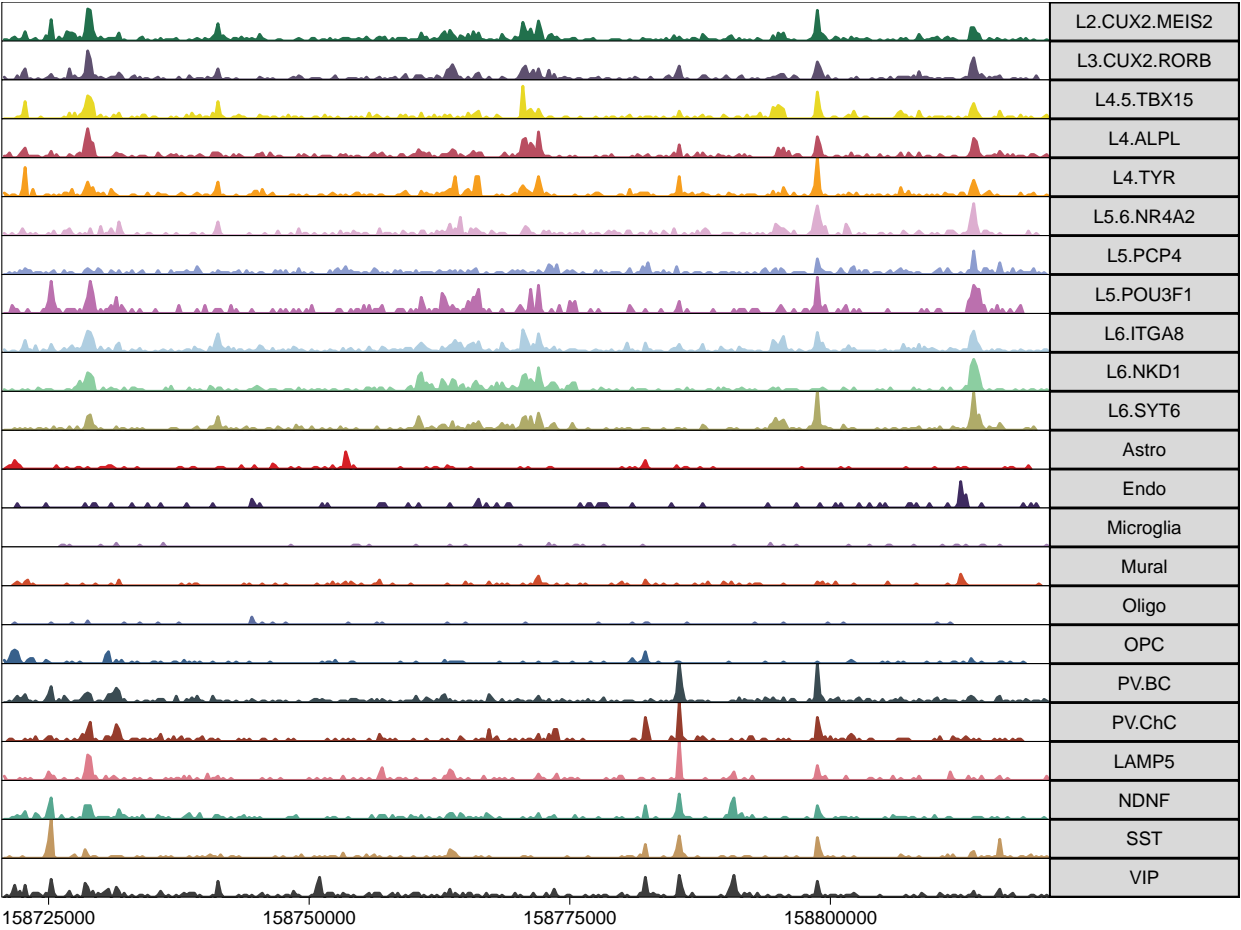
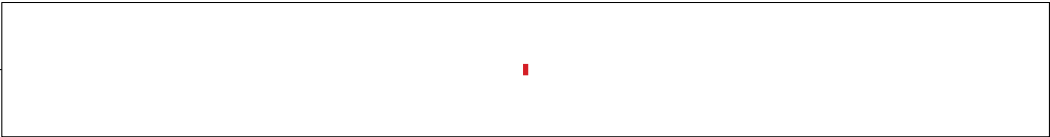


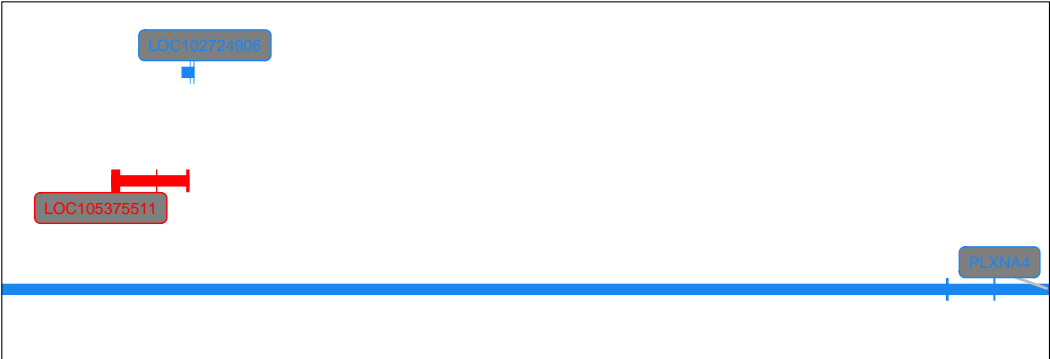
Coverage
(Norm. ATAC Signal Range (0–0.71) by ReadsInTSS)



L4.5.TBX15



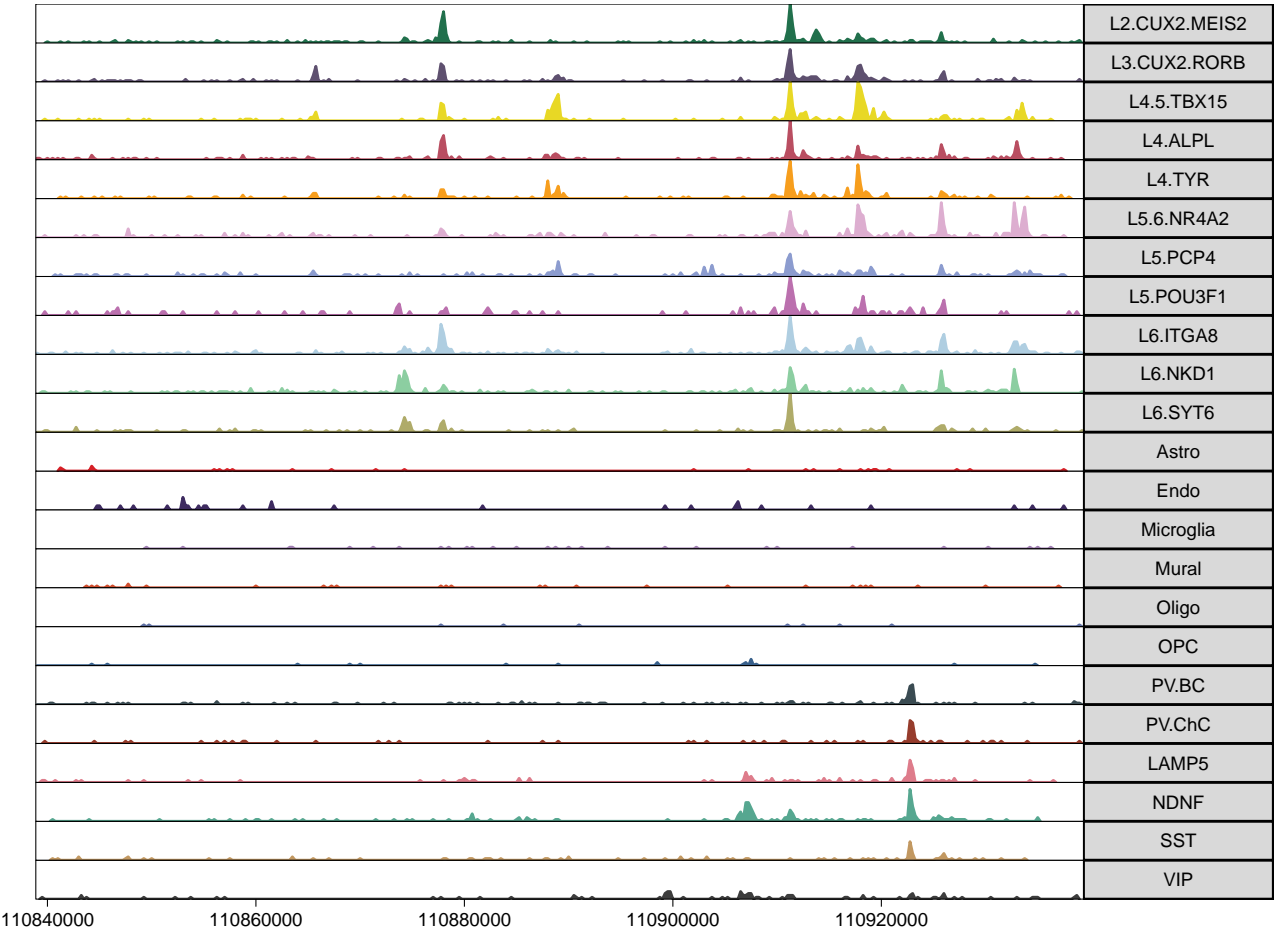
Peaks



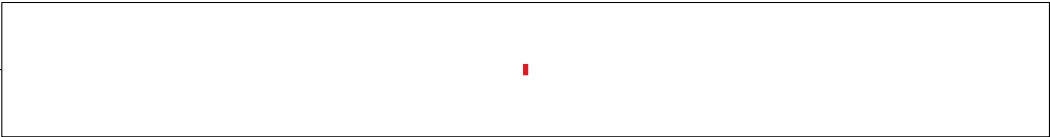
Genes

chr1:110838885–110939386 Marker Peak Tracks for: L4.5.TBX15

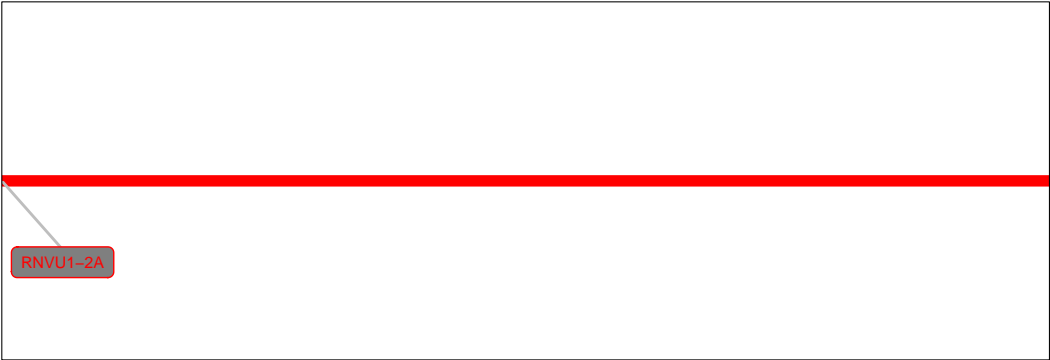
Coverage
(Norm. ATAC Signal Range (0–0.74) by ReadsInTSS)



L4.5.TBX15



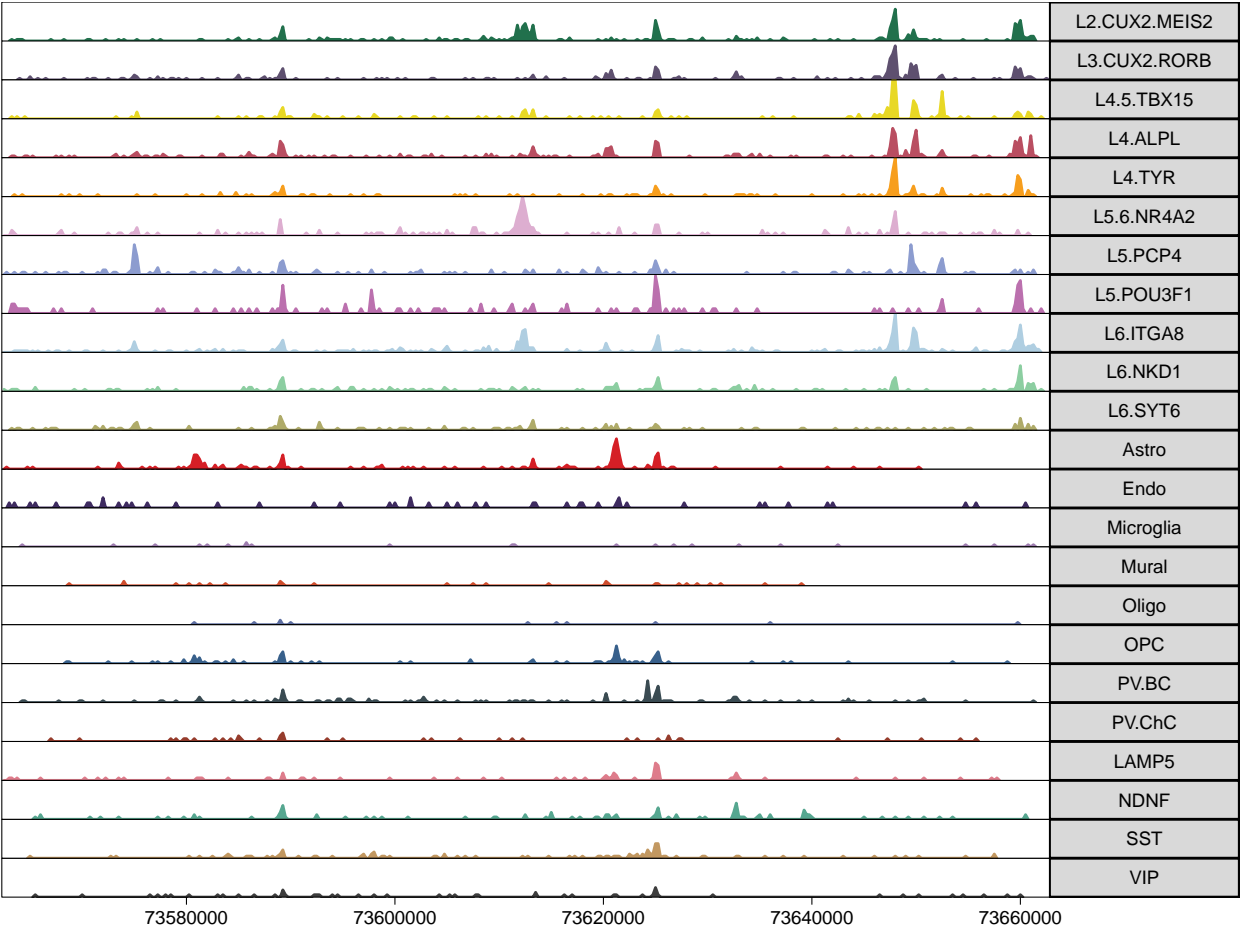
Peaks



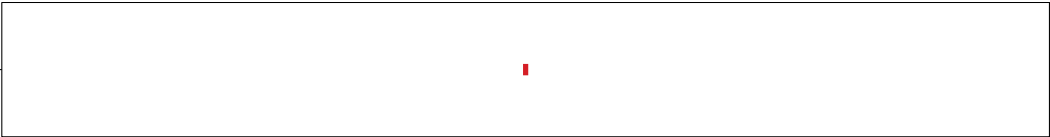
Genes

chr17:73562288–73662789 Marker Peak Tracks for: L4.5.TBX15

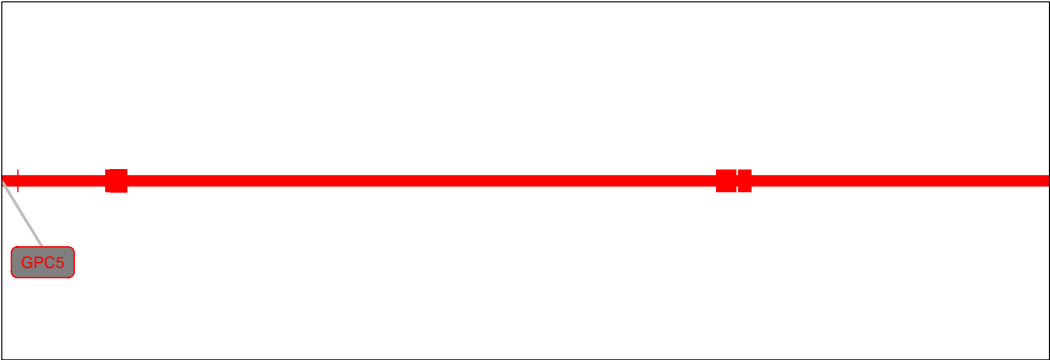
Coverage
(Norm. ATAC Signal Range (0–0.61) by ReadsInTSS)



L4.5.TBX15

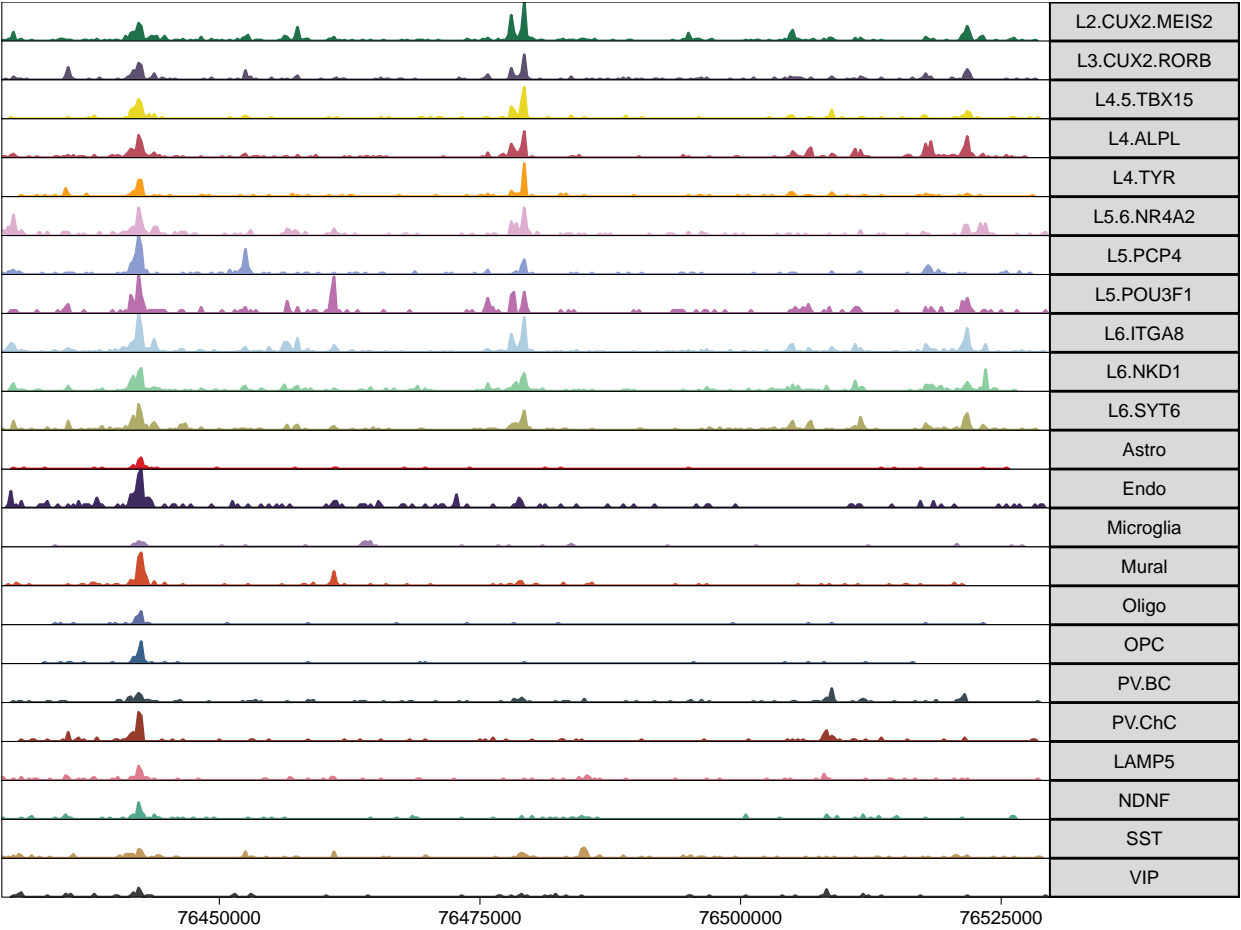


Peaks

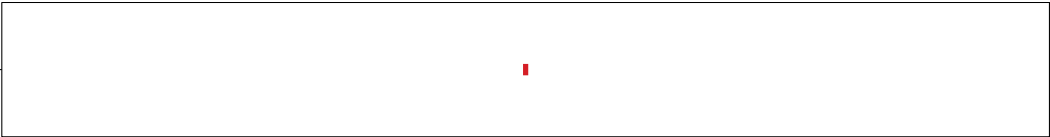


Genes

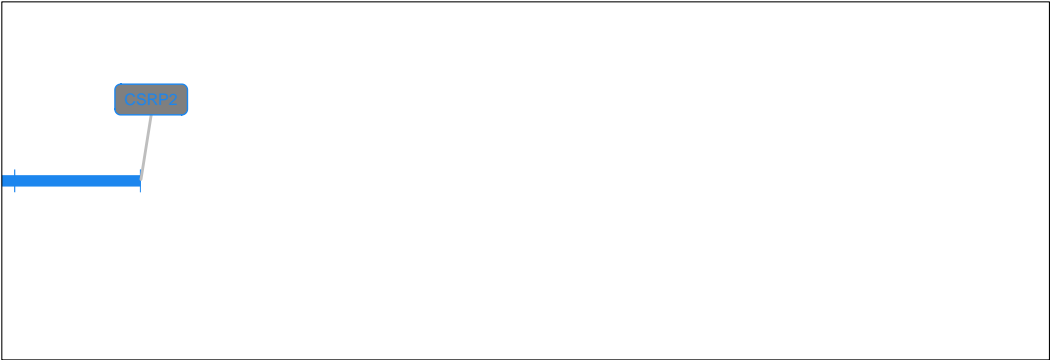
Coverage
(Norm. ATAC Signal Range (0–0.94) by ReadsInTSS)



L4.5.TBX15



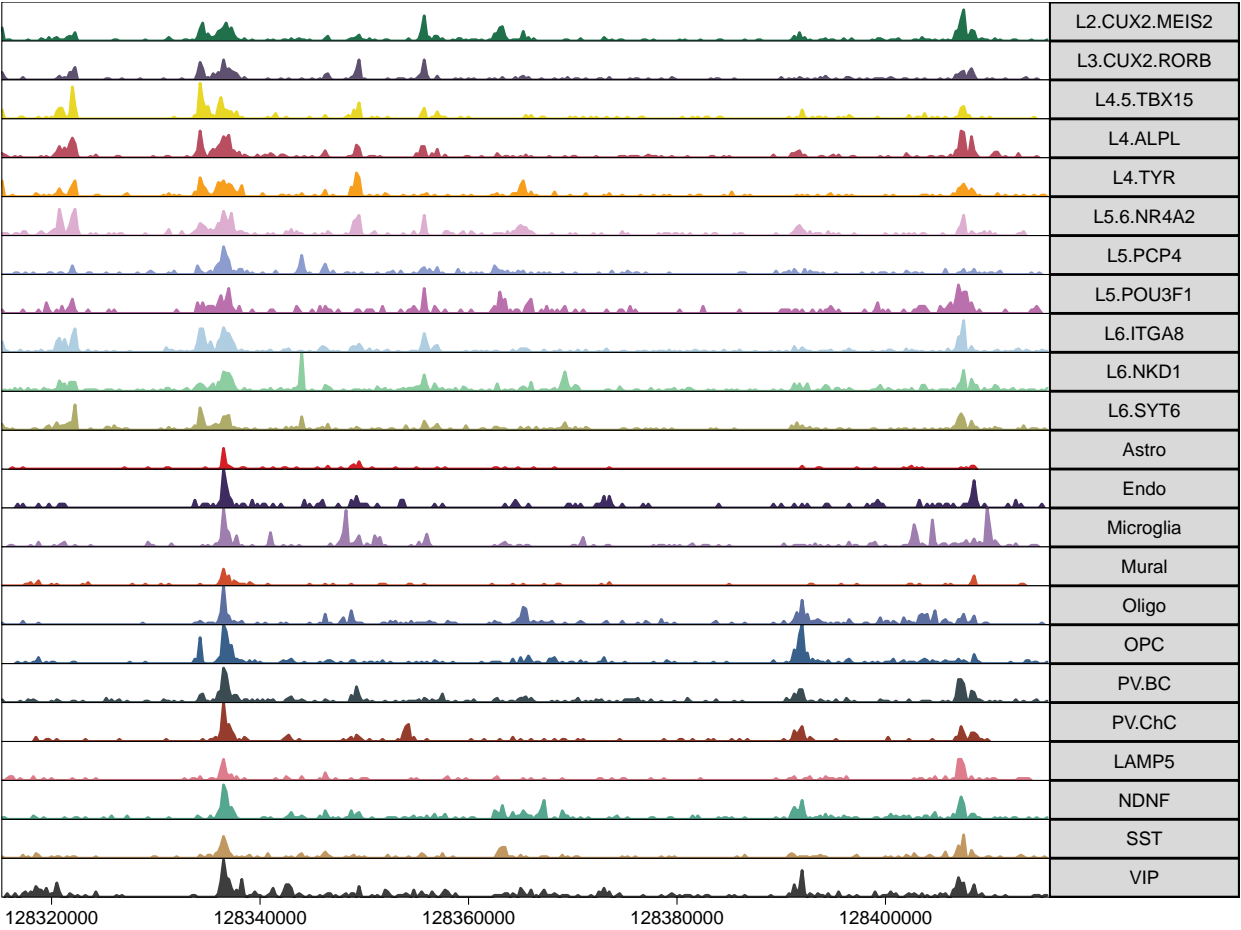
Peaks



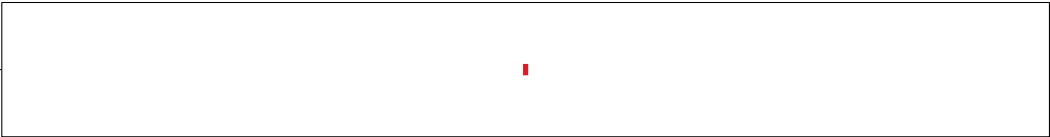
Genes

chr9:128315227–128415728 Marker Peak Tracks for: L4.5.TBX15

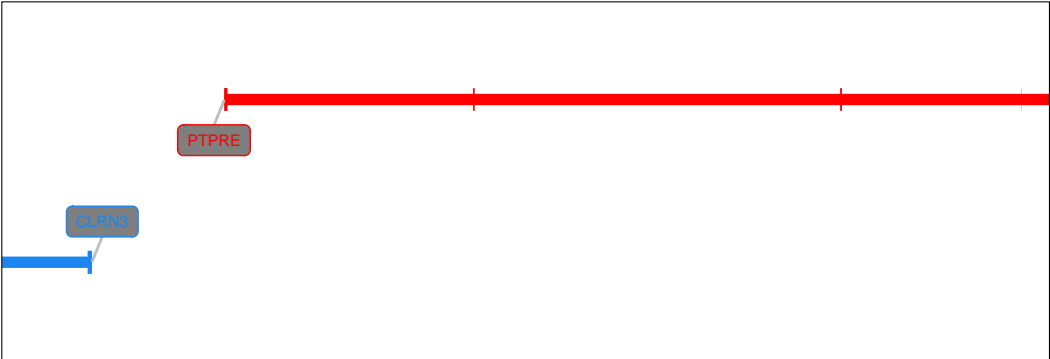
Coverage
(Norm. ATAC Signal Range (0–0.8) by ReadsInTSS)



L4.5.TBX15



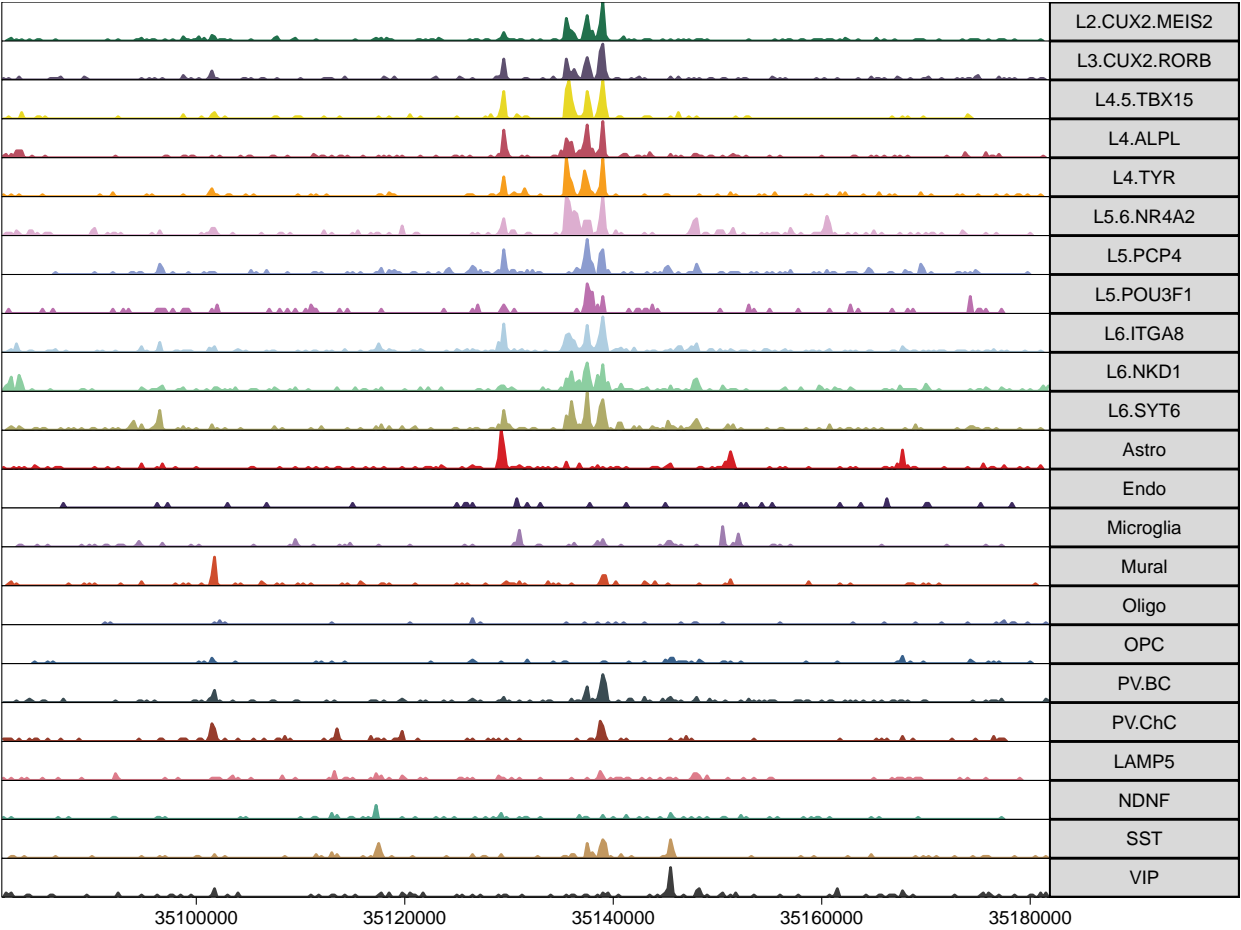
Peaks



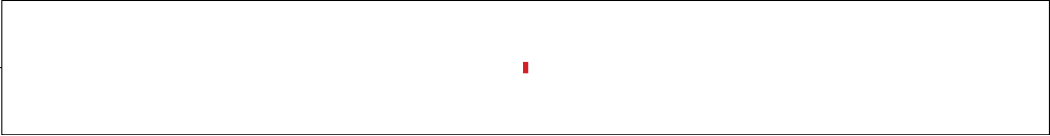
Genes

chr9:35081344–35181845 Marker Peak Tracks for: L4.5.TBX15

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



L4.5.TBX15

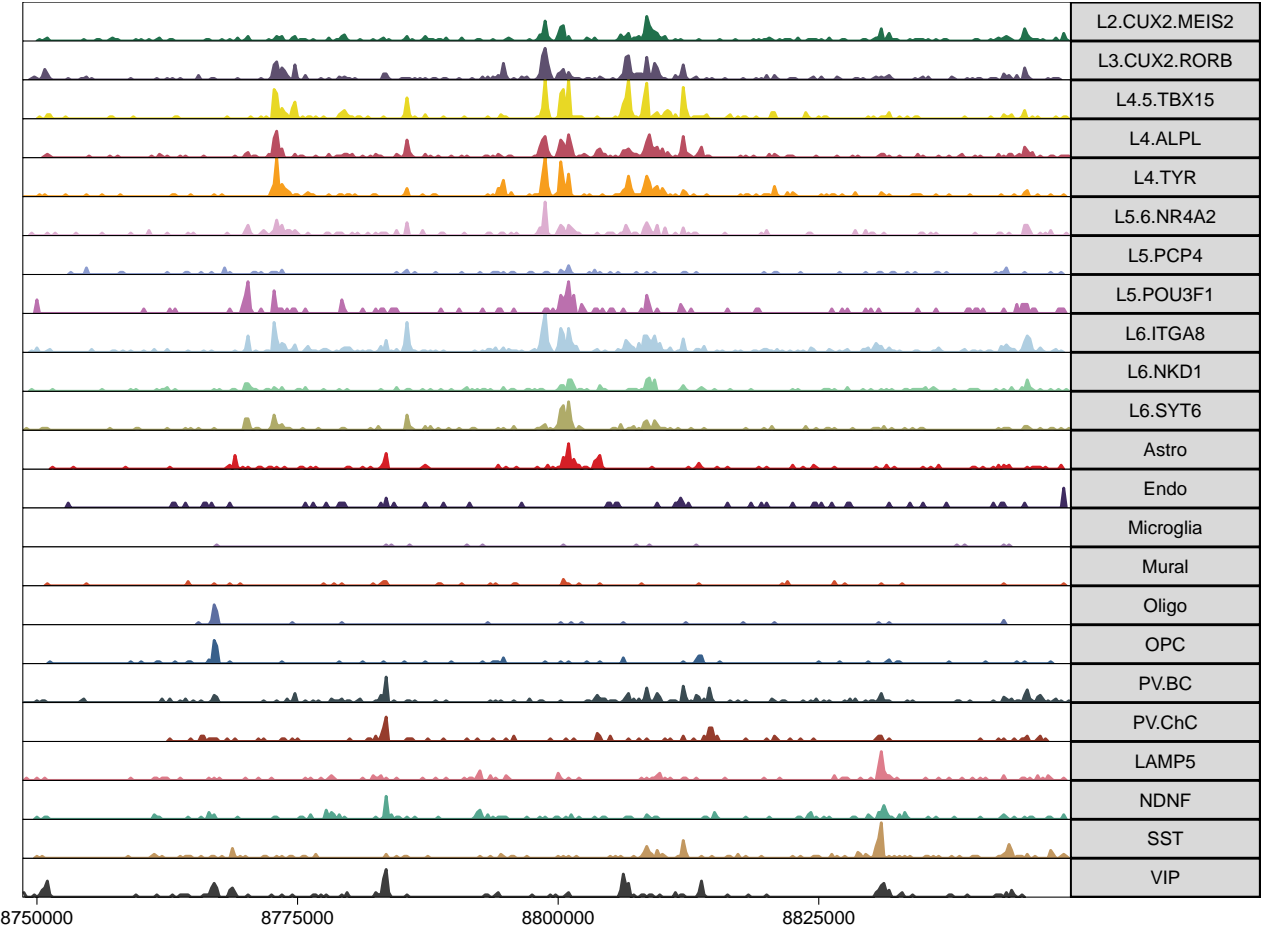


Peaks

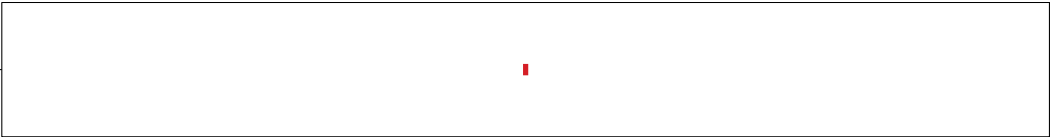


Genes

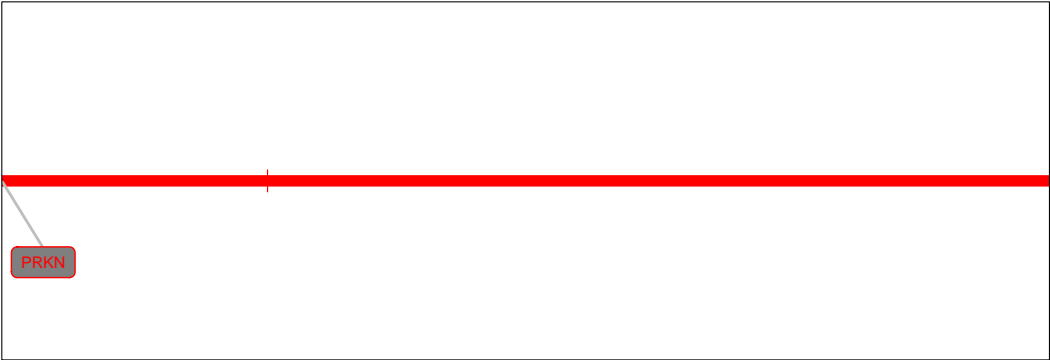
Coverage
(Norm. ATAC Signal Range (0–0.63) by ReadsInTSS)



L4.5.TBX15

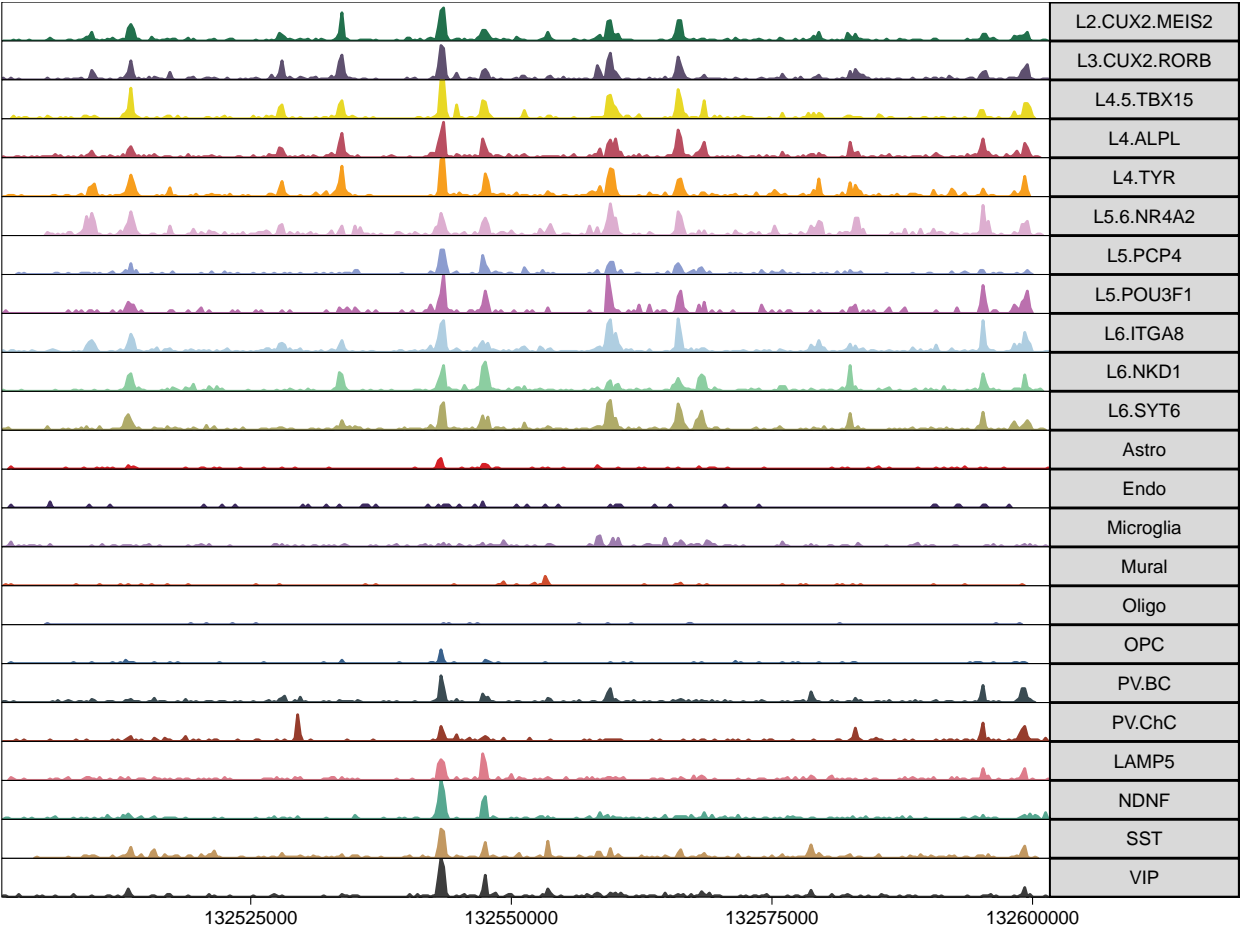


Peaks

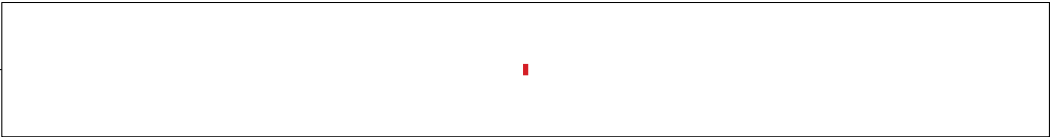


Genes

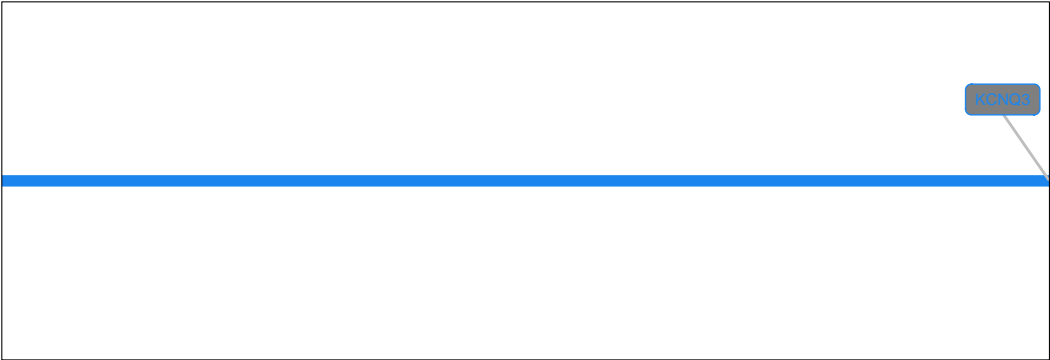
Coverage
(Norm. ATAC Signal Range (0–1.02) by ReadsInTSS)



L4.5.TBX15

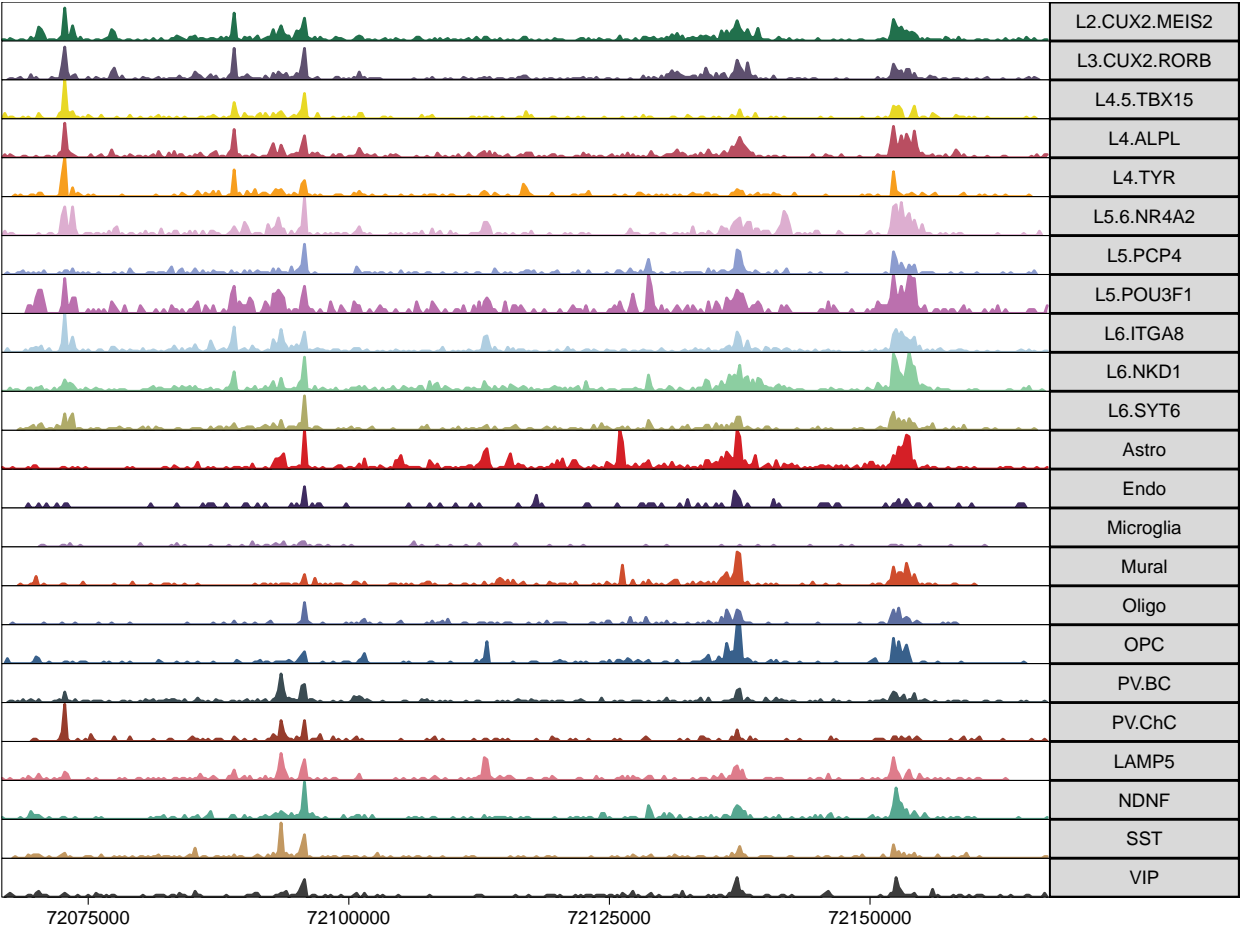


Peaks

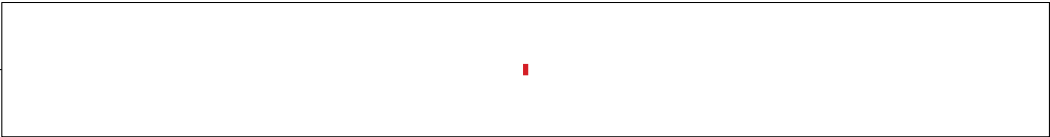


Genes

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



L4.5.TBX15

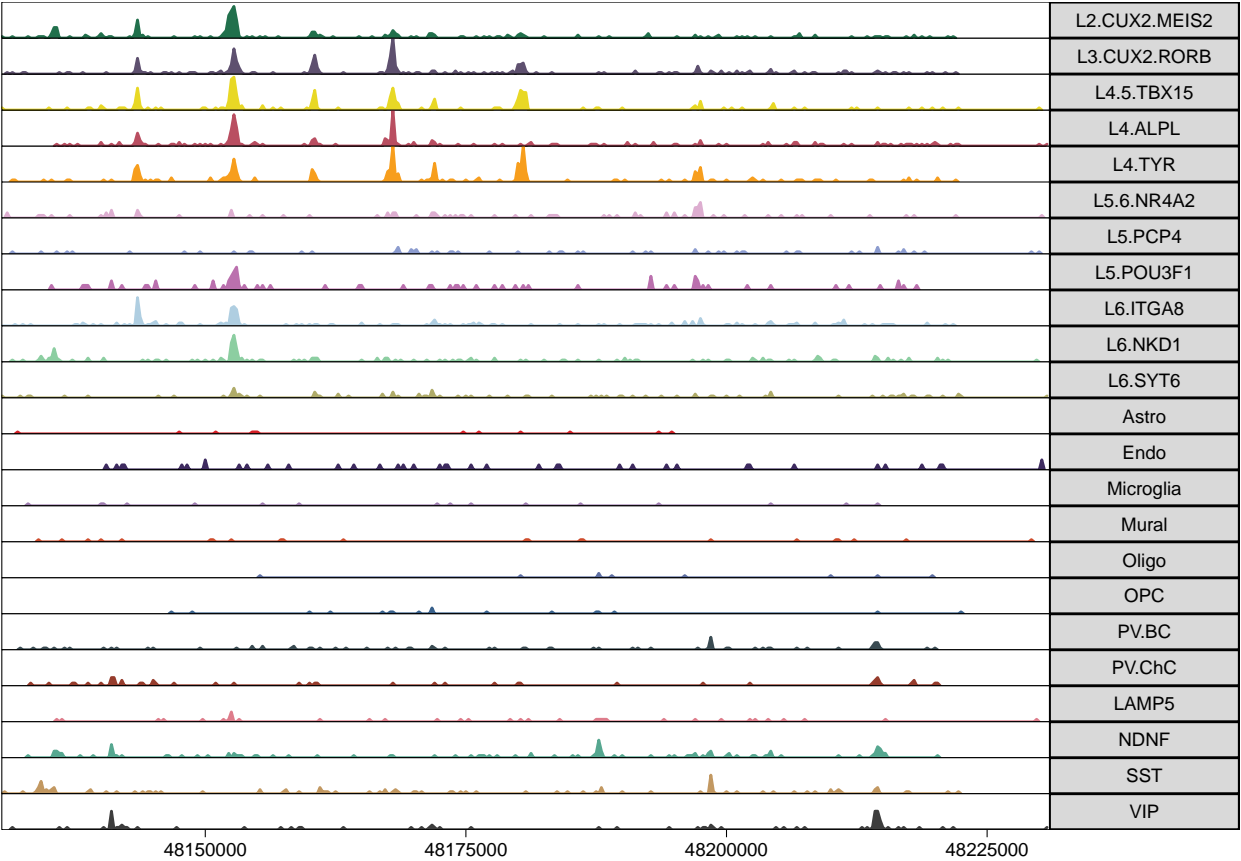


Peaks

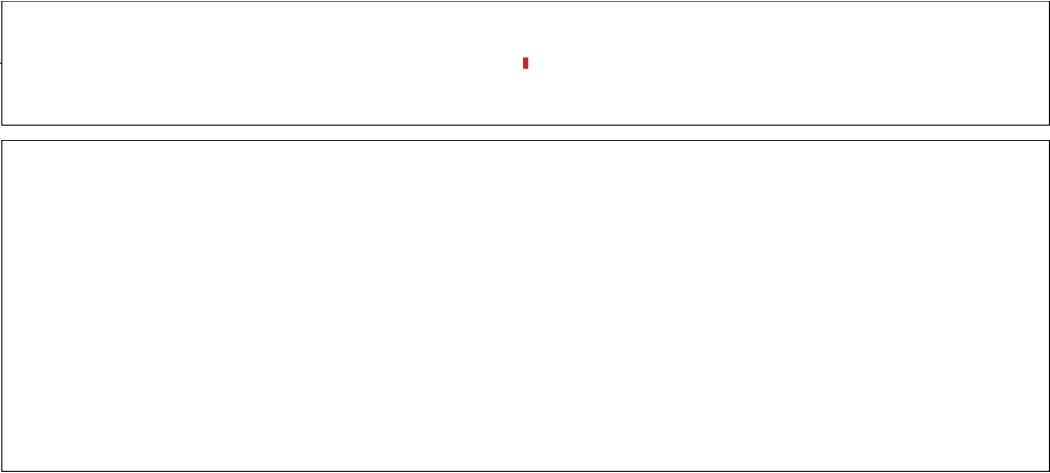


Genes

Coverage
(Norm. ATAC Signal Range (0–0.58) by ReadsInTSS)



L4.5.TBX15

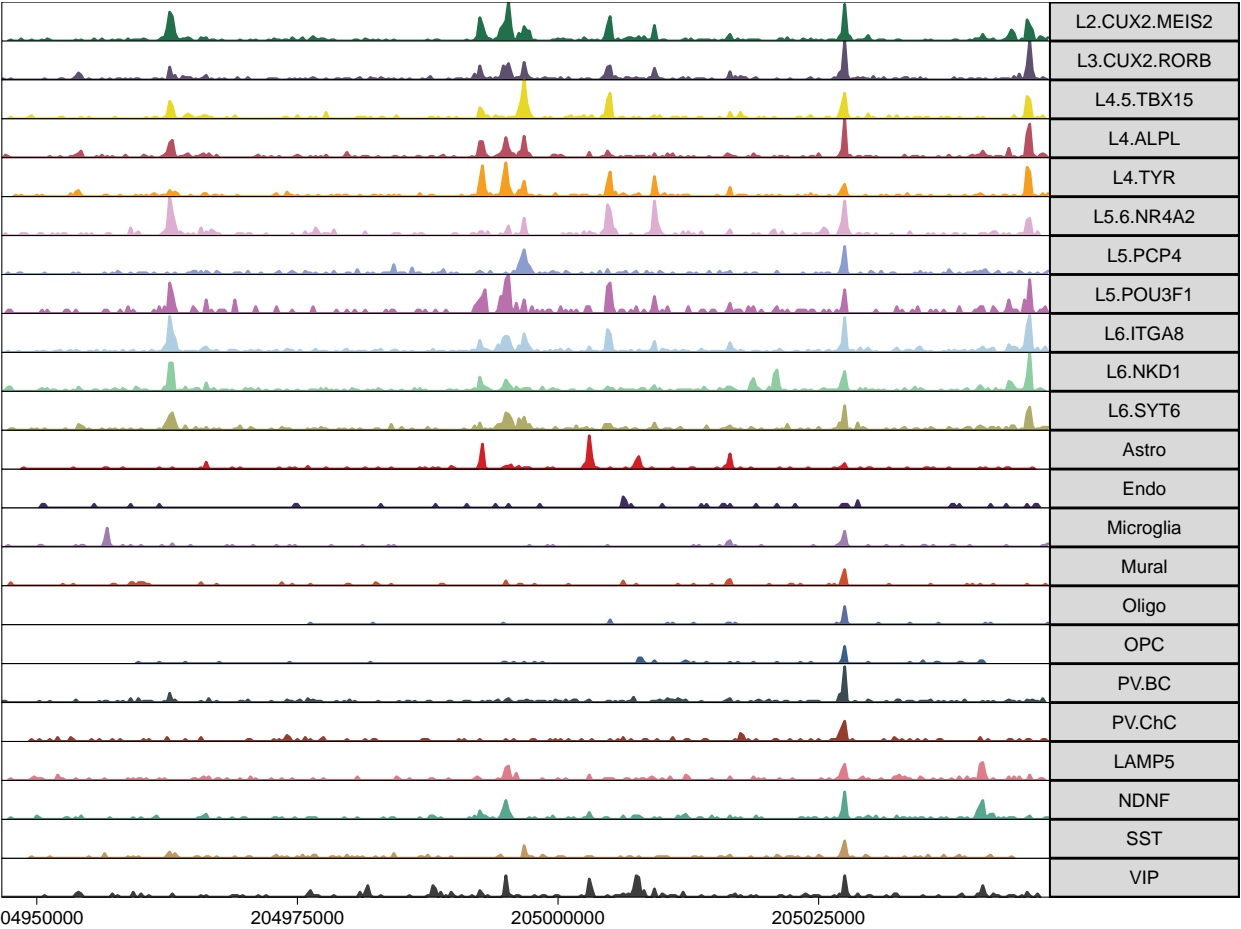


Peaks

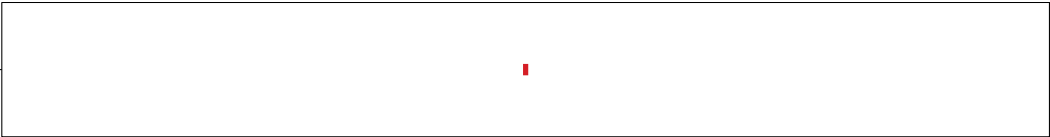
GeneTrack

chr1:204946642–205047143 Marker Peak Tracks for: L4.5.TBX15

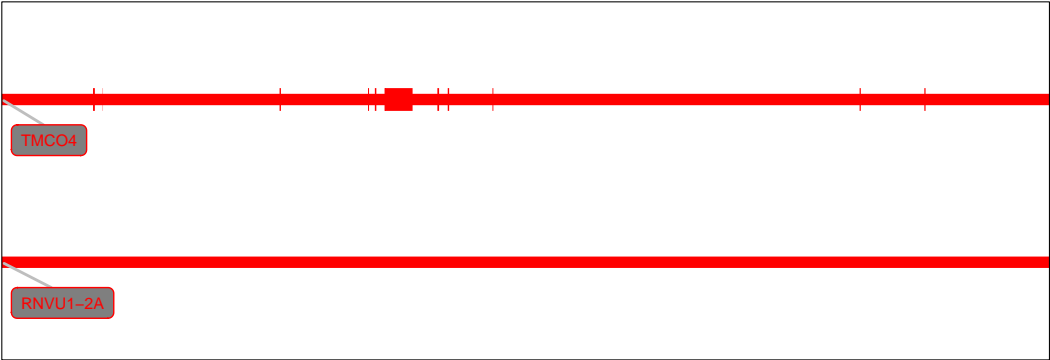
Coverage
(Norm. ATAC Signal Range (0–0.85) by ReadsInTSS)



L4.5.TBX15

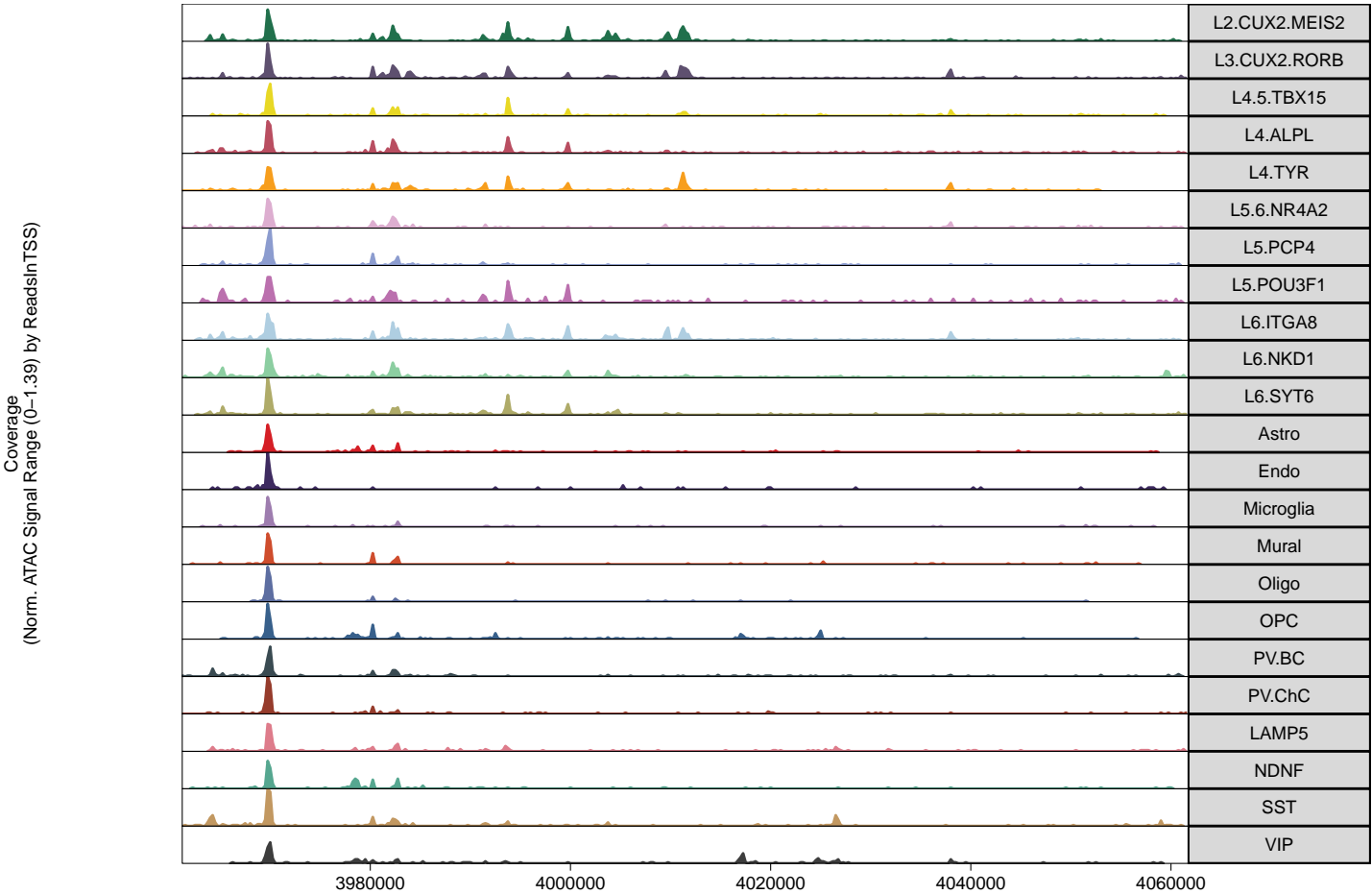


Peaks

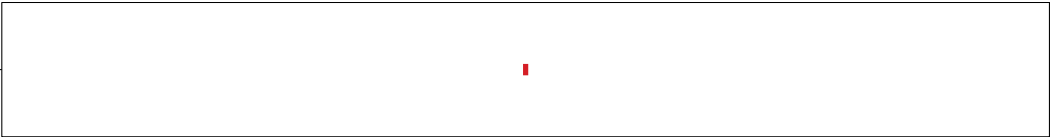


Genes

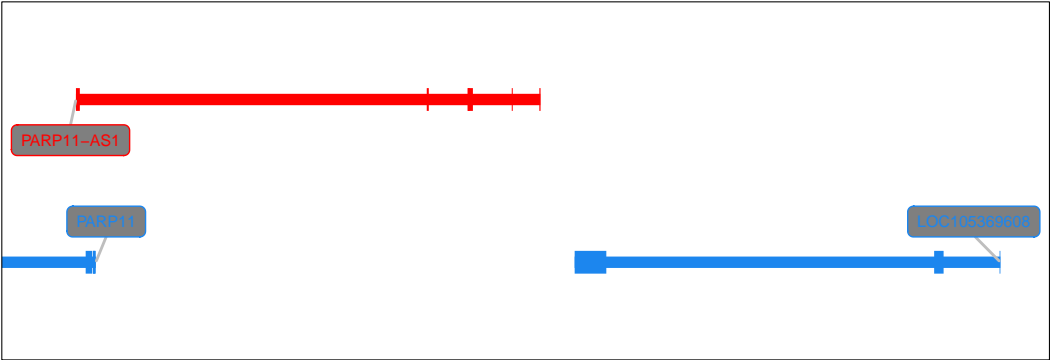
chr11:3961209-4061710 Marker Peak Tracks for: L4.5.TBX15



L4.5.TBX15

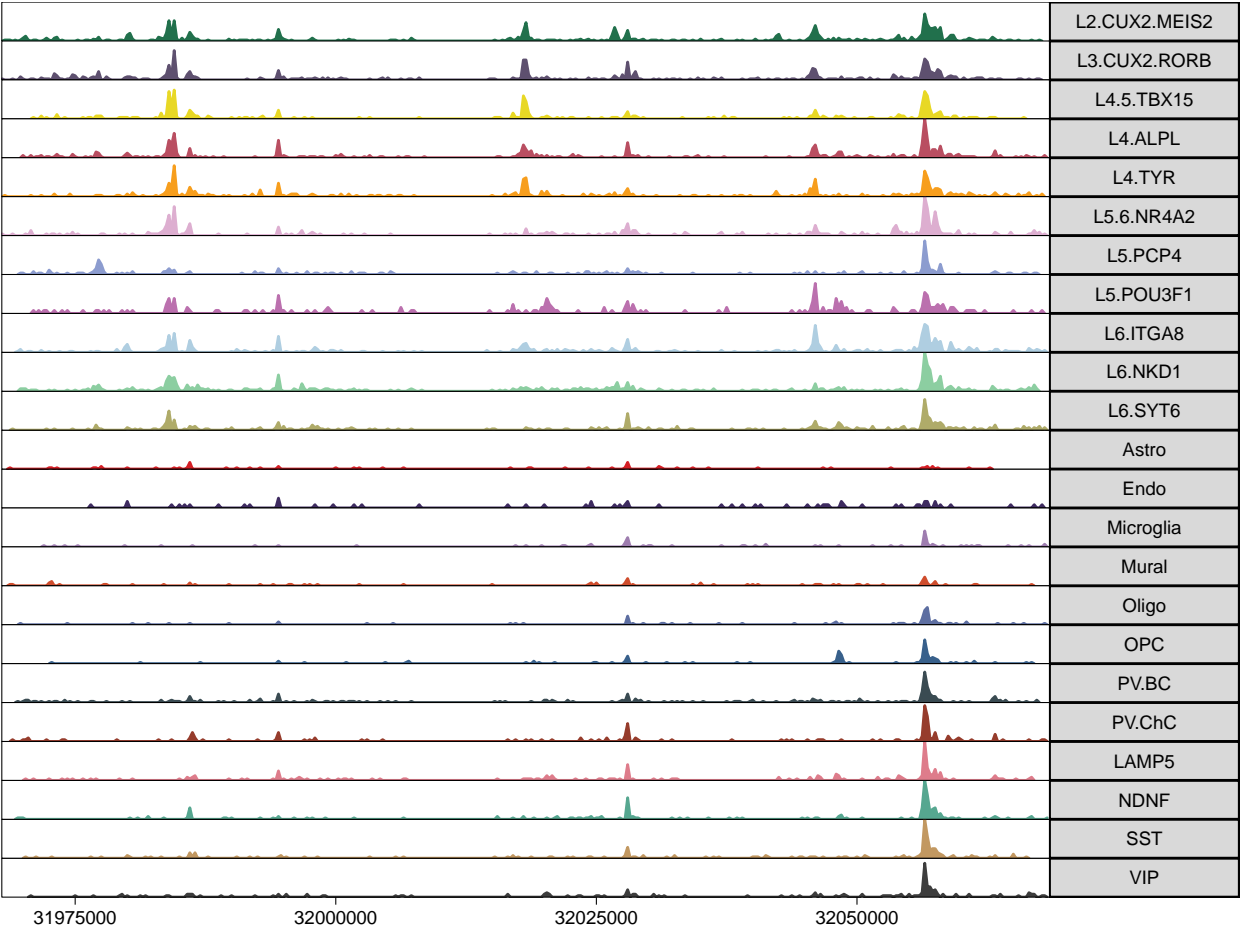


Peaks

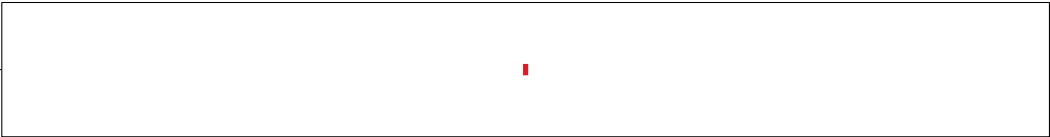


Genes

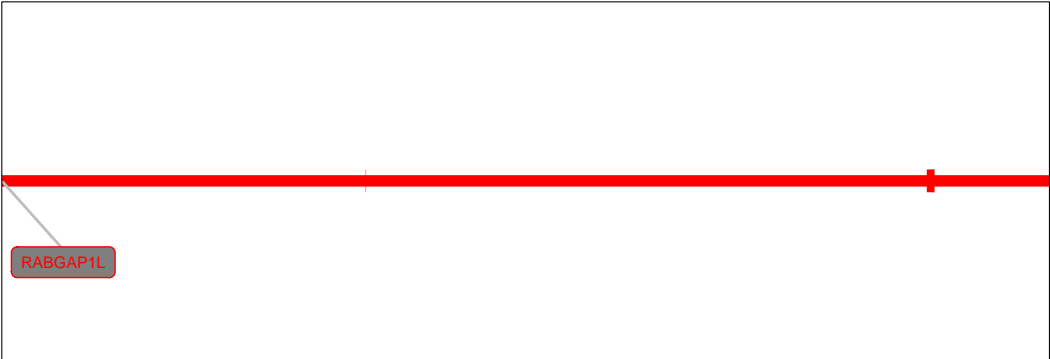
Coverage
(Norm. ATAC Signal Range (0-0.95) by ReadsInTSS)



L4.5.TBX15

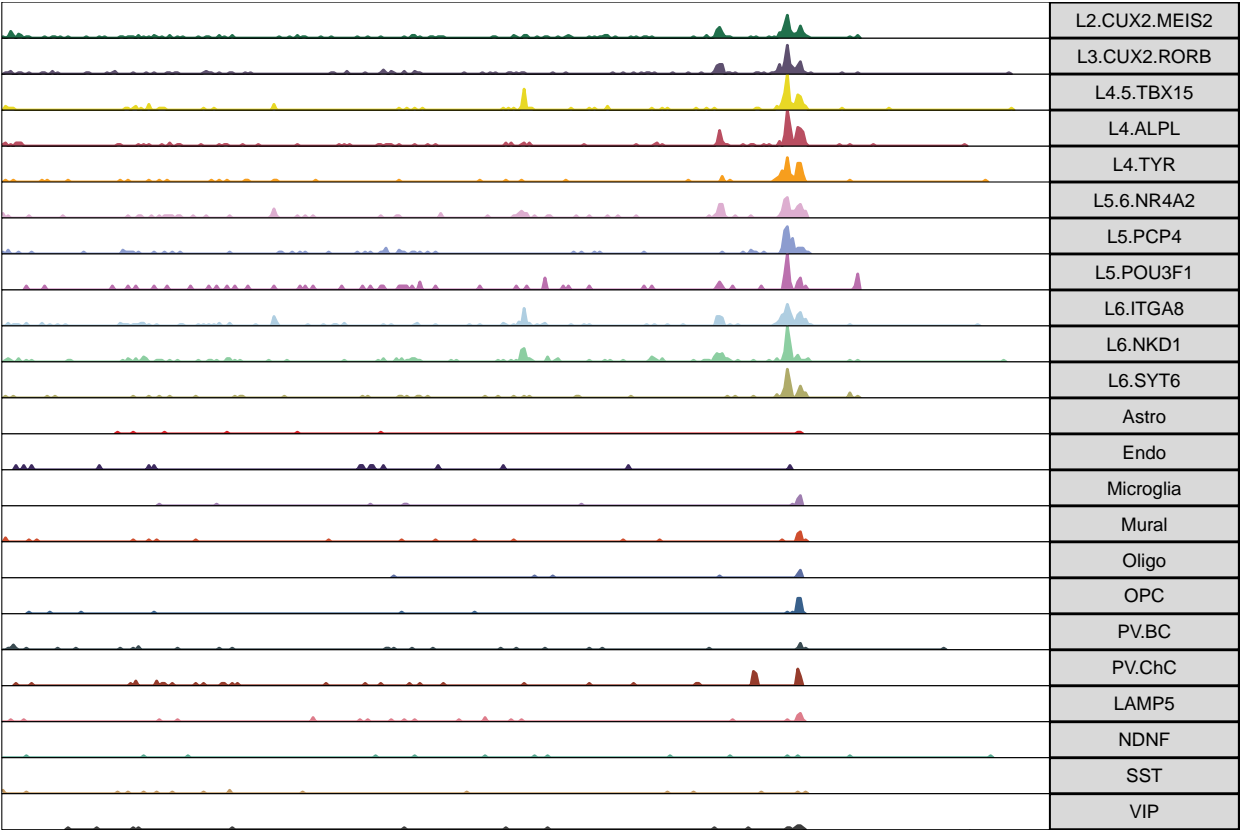


Peaks

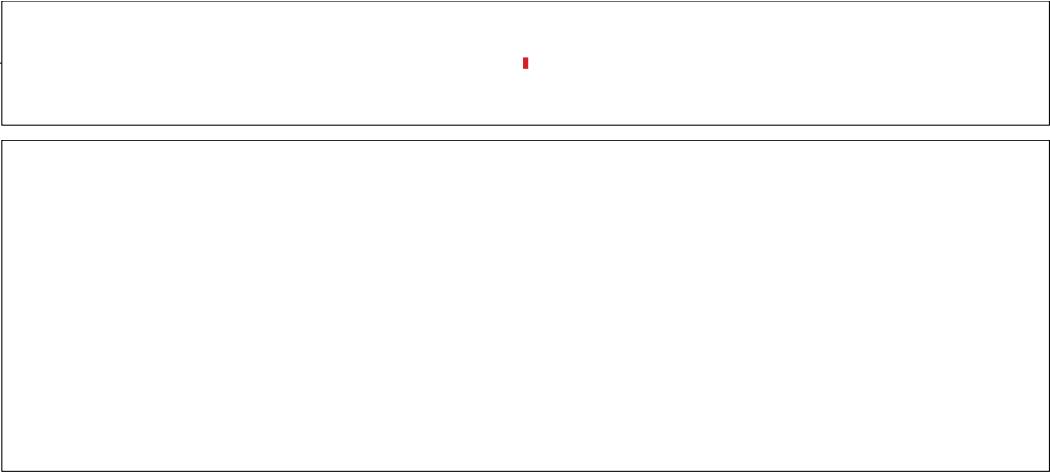


Genes

Coverage
(Norm. ATAC Signal Range (0–0.66) by ReadsInTSS)



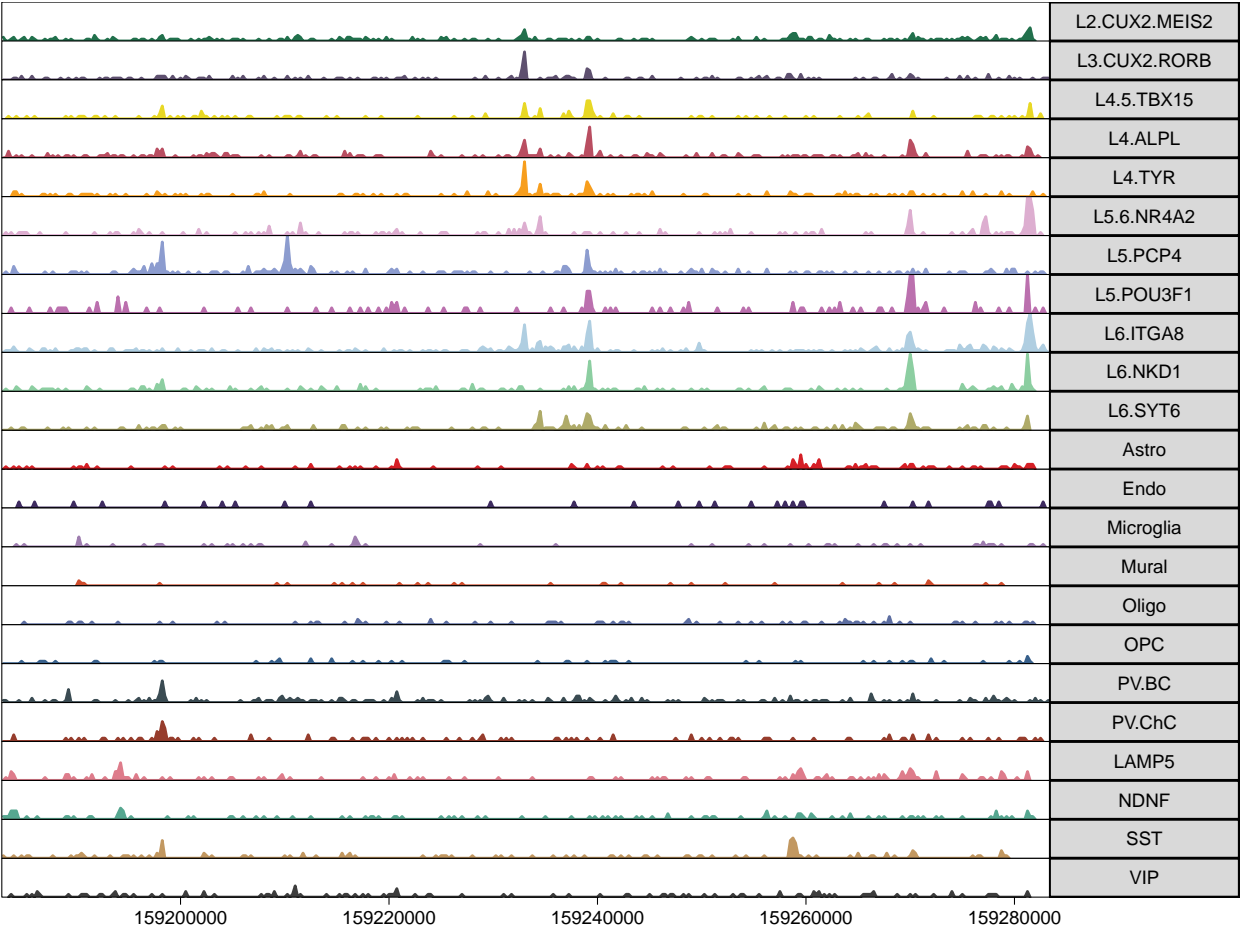
L4.5.TBX15



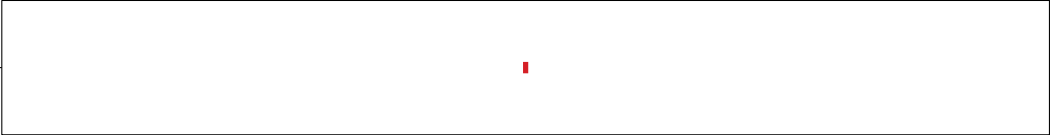
Peaks

GeneTrack

Coverage
(Norm. ATAC Signal Range (0–0.51) by ReadsInTSS)



L4.5.TBX15

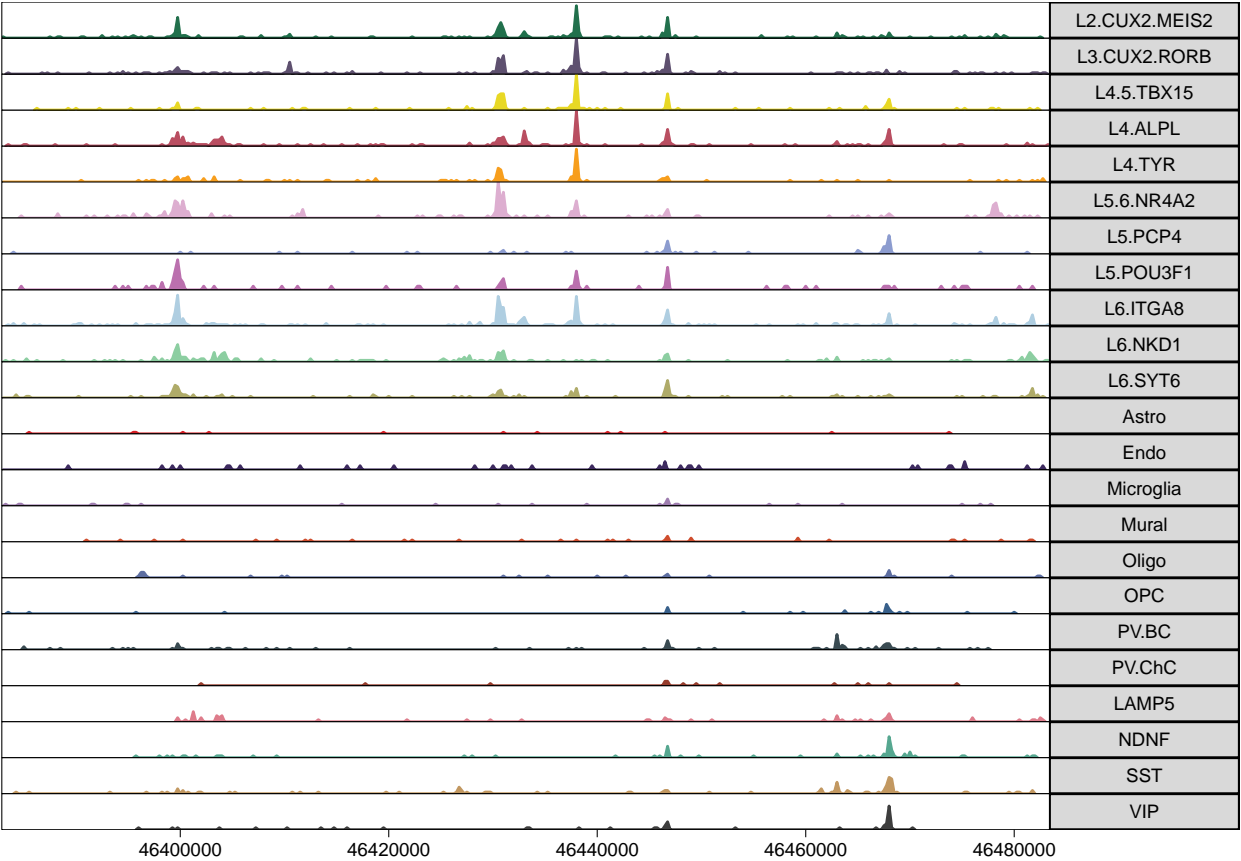


Peaks

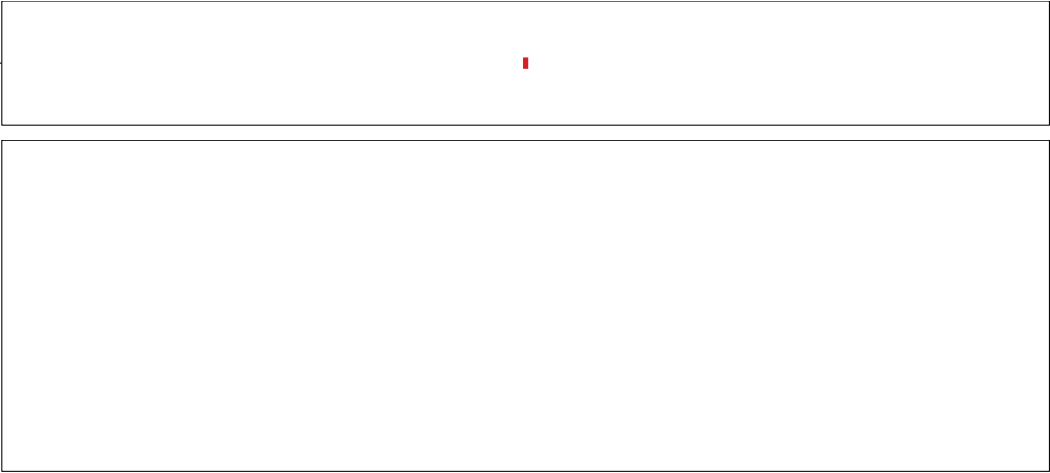


Genes

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



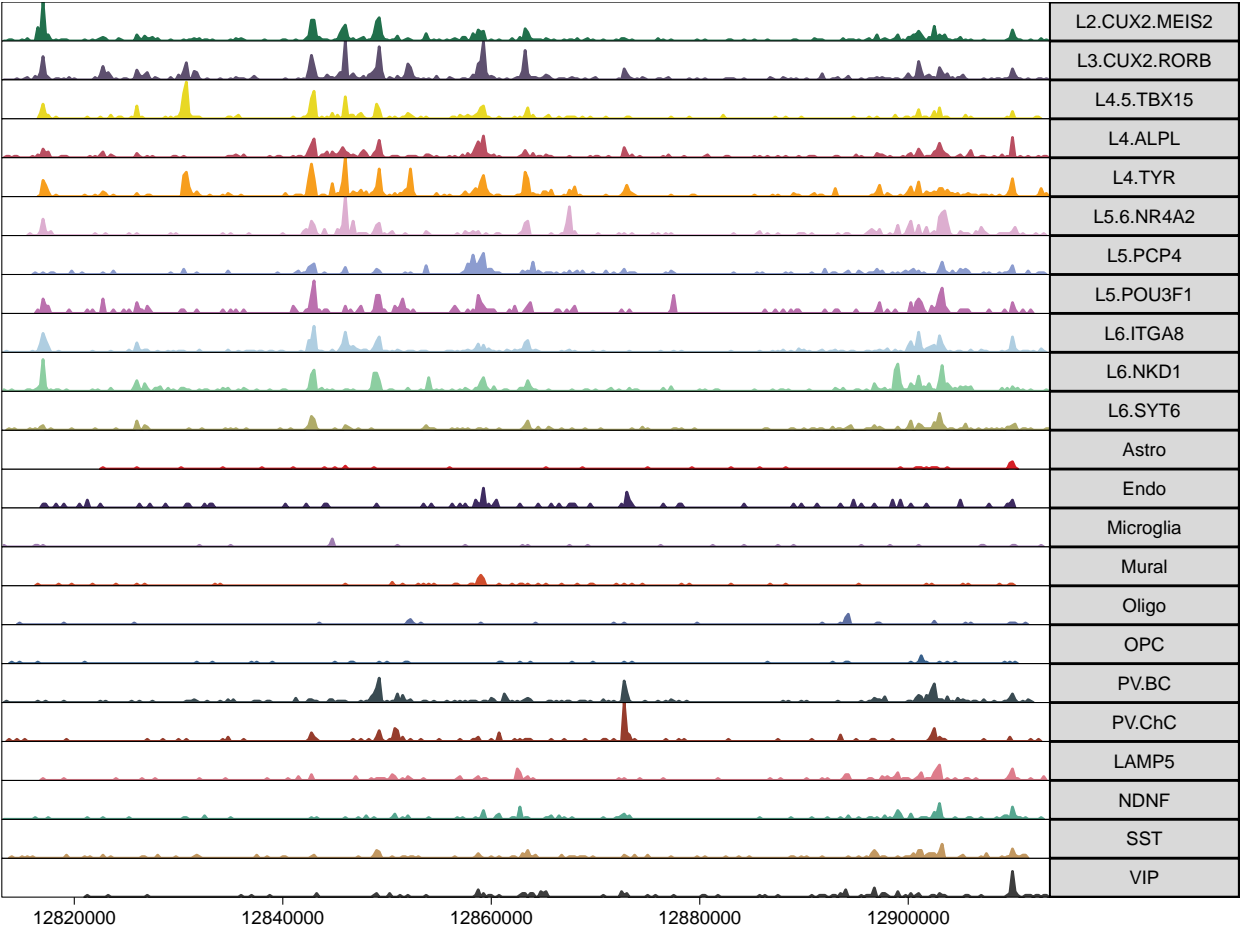
L4.5.TBX15



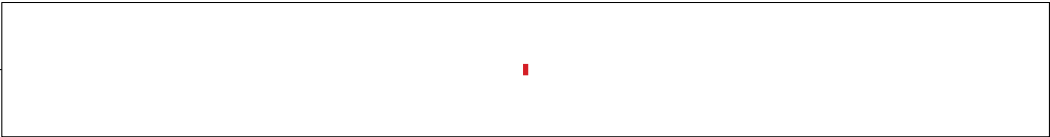
Peaks

GeneTrack

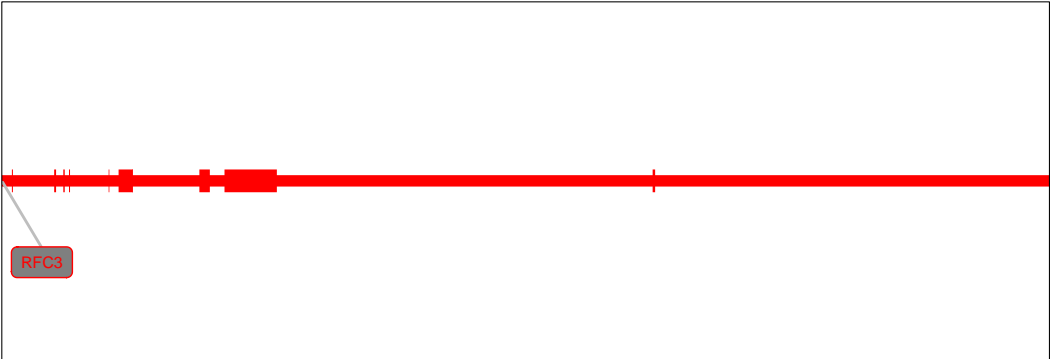
Coverage
(Norm. ATAC Signal Range (0–0.79) by ReadsInTSS)



L4.5.TBX15

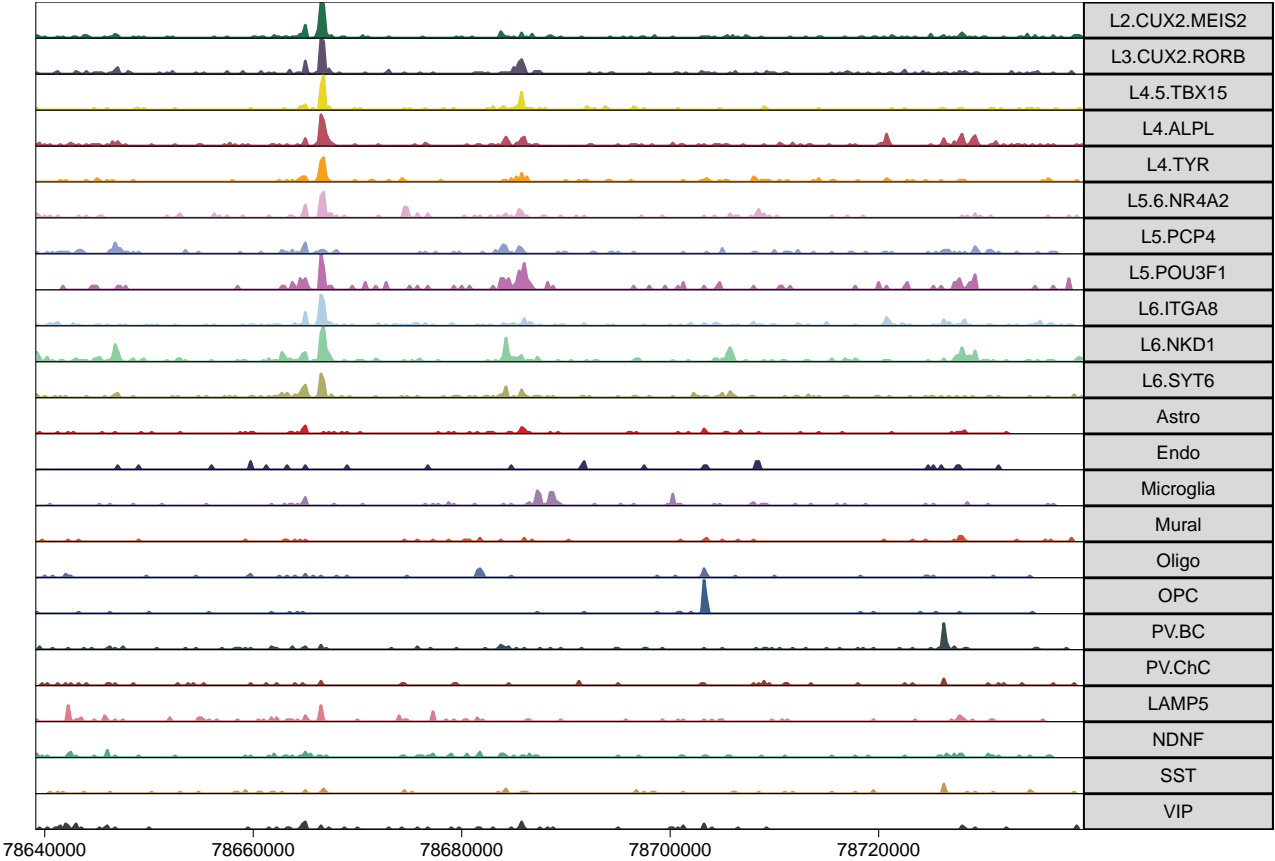


Peaks



Genes

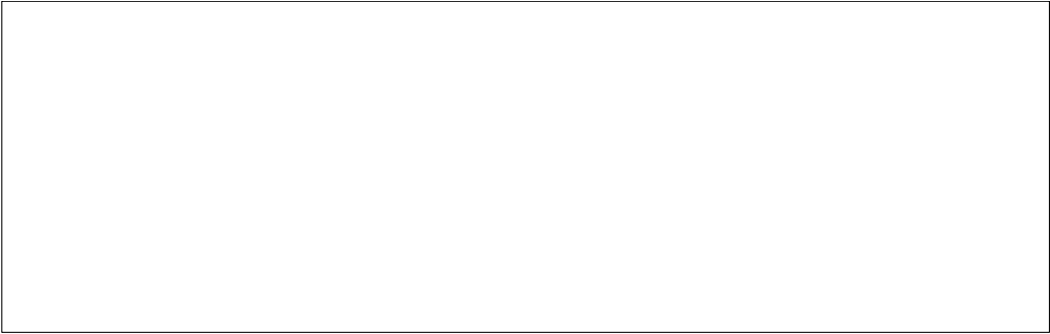
Coverage
(Norm. ATAC Signal Range (0–0.7) by ReadsInTSS)



L4.5.TBX15

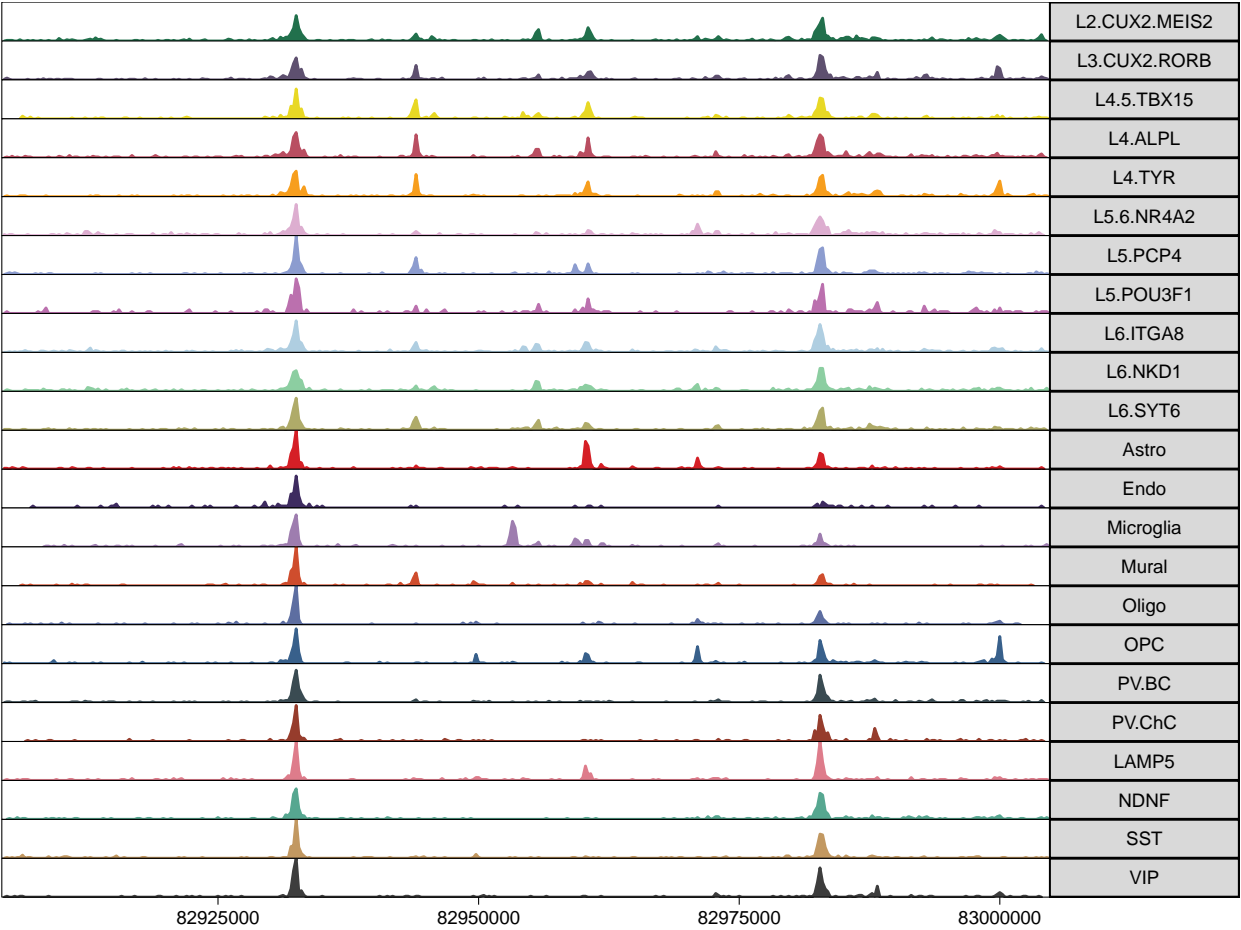


Peaks

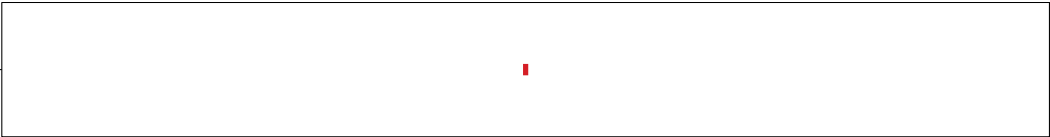


GeneTrack

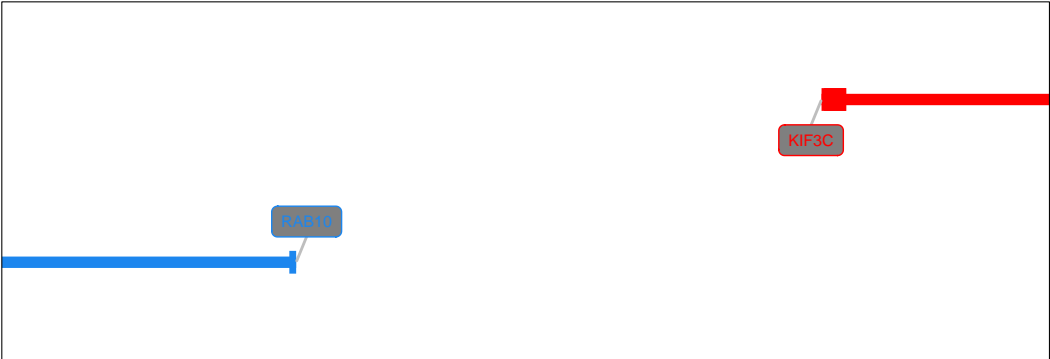
Coverage
(Norm. ATAC Signal Range (0–1.57) by ReadsInTSS)



L4.5.TBX15

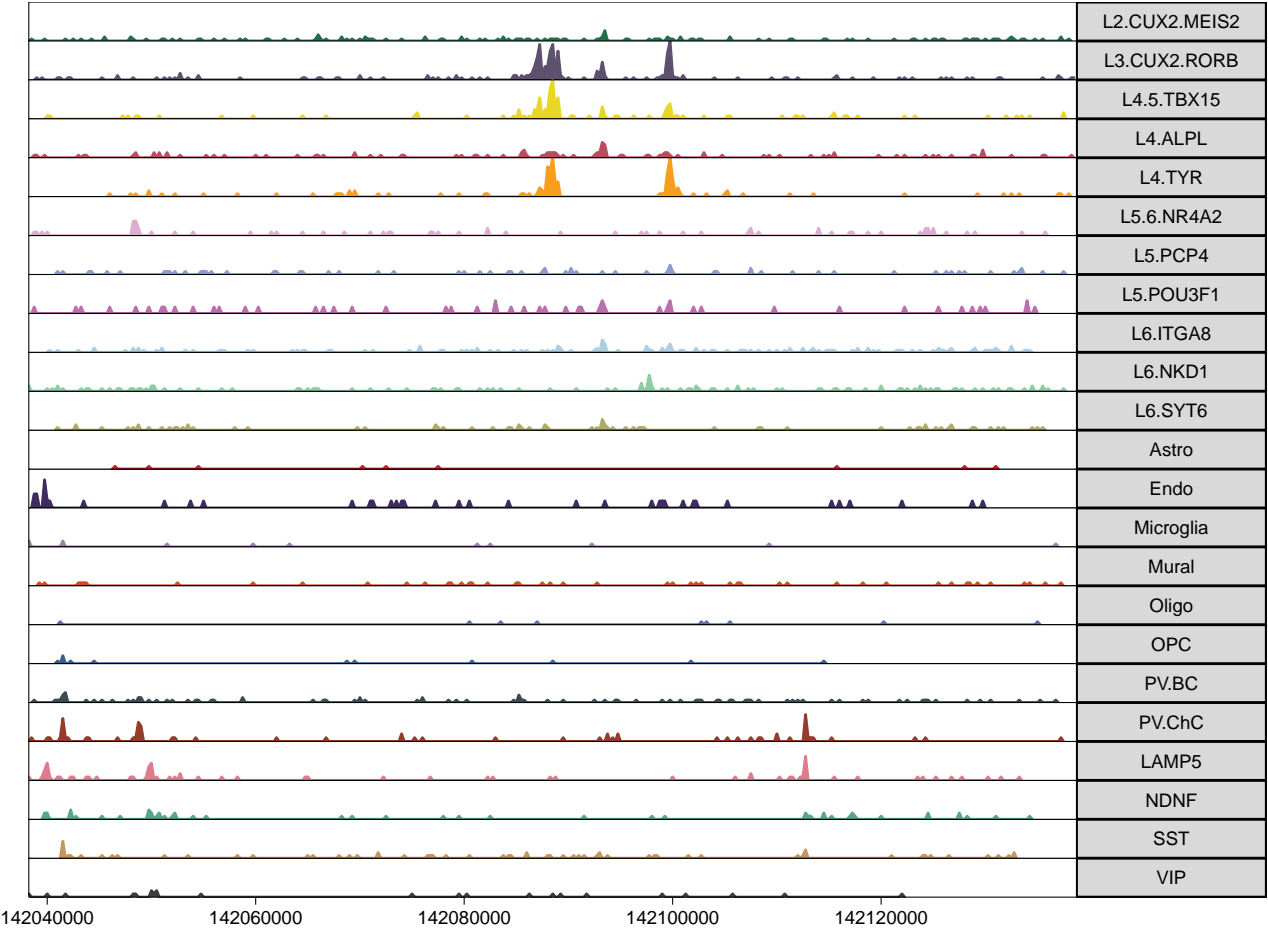


Peaks

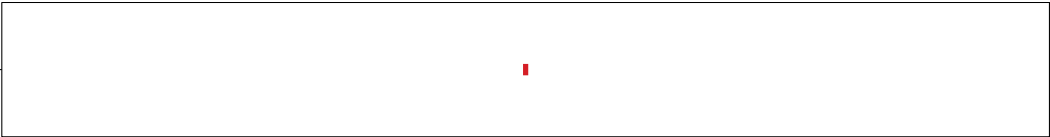


Genes

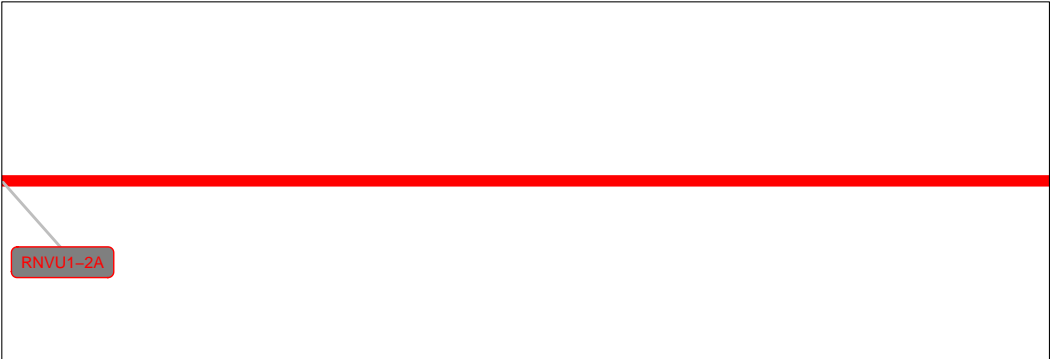
Coverage
(Norm. ATAC Signal Range (0–0.45) by ReadsInTSS)



L4.5.TBX15

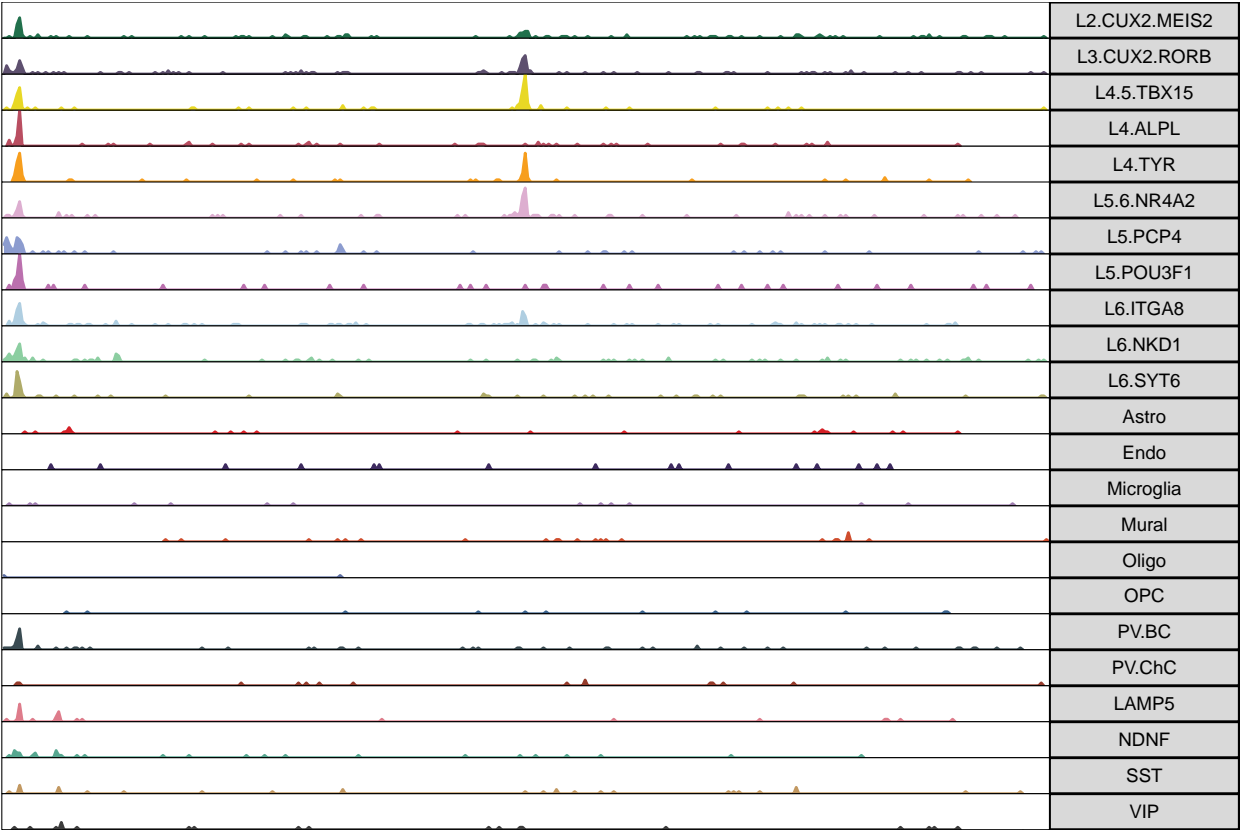


Peaks



Genes

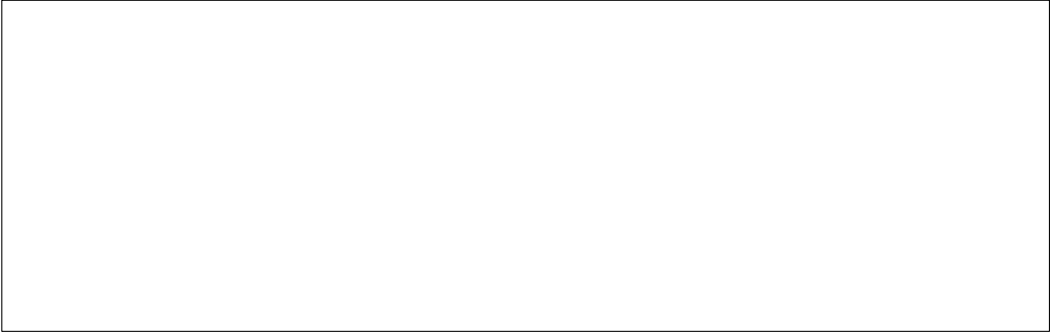
Coverage
(Norm. ATAC Signal Range (0–0.55) by ReadsInTSS)



L4.5.TBX15



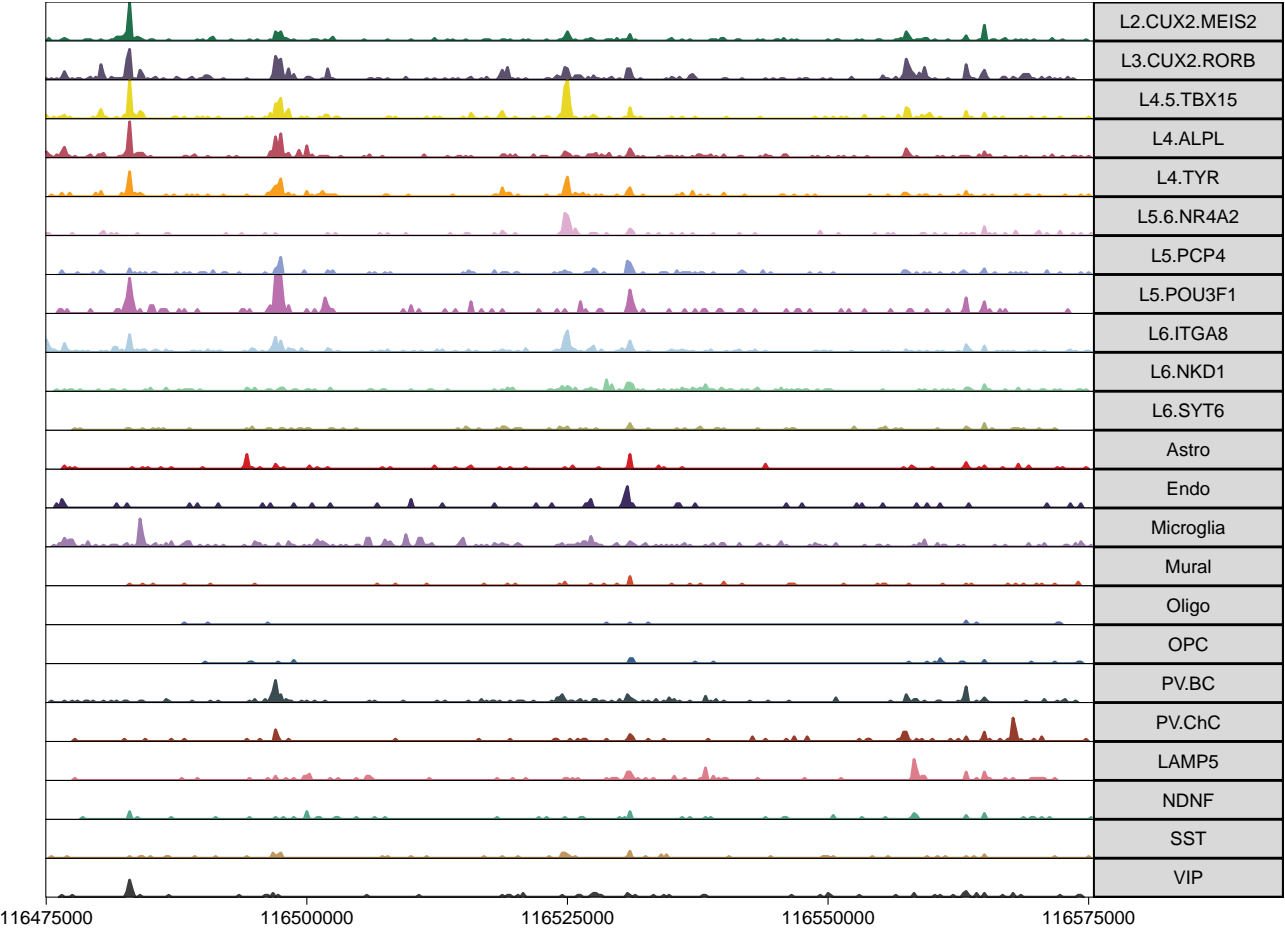
Peaks



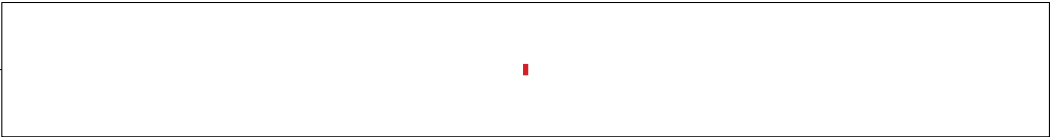
GeneTrack

chr1:116474960–116575461 Marker Peak Tracks for: L4.5.TBX15

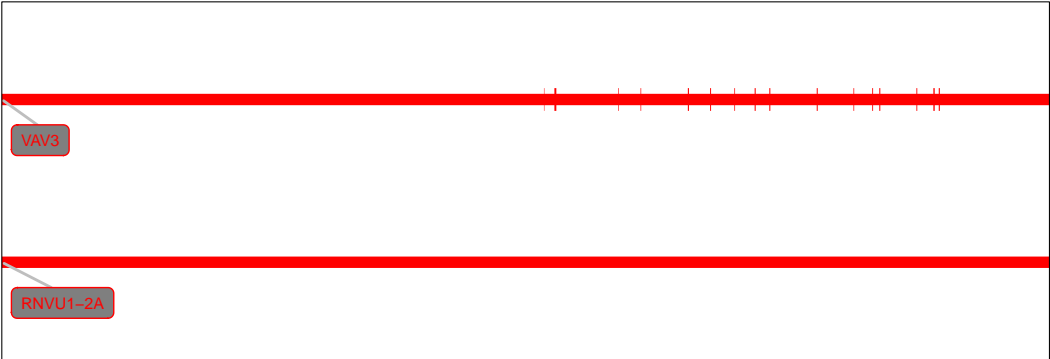
Coverage
(Norm. ATAC Signal Range (0–0.73) by ReadsInTSS)



L4.5.TBX15

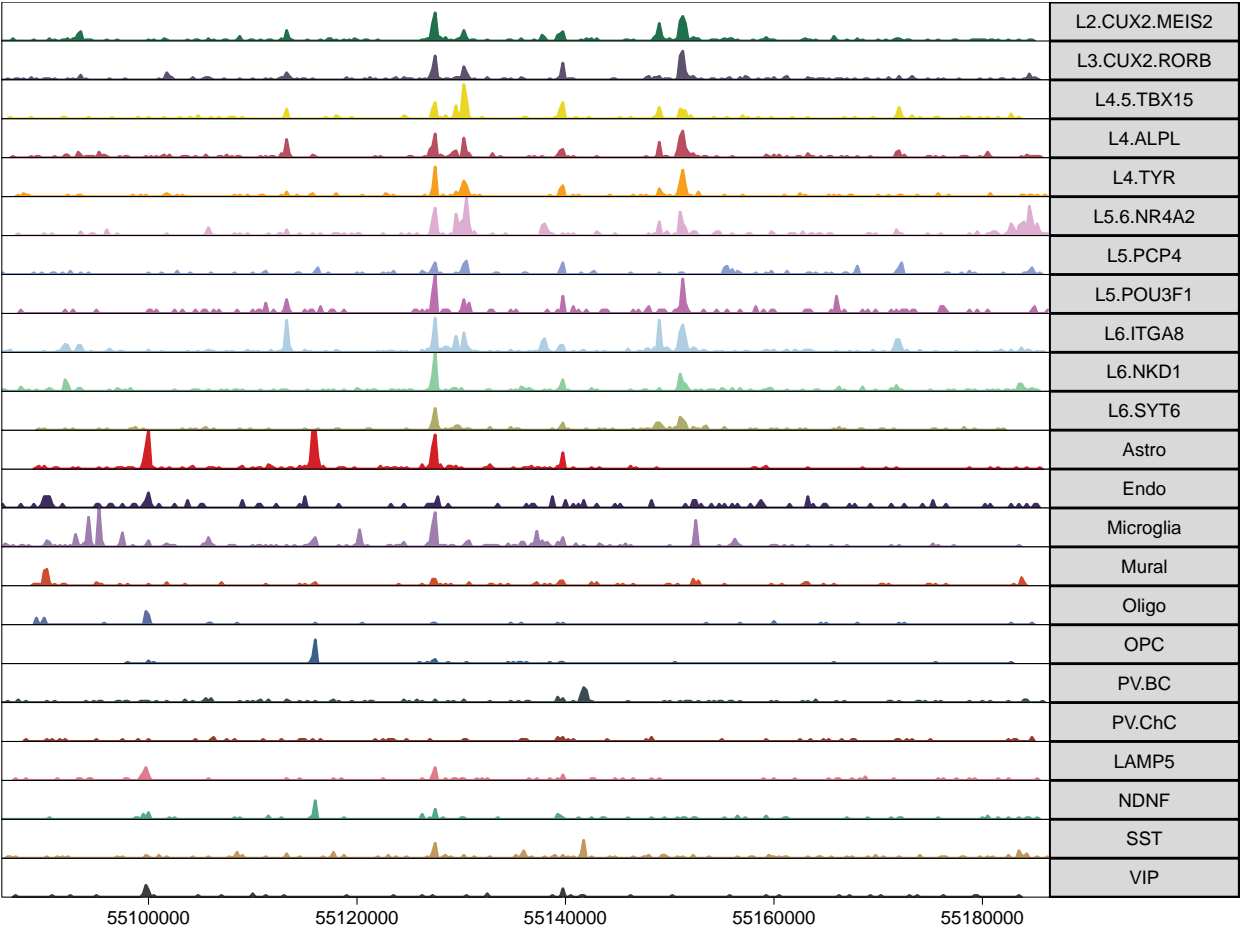


Peaks

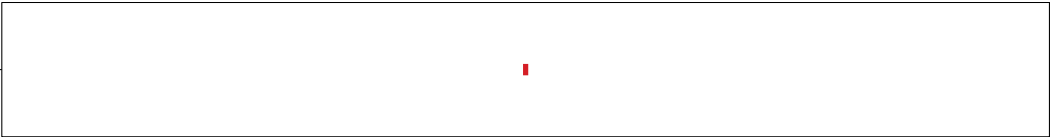


Genes

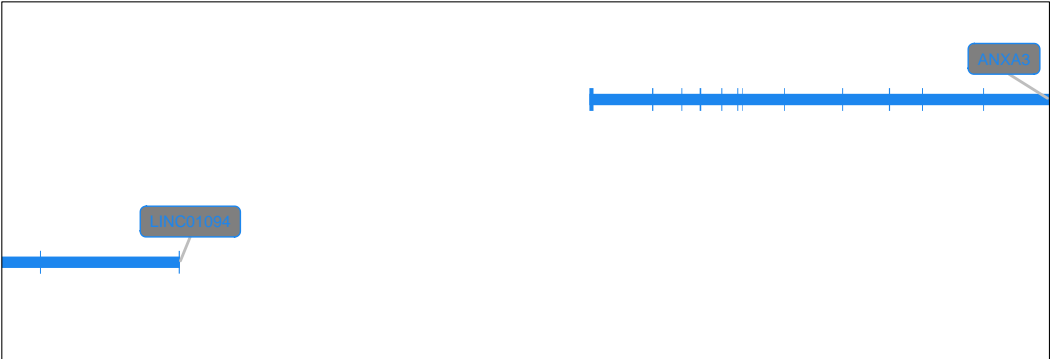
Coverage
(Norm. ATAC Signal Range (0–0.83) by ReadsInTSS)



L4.5.TBX15

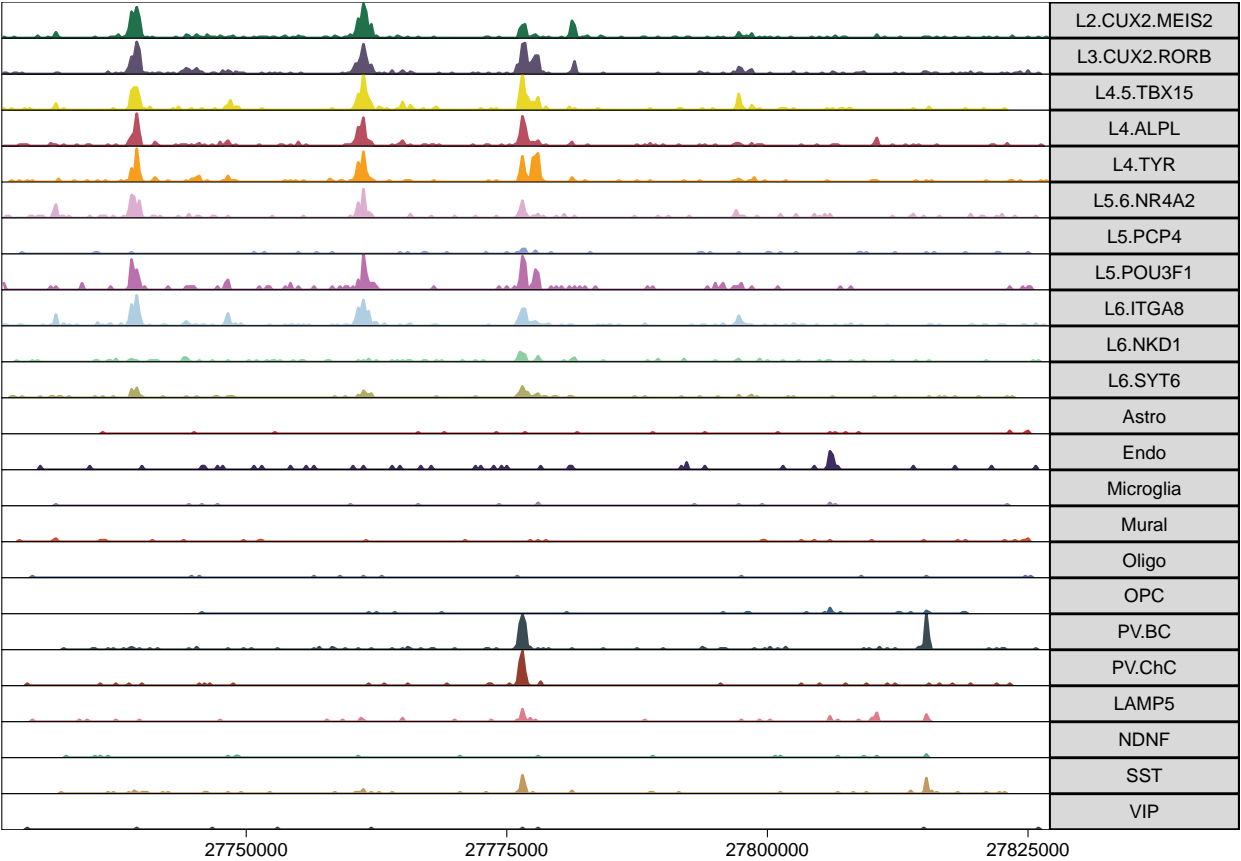


Peaks

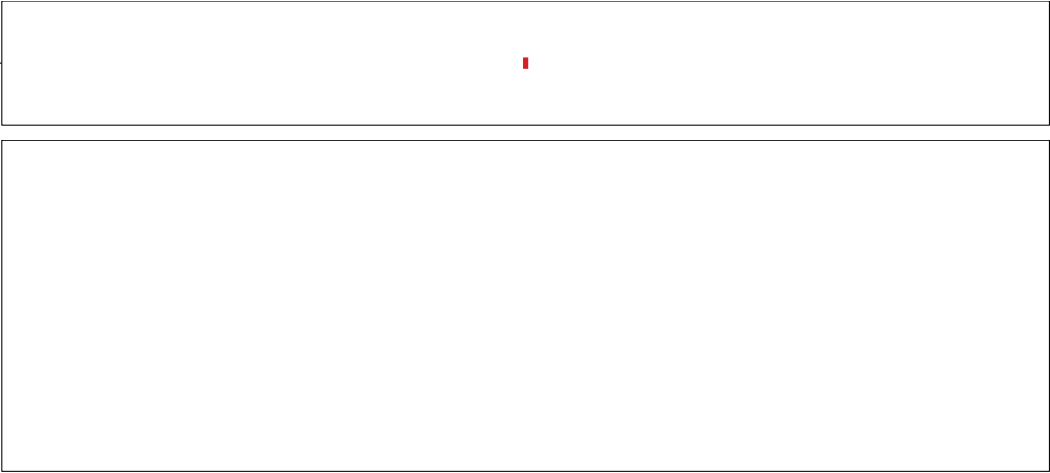


Genes

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



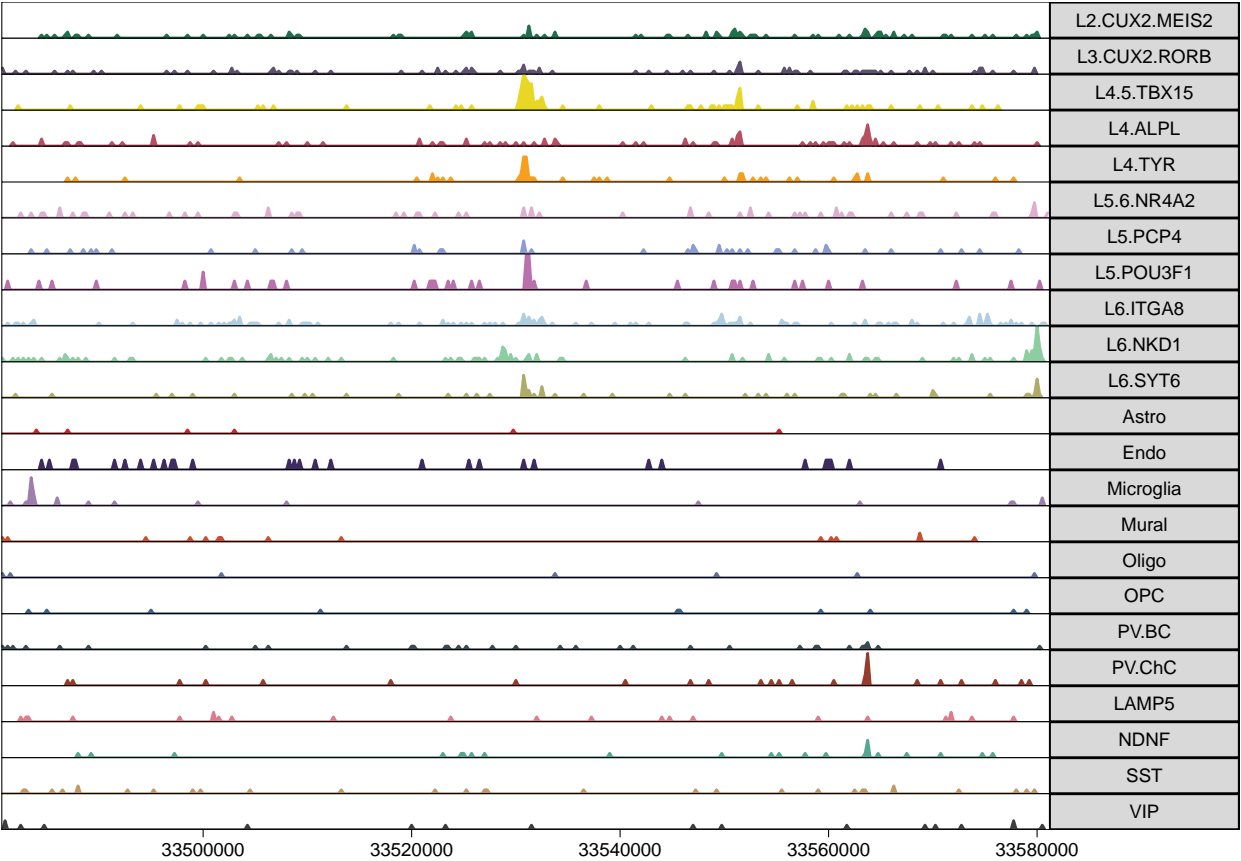
L4.5.TBX15



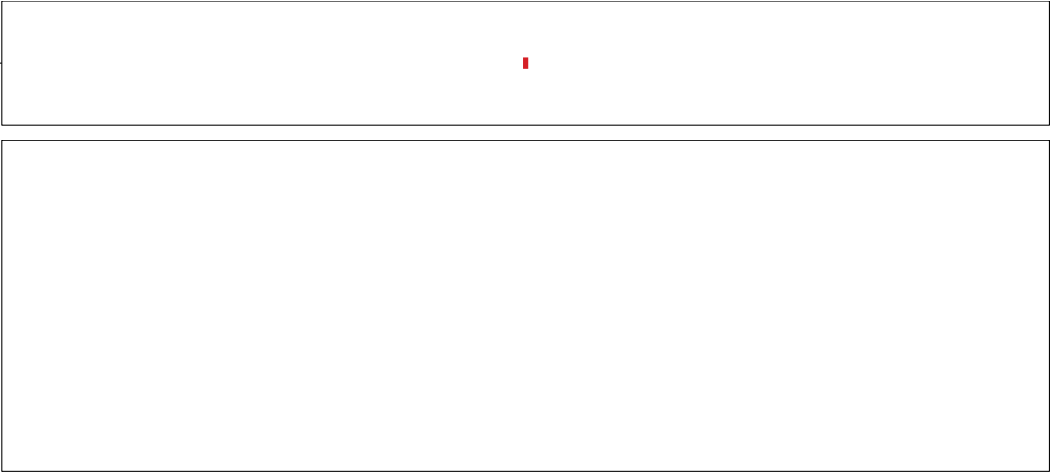
Peaks

GeneTrack

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



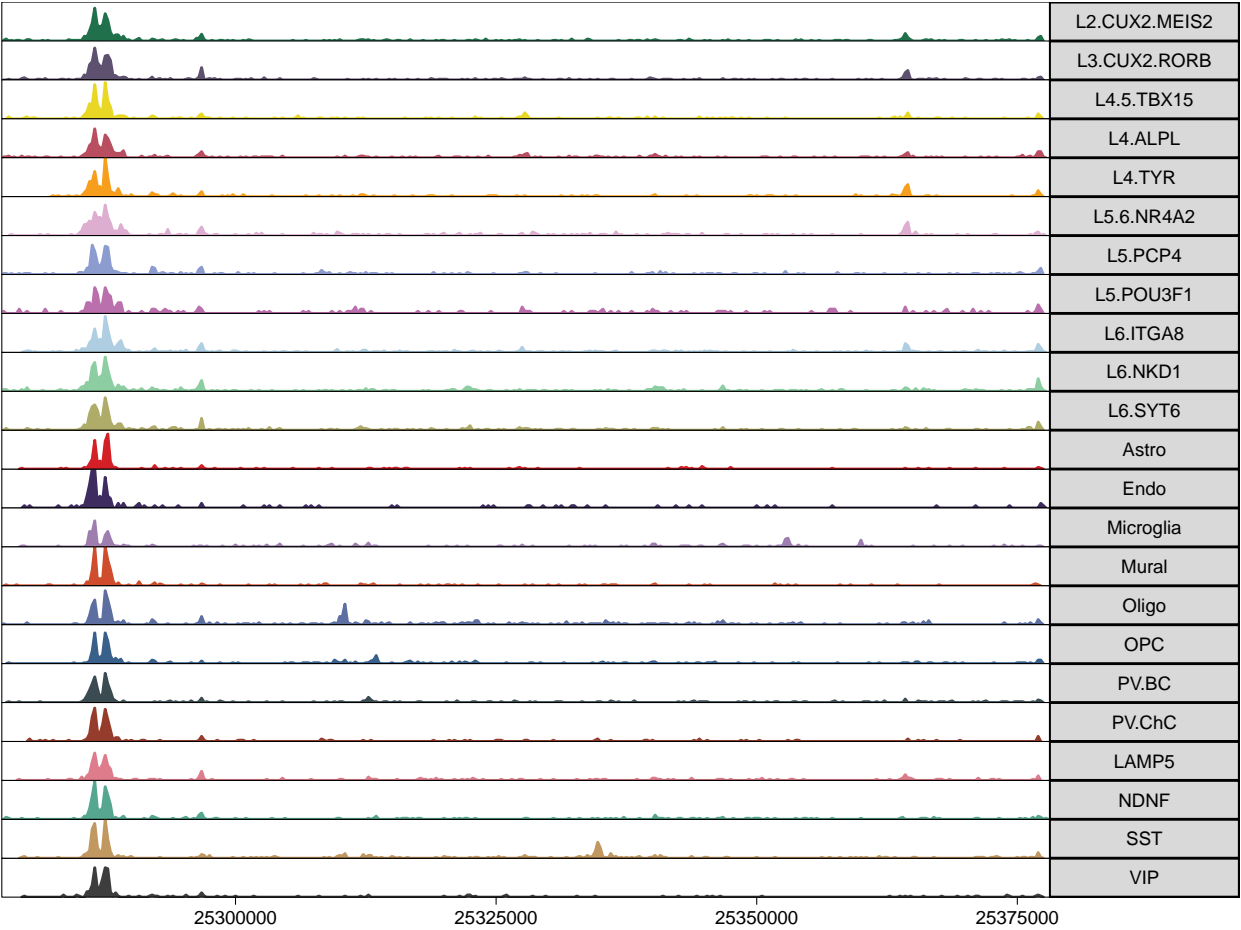
L4.5.TBX15



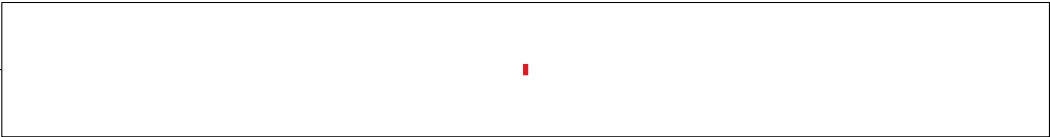
Peaks

GeneTrack

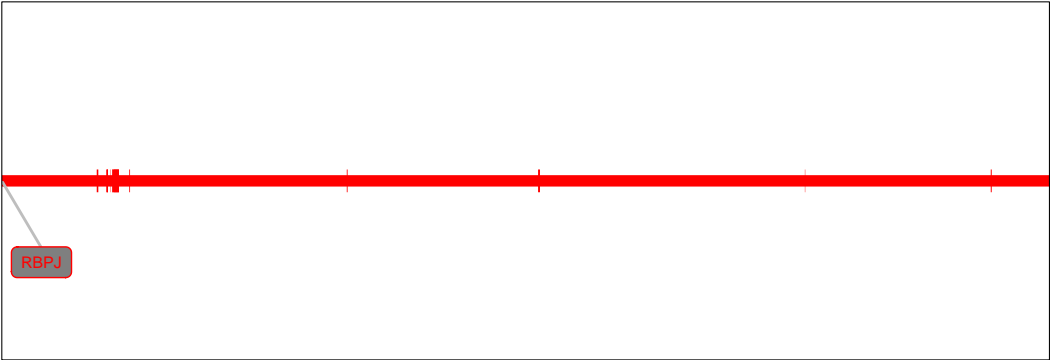
Coverage
(Norm. ATAC Signal Range (0-1.33) by ReadsInTSS)



L4.5.TBX15



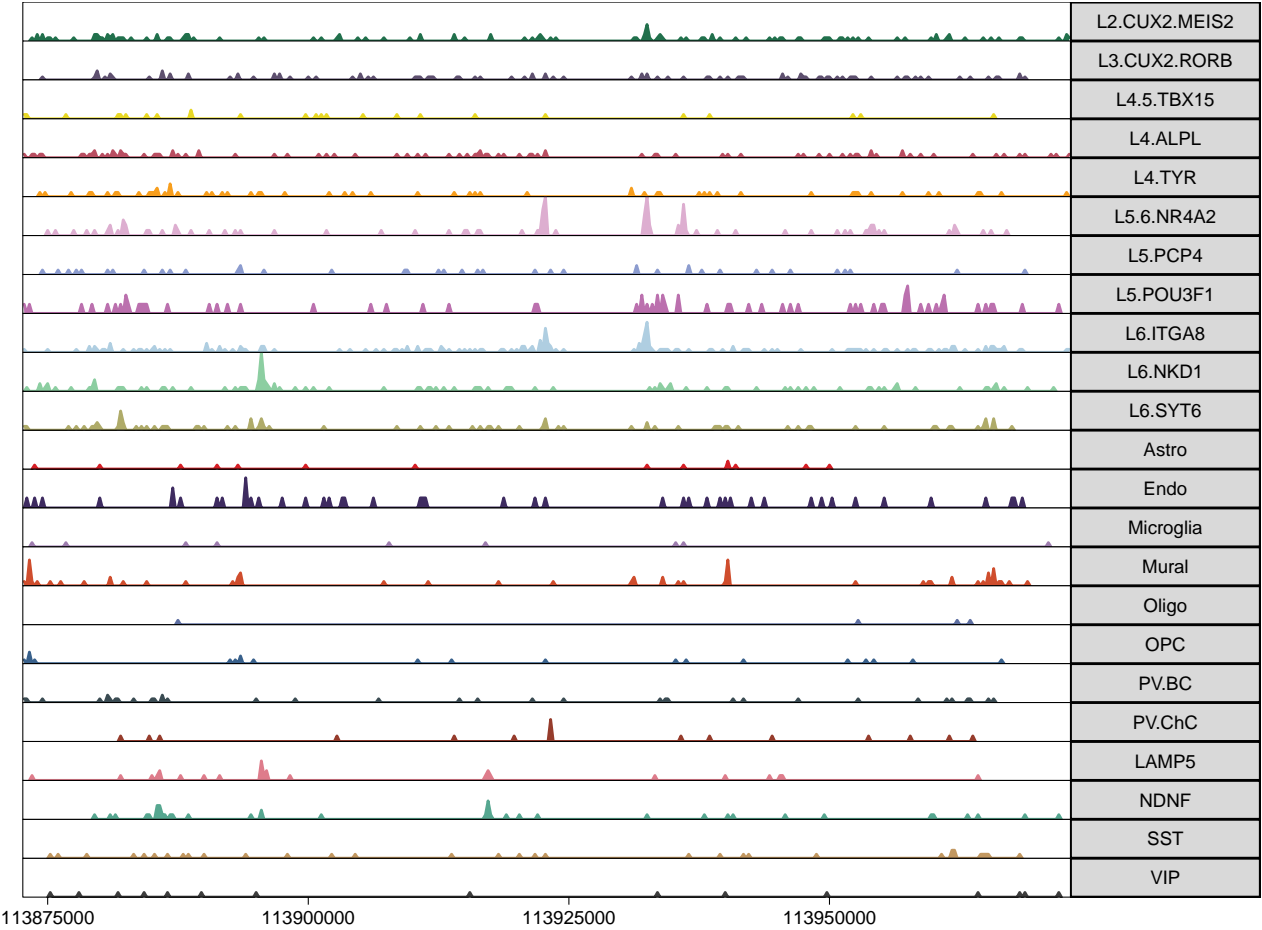
Peaks



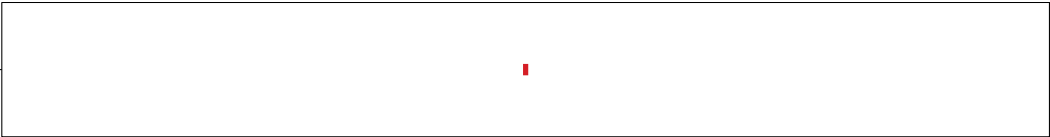
Genes

chr4:113872615–113973116 Marker Peak Tracks for: L4.5.TBX15

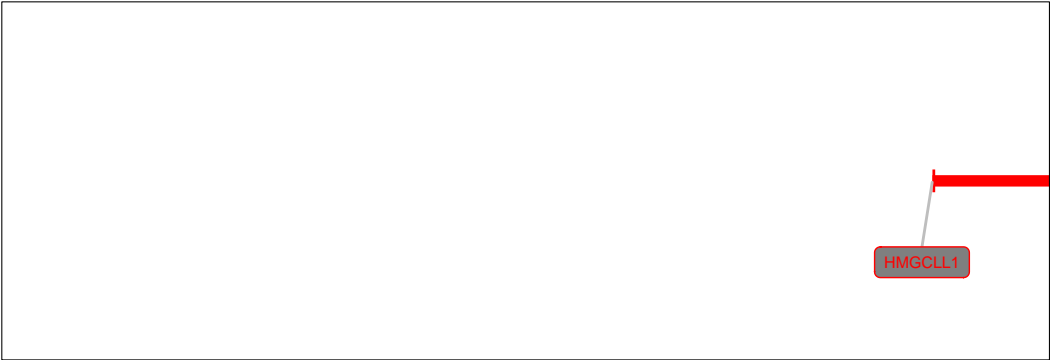
Coverage
(Norm. ATAC Signal Range (0–0.31) by ReadsInTSS)



L4.5.TBX15



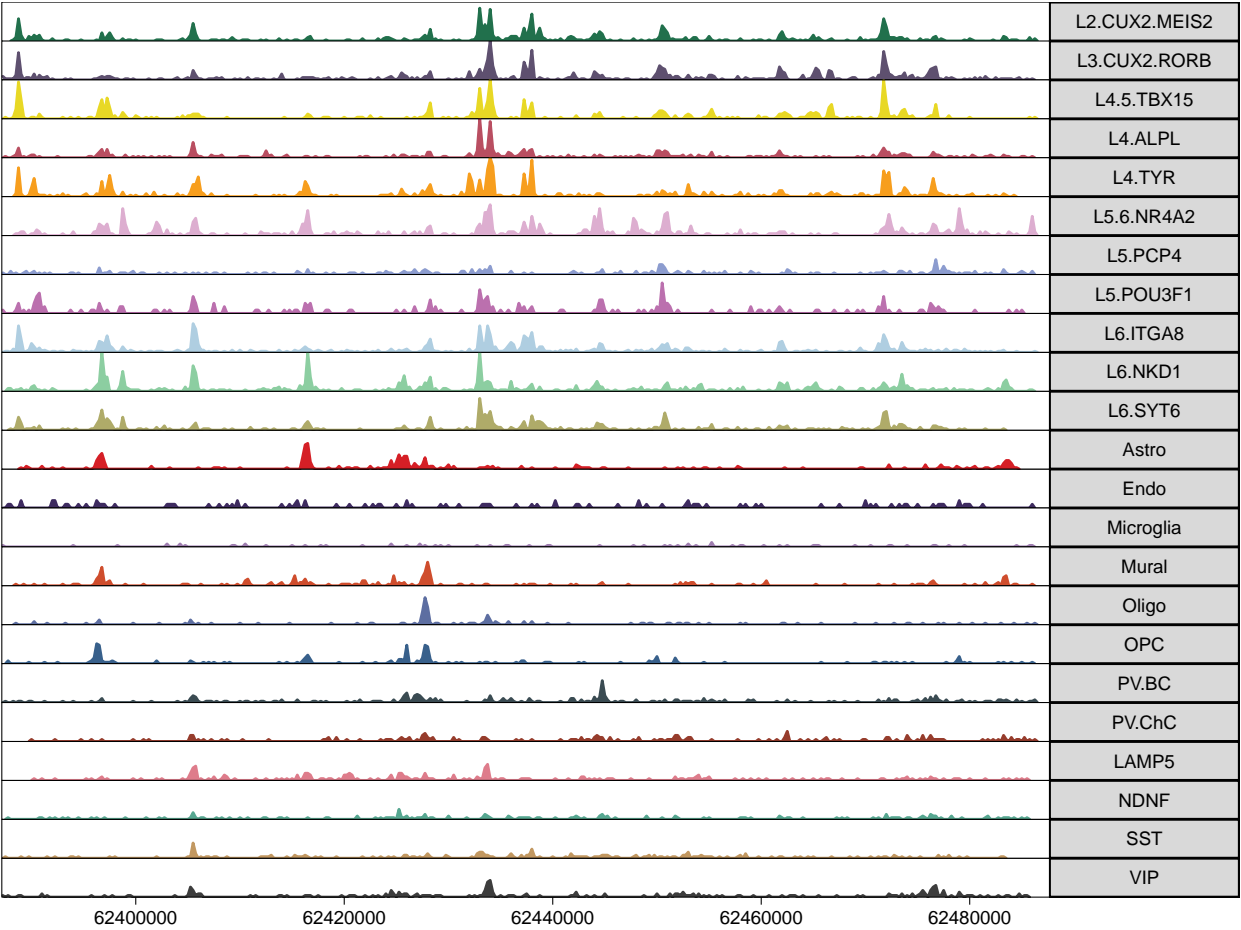
Peaks



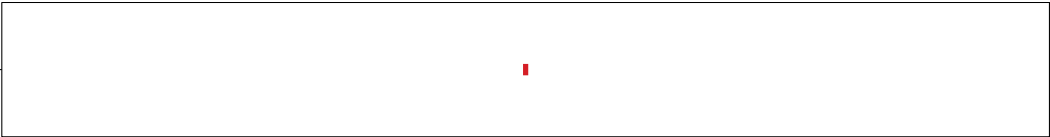
Genes

chr11:62387149–62487650 Marker Peak Tracks for: L4.5.TBX15

Coverage
(Norm. ATAC Signal Range (0–0.84) by ReadsInTSS)



L4.5.TBX15

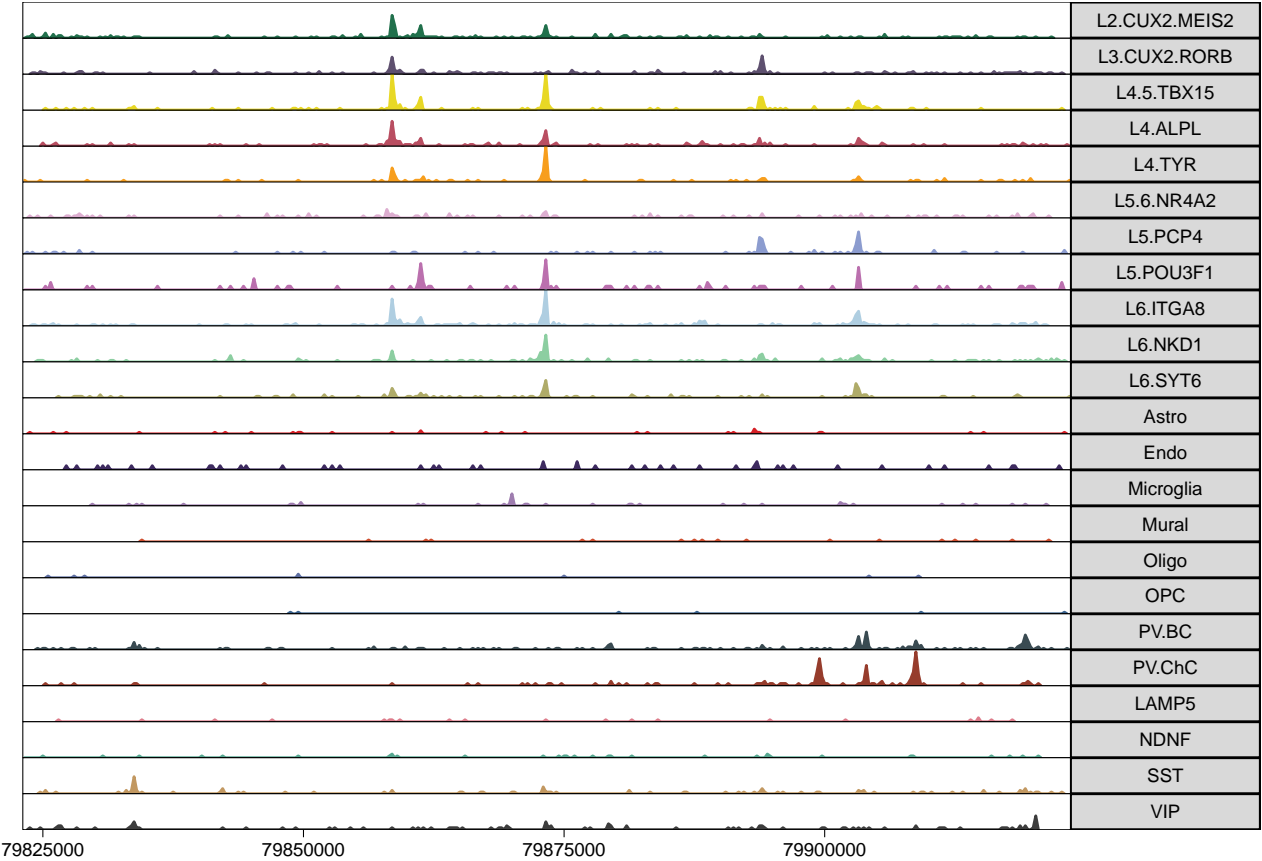


Peaks

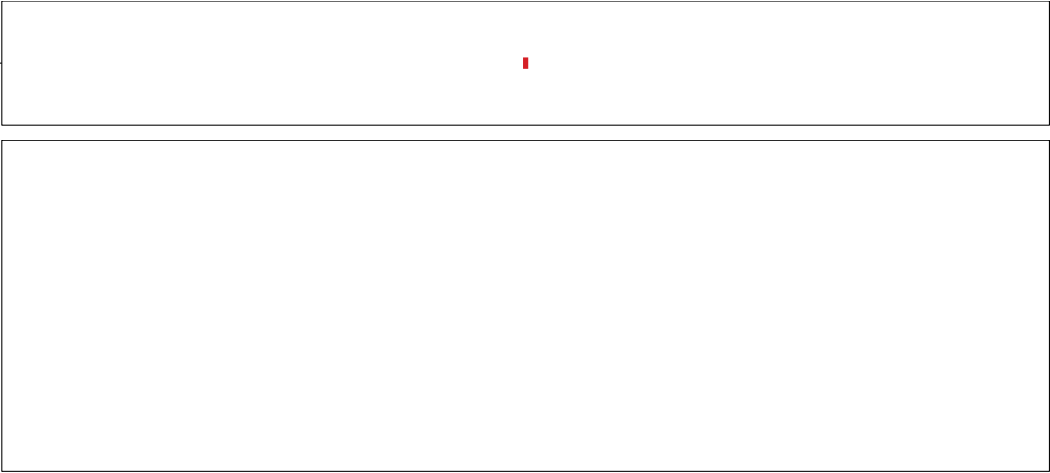


Genes

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



L4.5.TBX15

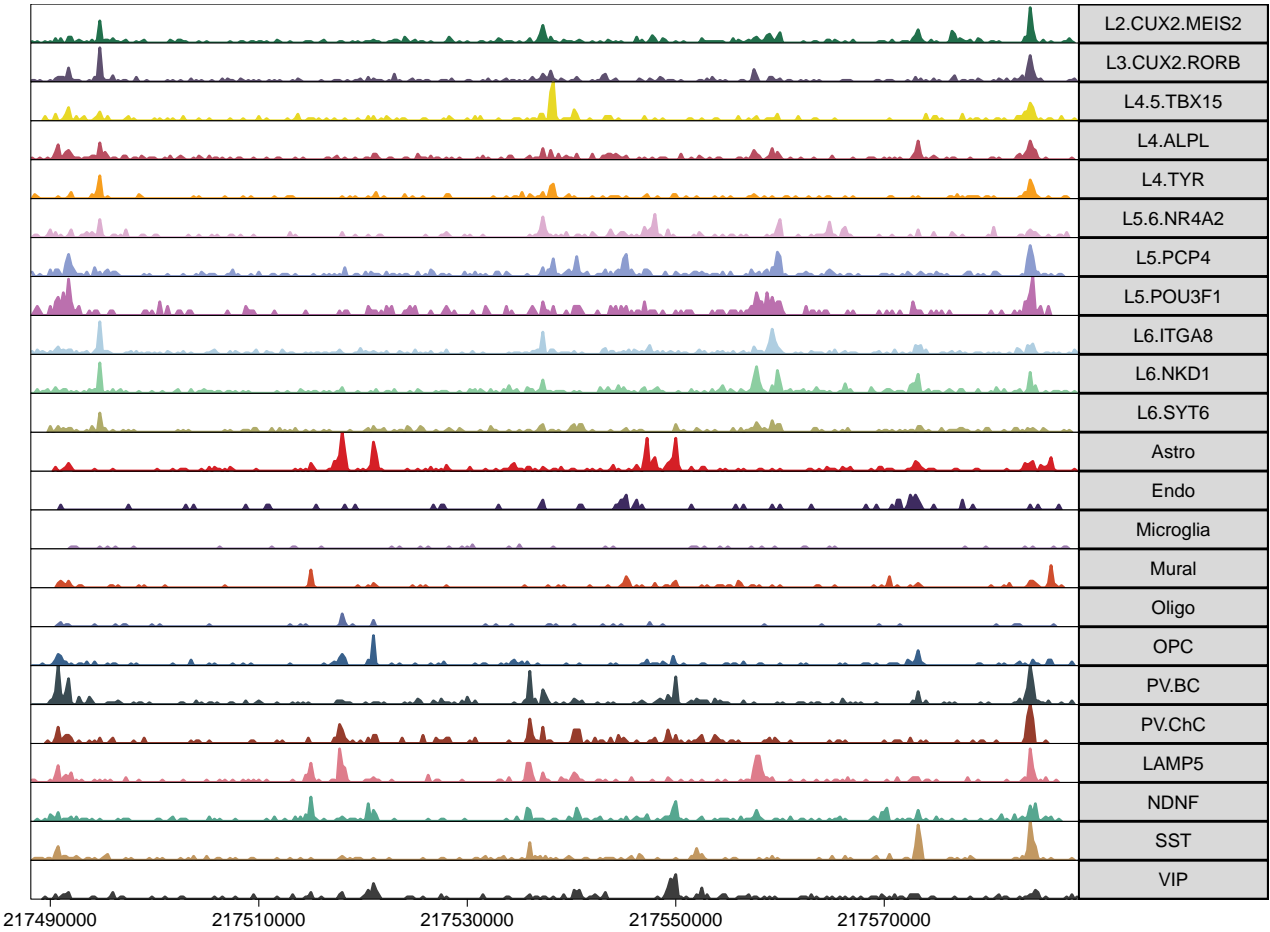


Peaks

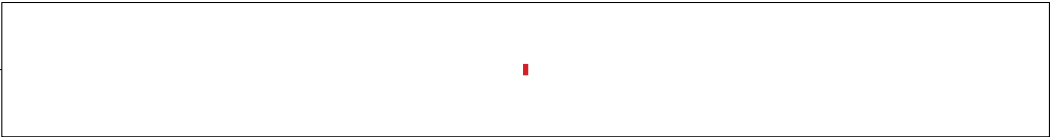
GeneTrack

chr1:217488128–217588629 Marker Peak Tracks for: L4.5.TBX15

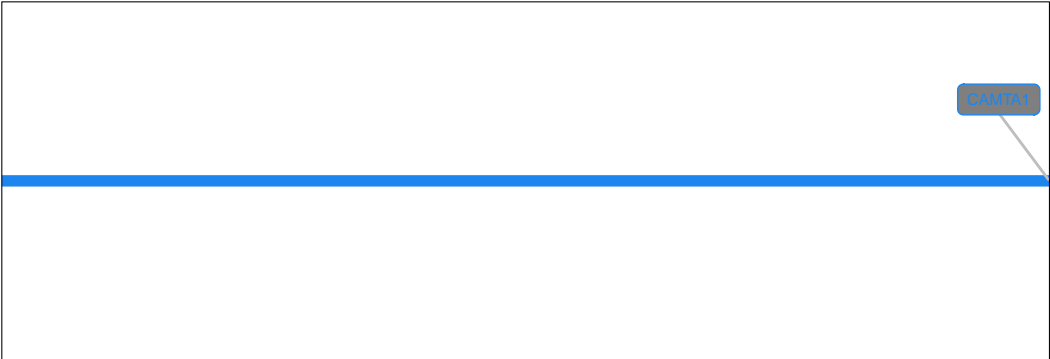
Coverage
(Norm. ATAC Signal Range (0–0.63) by ReadsInTSS)



L4.5.TBX15

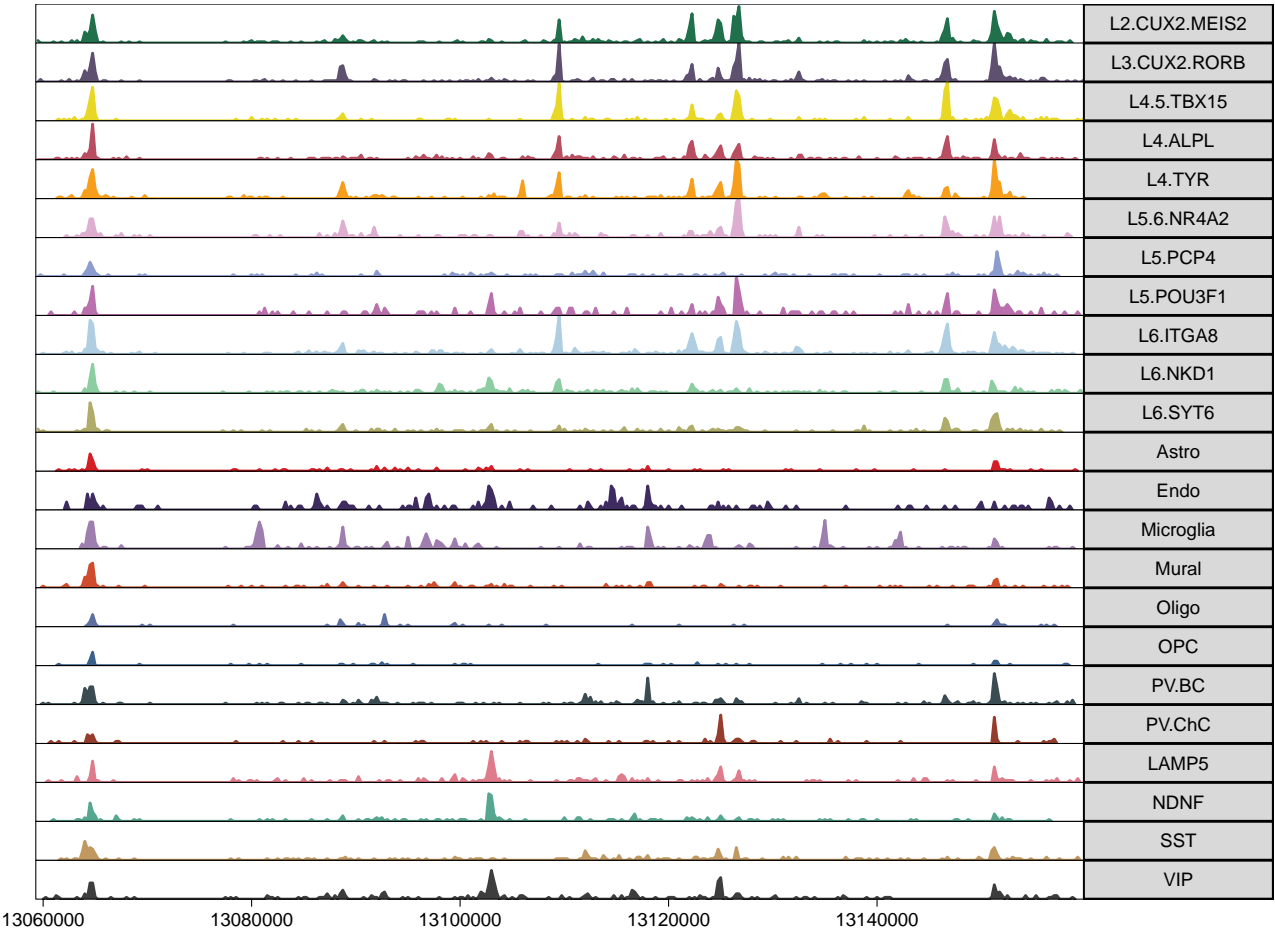


Peaks

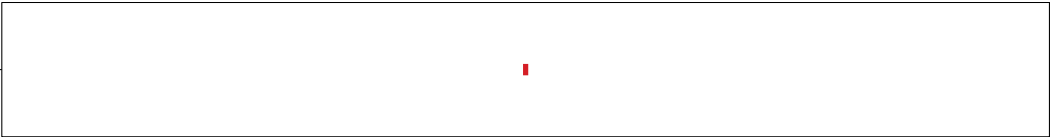


Genes

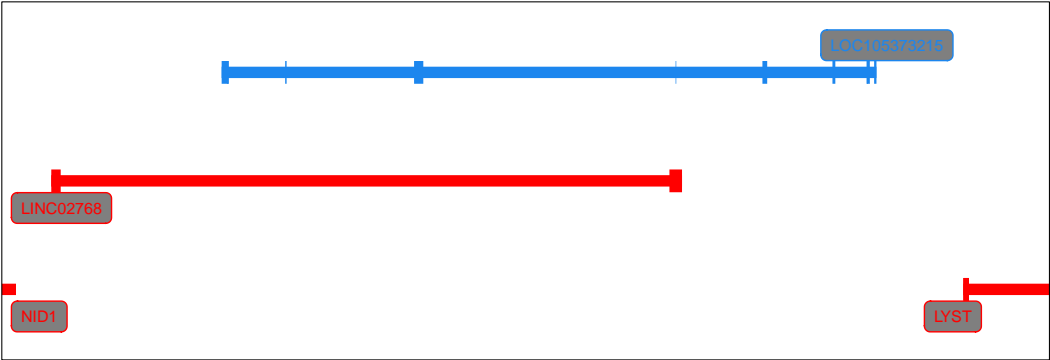
Coverage
(Norm. ATAC Signal Range (0-0.78) by ReadsInTSS)



L4.5.TBX15

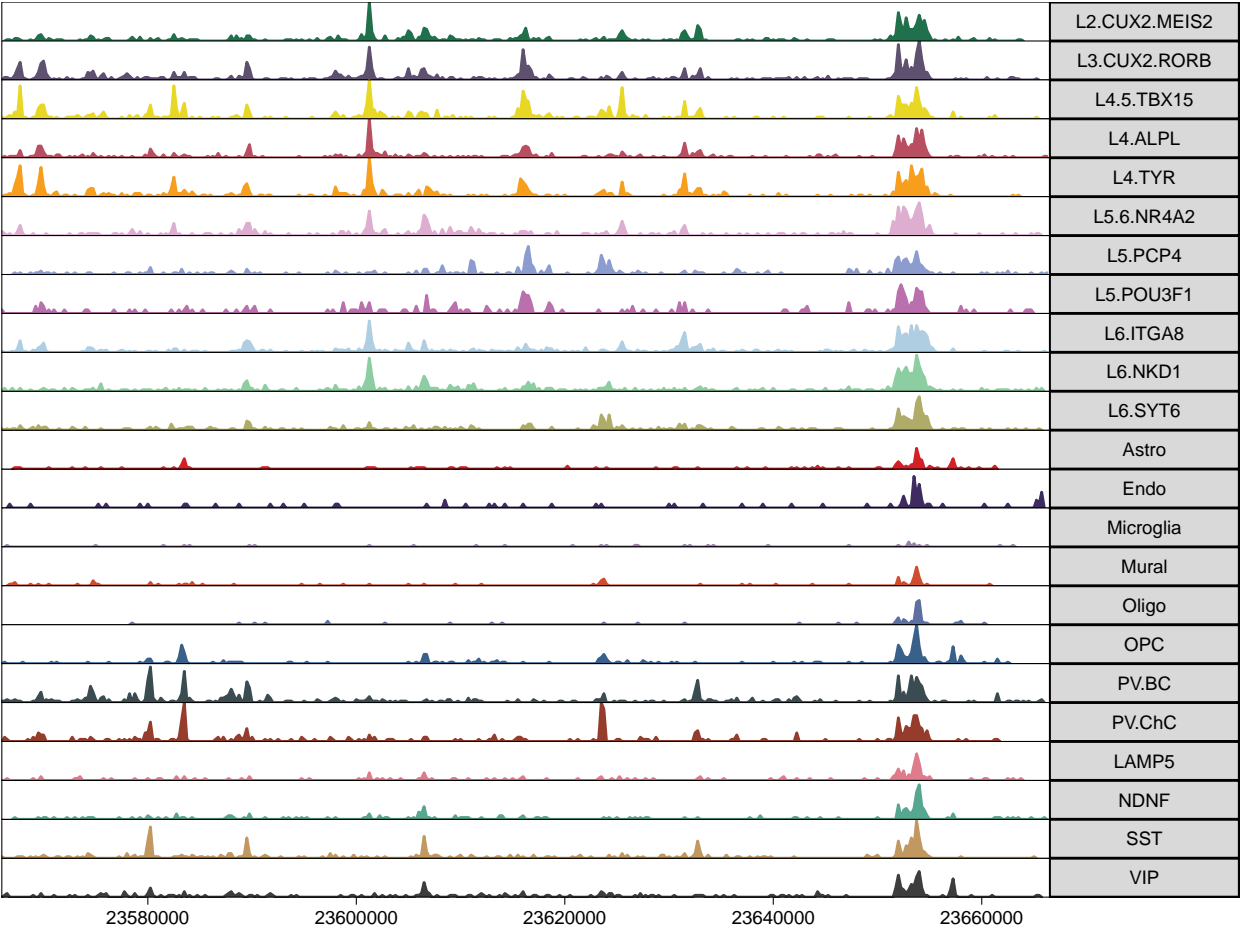


Peaks

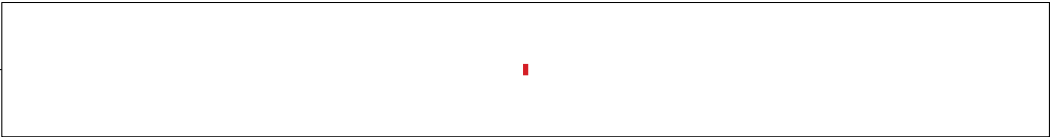


Genes

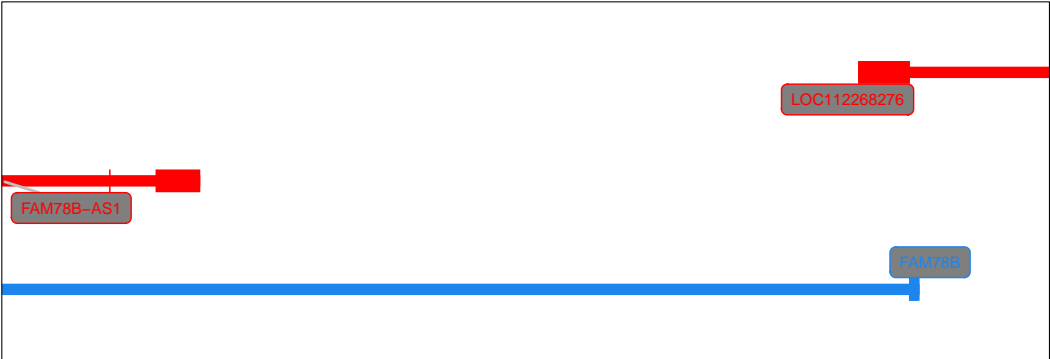
Coverage
(Norm. ATAC Signal Range (0–0.79) by ReadsInTSS)



L4.5.TBX15

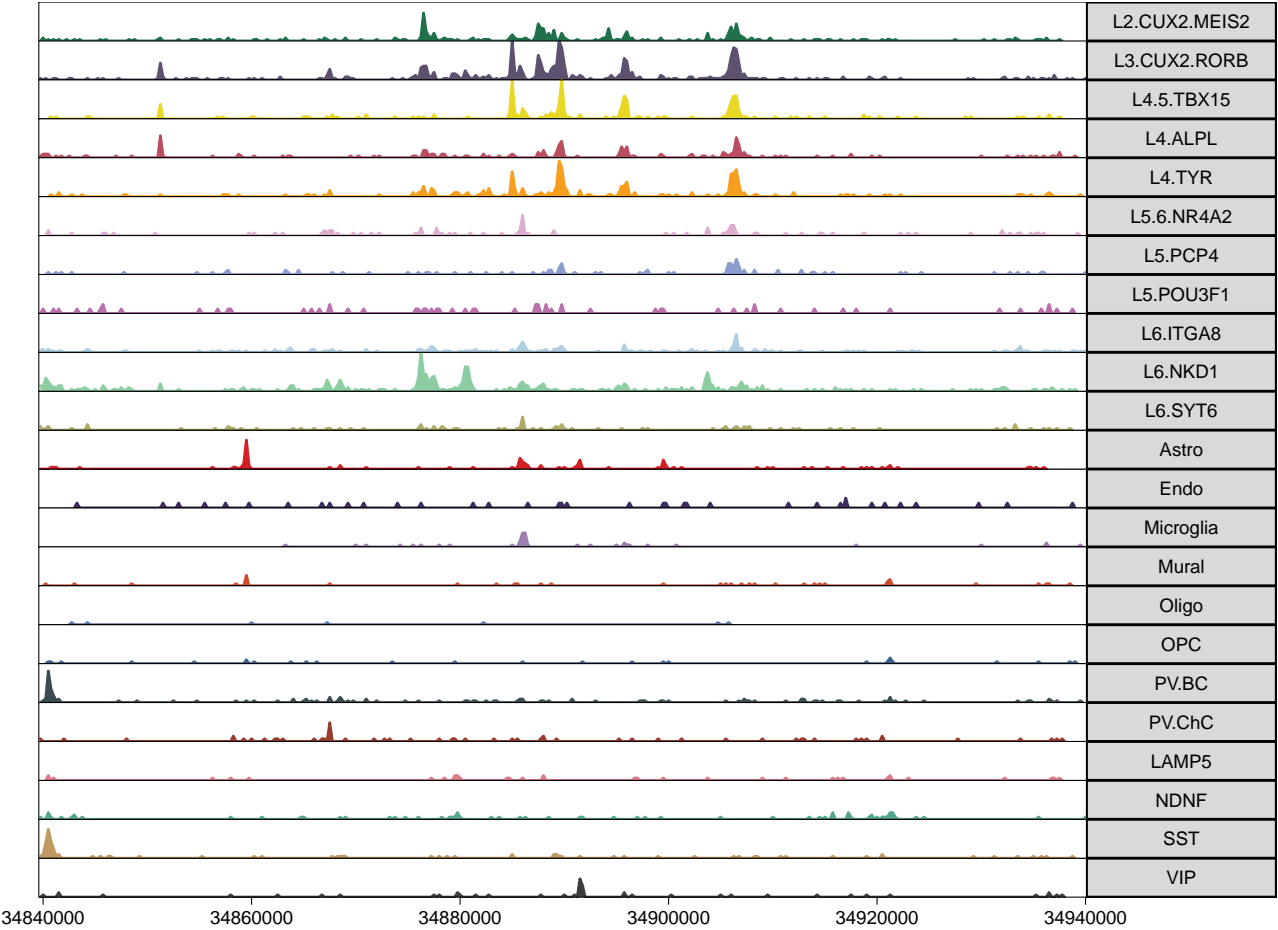


Peaks

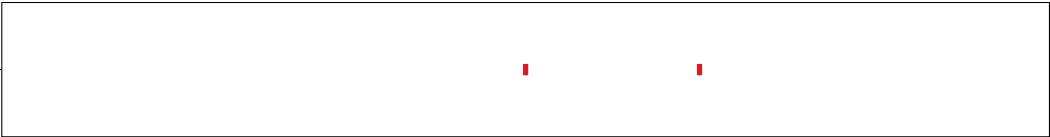


Genes

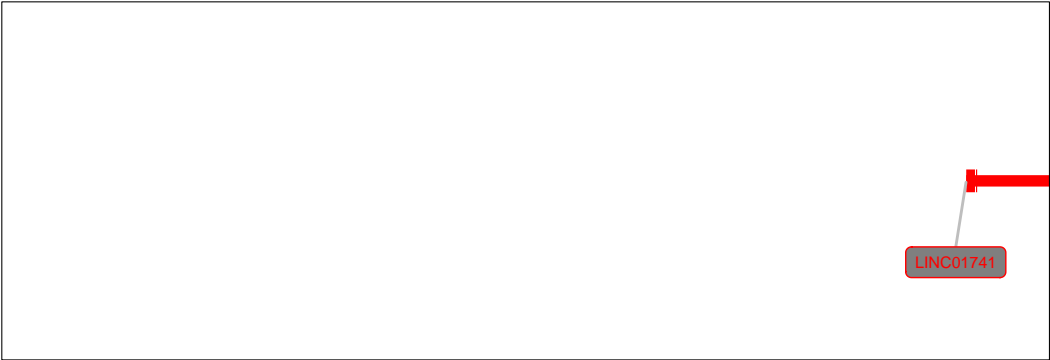
Coverage
(Norm. ATAC Signal Range (0-0.64) by ReadsInTSS)



L4.5.TBX15

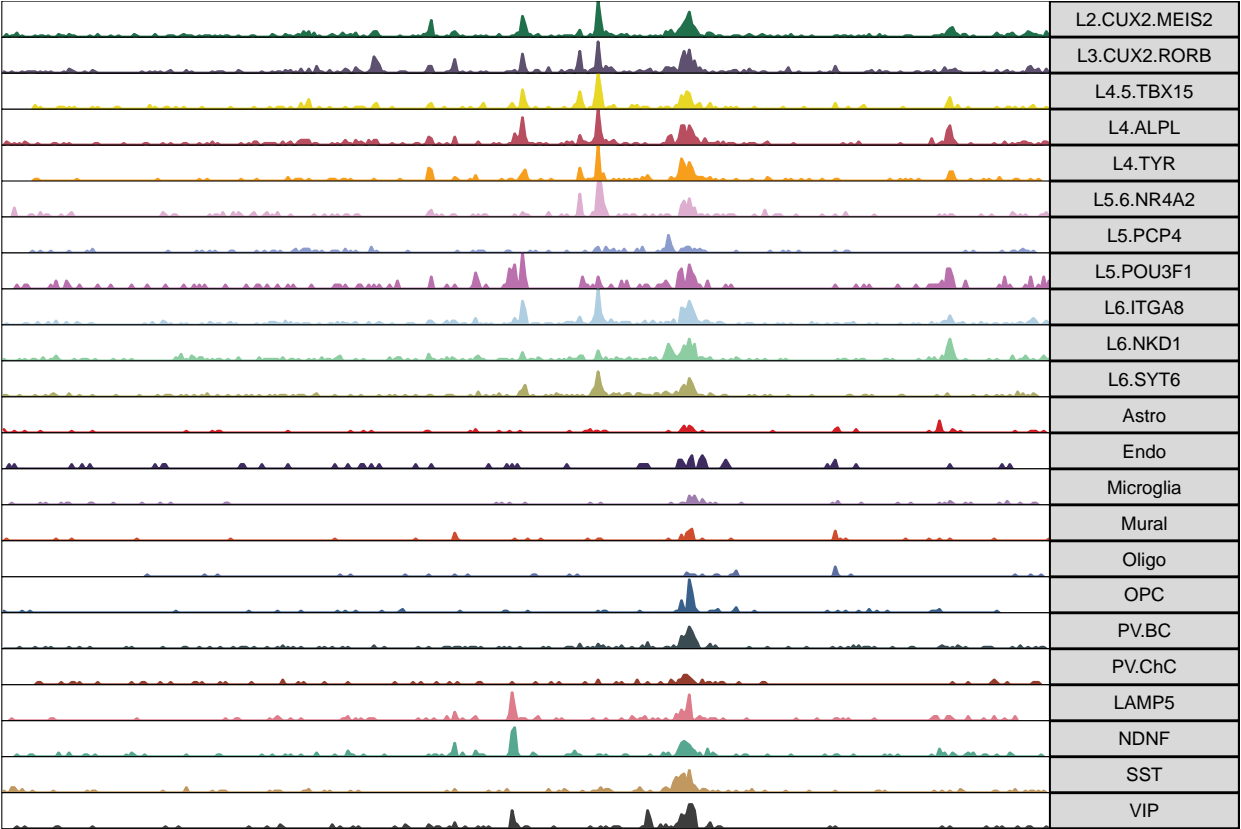


Peaks

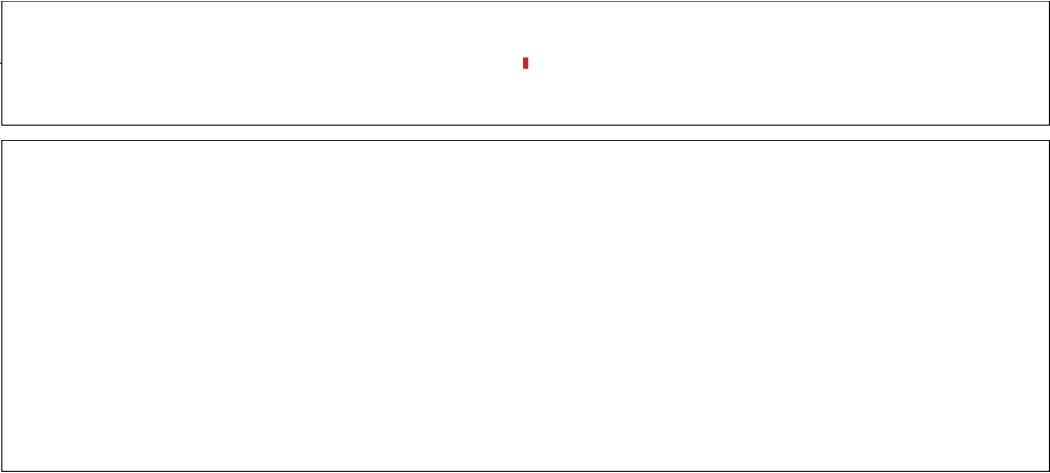


Genes

Coverage
(Norm. ATAC Signal Range (0–0.66) by ReadsInTSS)



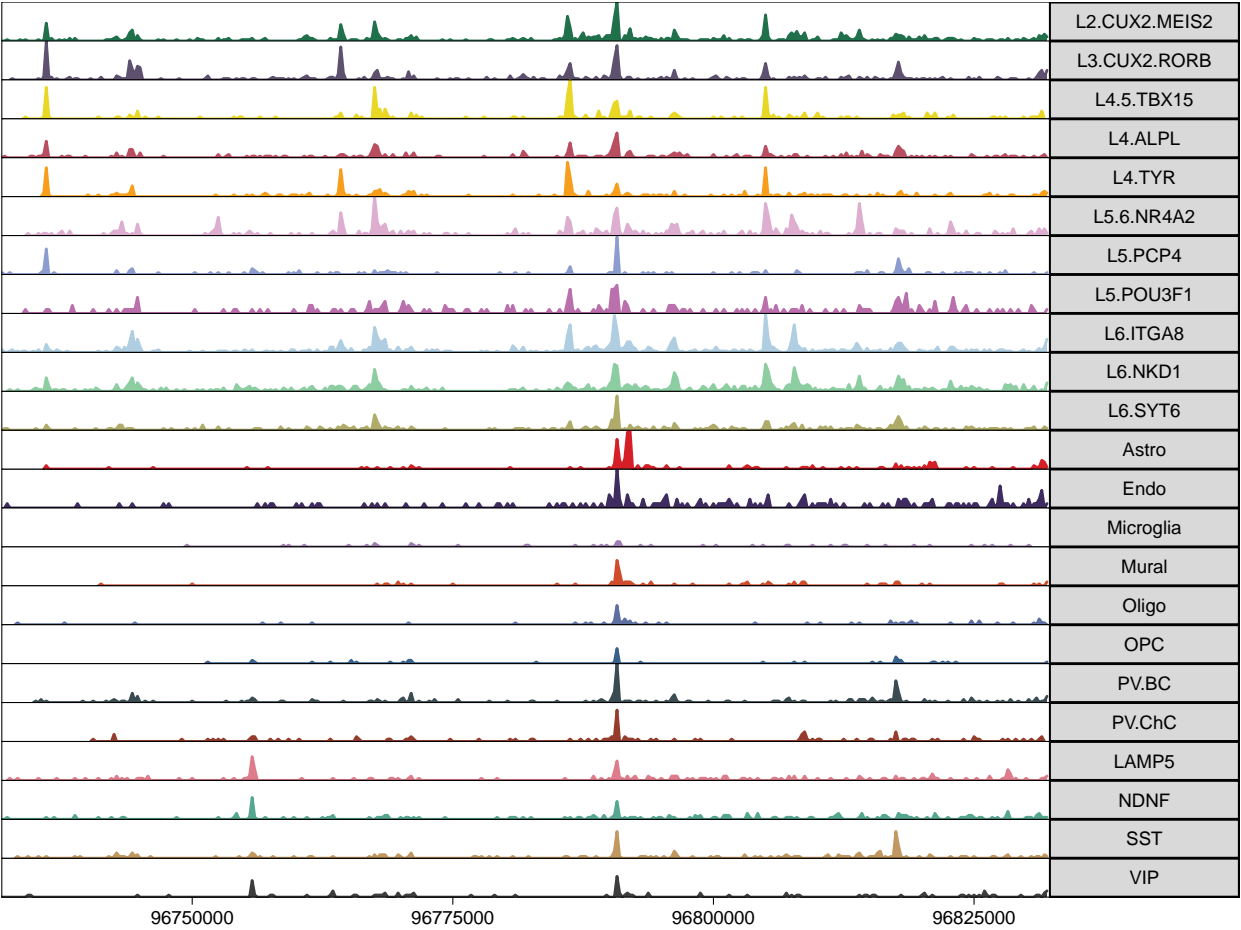
L4.5.TBX15



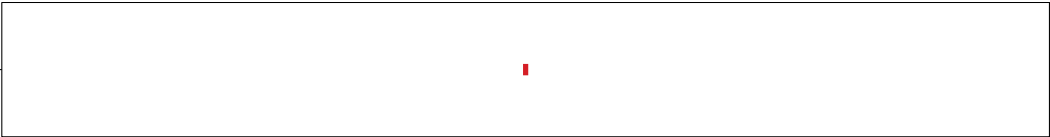
Peaks

GeneTrack

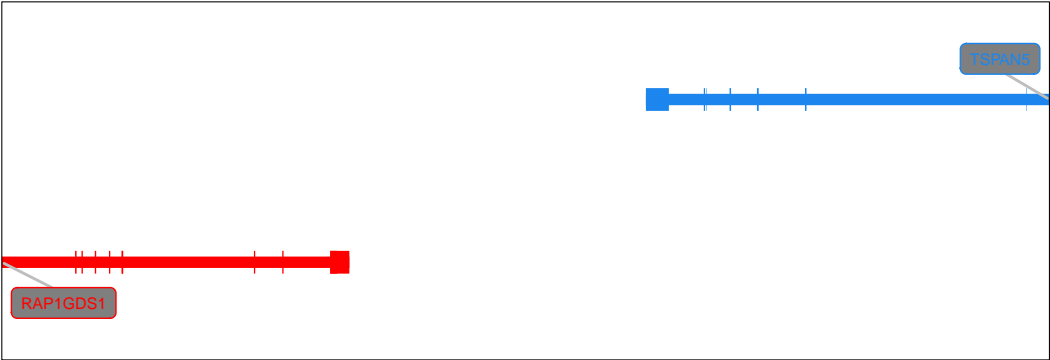
Coverage
(Norm. ATAC Signal Range (0-0.71) by ReadsInTSS)



L4.5.TBX15



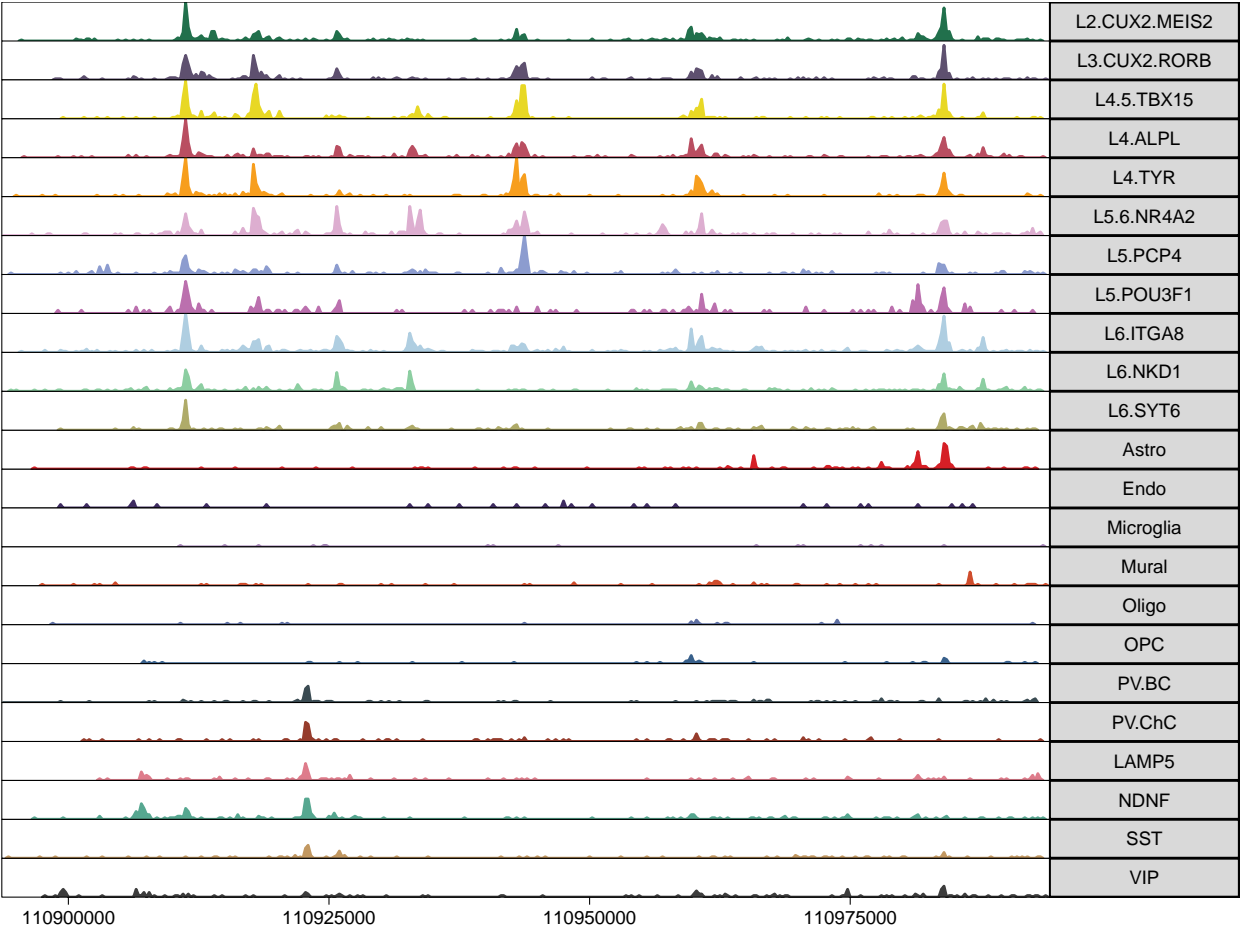
Peaks



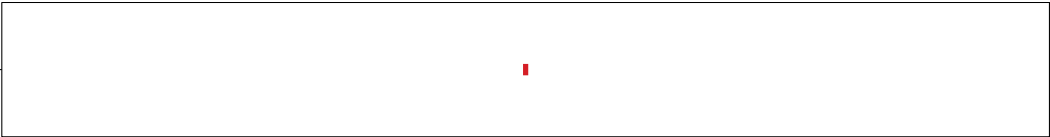
Genes

chr1:110893613–110994114 Marker Peak Tracks for: L4.5.TBX15

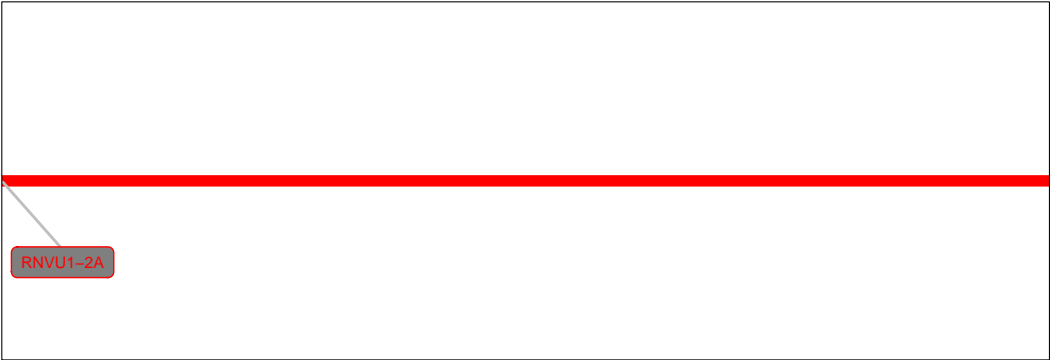
Coverage
(Norm. ATAC Signal Range (0–0.89) by ReadsInTSS)



L4.5.TBX15

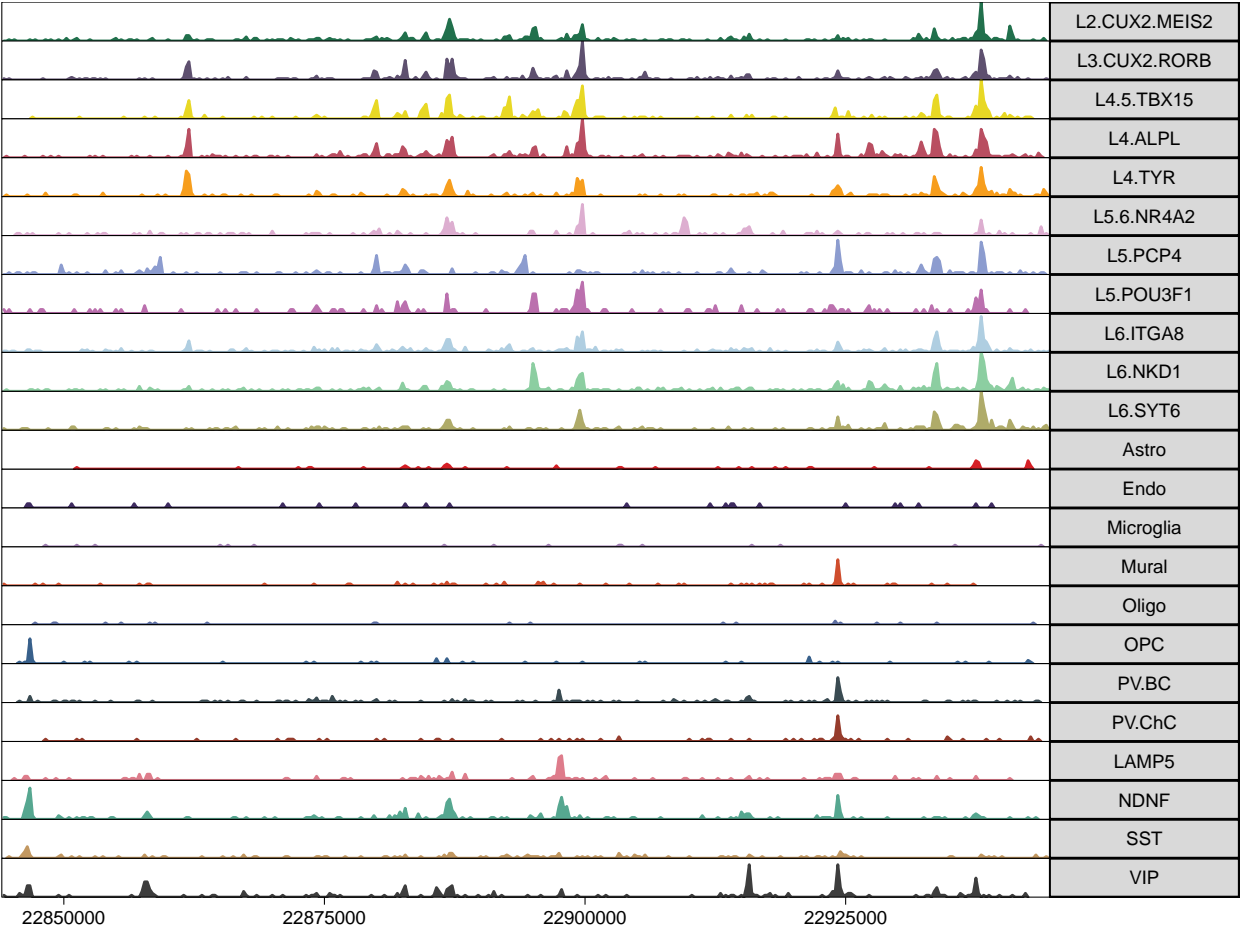


Peaks

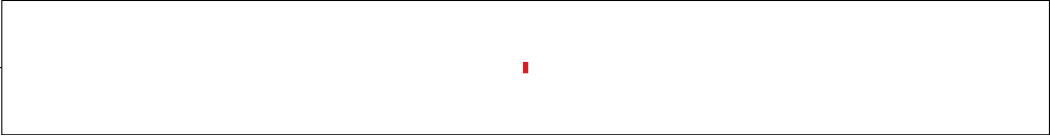


Genes

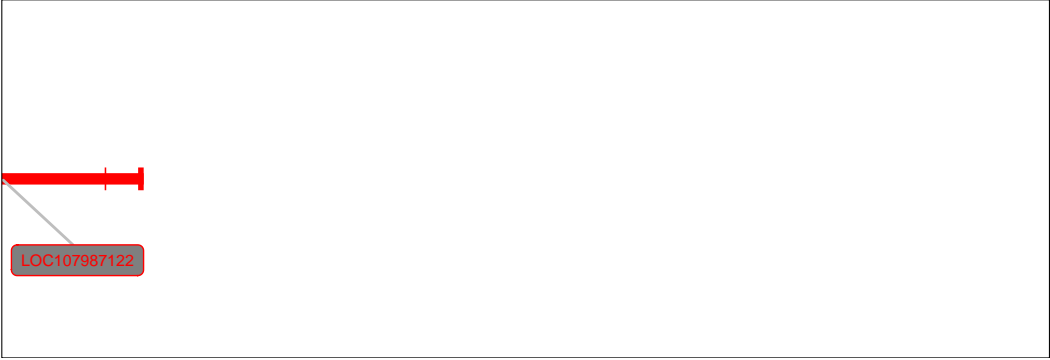
Coverage
(Norm. ATAC Signal Range (0–0.73) by ReadsInTSS)



L4.5.TBX15



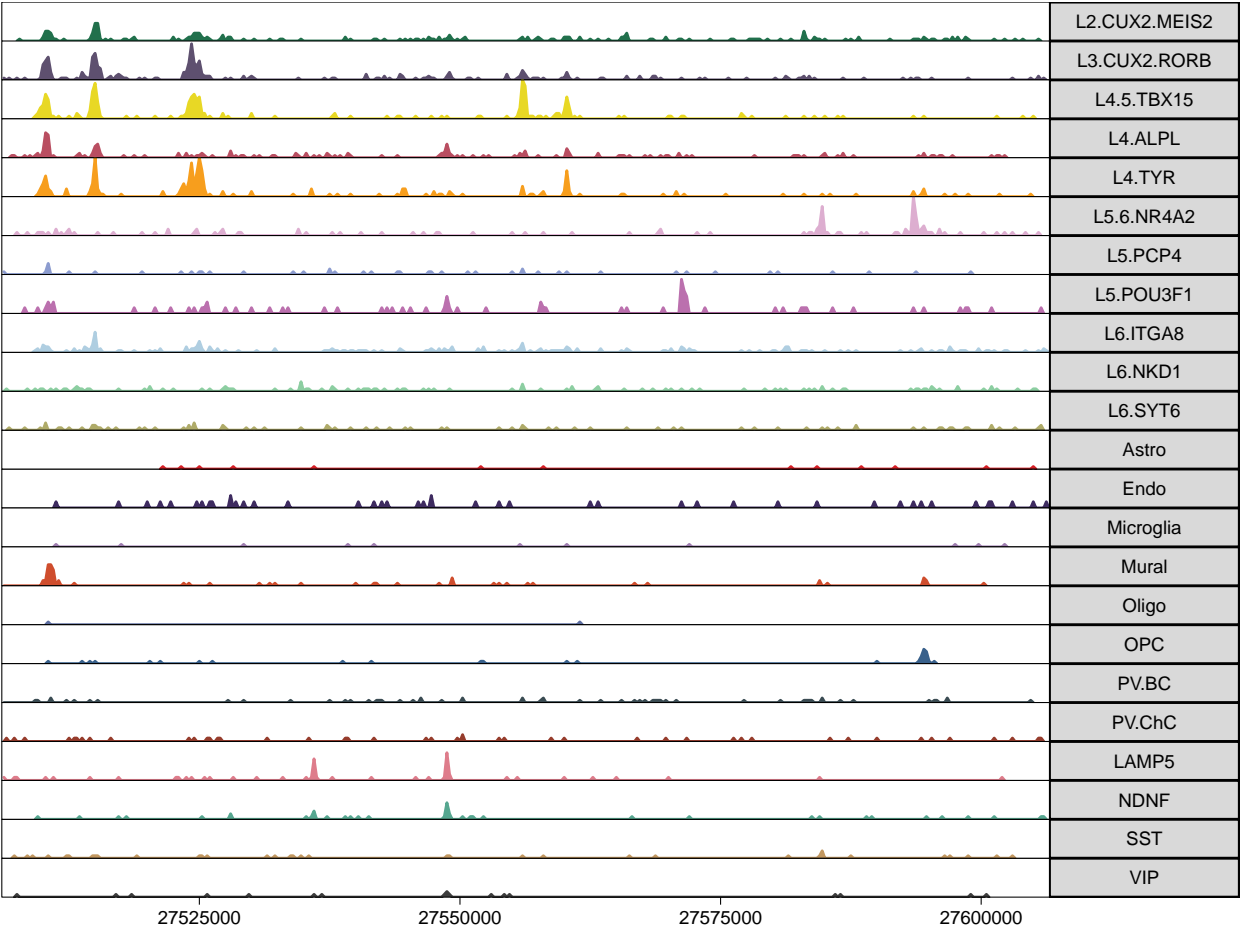
Peaks



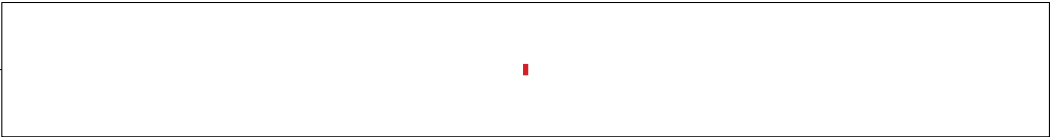
Genes

chr1:27506048–27606549 Marker Peak Tracks for: L4.5.TBX15

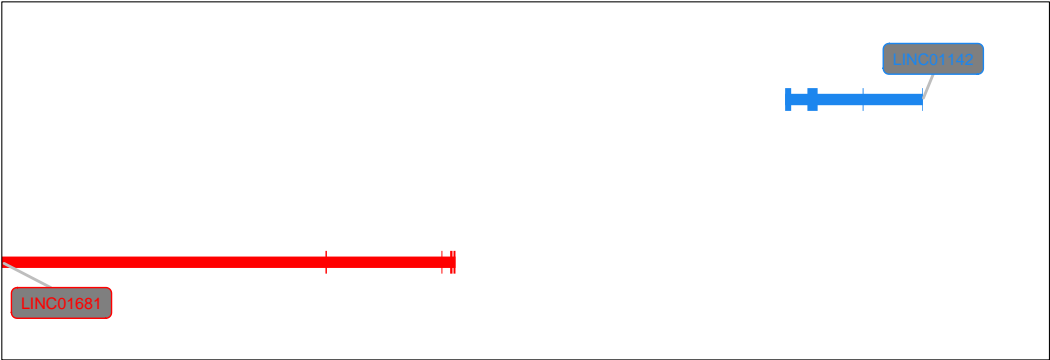
Coverage
(Norm. ATAC Signal Range (0–0.5) by ReadsInTSS)



L4.5.TBX15

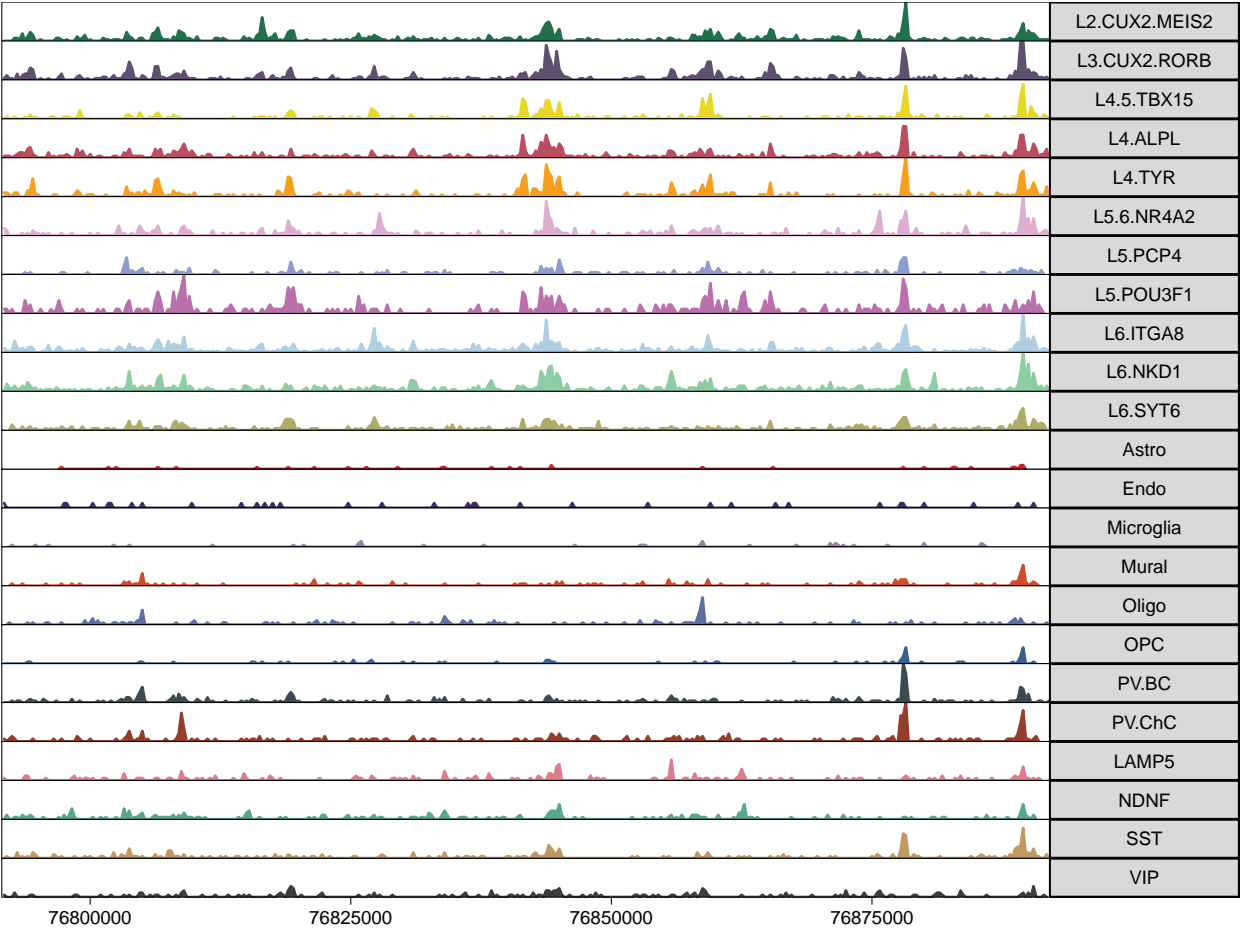


Peaks

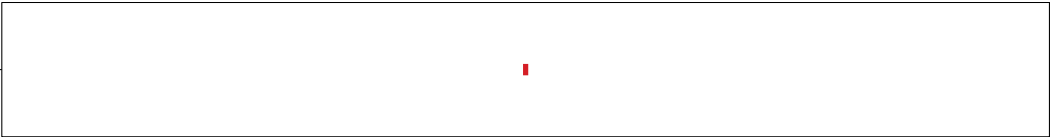


Genes

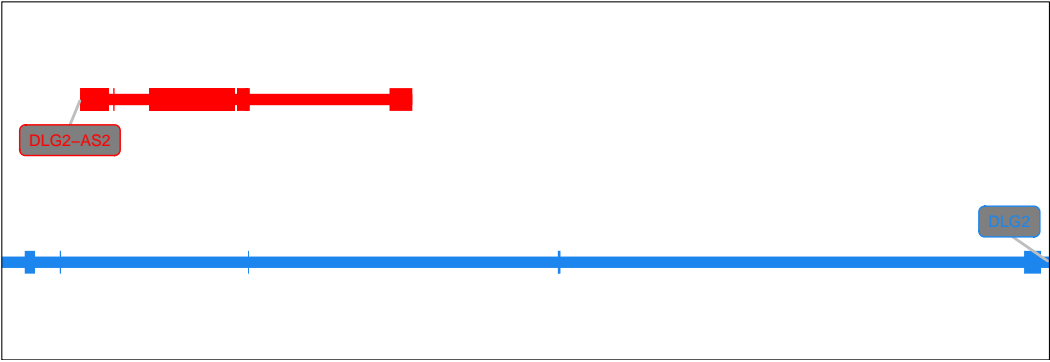
Coverage
(Norm. ATAC Signal Range (0–0.66) by ReadsInTSS)



L4.5.TBX15

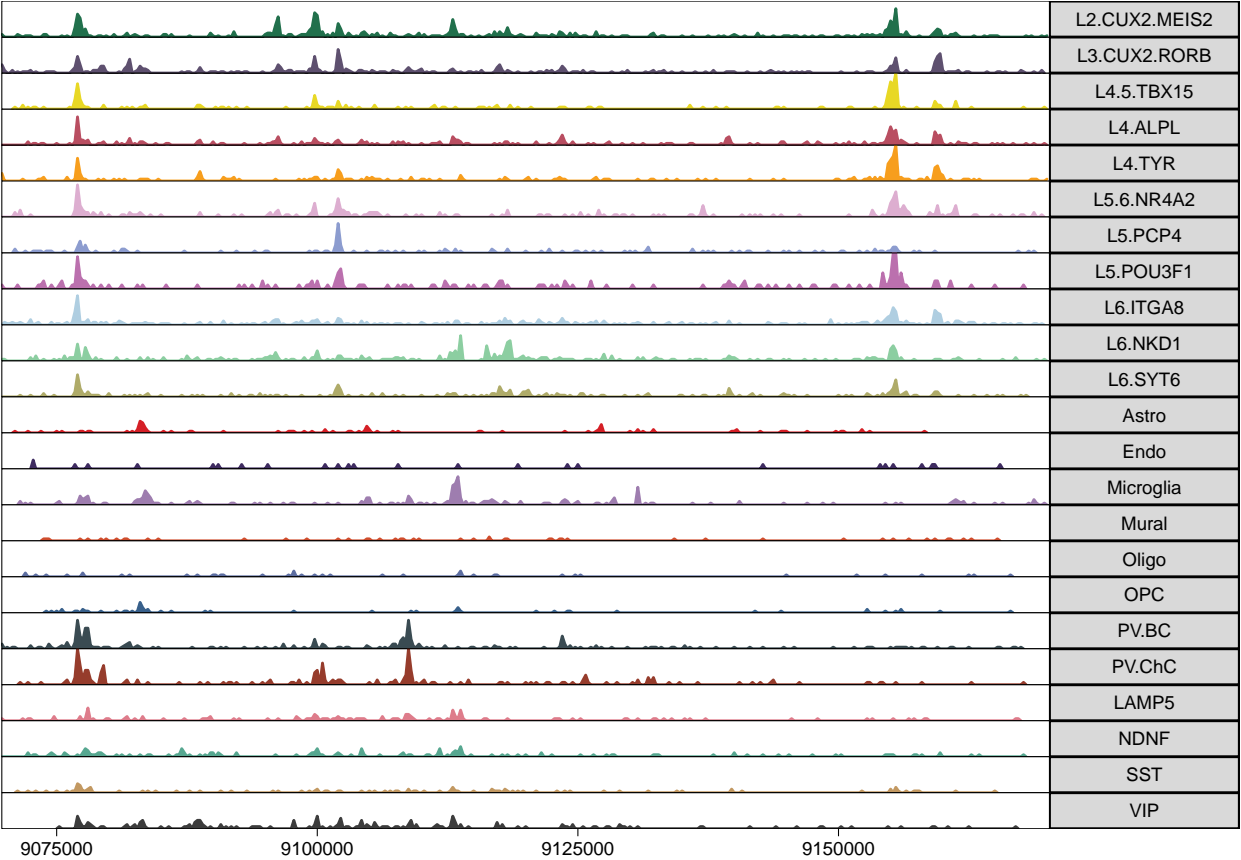


Peaks

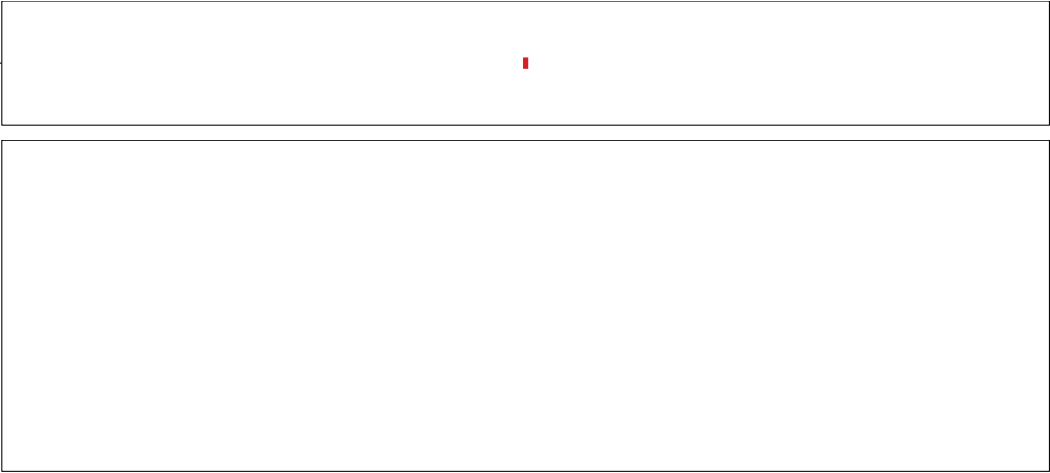


Genes

Coverage
(Norm. ATAC Signal Range (0–0.66) by ReadsInTSS)



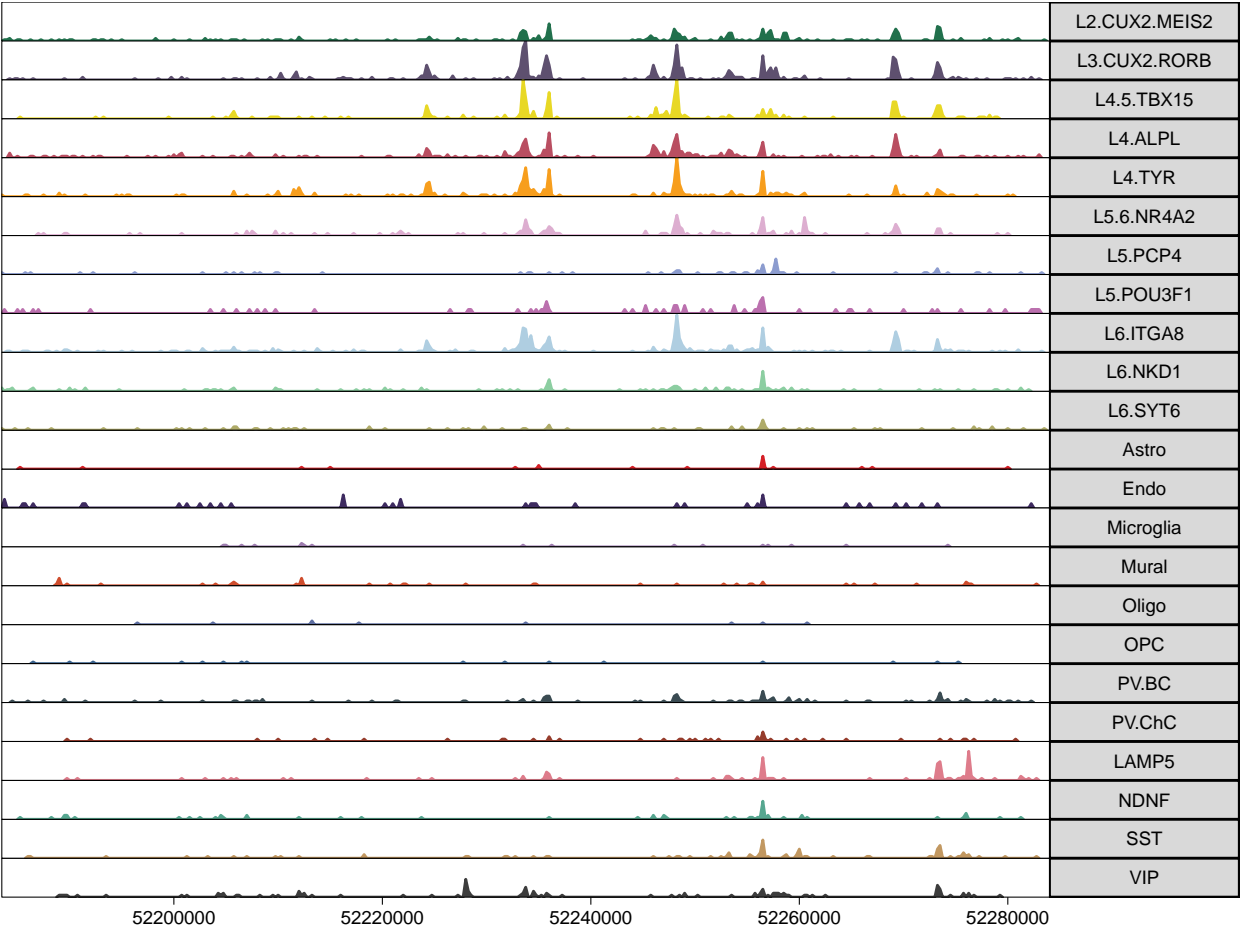
L4.5.TBX15



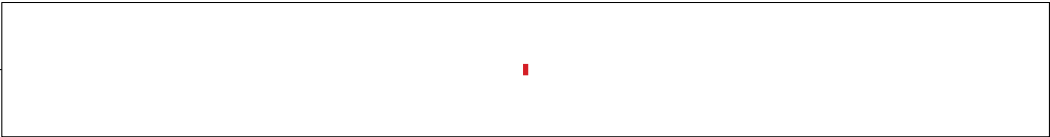
Peaks

GeneTrack

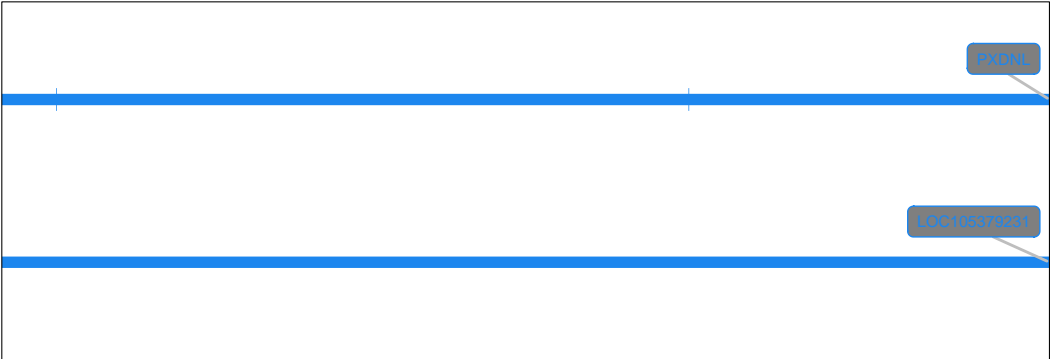
Coverage
(Norm. ATAC Signal Range (0–0.72) by ReadsInTSS)



L4.5.TBX15

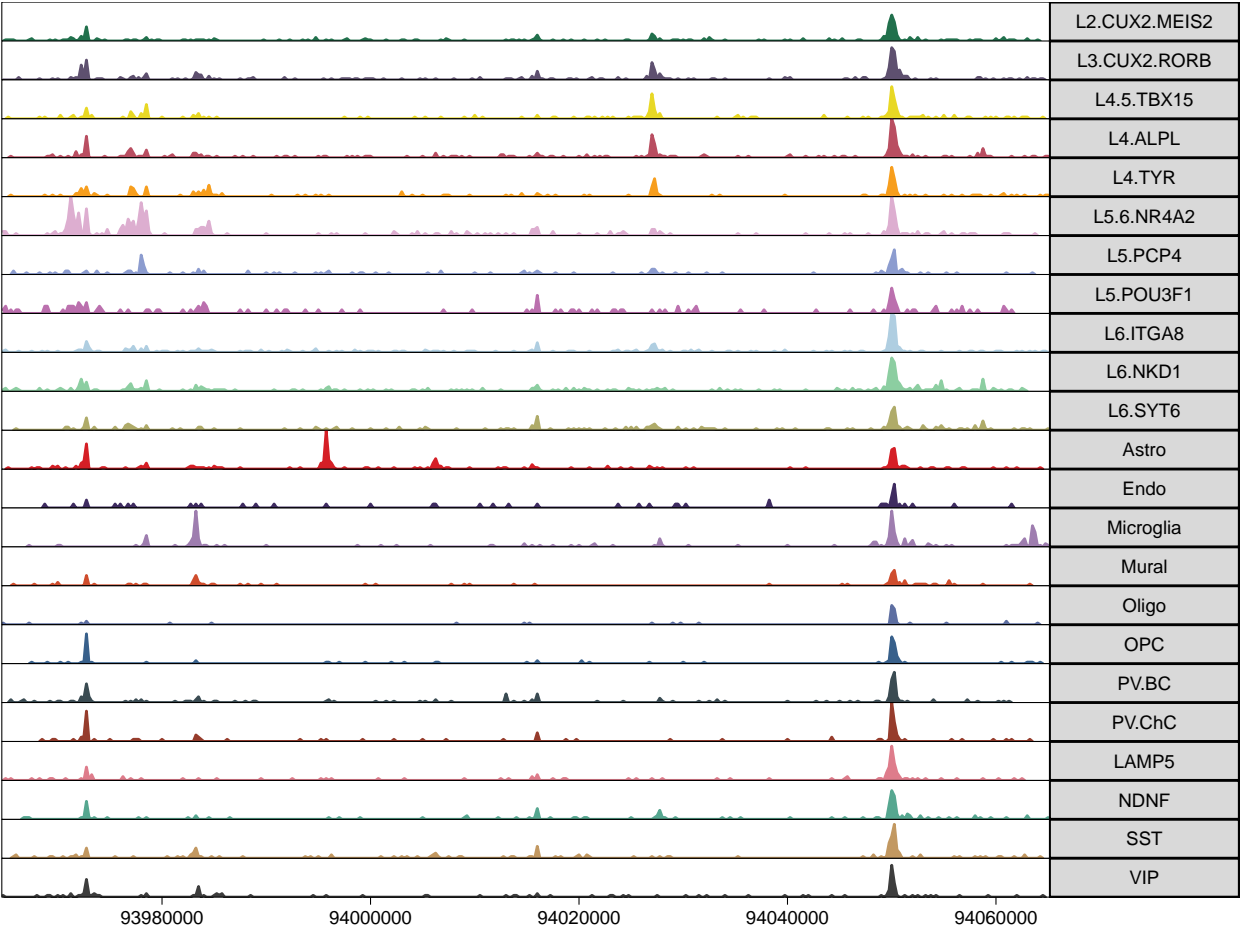


Peaks

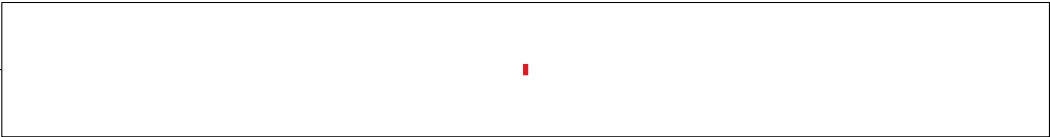


Genes

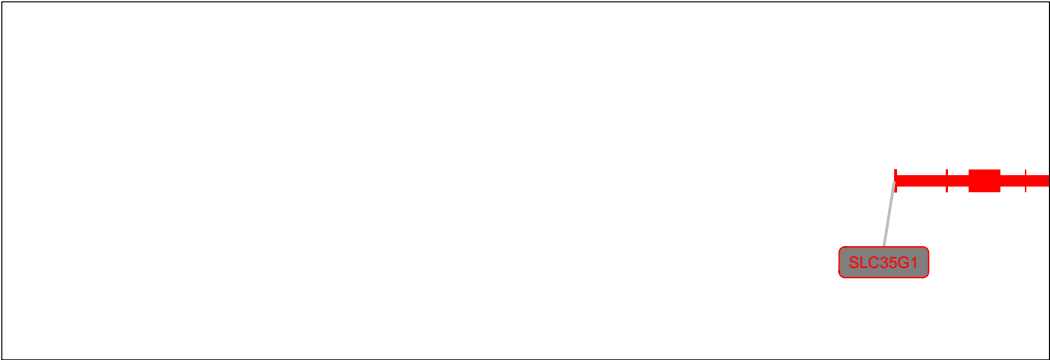
Coverage
(Norm. ATAC Signal Range (0-0.79) by ReadsInTSS)



L4.5.TBX15

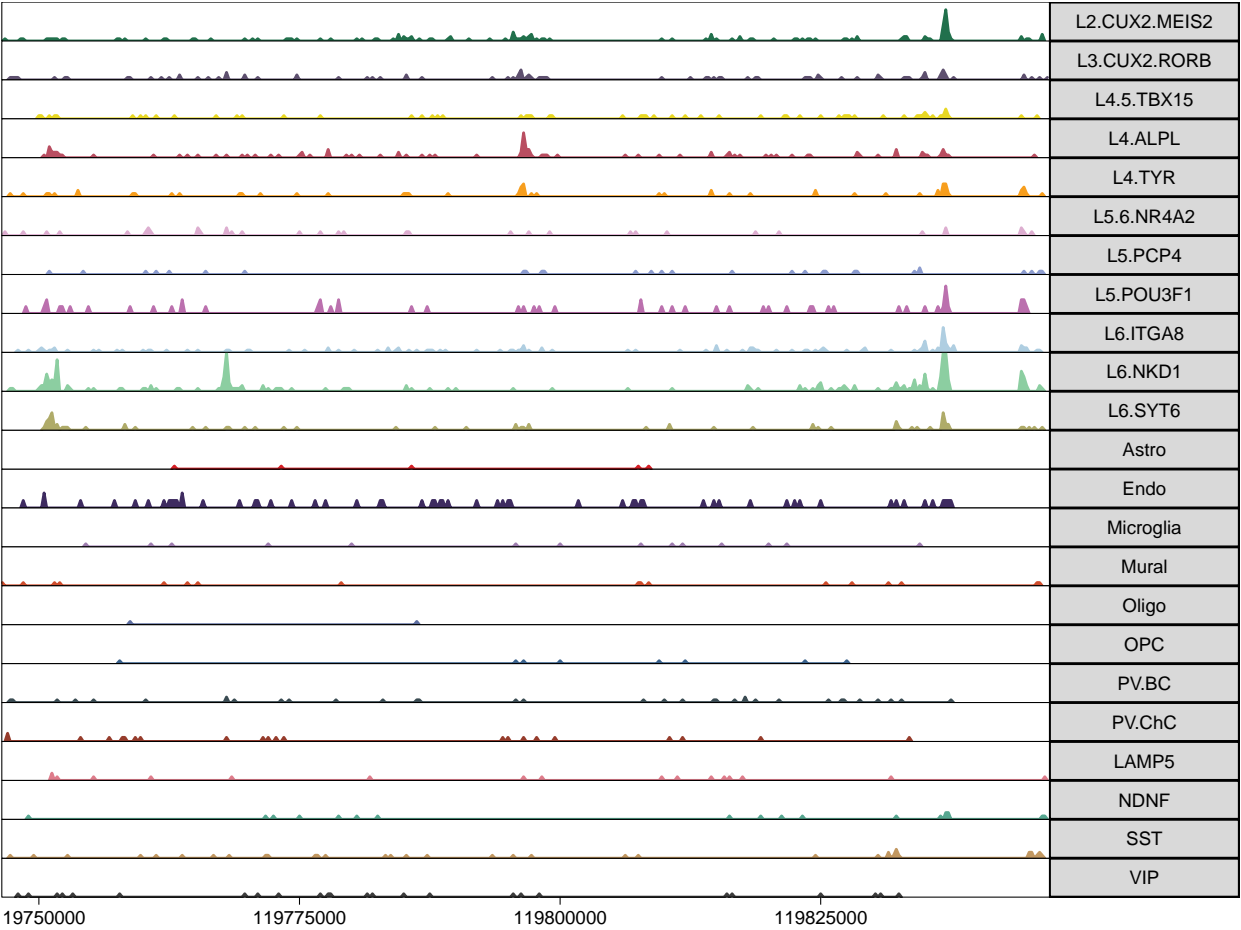


Peaks

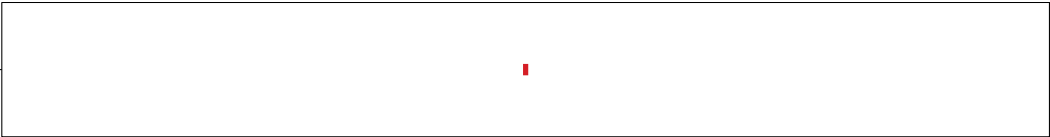


Genes

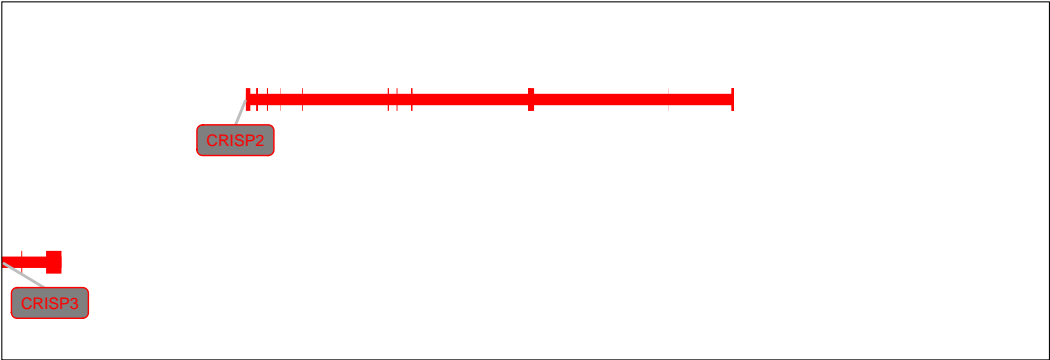
Coverage
(Norm. ATAC Signal Range (0–0.42) by ReadsInTSS)



L4.5.TBX15

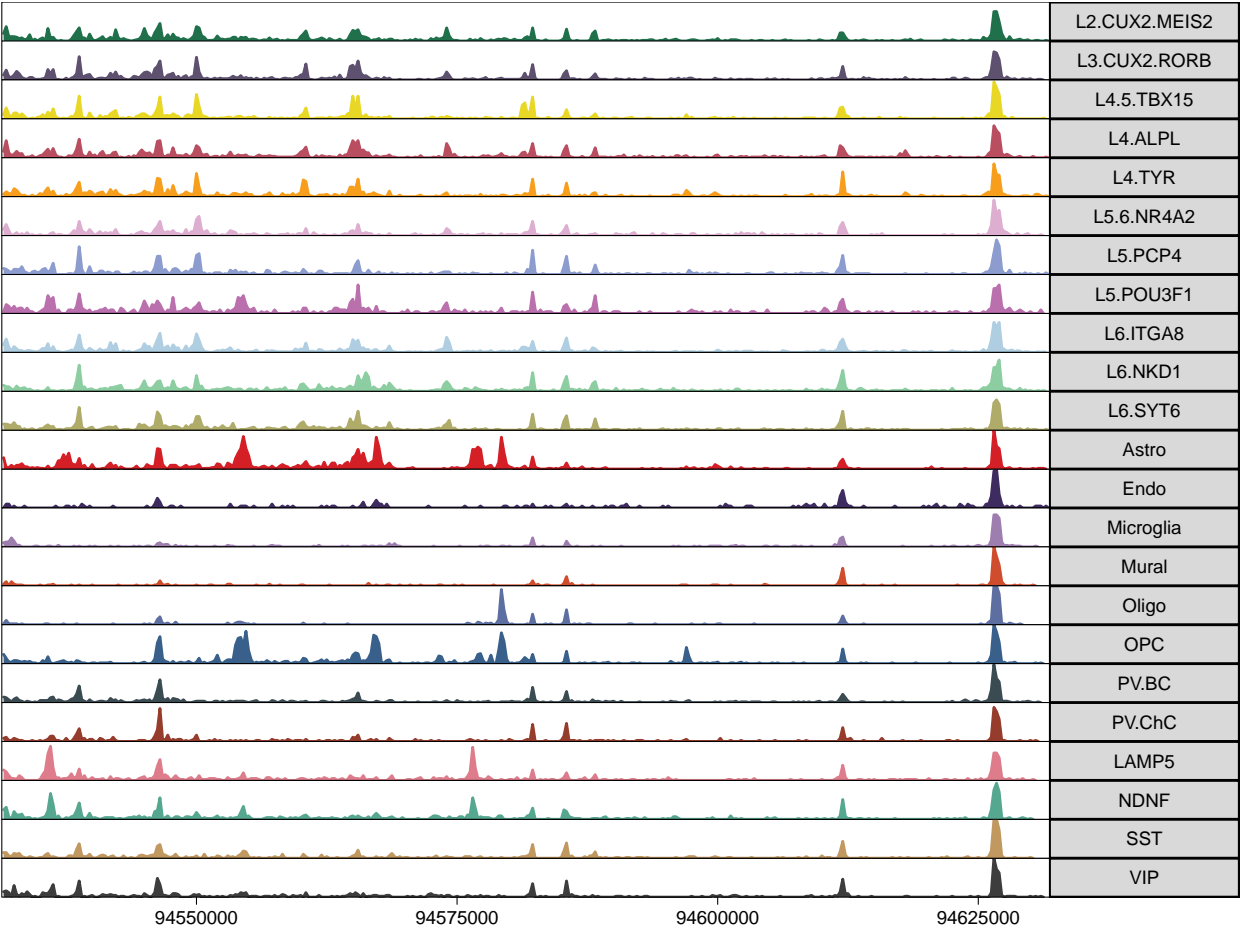


Peaks

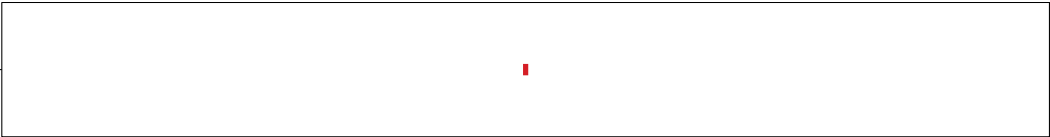


Genes

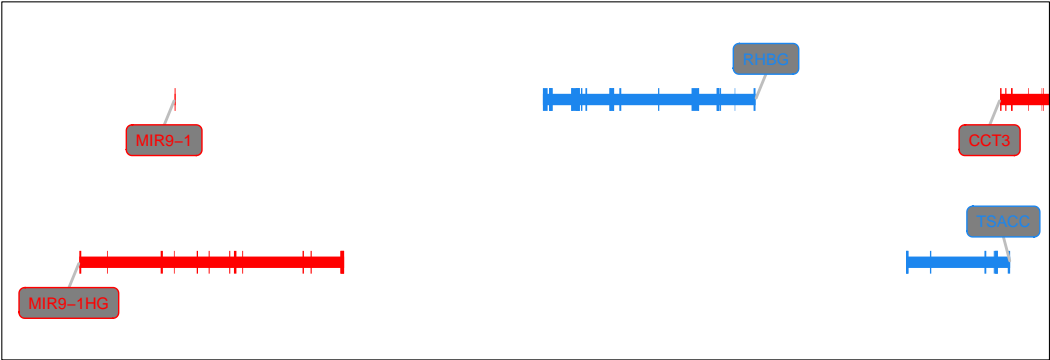
Coverage
(Norm. ATAC Signal Range (0–1.61) by ReadsInTSS)



L4.5.TBX15



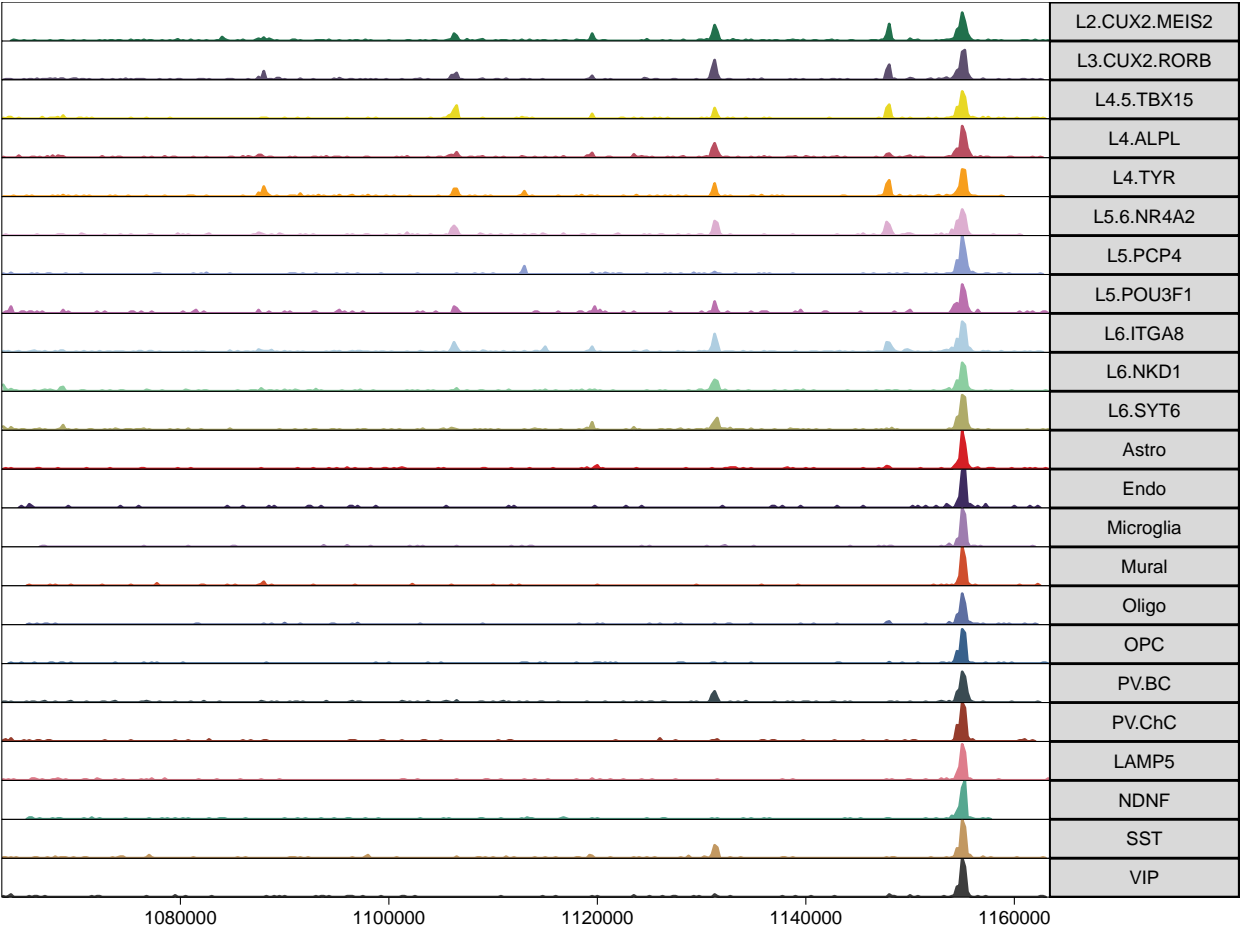
Peaks



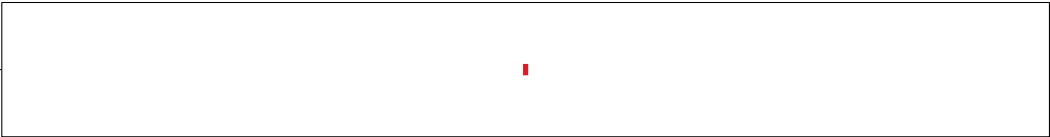
Genes

chr18:1062868–1163369 Marker Peak Tracks for: L4.5.TBX15

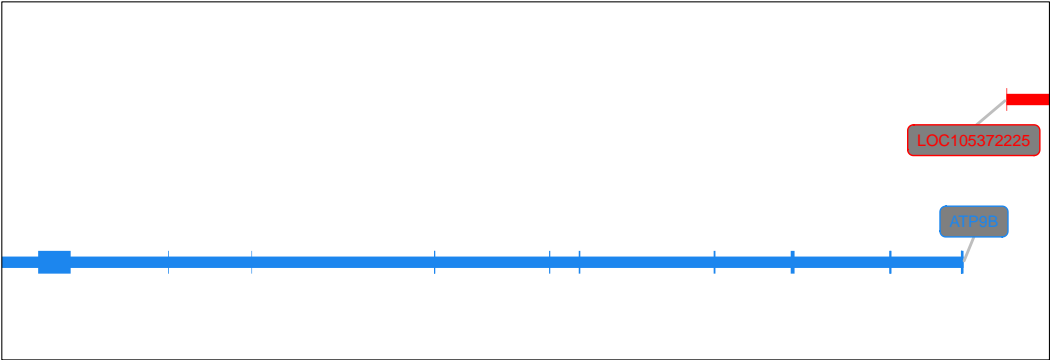
Coverage
(Norm. ATAC Signal Range (0–1.68) by ReadsInTSS)



L4.5.TBX15



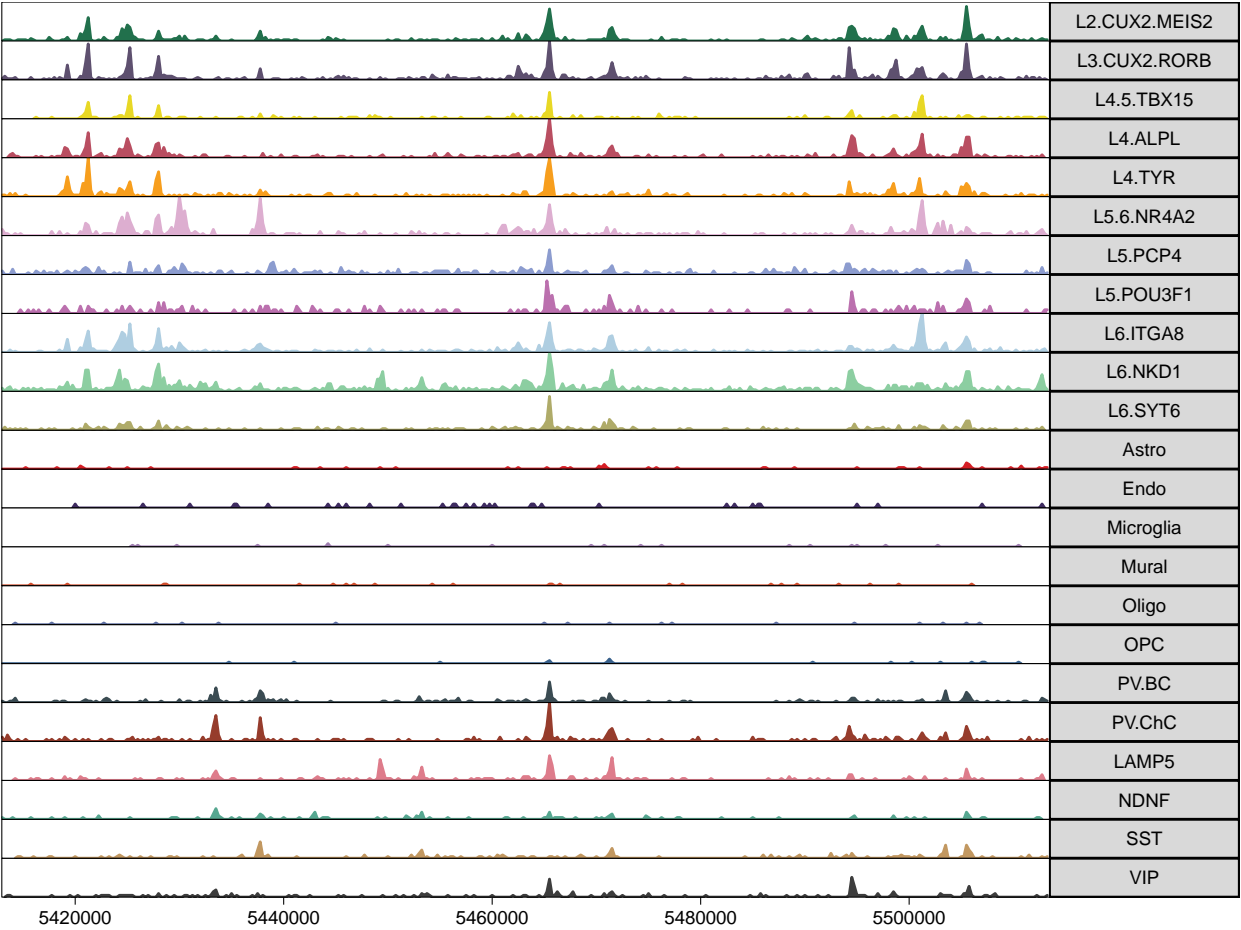
Peaks



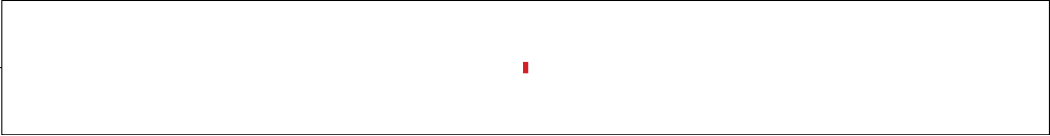
Genes

chr20:5412959–5513460 Marker Peak Tracks for: L4.5.TBX15

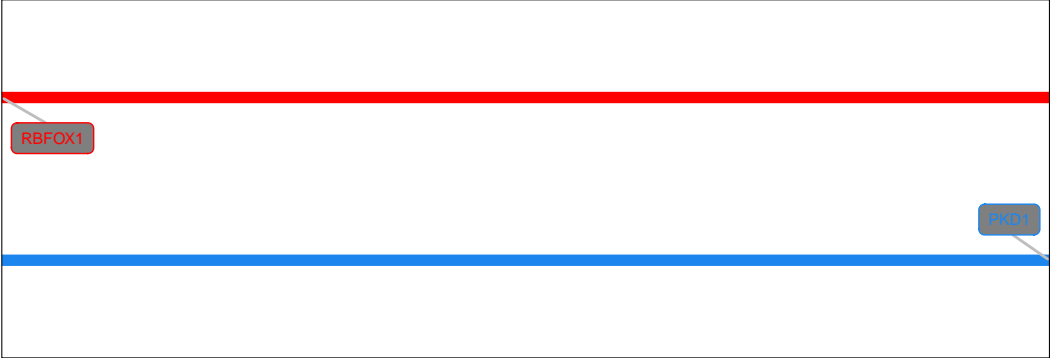
Coverage
(Norm. ATAC Signal Range (0–0.79) by ReadsInTSS)



L4.5.TBX15

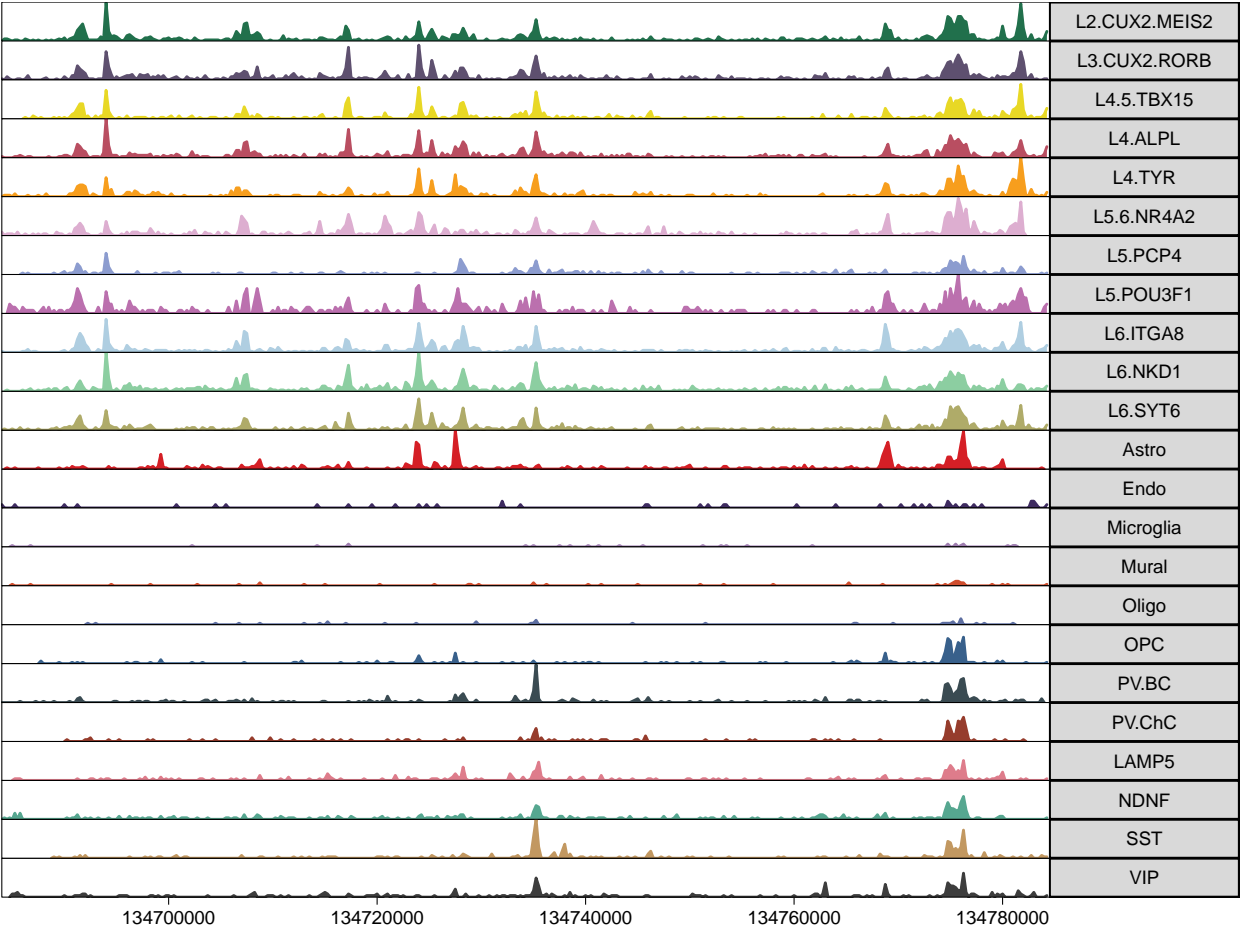


Peaks

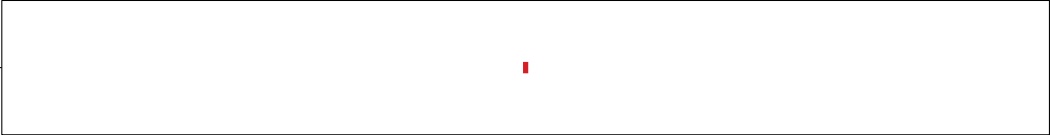


Genes

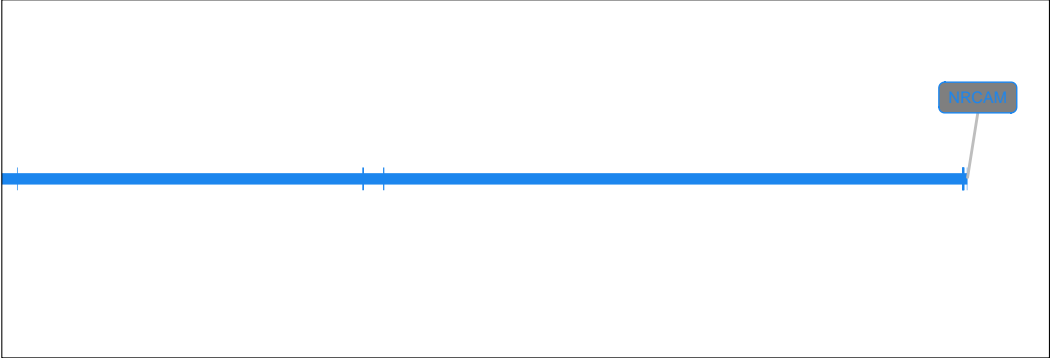
Coverage
(Norm. ATAC Signal Range (0–0.92) by ReadsInTSS)



L4.5.TBX15

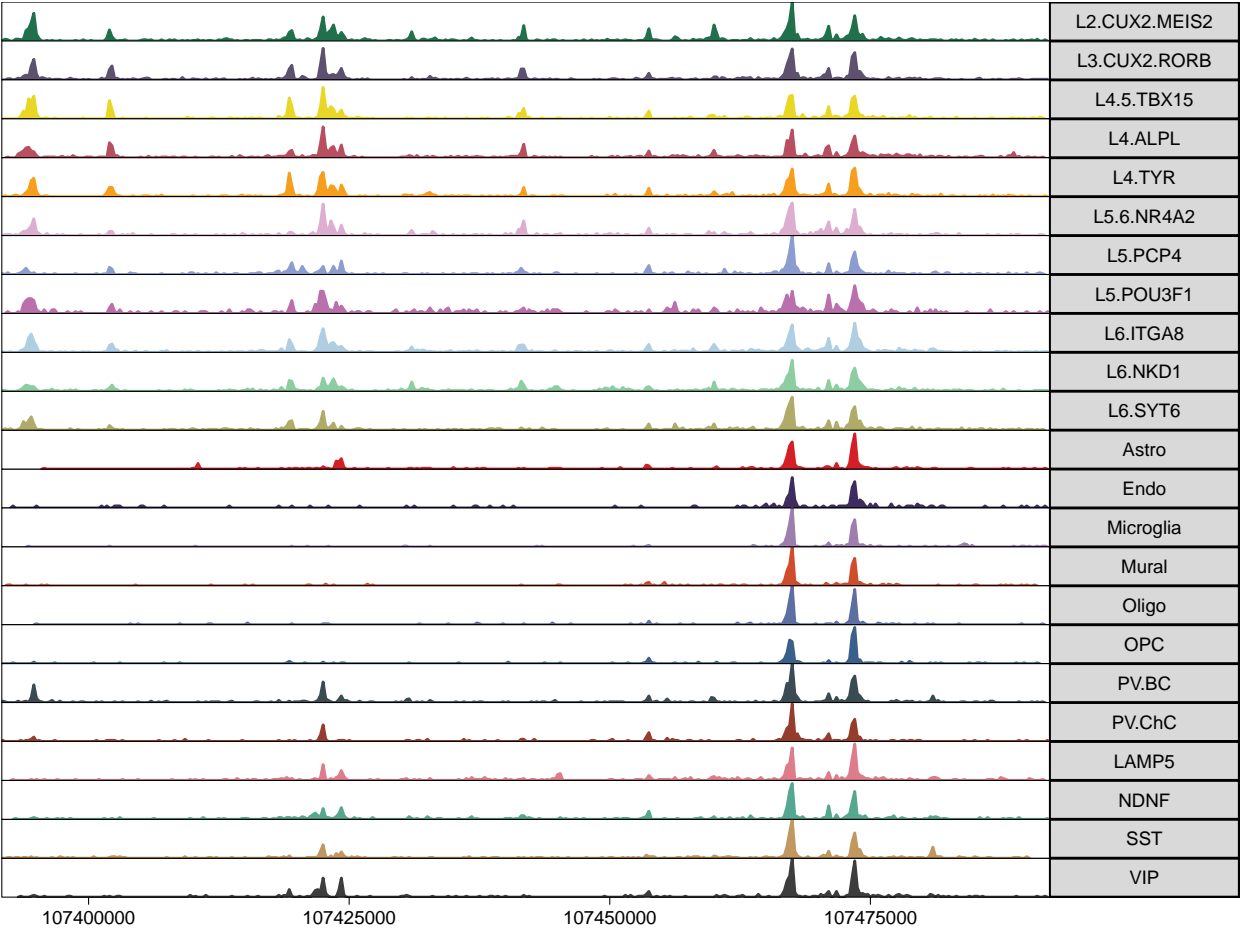


Peaks

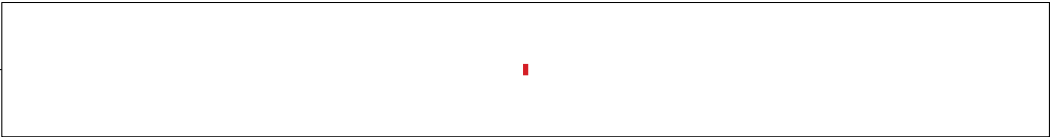


Genes

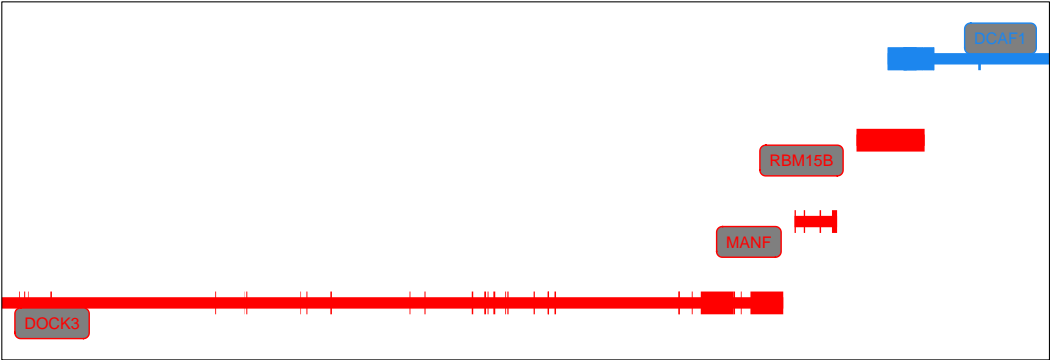
Coverage
(Norm. ATAC Signal Range (0–1.54) by ReadsInTSS)



L4.5.TBX15

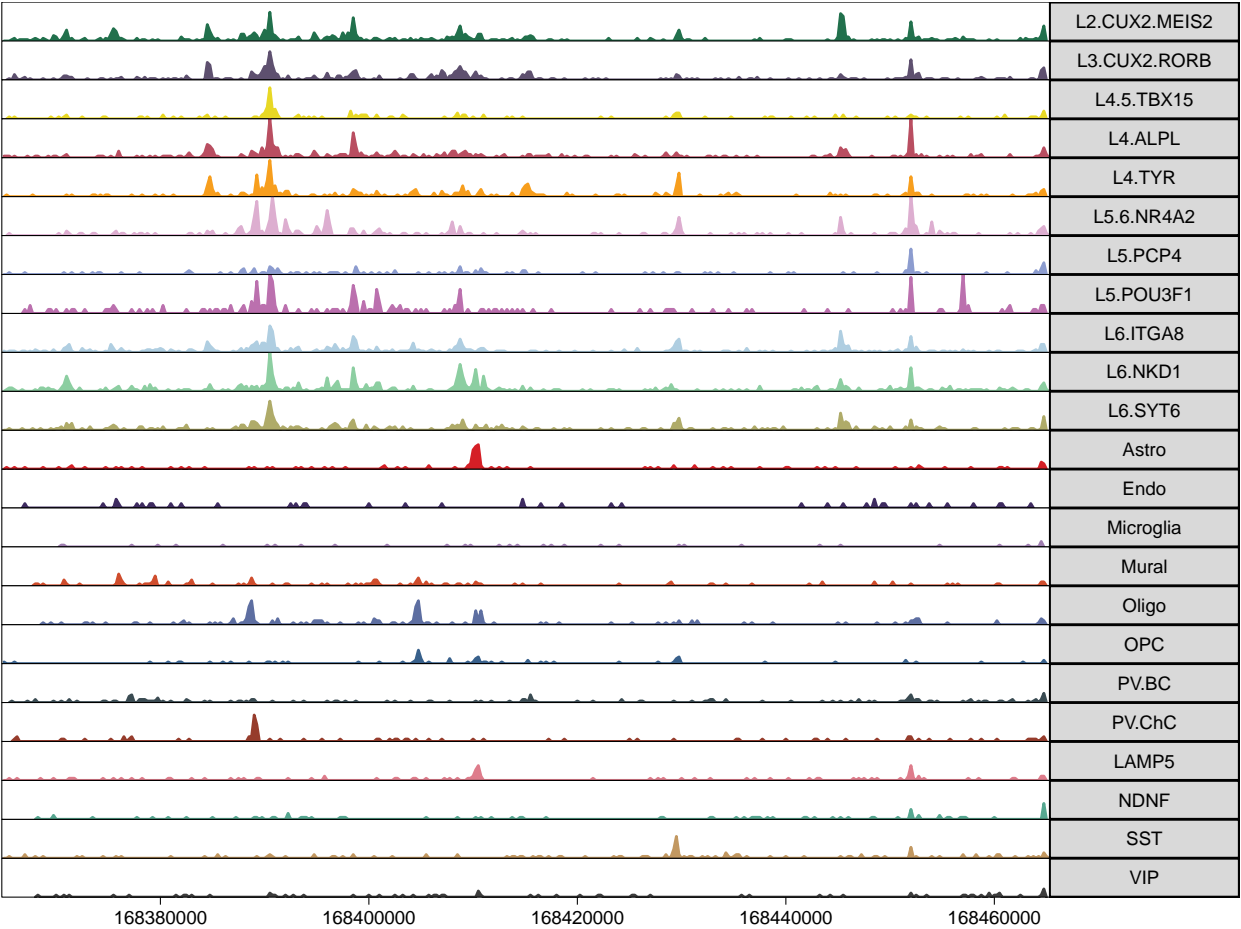


Peaks

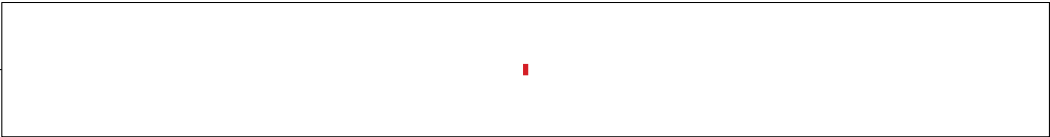


Genes

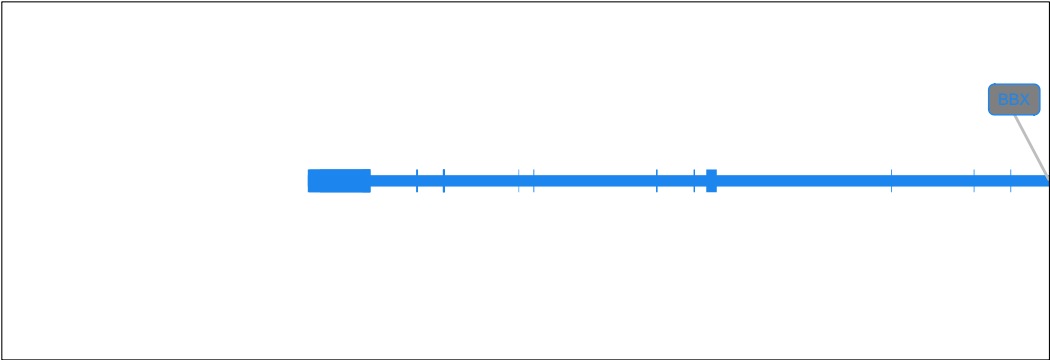
Coverage
(Norm. ATAC Signal Range (0–0.71) by ReadsInTSS)



L4.5.TBX15

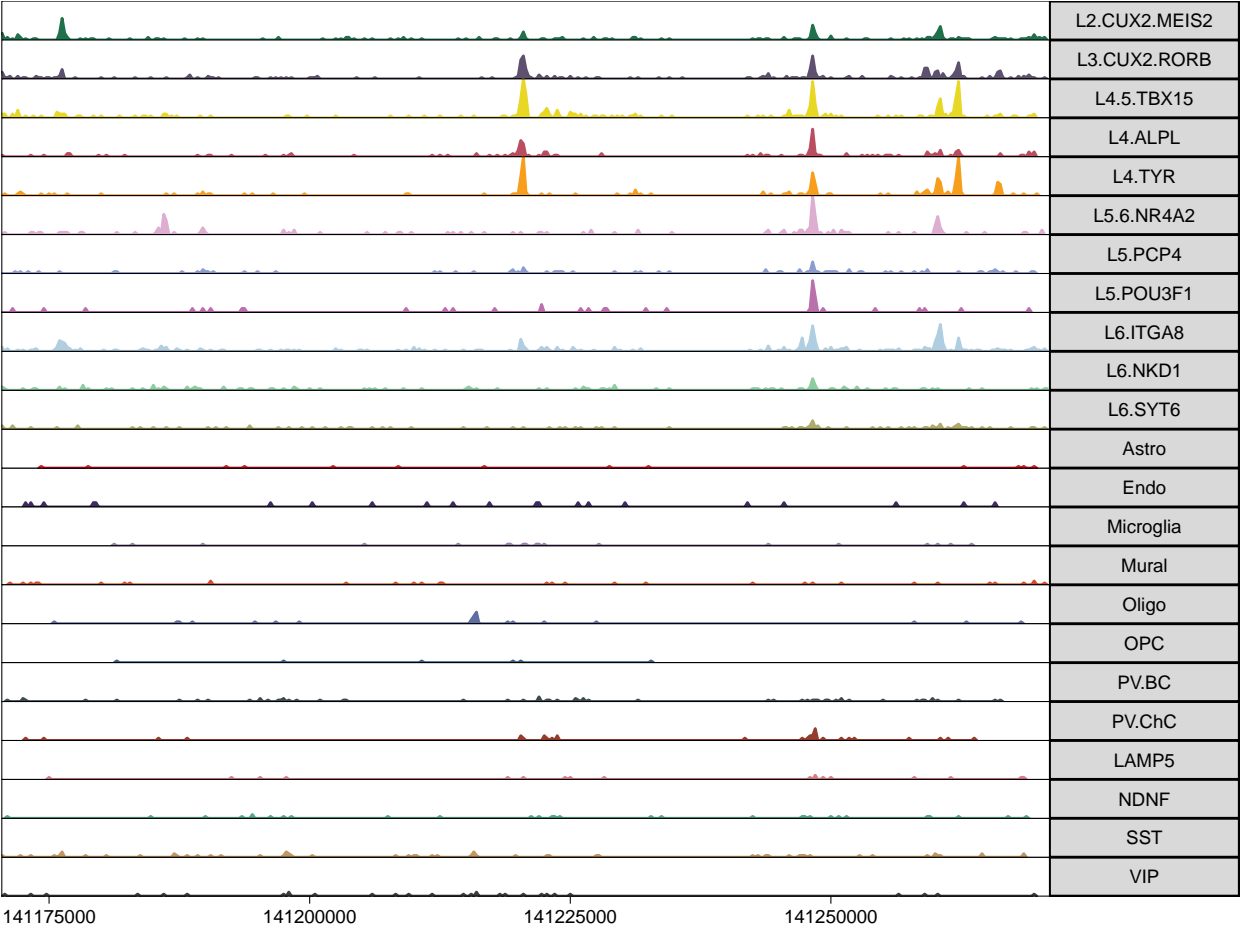


Peaks

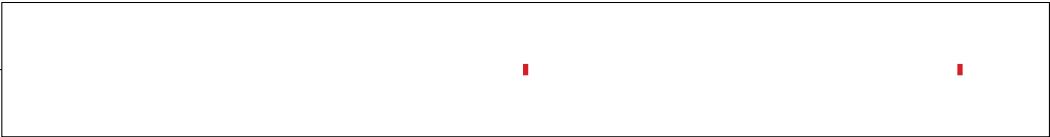


Genes

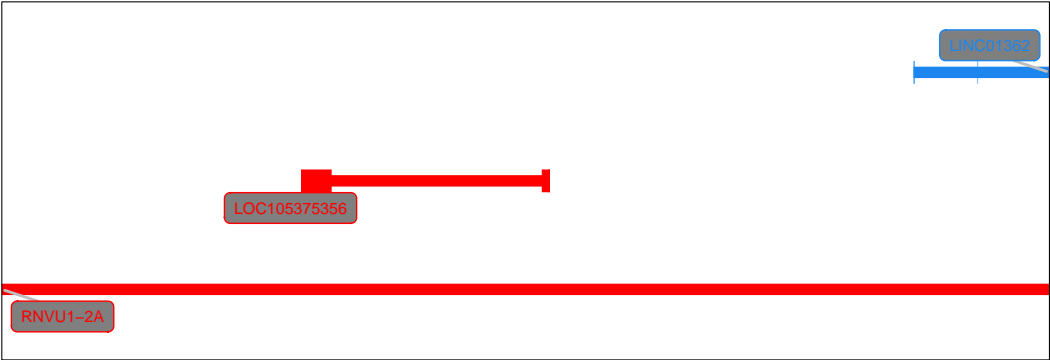
Coverage
(Norm. ATAC Signal Range (0–0.72) by ReadsInTSS)



L4.5.TBX15

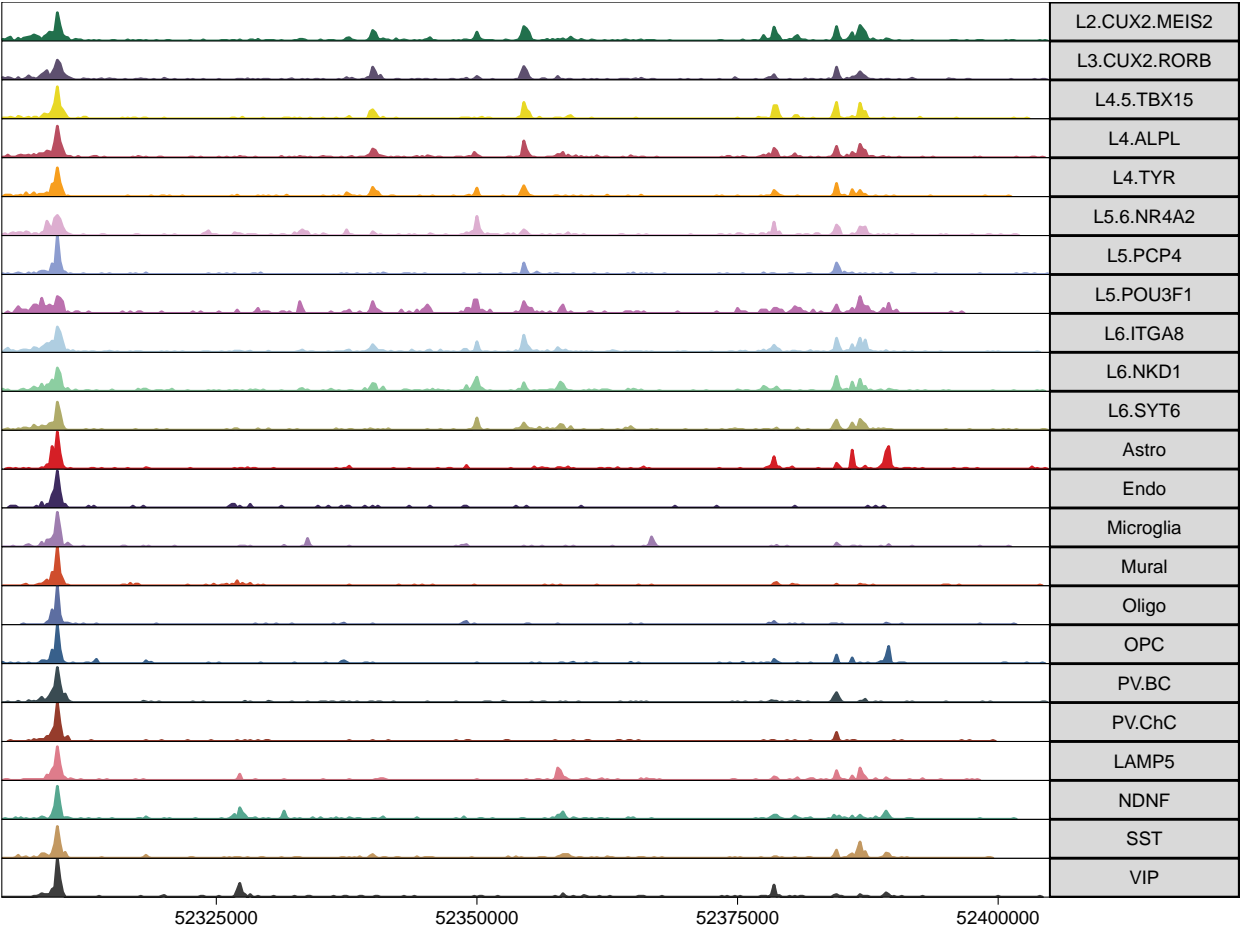


Peaks

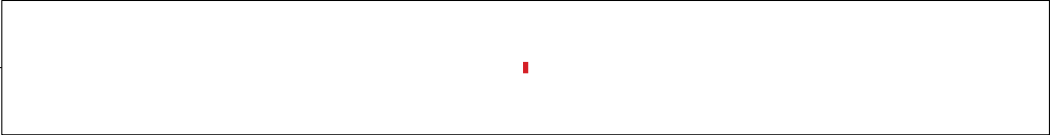


Genes

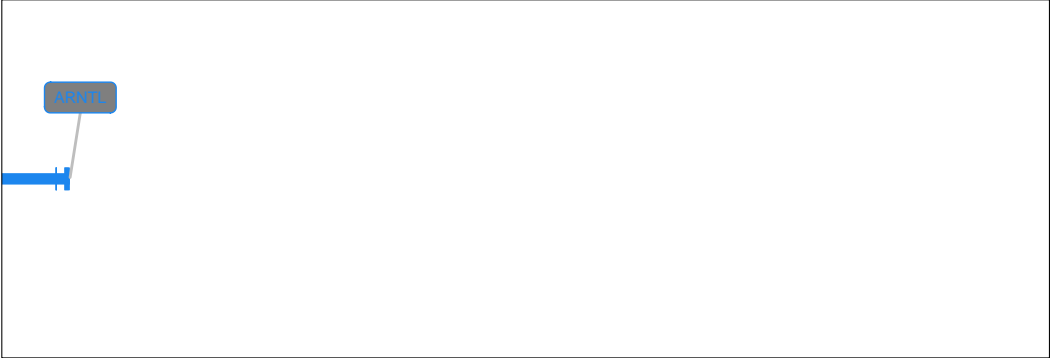
Coverage
(Norm. ATAC Signal Range (0–1.69) by ReadsInTSS)



L4.5.TBX15

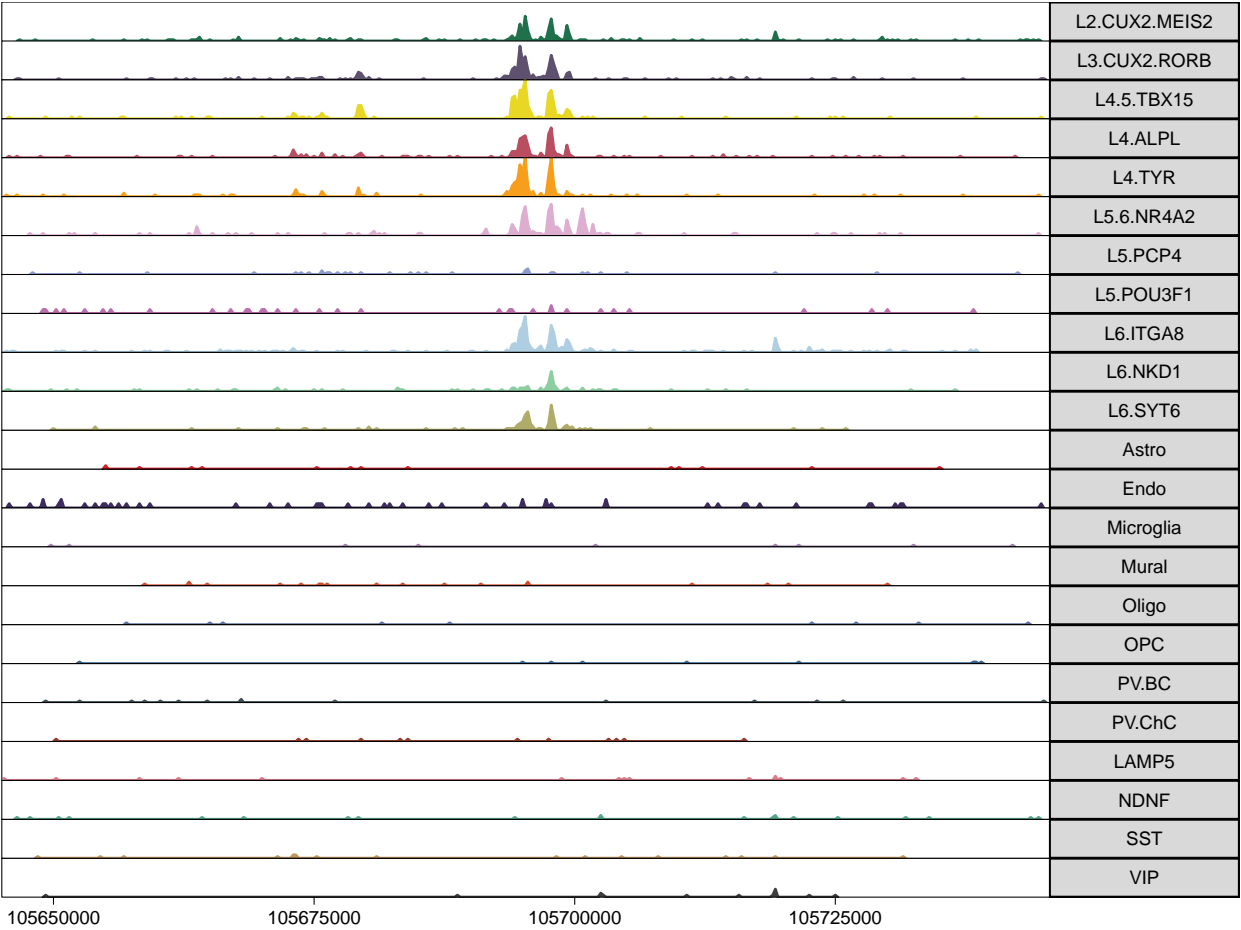


Peaks

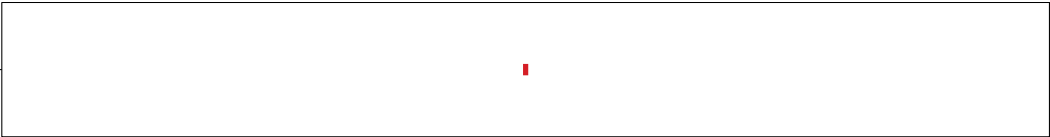


Genes

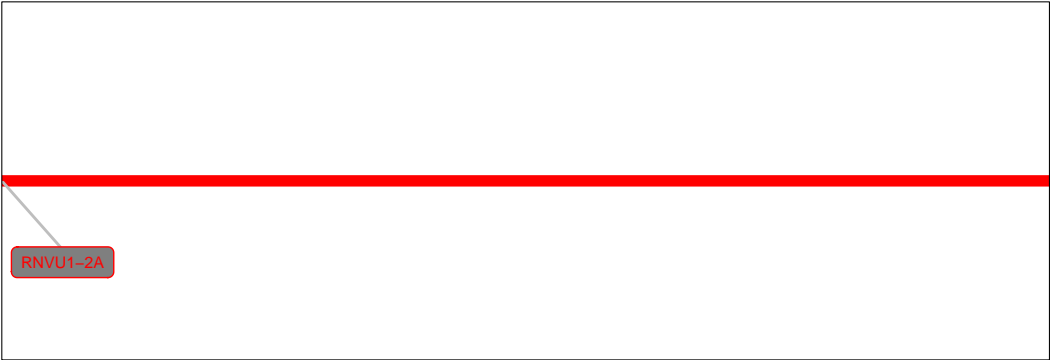
Coverage
(Norm. ATAC Signal Range (0–0.73) by ReadsInTSS)



L4.5.TBX15



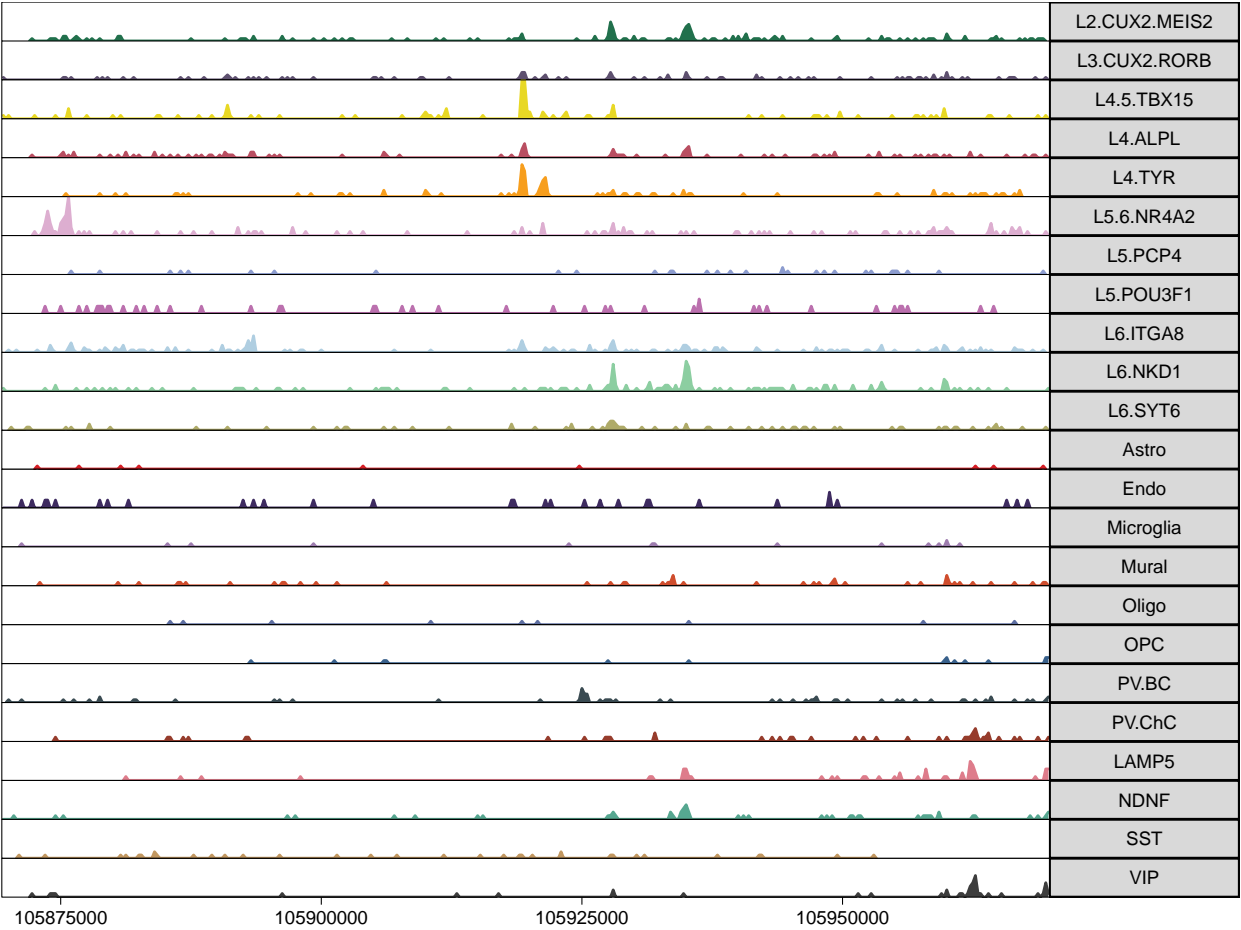
Peaks



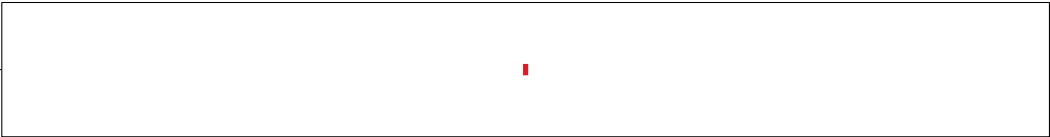
Genes

chr1:105869350–105969851 Marker Peak Tracks for: L4.5.TBX15

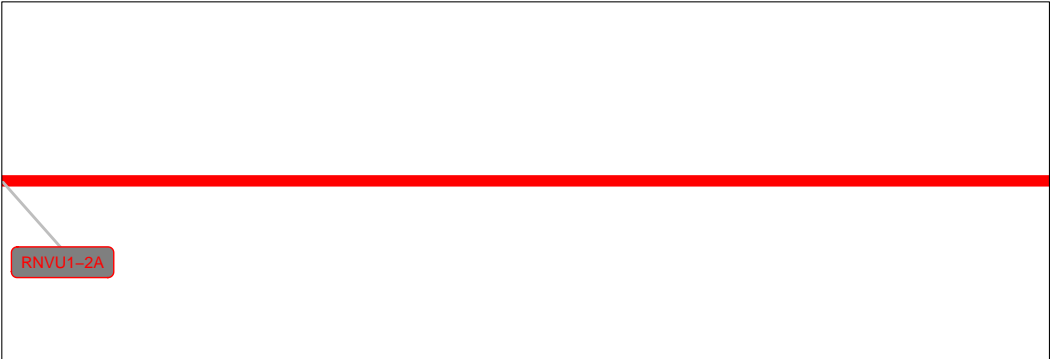
Coverage
(Norm. ATAC Signal Range (0–0.4) by ReadsInTSS)



L4.5.TBX15

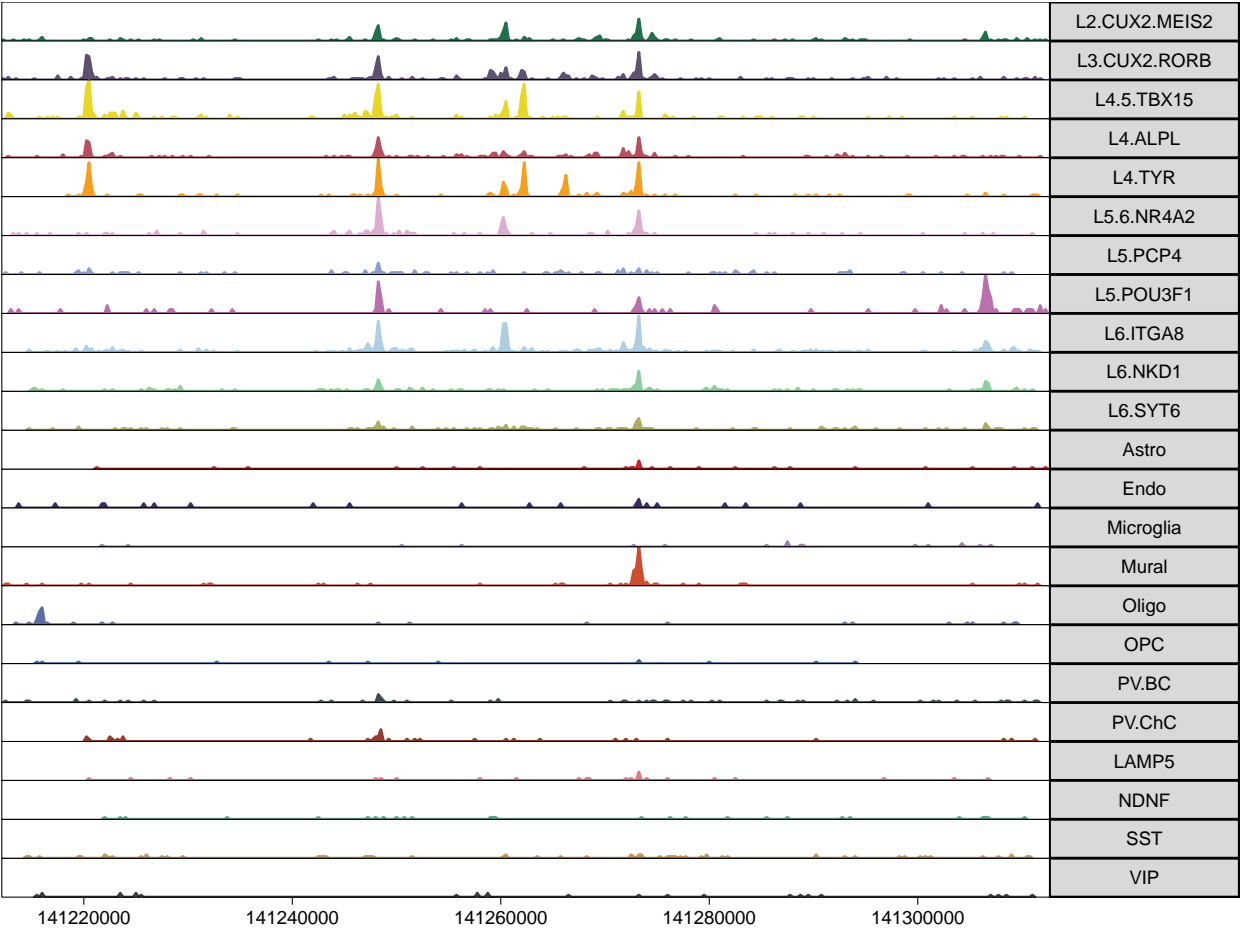


Peaks

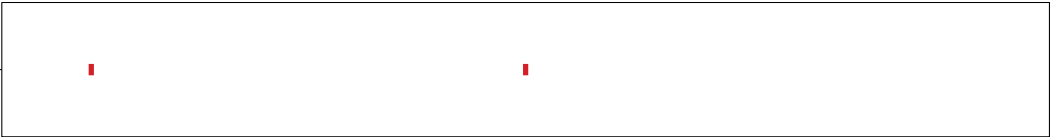


Genes

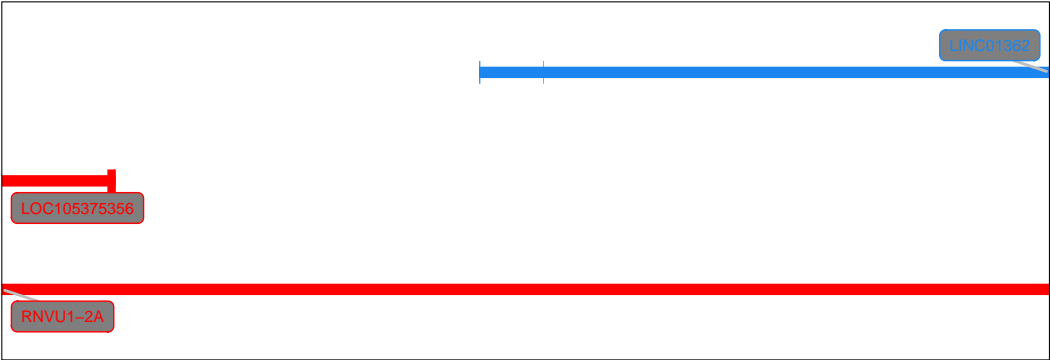
Coverage
(Norm. ATAC Signal Range (0–0.72) by ReadsInTSS)



L4.5.TBX15

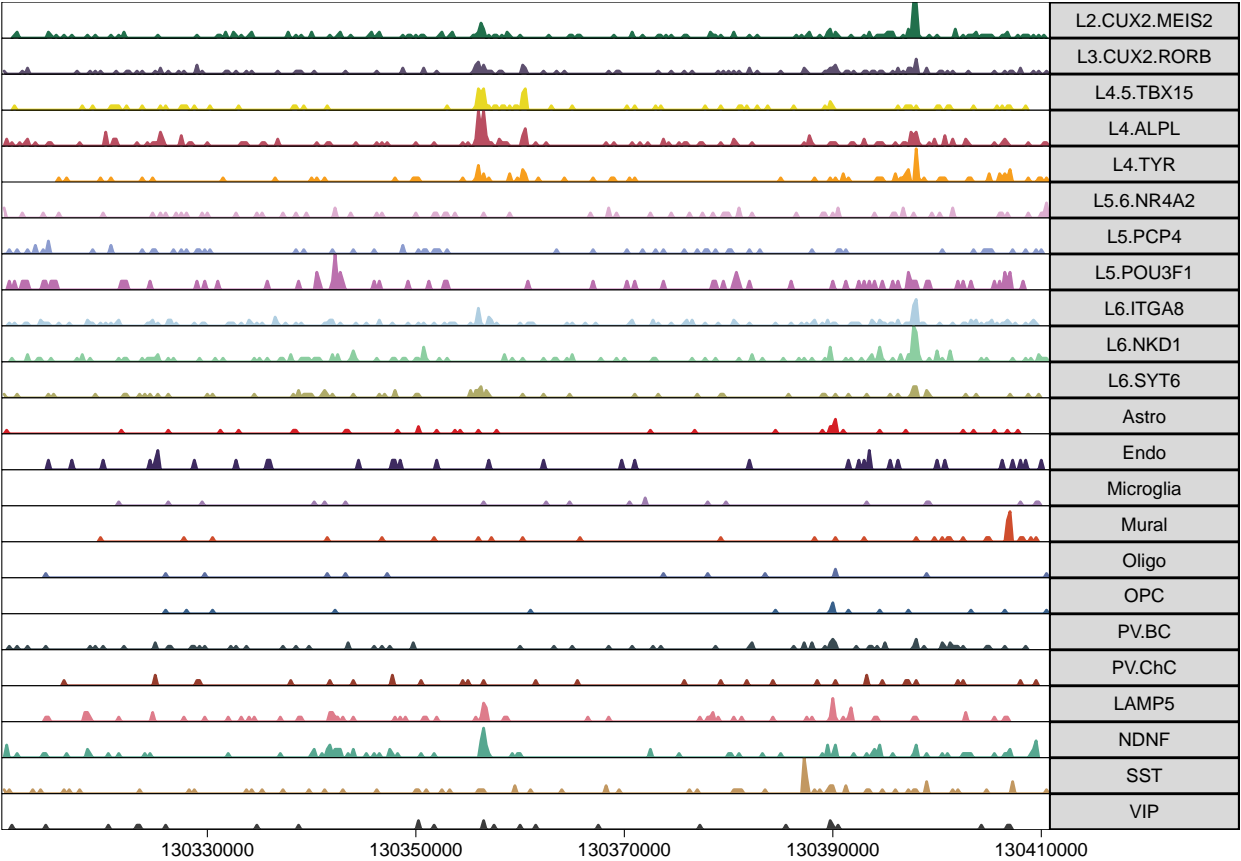


Peaks

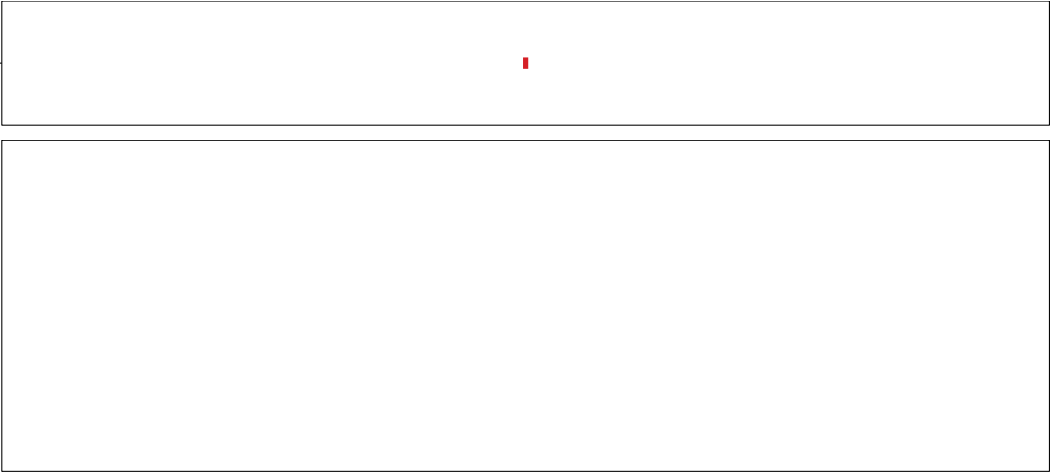


Genes

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



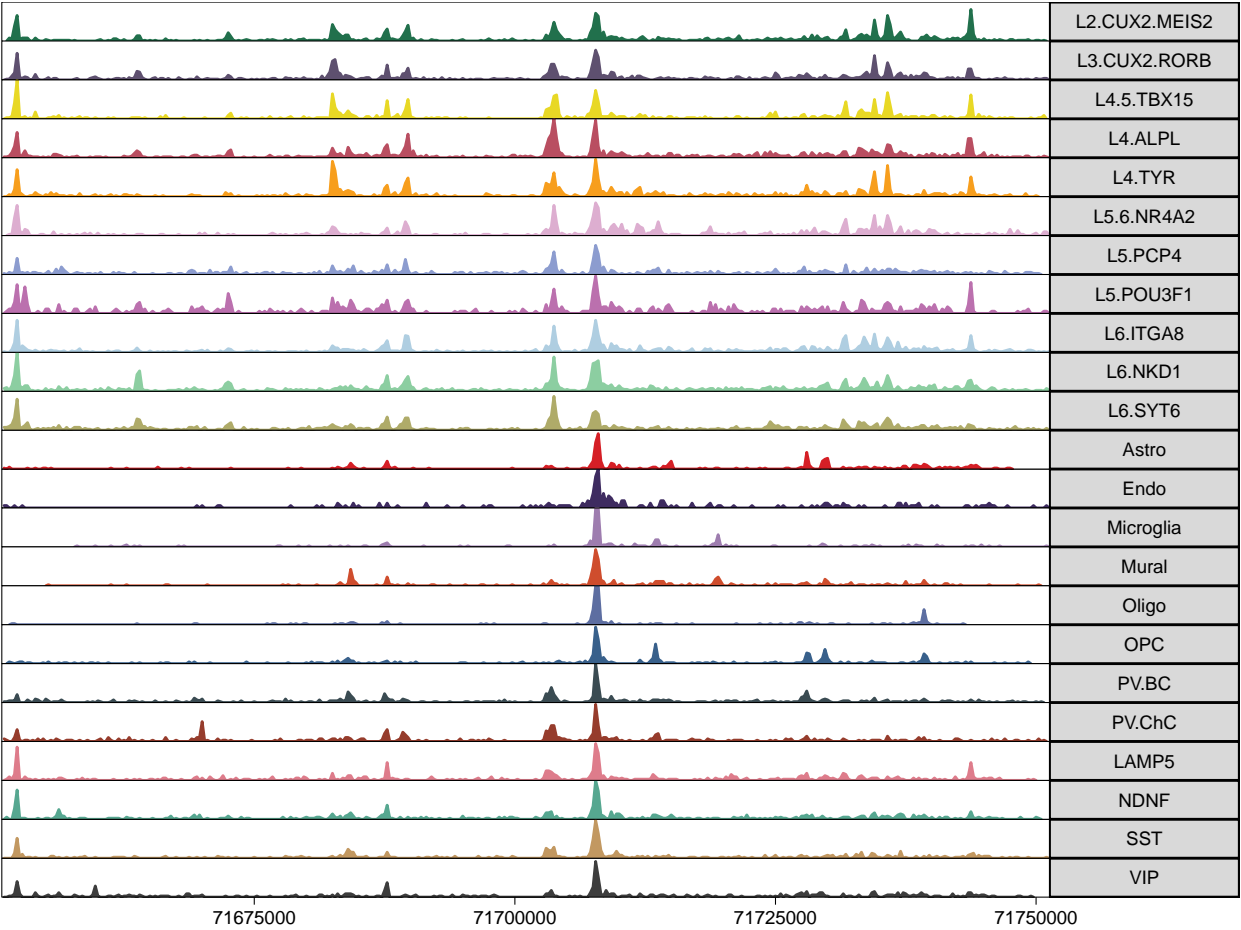
L4.5.TBX15



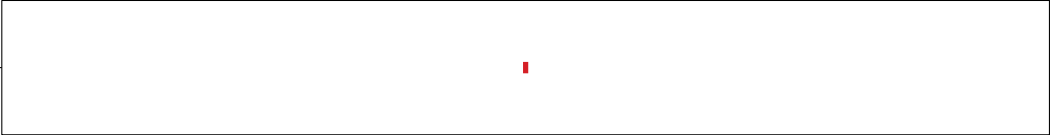
Peaks

GeneTrack

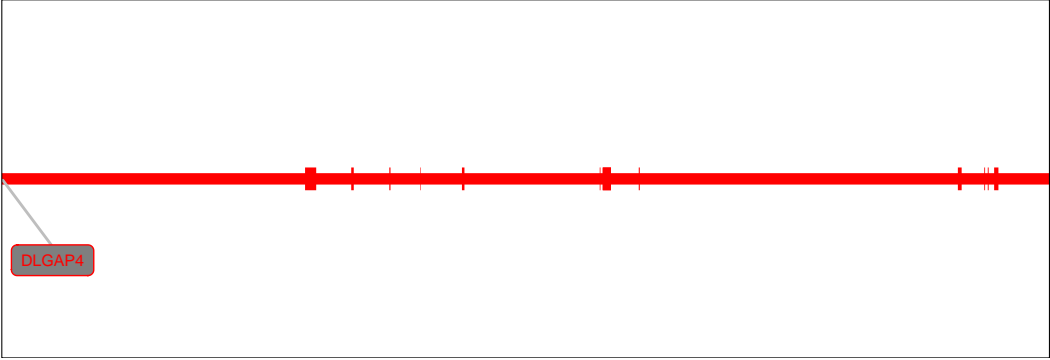
Coverage
(Norm. ATAC Signal Range (0–1.29) by ReadsInTSS)



L4.5.TBX15

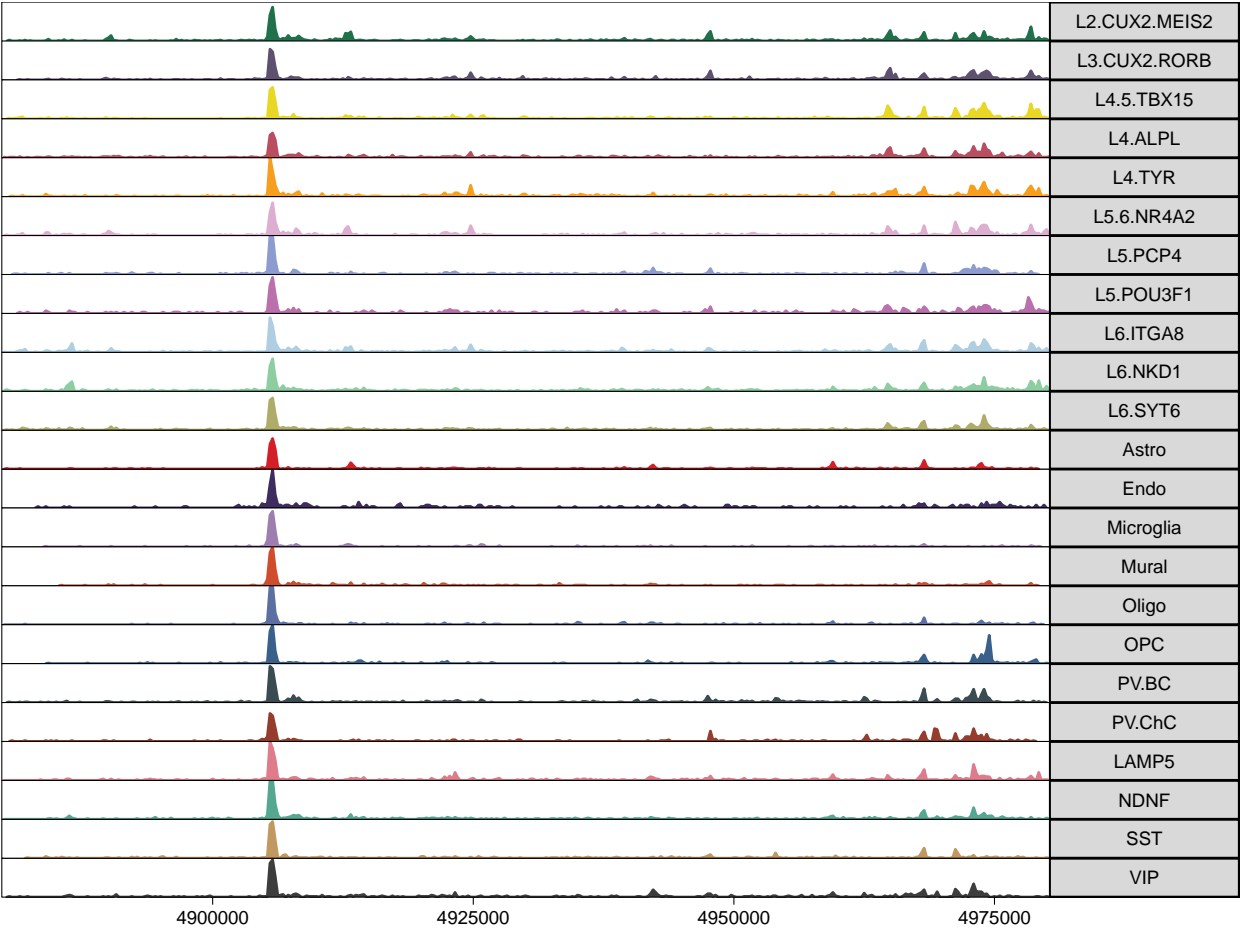


Peaks

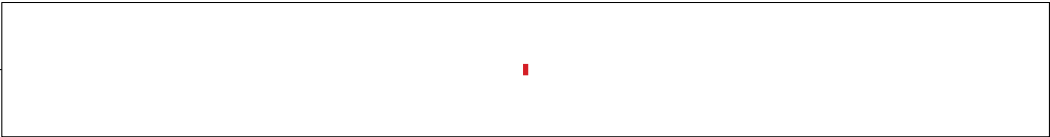


Genes

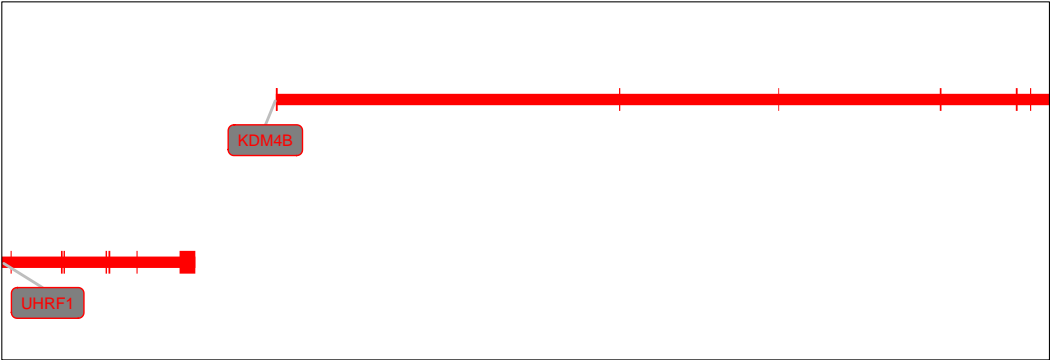
Coverage
(Norm. ATAC Signal Range (0-2.14) by ReadslnTSS)



L4.5.TBX15

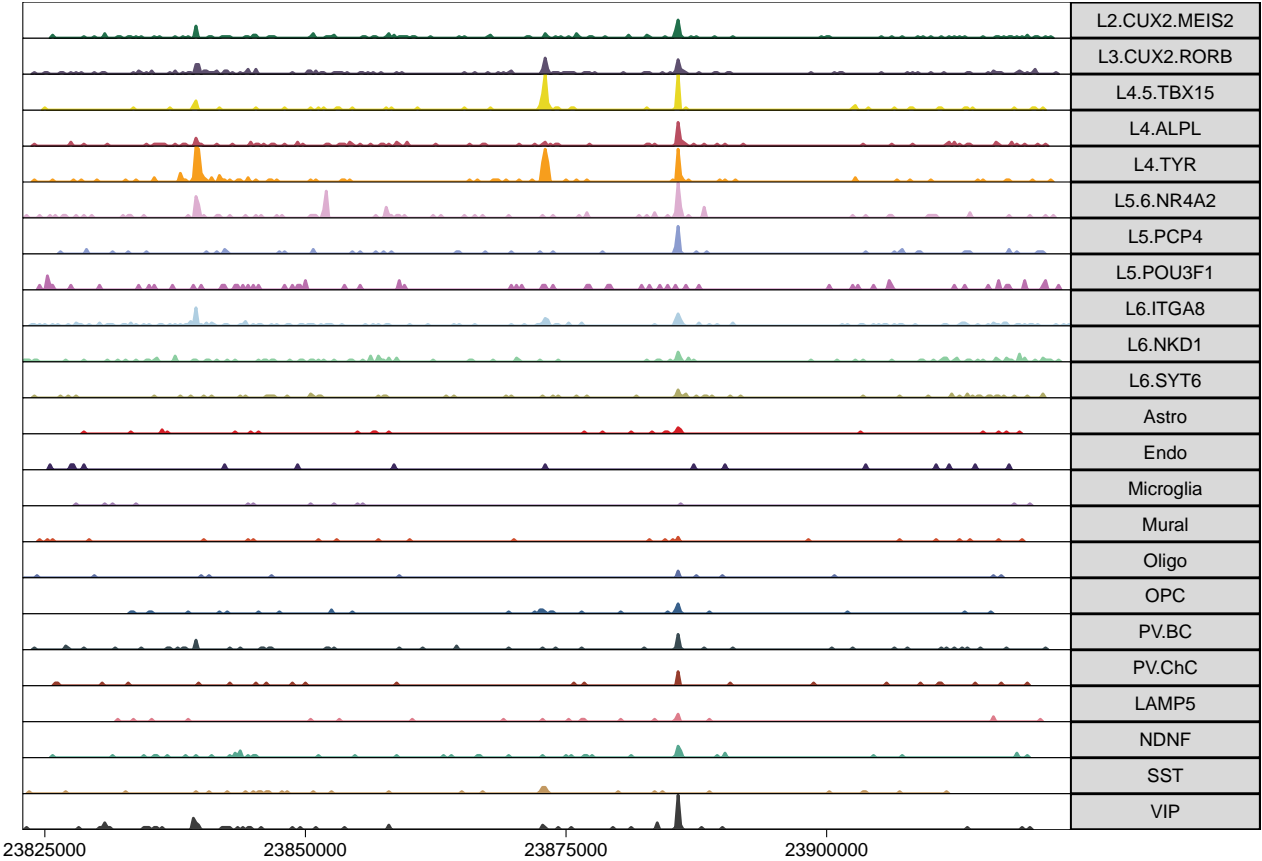


Peaks

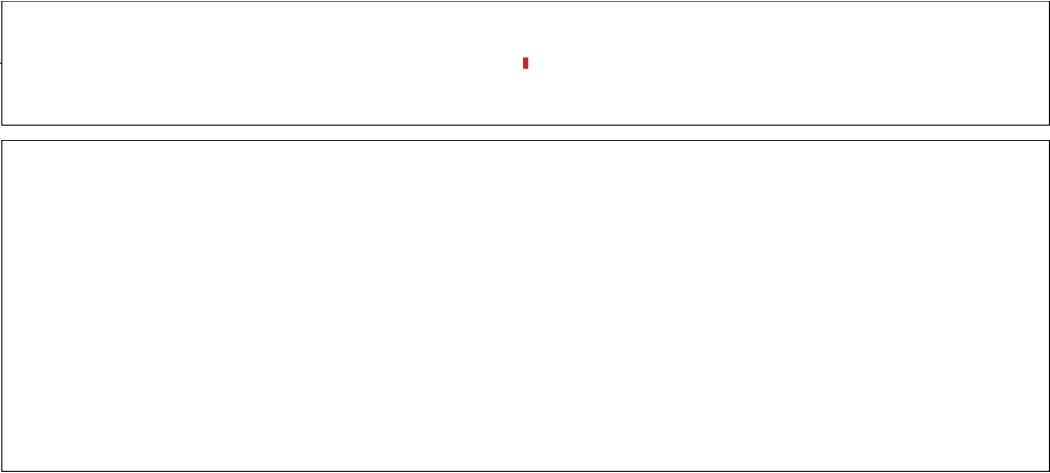


Genes

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



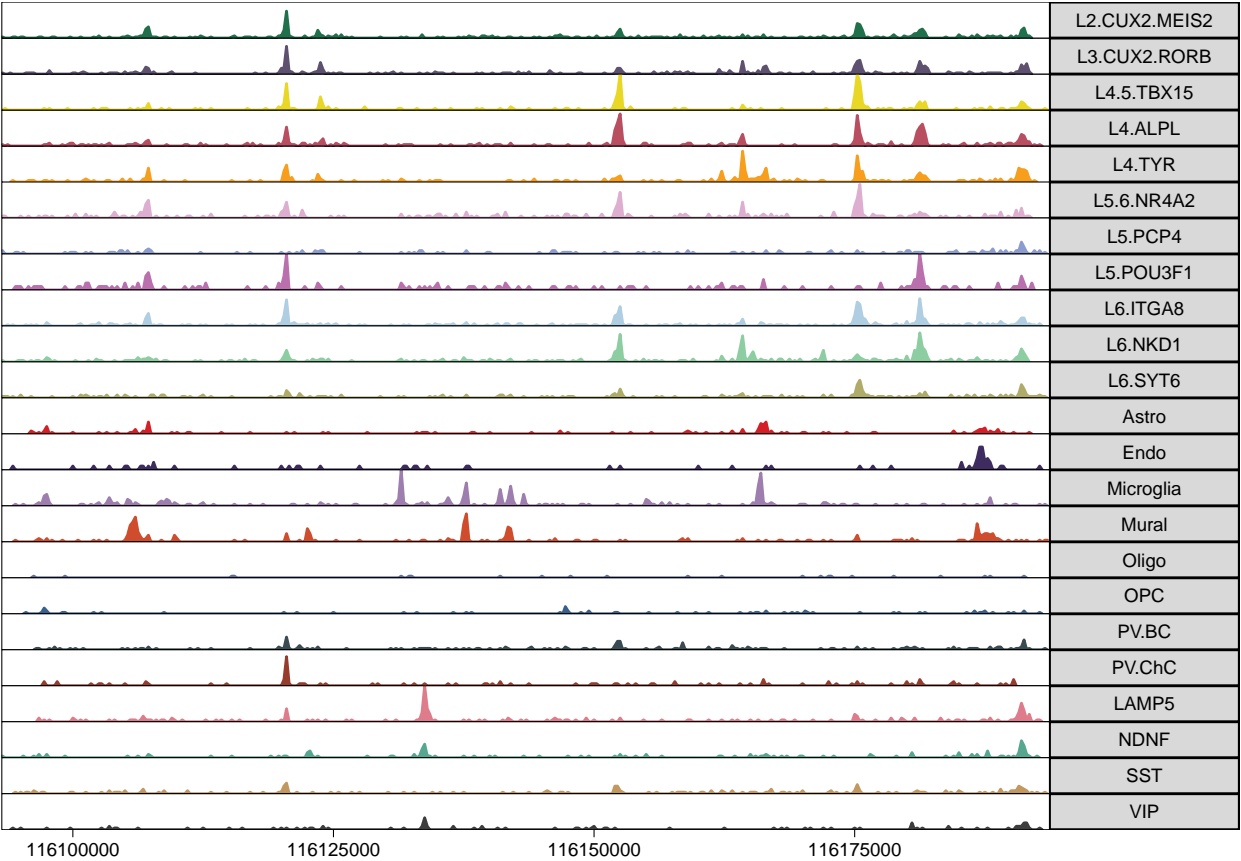
L4.5.TBX15



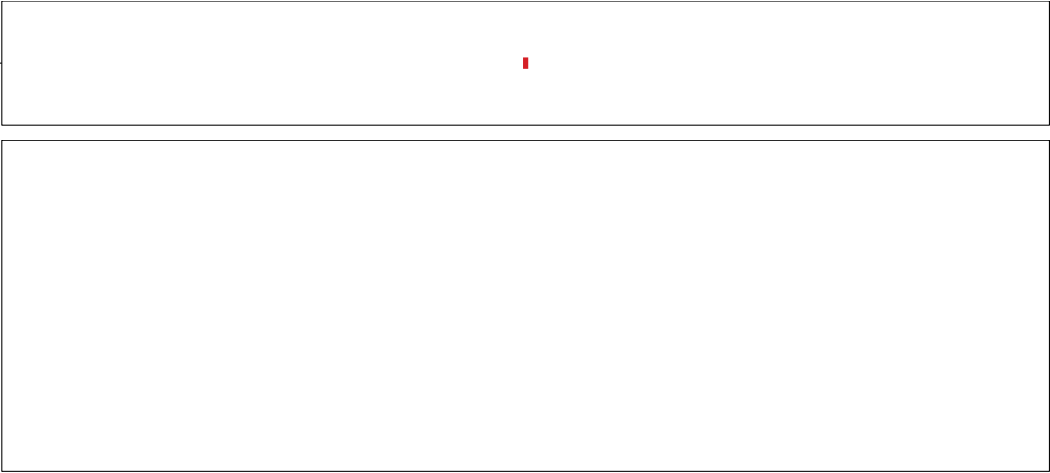
Peaks

GeneTrack

Coverage
(Norm. ATAC Signal Range (0–0.76) by ReadsInTSS)



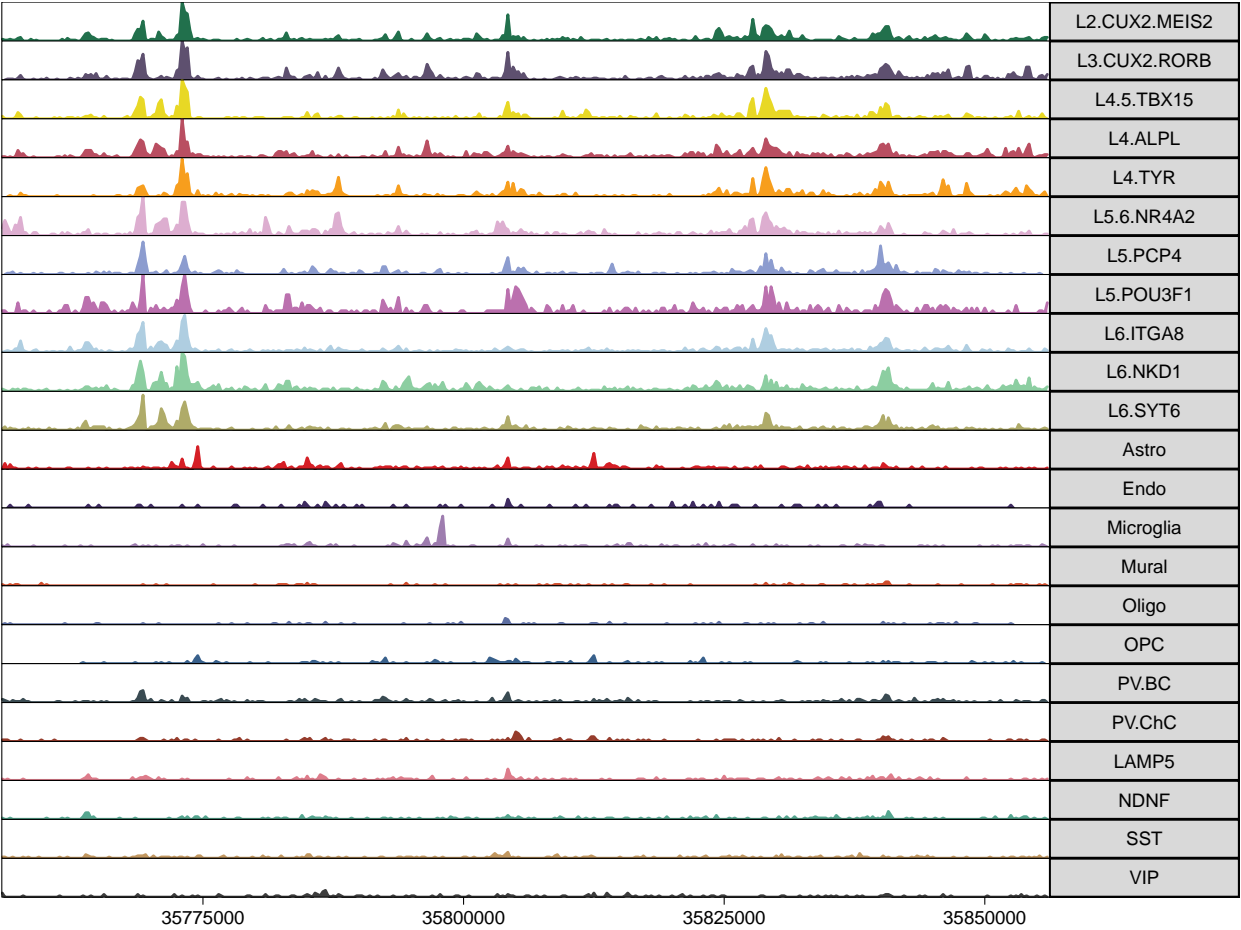
L4.5.TBX15



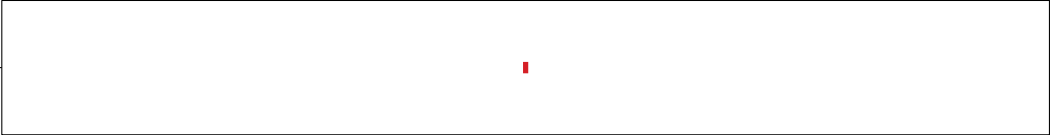
Peaks

GeneTrack

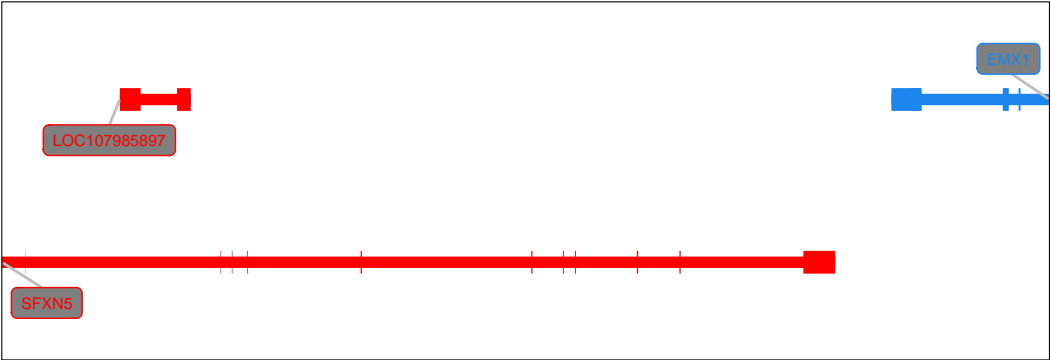
Coverage
(Norm. ATAC Signal Range (0–1.07) by ReadsInTSS)



L4.5.TBX15

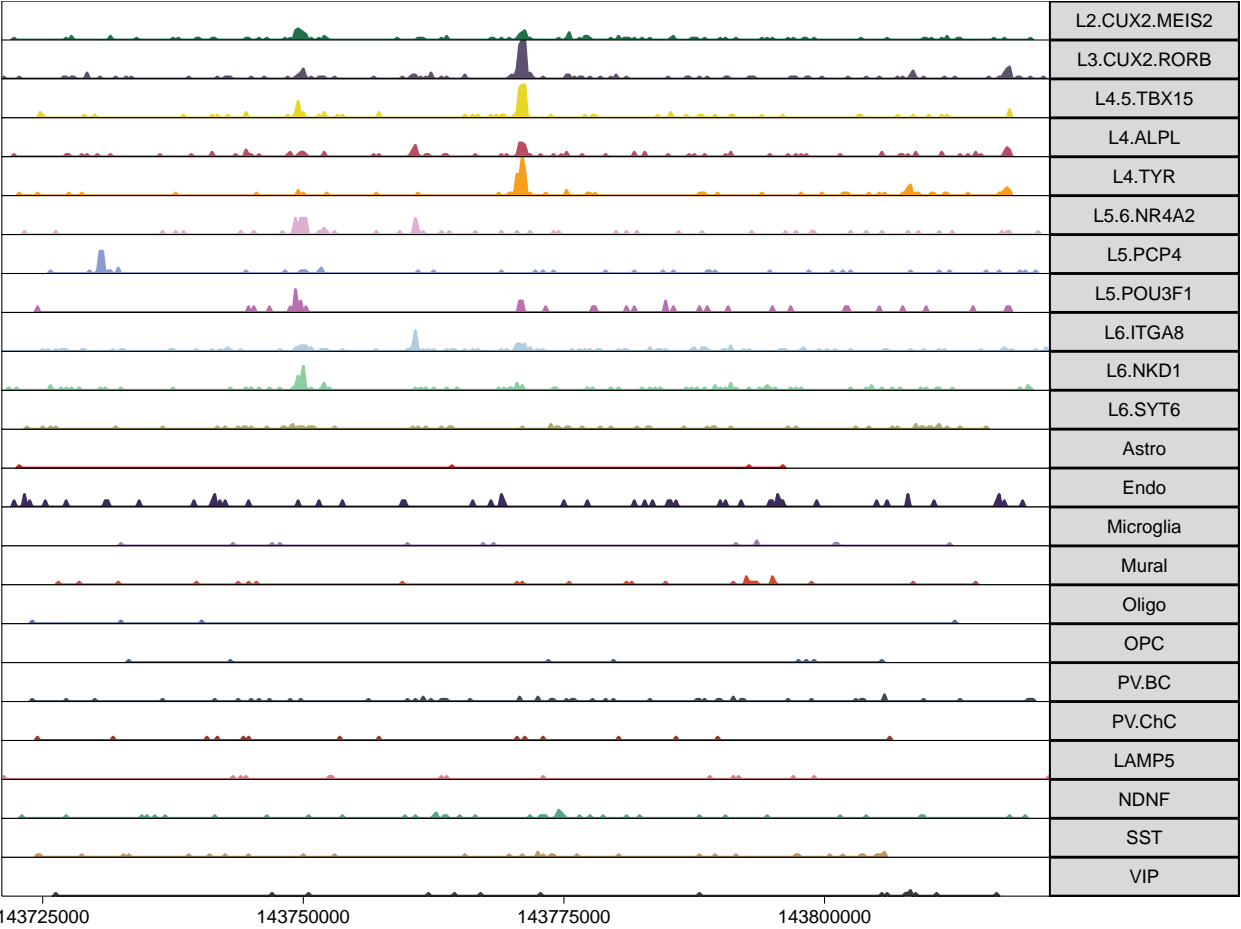


Peaks

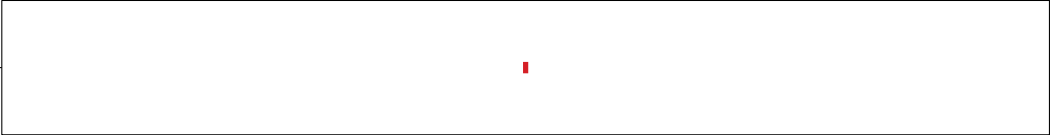


Genes

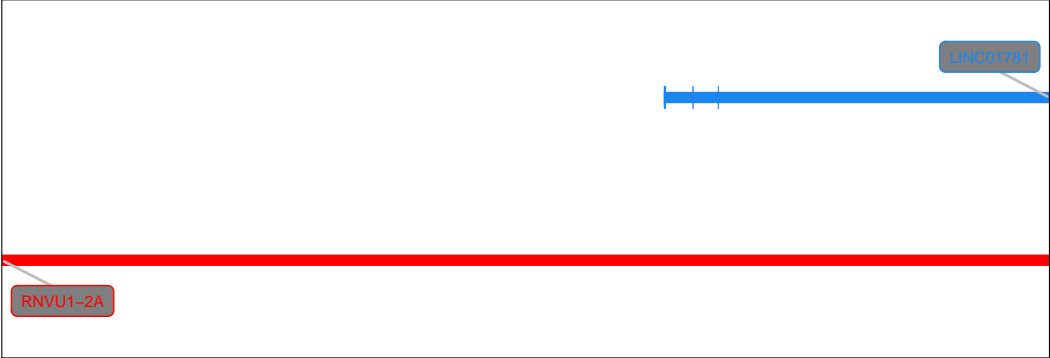
Coverage
(Norm. ATAC Signal Range (0–0.5) by ReadsInTSS)



L4.5.TBX15

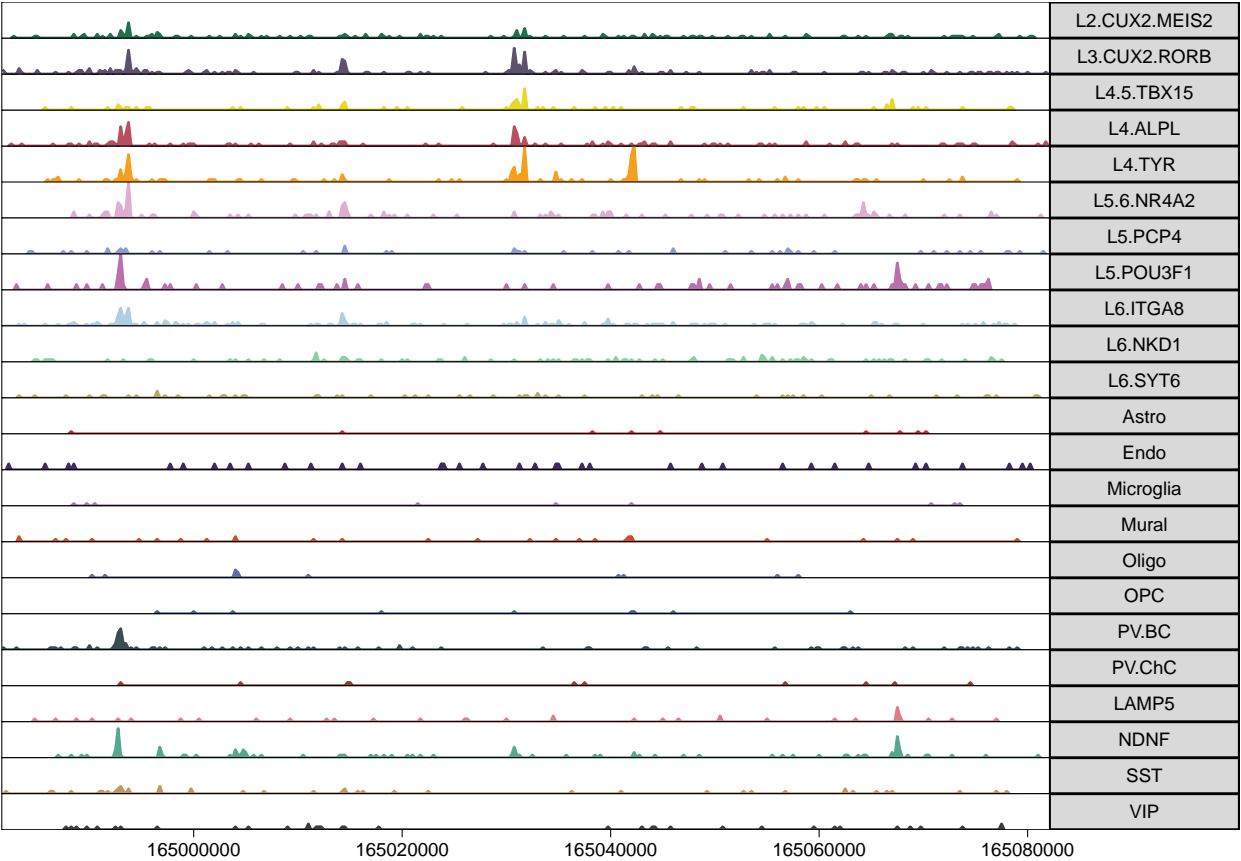


Peaks

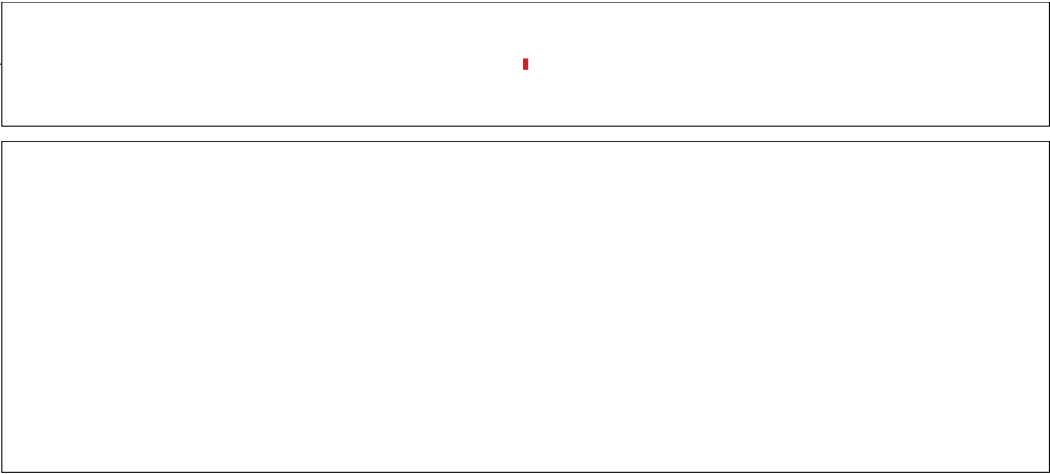


Genes

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



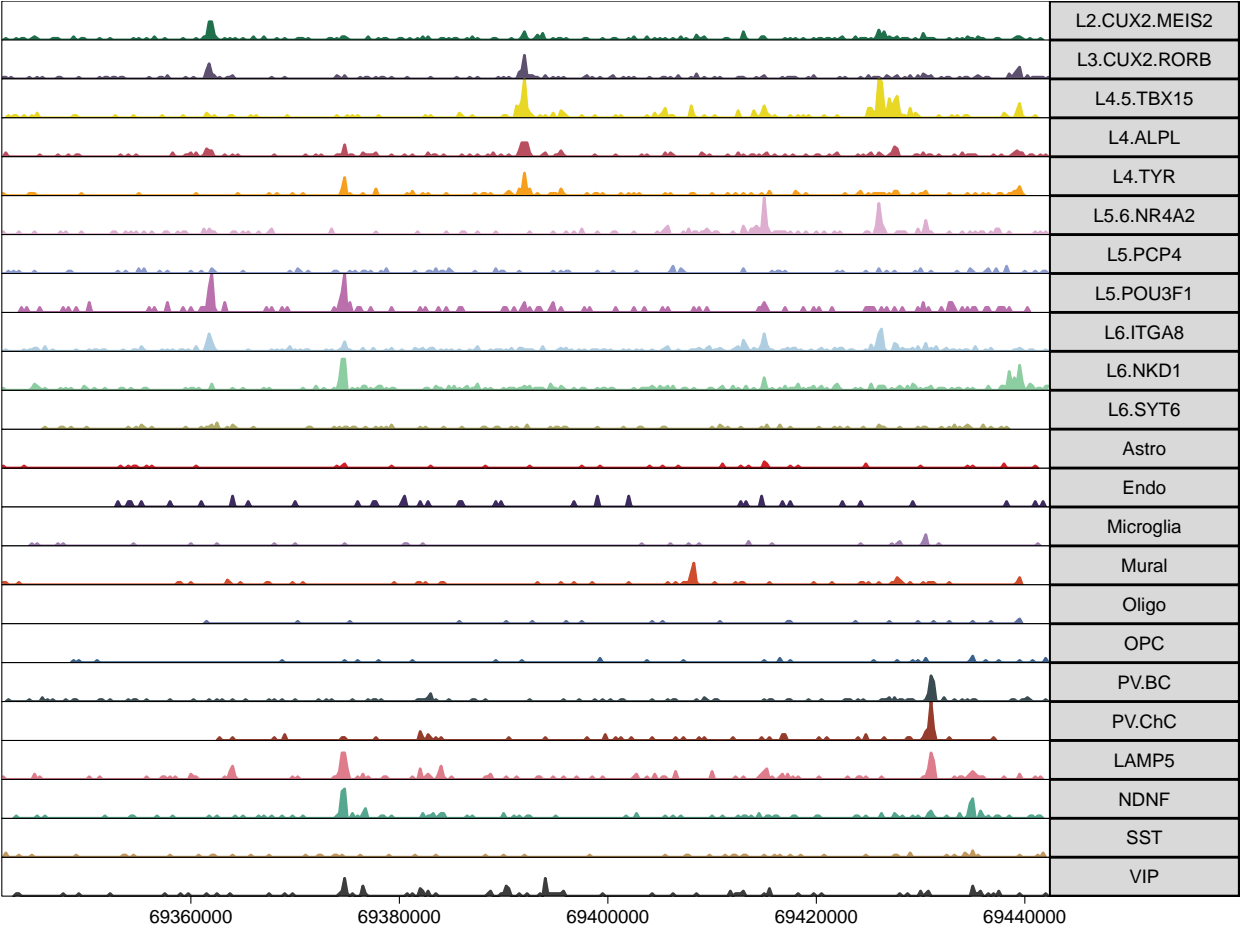
L4.5.TBX15



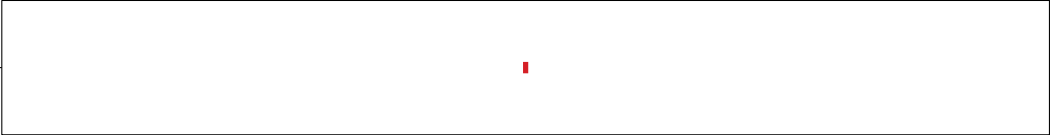
Peaks

GeneTrack

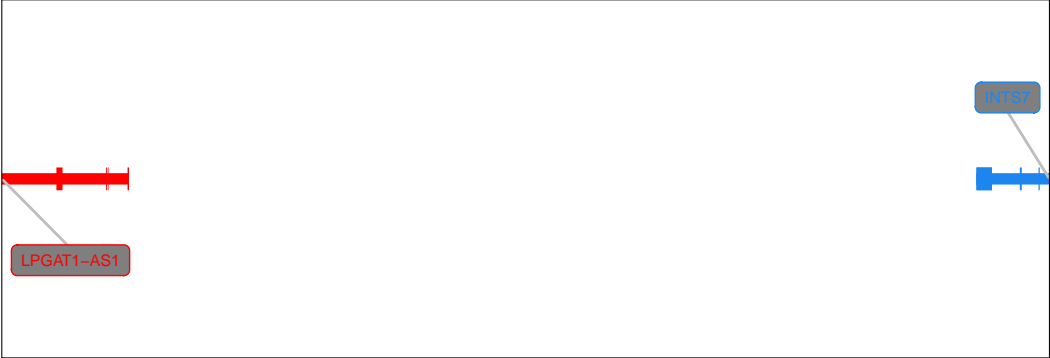
Coverage
(Norm. ATAC Signal Range (0–0.57) by ReadsInTSS)



L4.5.TBX15

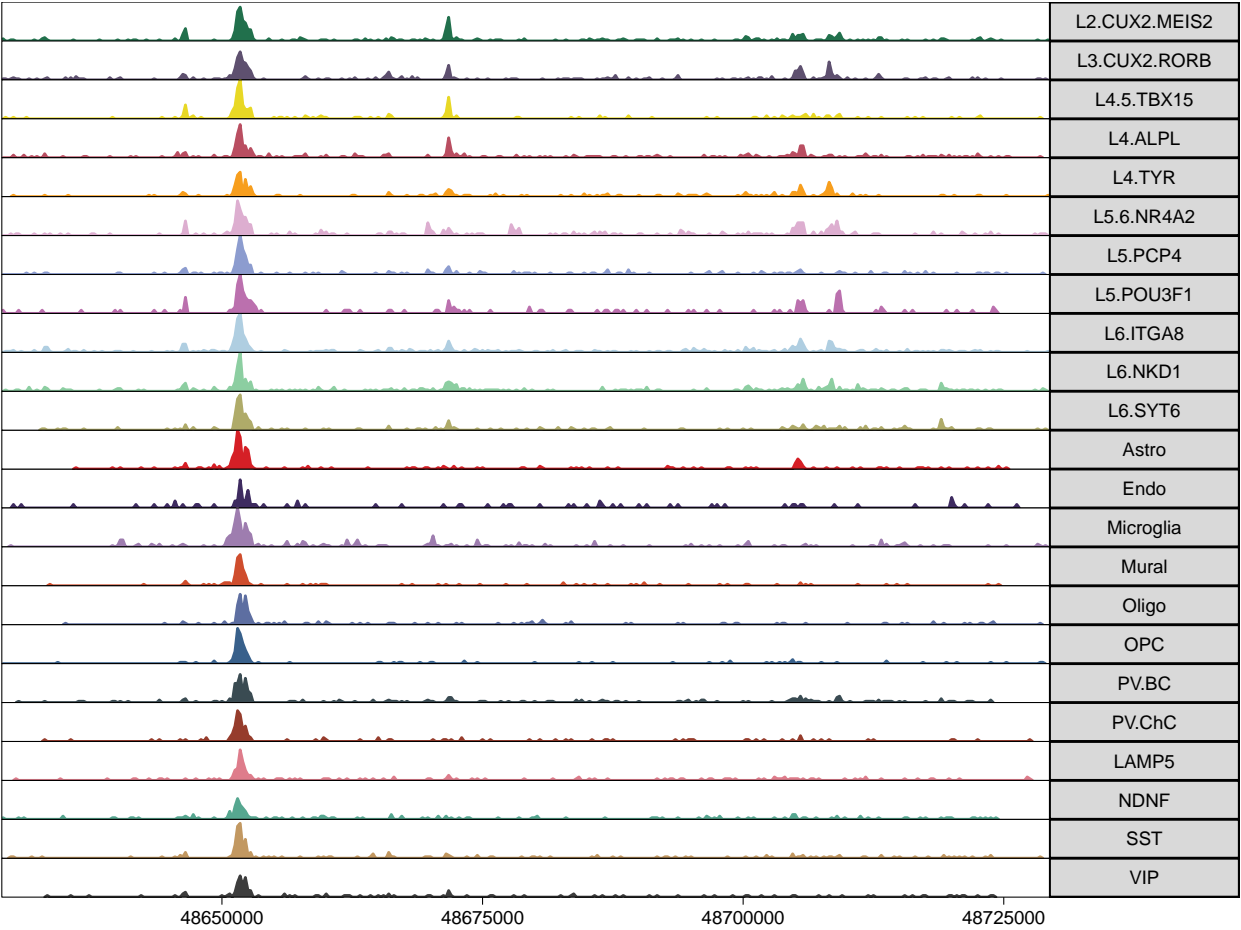


Peaks

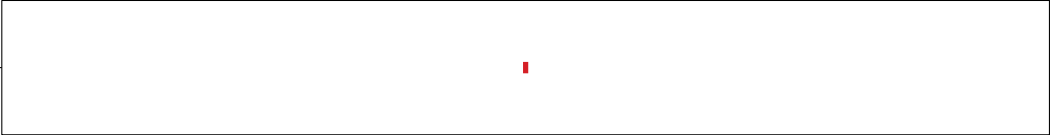


Genes

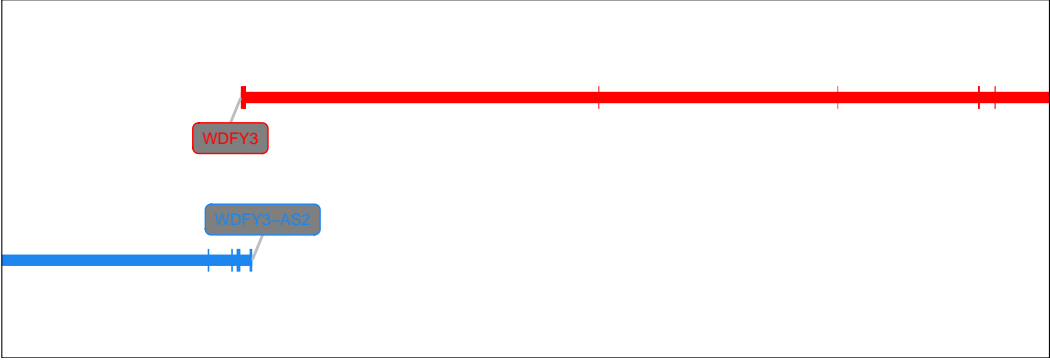
Coverage
(Norm. ATAC Signal Range (0–0.89) by ReadsInTSS)



L4.5.TBX15

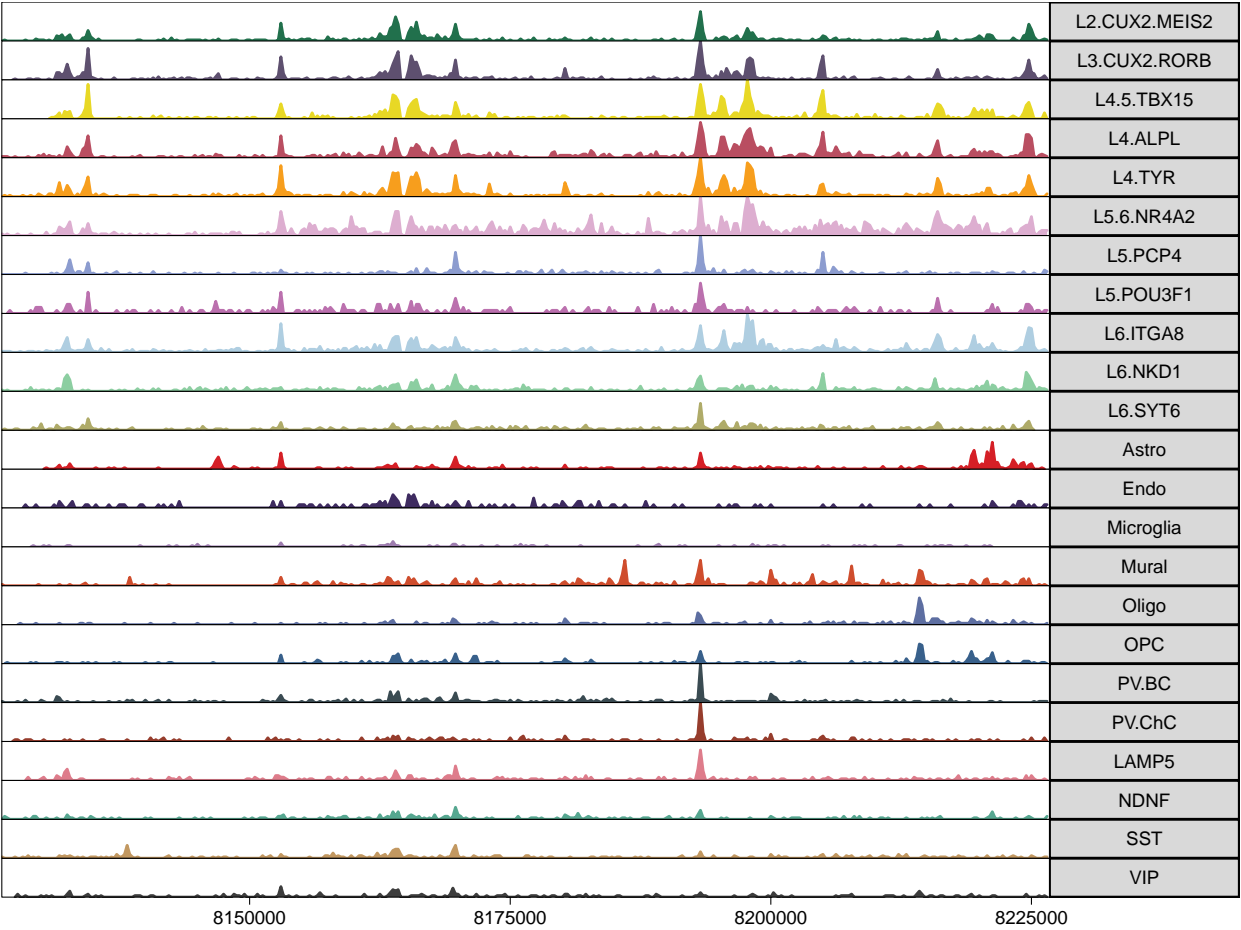


Peaks

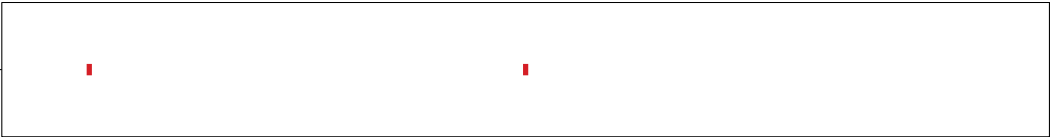


Genes

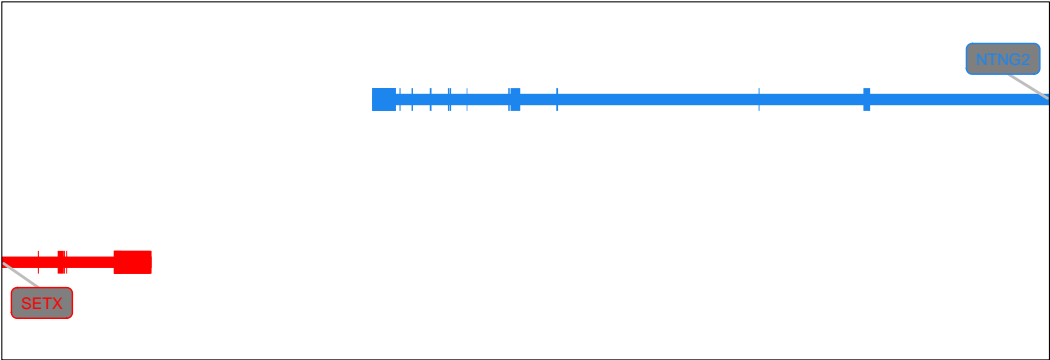
Coverage
(Norm. ATAC Signal Range (0-0.94) by ReadsInTSS)



L4.5.TBX15

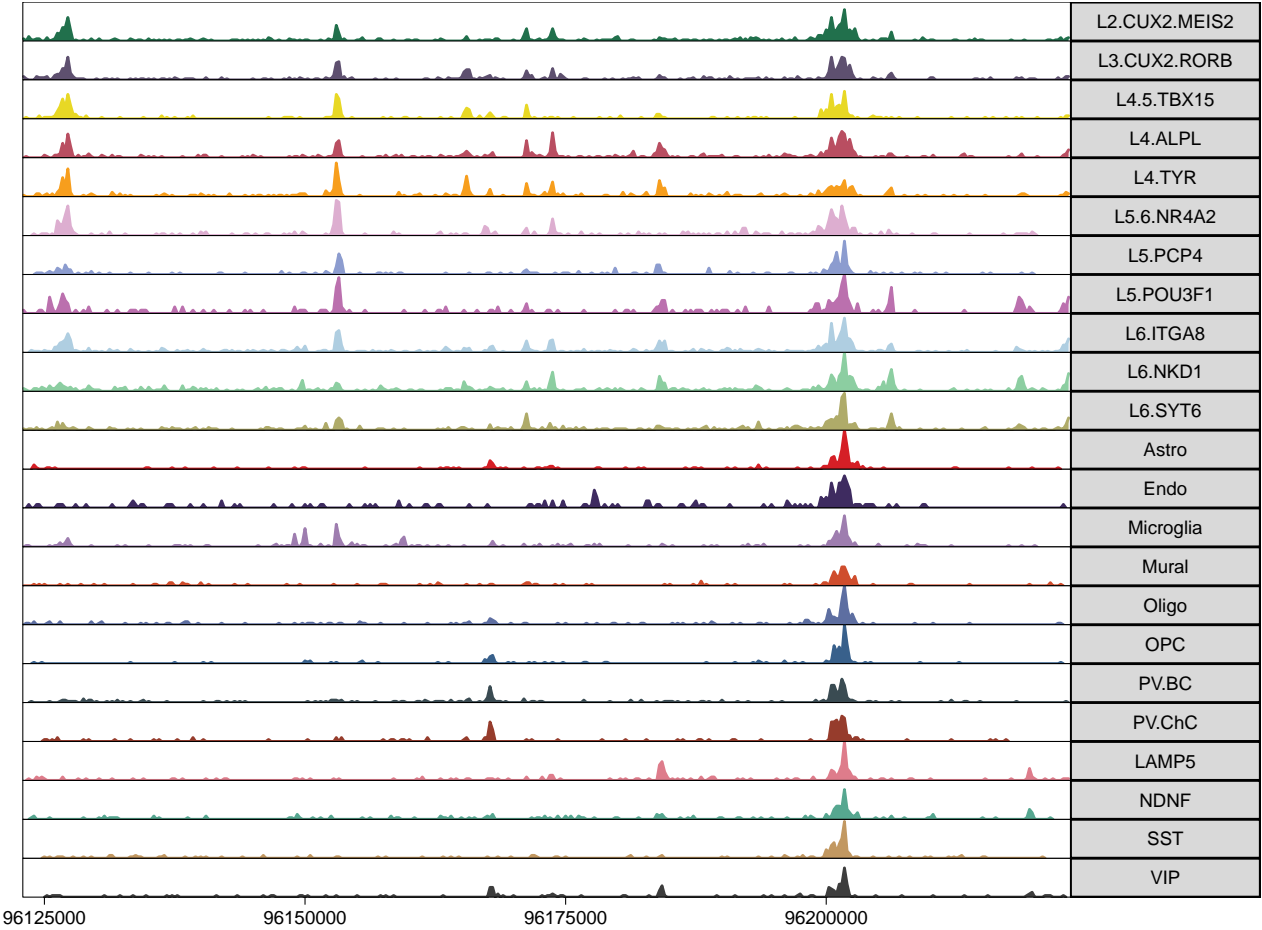


Peaks

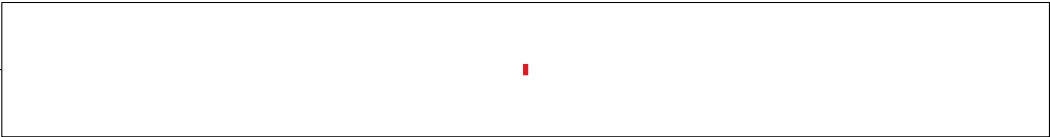


Genes

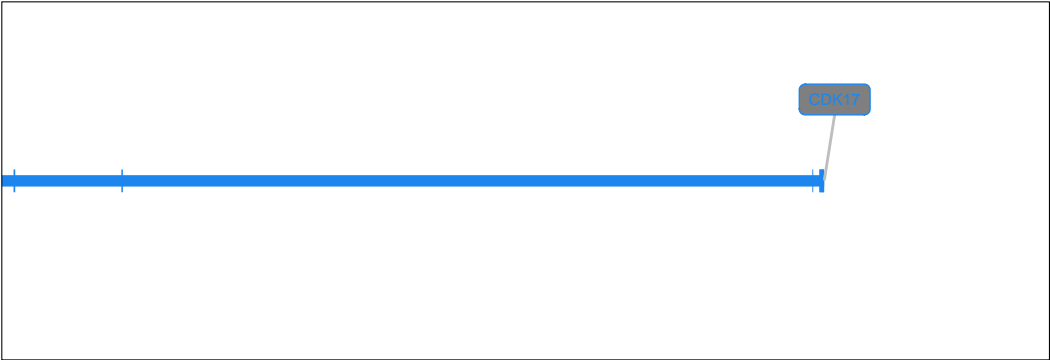
Coverage
(Norm. ATAC Signal Range (0-0.88) by ReadsInTSS)



L4.5.TBX15

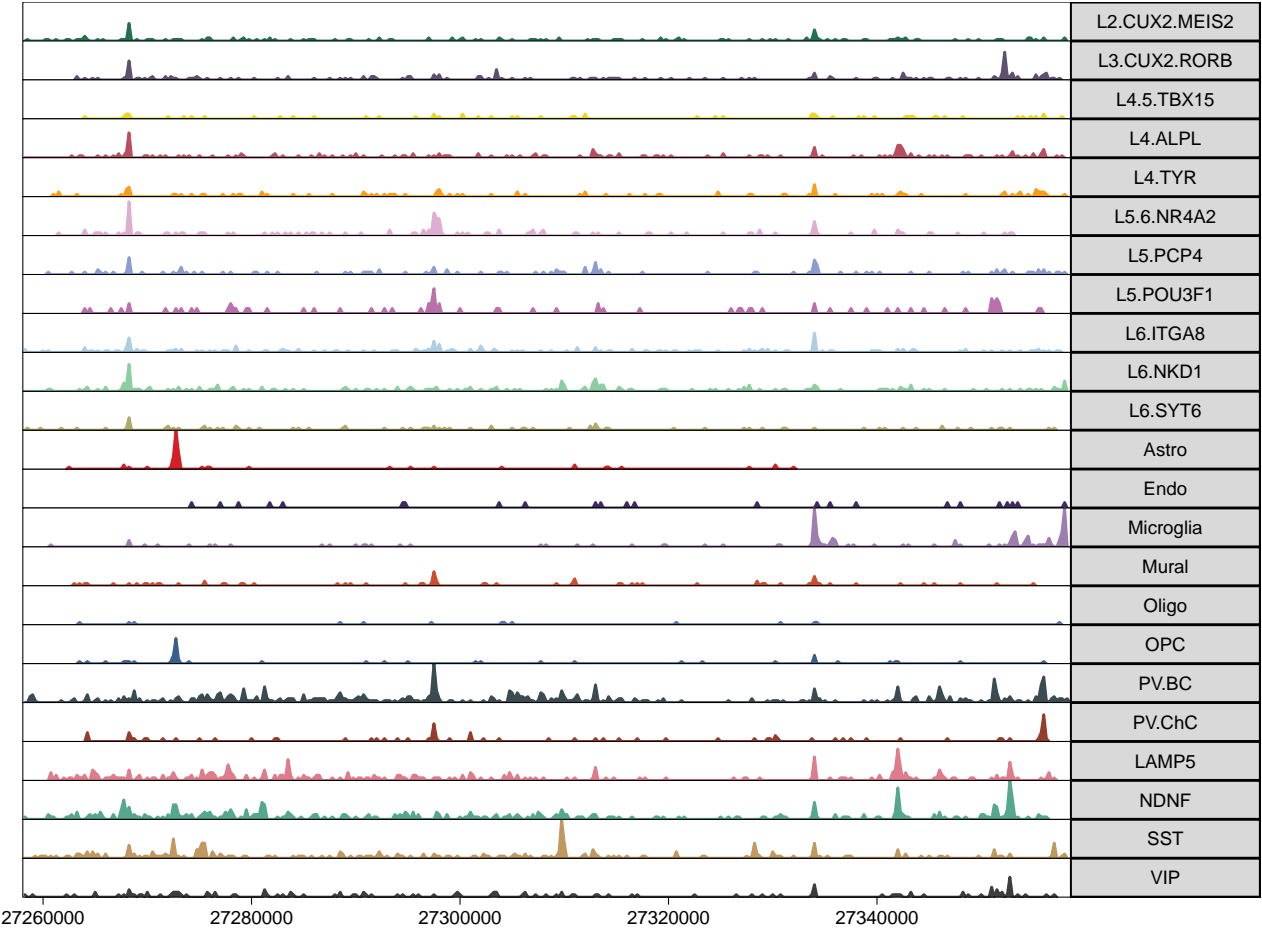


Peaks

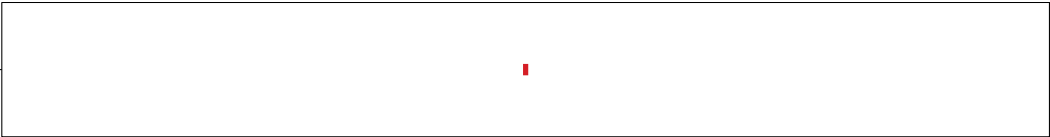


Genes

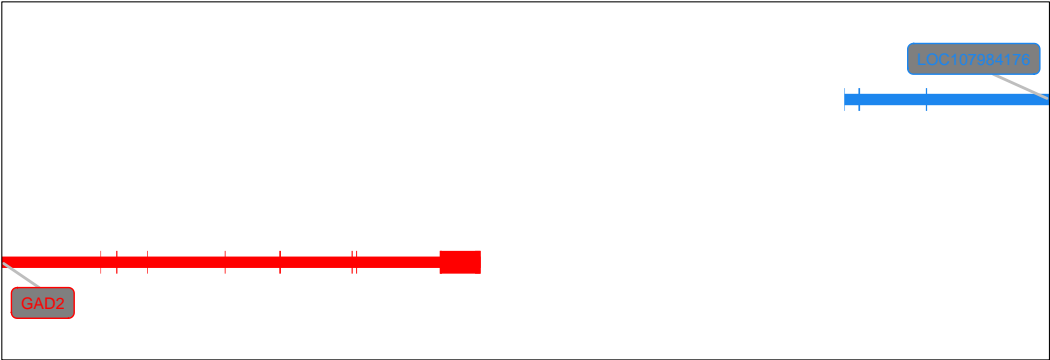
Coverage
(Norm. ATAC Signal Range (0–0.58) by ReadsInTSS)



L4.5.TBX15

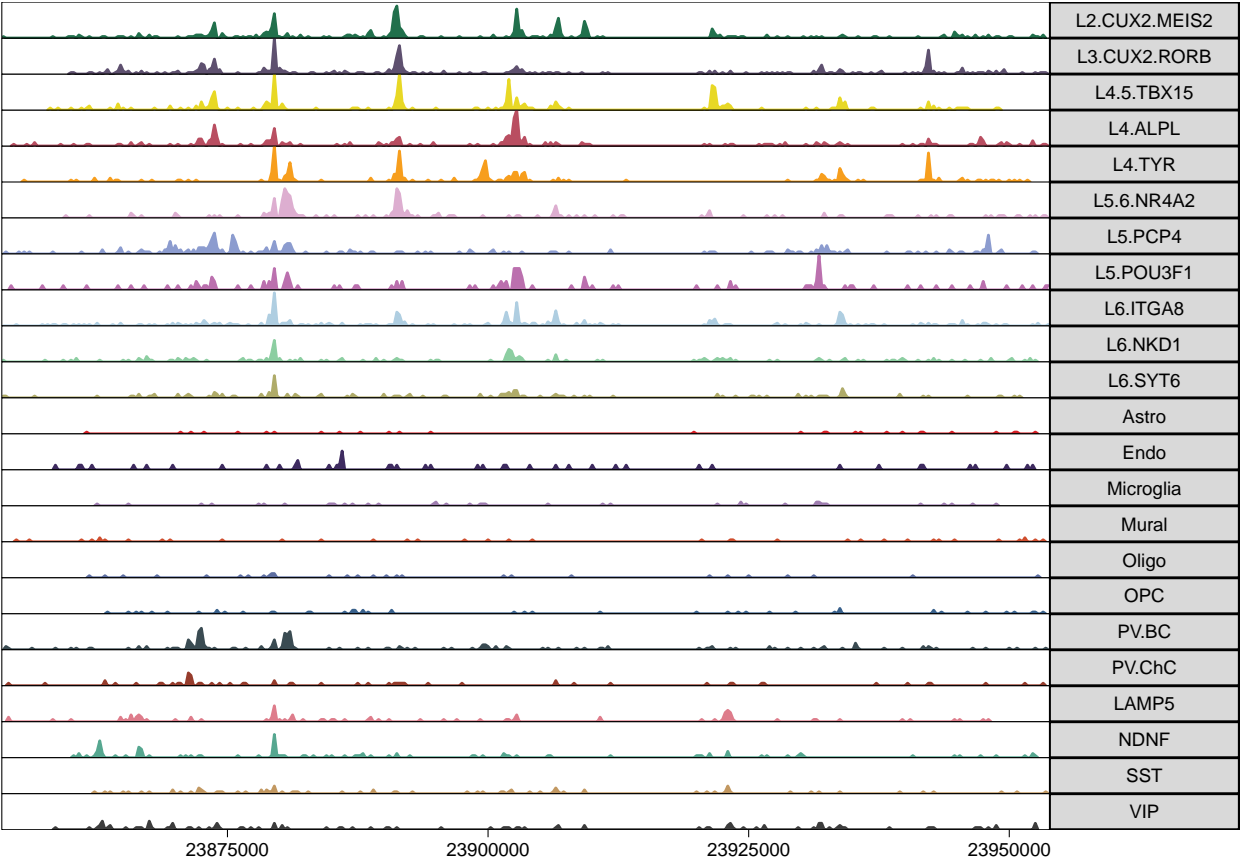


Peaks

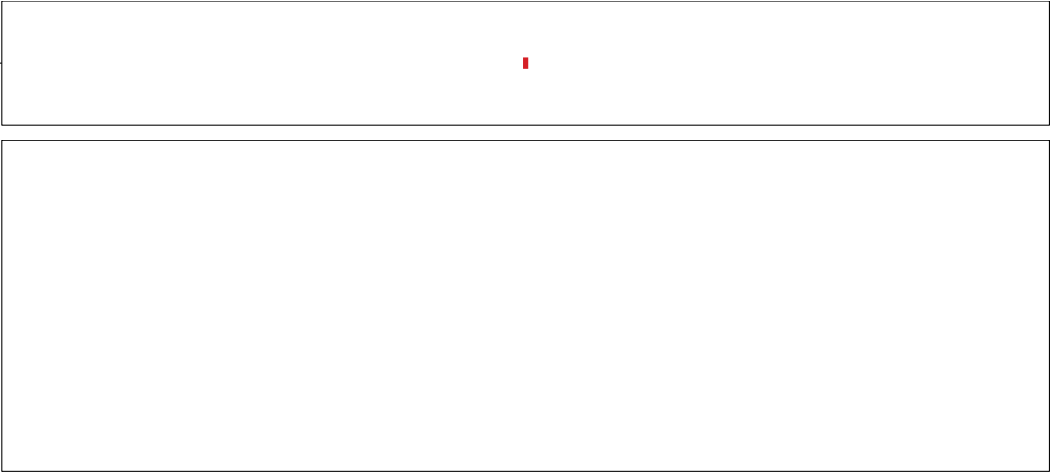


Genes

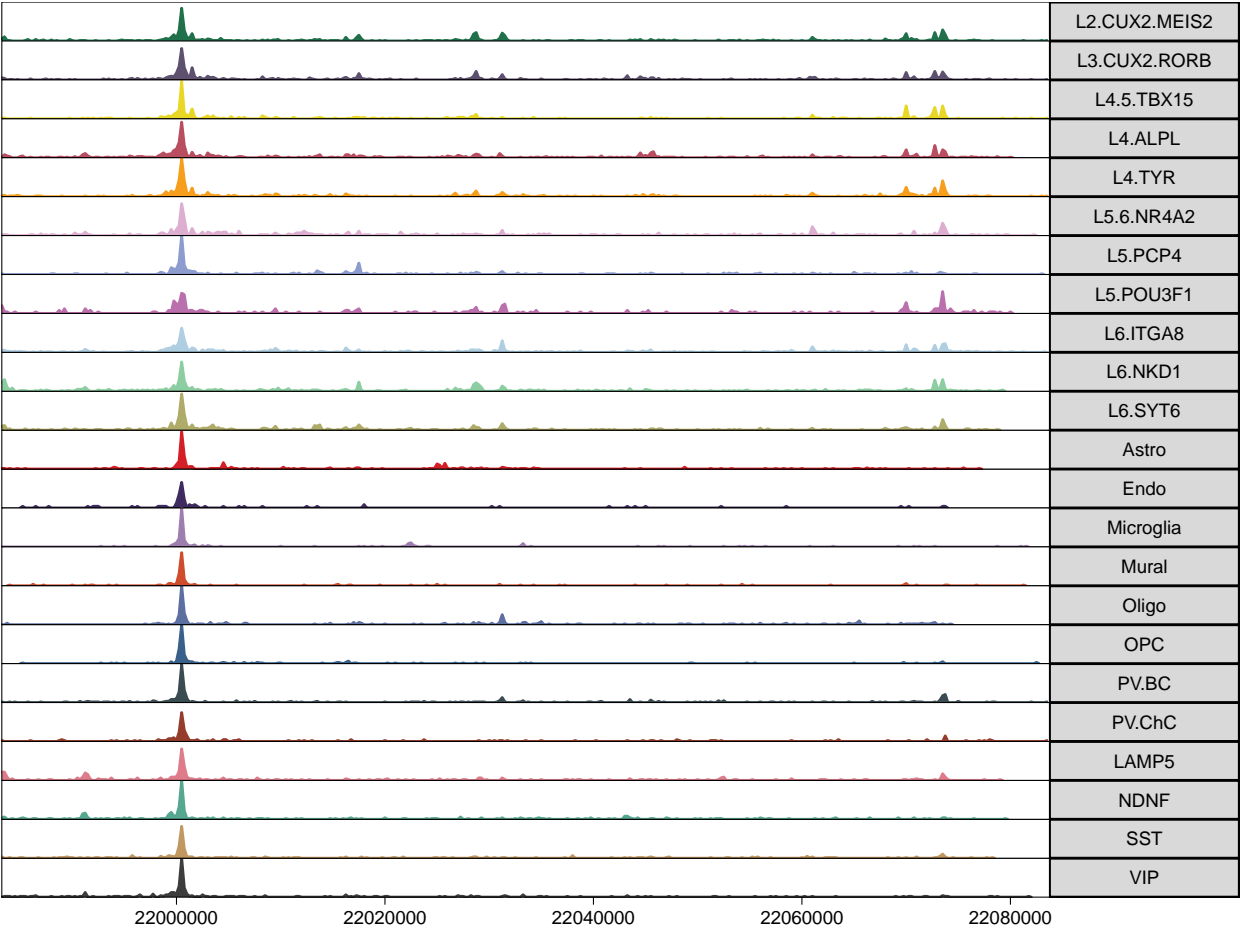
Coverage
(Norm. ATAC Signal Range (0–0.62) by ReadsInTSS)



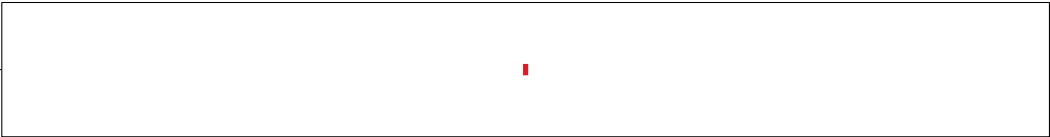
L4.5.TBX15



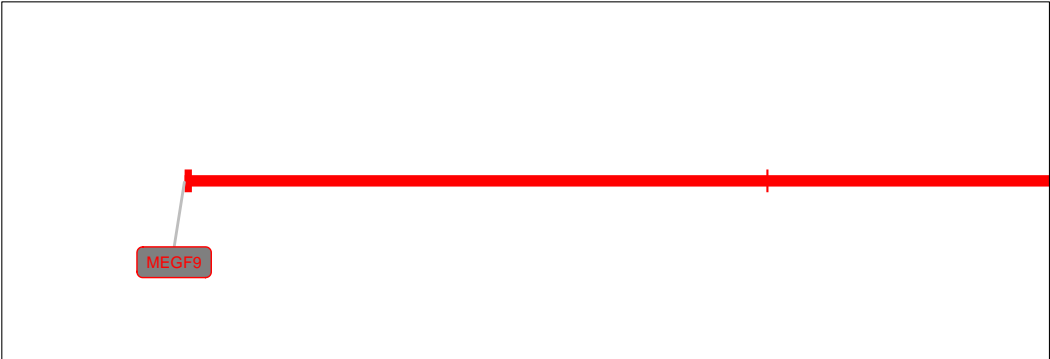
Coverage
(Norm. ATAC Signal Range (0–1.83) by ReadsInTSS)



L4.5.TBX15



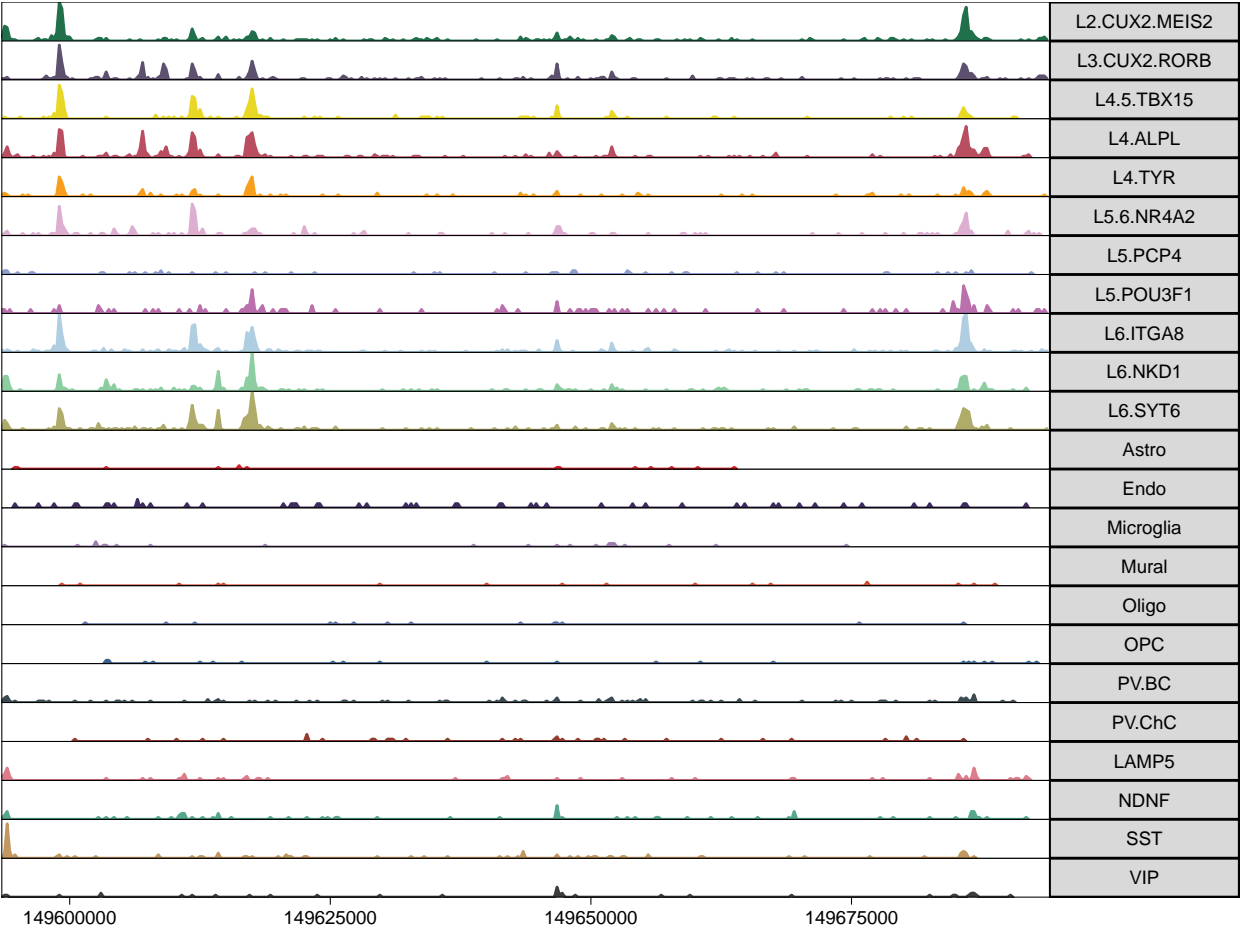
Peaks



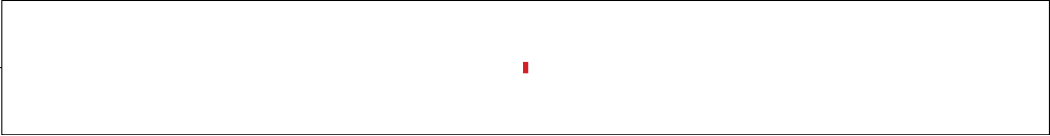
Genes

chr7:149593487–149693988 Marker Peak Tracks for: L4.5.TBX15

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



L4.5.TBX15

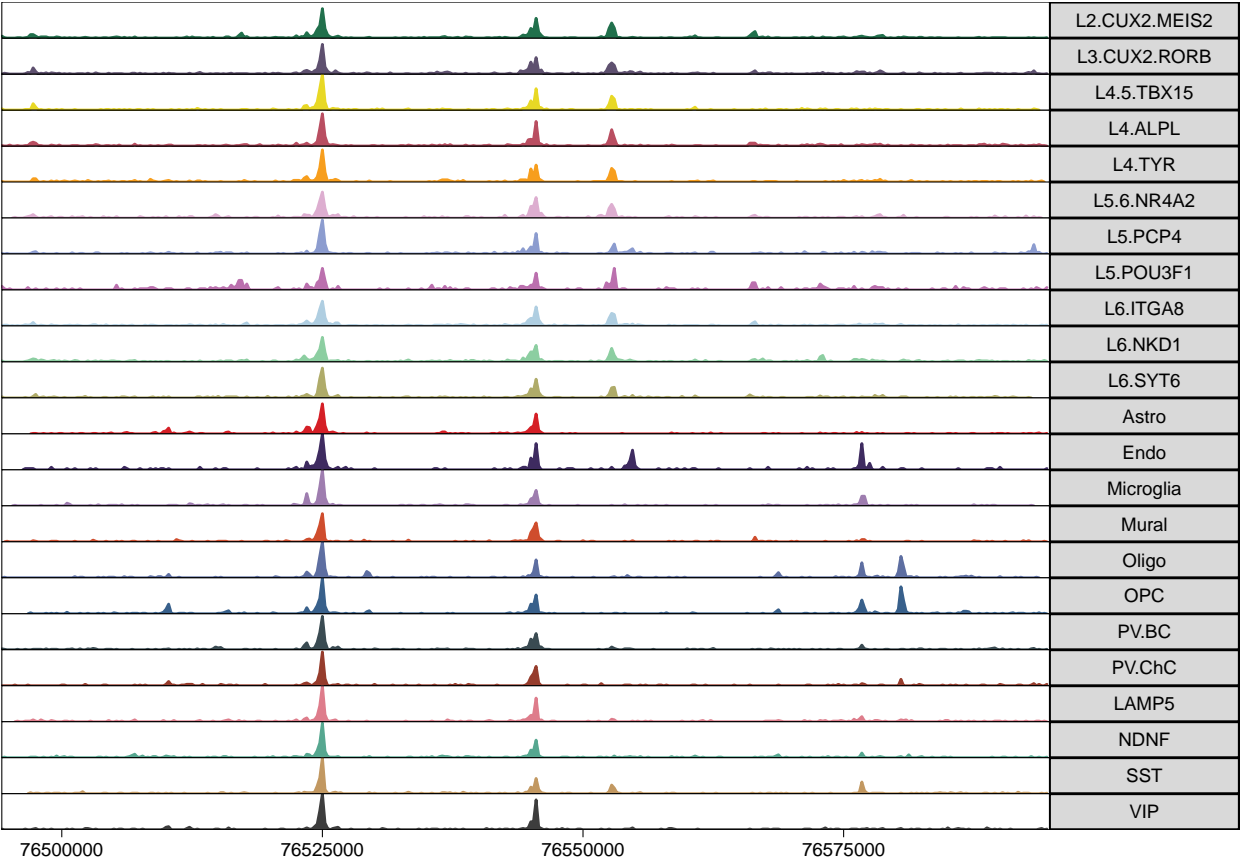


Peaks

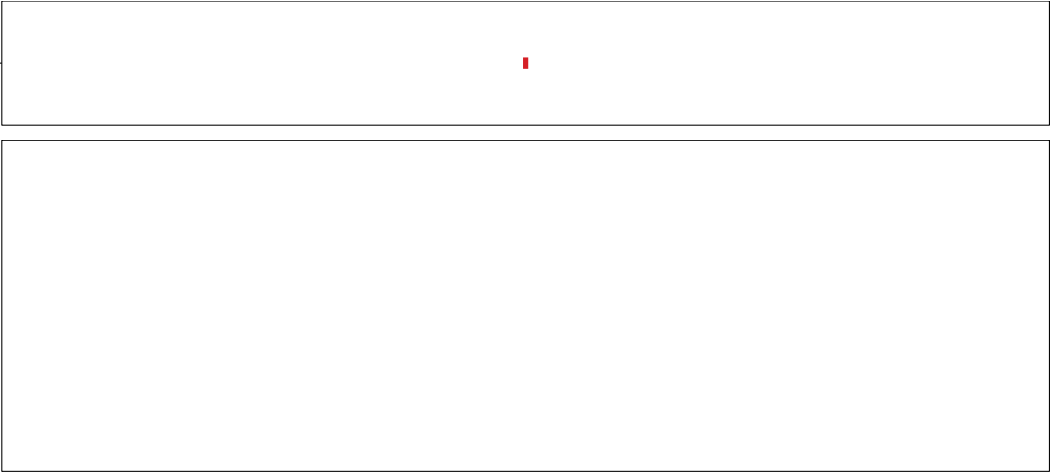


Genes

Coverage
(Norm. ATAC Signal Range (0–2.2) by ReadsInTSS)



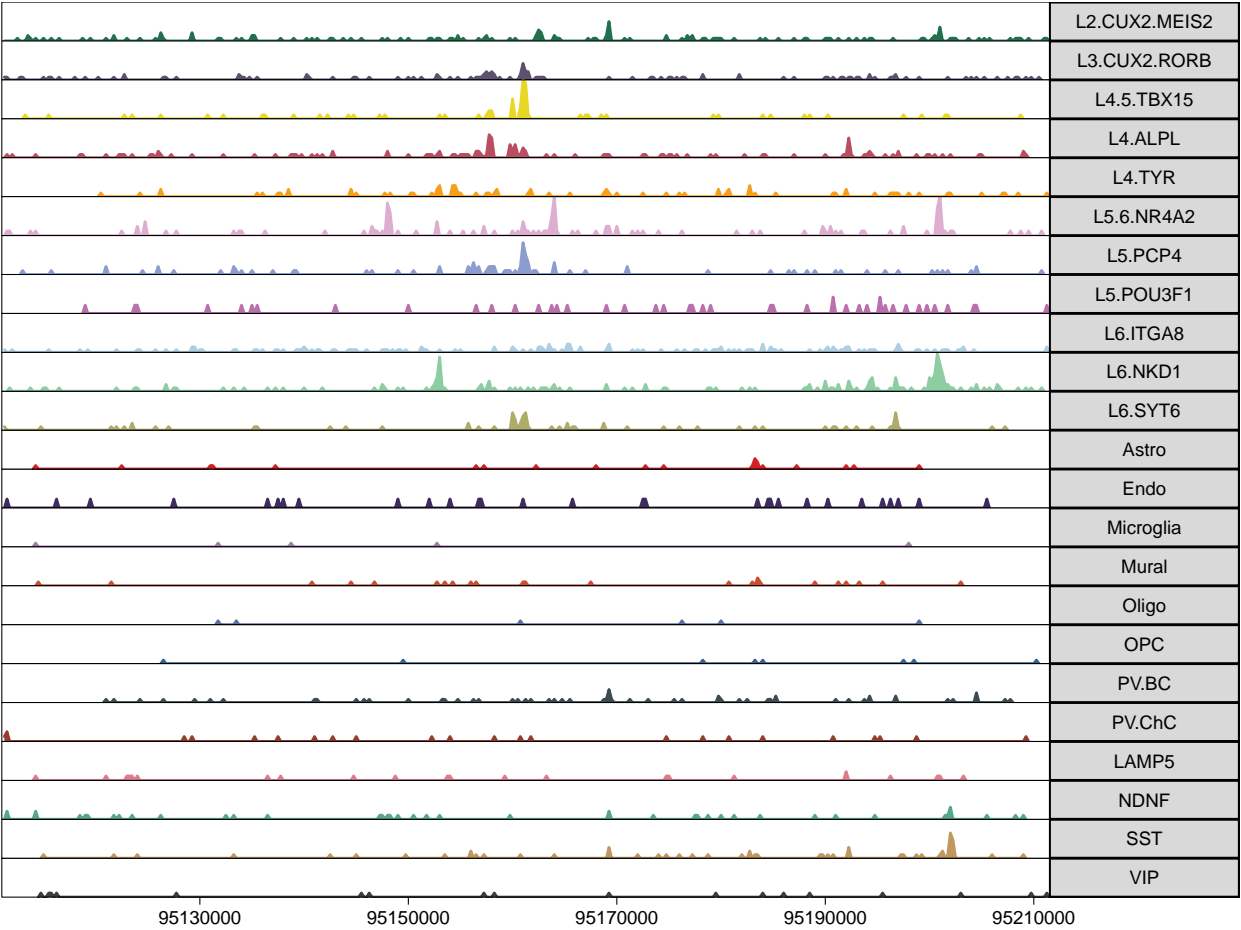
L4.5.TBX15



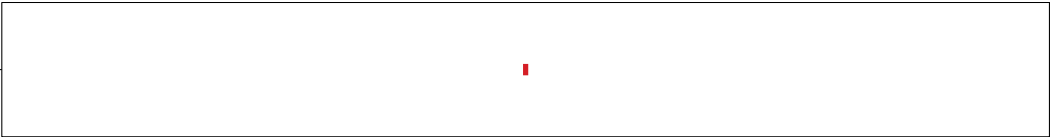
Peaks

GeneTrack

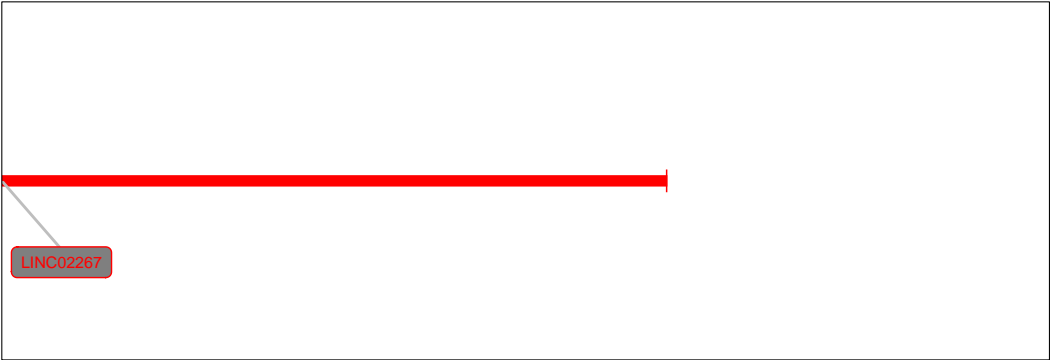
Coverage
(Norm. ATAC Signal Range (0–0.35) by ReadsInTSS)



L4.5.TBX15

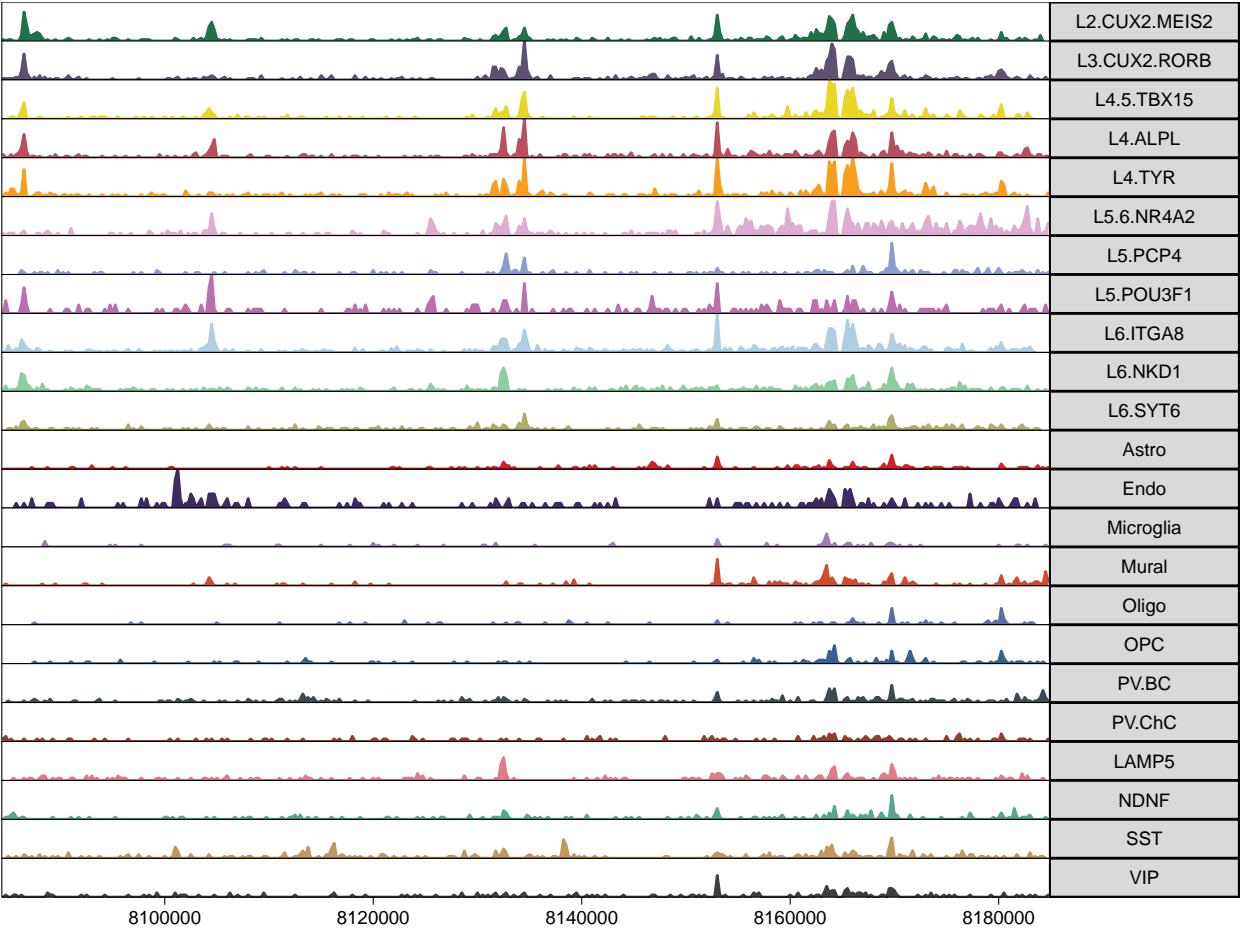


Peaks

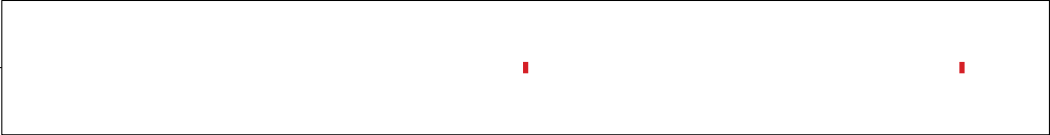


Genes

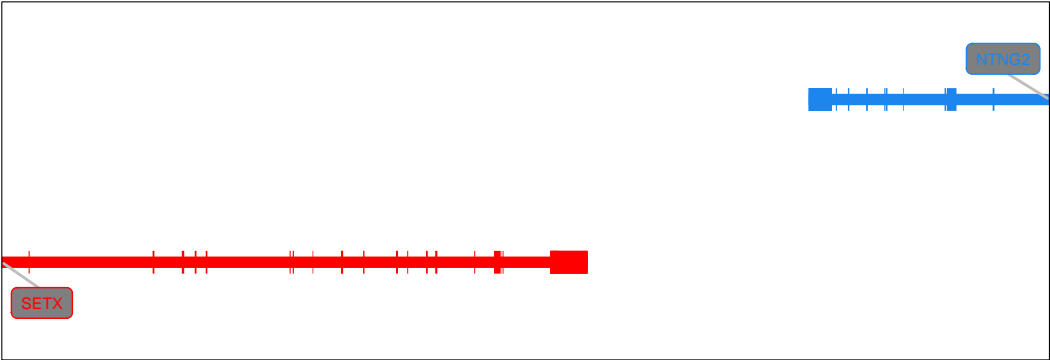
Coverage
(Norm. ATAC Signal Range (0–0.67) by ReadsInTSS)



L4.5.TBX15



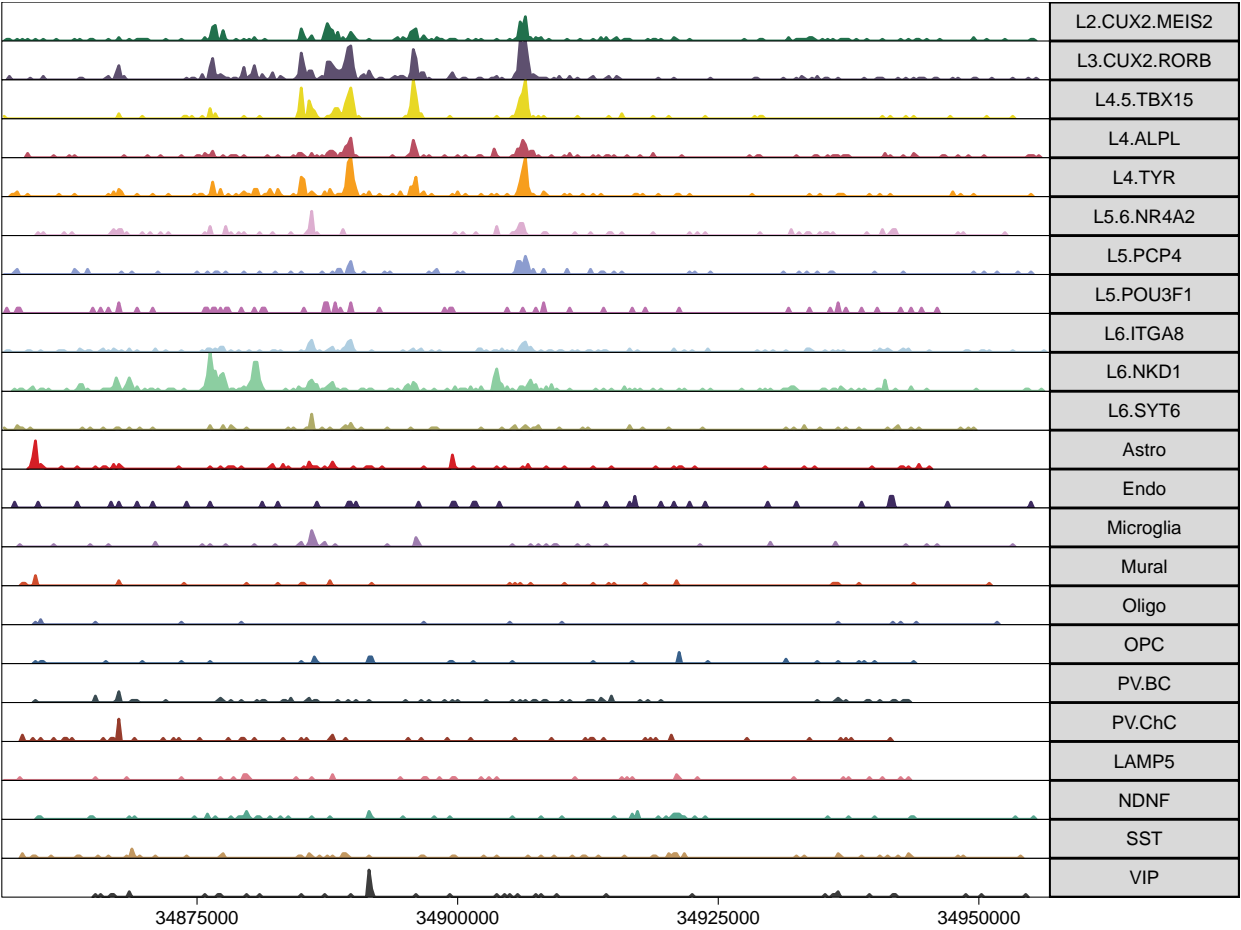
Peaks



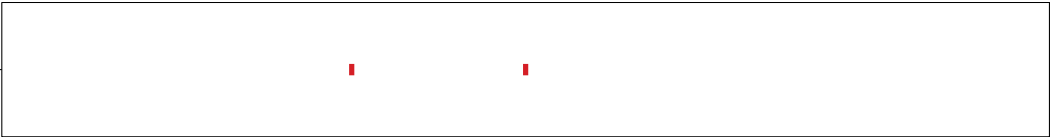
Genes

chr1:34856271–34956772 Marker Peak Tracks for: L4.5.TBX15

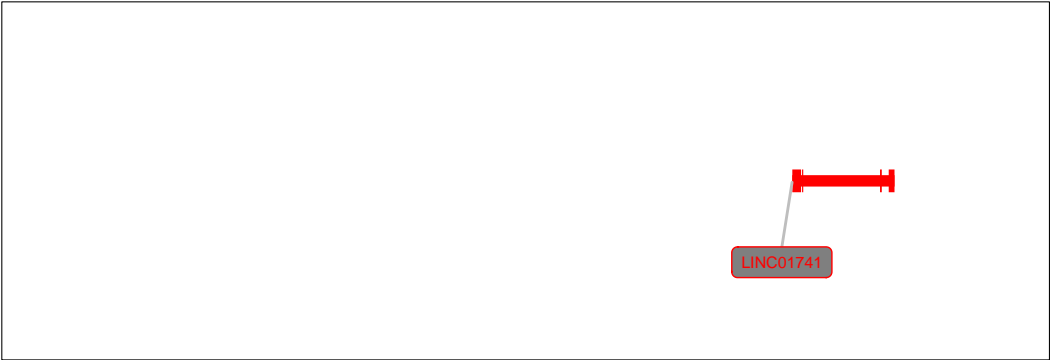
Coverage
(Norm. ATAC Signal Range (0–0.54) by ReadsInTSS)



L4.5.TBX15

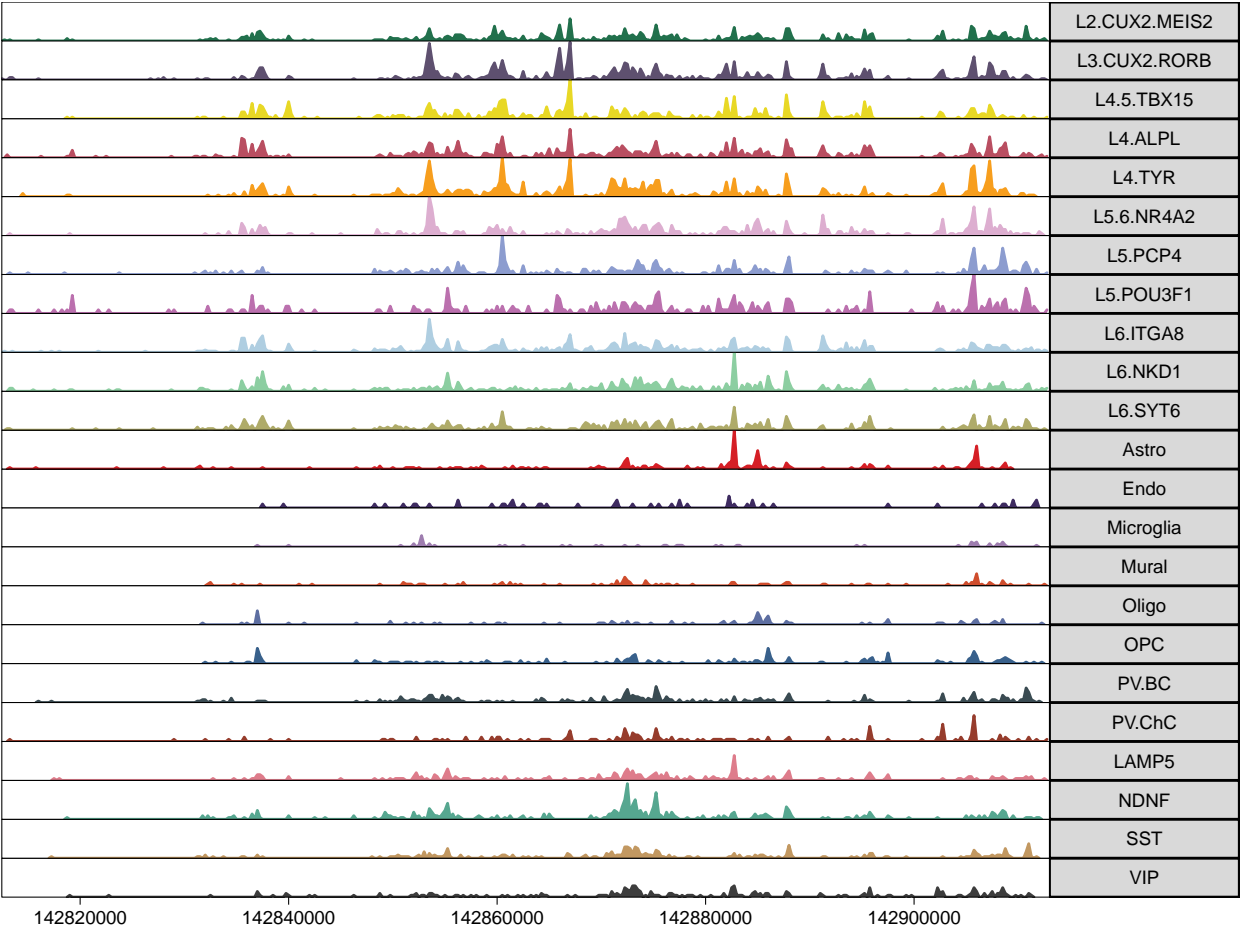


Peaks

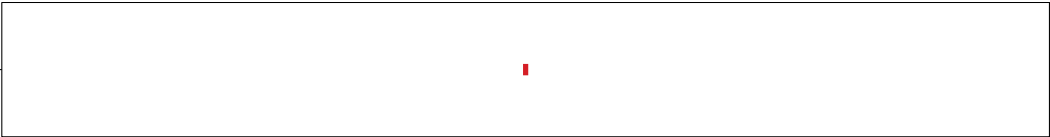


Genes

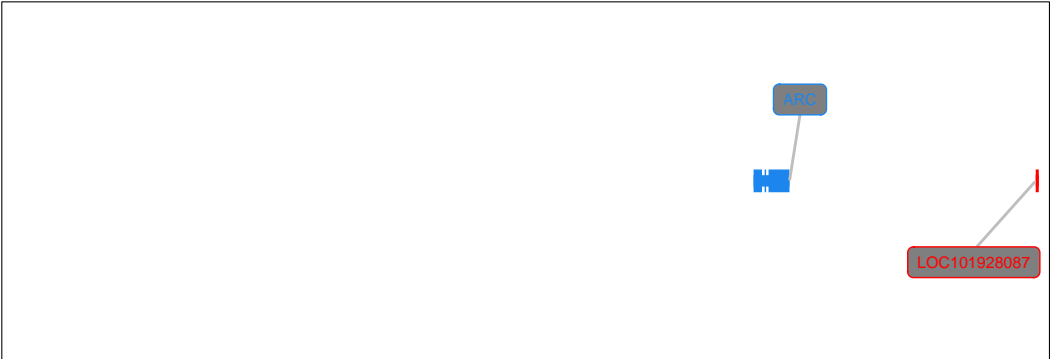
Coverage
(Norm. ATAC Signal Range (0–0.79) by ReadsInTSS)



L4.5.TBX15

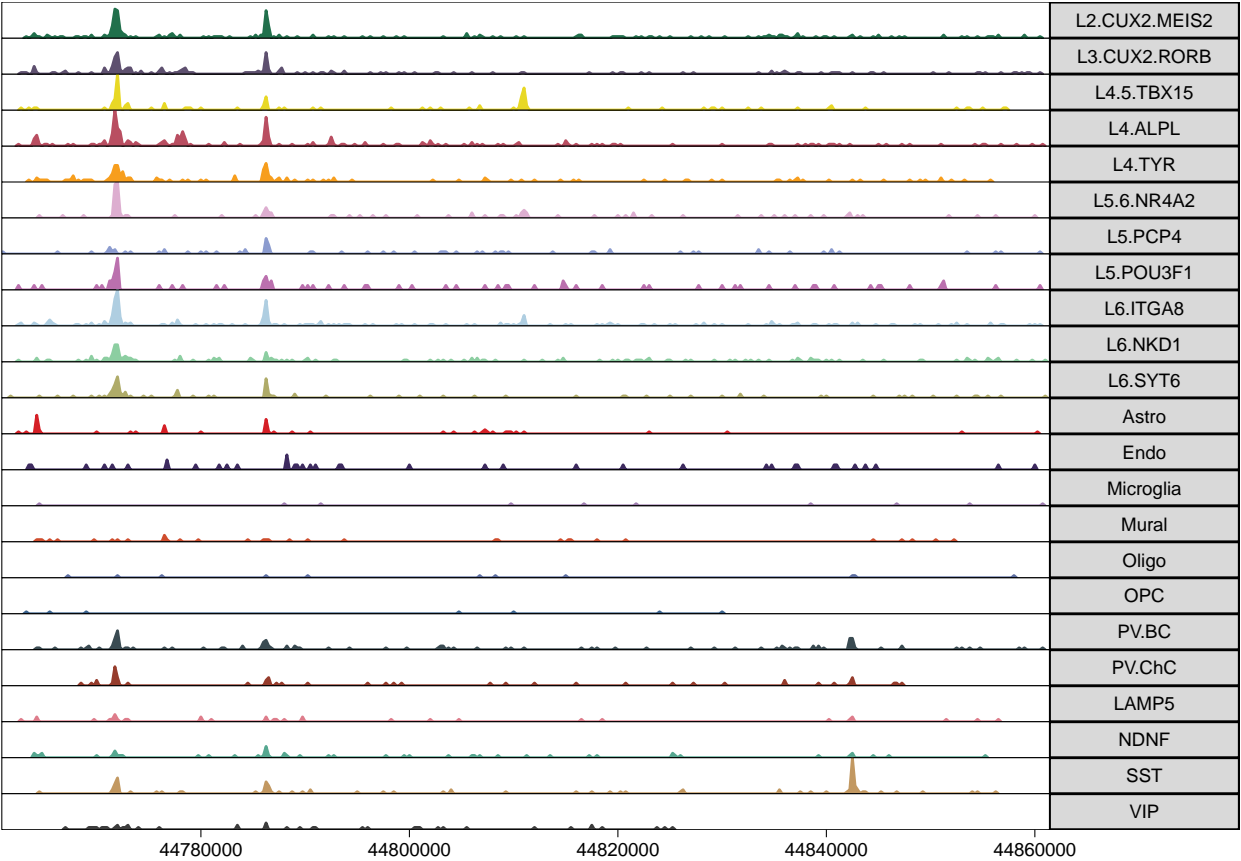


Peaks

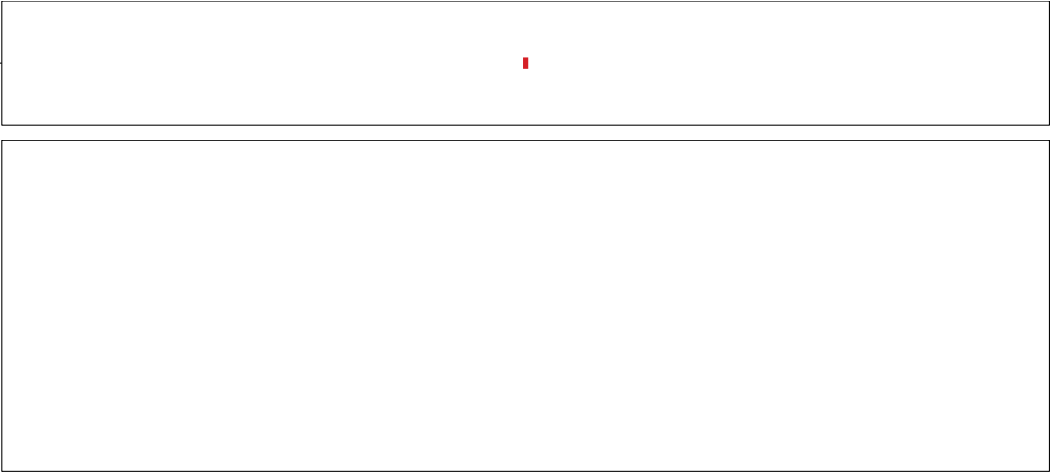


Genes

Coverage
(Norm. ATAC Signal Range (0-0.58) by ReadsInTSS)

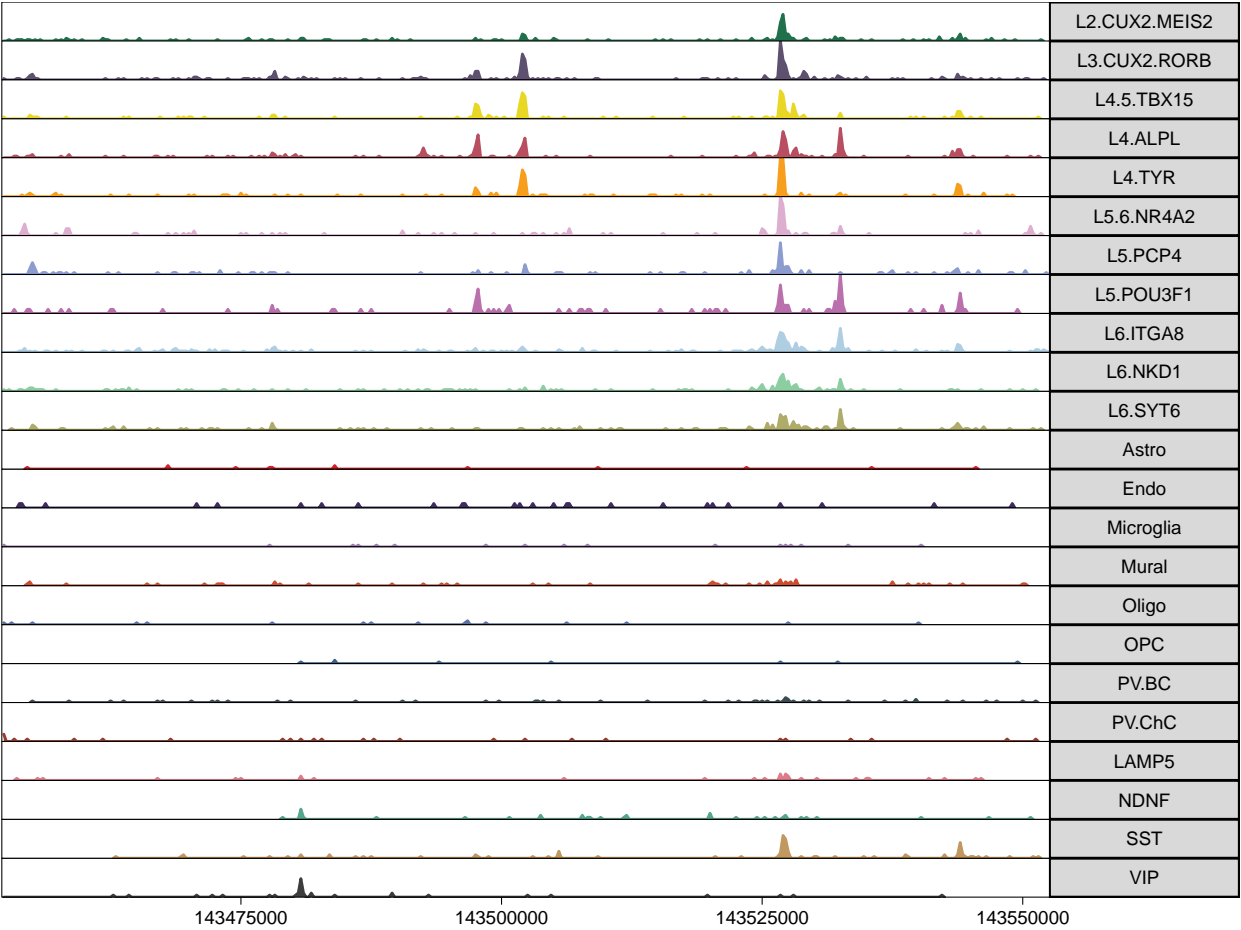


L4.5.TBX15

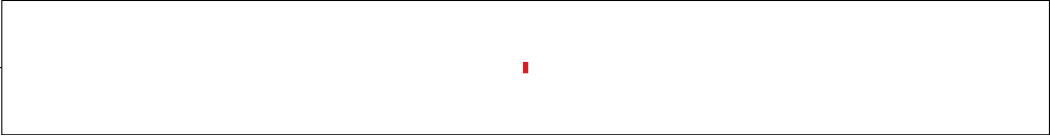


Peaks

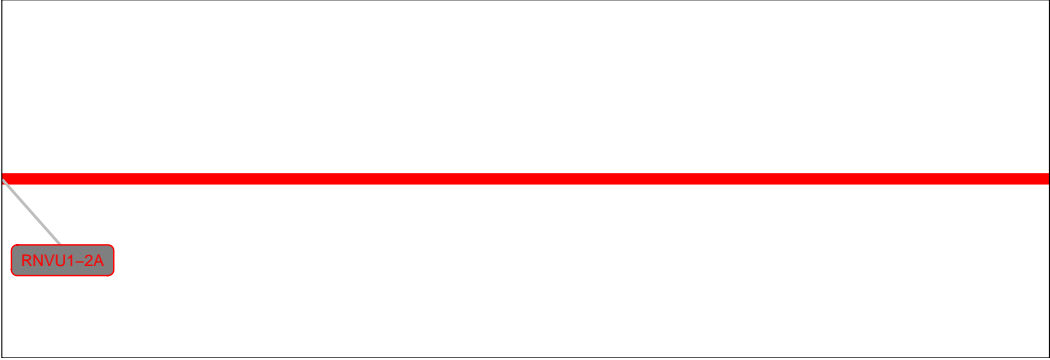
Coverage
(Norm. ATAC Signal Range (0–0.7) by ReadsInTSS)



L4.5.TBX15

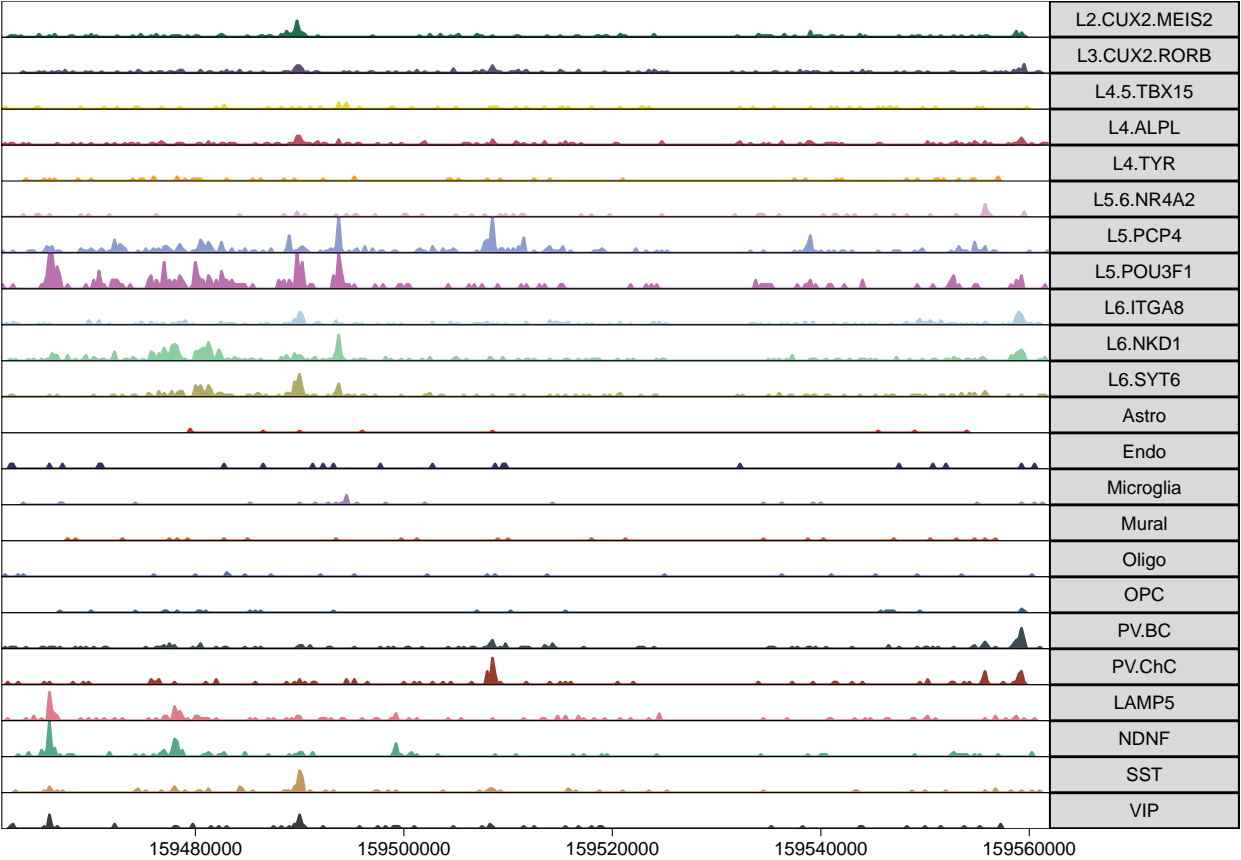


Peaks

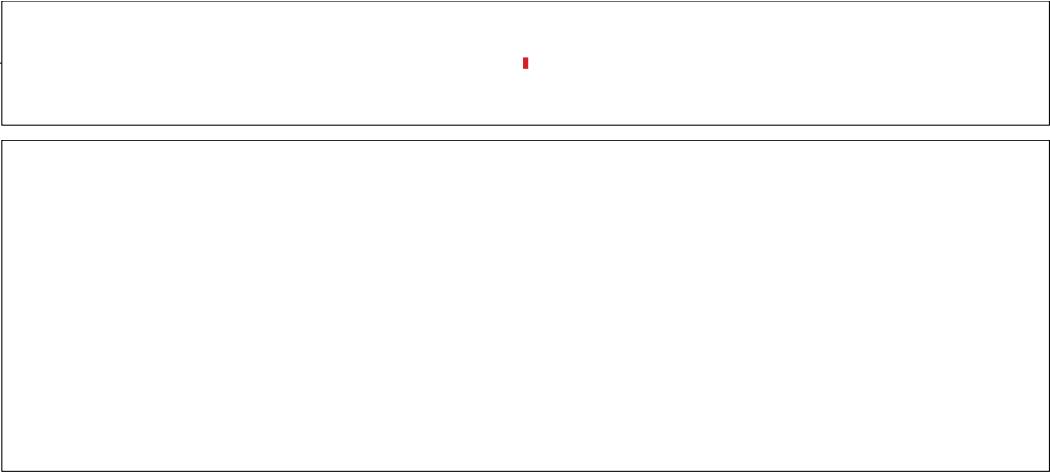


Genes

Coverage
(Norm. ATAC Signal Range (0–0.59) by ReadsInTSS)



L4.5.TBX15

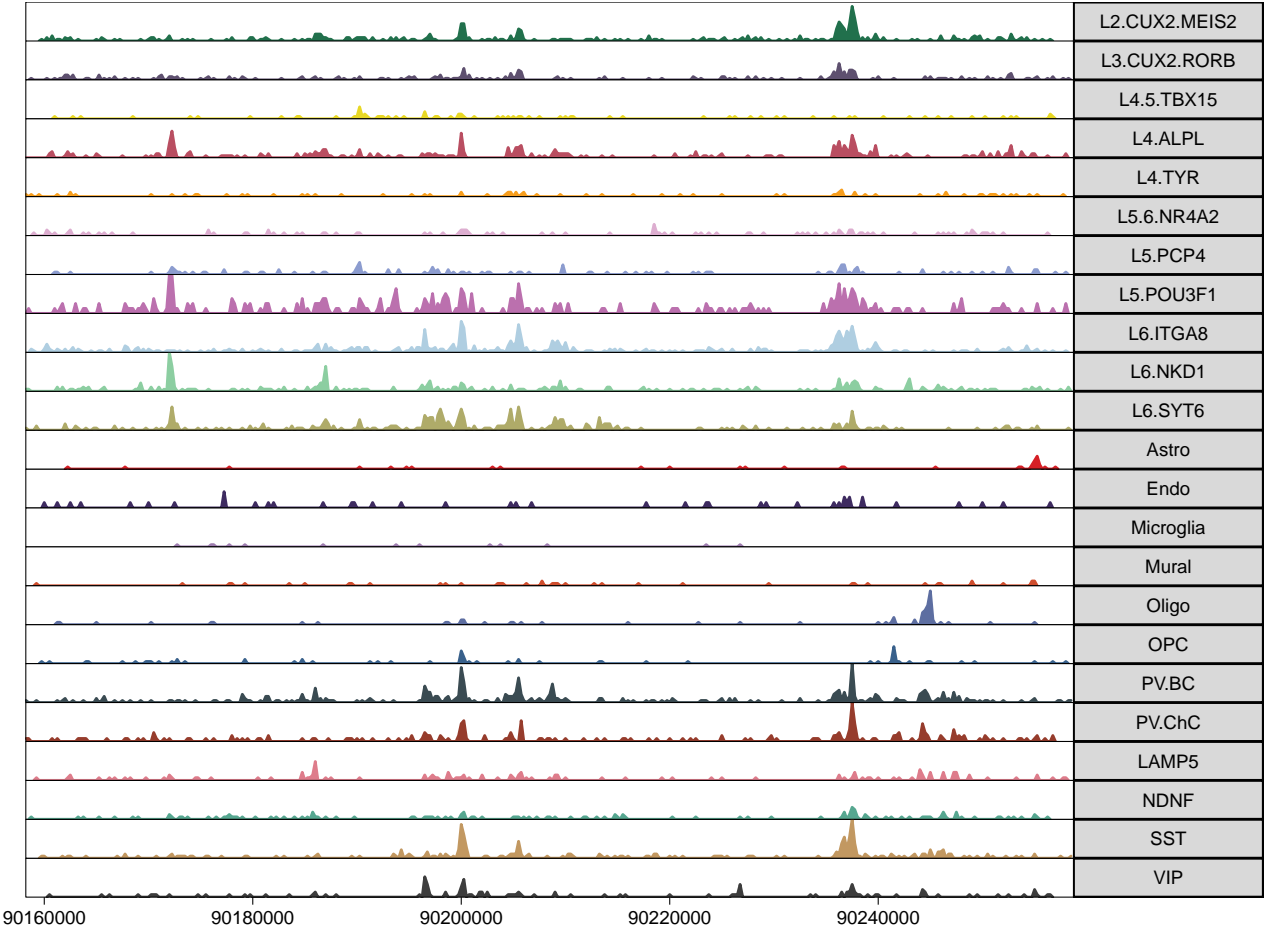


Peaks

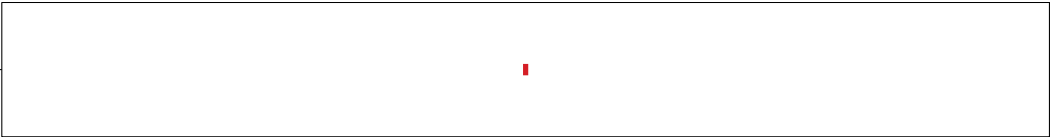
GeneTrack

chr17:90158228–90258729 Marker Peak Tracks for: L4.5.TBX15

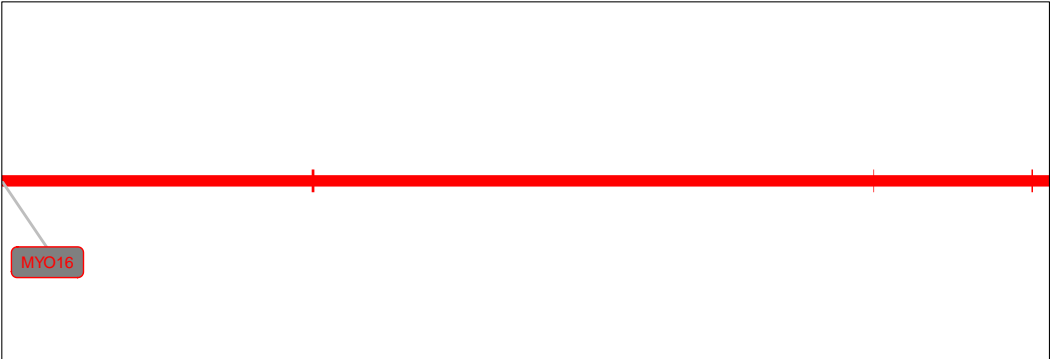
Coverage
(Norm. ATAC Signal Range (0–0.58) by ReadsInTSS)



L4.5.TBX15

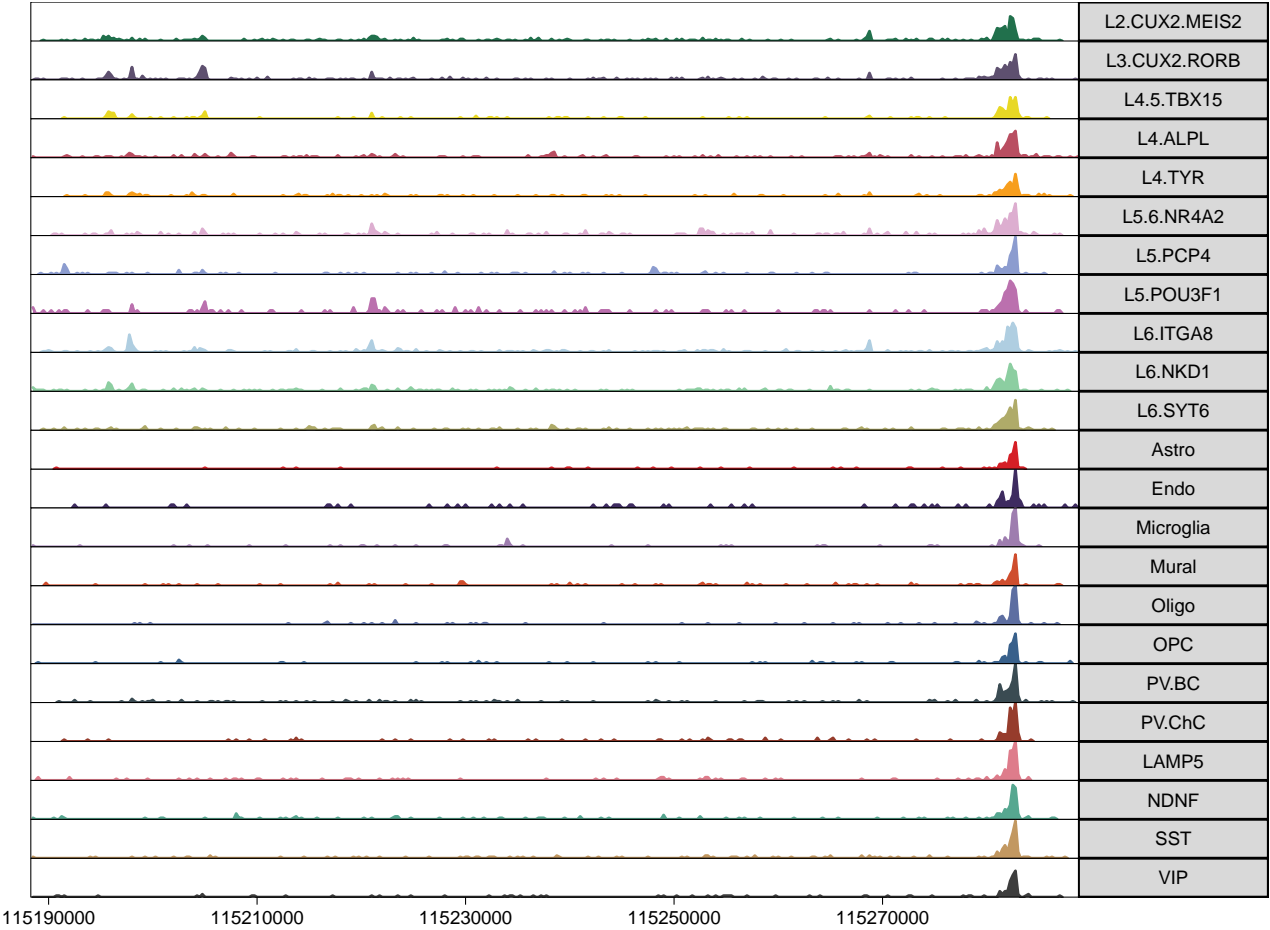


Peaks

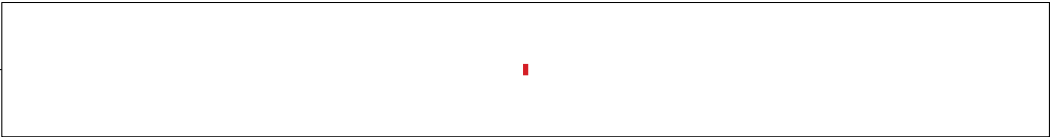


Genes

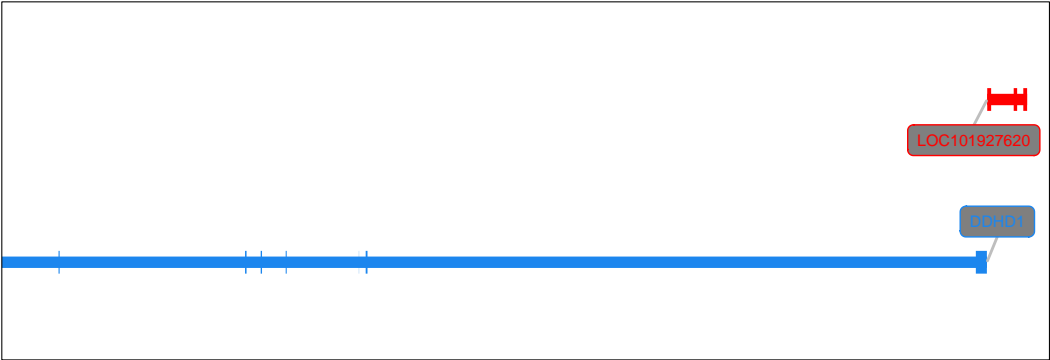
Coverage
(Norm. ATAC Signal Range (0–0.97) by ReadsInTSS)



L4.5.TBX15

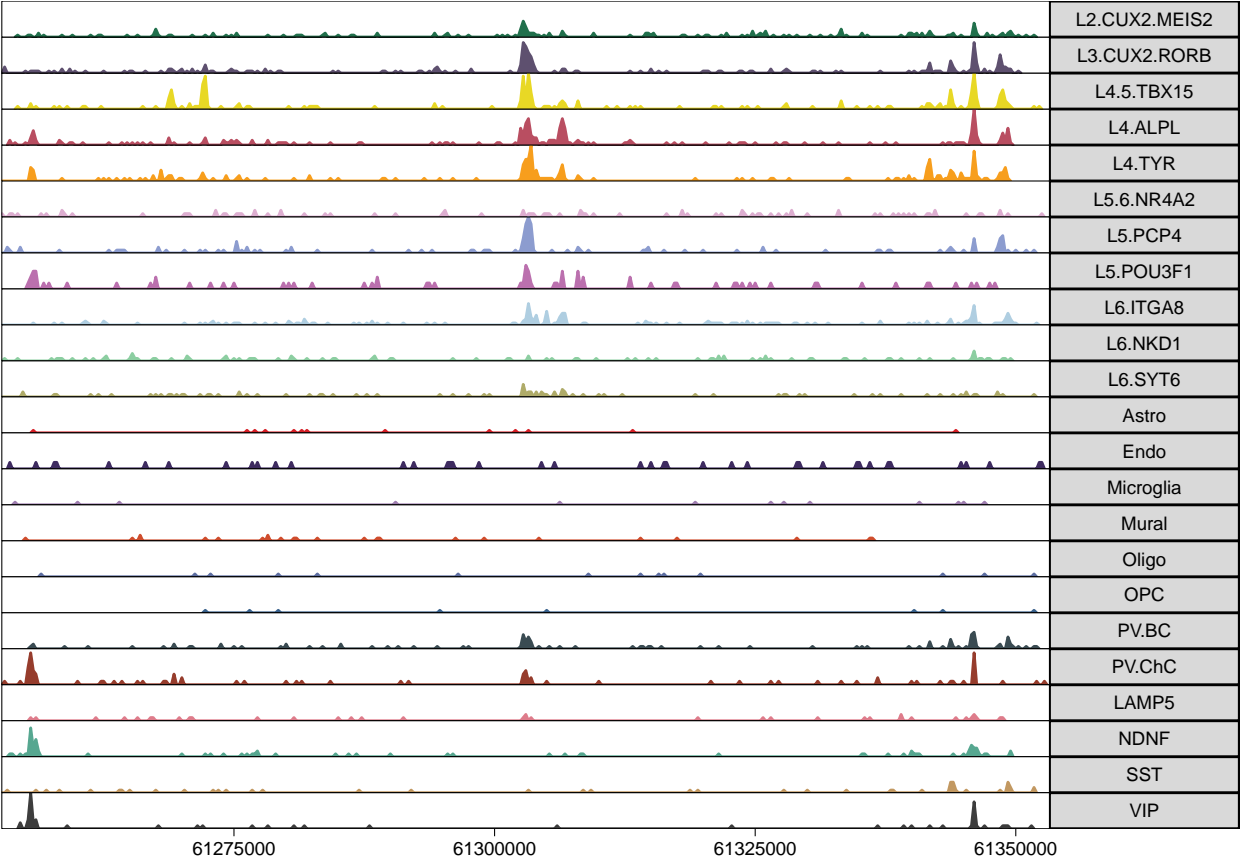


Peaks



Genes

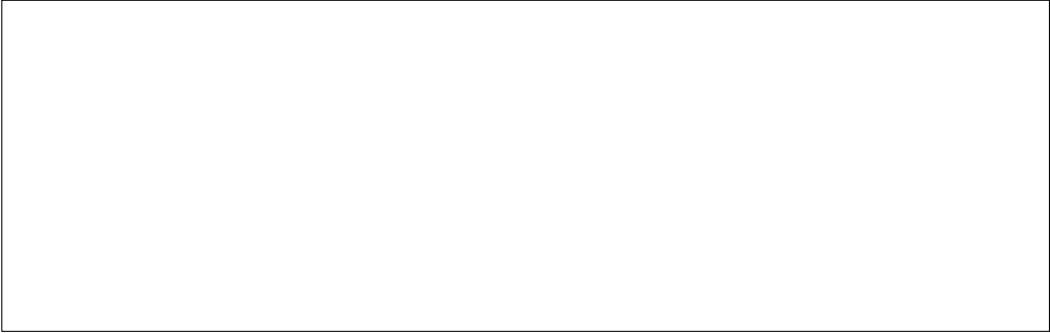
Coverage
(Norm. ATAC Signal Range (0–0.45) by ReadsInTSS)



L4.5.TBX15

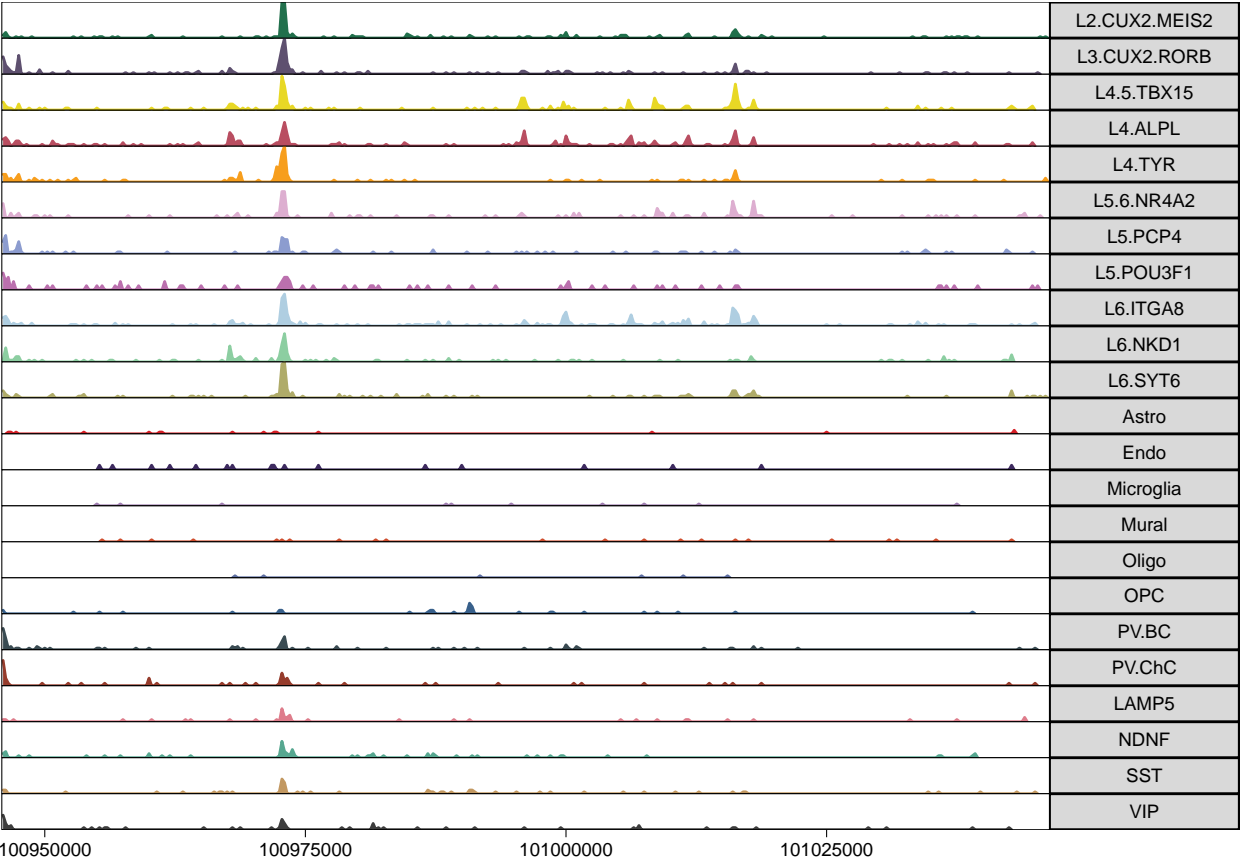


Peaks

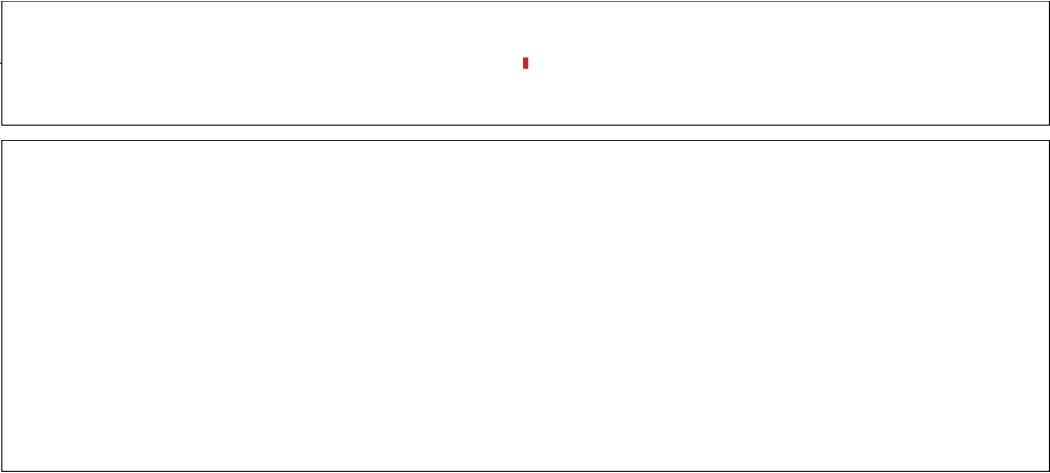


GeneTrack

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

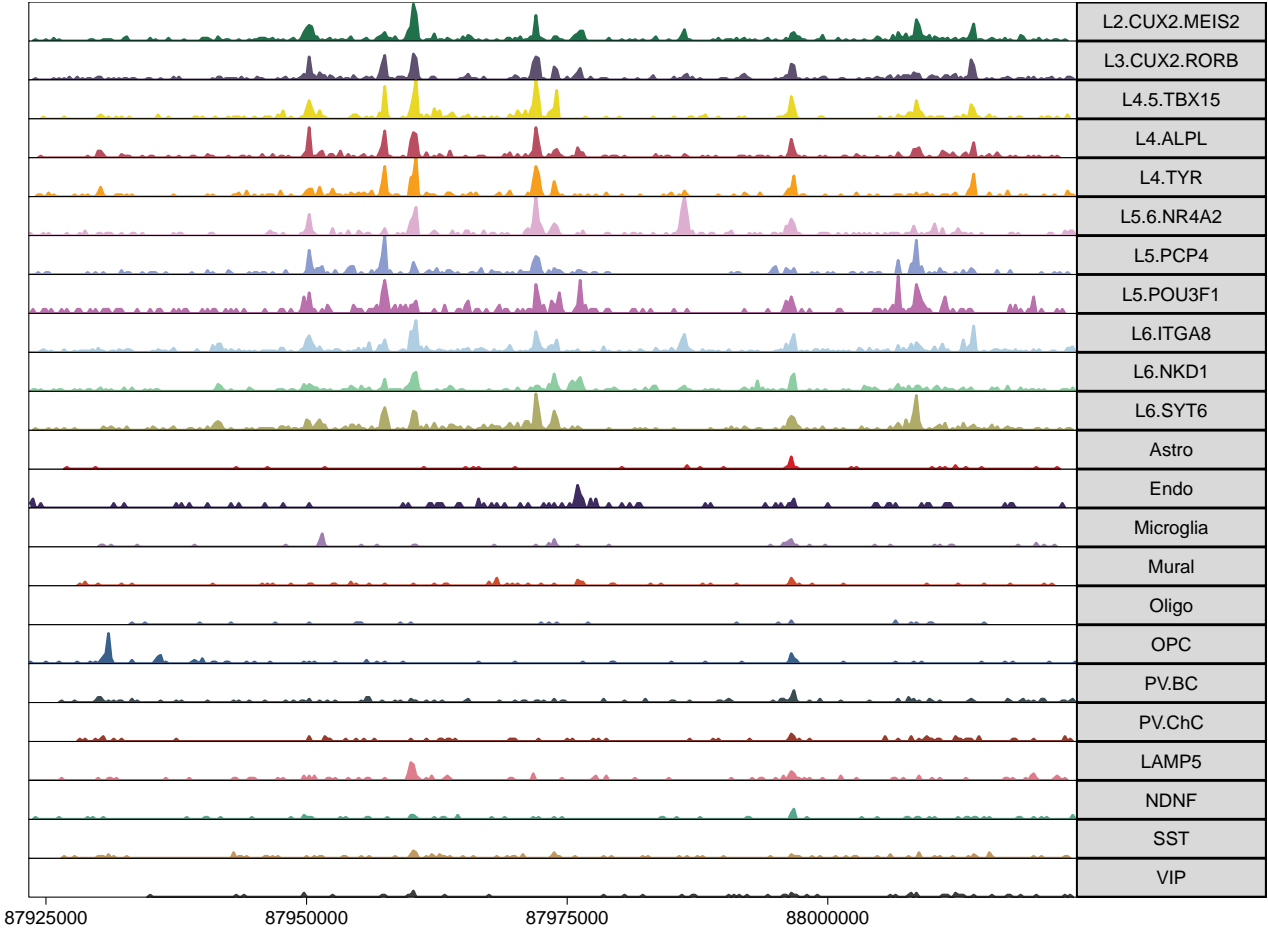


L4.5.TBX15

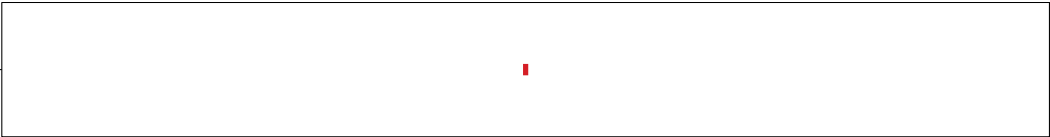


GeneTrack

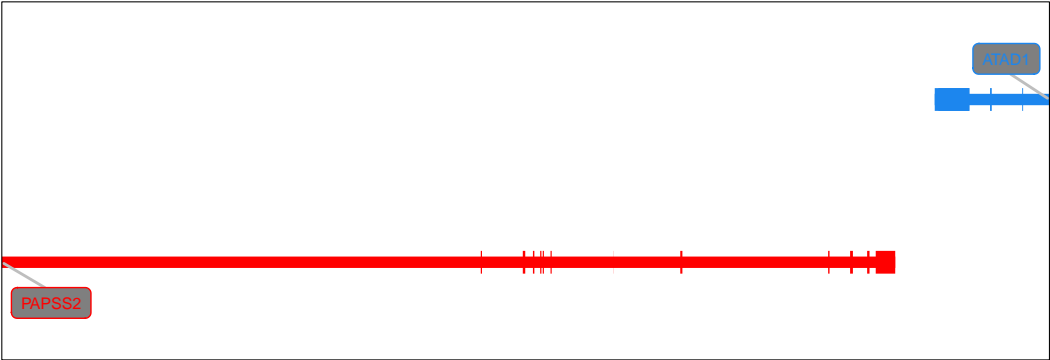
Coverage
(Norm. ATAC Signal Range (0–0.69) by ReadsInTSS)



L4.5.TBX15



Peaks



Genes