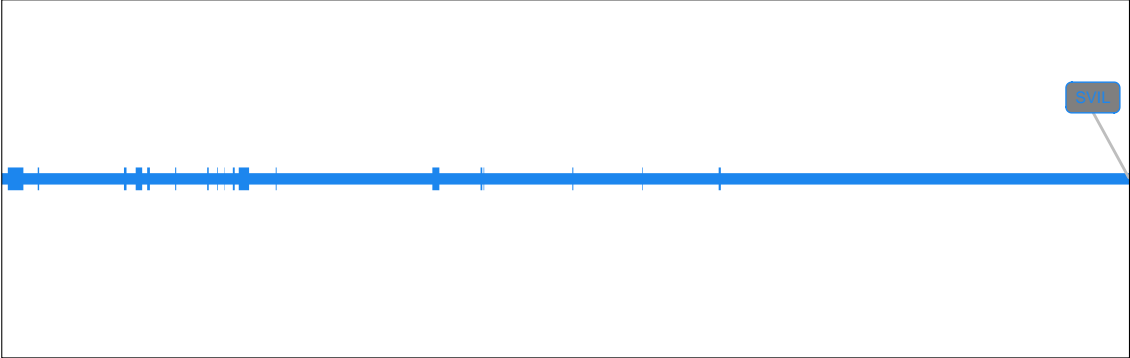
Coverage  
(Norm. ATAC Signal Range (0–0.68) by ReadsInTSS)



SST

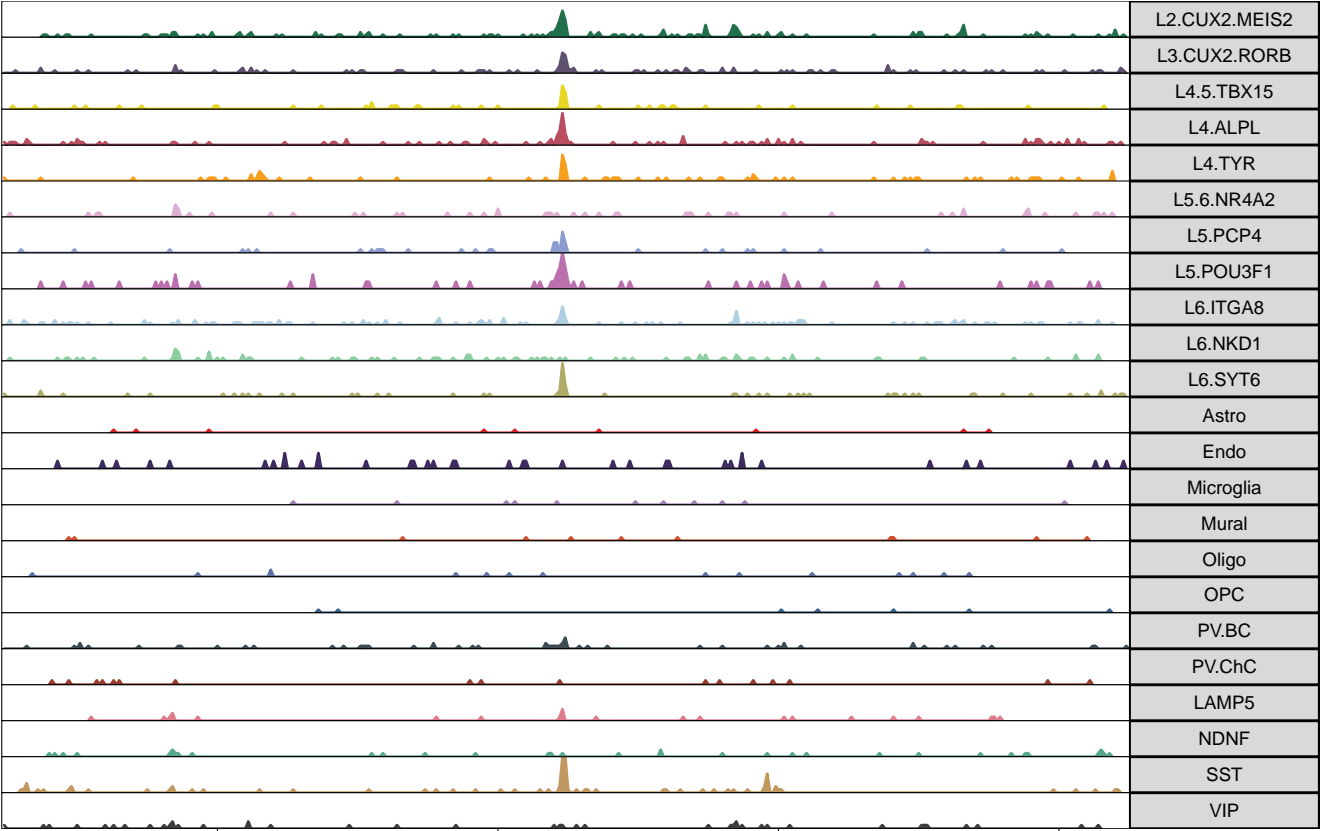


Peaks

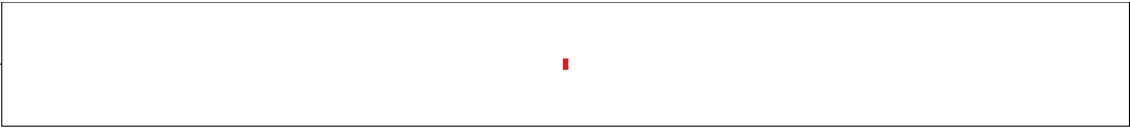


Genes

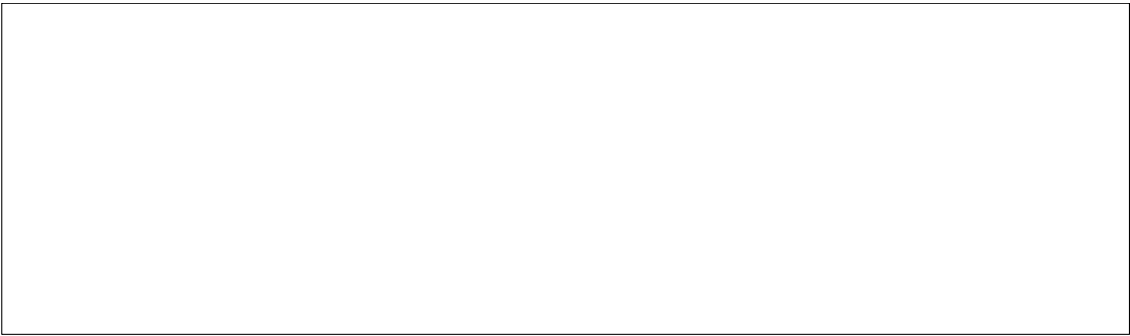
Coverage  
(Norm. ATAC Signal Range (0-0.37) by ReadsInTSS)



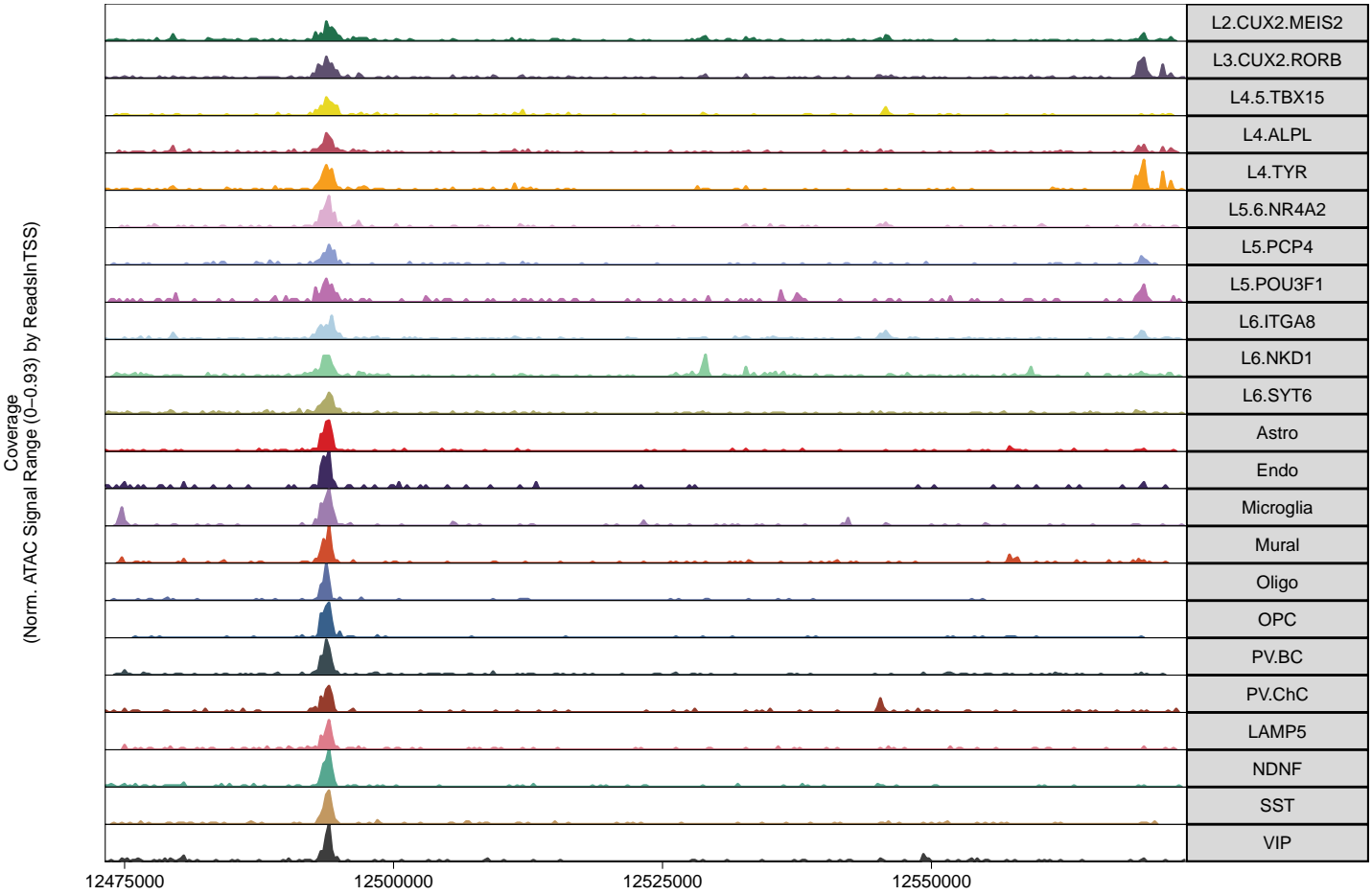
SST



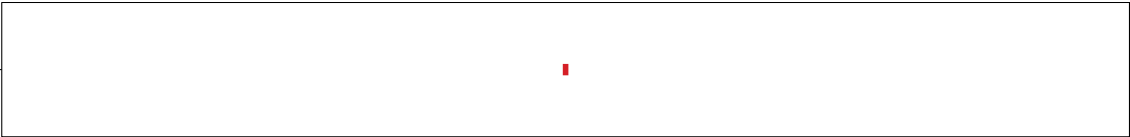
Peaks



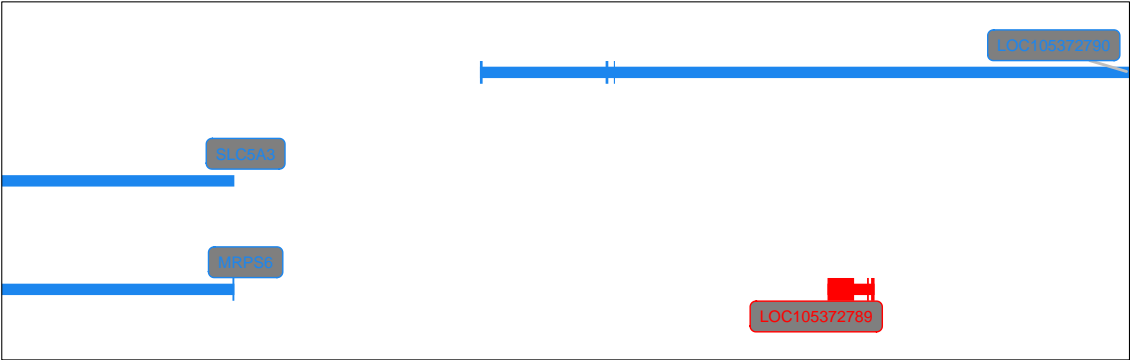
GeneTrack



SST

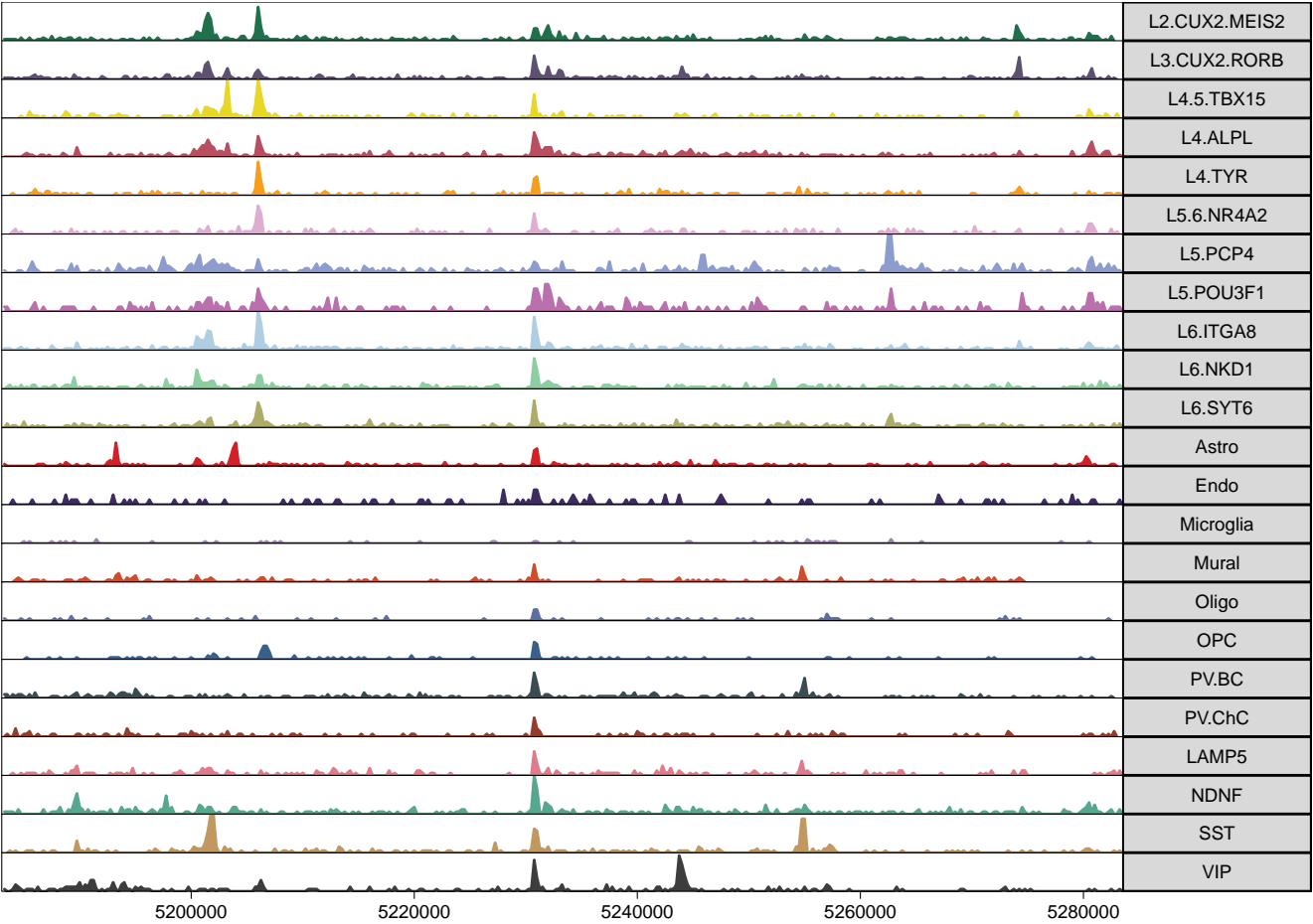


Peaks

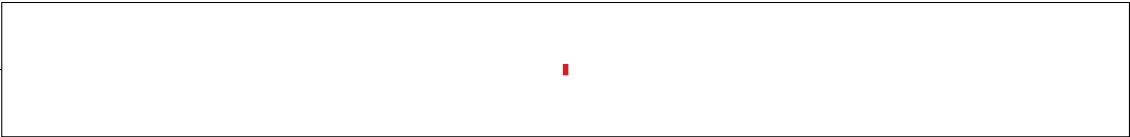


Genes

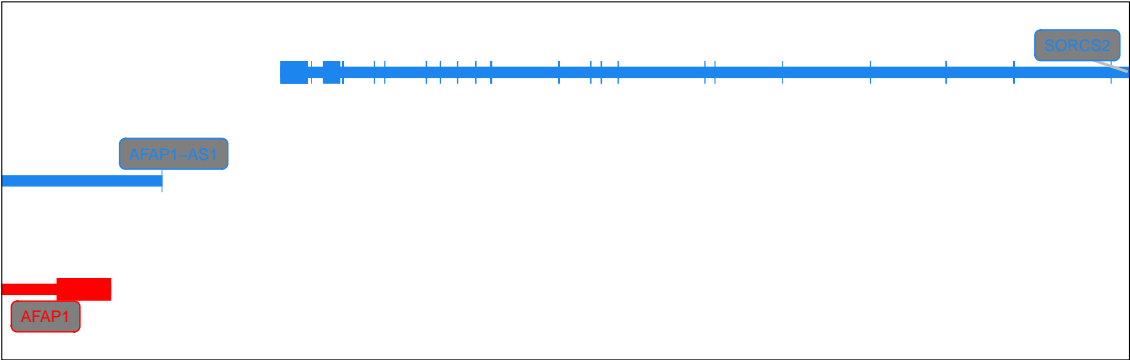
Coverage  
(Norm. ATAC Signal Range (0–0.63) by ReadsInTSS)



SST



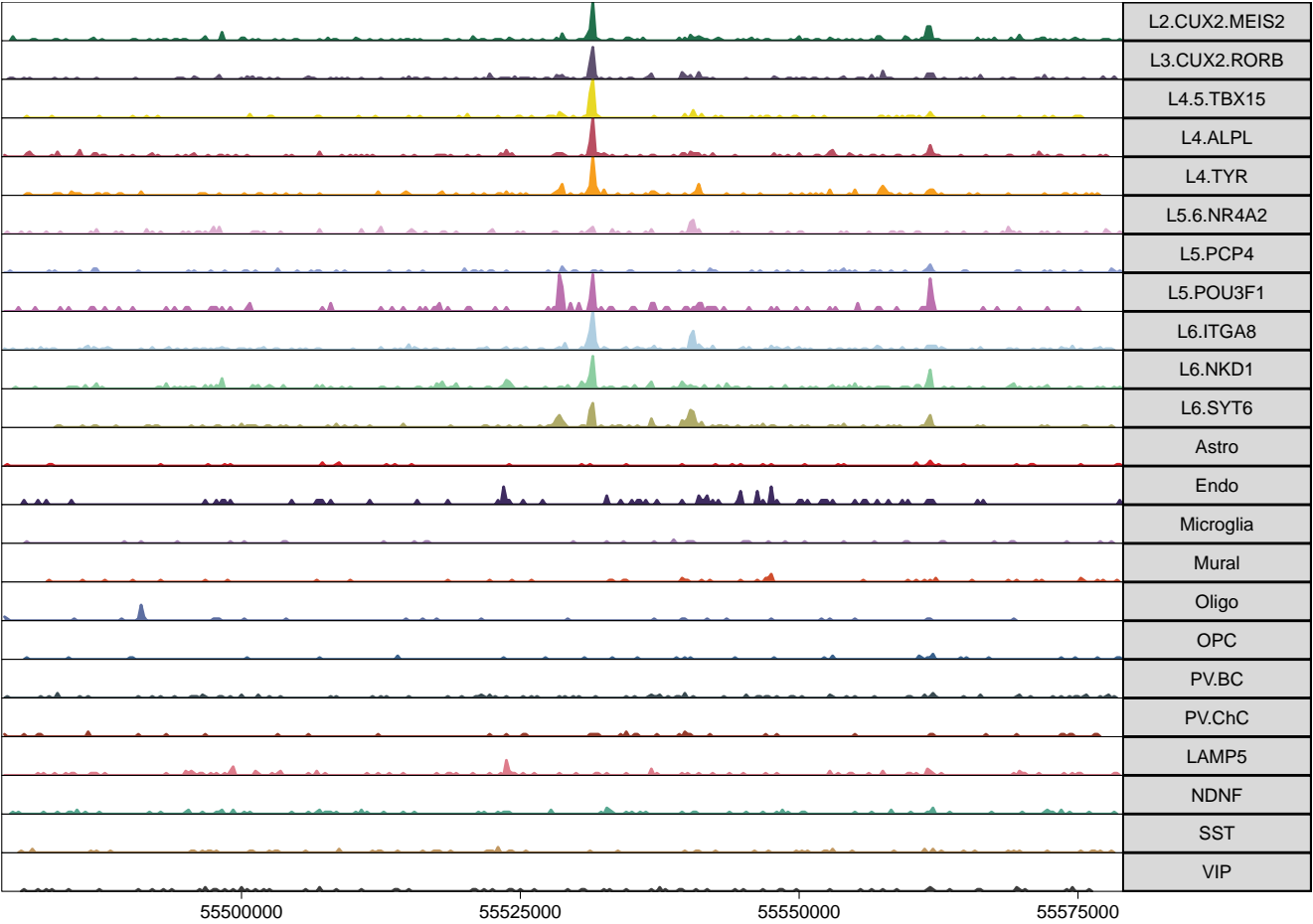
Peaks



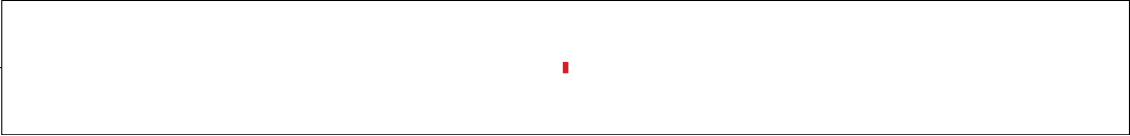
Genes

chr14:55478500–55579001 Marker Peak Tracks for: SST

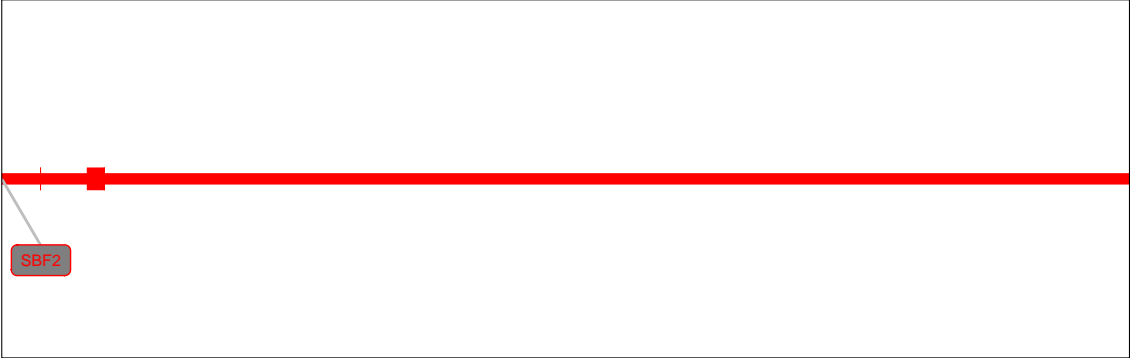
Coverage  
(Norm. ATAC Signal Range (0–0.69) by ReadsInTSS)



SST



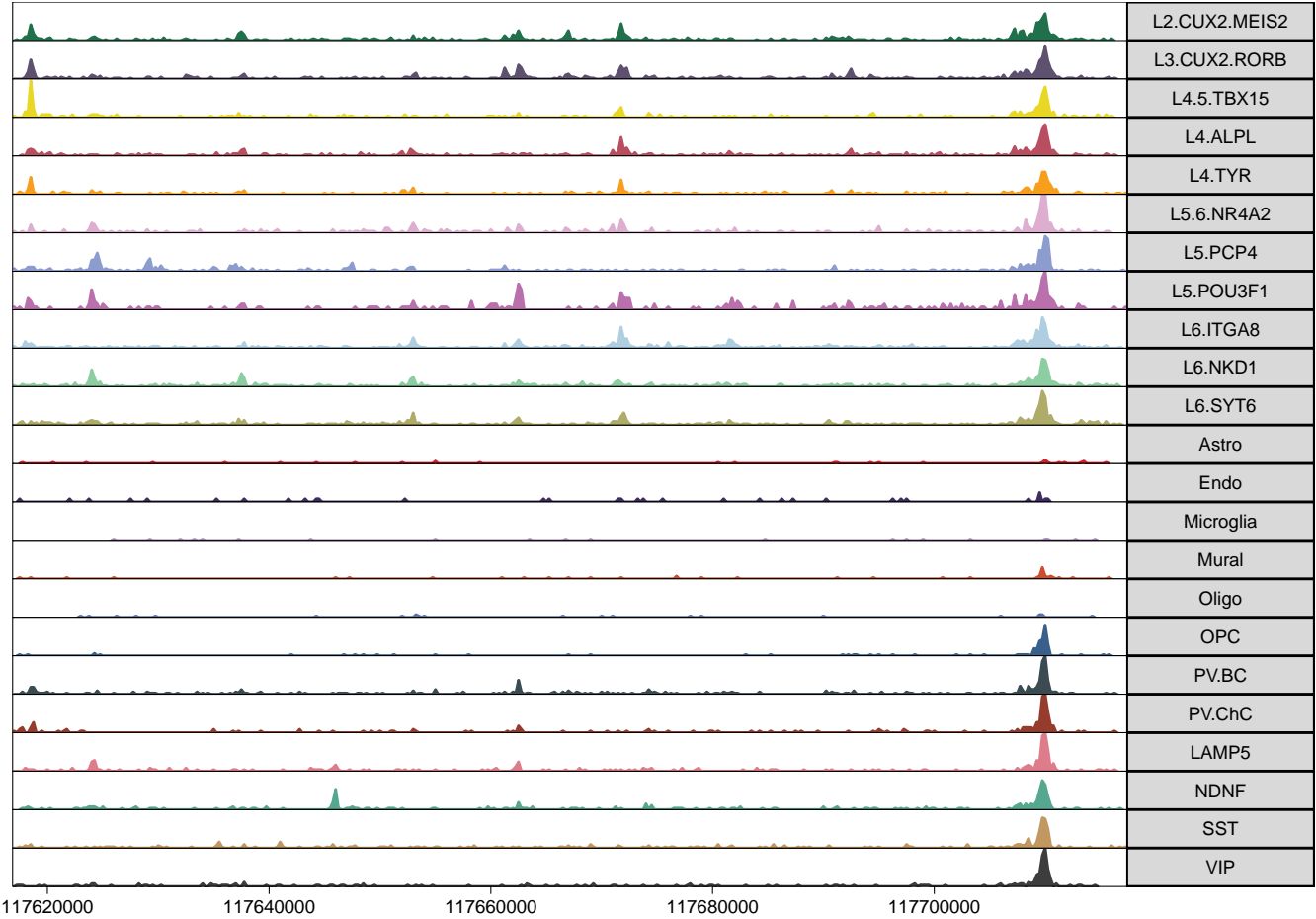
Peaks



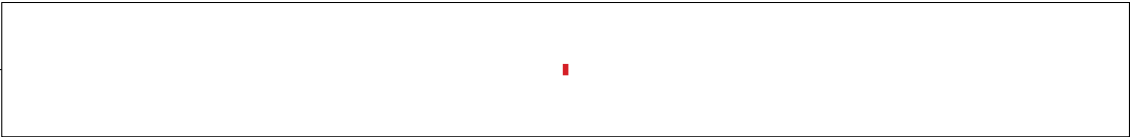
Genes

chr11:117616861–117717362 Marker Peak Tracks for: SST

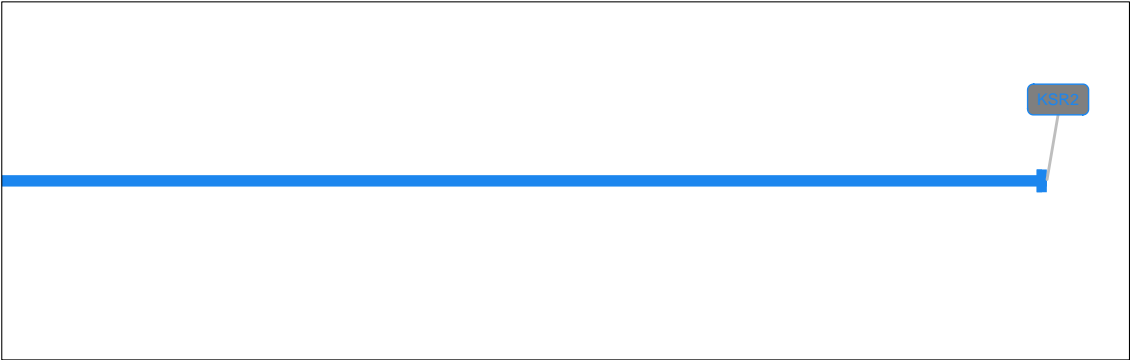
Coverage  
(Norm. ATAC Signal Range (0–0.98) by ReadsInTSS)



SST

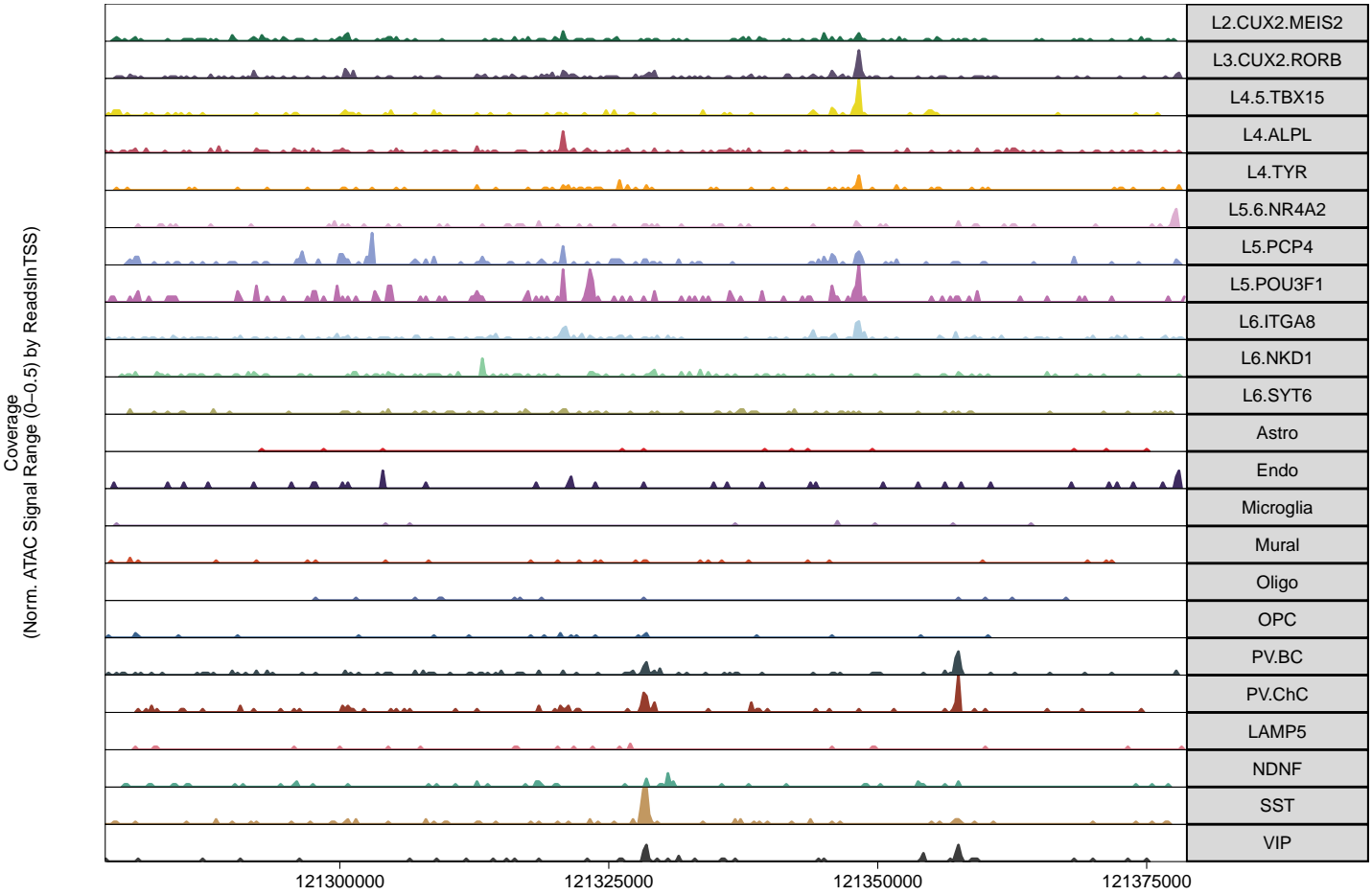


Peaks



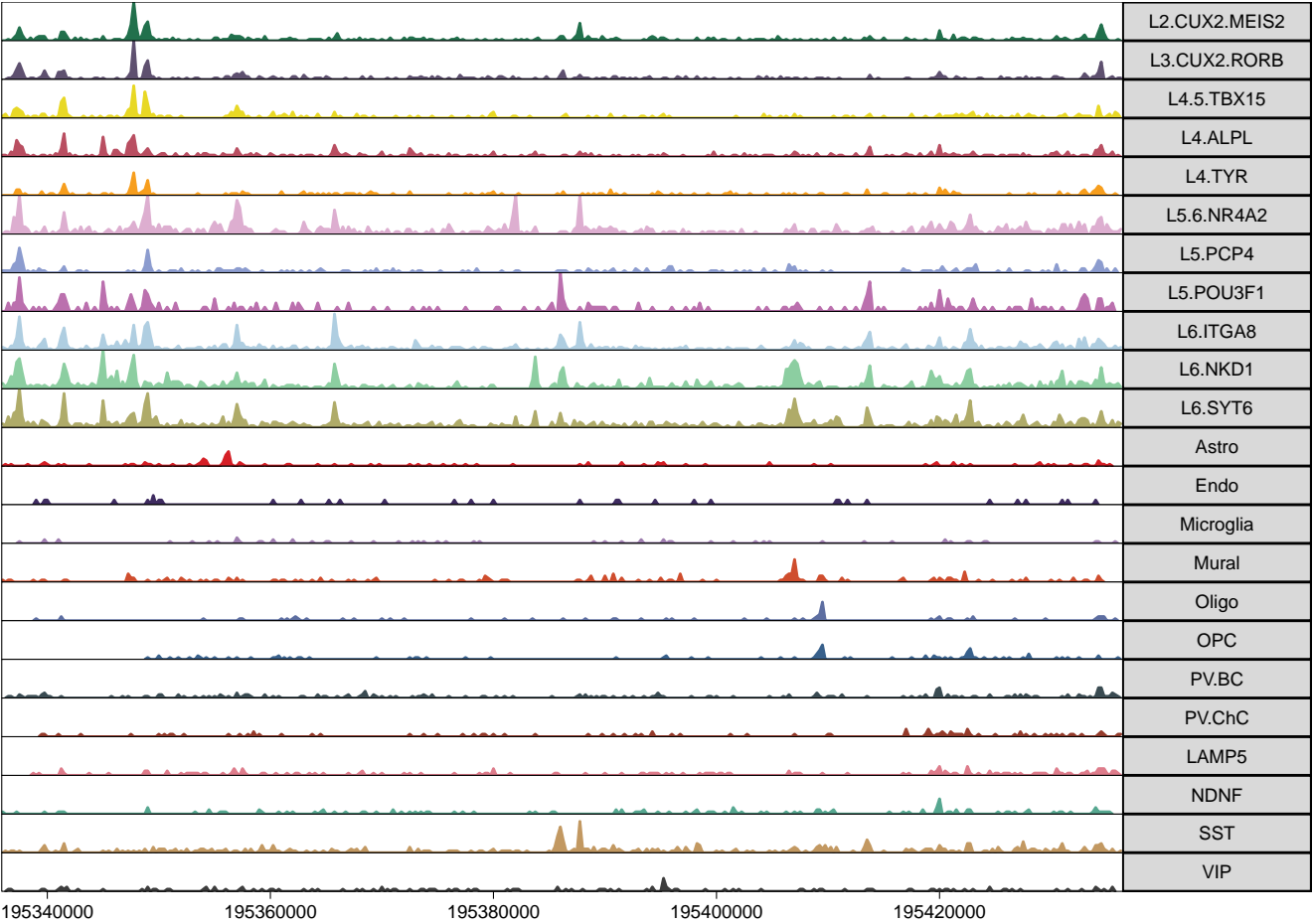
Genes

chr1:121278193–121378694 Marker Peak Tracks for: SST

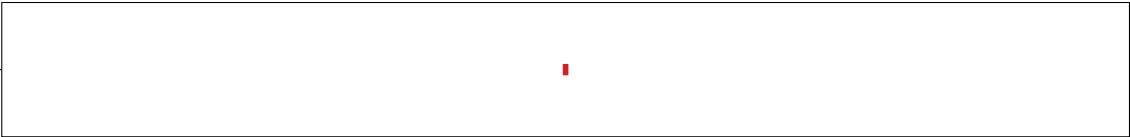


chr1:195335912–195436413 Marker Peak Tracks for: SST

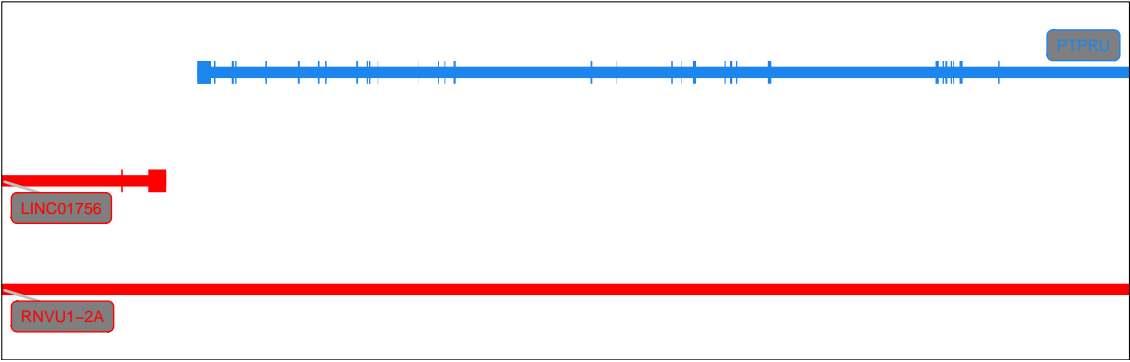
Coverage  
(Norm. ATAC Signal Range (0–0.67) by ReadsInTSS)



SST



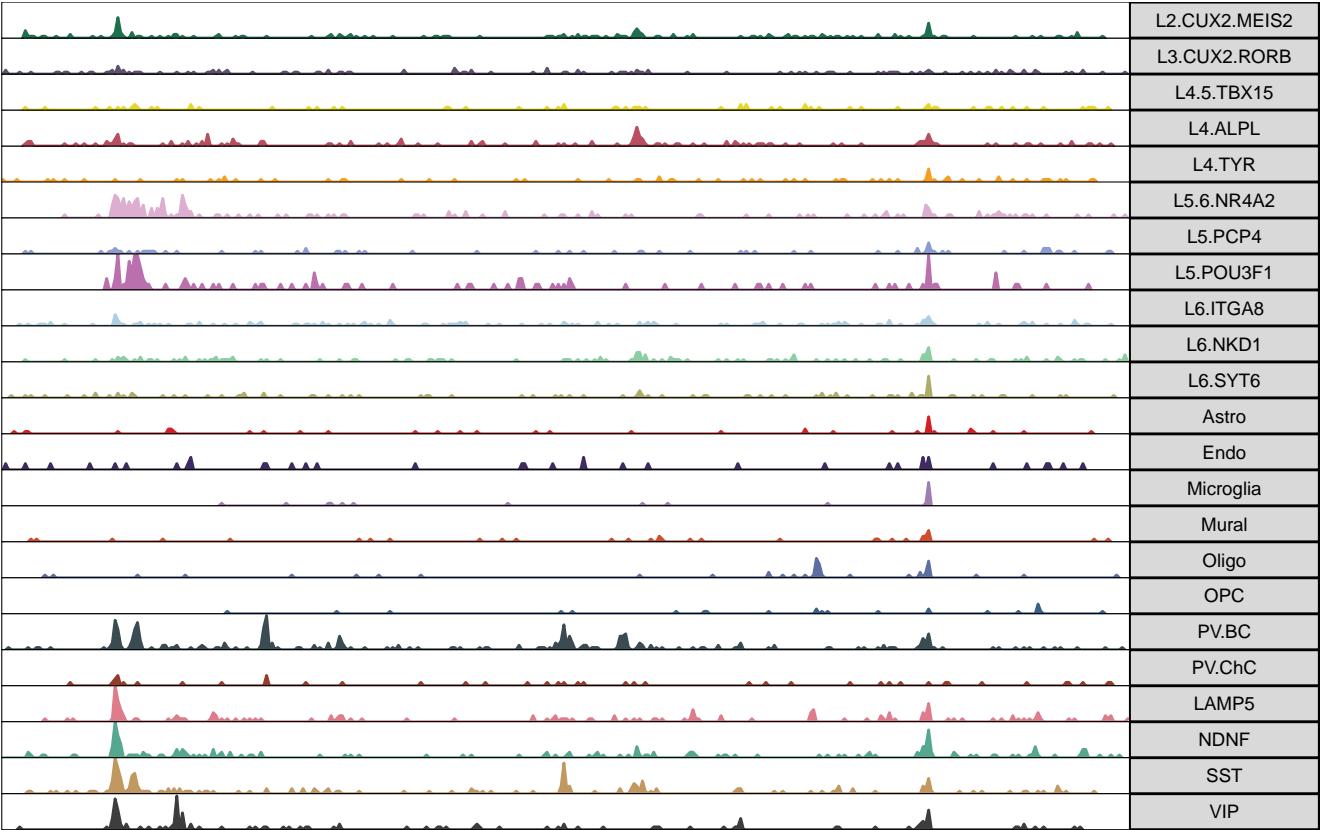
Peaks



Genes



Coverage  
(Norm. ATAC Signal Range (0-0.46) by ReadsInTSS)



SST



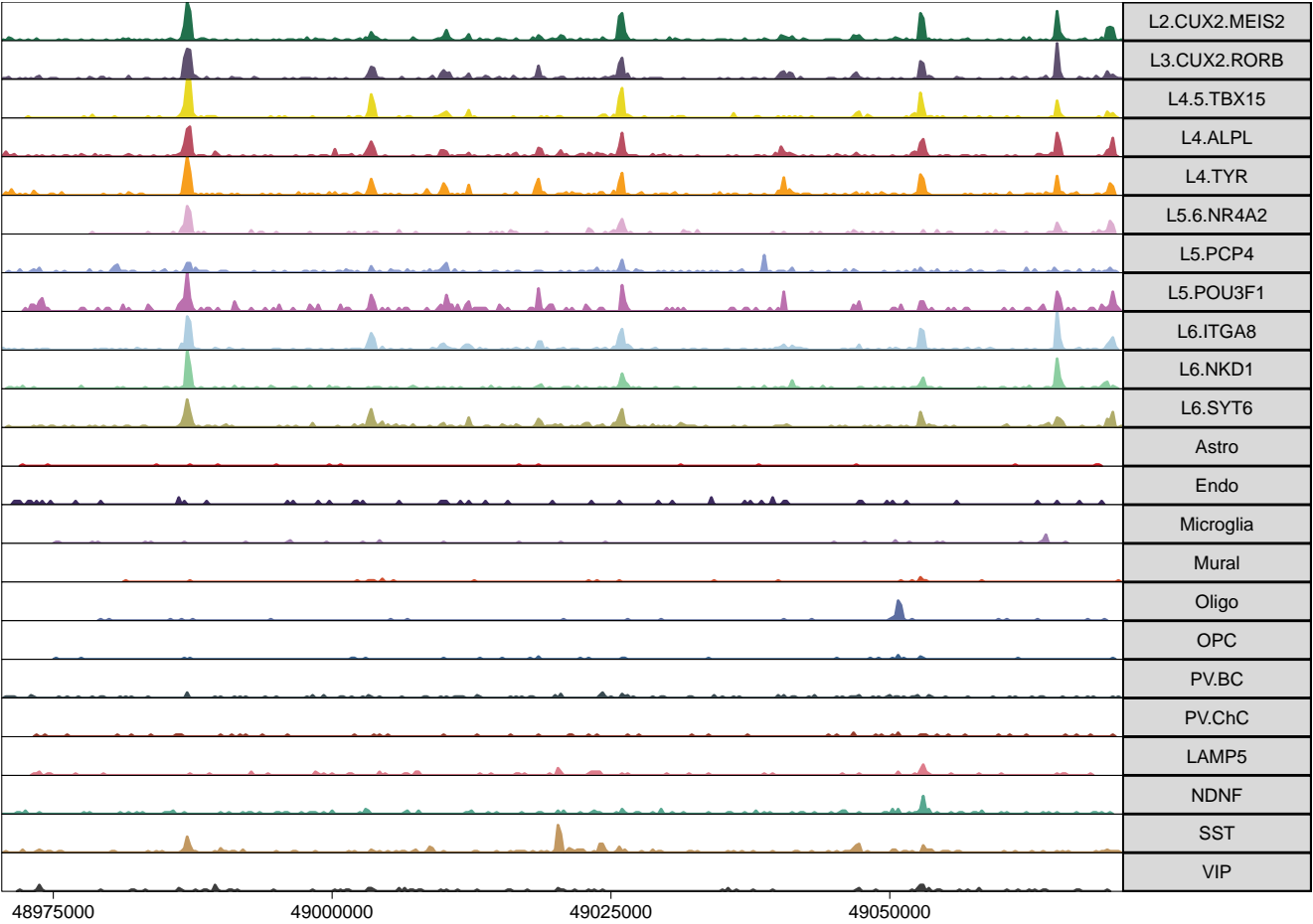
Peaks

GeneTrack

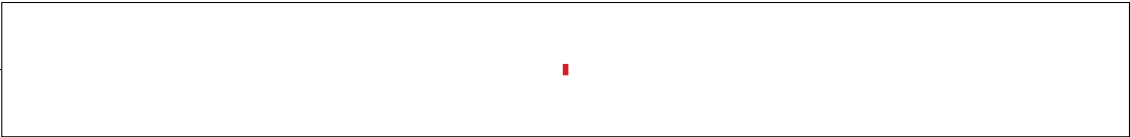
strand ● ·

chr20:48970375–49070876 Marker Peak Tracks for: SST

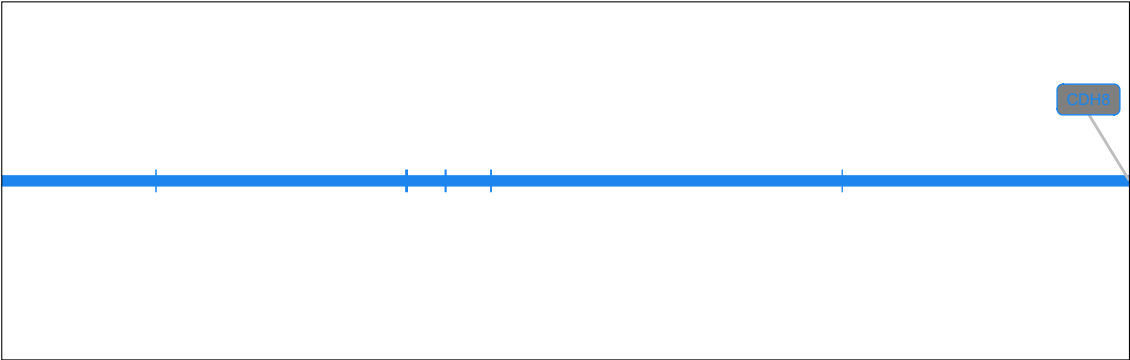
Coverage  
(Norm. ATAC Signal Range (0–0.87) by ReadsInTSS)



SST



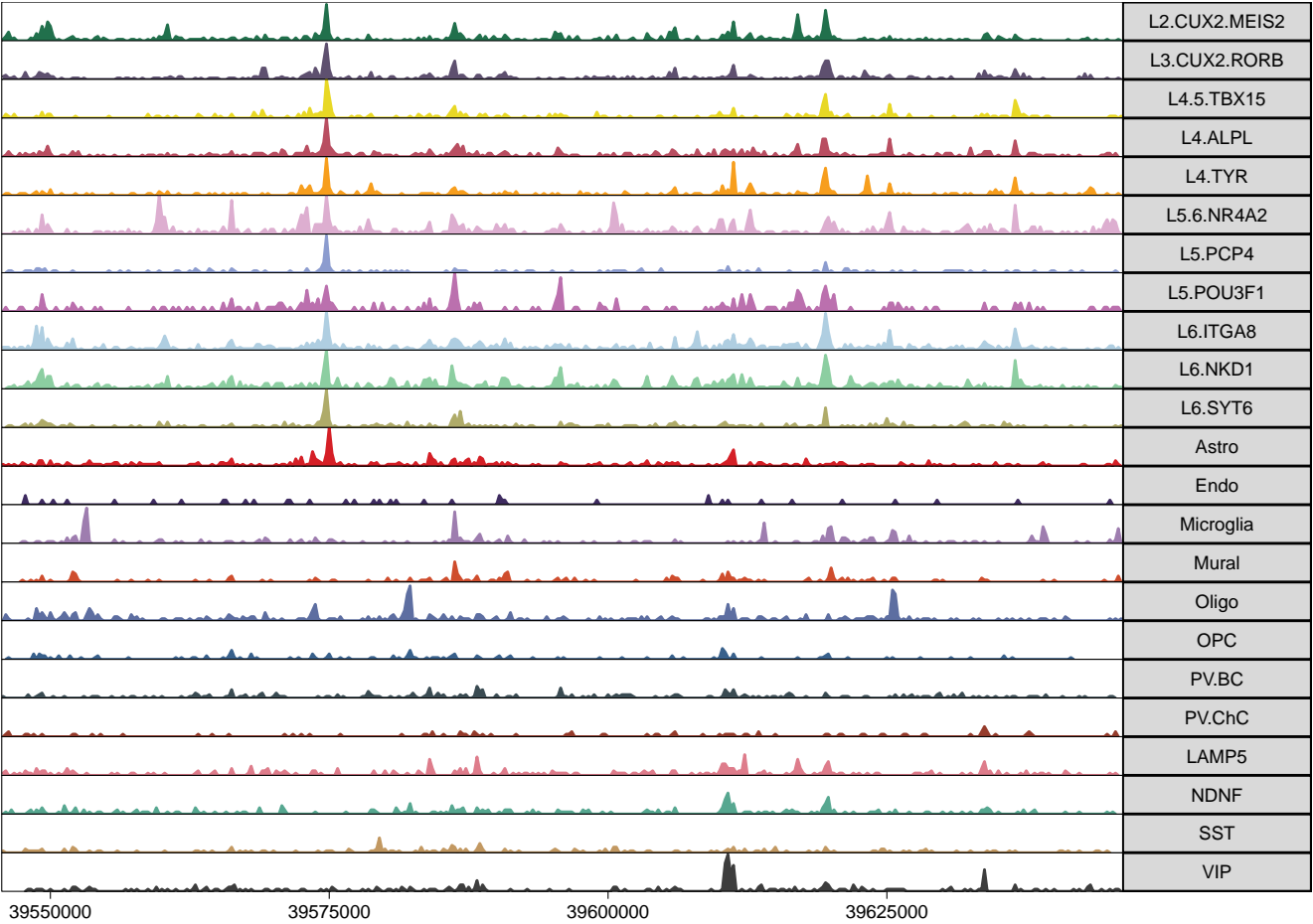
Peaks



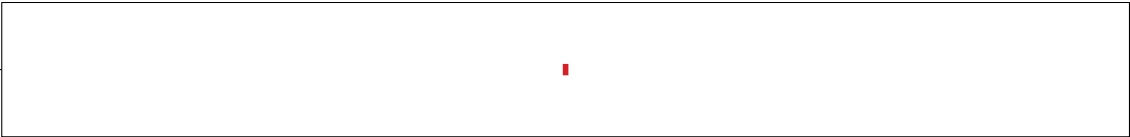
Genes

chr12:39545634–39646135 Marker Peak Tracks for: SST

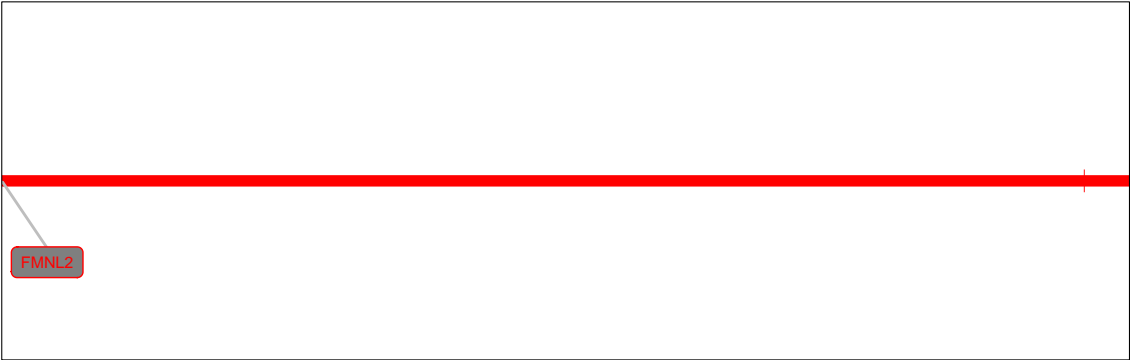
Coverage  
(Norm. ATAC Signal Range (0–0.67) by ReadsInTSS)



SST

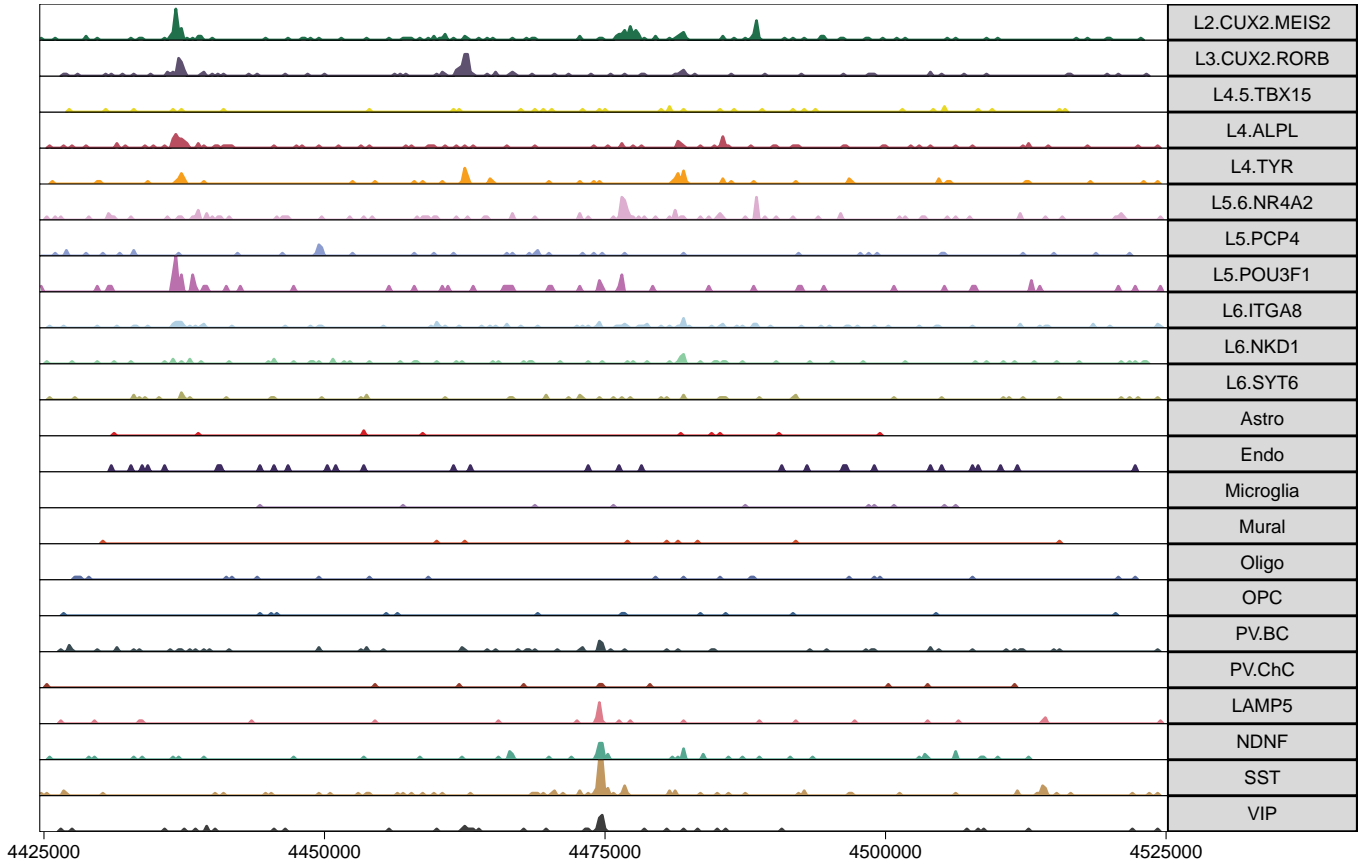


Peaks



Genes

Coverage  
(Norm. ATAC Signal Range (0–0.46) by ReadsInTSS)



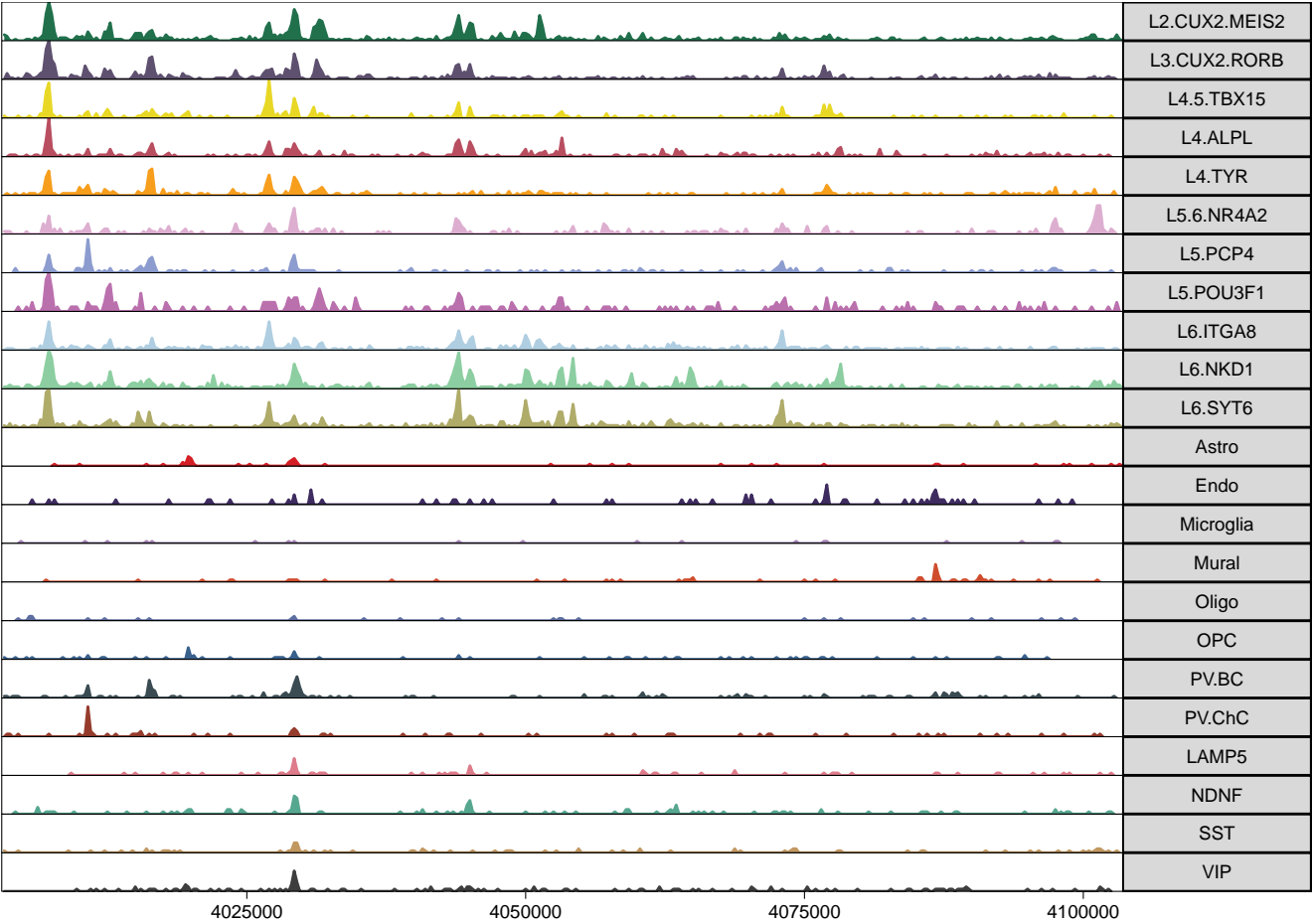
SST



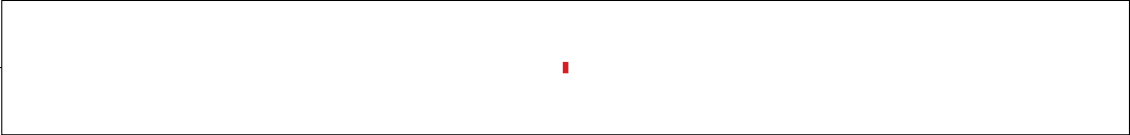
Peaks

GeneTrack

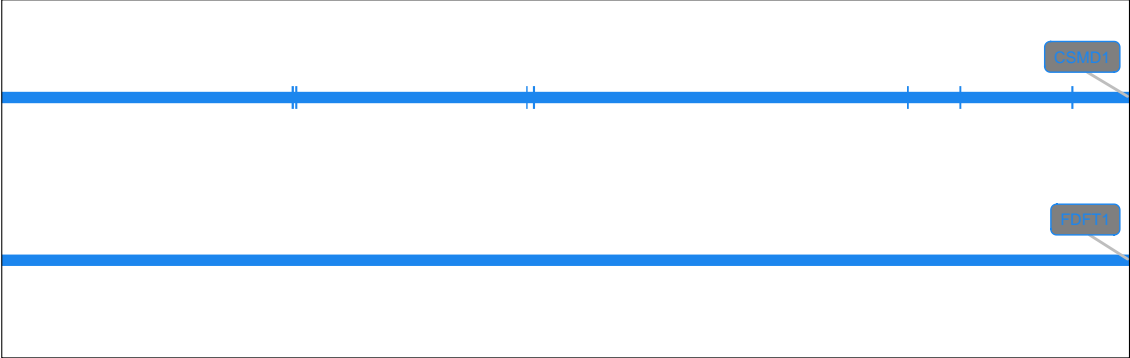
Coverage  
(Norm. ATAC Signal Range (0–0.62) by ReadsInTSS)



SST

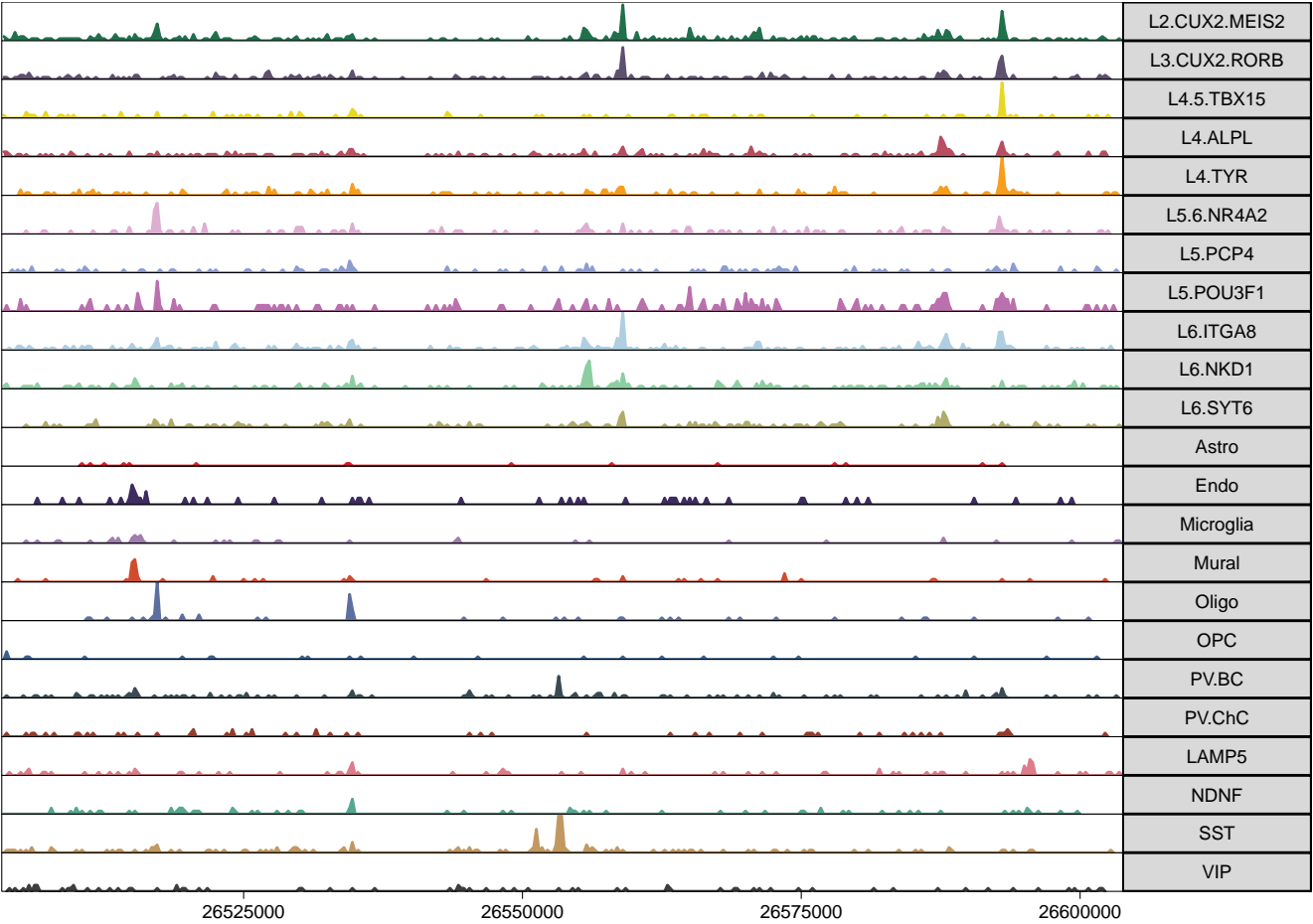


Peaks

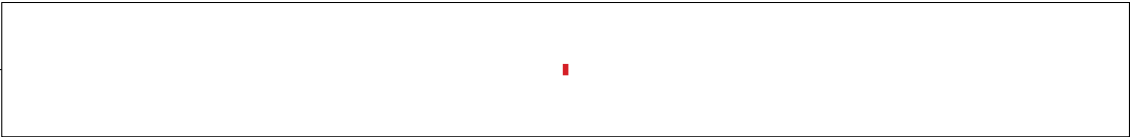


Genes

Coverage  
(Norm. ATAC Signal Range (0-0.47) by ReadsInTSS)



SST



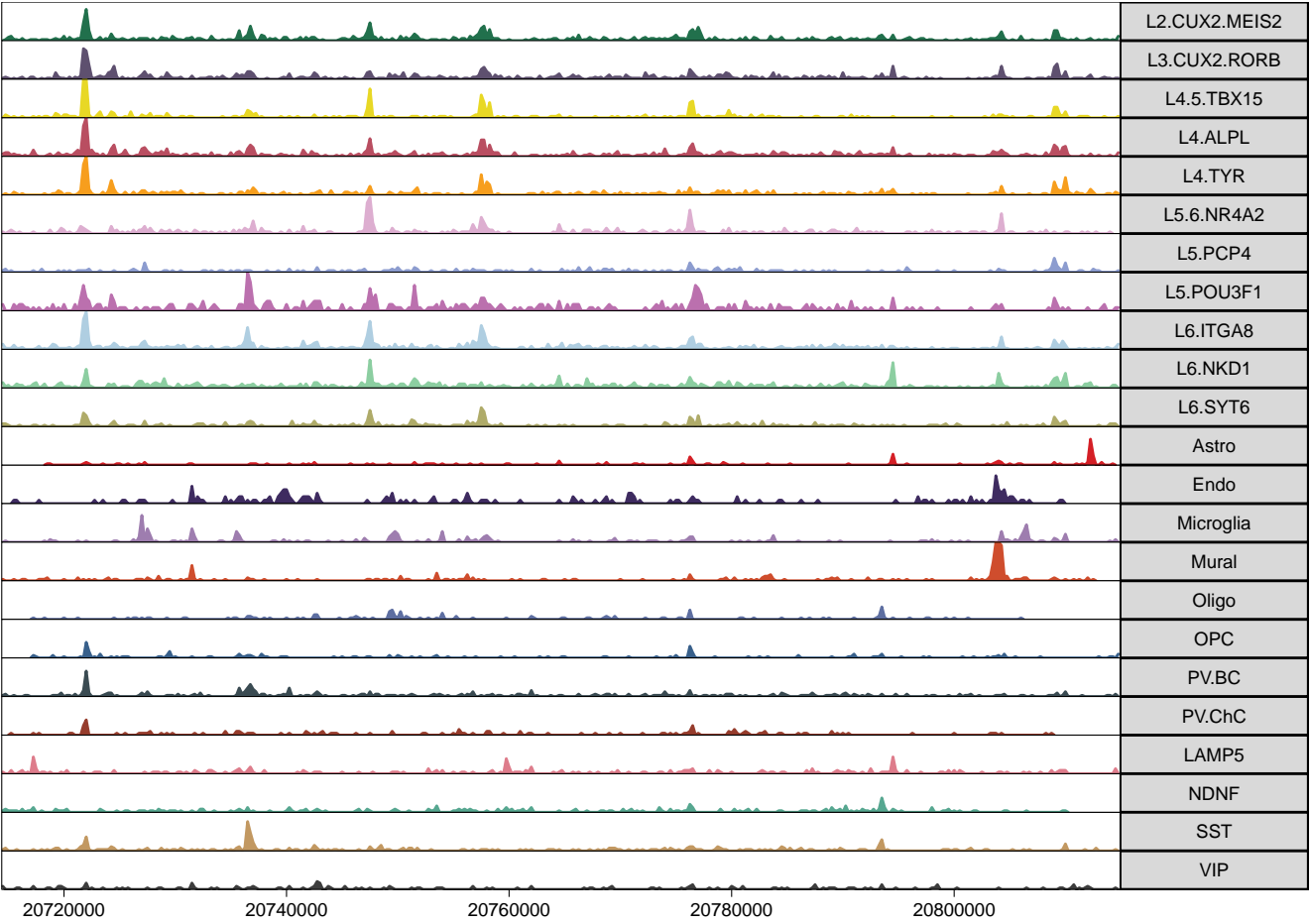
Peaks



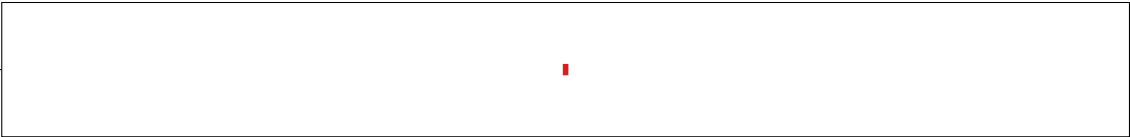
Genes

chr3:20714405–20814906 Marker Peak Tracks for: SST

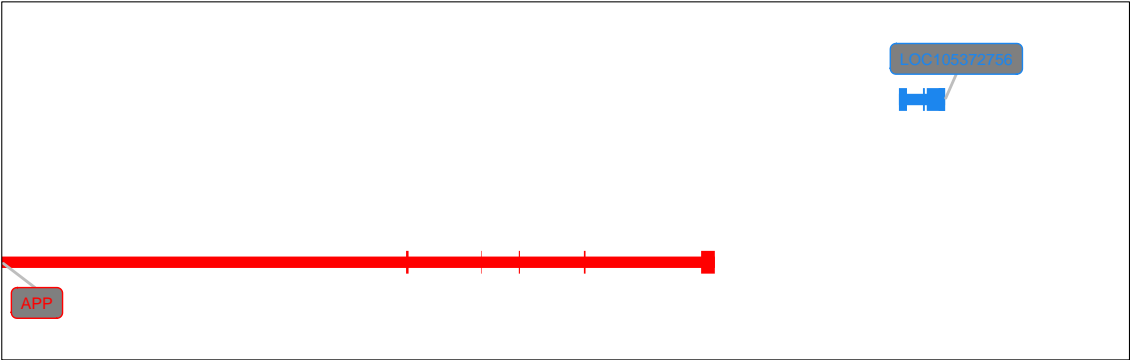
Coverage  
(Norm. ATAC Signal Range (0–0.89) by ReadsInTSS)



SST

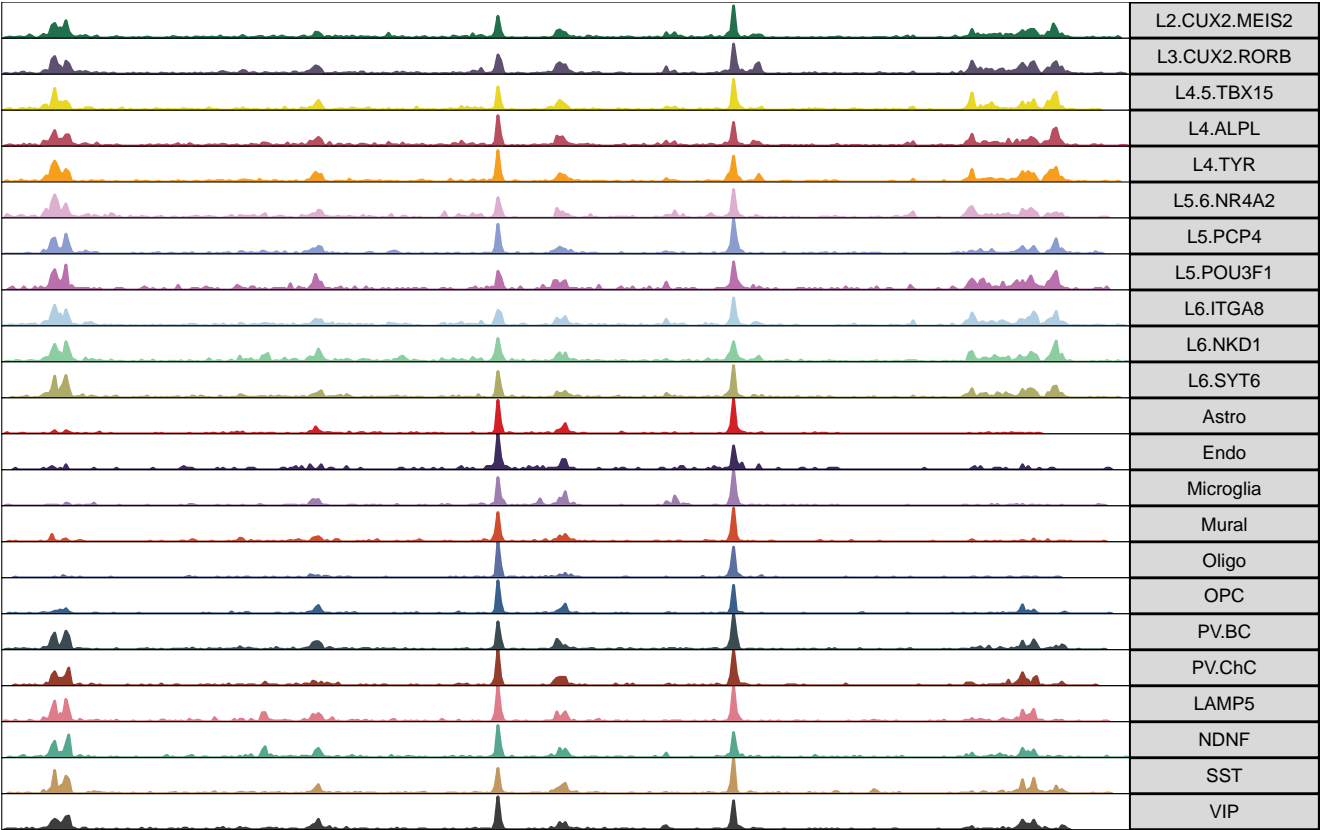


Peaks



Genes

Coverage  
(Norm. ATAC Signal Range (0–1.69) by ReadsInTSS)



SST

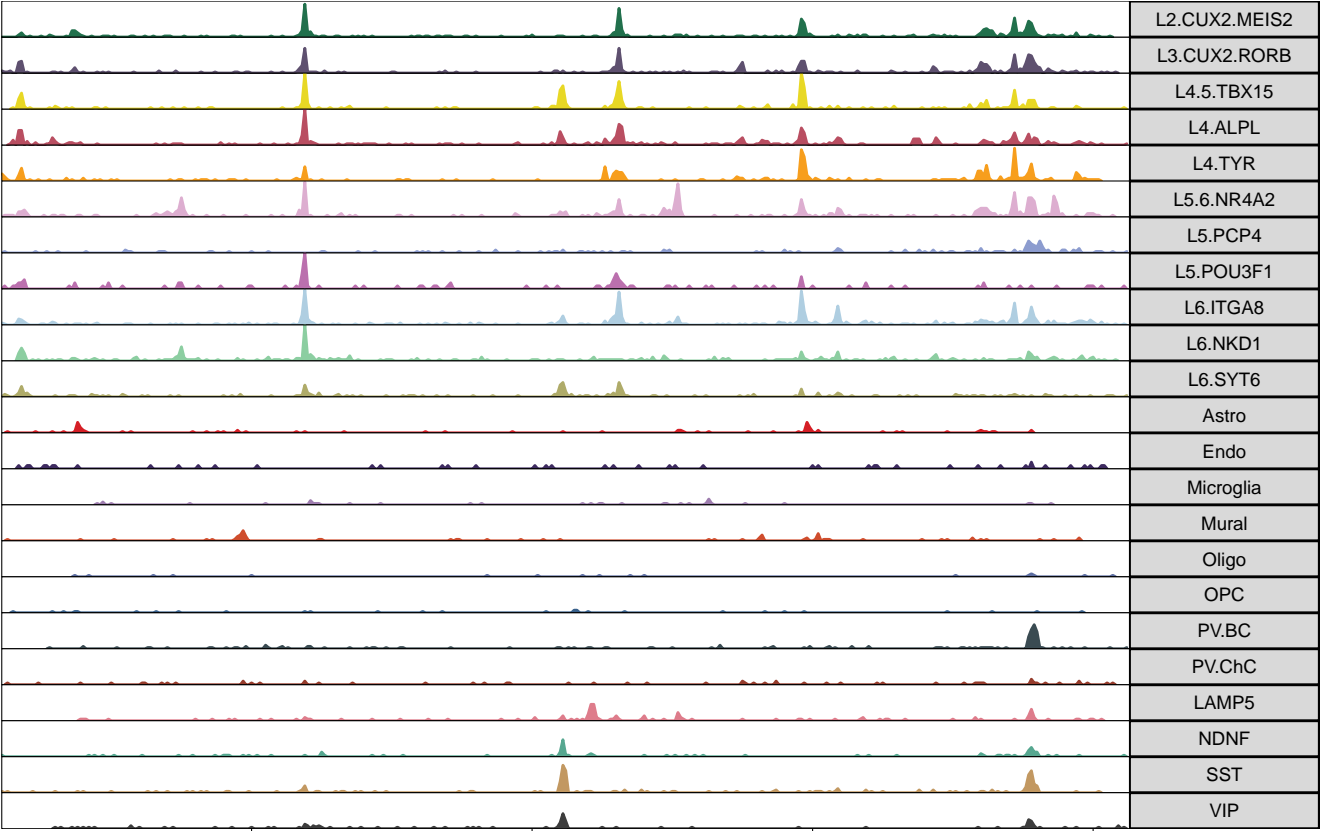


Peaks

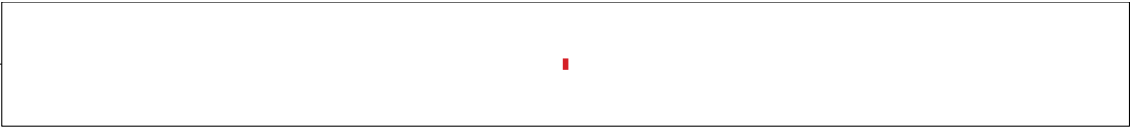
GeneTrack



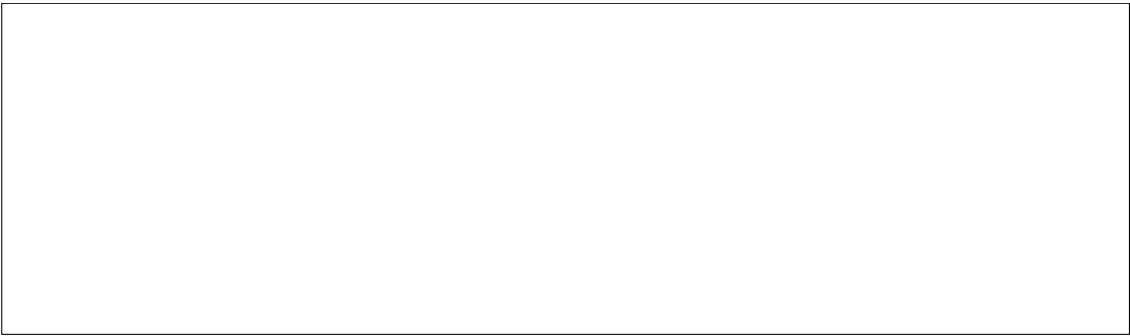
Coverage  
(Norm. ATAC Signal Range (0–0.86) by ReadsInTSS)



SST

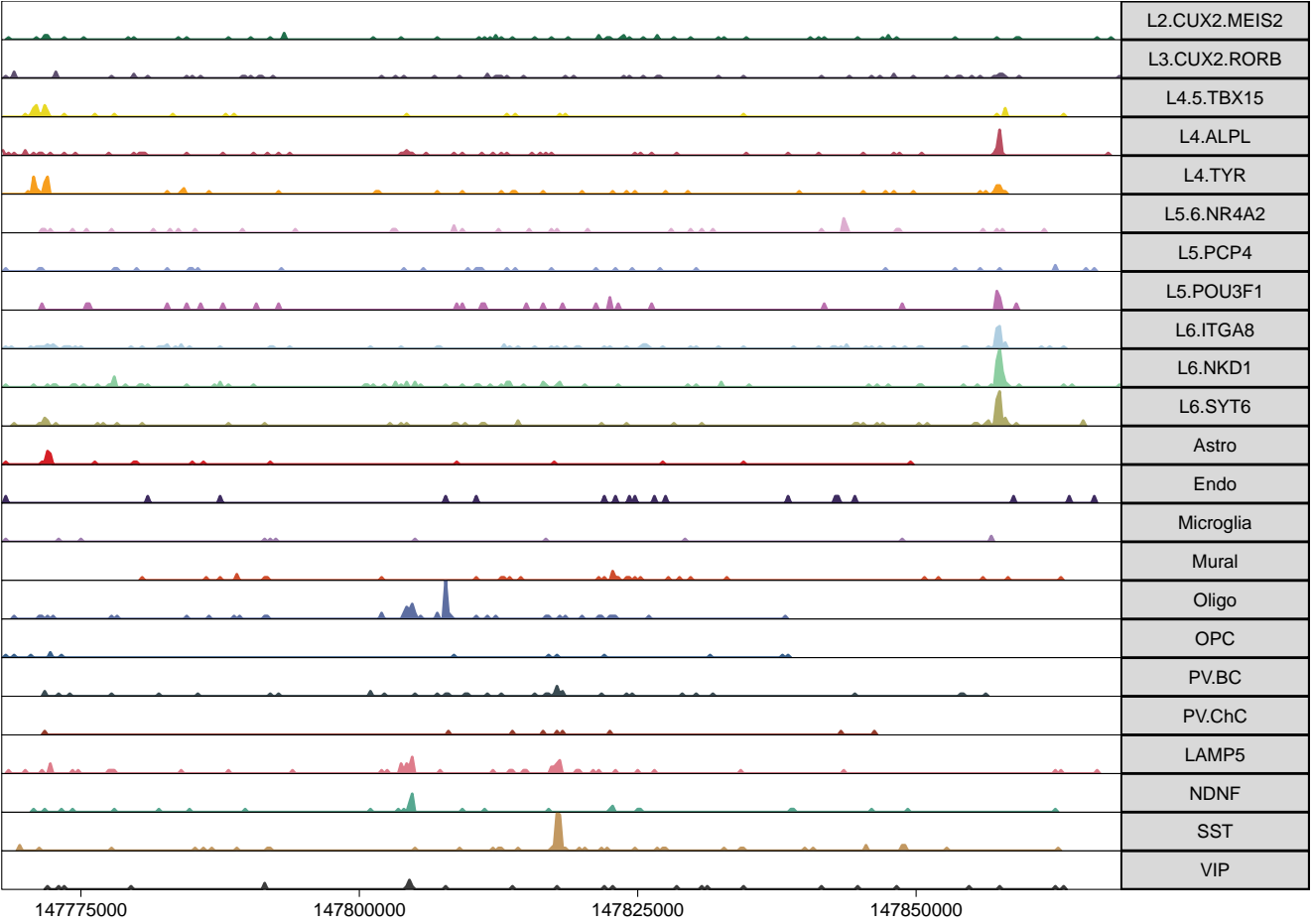


Peaks

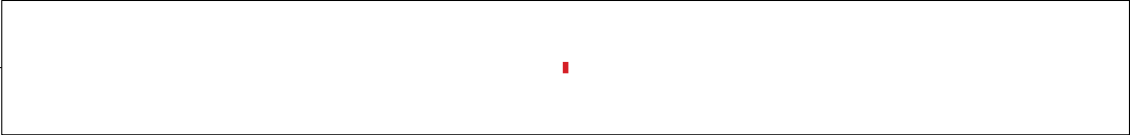


GeneTrack

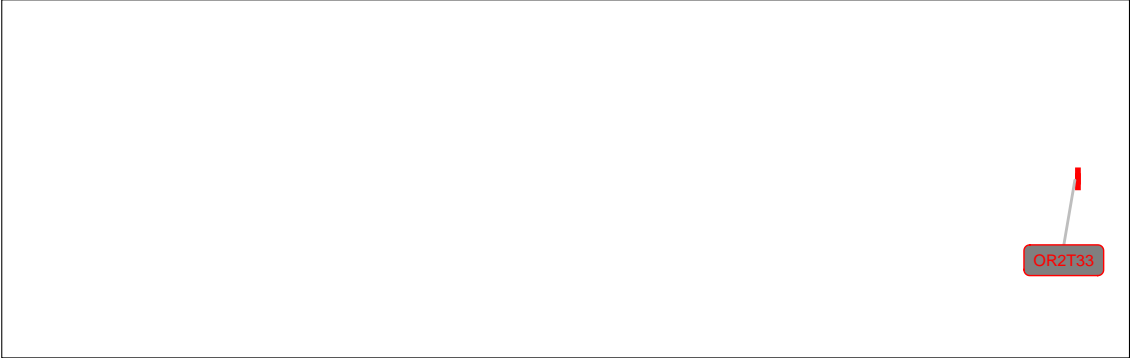
Coverage  
(Norm. ATAC Signal Range (0–0.45) by ReadsInTSS)



SST



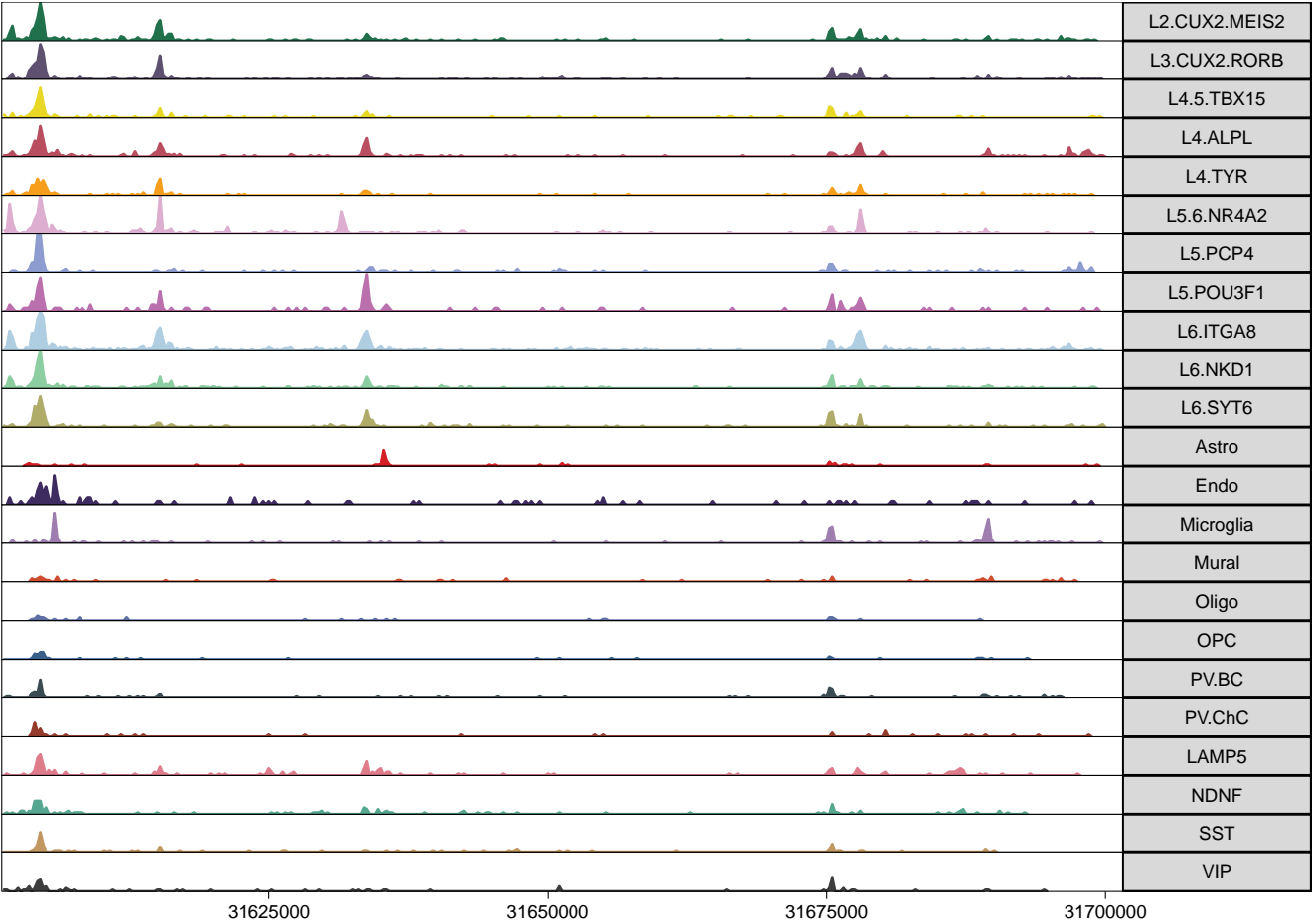
Peaks



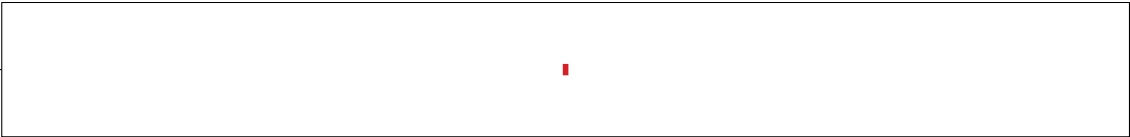
Genes

chr16:31601036–31701537 Marker Peak Tracks for: SST

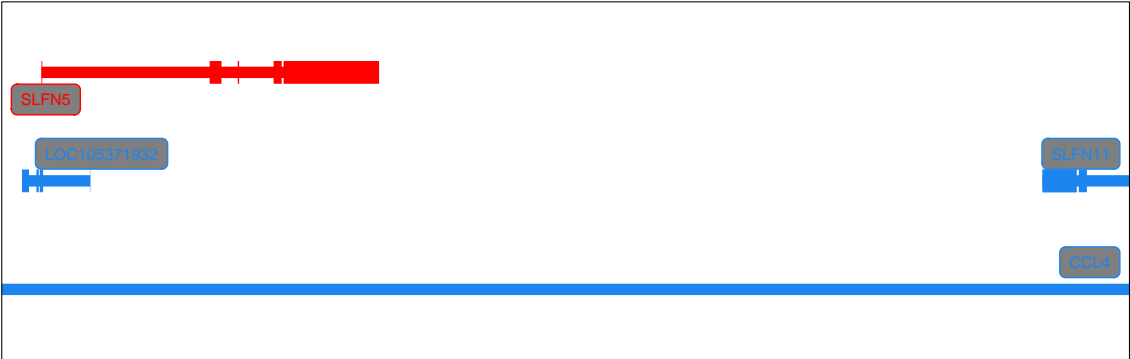
Coverage  
(Norm. ATAC Signal Range (0–0.85) by ReadsInTSS)



SST

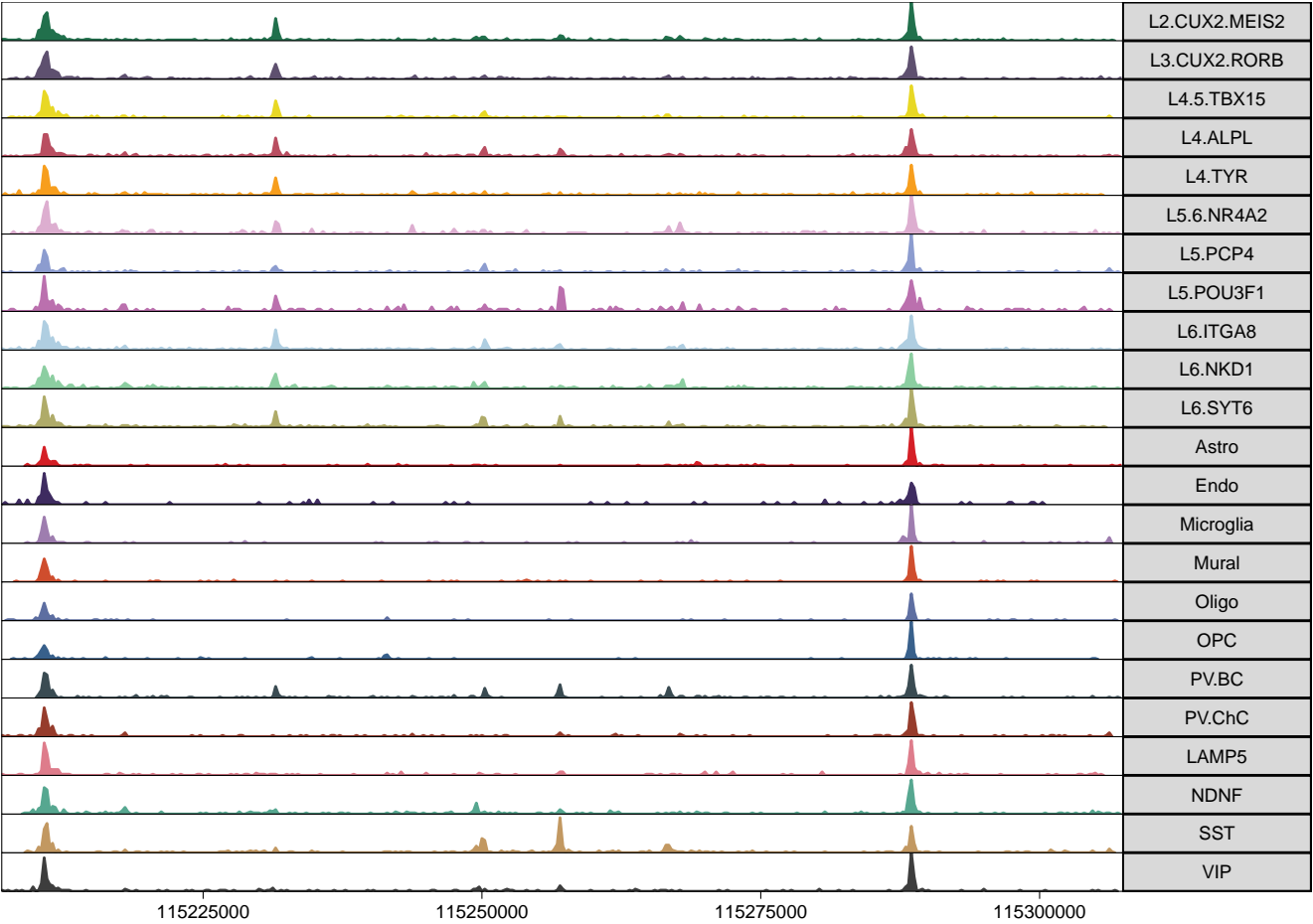


Peaks

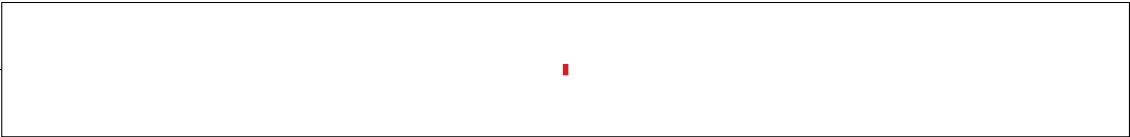


Genes

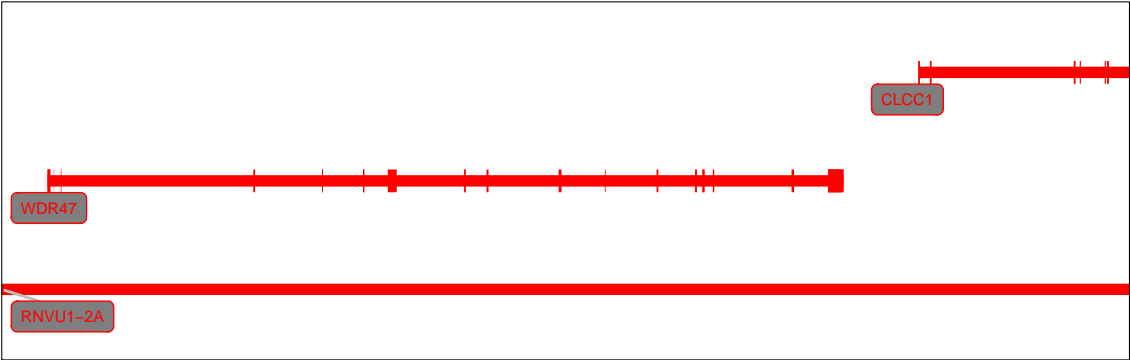
Coverage  
(Norm. ATAC Signal Range (0–1.29) by ReadsInTSS)



SST

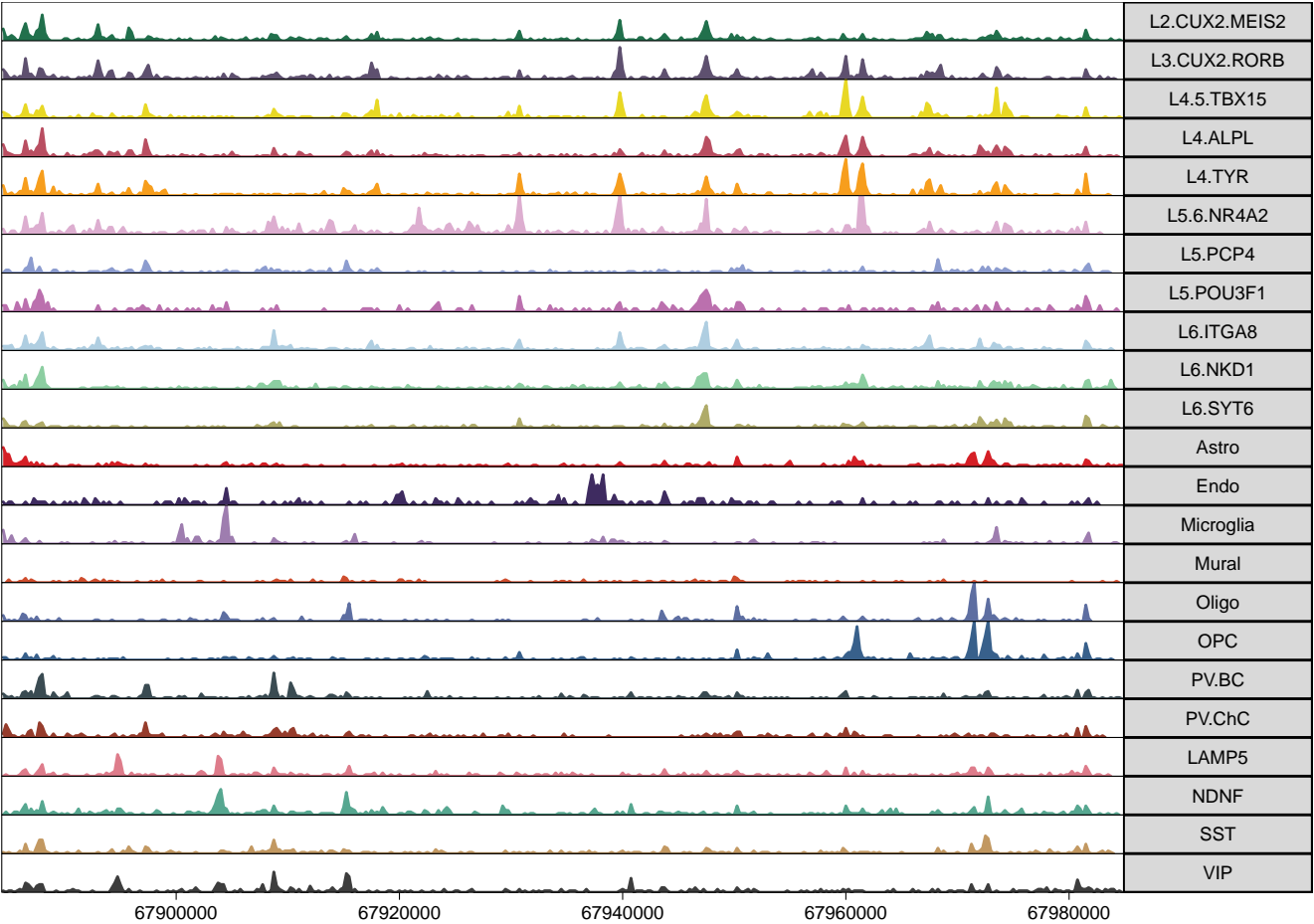


Peaks

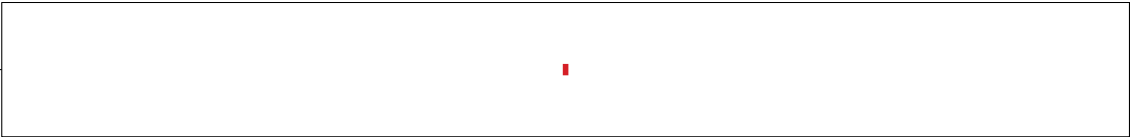


Genes

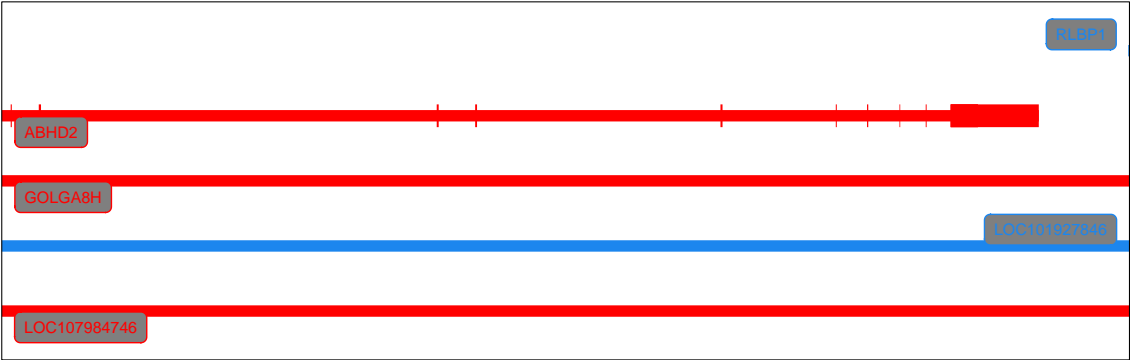
Coverage  
(Norm. ATAC Signal Range (0-0.92) by ReadsInTSS)



SST

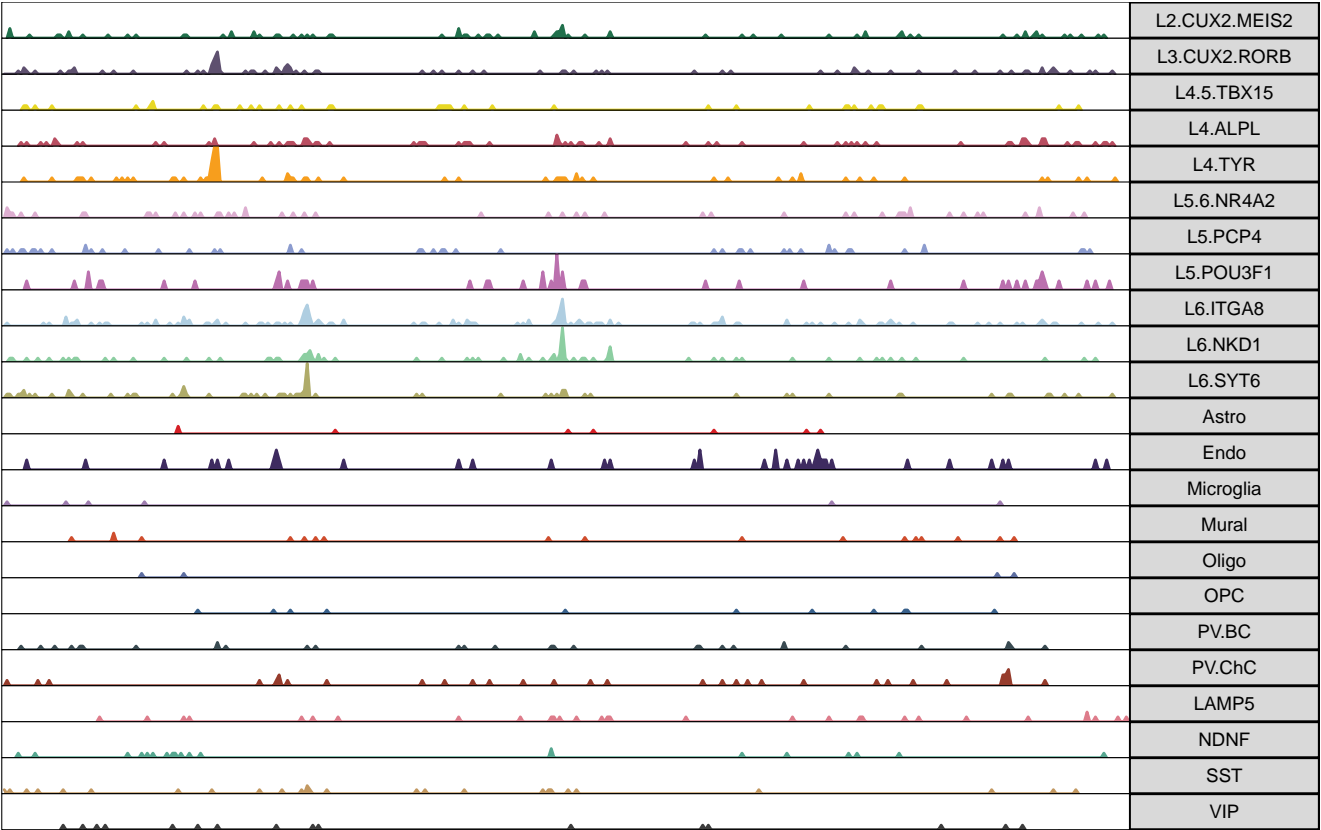


Peaks



Genes

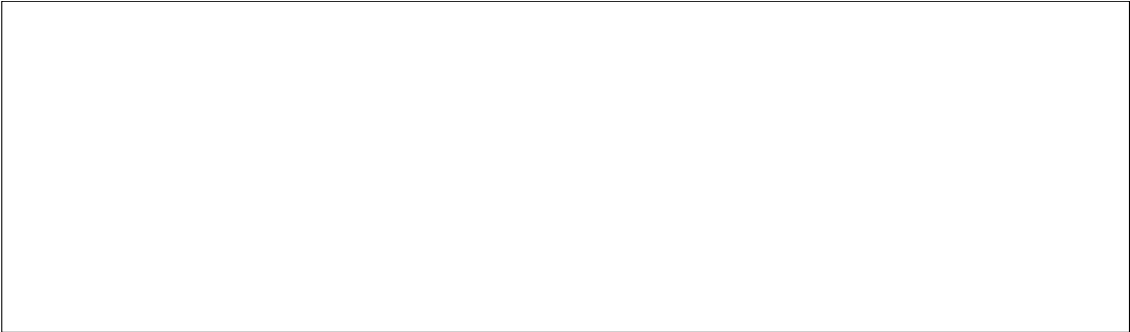
Coverage  
(Norm. ATAC Signal Range (0-0.29) by ReadsInTSS)



SST

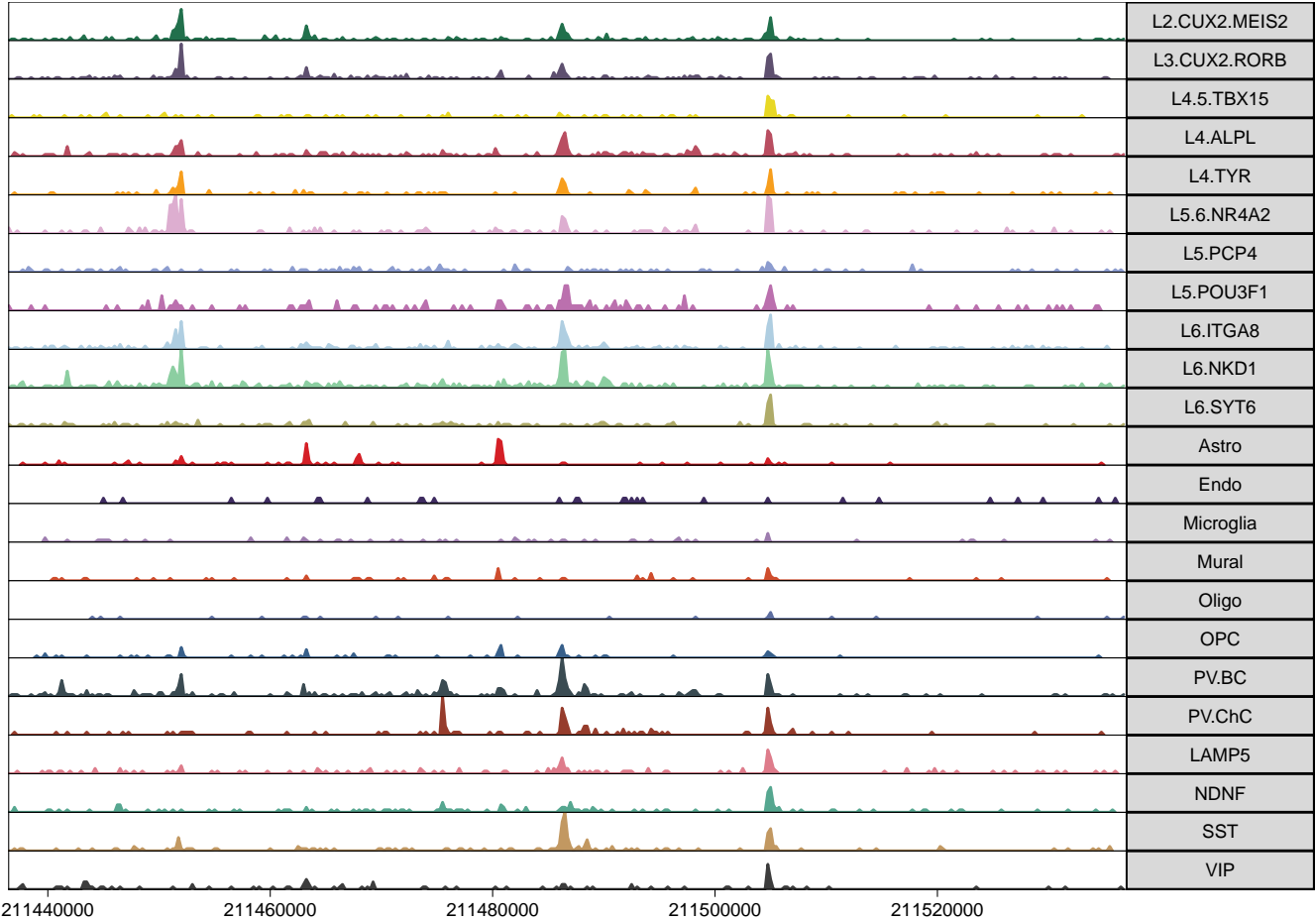


Peaks

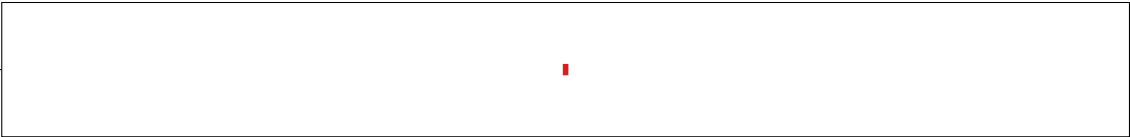


GeneTrack

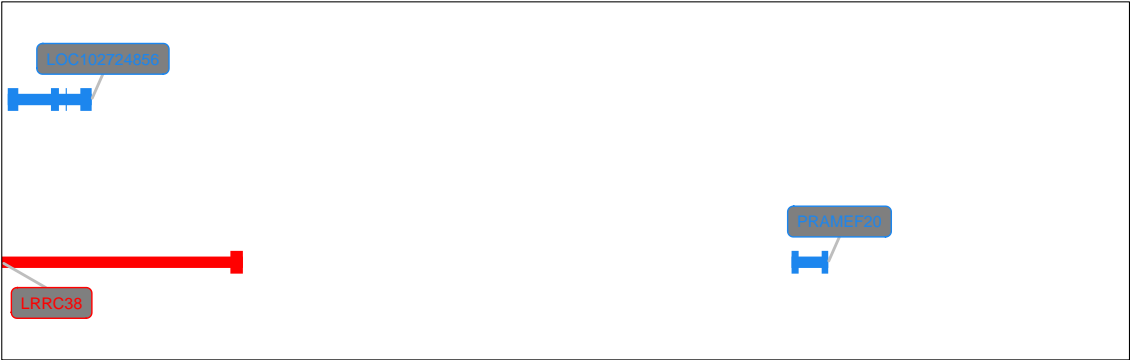
Coverage  
(Norm. ATAC Signal Range (0–0.57) by ReadsInTSS)



SST

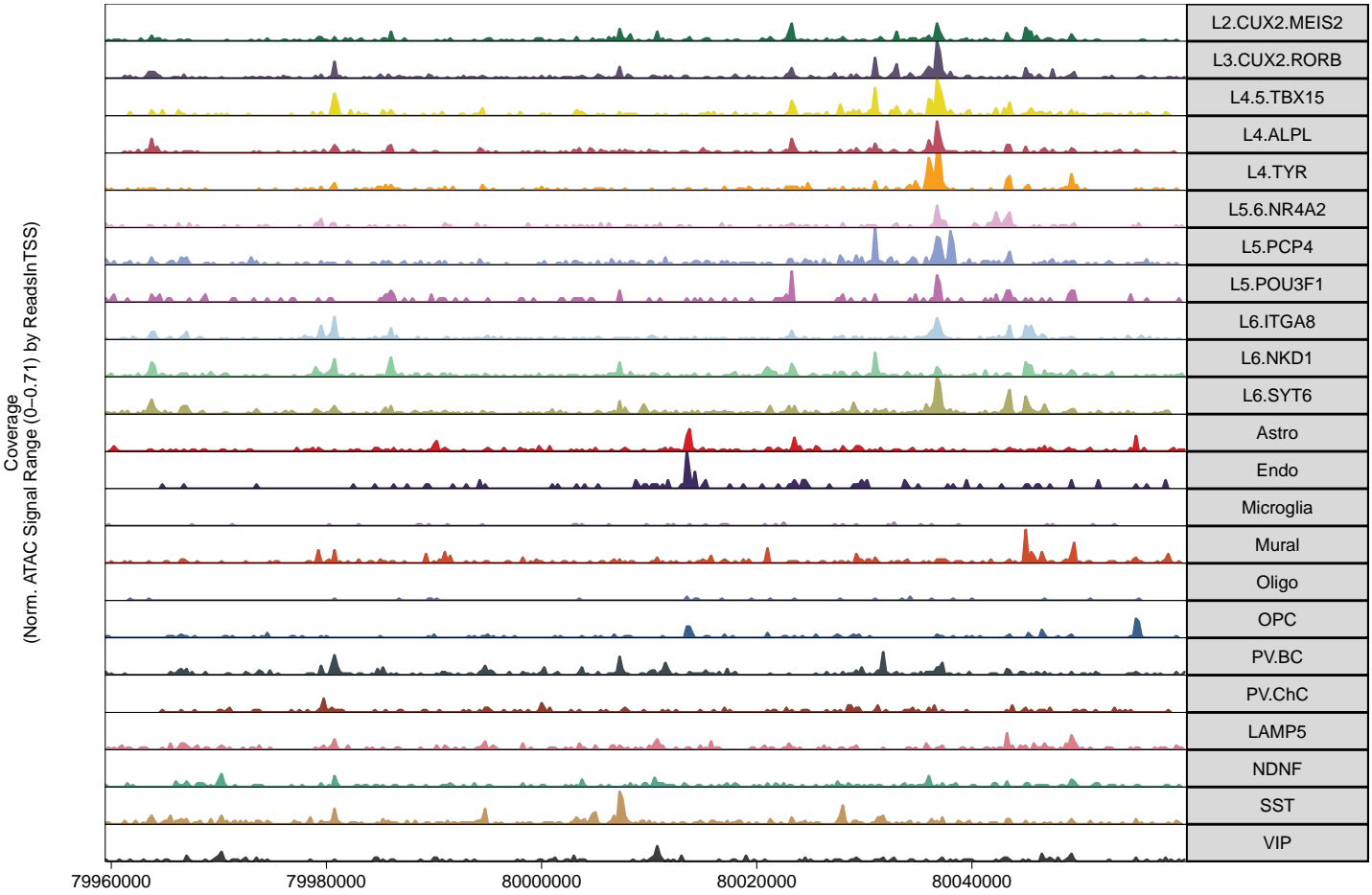


Peaks

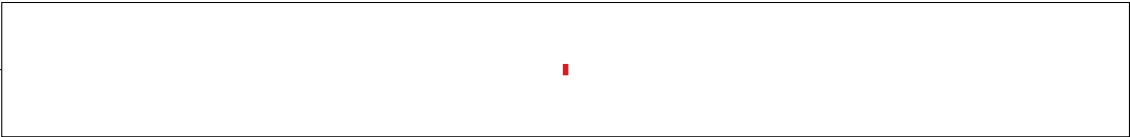


Genes

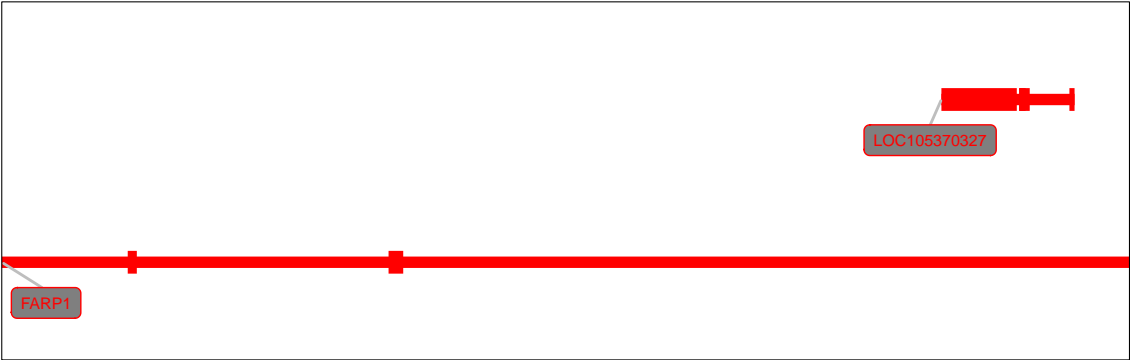
chr17:79959436–80059937 Marker Peak Tracks for: SST



SST



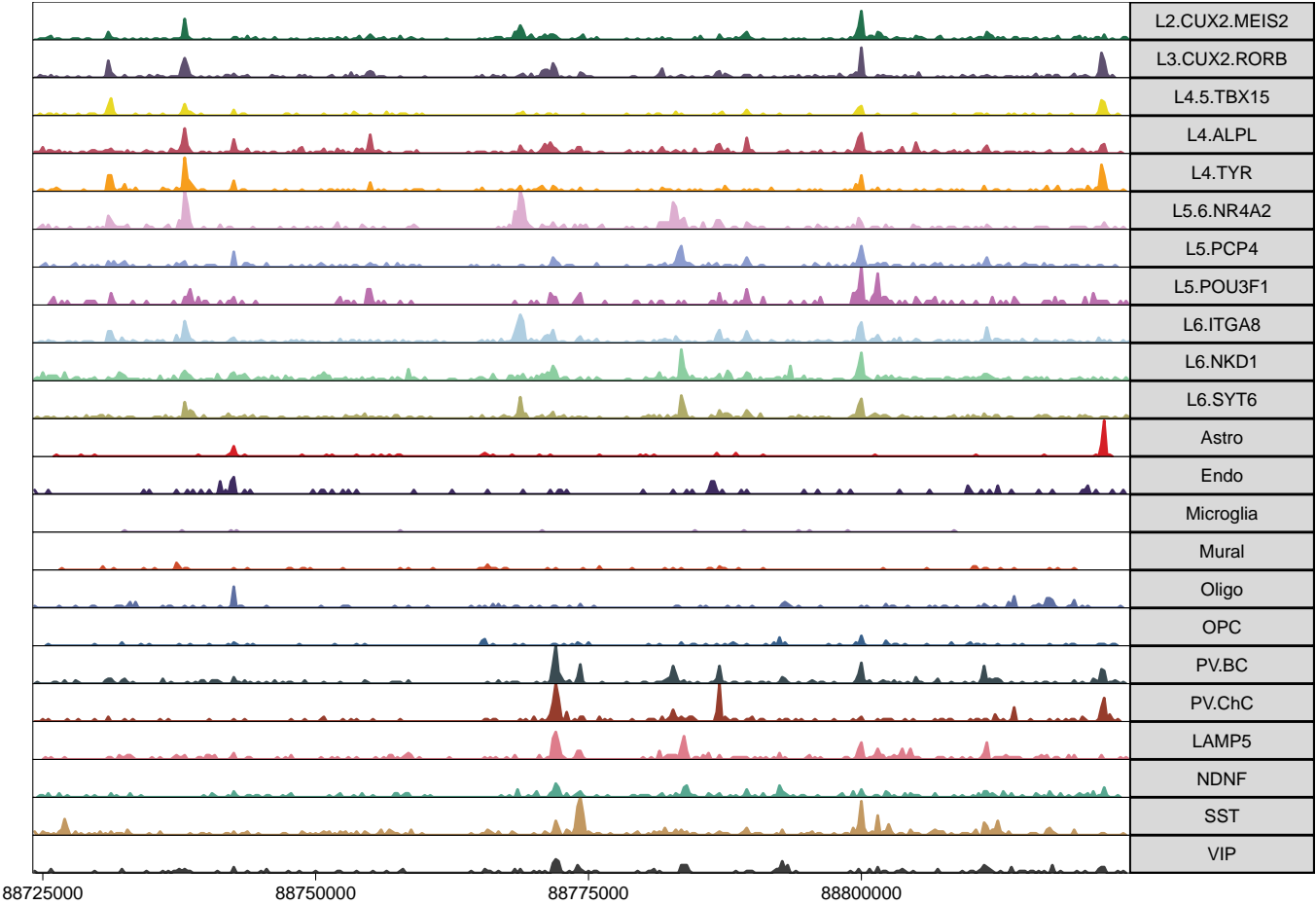
Peaks



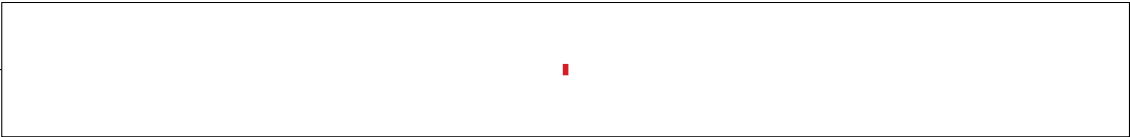
Genes



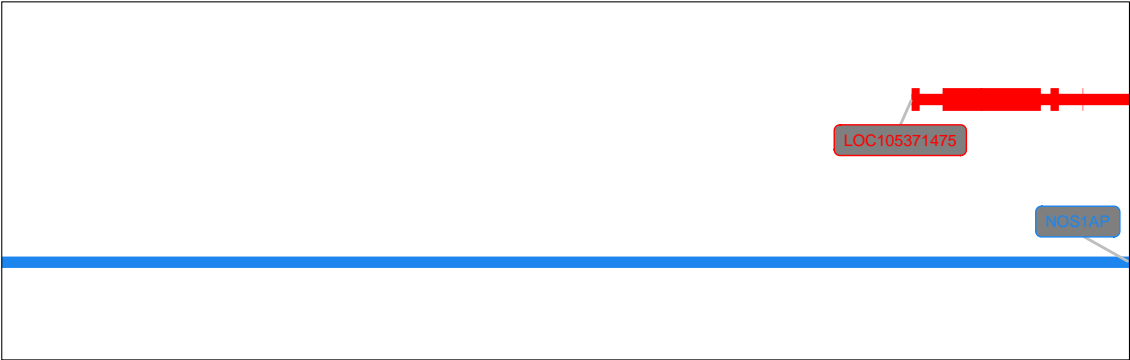
Coverage  
(Norm. ATAC Signal Range (0-0.71) by ReadsInTSS)



SST

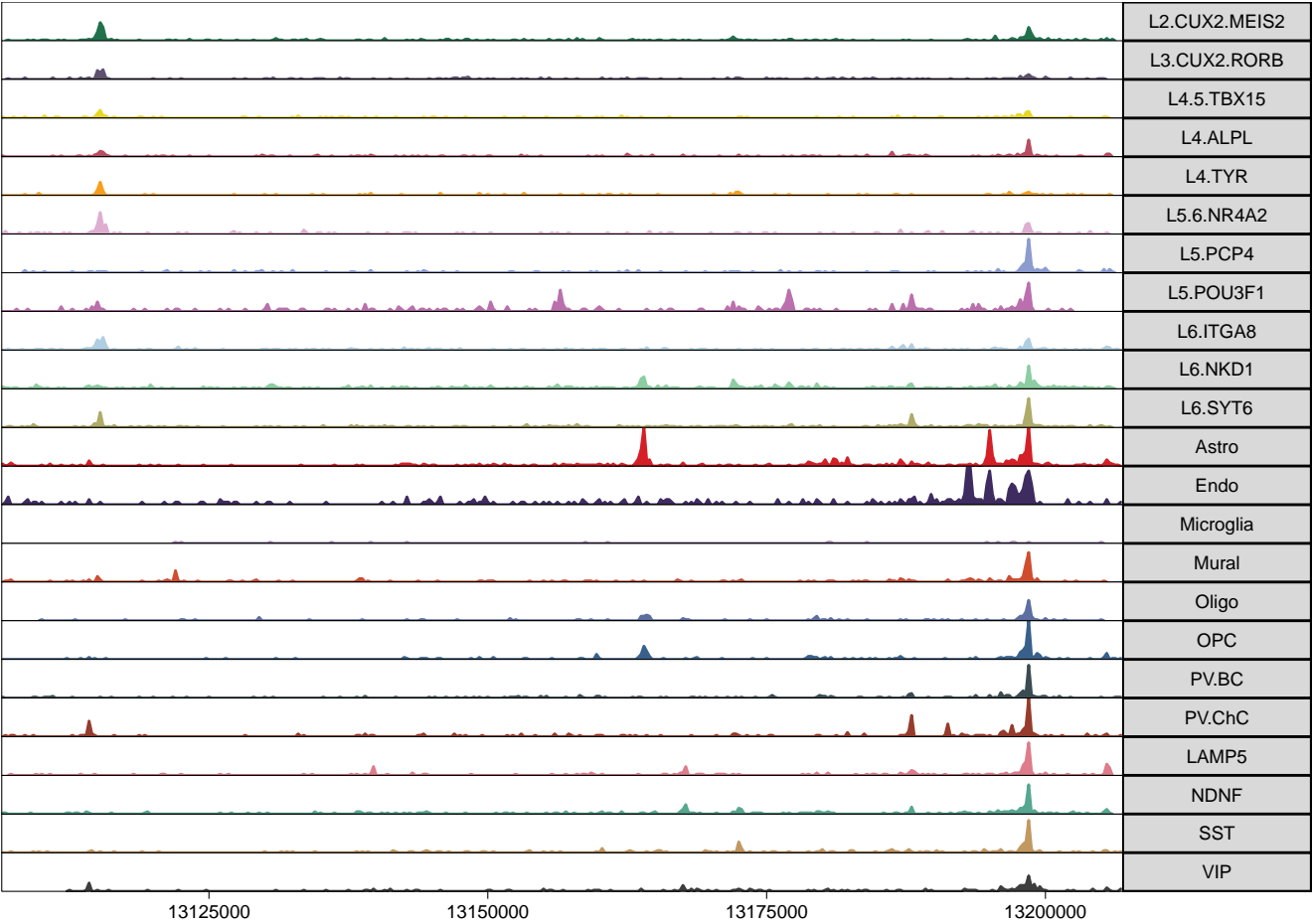


Peaks

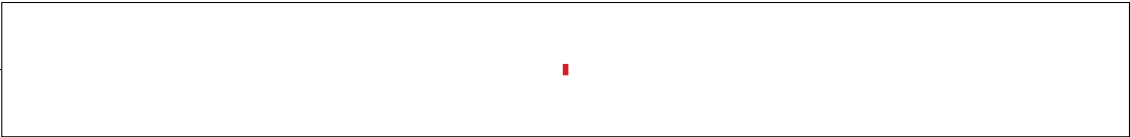


Genes

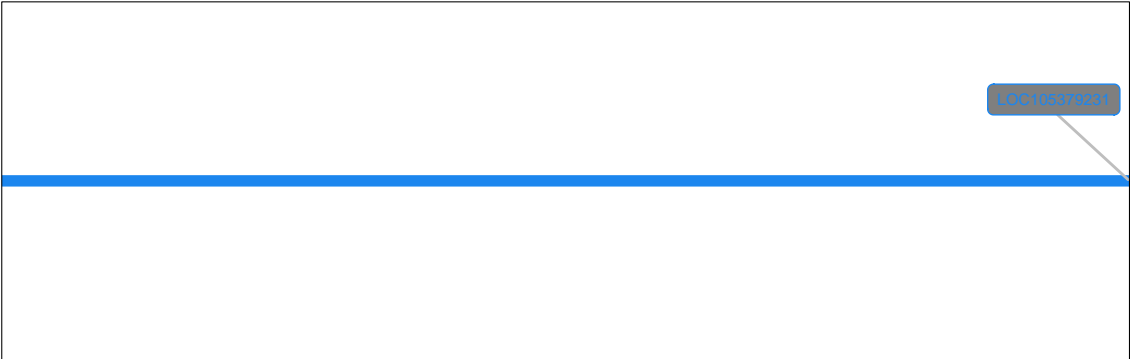
Coverage  
(Norm. ATAC Signal Range (0–1.2) by ReadsInTSS)



SST

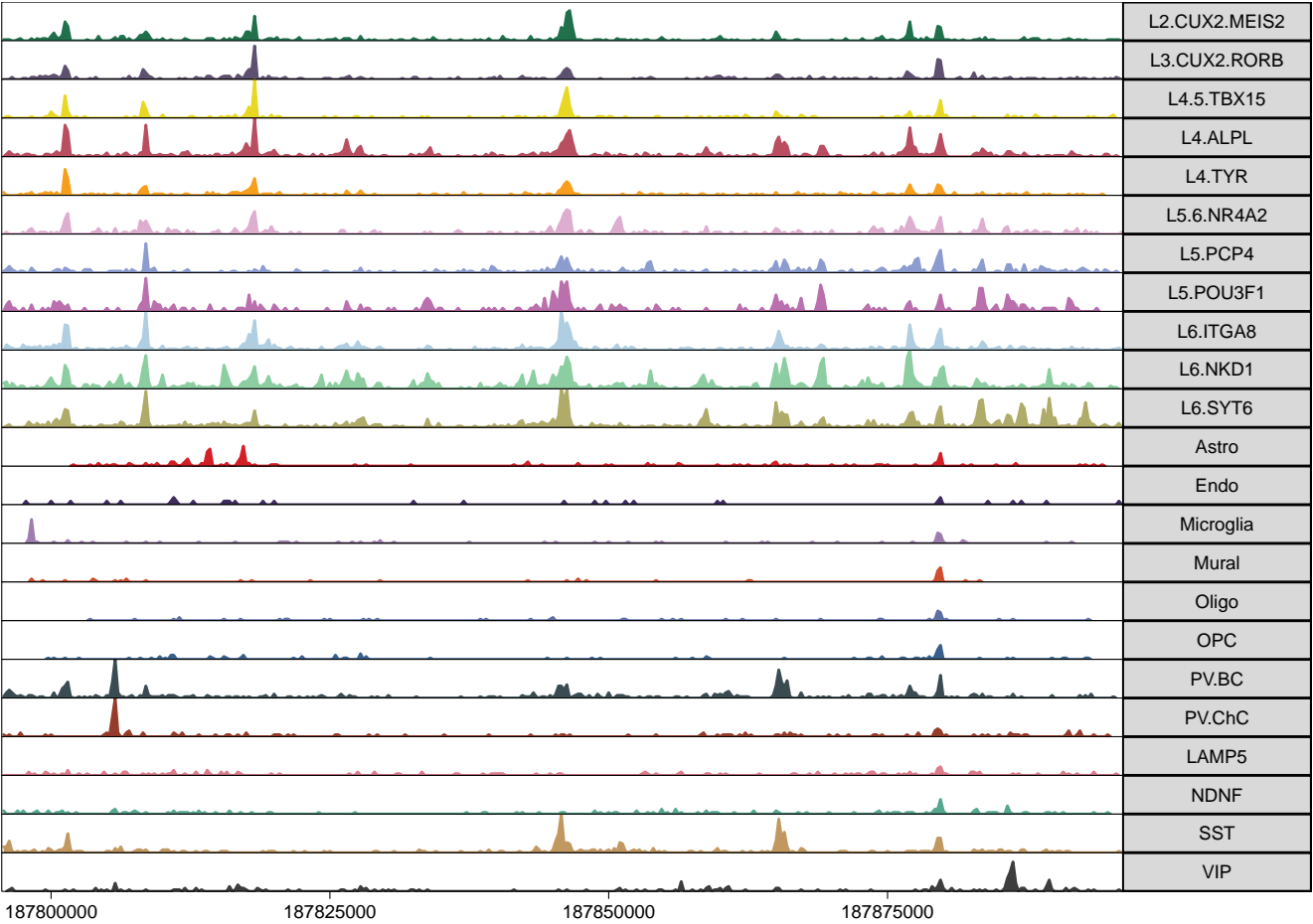


Peaks

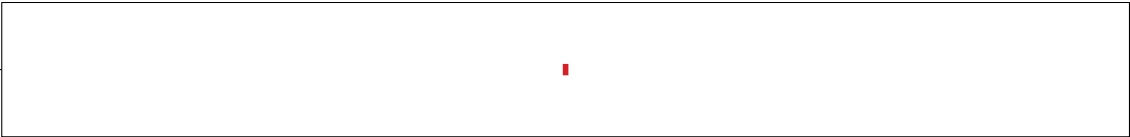


Genes

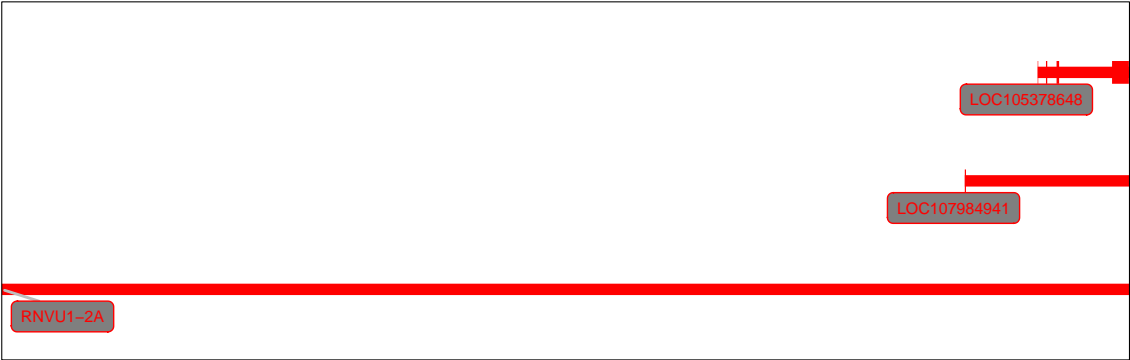
Coverage  
(Norm. ATAC Signal Range (0–0.86) by ReadsInTSS)



SST

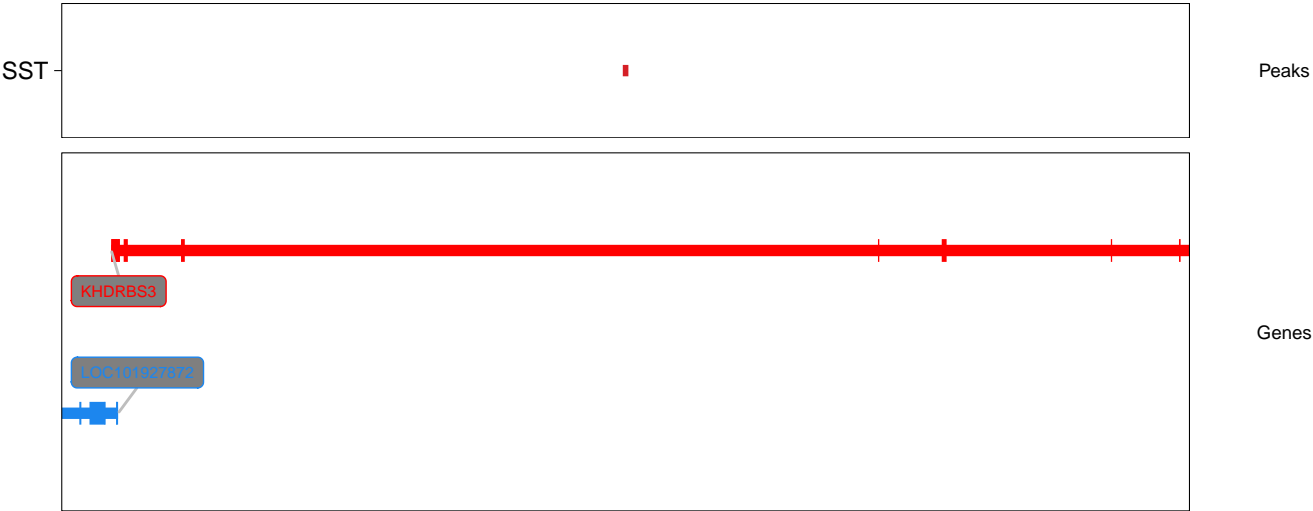
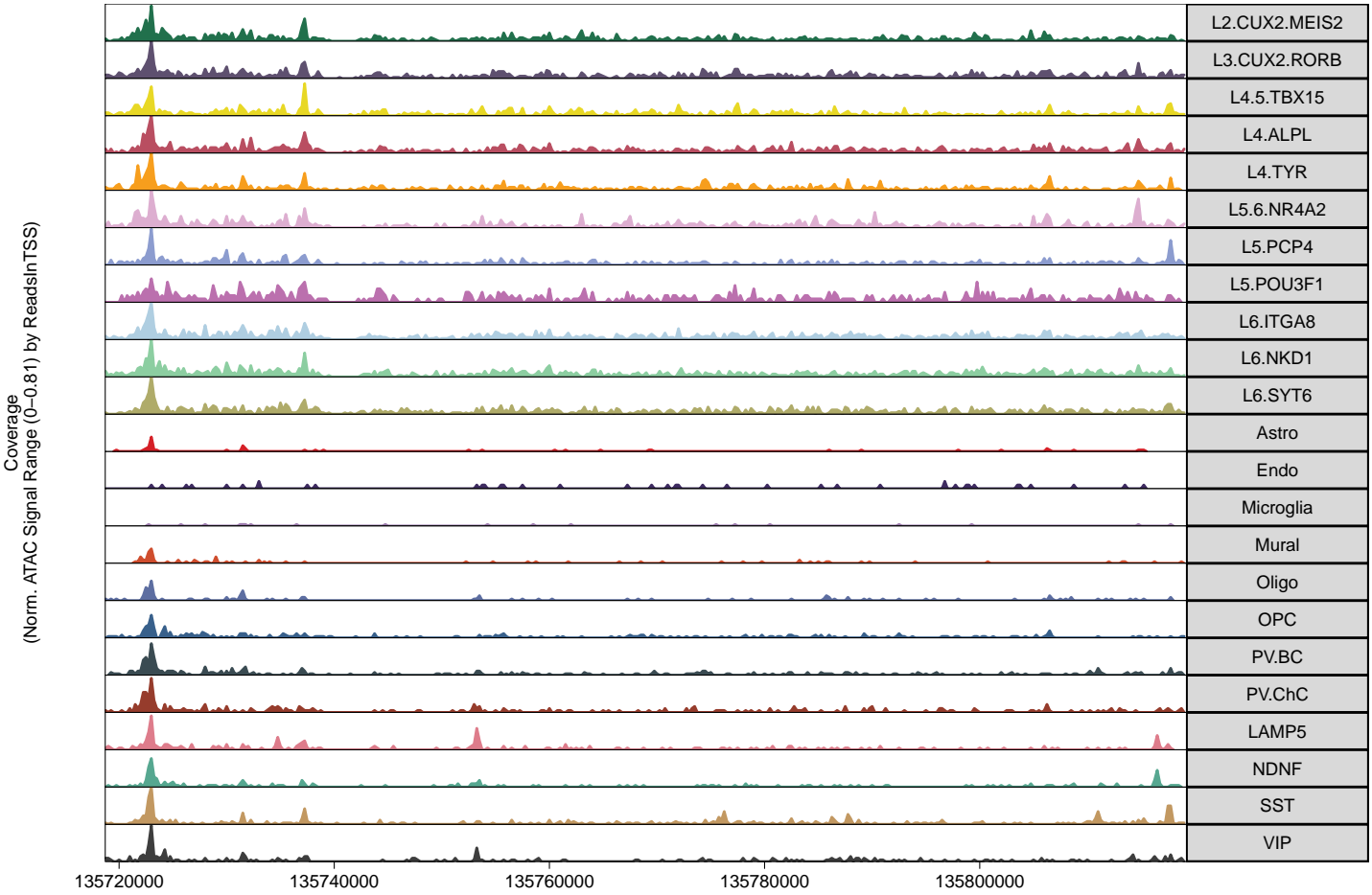


Peaks

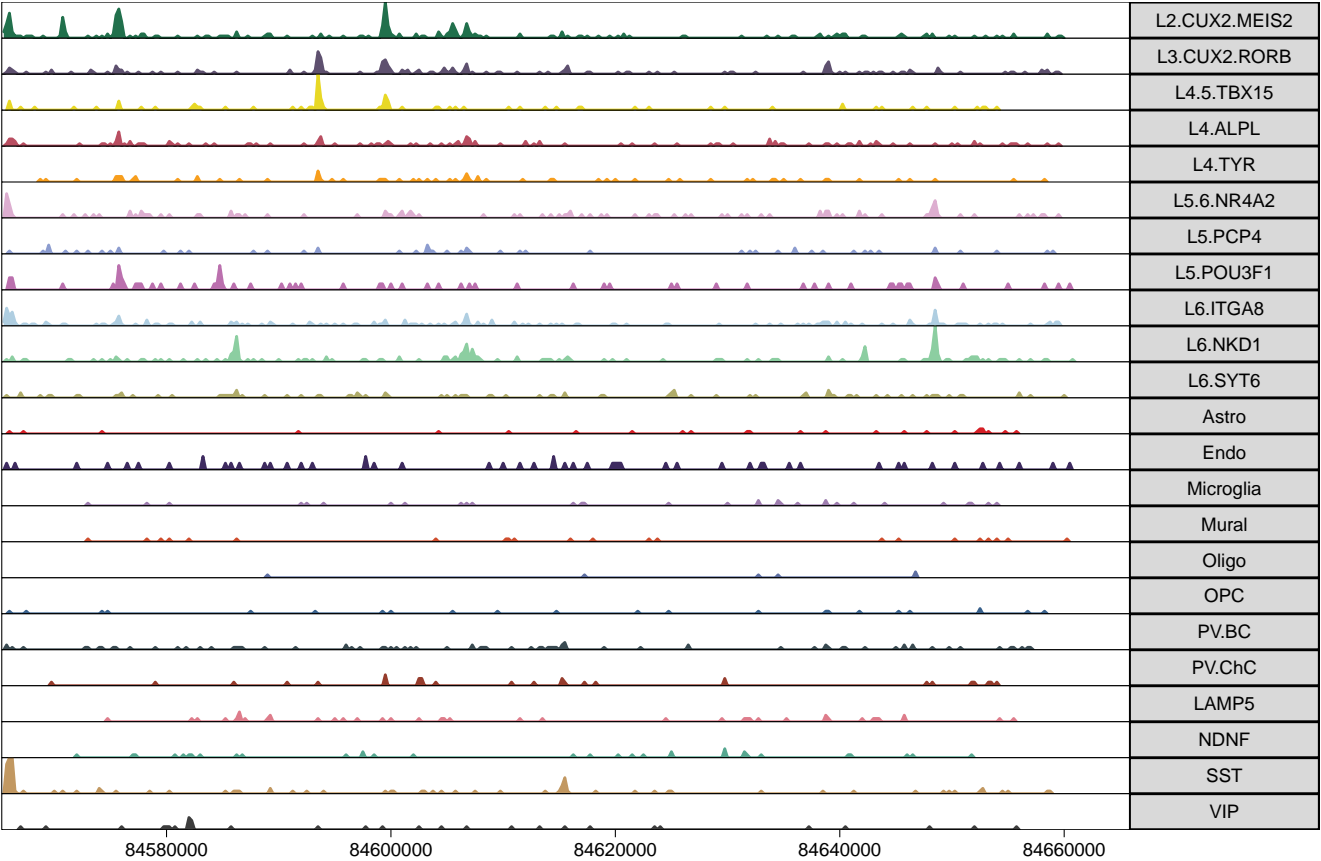


Genes

chr8:135718711–135819212 Marker Peak Tracks for: SST



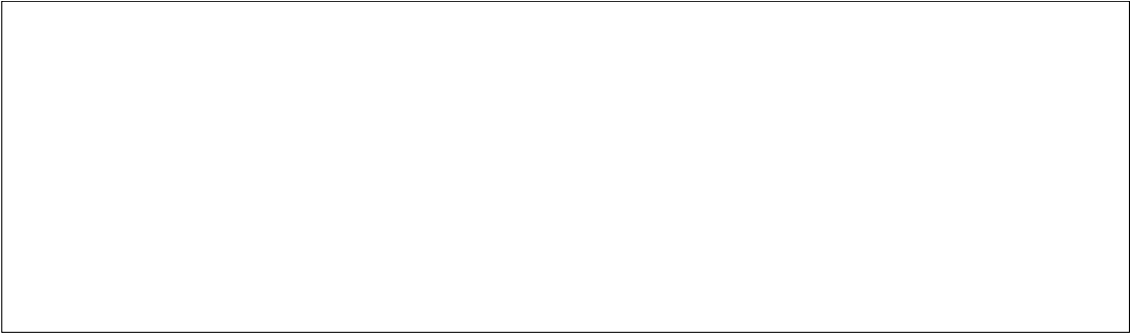
Coverage  
(Norm. ATAC Signal Range (0-0.43) by ReadsInTSS)



SST

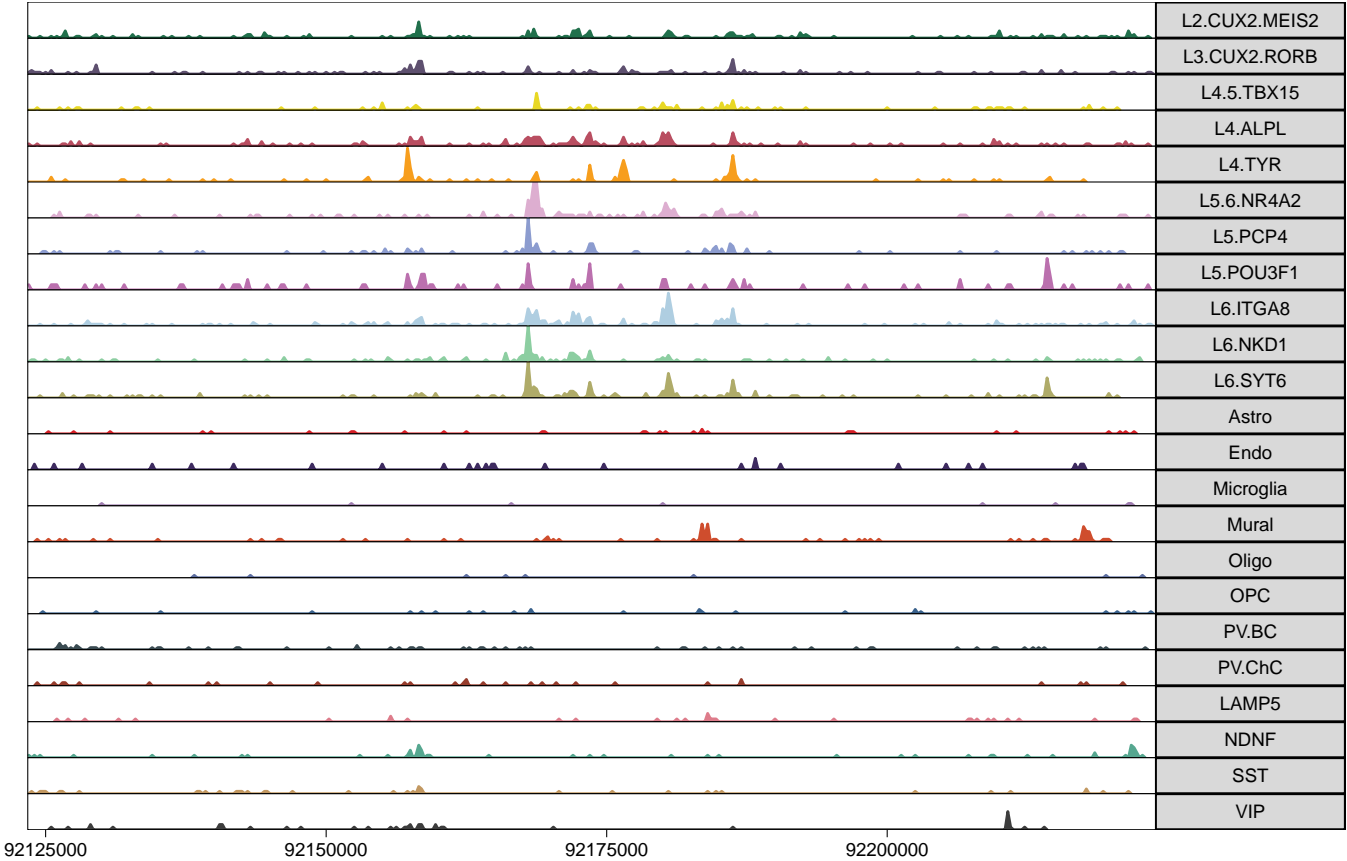


Peaks



GeneTrack

Coverage  
(Norm. ATAC Signal Range (0–0.5) by ReadsInTSS)



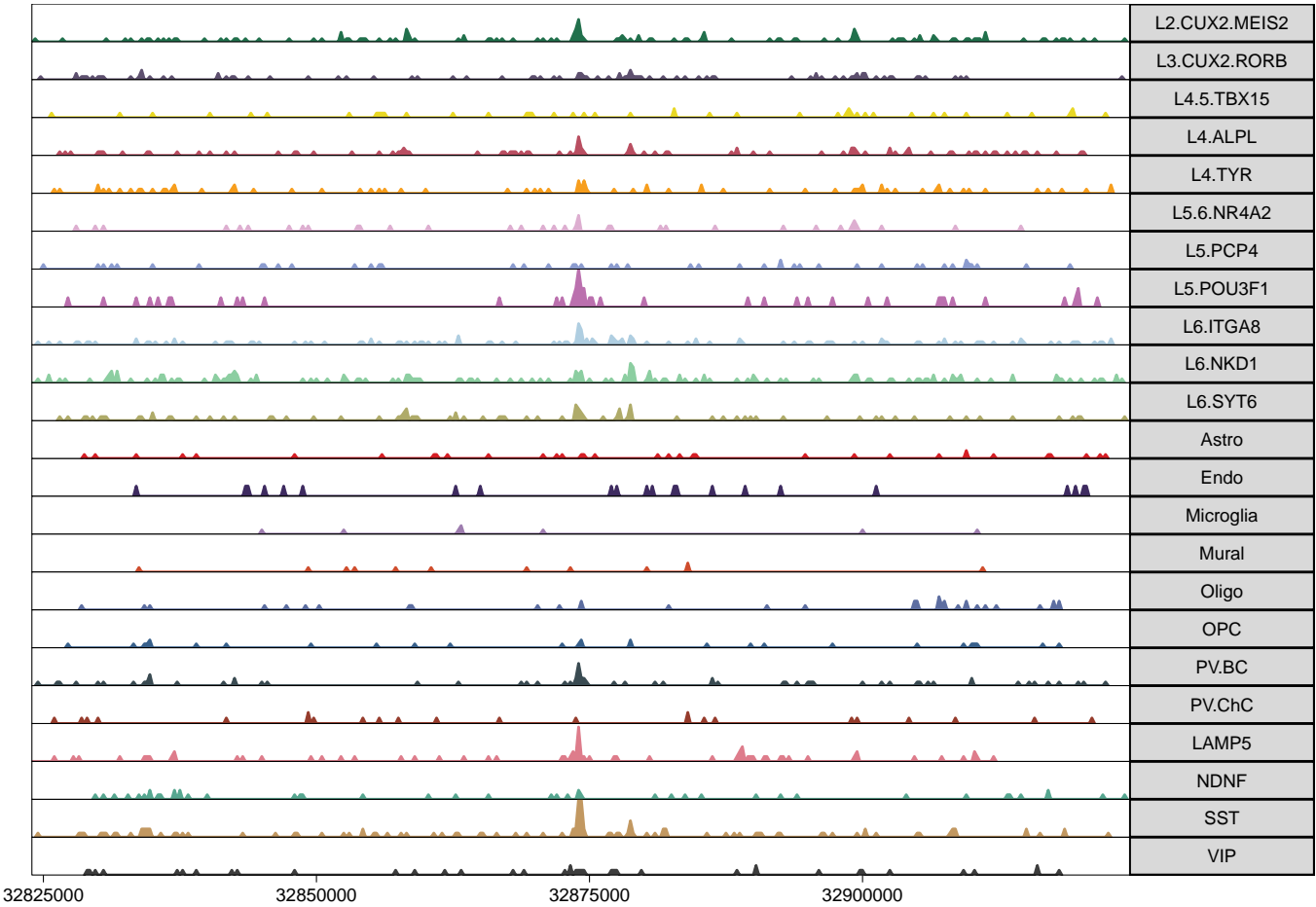
SST

Peaks

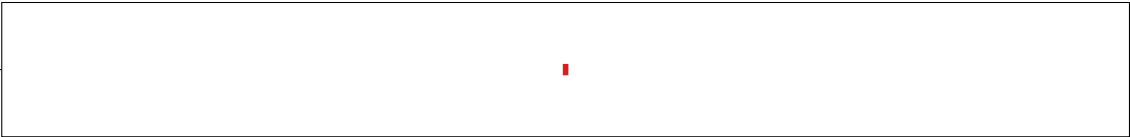
GeneTrack

strand ● ·

Coverage  
(Norm. ATAC Signal Range (0–0.3) by ReadsInTSS)



SST

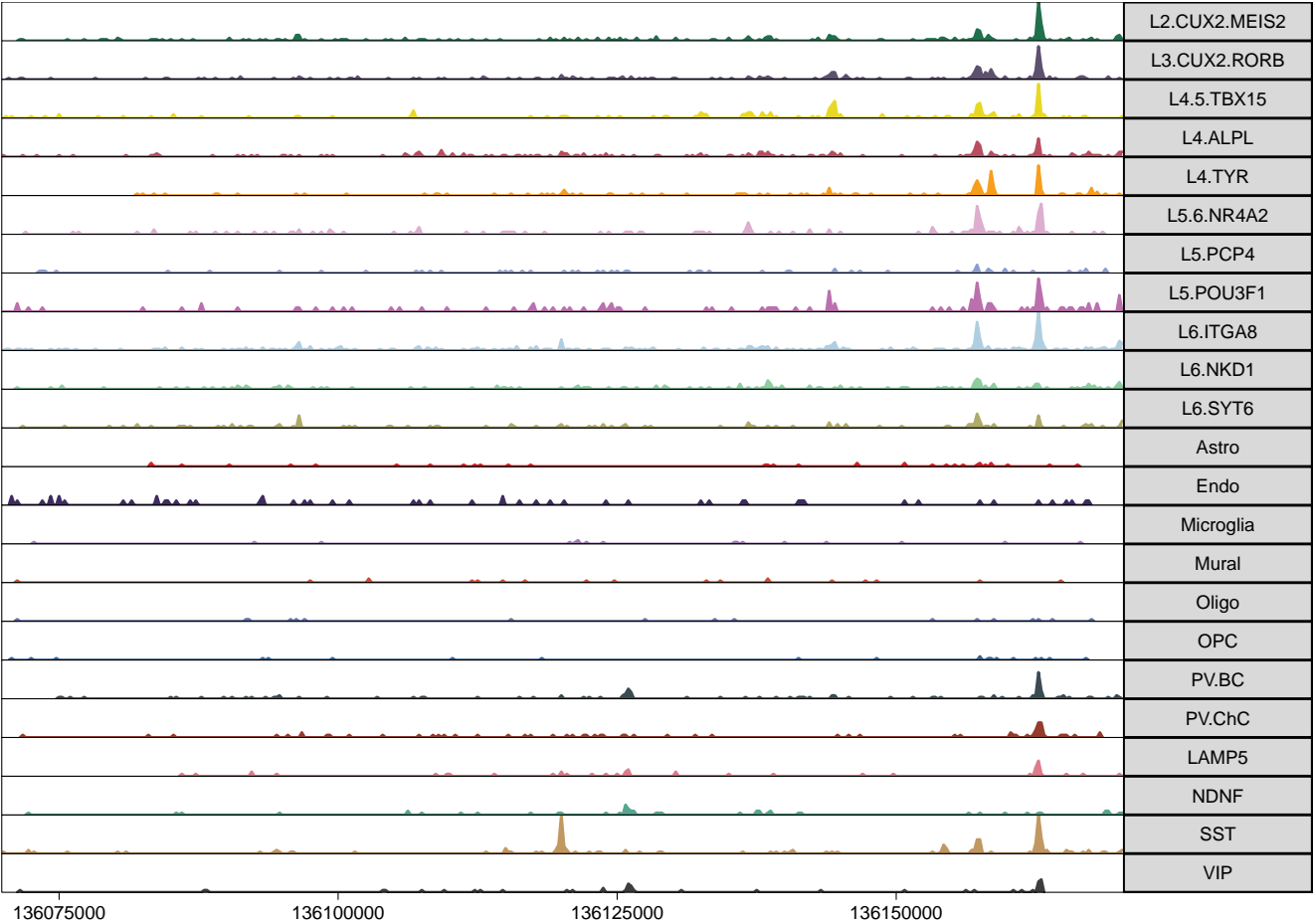


Peaks

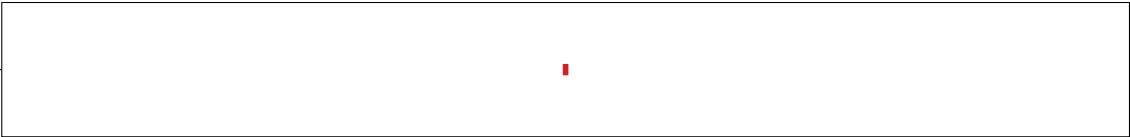


Genes

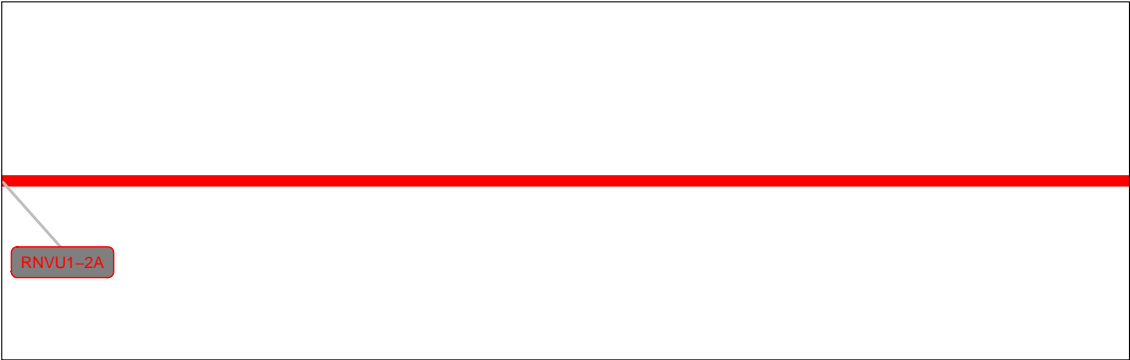
Coverage  
(Norm. ATAC Signal Range (0–0.69) by ReadsInTSS)



SST



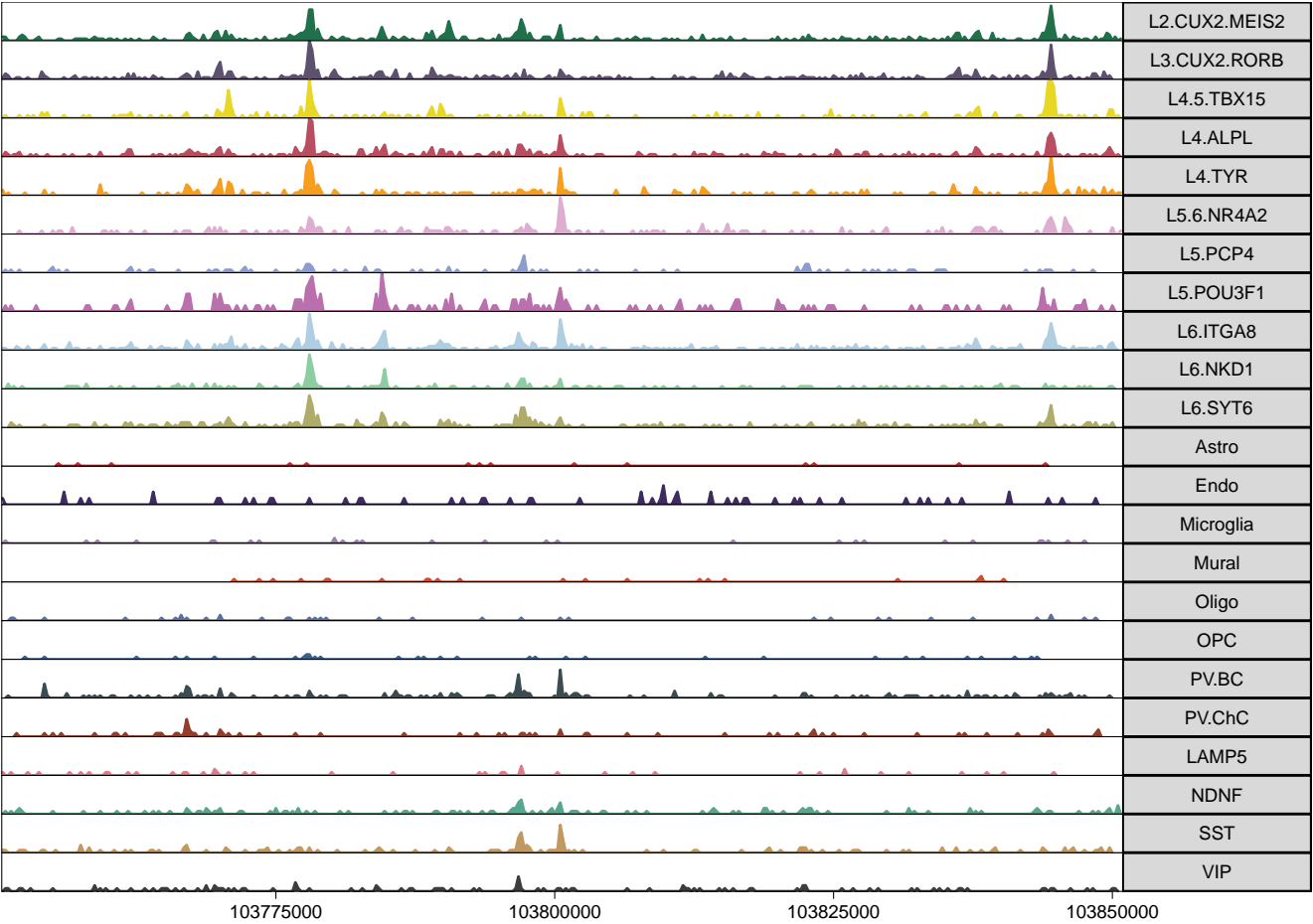
Peaks



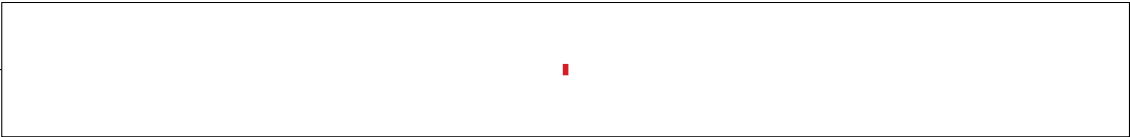
Genes



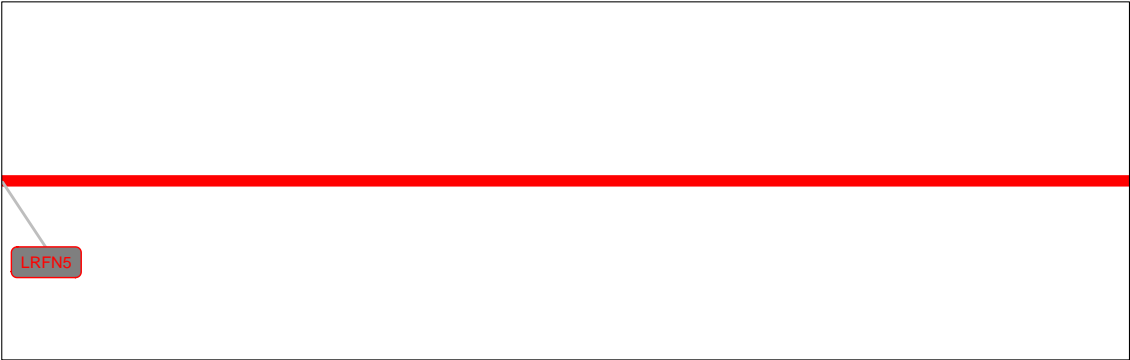
Coverage  
(Norm. ATAC Signal Range (0-0.49) by ReadsInTSS)



SST

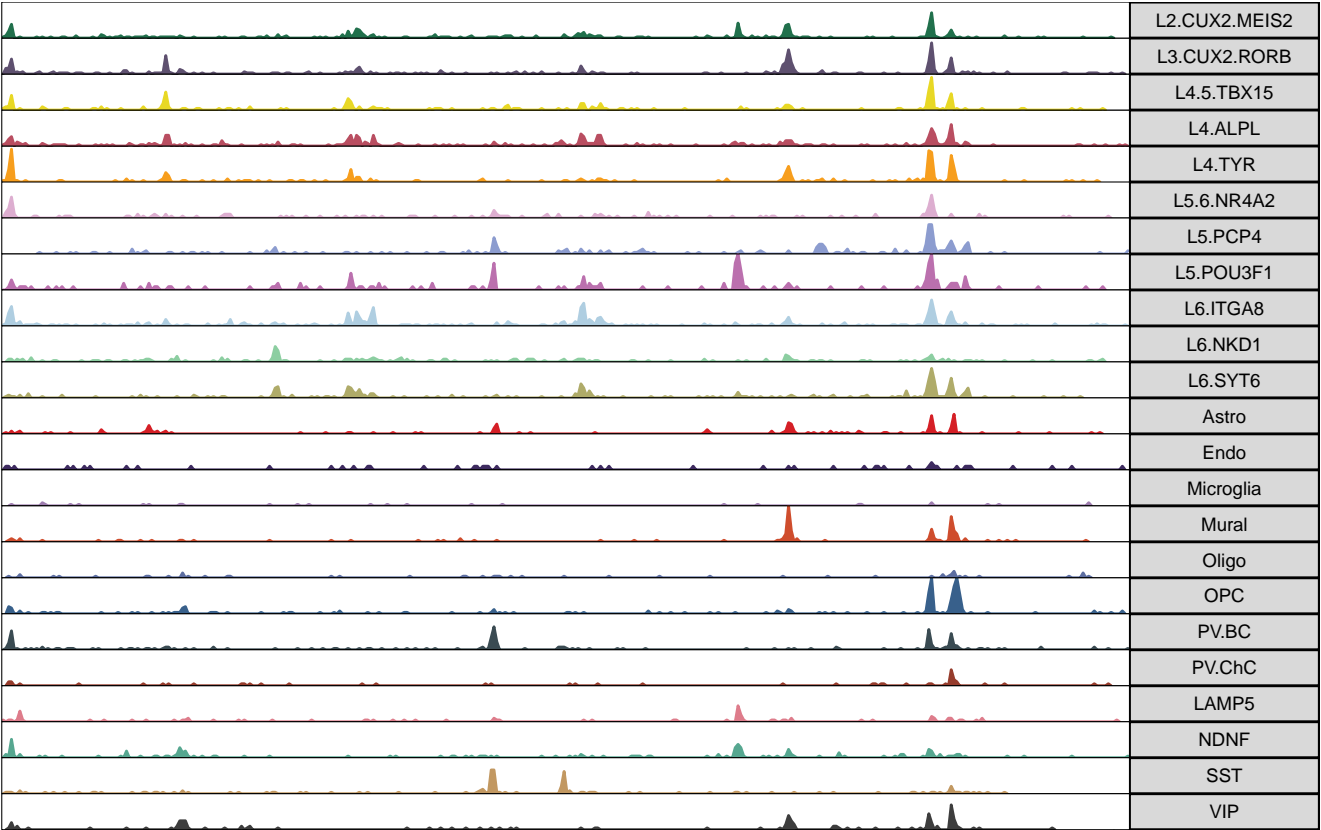


Peaks



Genes

Coverage  
(Norm. ATAC Signal Range (0–0.79) by ReadsInTSS)

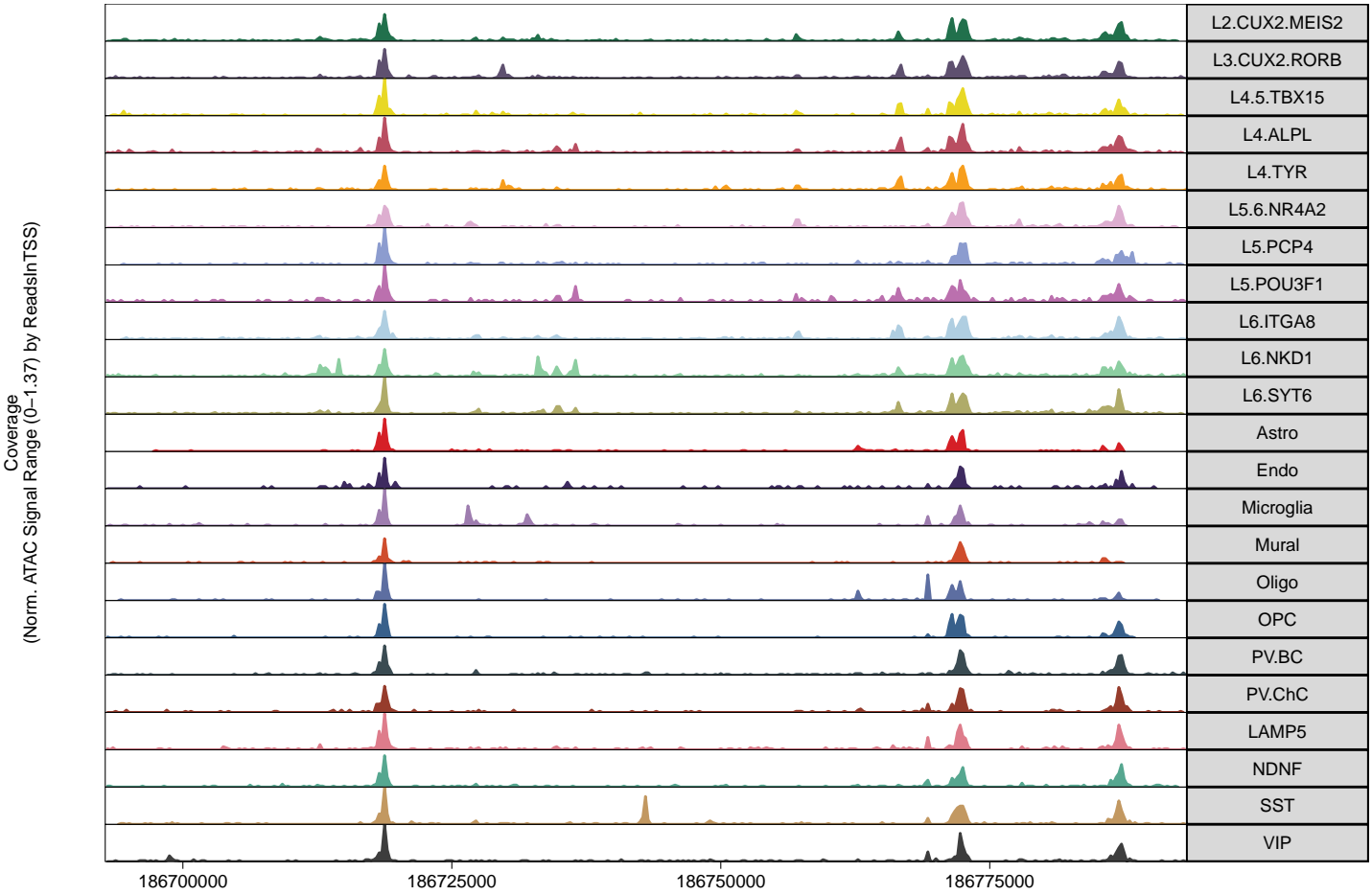


SST

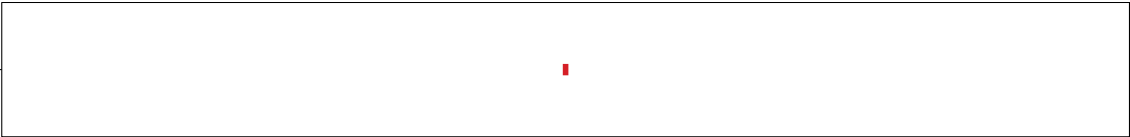


Peaks

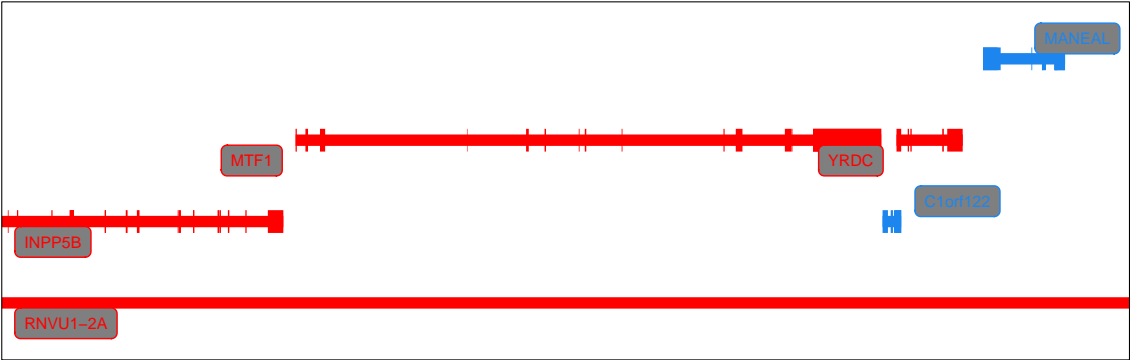
GeneTrack



SST



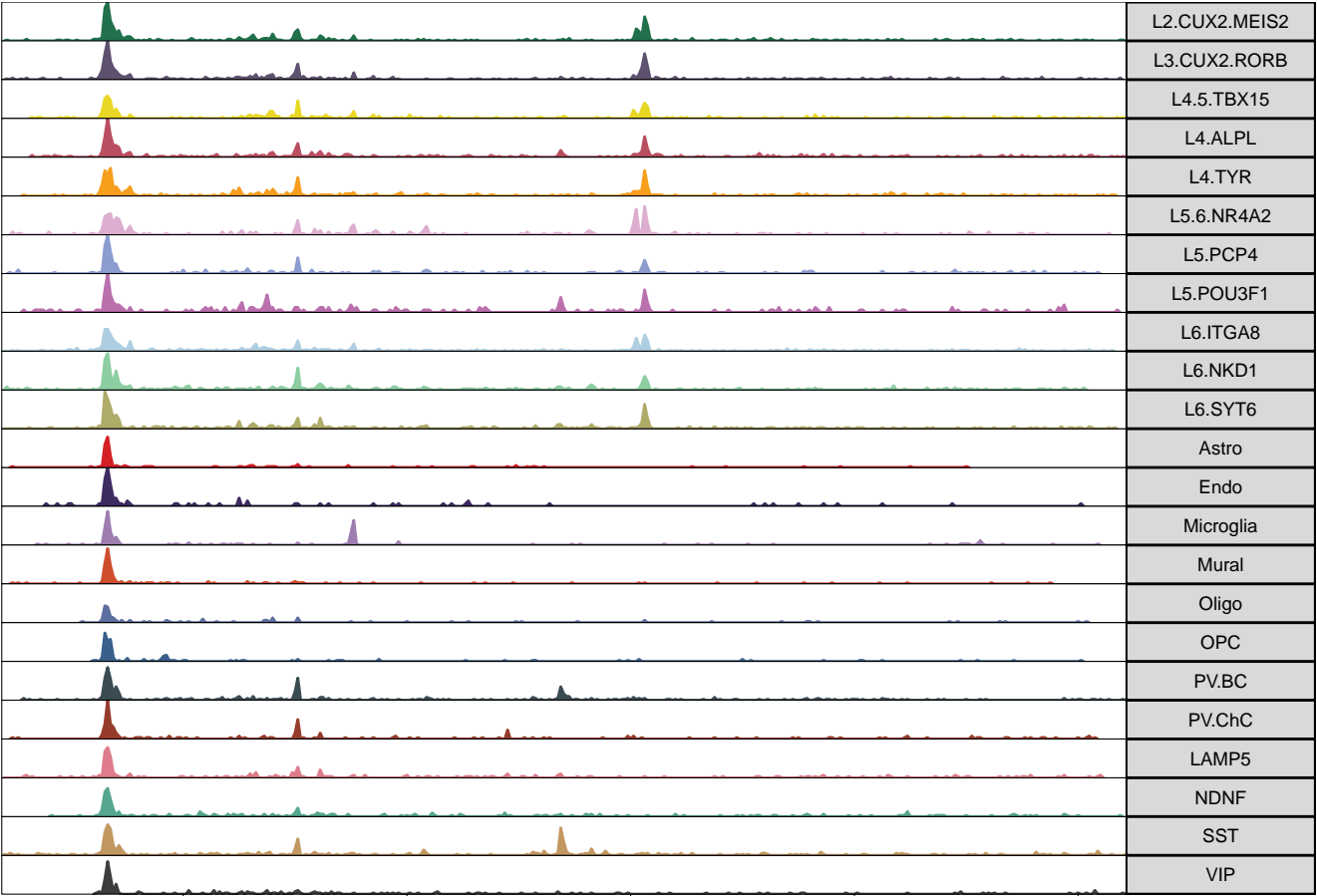
Peaks



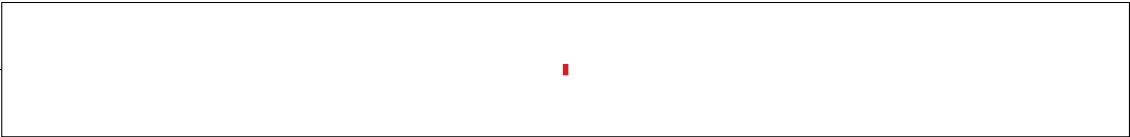
Genes

chr1:172083780–172184281 Marker Peak Tracks for: SST

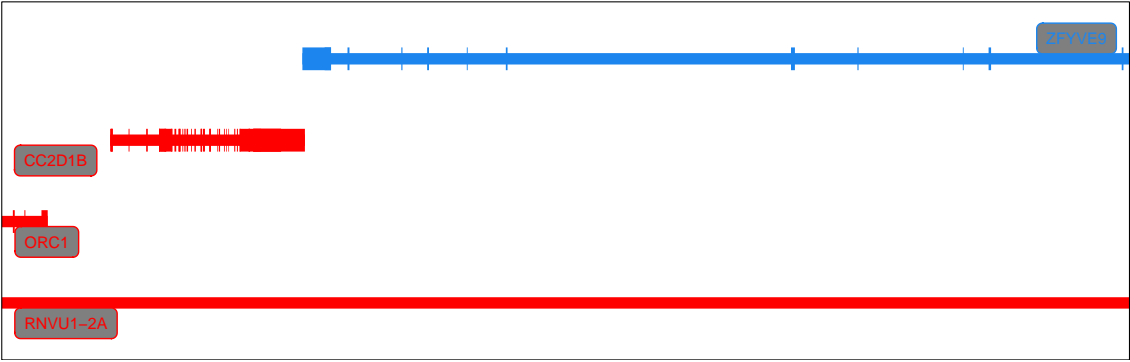
Coverage  
(Norm. ATAC Signal Range (0–1.4) by ReadsInTSS)



SST

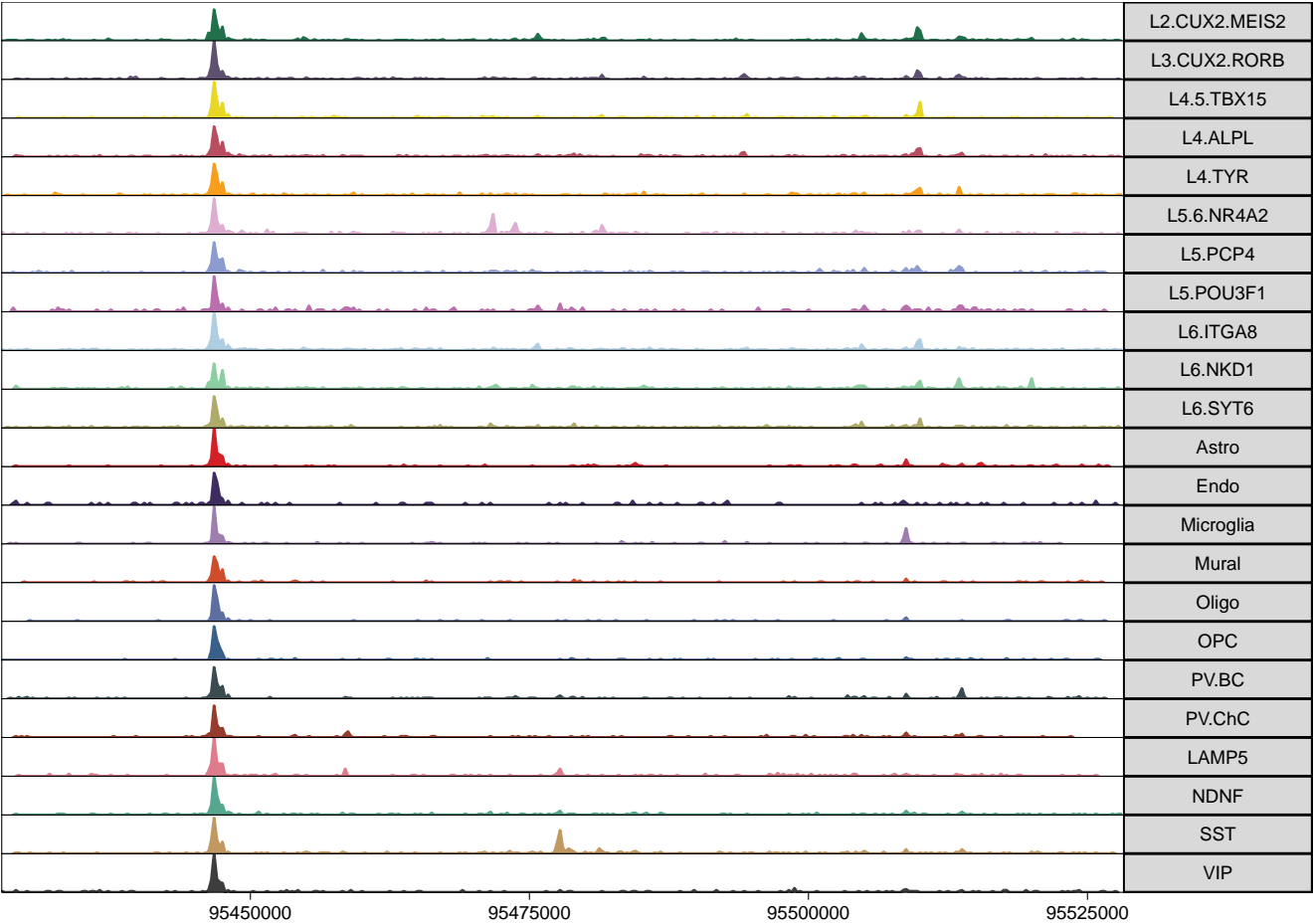


Peaks

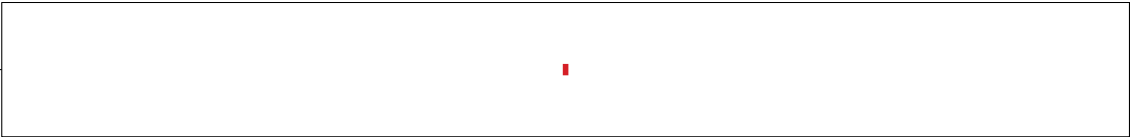


Genes

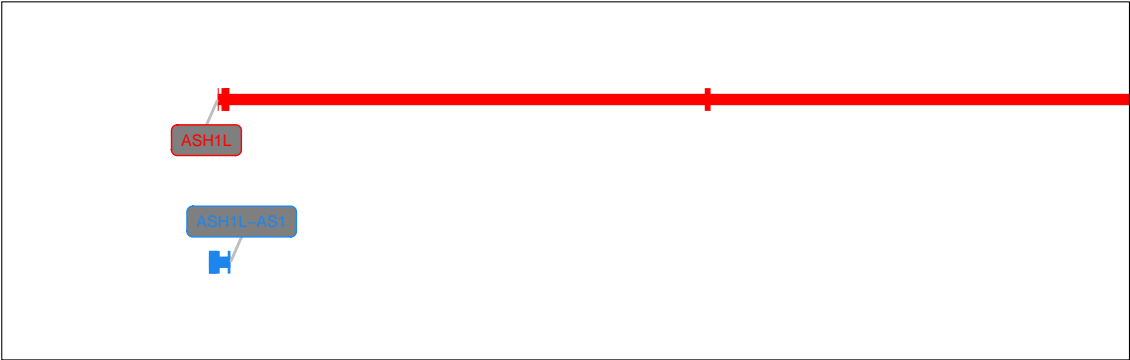
Coverage  
(Norm. ATAC Signal Range (0-1.45) by ReadsInTSS)



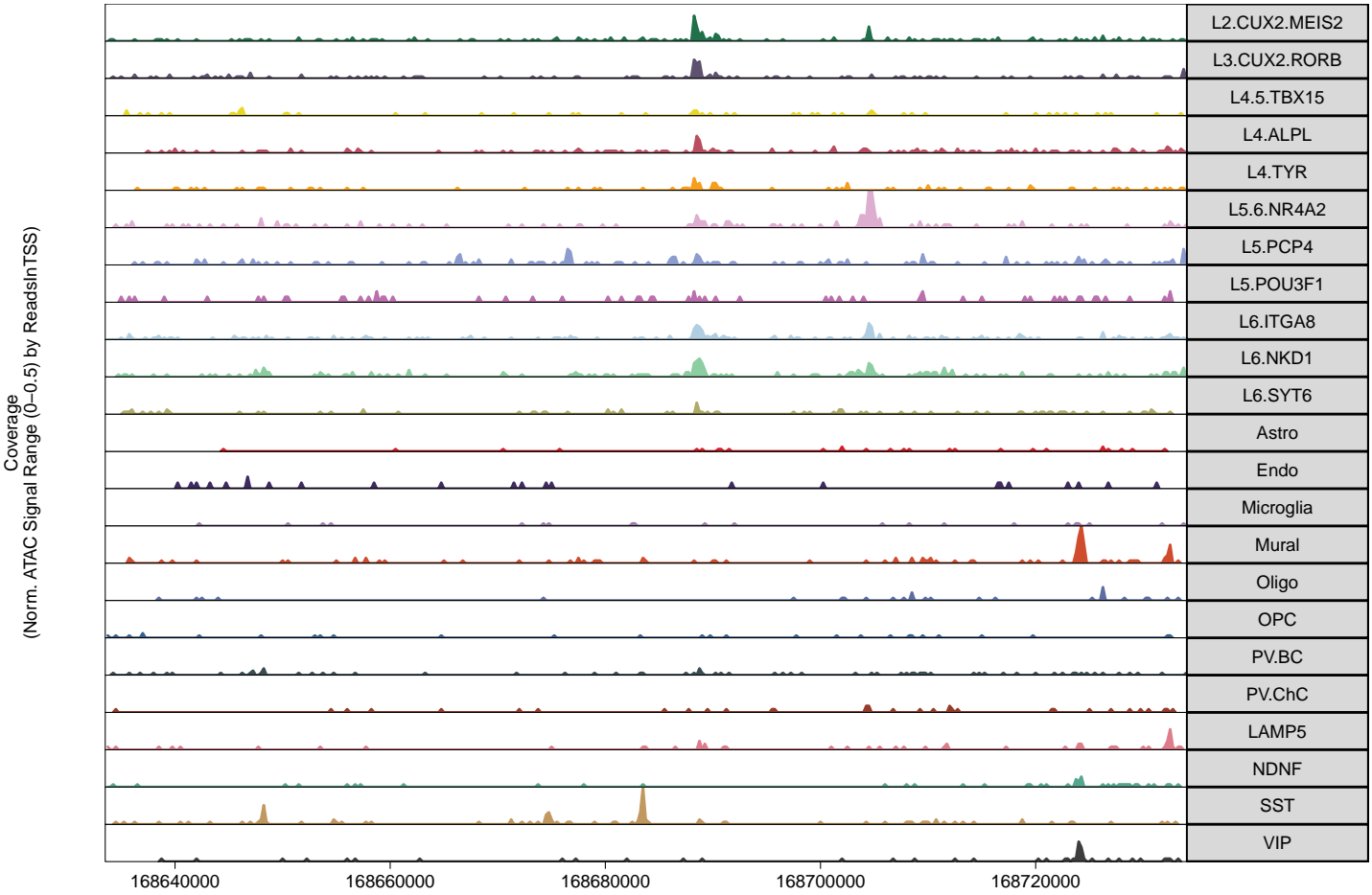
SST



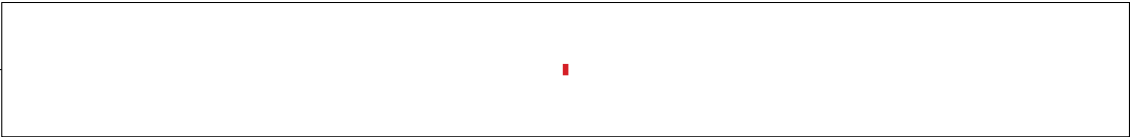
Peaks



Genes



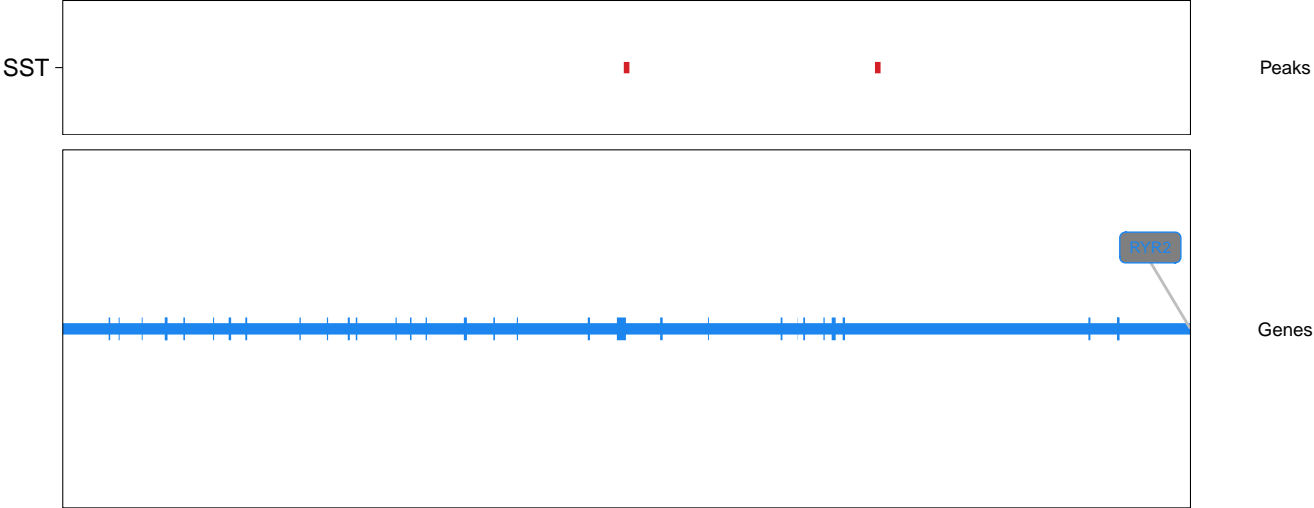
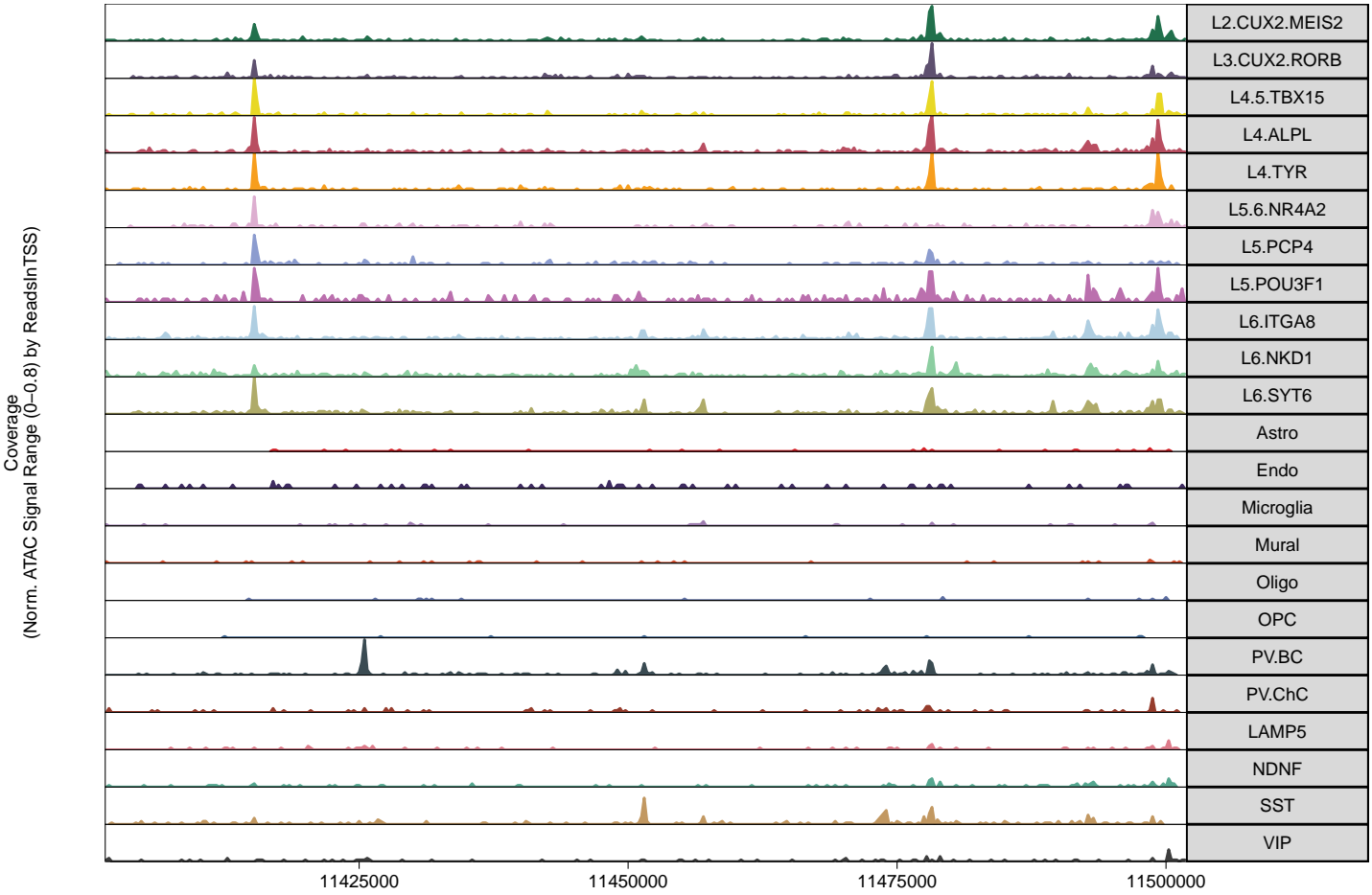
SST



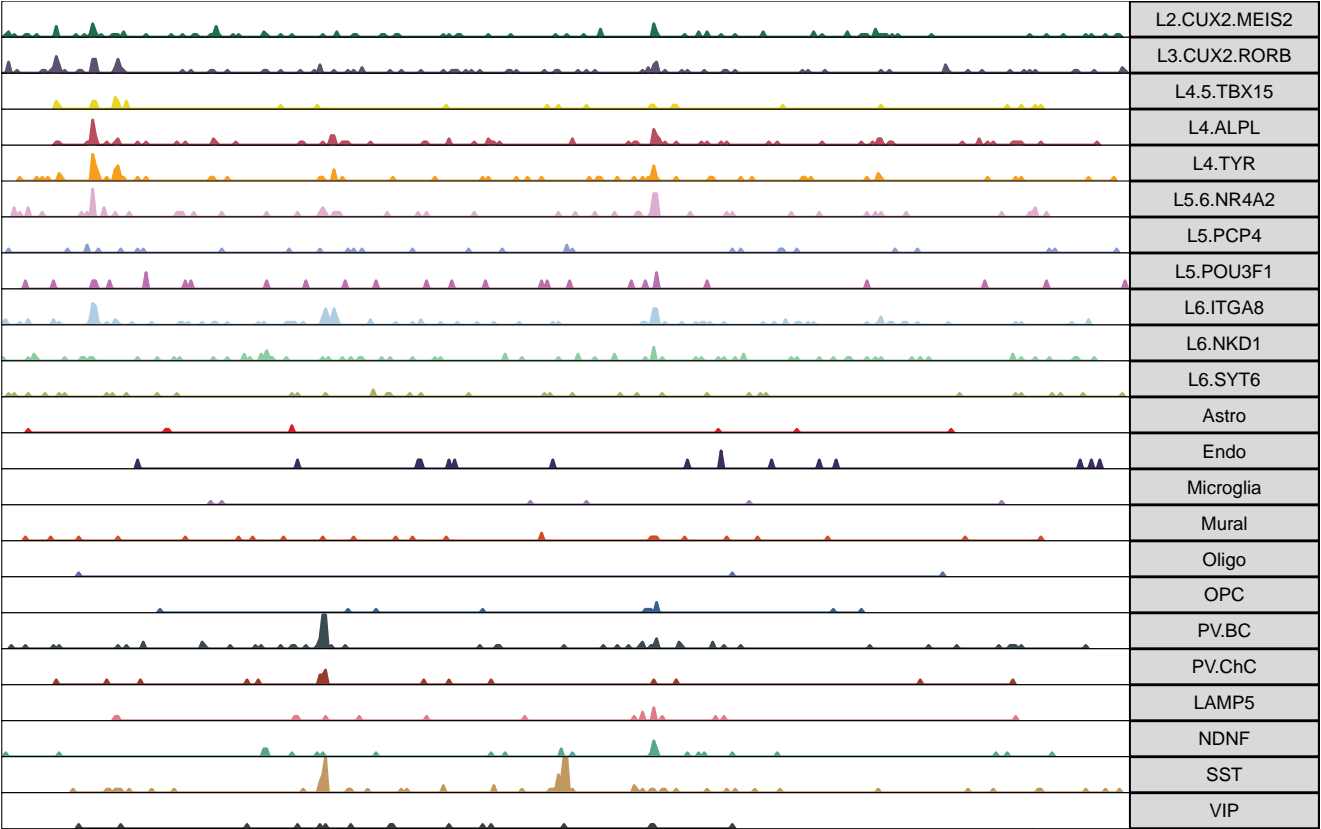
Peaks



Genes



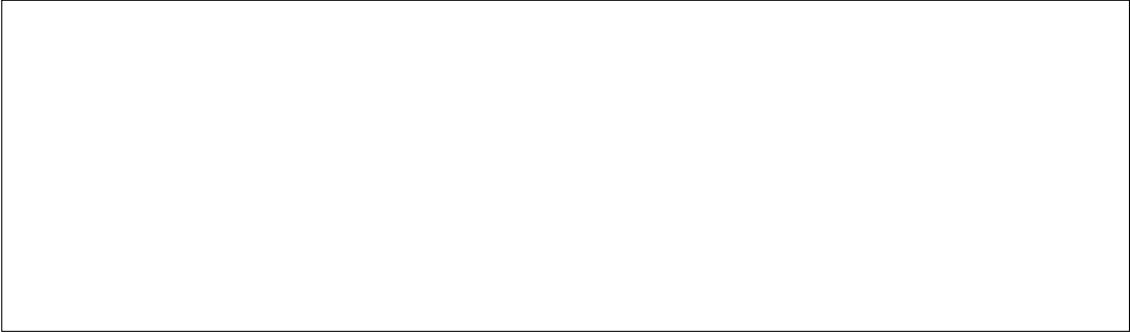
Coverage  
(Norm. ATAC Signal Range (0–0.32) by ReadsInTSS)



SST



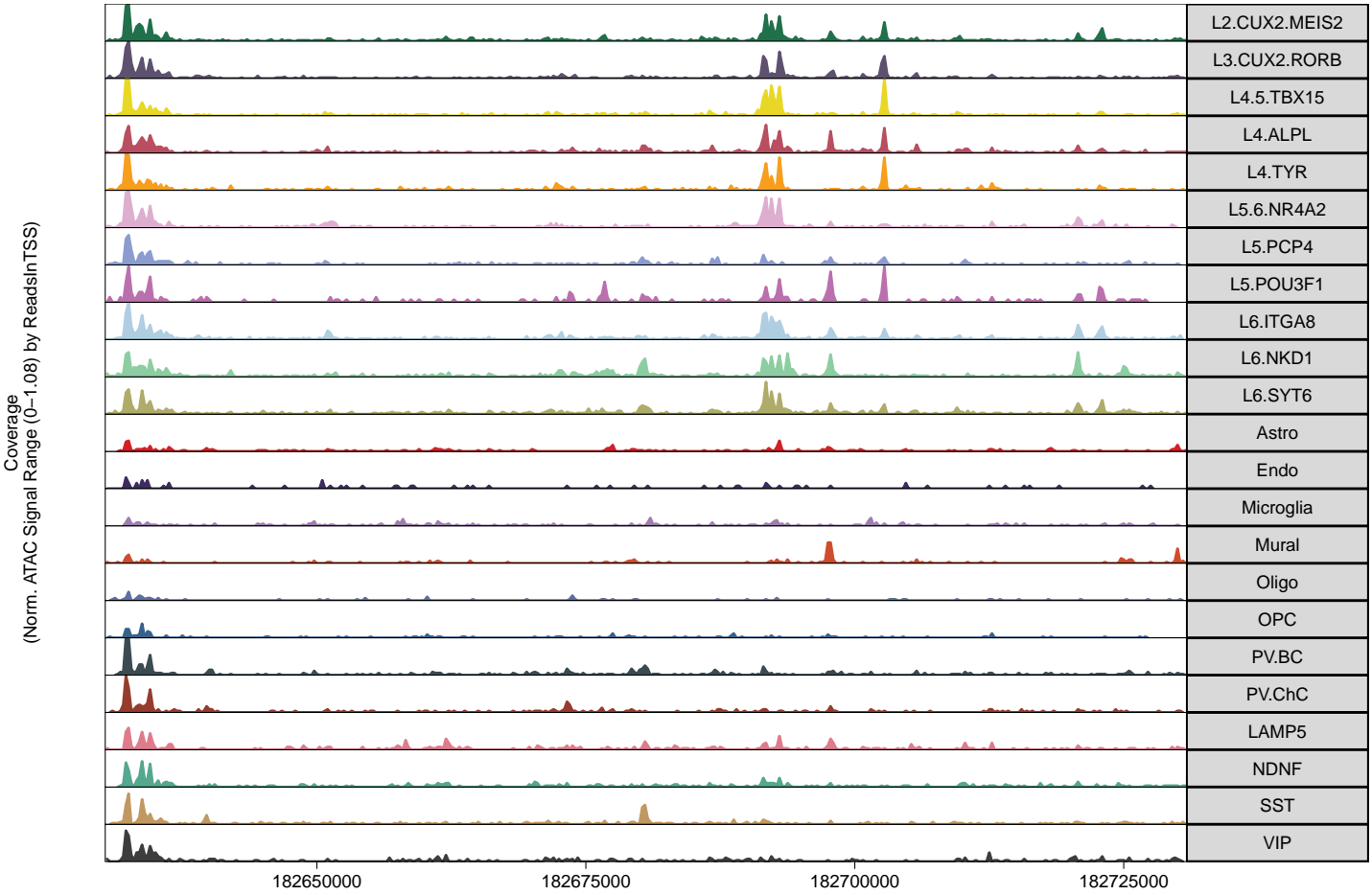
Peaks



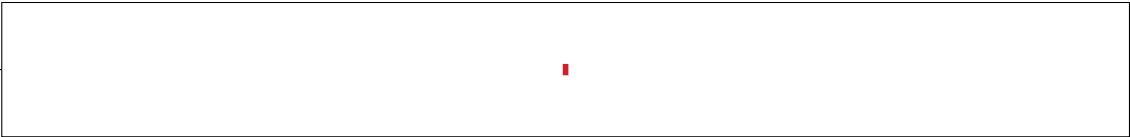
GeneTrack



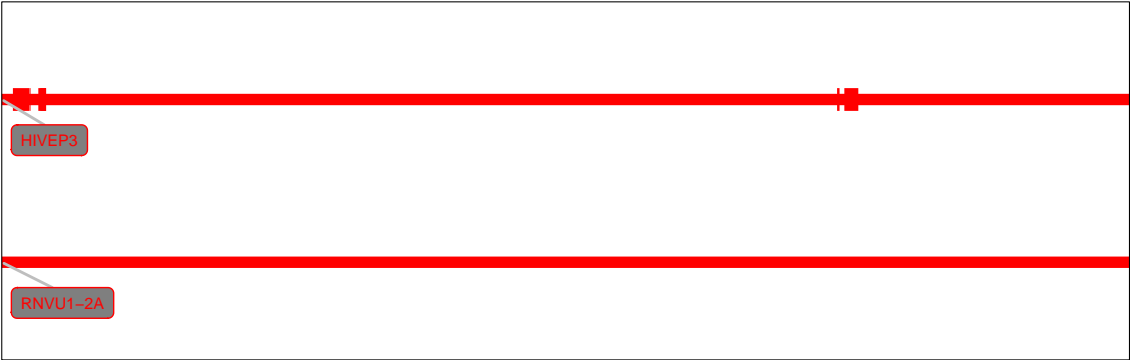
chr1:182630324–182730825 Marker Peak Tracks for: SST



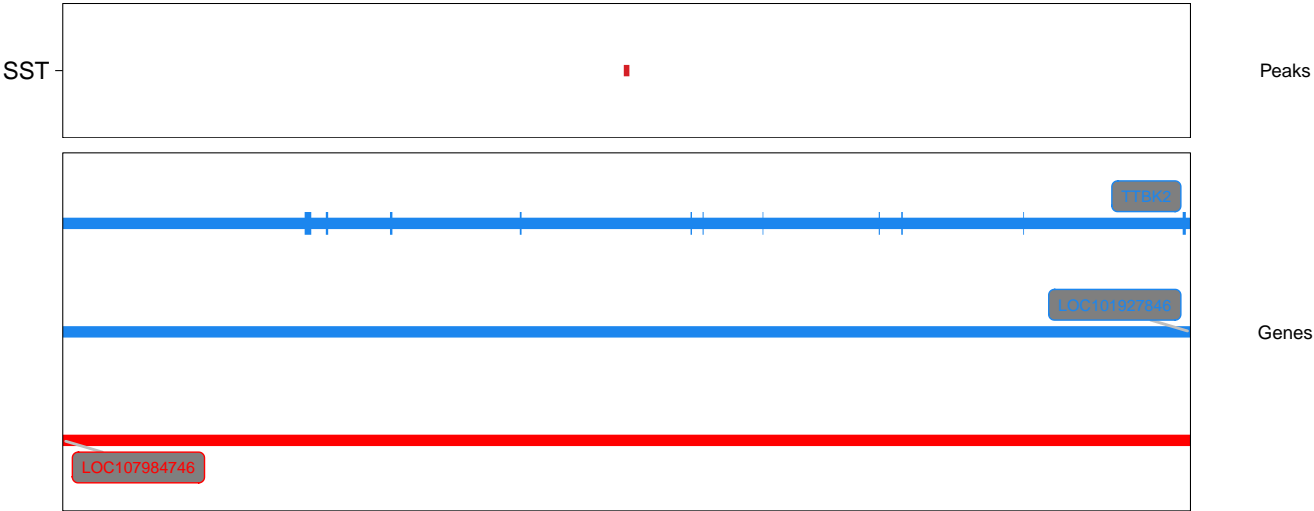
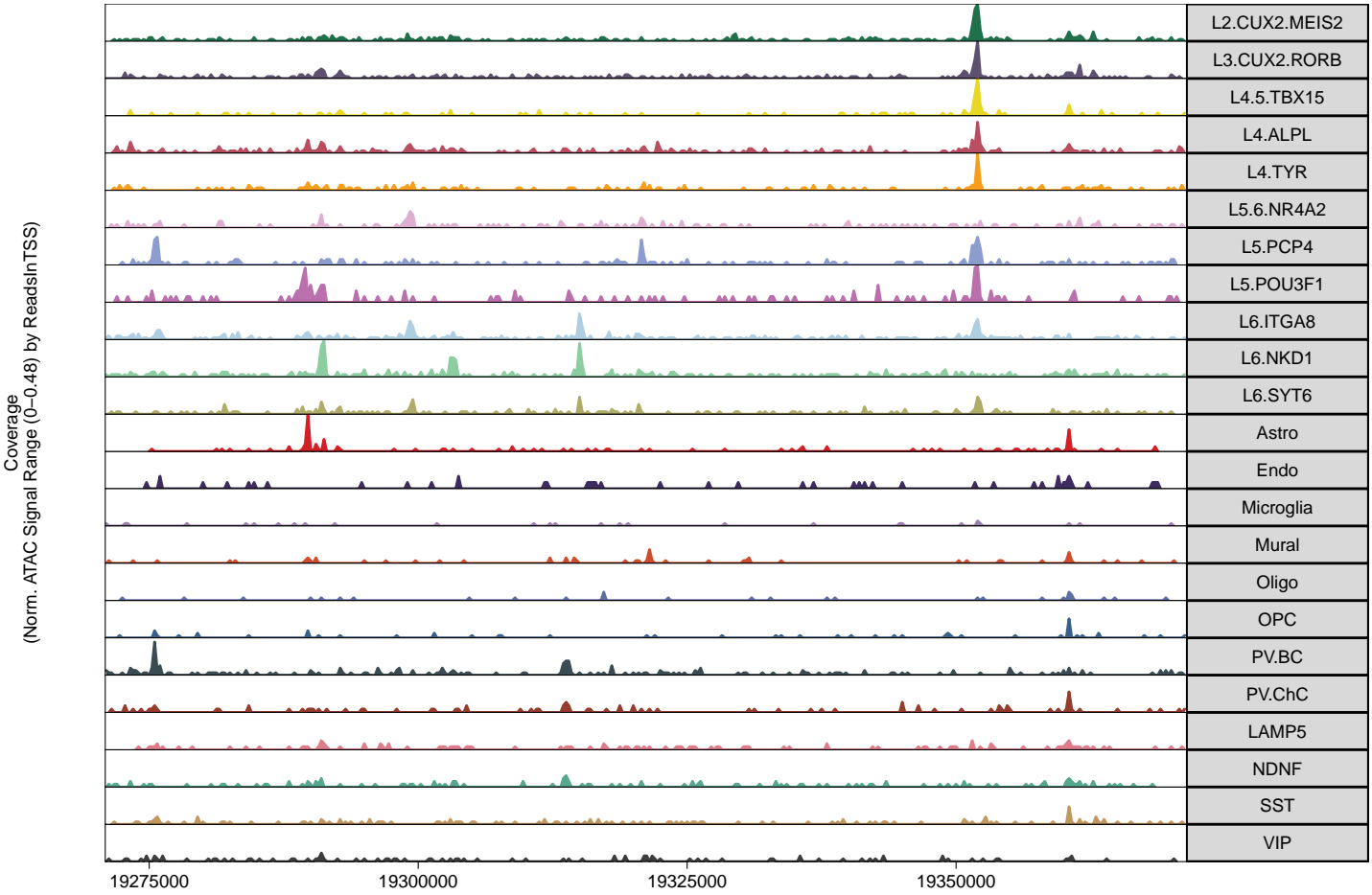
SST



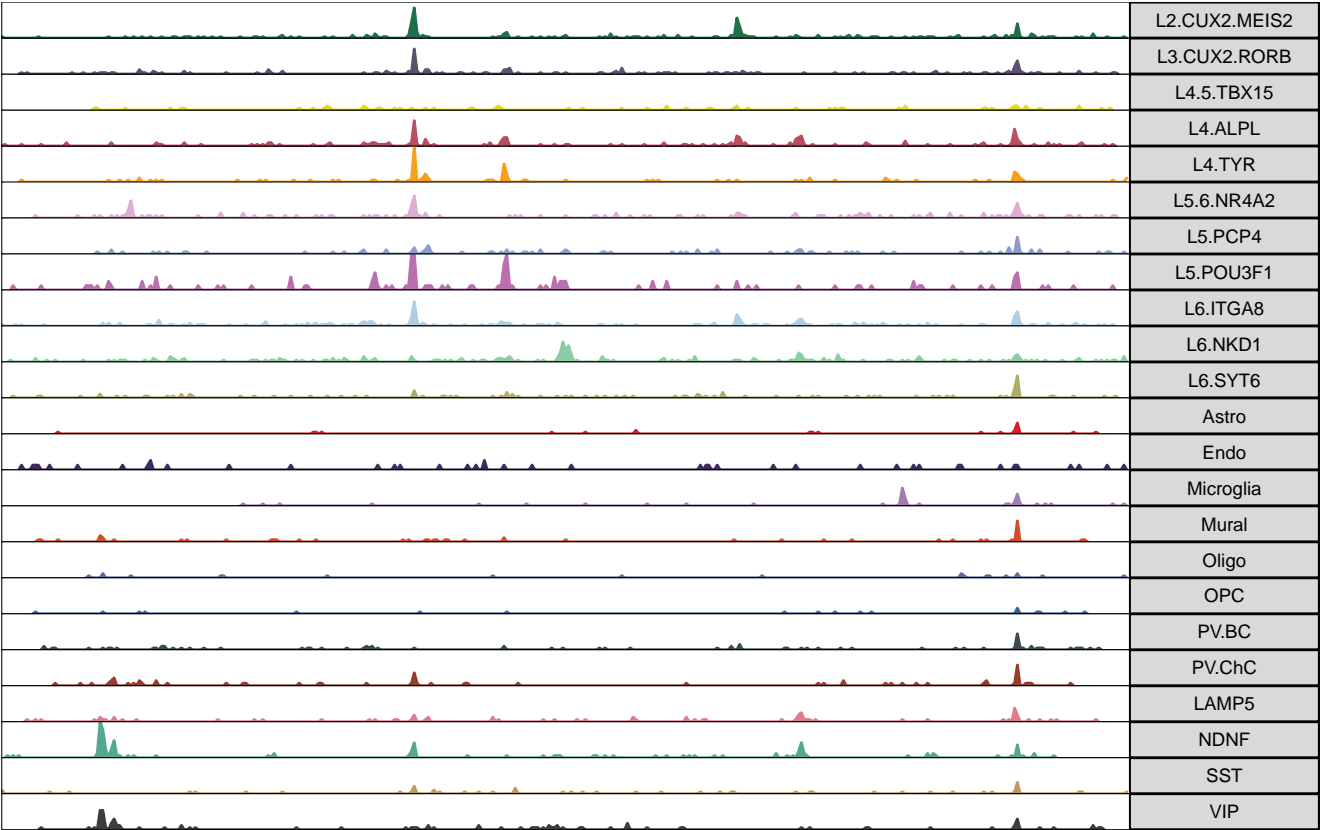
Peaks



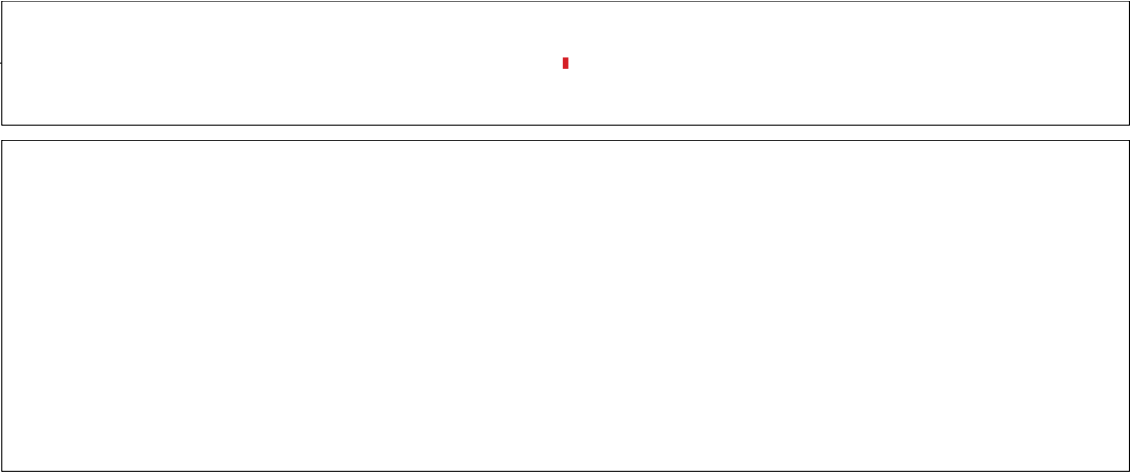
Genes



Coverage  
(Norm. ATAC Signal Range (0–0.61) by ReadsInTSS)

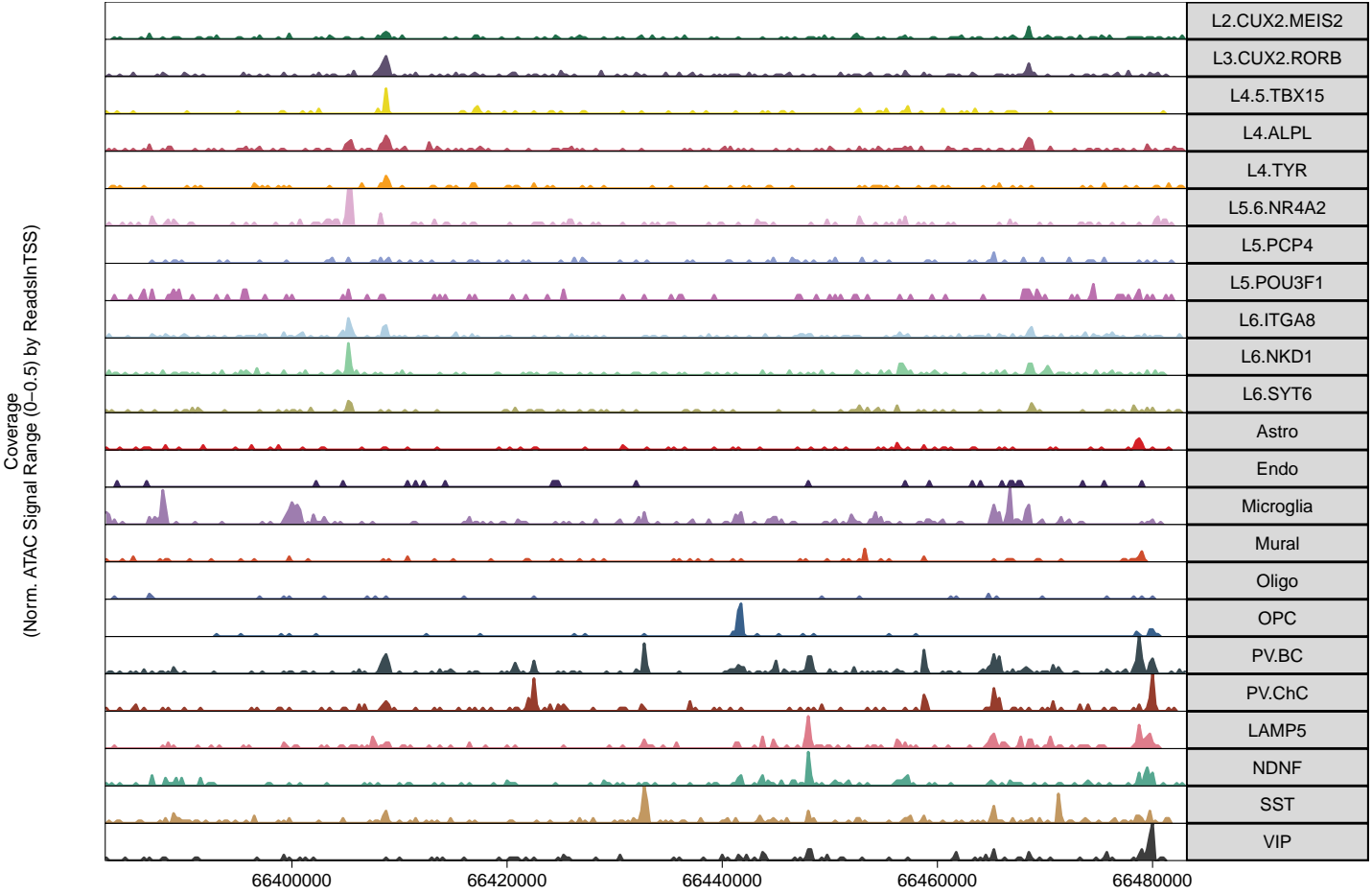


SST

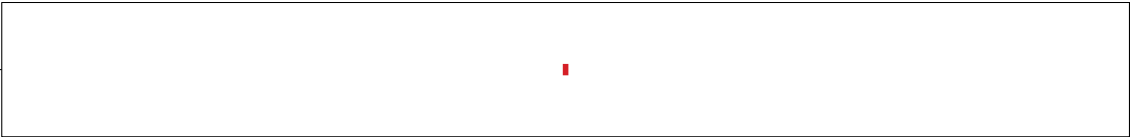


Peaks

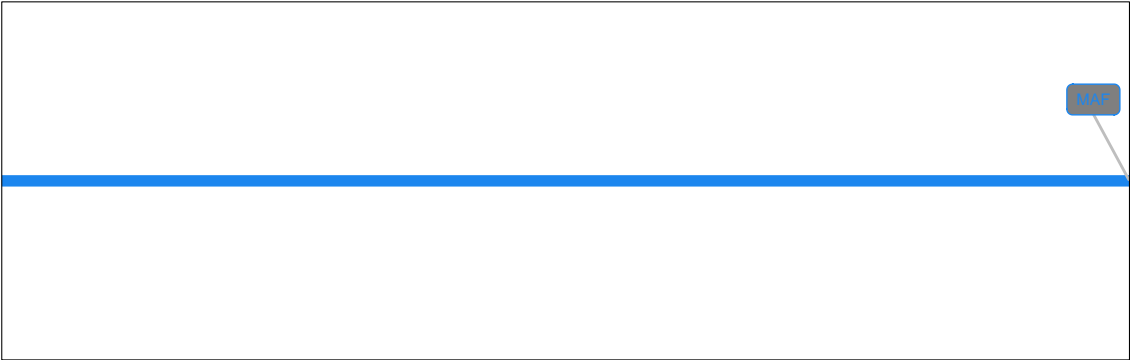
GeneTrack



SST

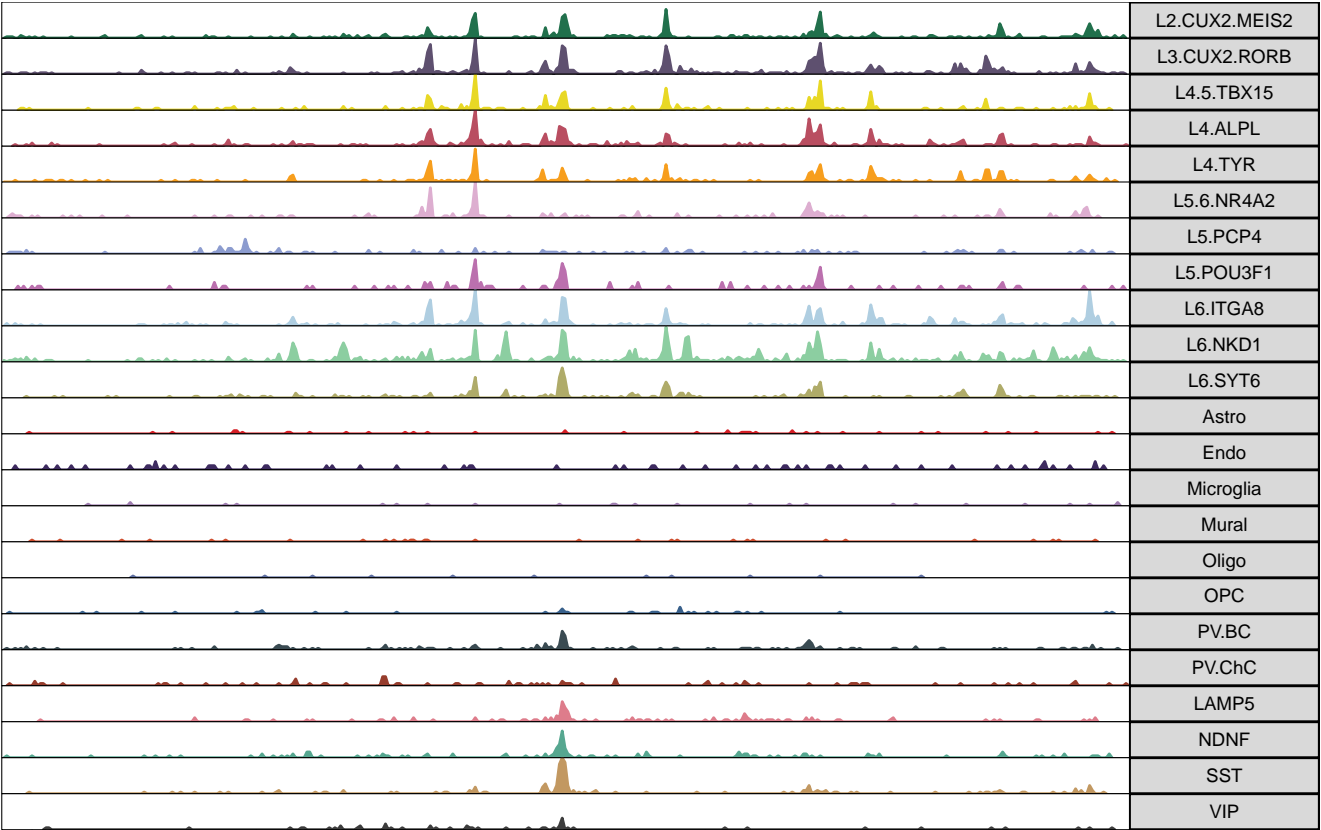


Peaks



Genes

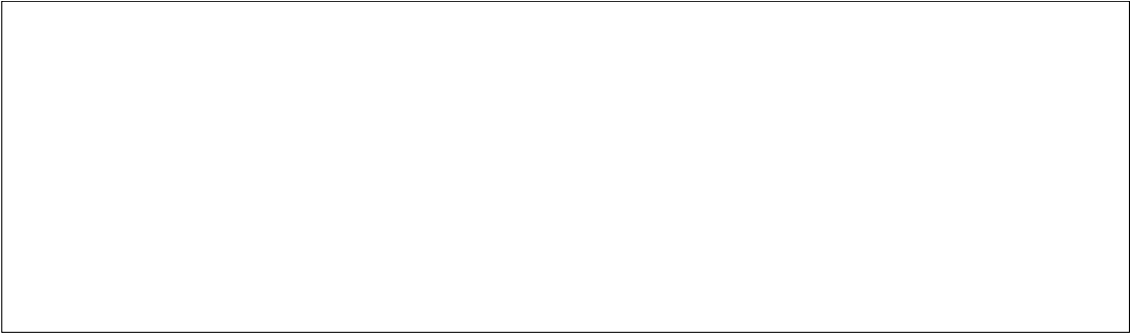
Coverage  
(Norm. ATAC Signal Range (0-0.7) by ReadsInTSS)



SST



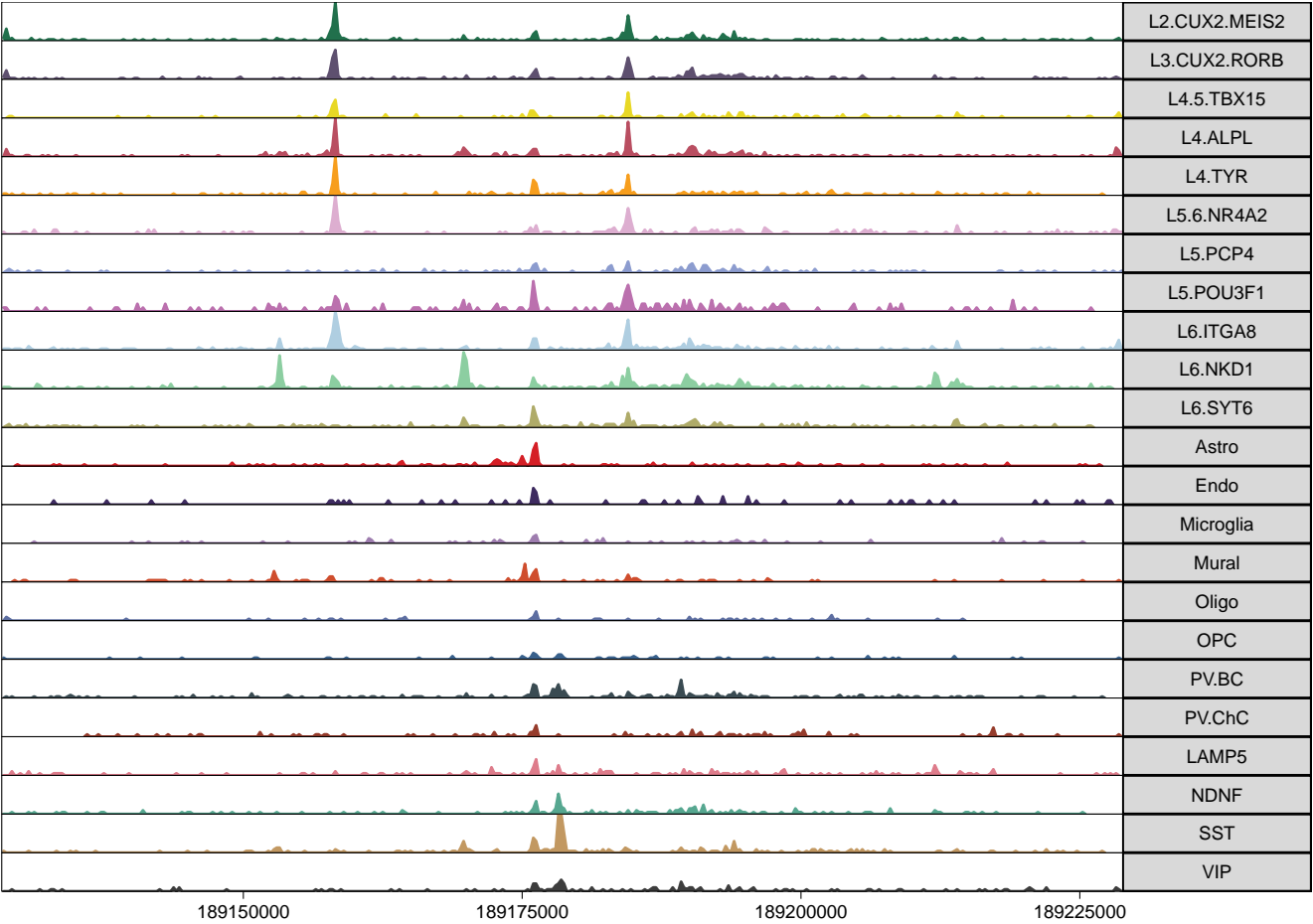
Peaks



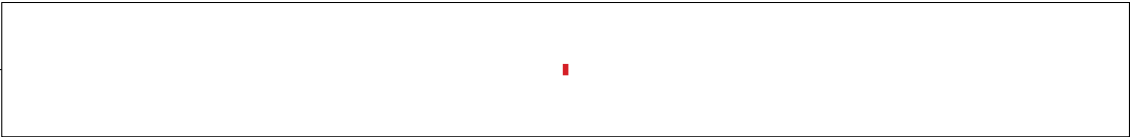
GeneTrack

chr1:189128336–189228837 Marker Peak Tracks for: SST

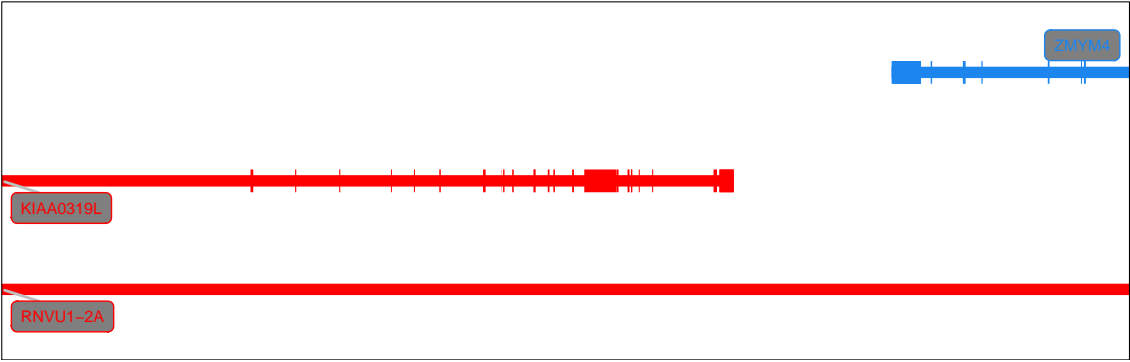
Coverage  
(Norm. ATAC Signal Range (0–0.75) by ReadsInTSS)



SST

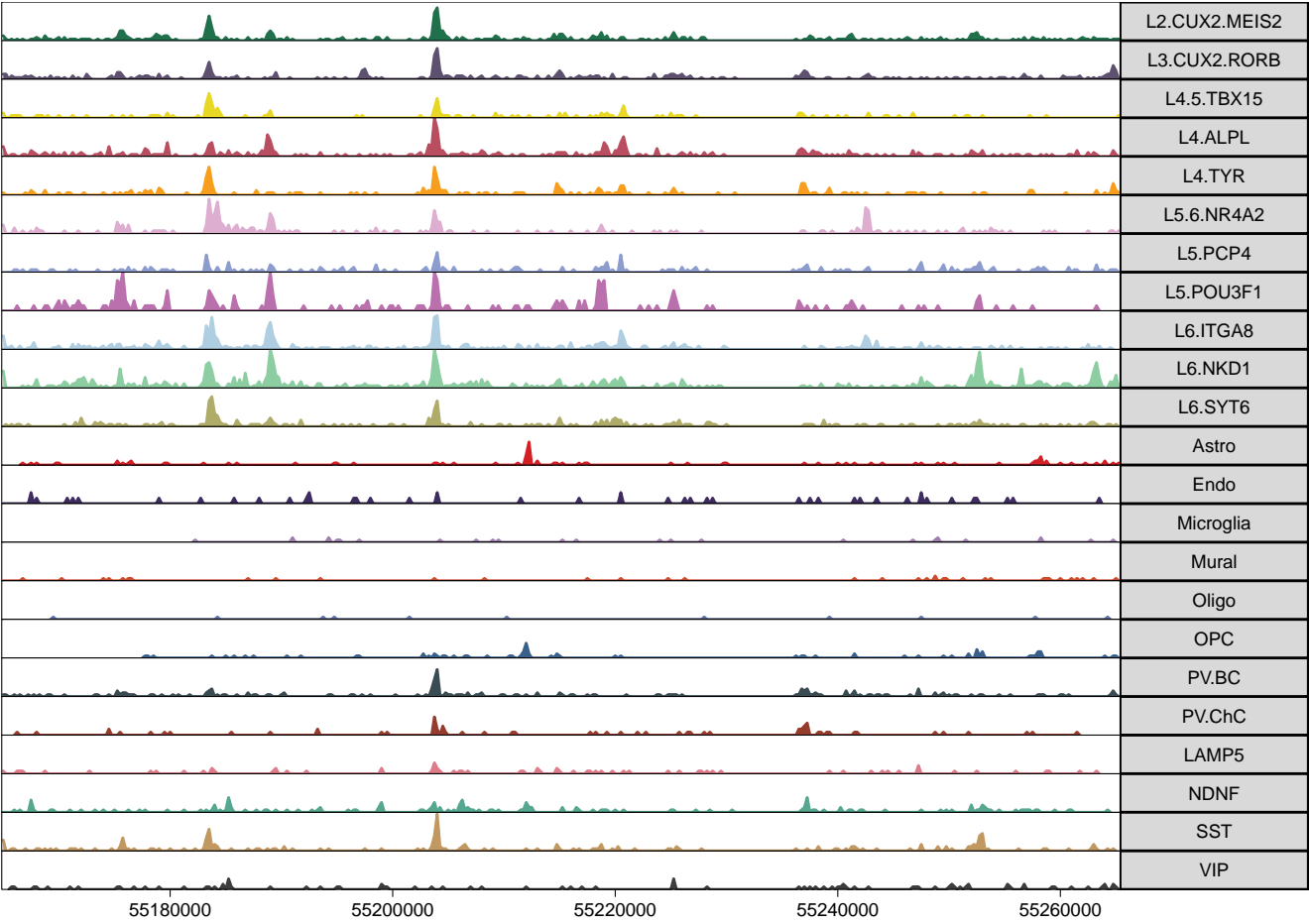


Peaks

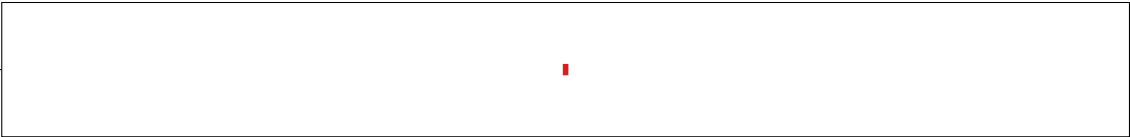


Genes

Coverage  
(Norm. ATAC Signal Range (0–0.56) by ReadsInTSS)



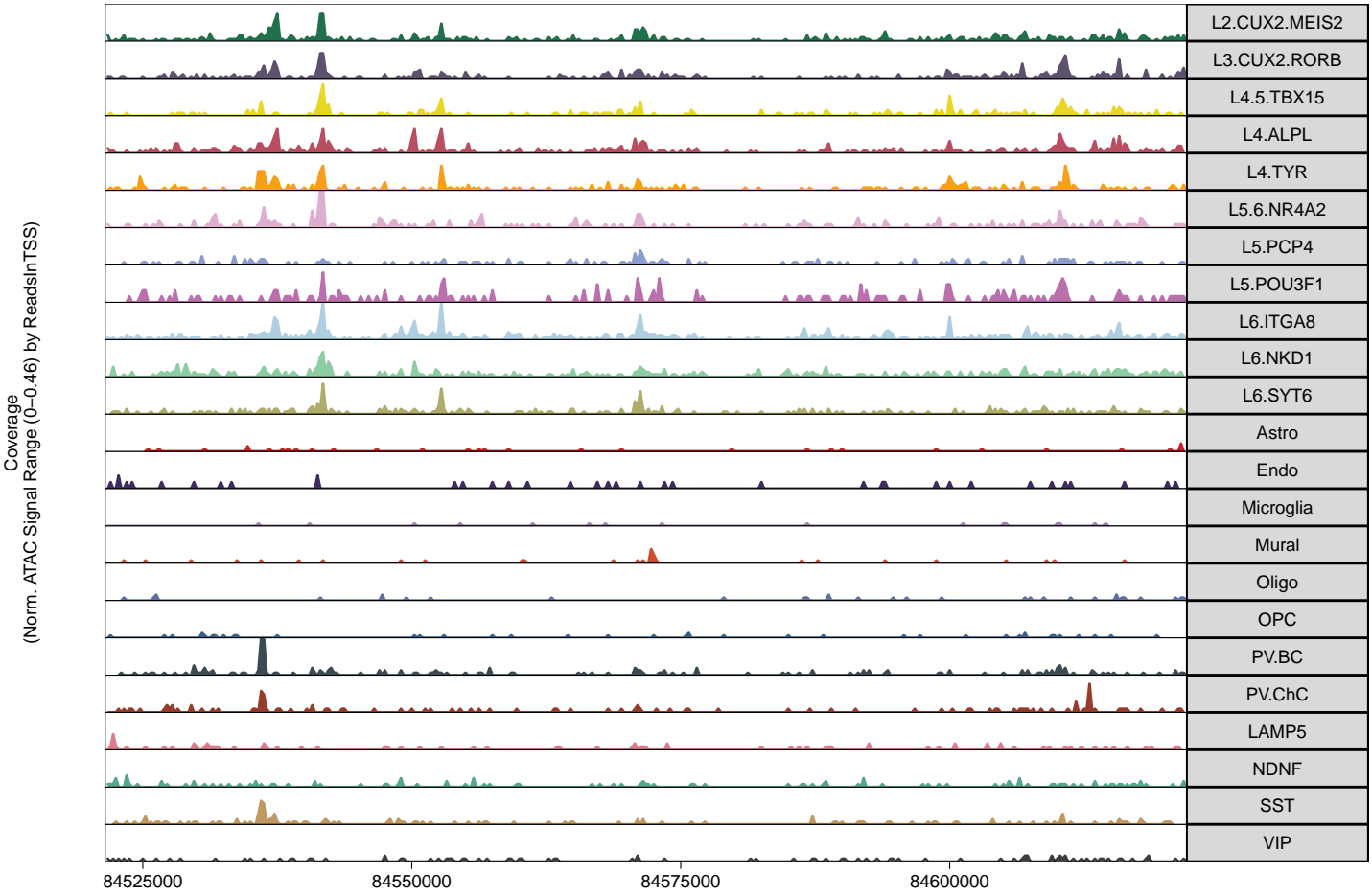
SST



Peaks

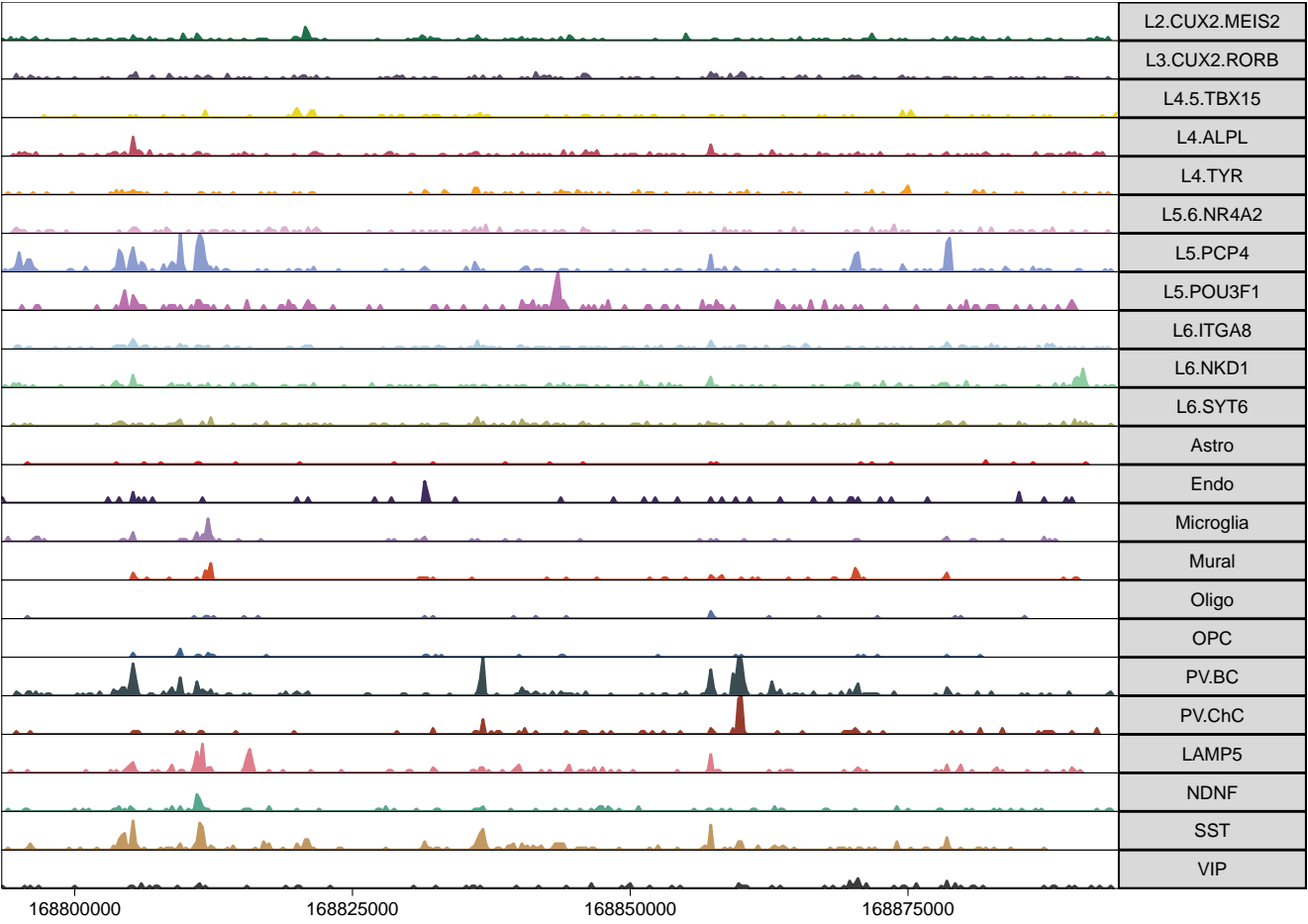


Genes

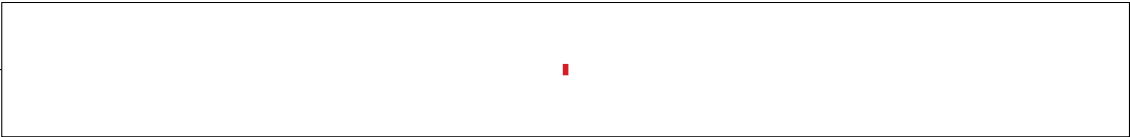




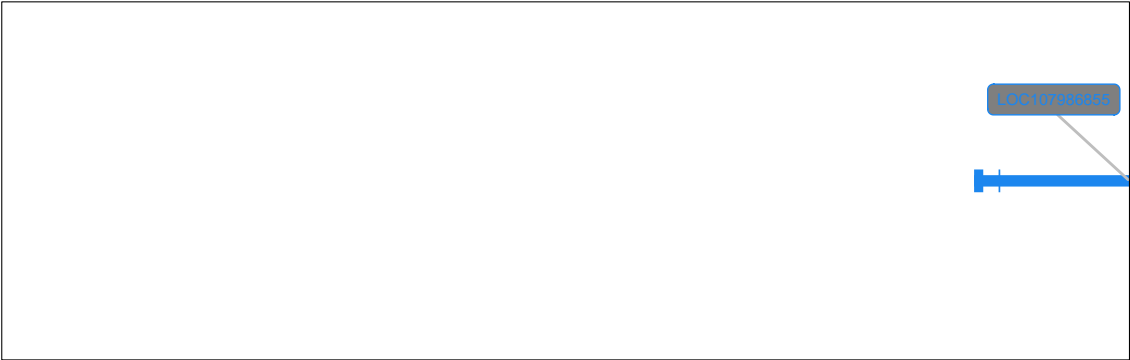
Coverage  
(Norm. ATAC Signal Range (0–0.57) by ReadsInTSS)



SST

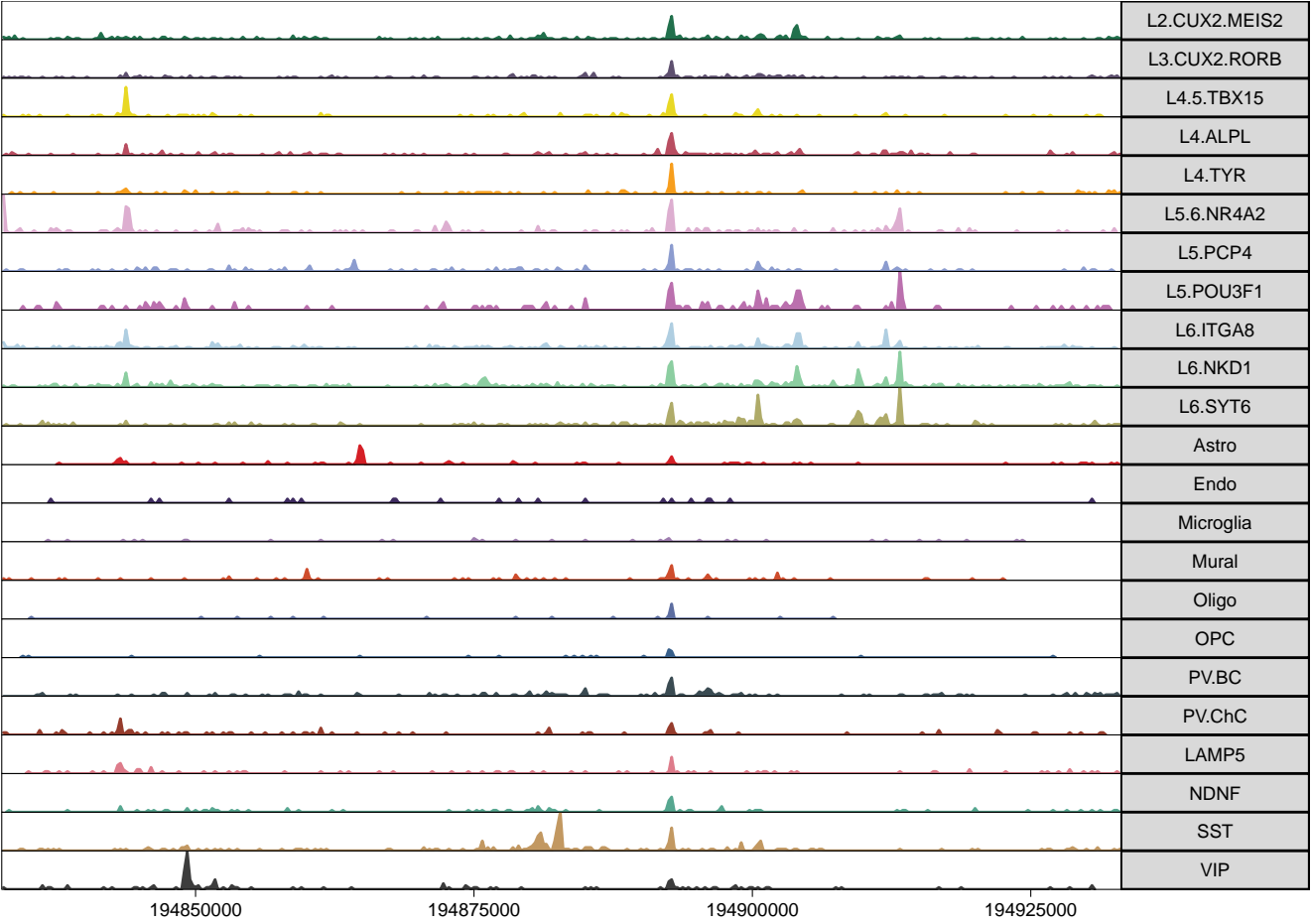


Peaks

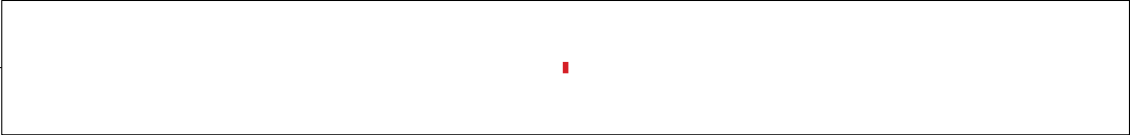


Genes

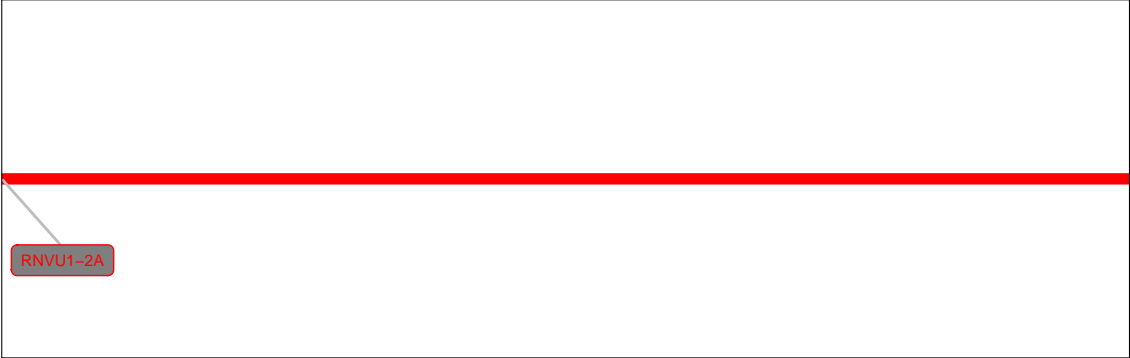
Coverage  
(Norm. ATAC Signal Range (0–0.74) by ReadsInTSS)



SST

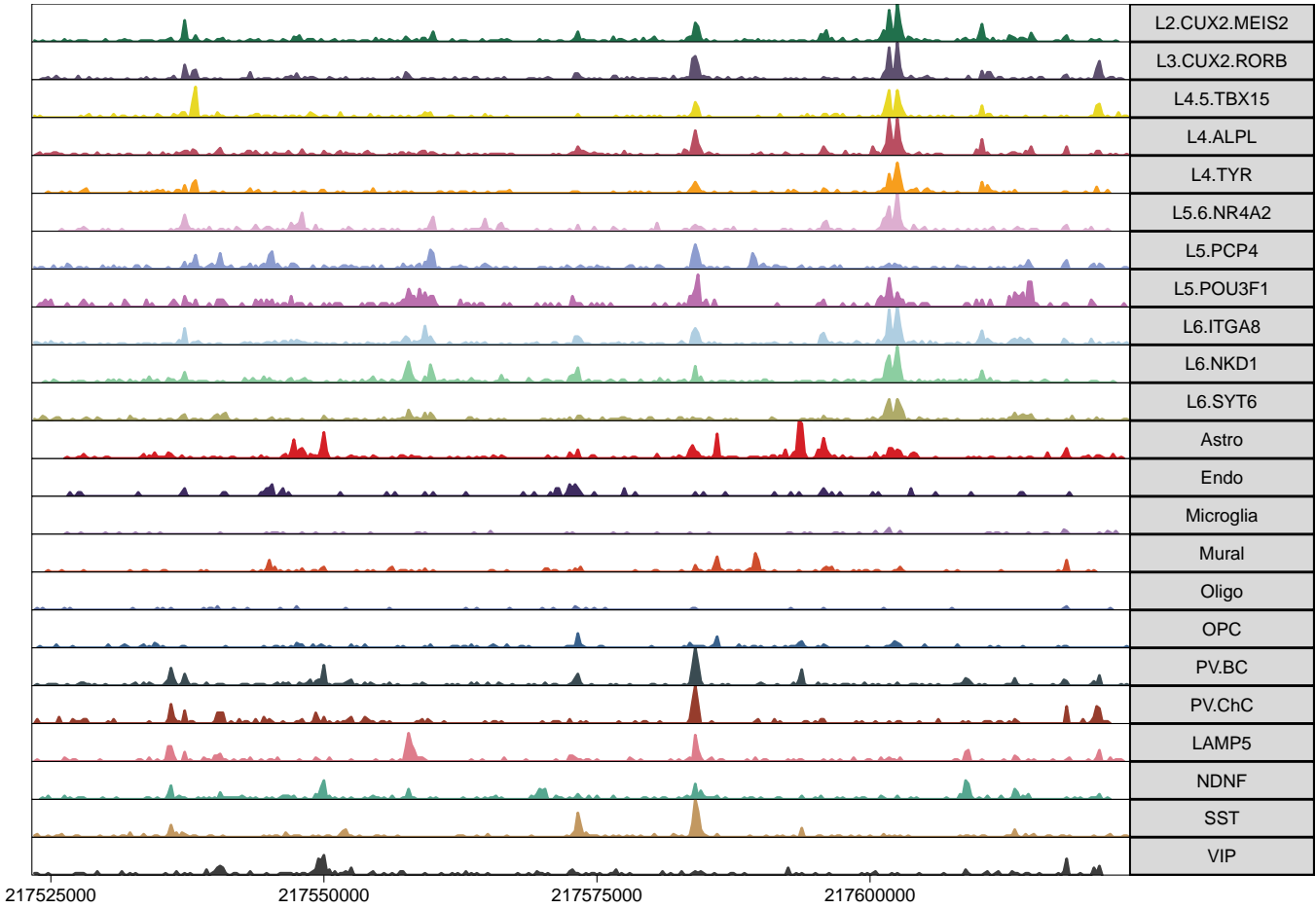


Peaks

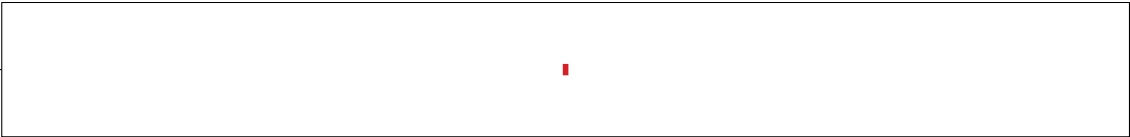


Genes

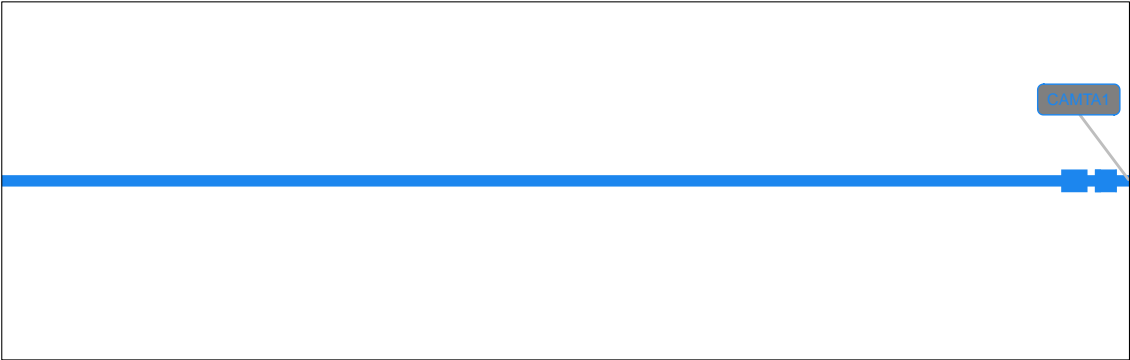
Coverage  
(Norm. ATAC Signal Range (0–0.78) by ReadsInTSS)



SST

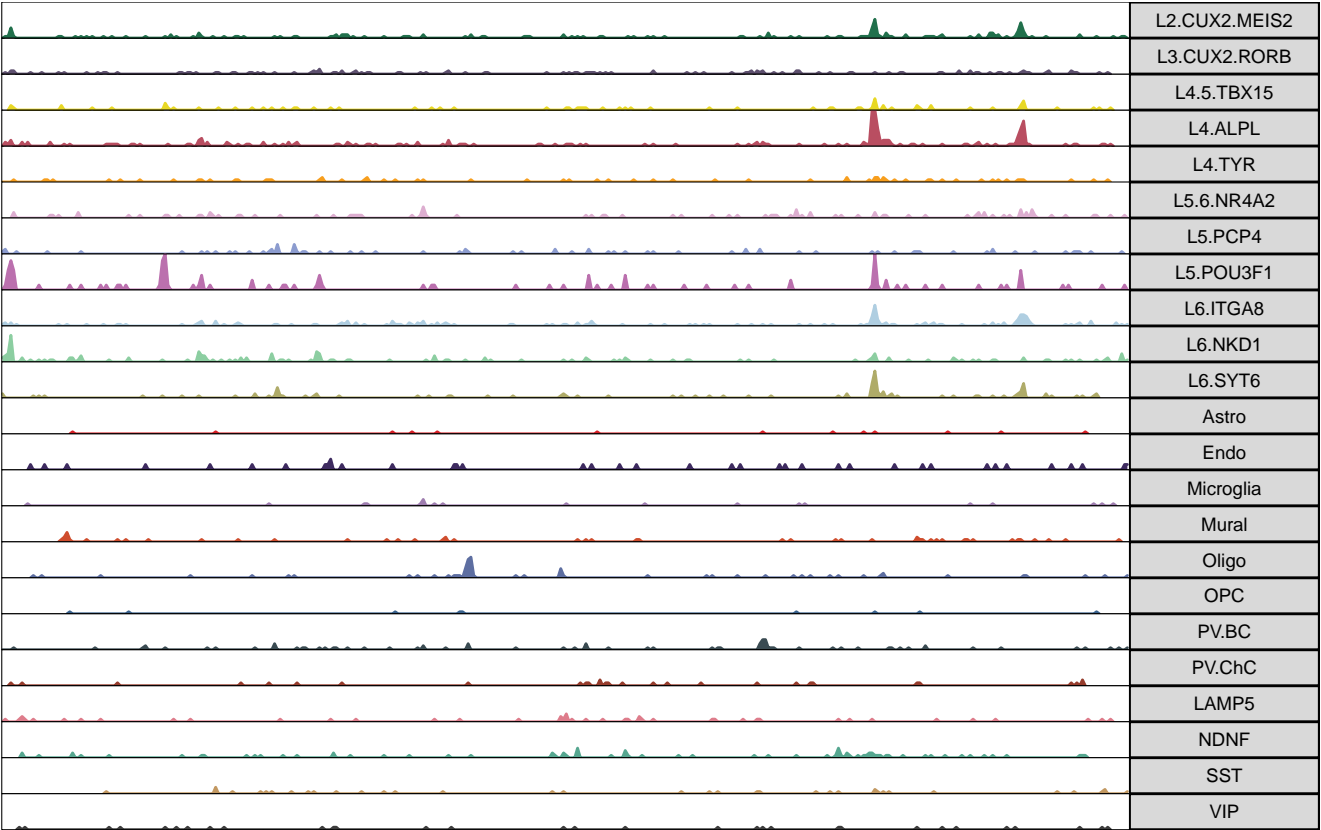


Peaks



Genes

Coverage  
(Norm. ATAC Signal Range (0–0.54) by ReadsInTSS)



SST

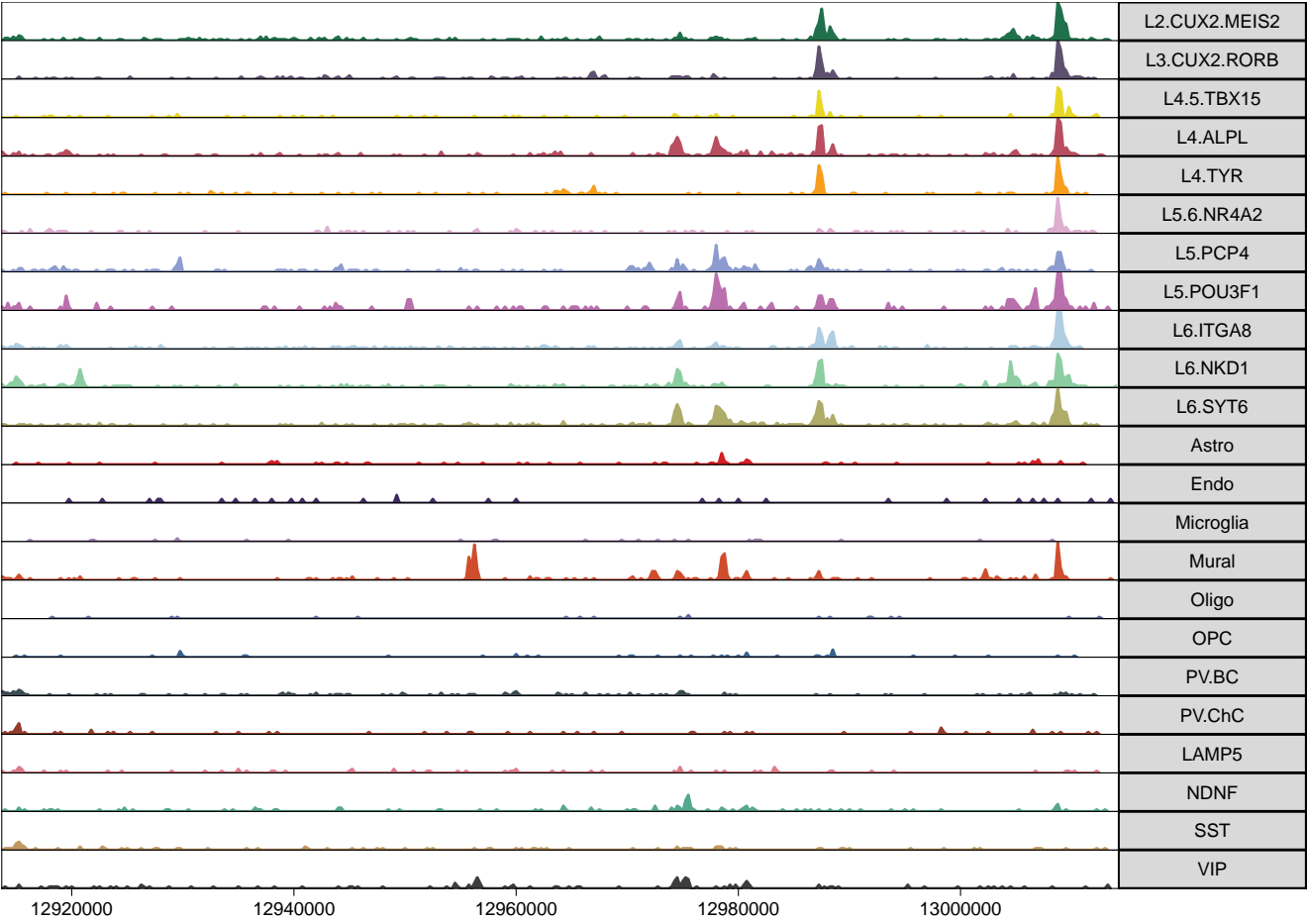


Peaks

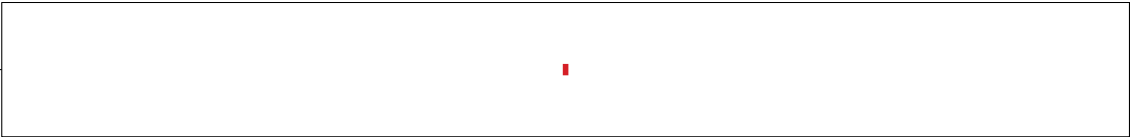
GeneTrack

strand ● ·

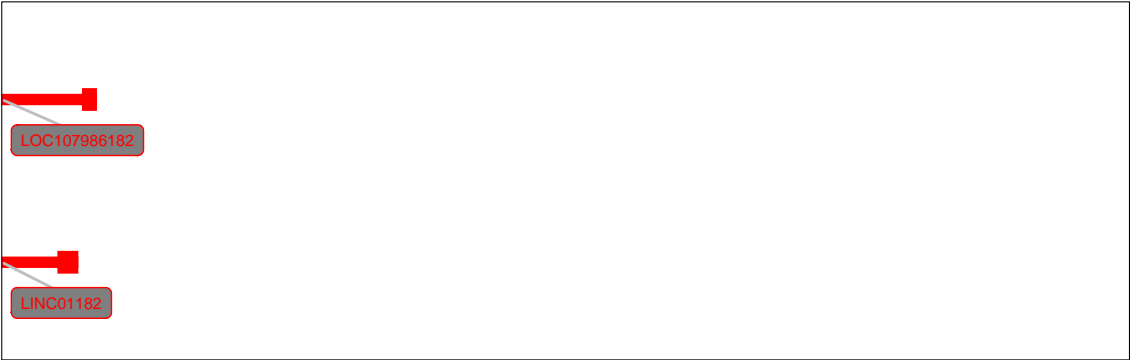
Coverage  
(Norm. ATAC Signal Range (0-0.77) by ReadsInTSS)



SST

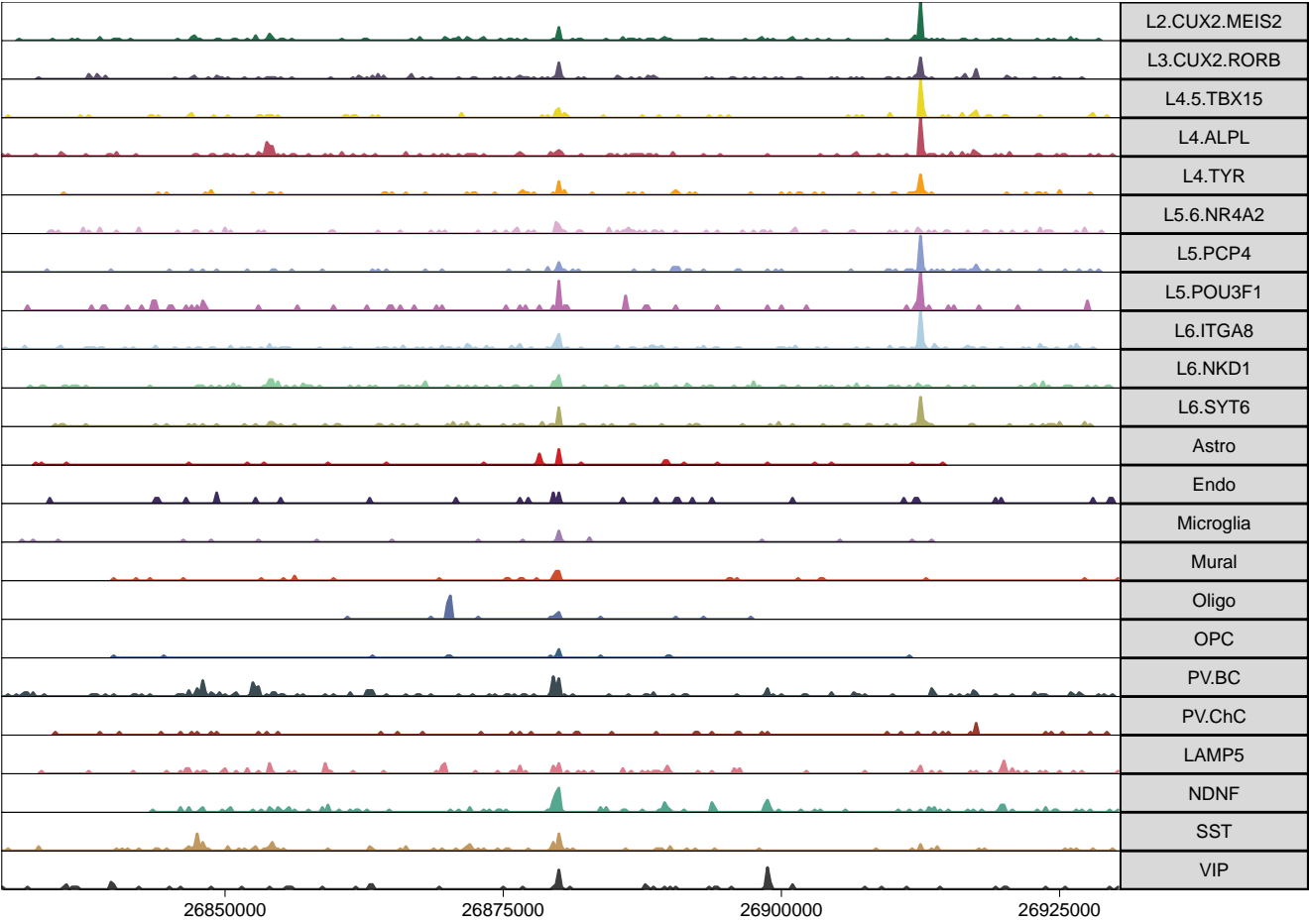


Peaks

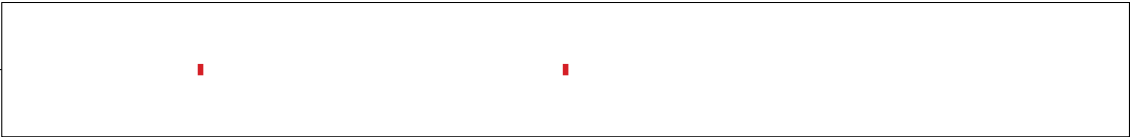


Genes

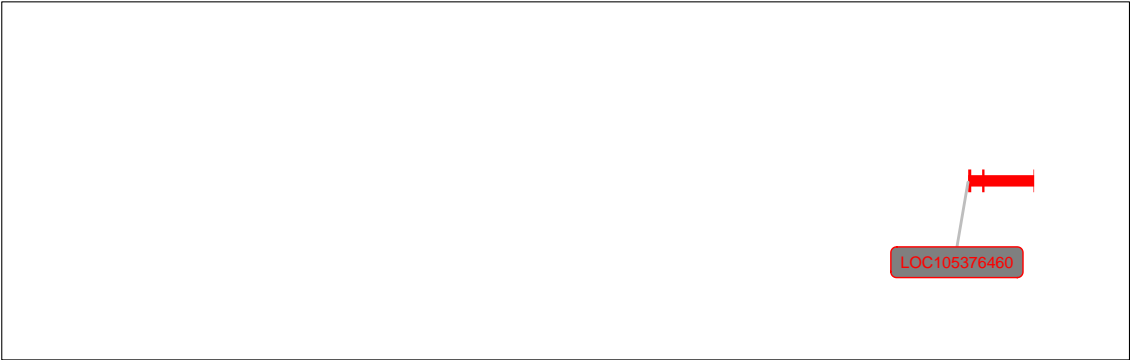
Coverage  
(Norm. ATAC Signal Range (0–0.57) by ReadsInTSS)



SST

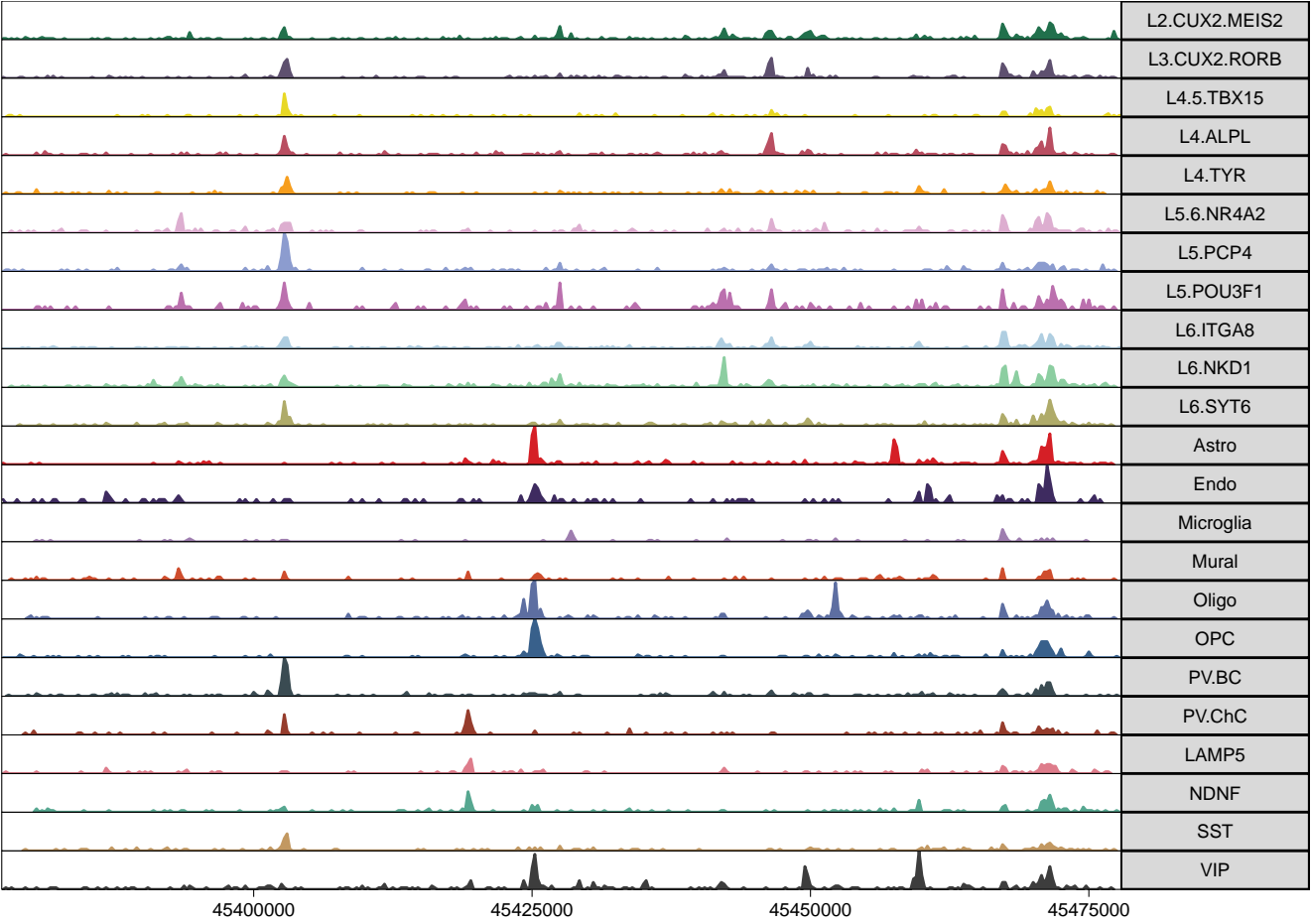


Peaks

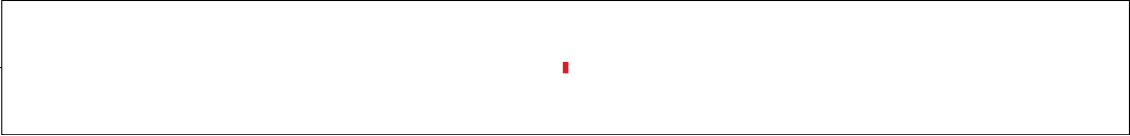


Genes

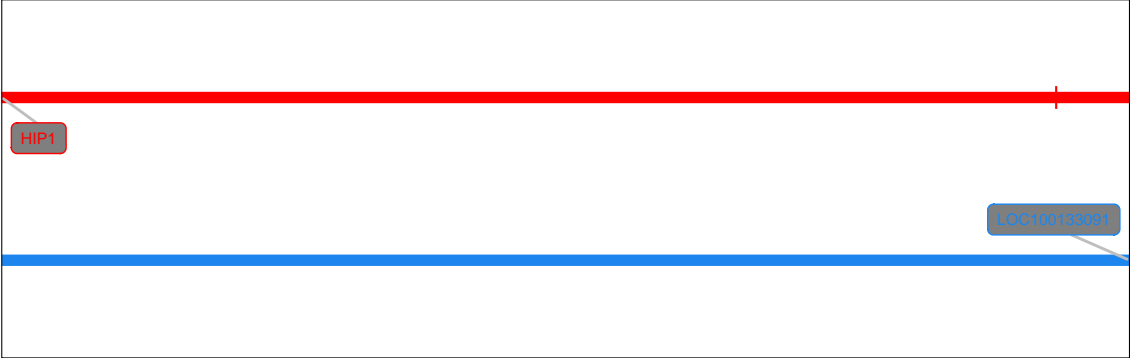
Coverage  
(Norm. ATAC Signal Range (0–0.83) by ReadsInTSS)



SST

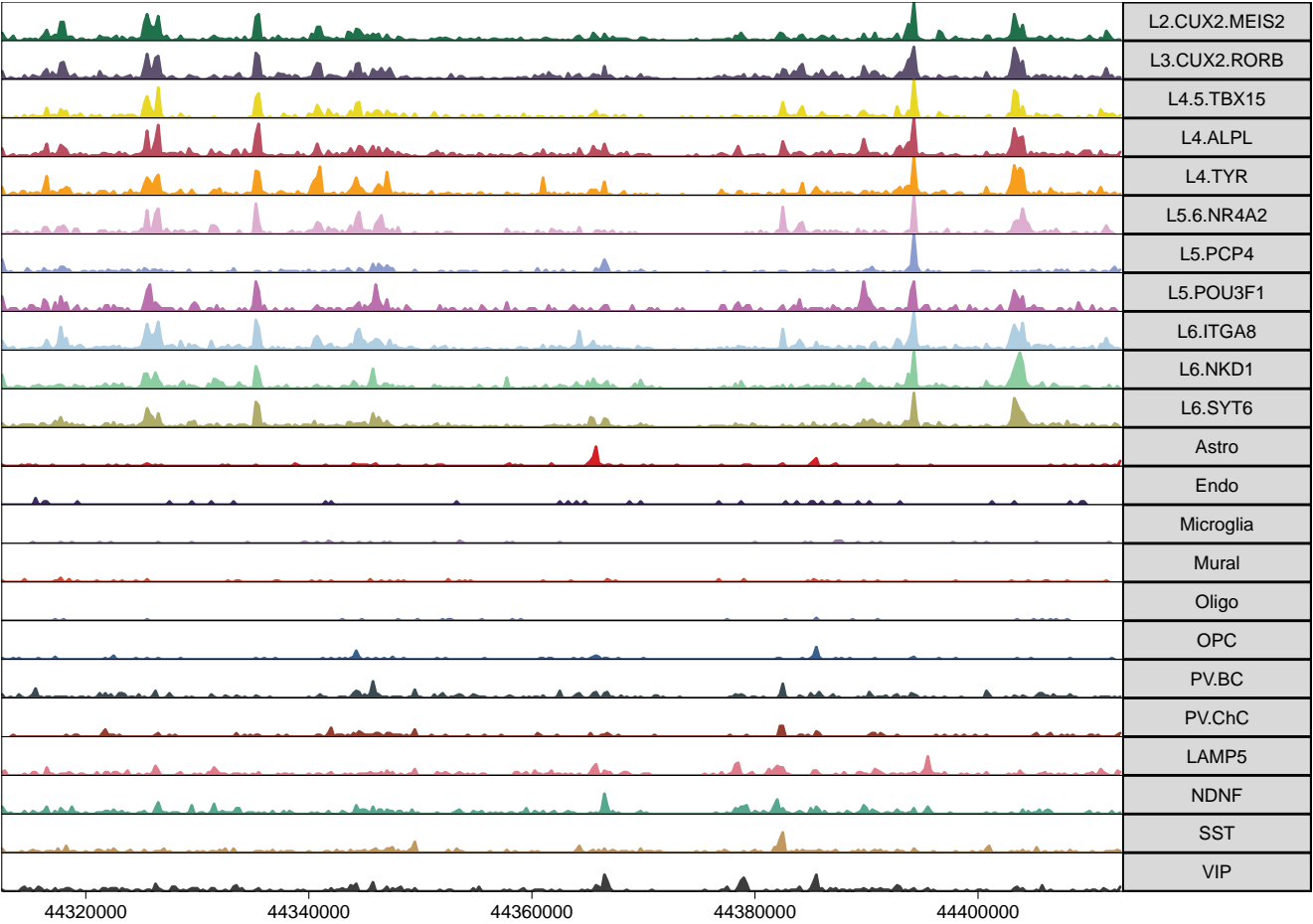


Peaks

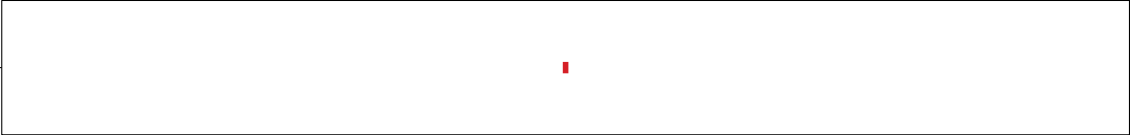


Genes

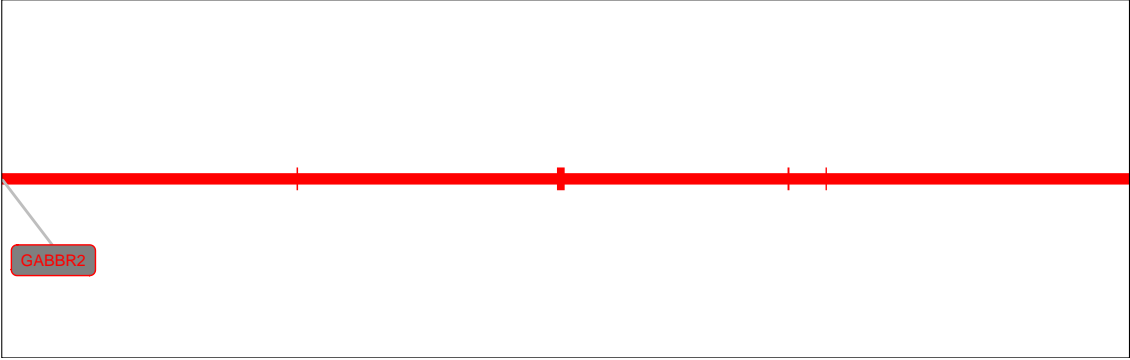
Coverage  
(Norm. ATAC Signal Range (0–0.95) by ReadsInTSS)



SST



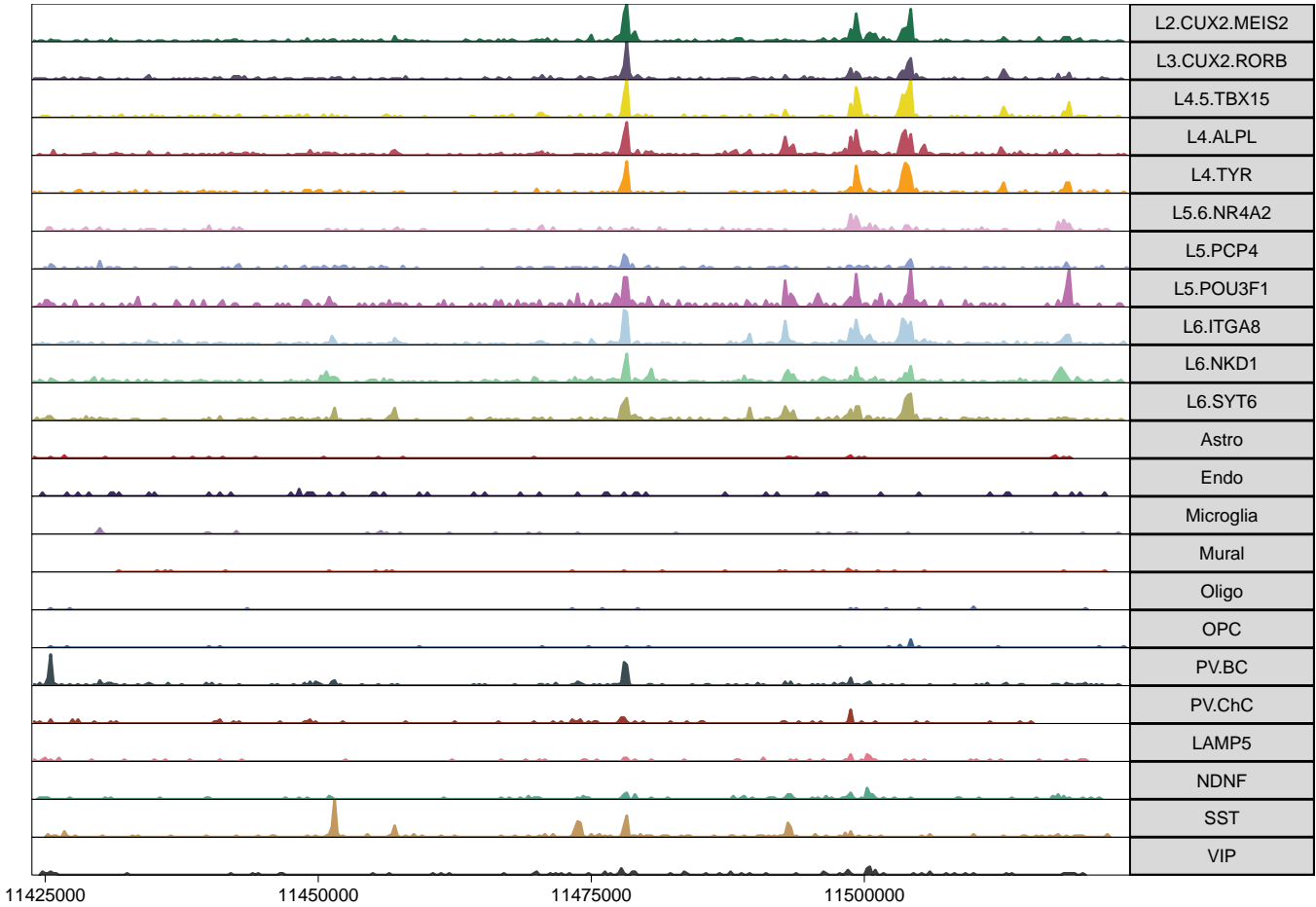
Peaks



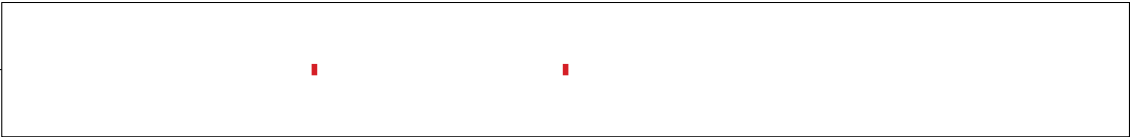
Genes



Coverage  
(Norm. ATAC Signal Range (0–0.85) by ReadsInTSS)



SST



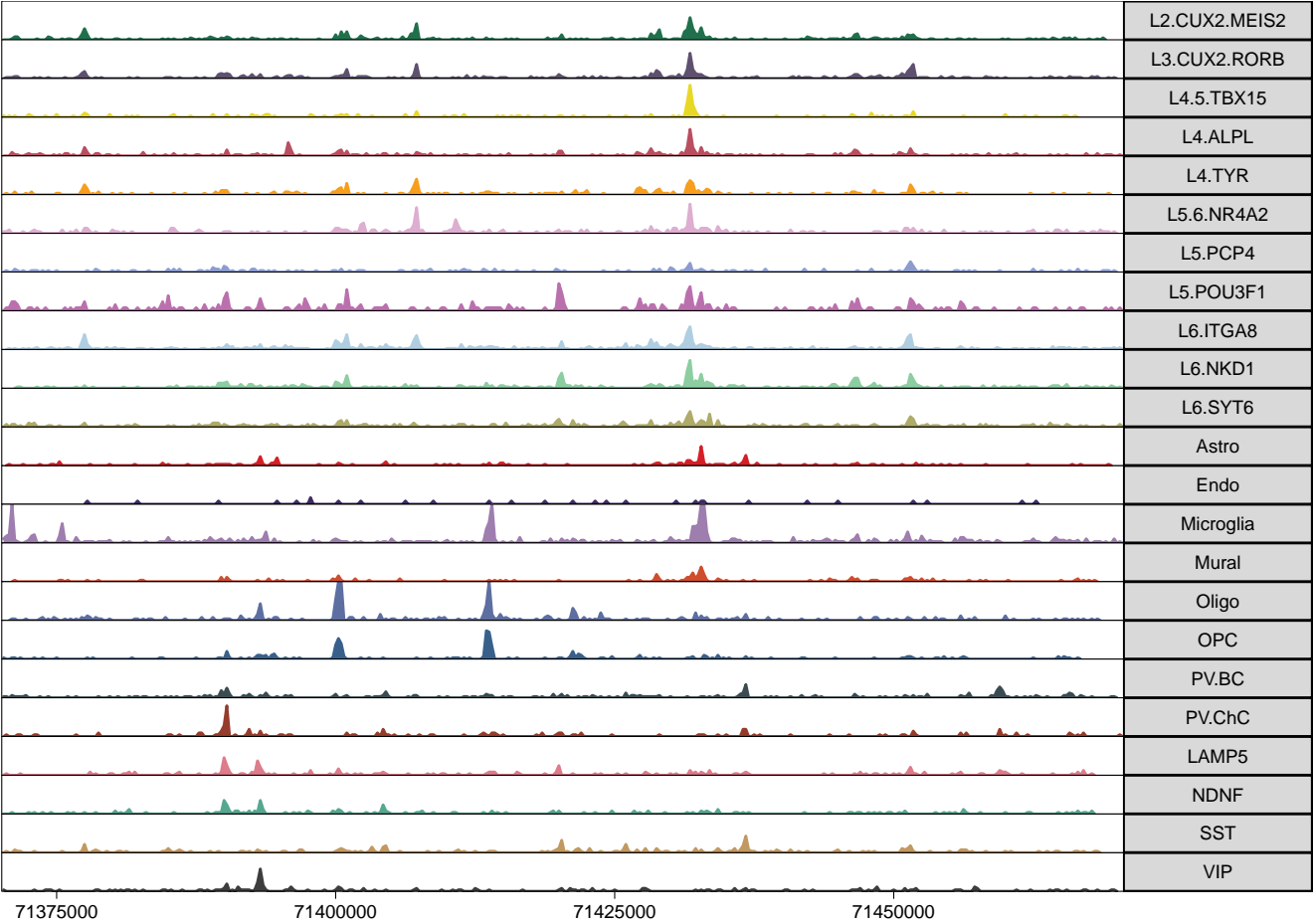
Peaks



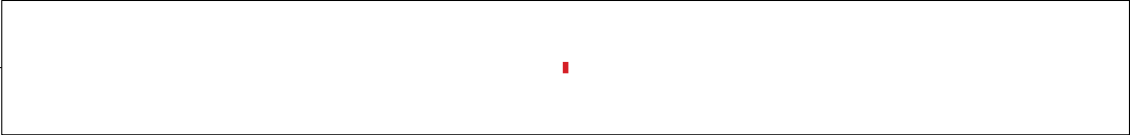
Genes

chr14:71370081–71470582 Marker Peak Tracks for: SST

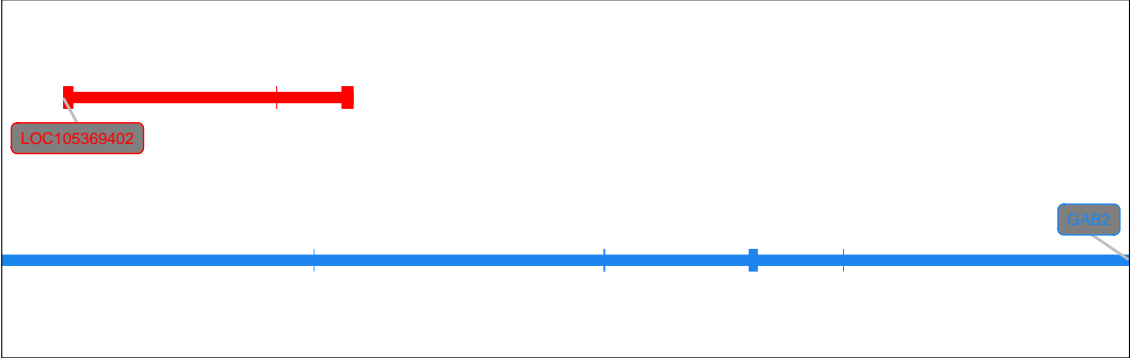
Coverage  
(Norm. ATAC Signal Range (0–0.94) by ReadsInTSS)



SST



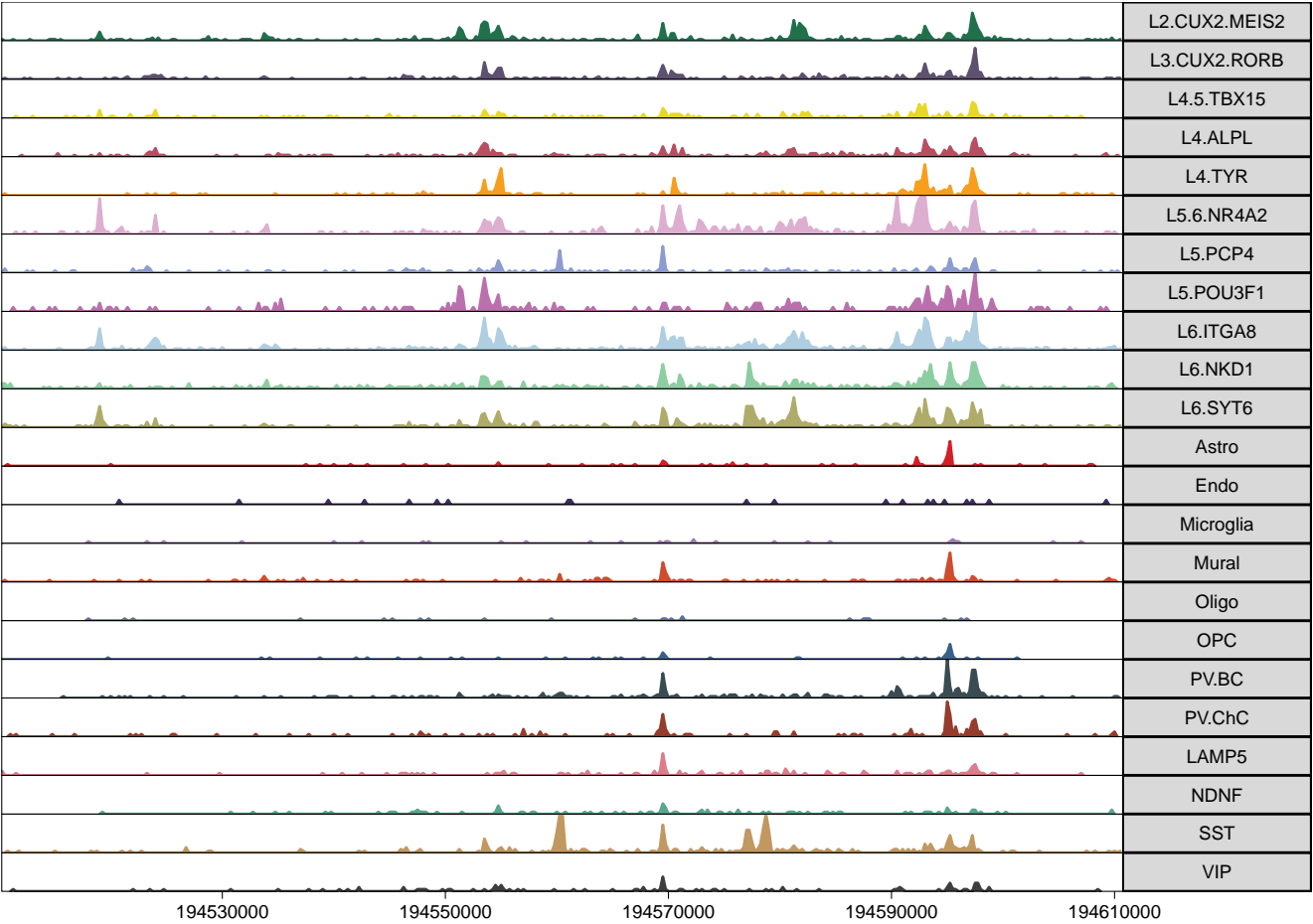
Peaks



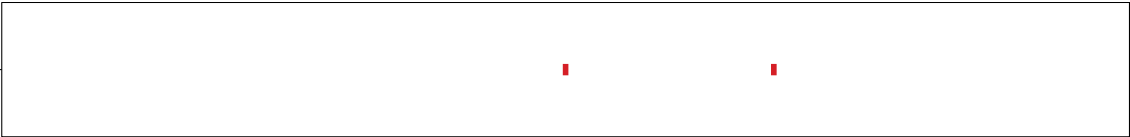
Genes

chr1:194510221–194610722 Marker Peak Tracks for: SST

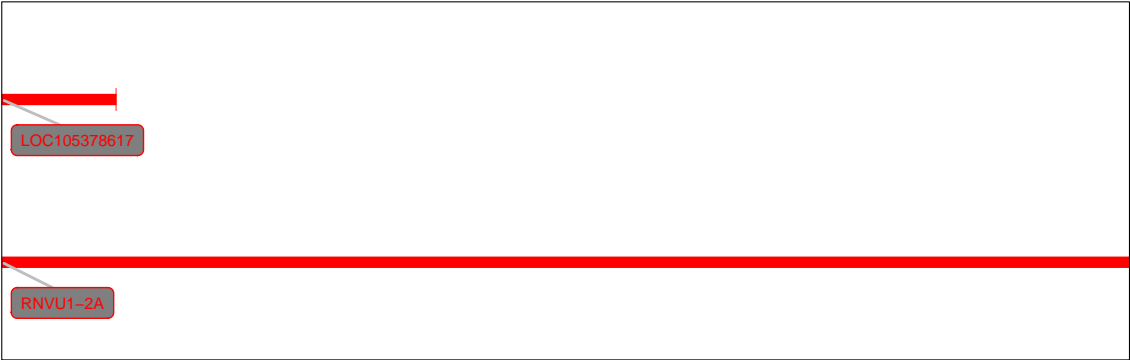
Coverage  
(Norm. ATAC Signal Range (0–0.68)) by ReadsInTSS



SST

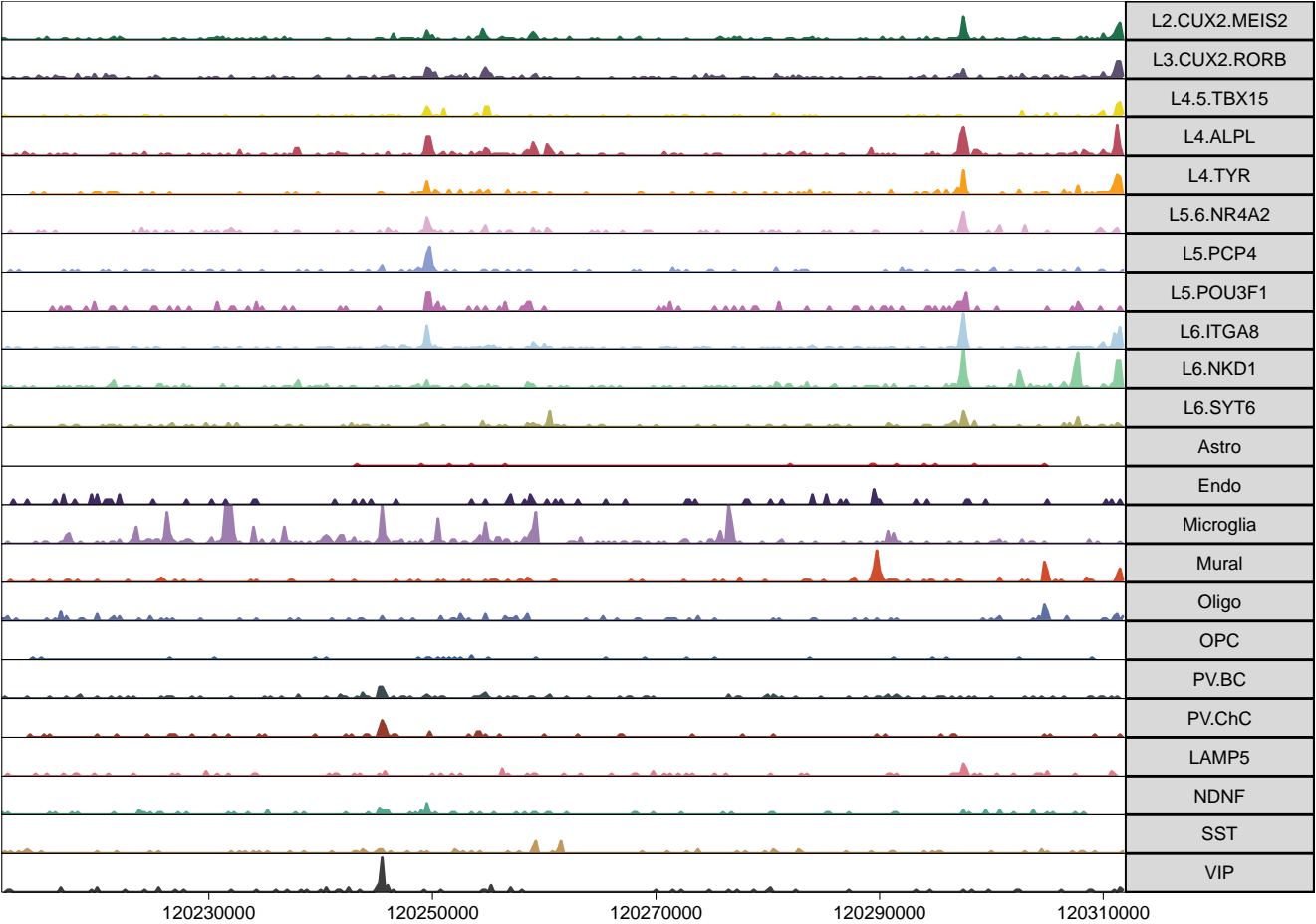


Peaks

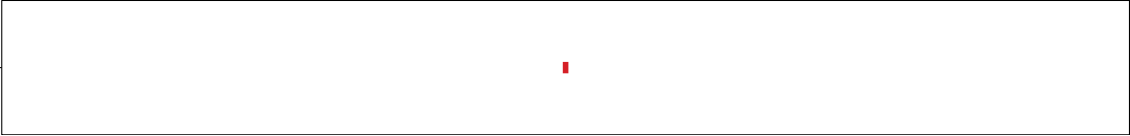


Genes

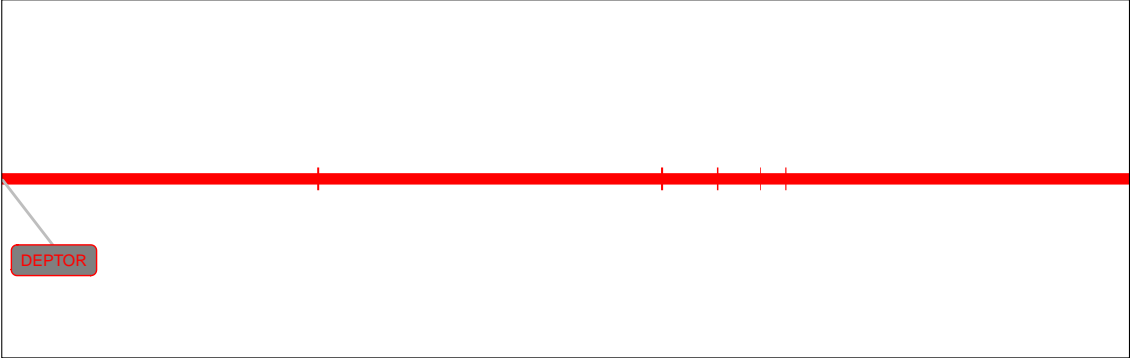
Coverage  
(Norm. ATAC Signal Range (0–0.61) by ReadsInTSS)



SST

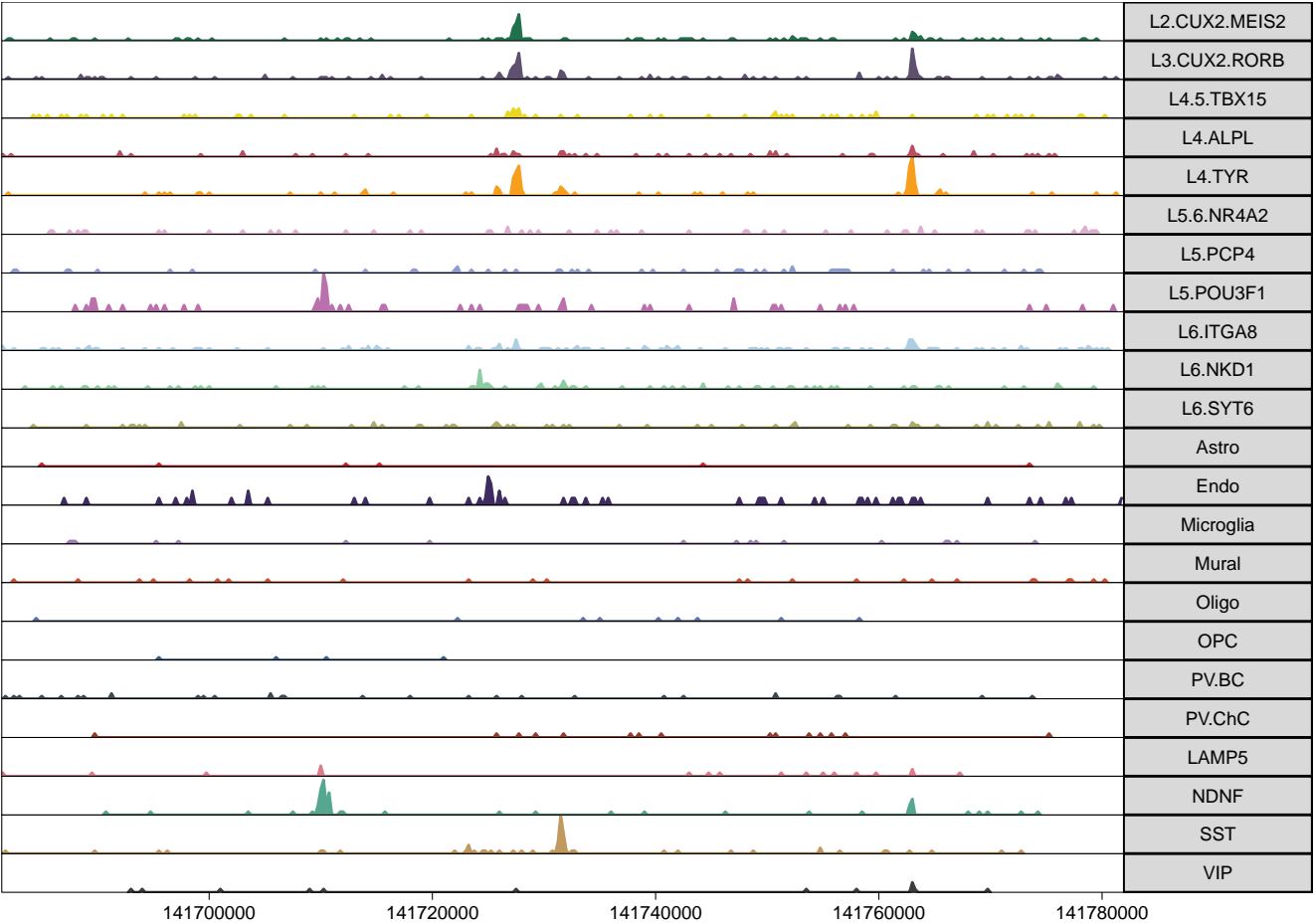


Peaks

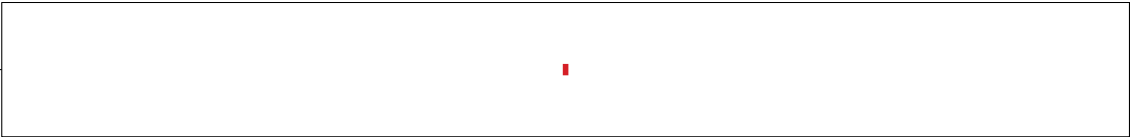


Genes

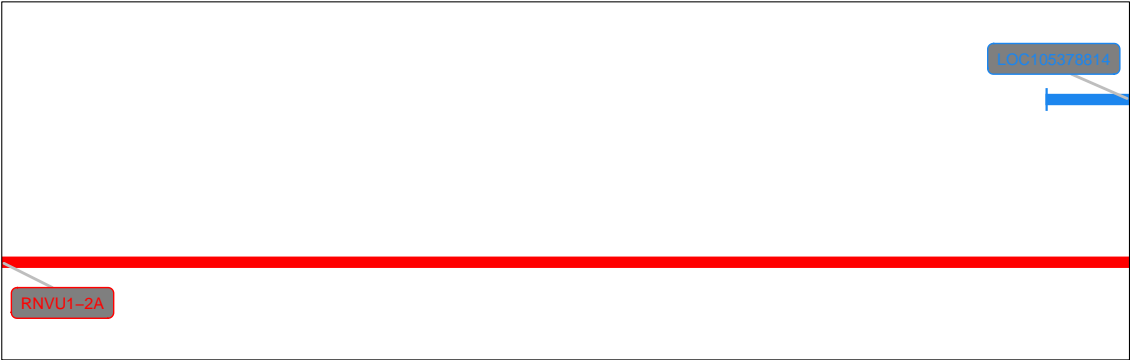
Coverage  
(Norm. ATAC Signal Range (0–0.43) by ReadsInTSS)



SST



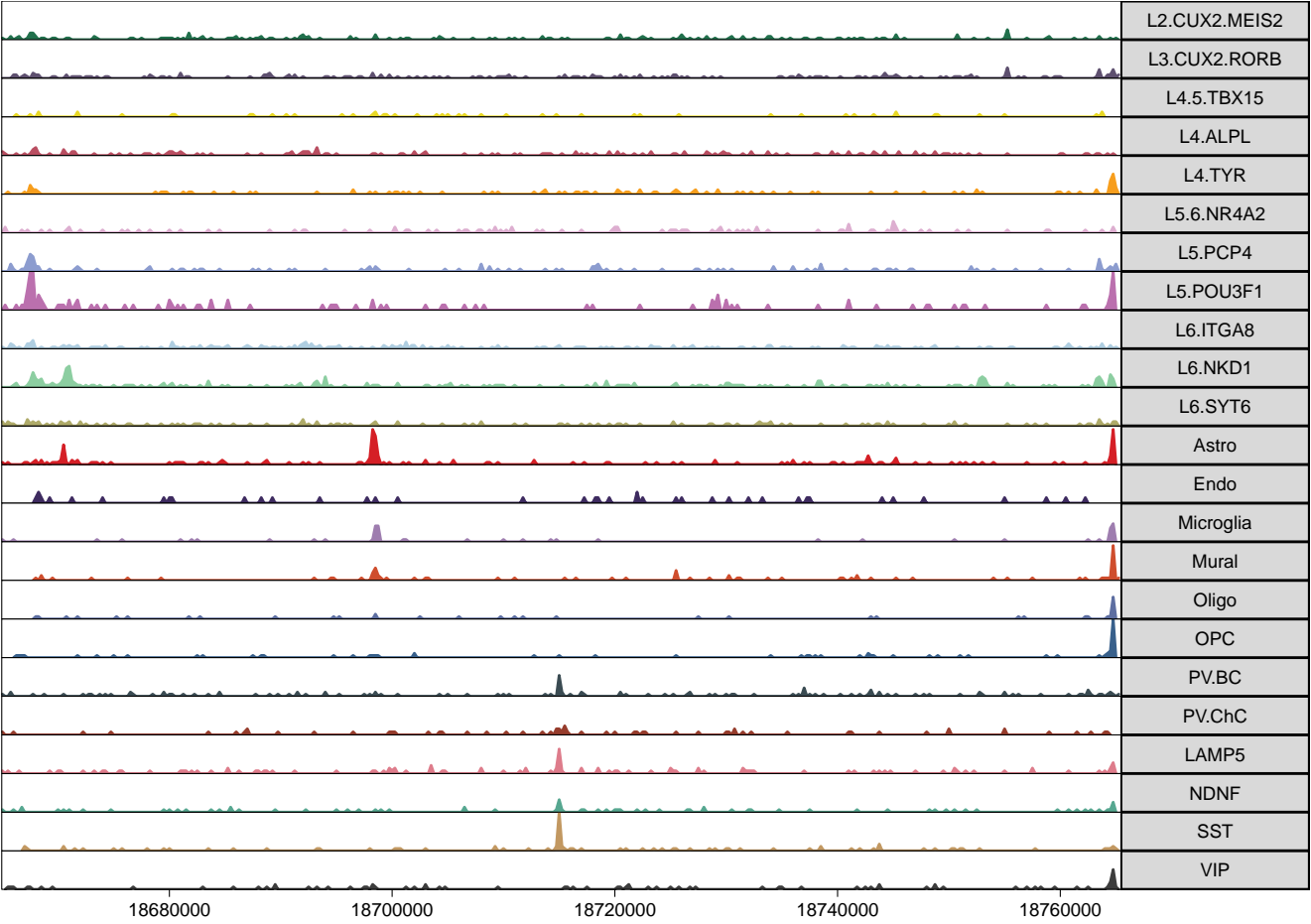
Peaks



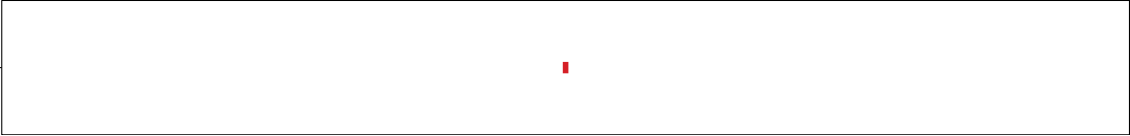
Genes

chr14:18664942–18765443 Marker Peak Tracks for: SST

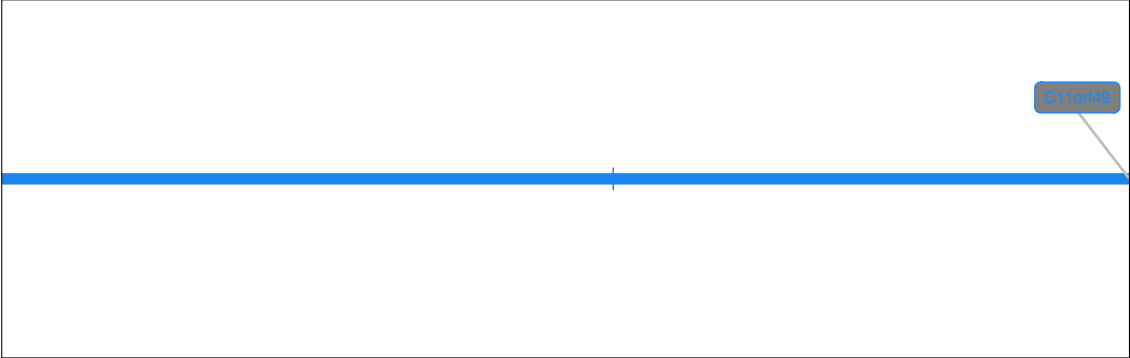
Coverage  
(Norm. ATAC Signal Range (0–0.56) by ReadsInTSS)



SST

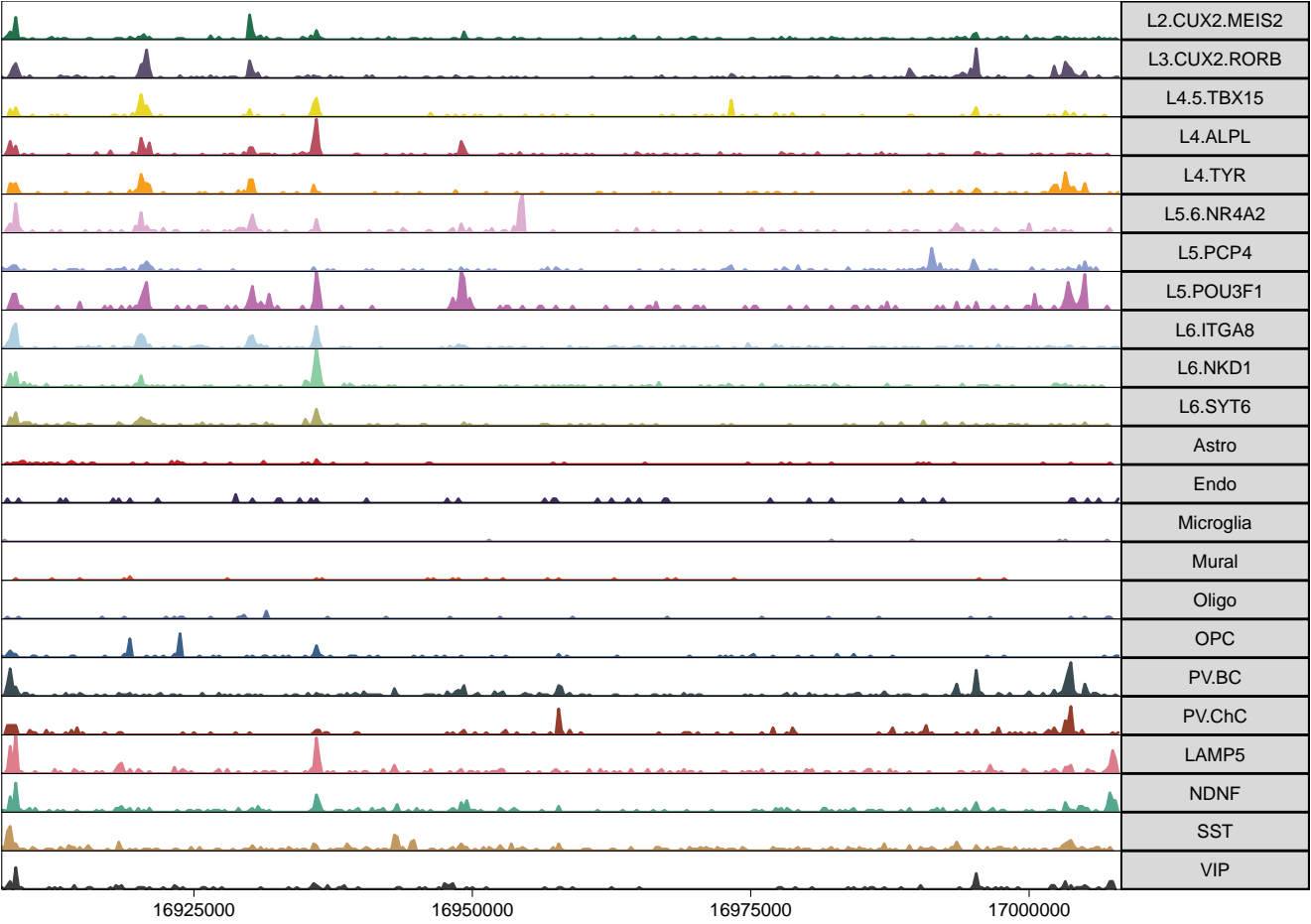


Peaks

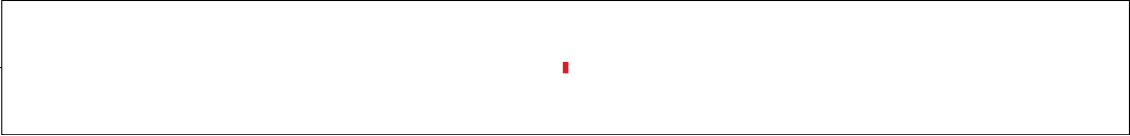


Genes

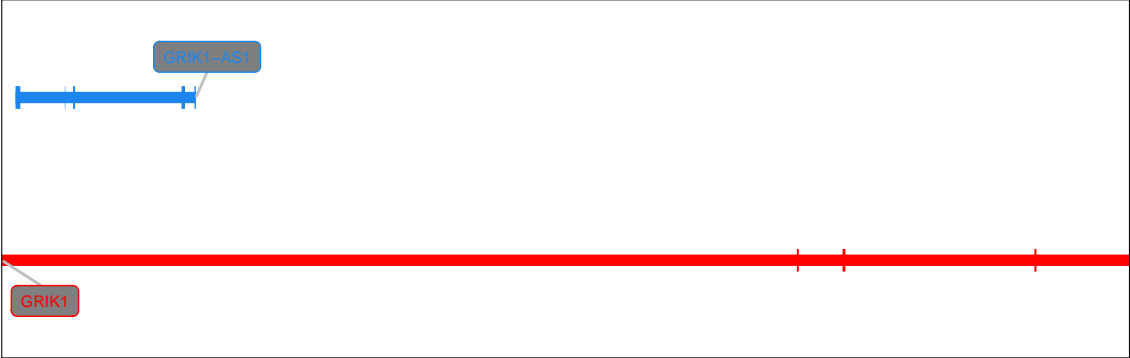
Coverage  
(Norm. ATAC Signal Range (0–0.72) by ReadsInTSS)



SST

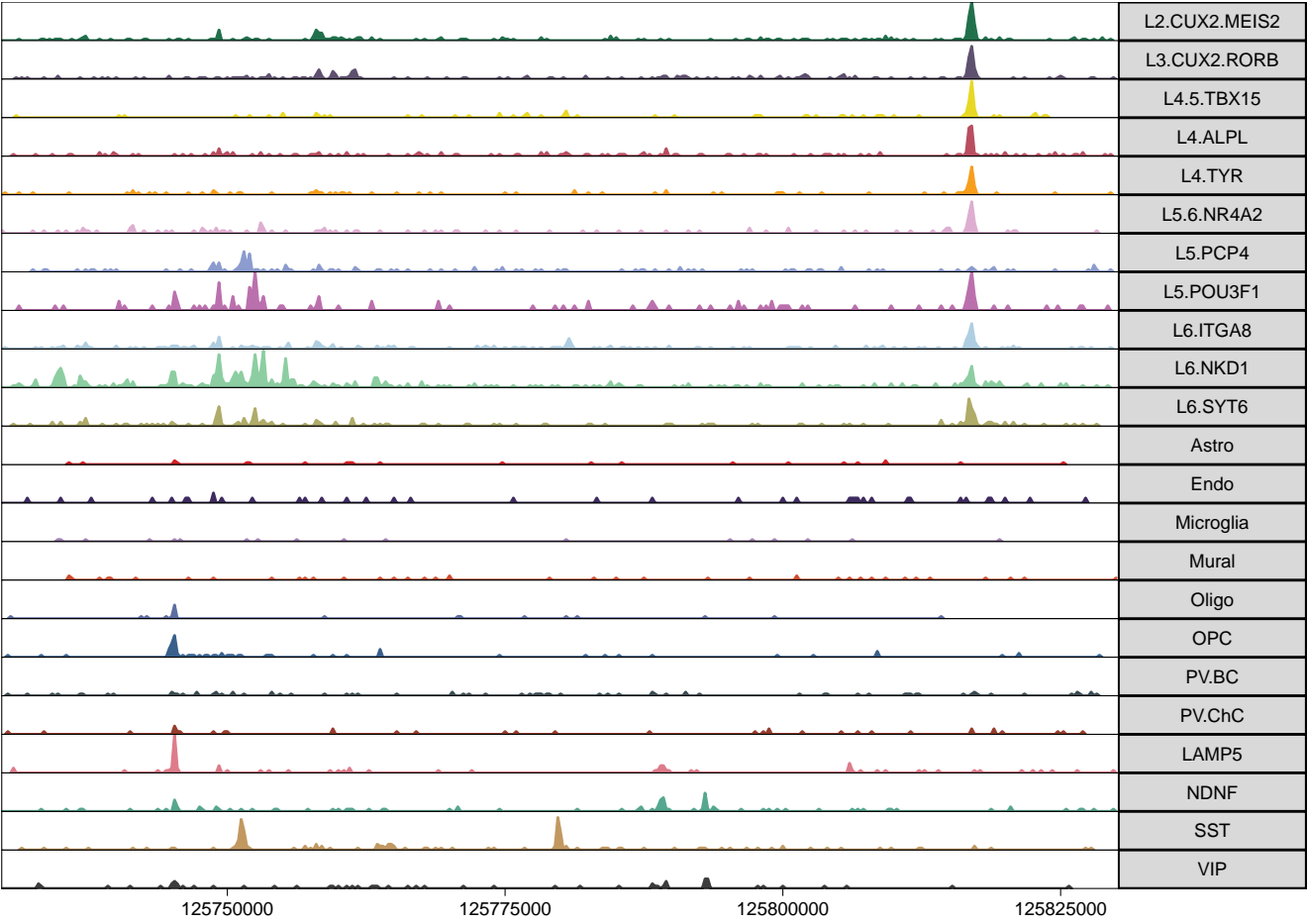


Peaks

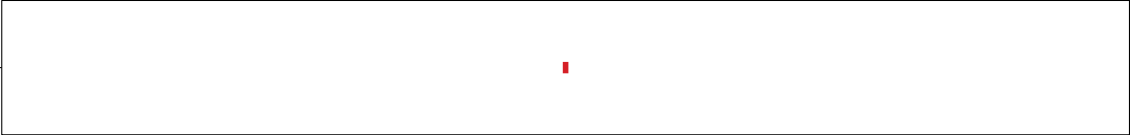


Genes

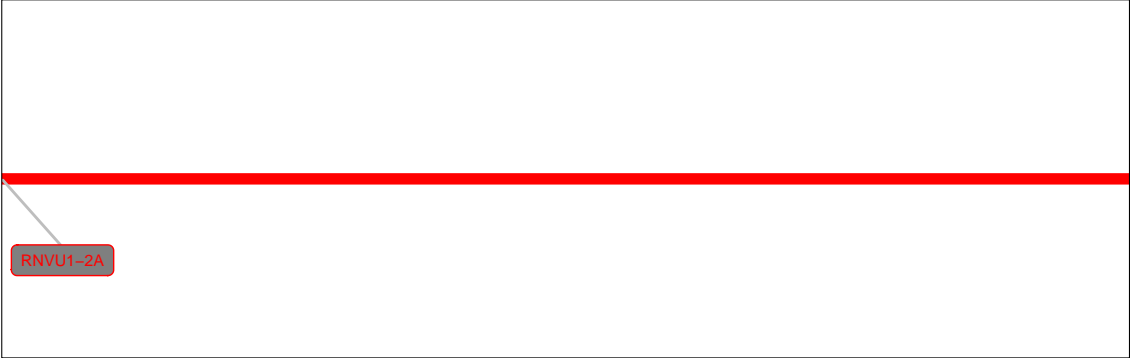
Coverage  
(Norm. ATAC Signal Range (0–0.61) by ReadsInTSS)



SST



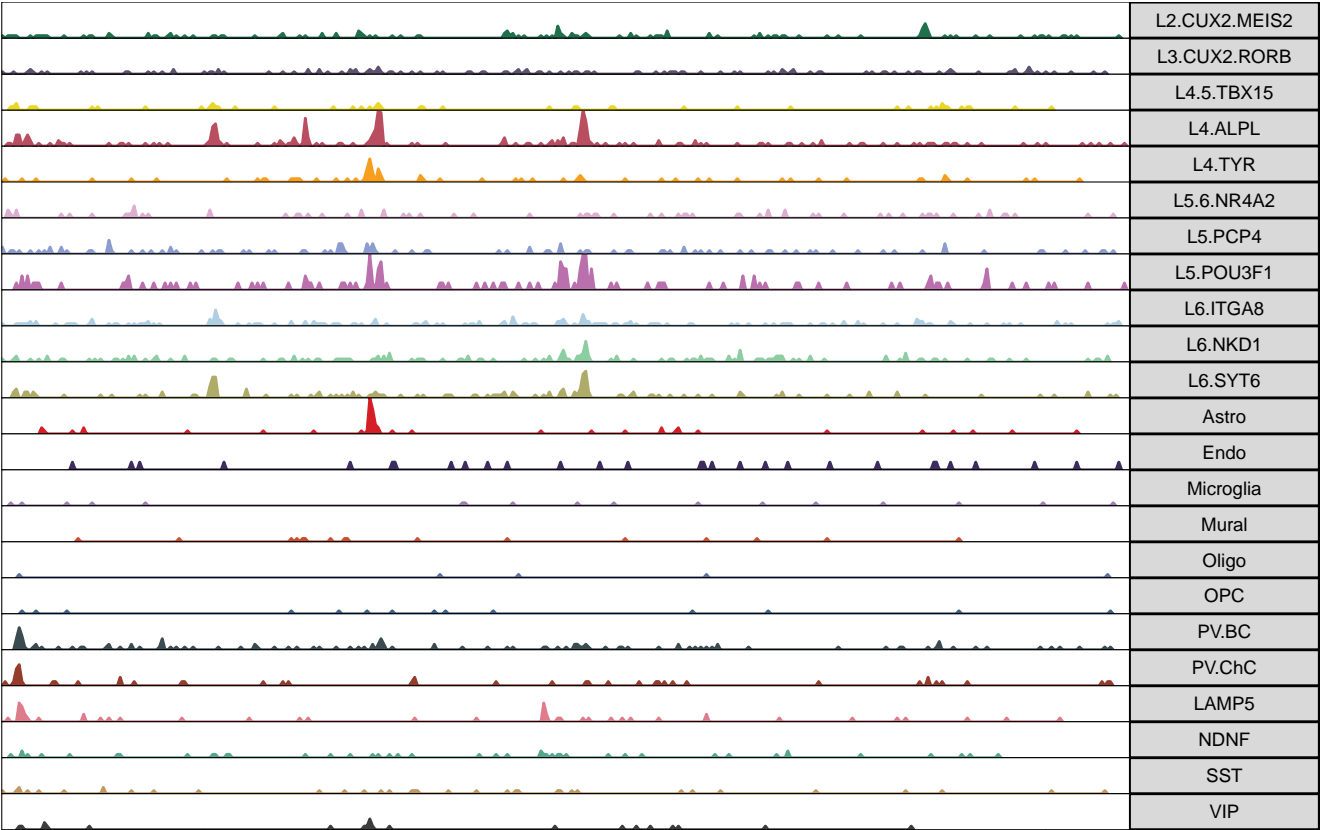
Peaks



Genes



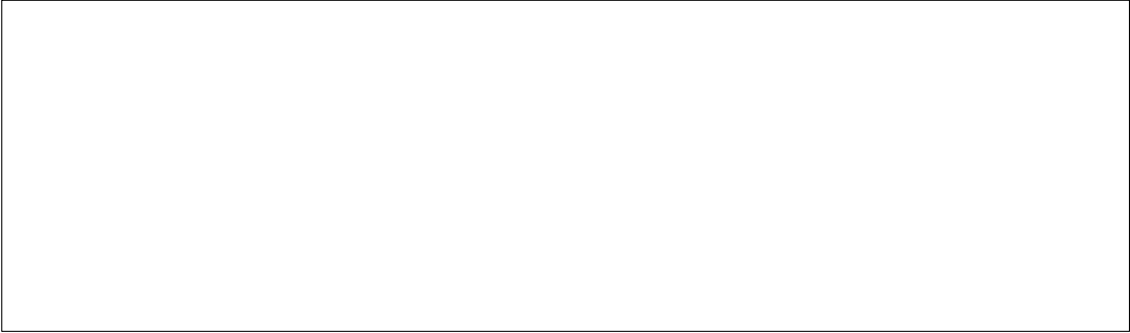
Coverage  
(Norm. ATAC Signal Range (0–0.37) by ReadsInTSS)



SST

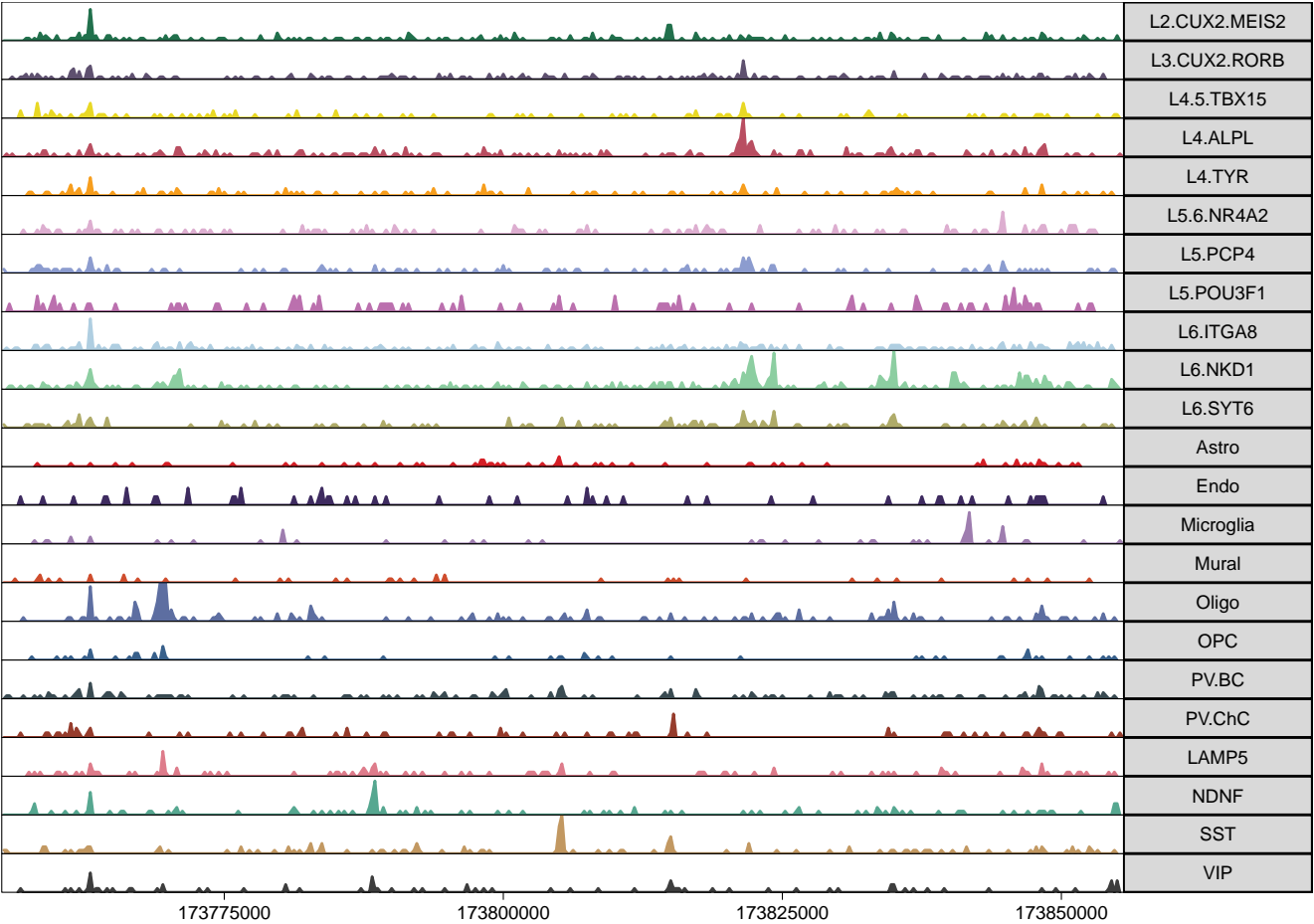


Peaks

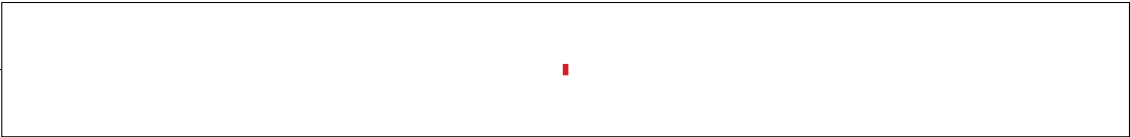


GeneTrack

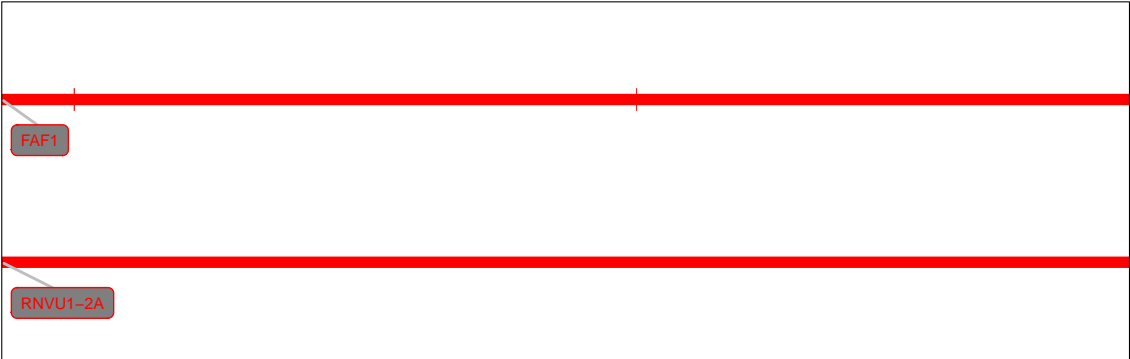
Coverage  
(Norm. ATAC Signal Range (0–0.36) by ReadsInTSS)



SST

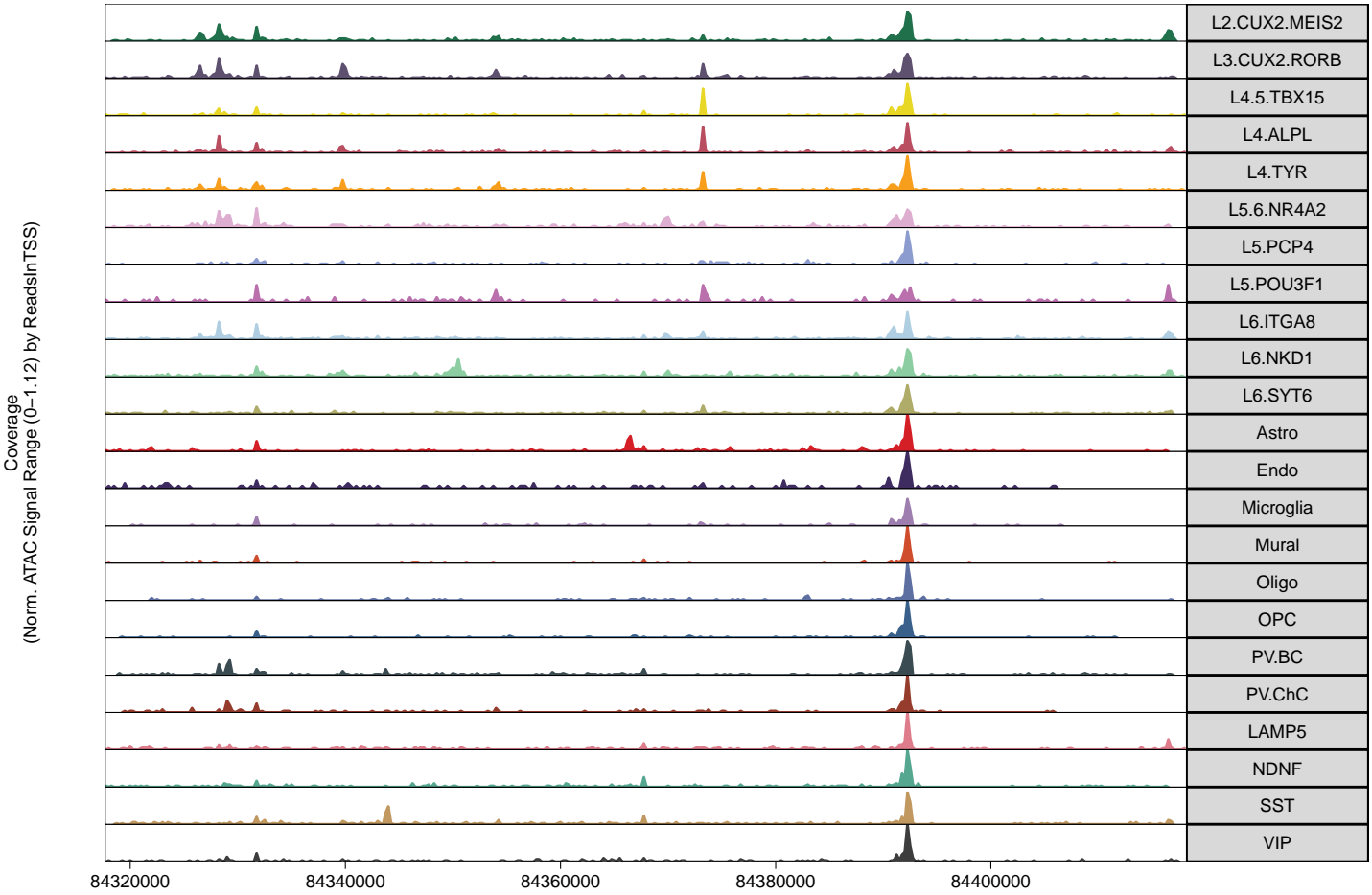


Peaks

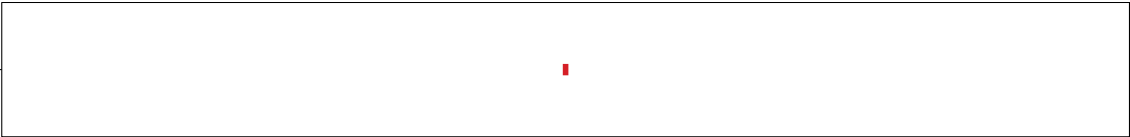


Genes

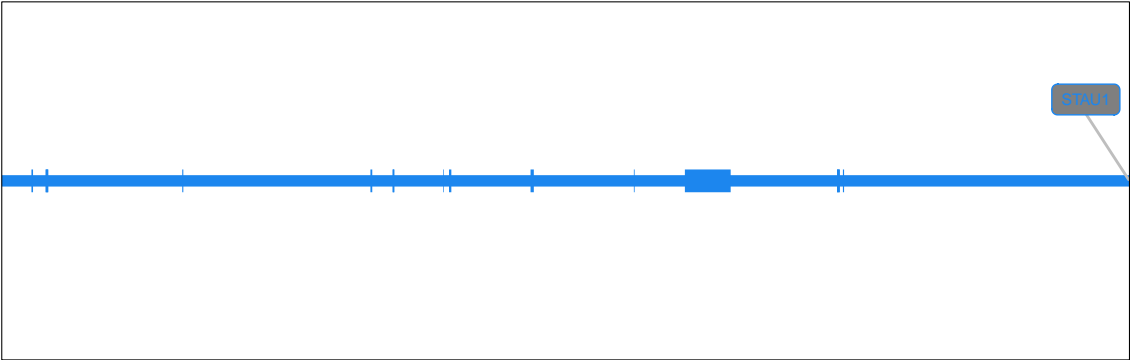
chr10:84317678–84418179 Marker Peak Tracks for: SST



SST



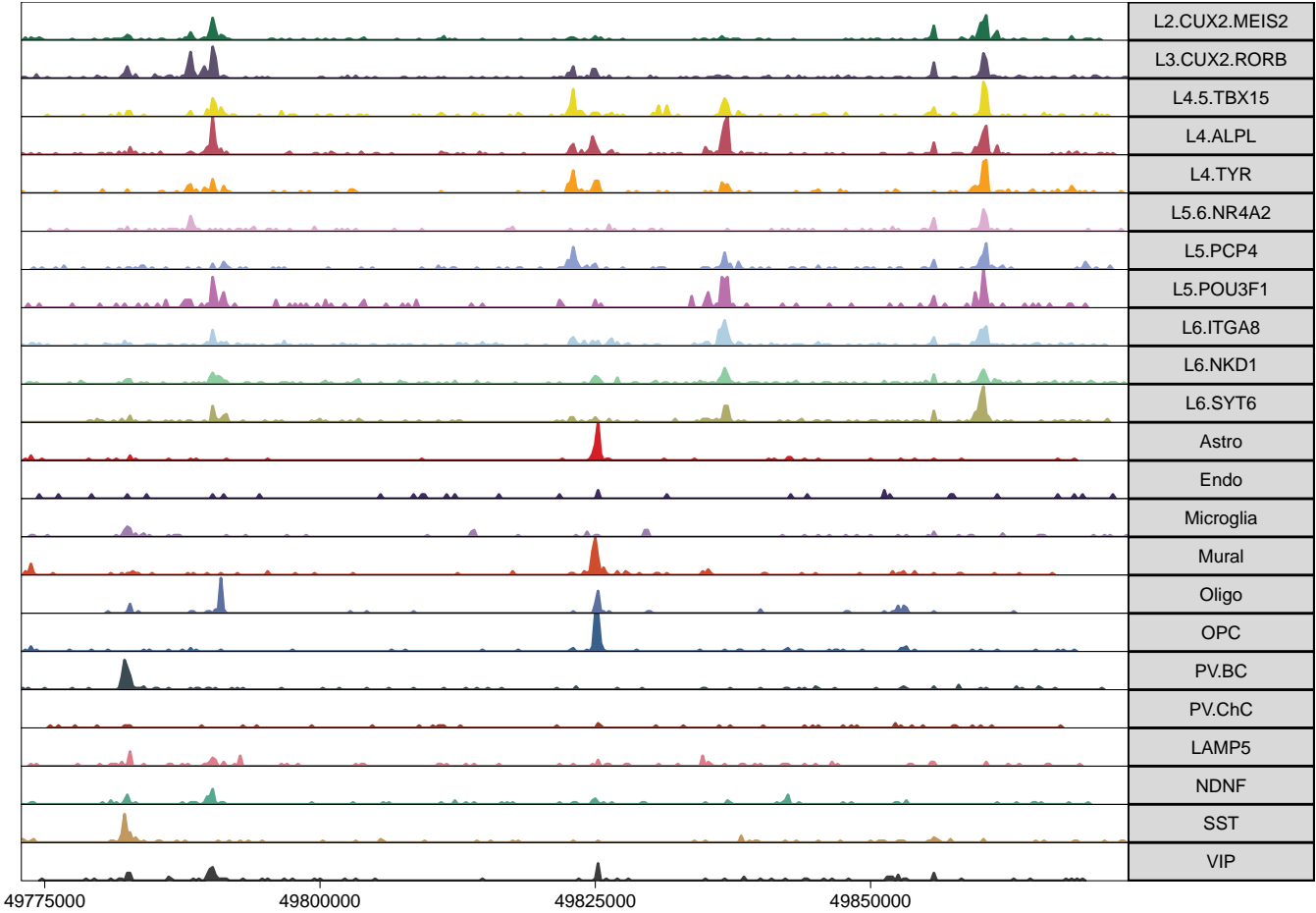
Peaks



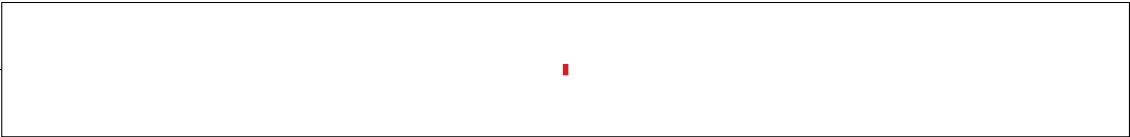
Genes

chr10:49772876–49873377 Marker Peak Tracks for: SST

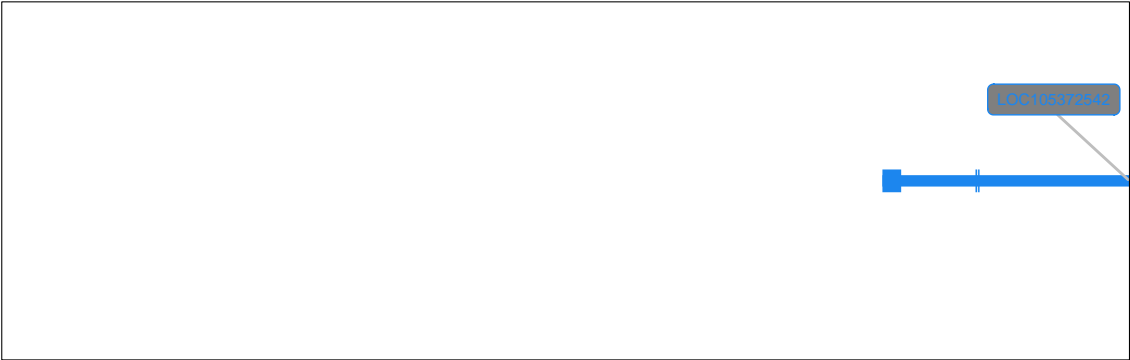
Coverage  
(Norm. ATAC Signal Range (0–0.73) by ReadsInTSS)



SST



Peaks



Genes