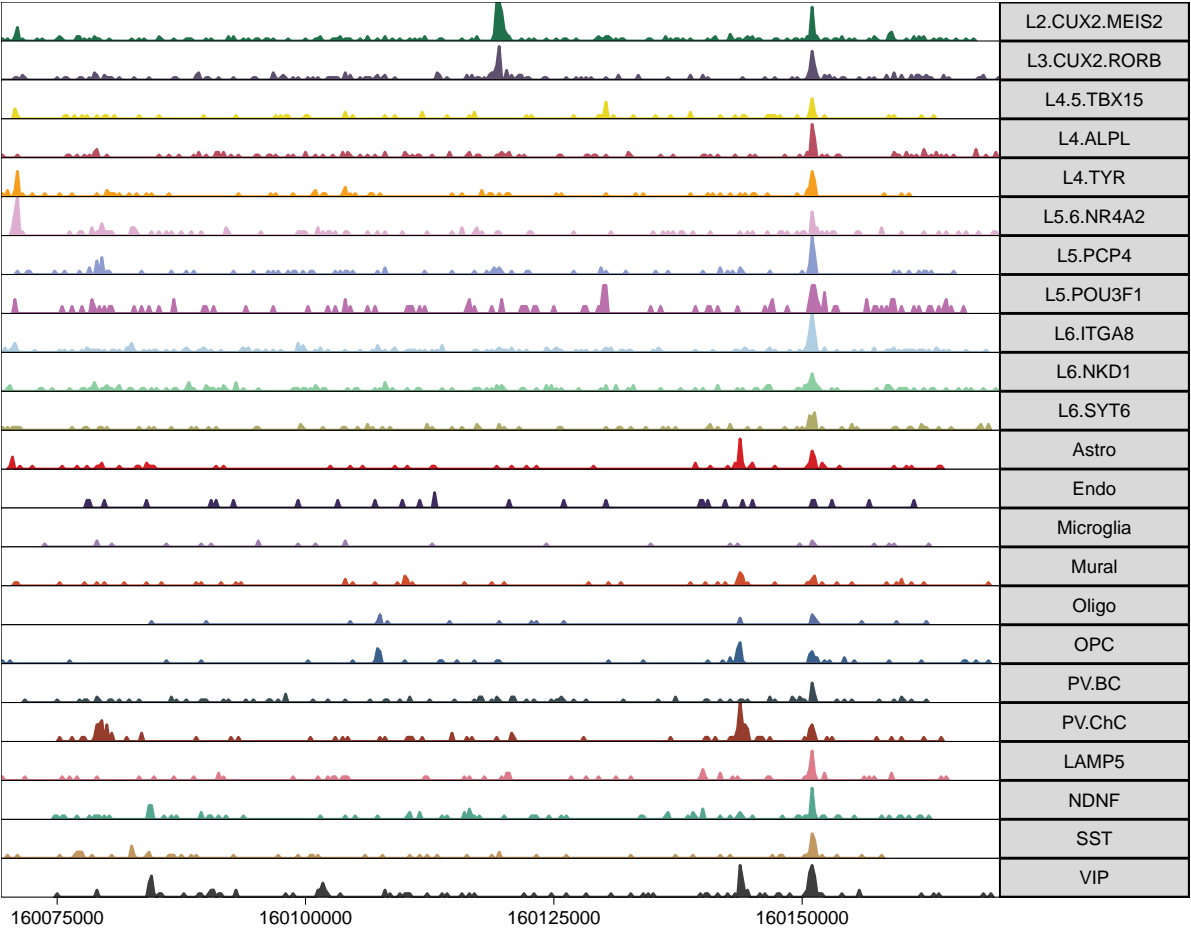
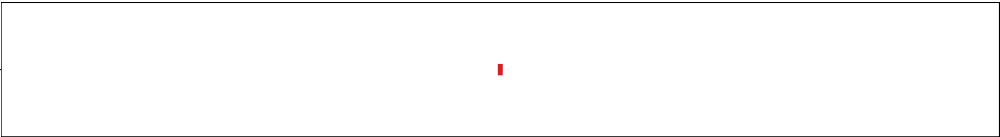


chr1:160069359–160169860 Marker Peak Tracks for: L2.CUX2.MEIS2

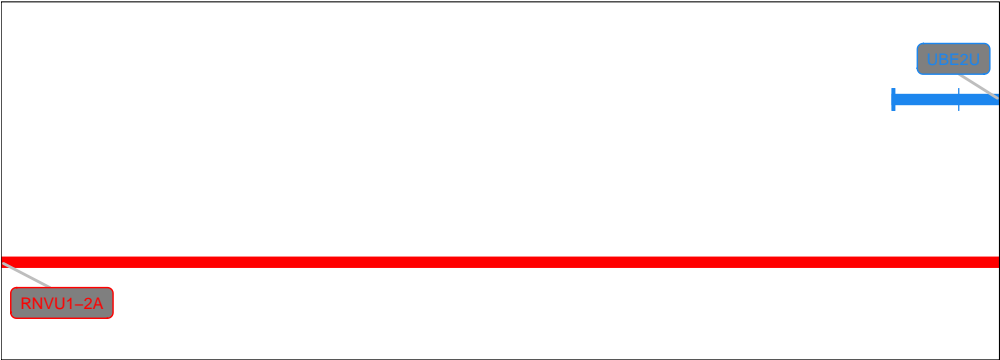
Coverage
(Norm. ATAC Signal Range (0–0.41) by ReadsInTSS)



L2.CUX2.MEIS2



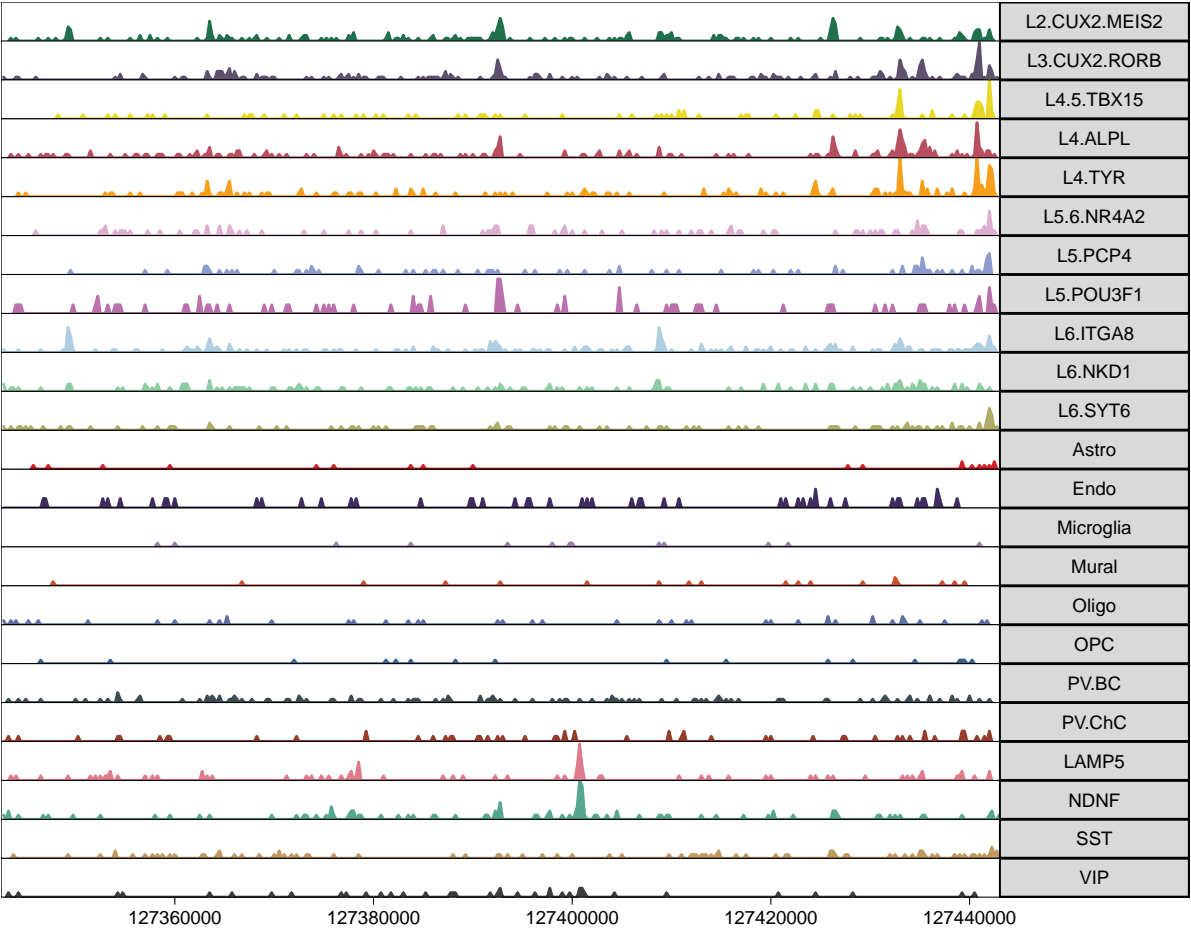
Peaks



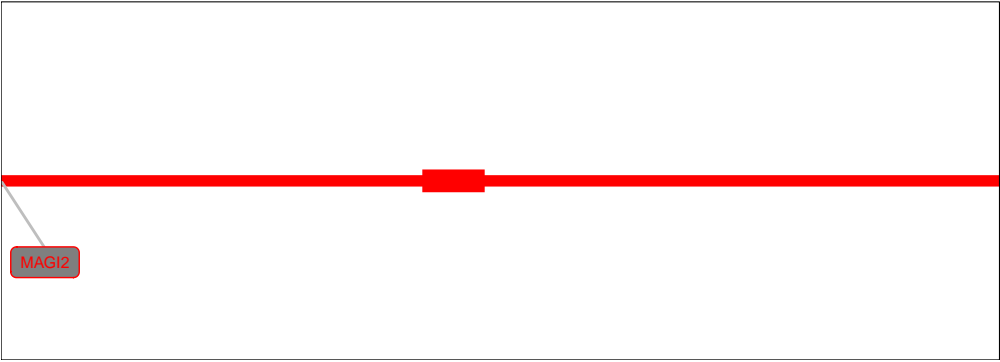
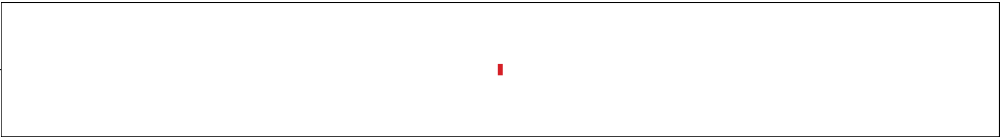
Genes

chr3:127342511–127443012 Marker Peak Tracks for: L2.CUX2.MEIS2

Coverage
(Norm. ATAC Signal Range (0–0.33) by ReadsInTSS)

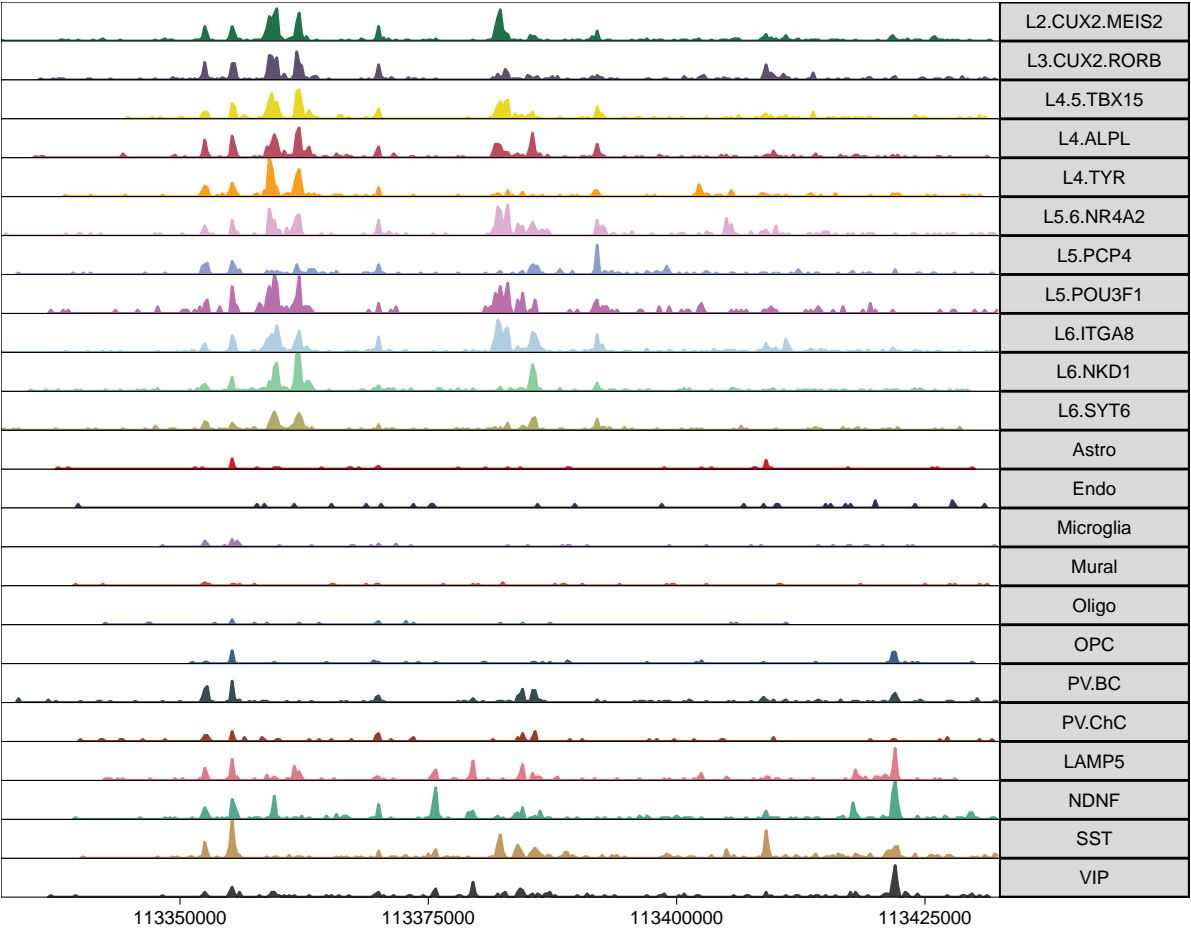


L2.CUX2.MEIS2

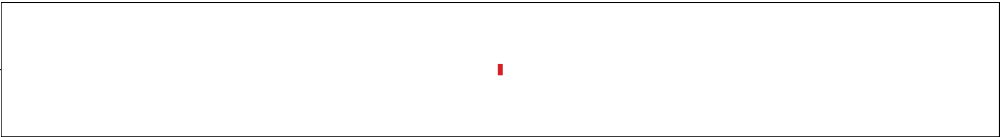


chr1:113331996–113432497 Marker Peak Tracks for: L2.CUX2.MEIS2

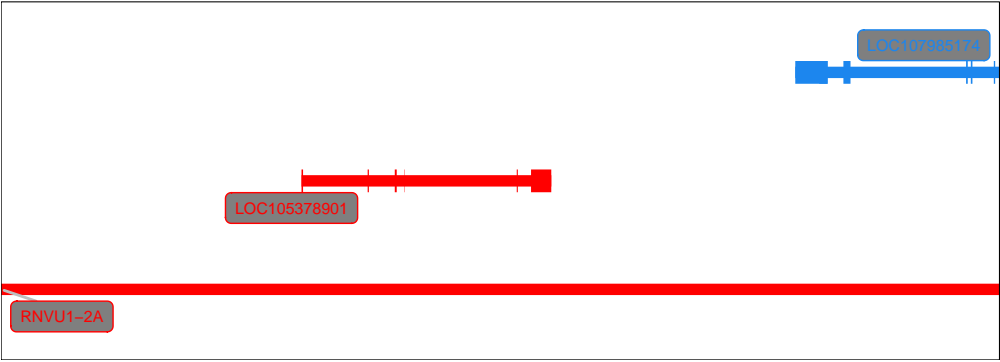
Coverage
(Norm. ATAC Signal Range (0–0.85) by ReadsInTSS)



L2.CUX2.MEIS2



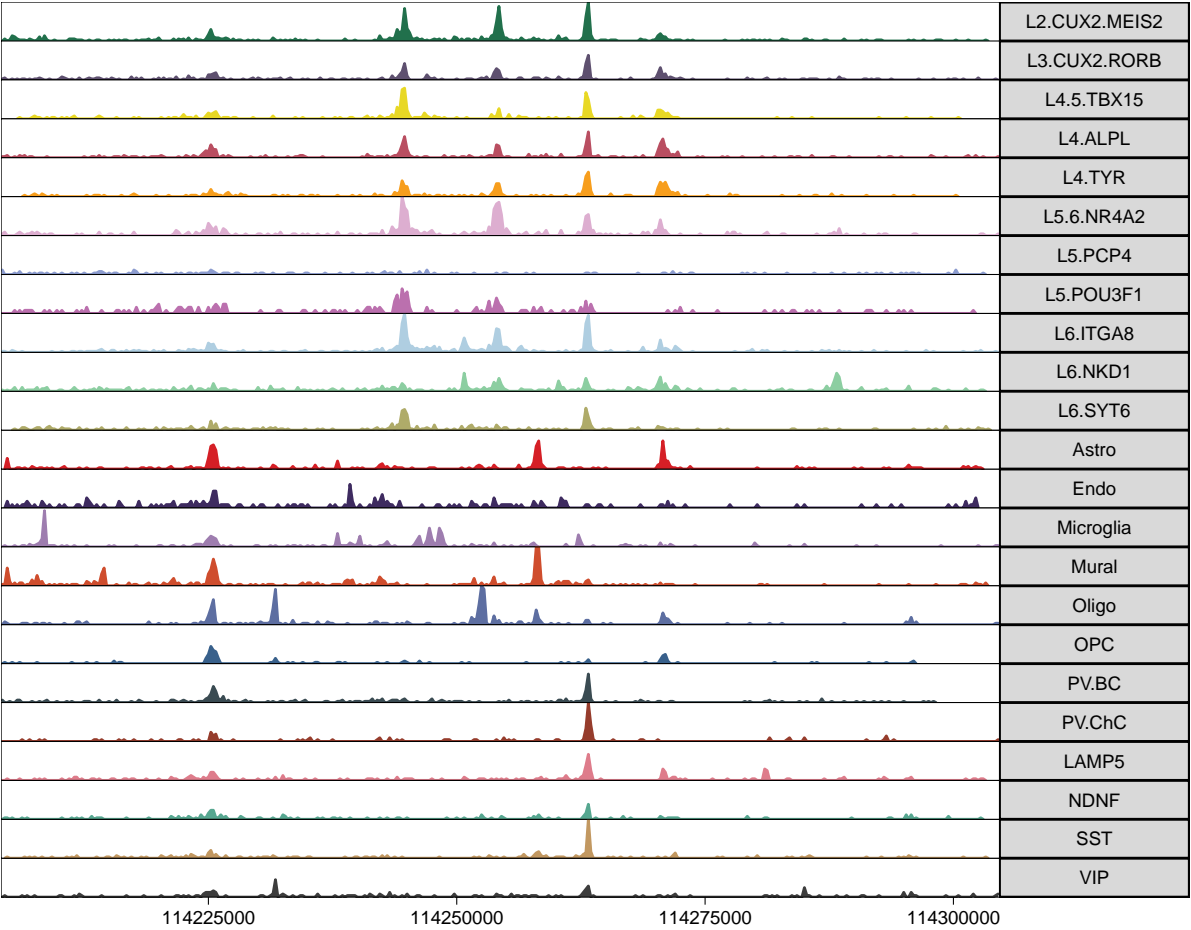
Peaks



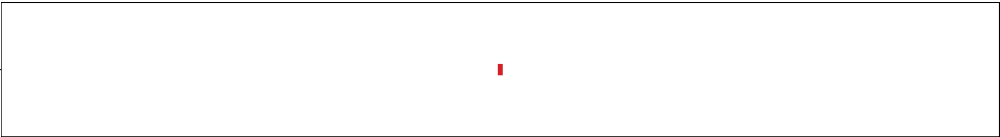
Genes

chr1:114204133–114304634 Marker Peak Tracks for: L2.CUX2.MEIS2

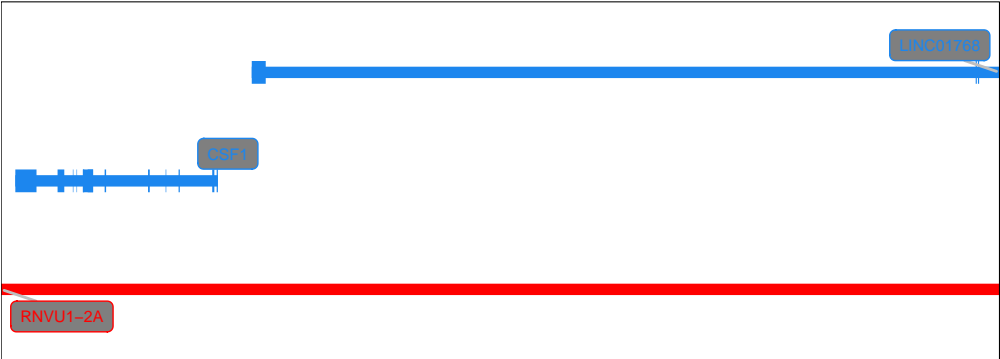
Coverage
(Norm. ATAC Signal Range (0–0.93) by ReadsInTSS)



L2.CUX2.MEIS2

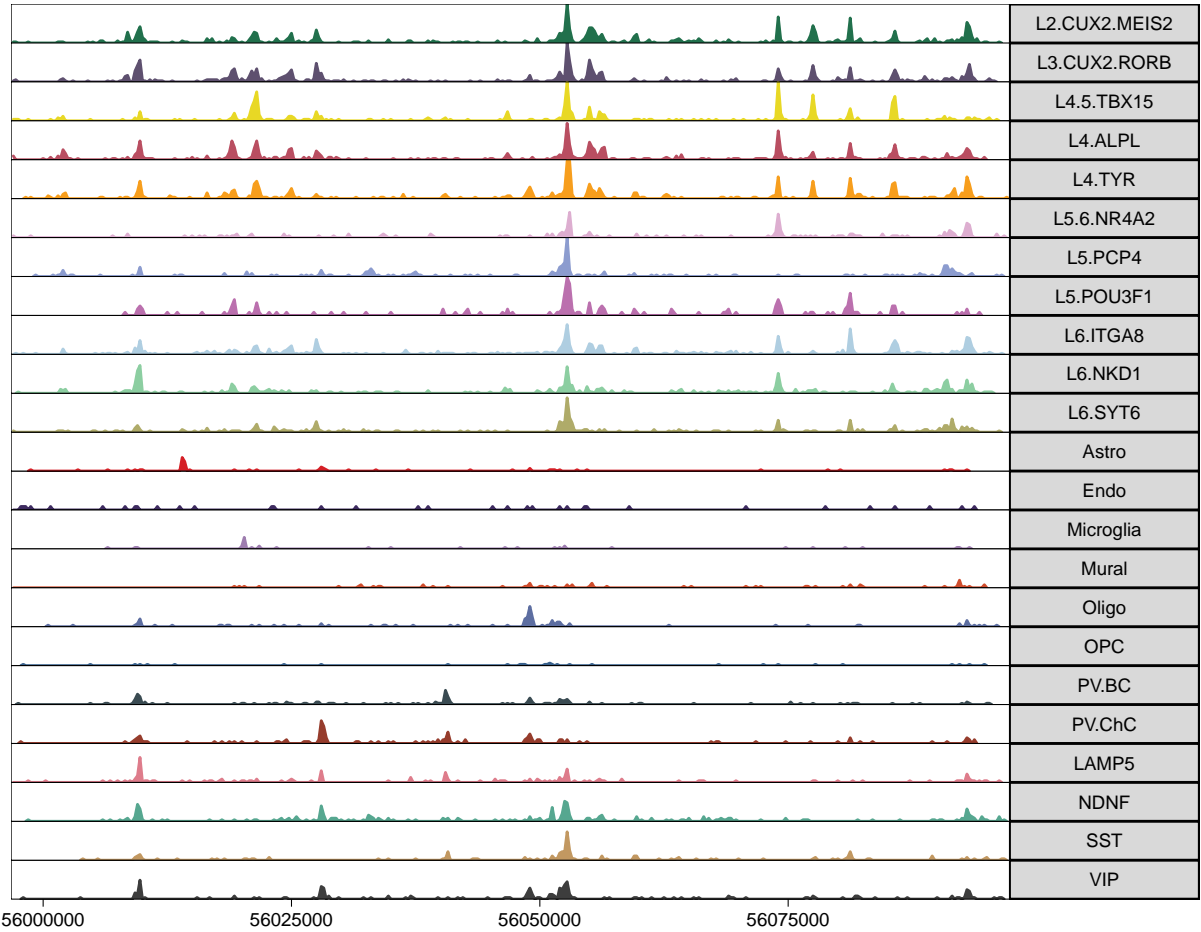


Peaks

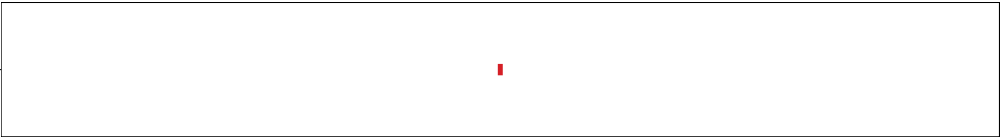


Genes

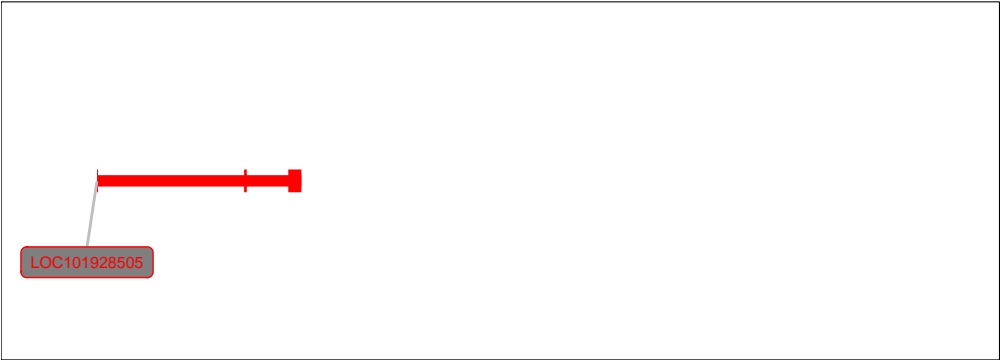
Coverage
(Norm. ATAC Signal Range (0-0.9) by ReadsInTSS)



L2.CUX2.MEIS2

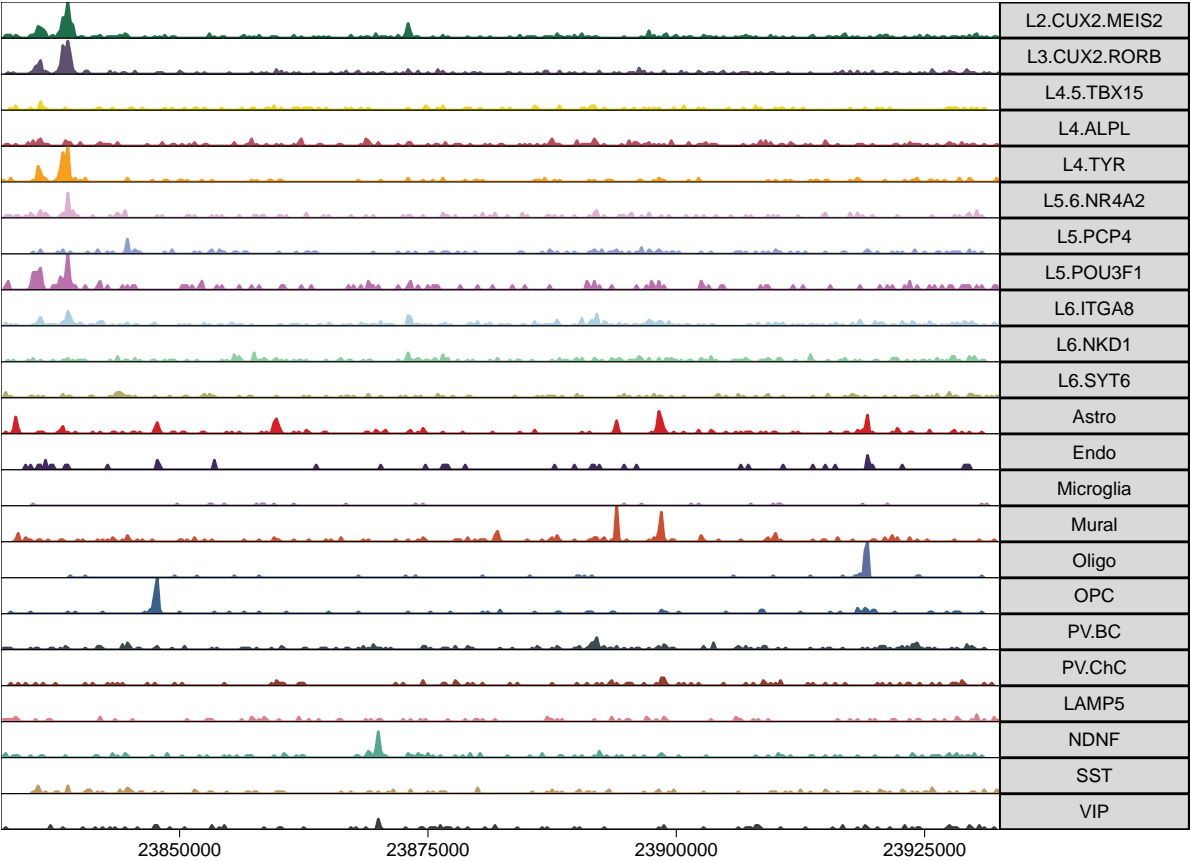


Peaks



Genes

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



L2.CUX2.MEIS2

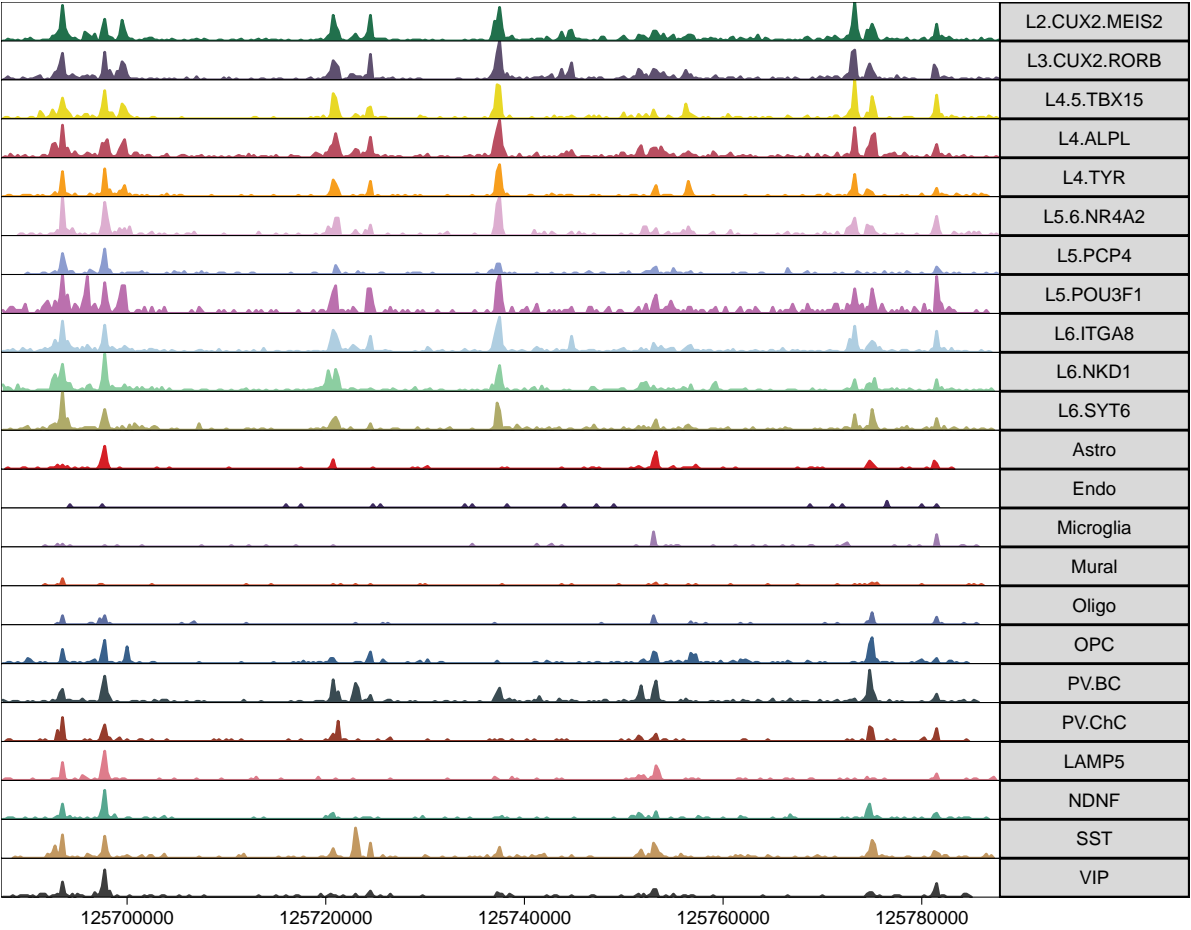


Peaks

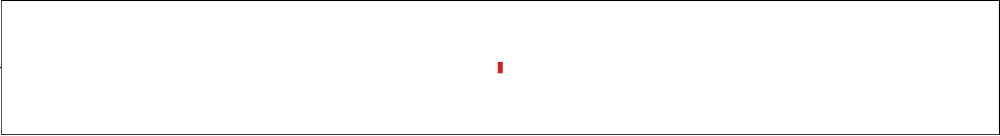
GeneTrack

chr14:125687320–125787821 Marker Peak Tracks for: L2.CUX2.MEIS2

Coverage
(Norm. ATAC Signal Range (0–0.93) by ReadsInTSS)



L2.CUX2.MEIS2



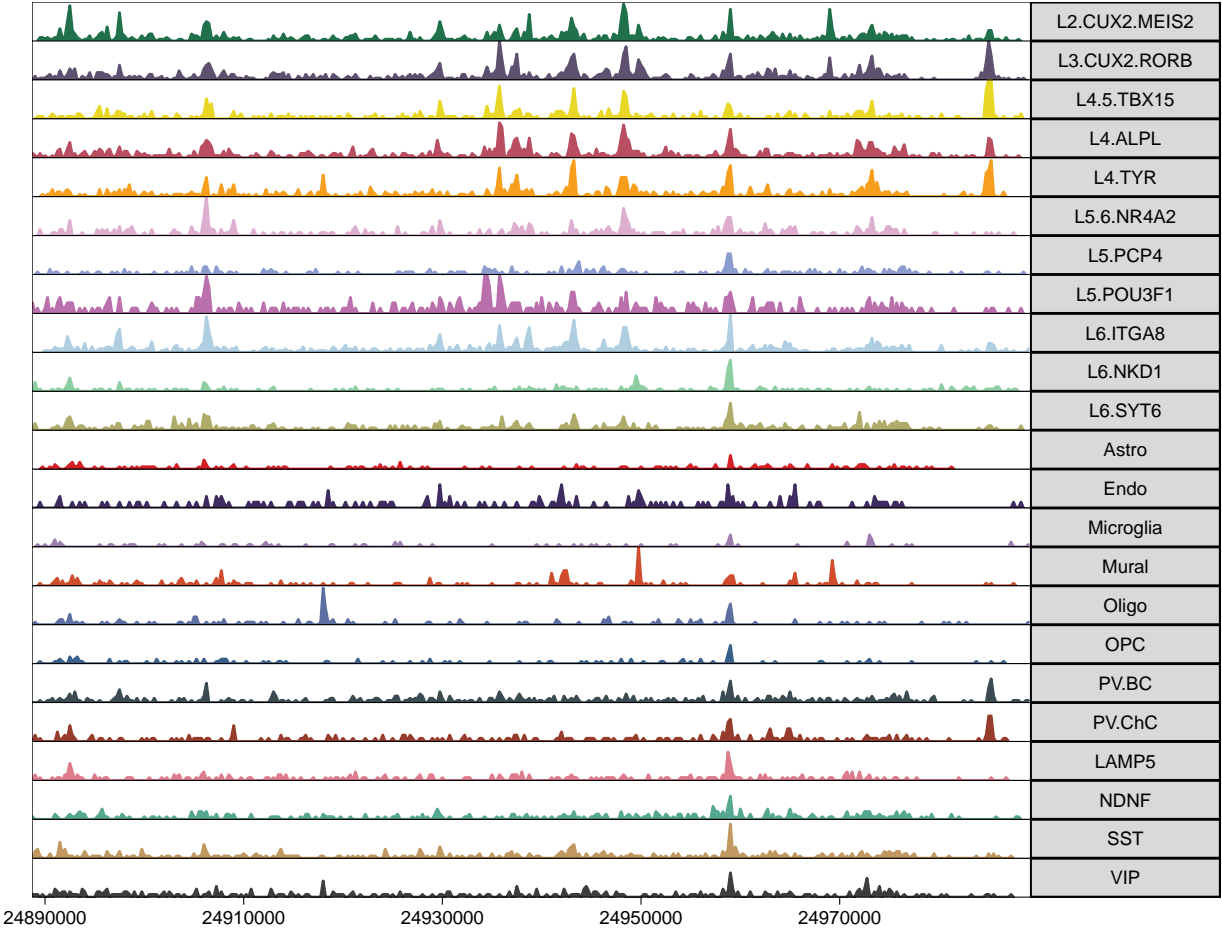
Peaks



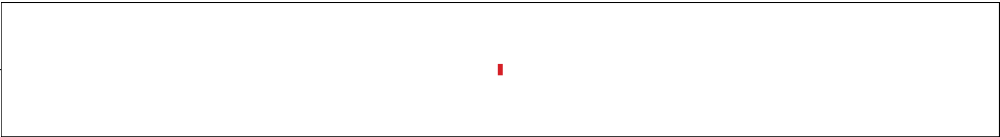
Genes

chr18:24888703–24989204 Marker Peak Tracks for: L2.CUX2.MEIS2

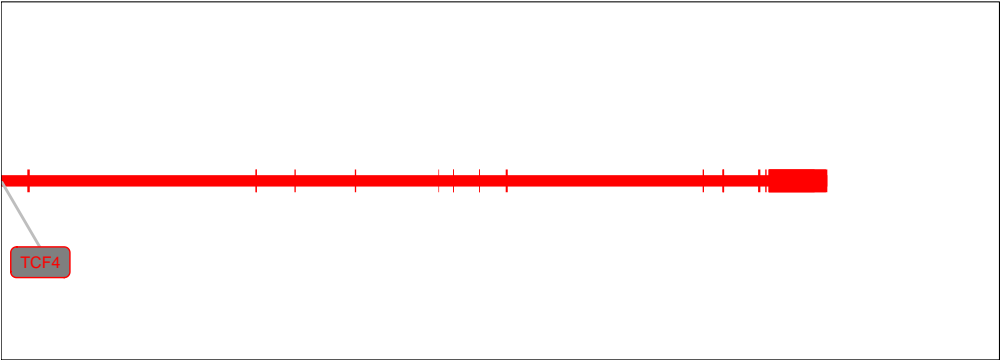
Coverage
(Norm. ATAC Signal Range (0–0.54) by ReadsInTSS)



L2.CUX2.MEIS2

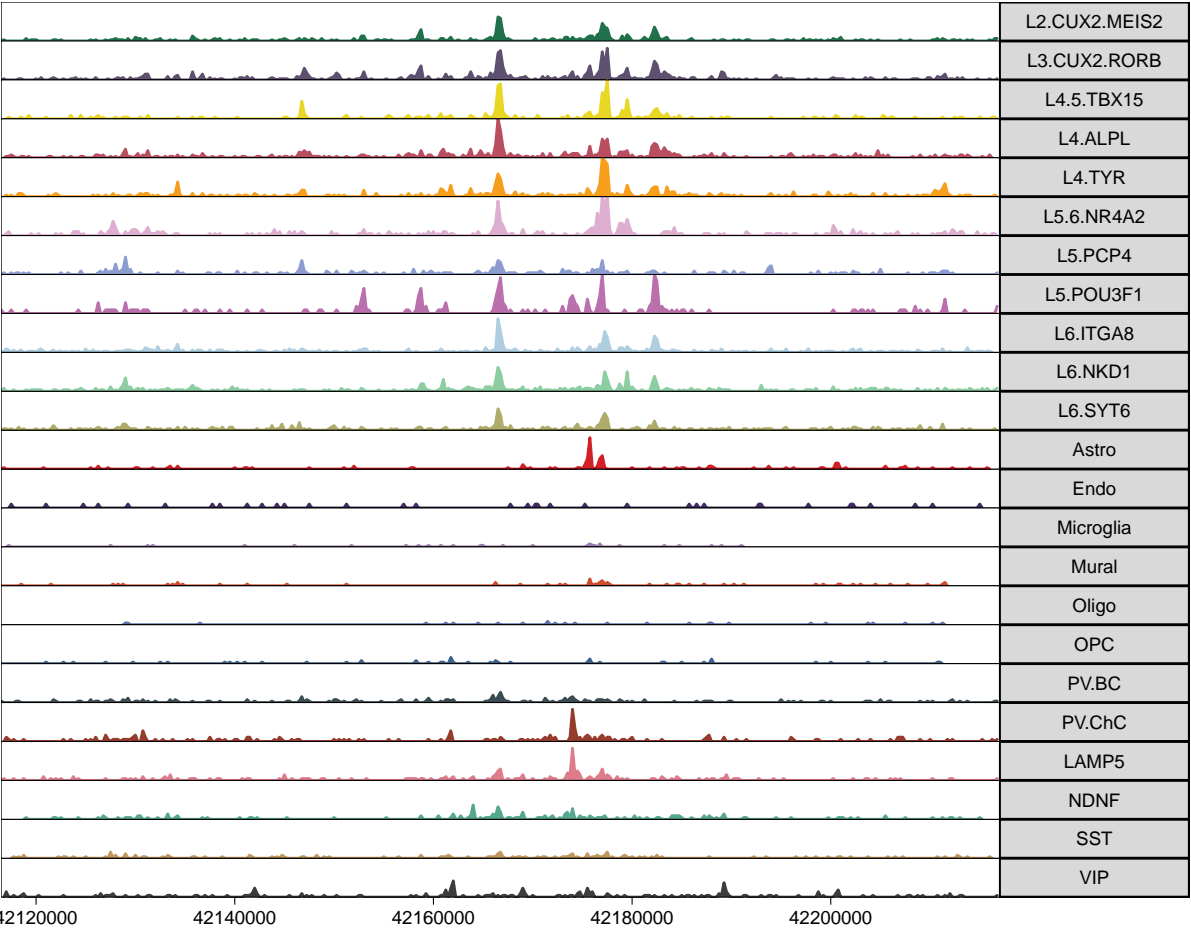


Peaks

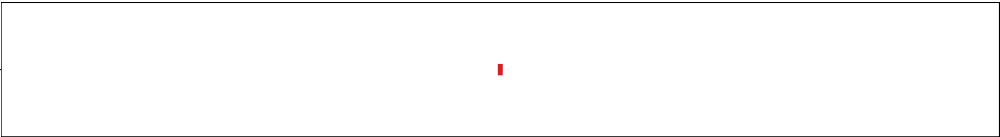


Genes

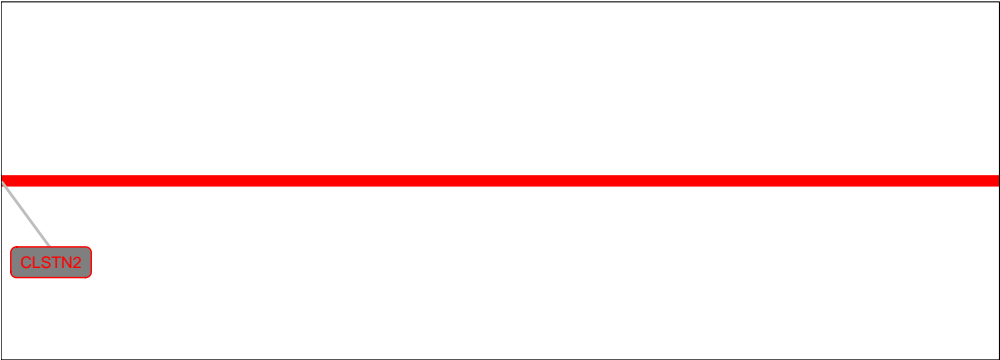
Coverage
(Norm. ATAC Signal Range (0–0.8) by ReadsInTSS)



L2.CUX2.MEIS2

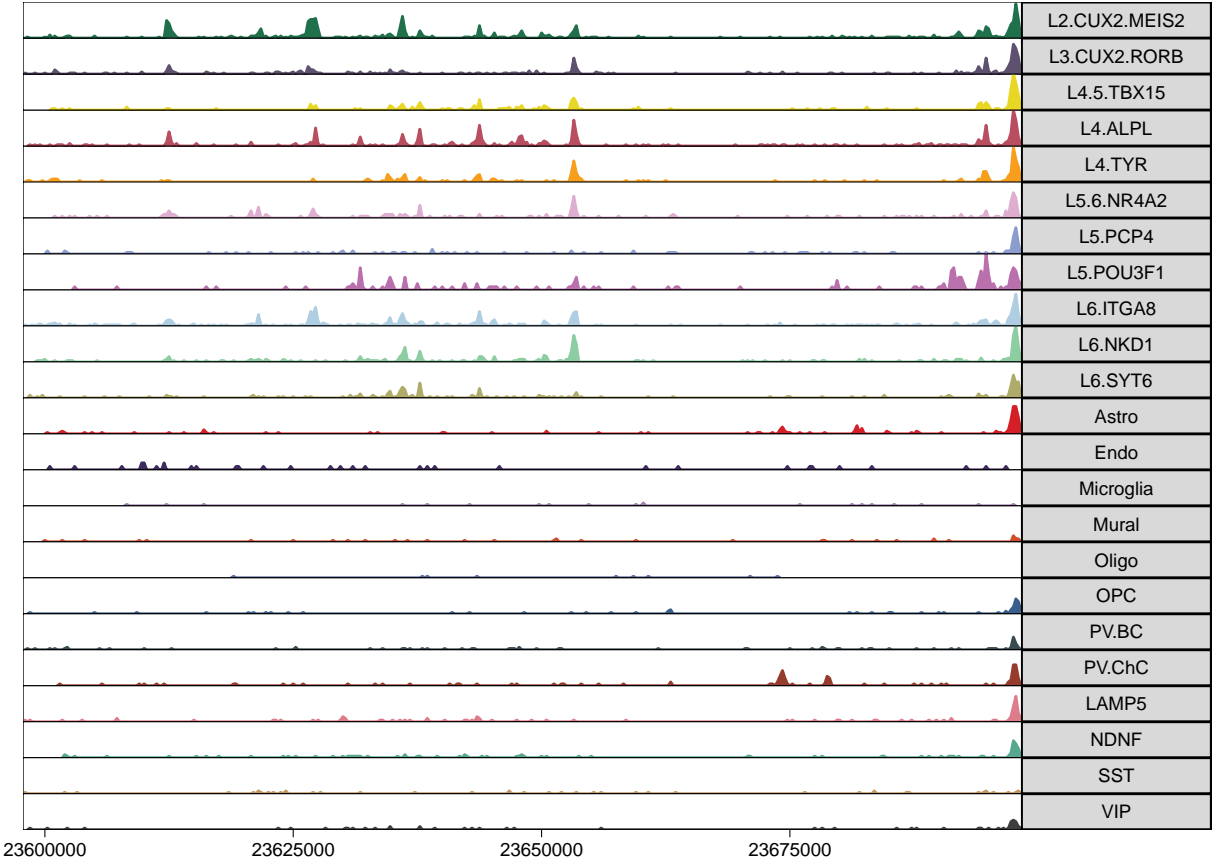


Peaks

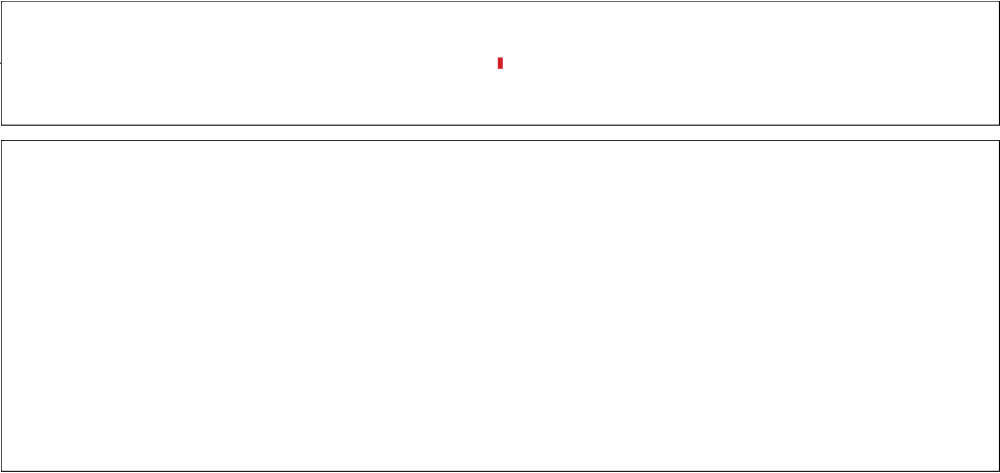


Genes

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



L2.CUX2.MEIS2

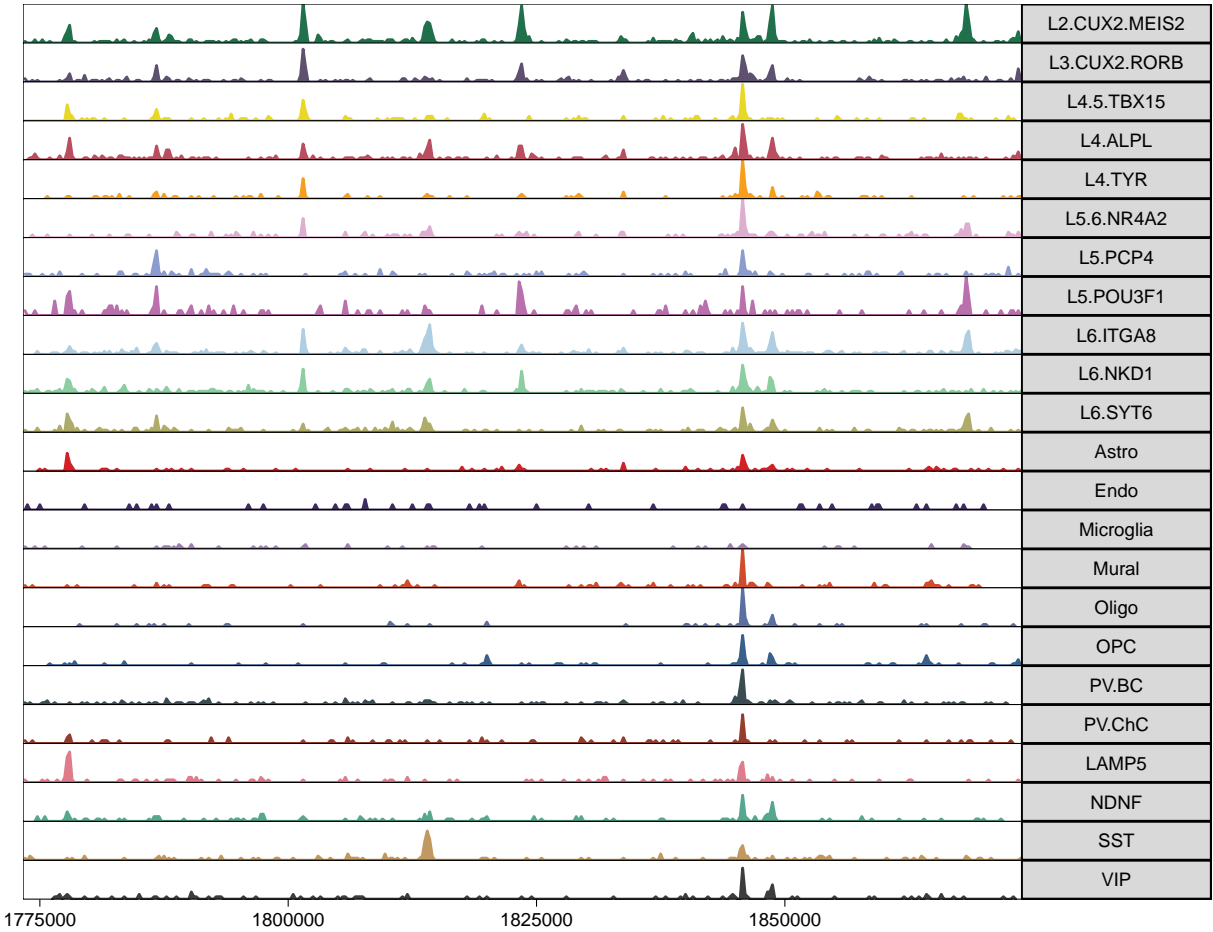


Peaks

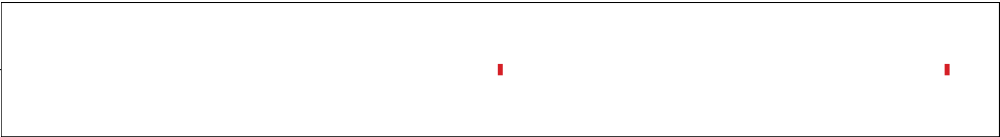
GeneTrack

chrX:1773316–1873817 Marker Peak Tracks for: L2.CUX2.MEIS2

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



L2.CUX2.MEIS2



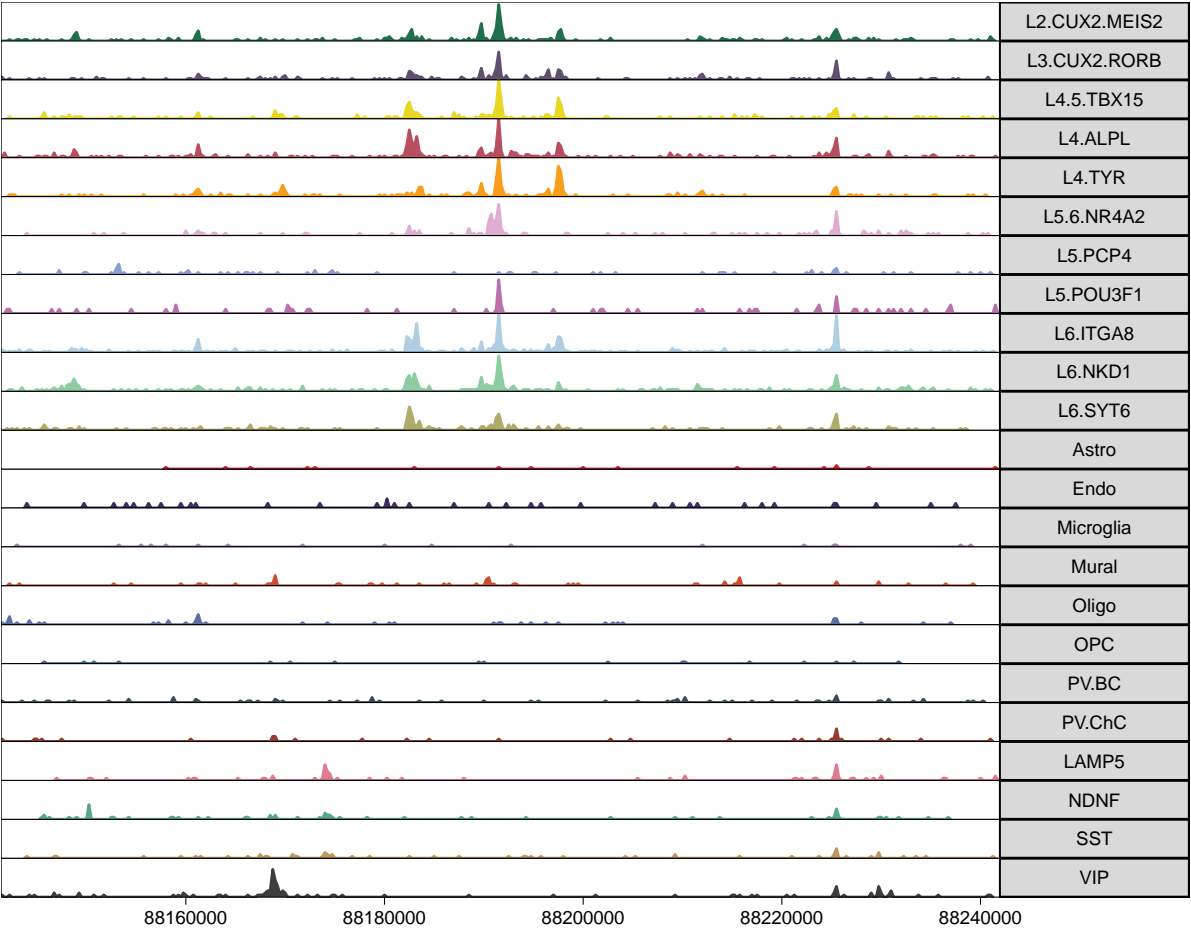
Peaks



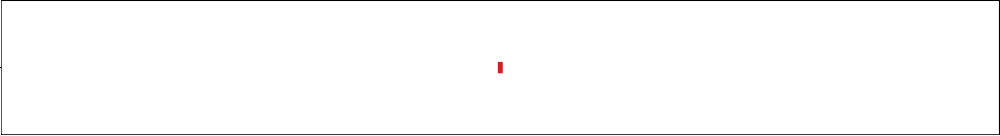
Genes

chr15:88141406–88241907 Marker Peak Tracks for: L2.CUX2.MEIS2

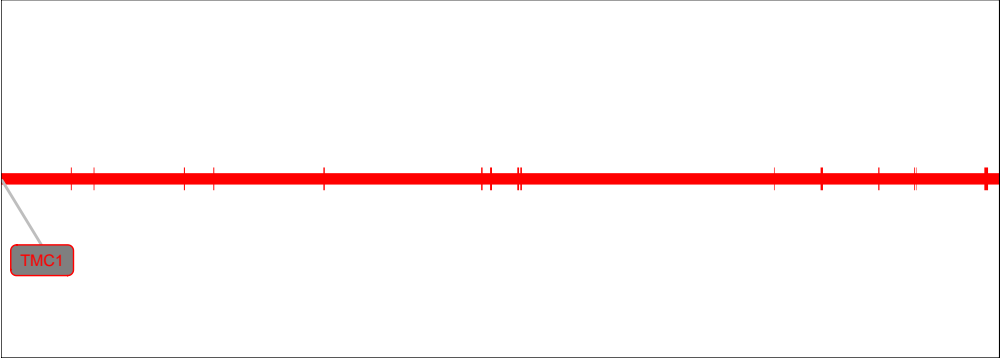
Coverage
(Norm. ATAC Signal Range (0–0.68) by ReadsInTSS)



L2.CUX2.MEIS2



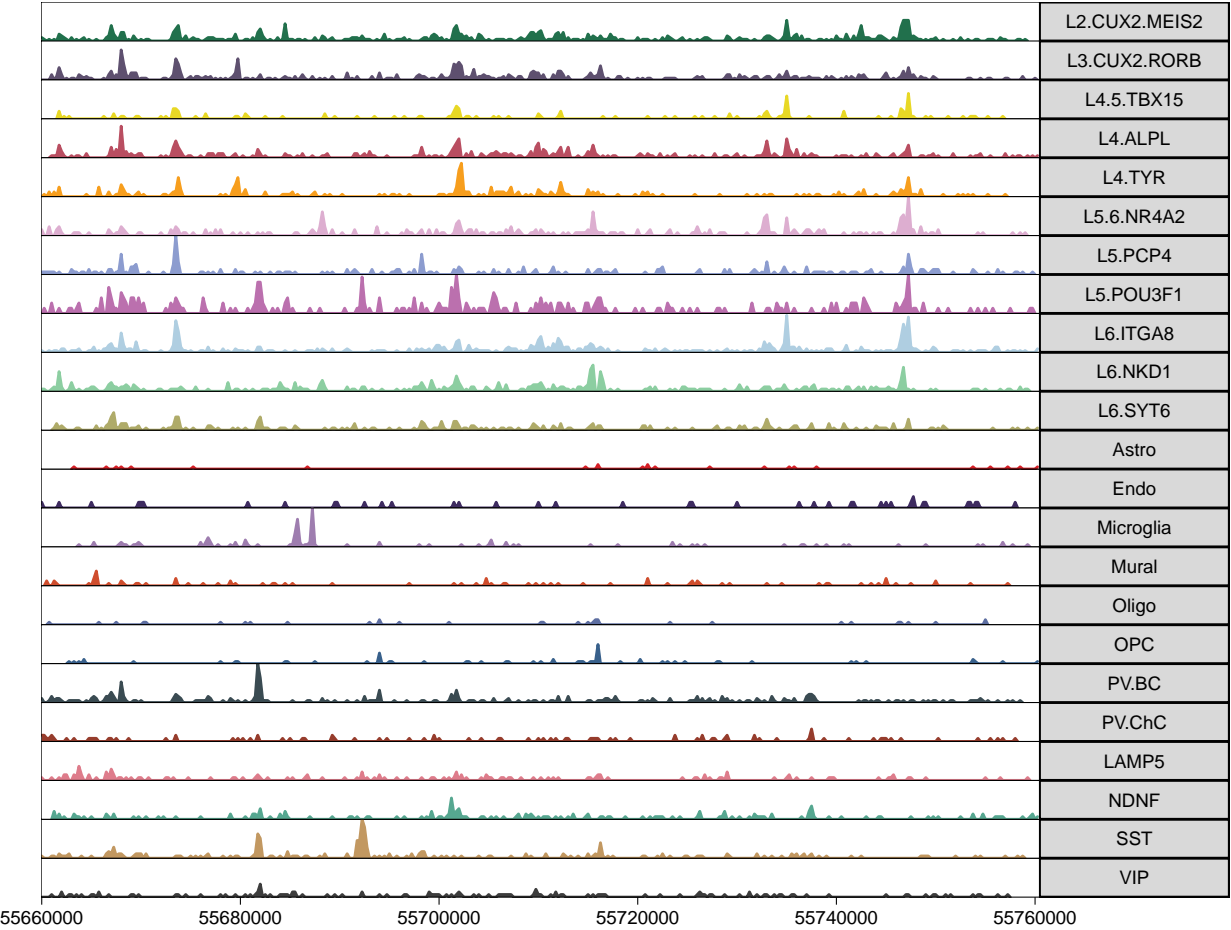
Peaks



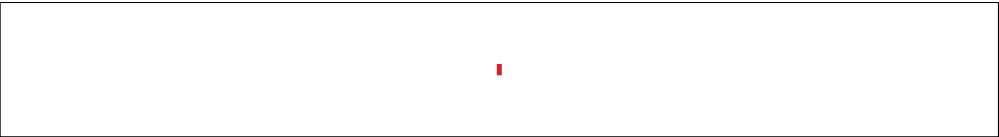
Genes

chr12:55659940–55760441 Marker Peak Tracks for: L2.CUX2.MEIS2

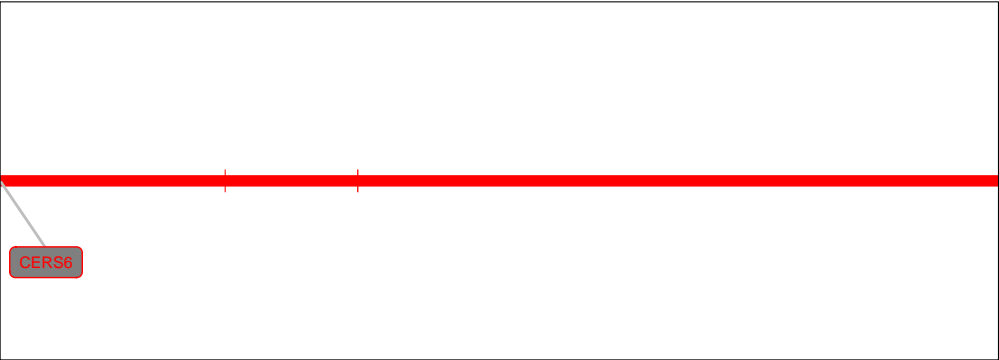
Coverage
(Norm. ATAC Signal Range (0–0.55) by ReadsInTSS)



L2.CUX2.MEIS2



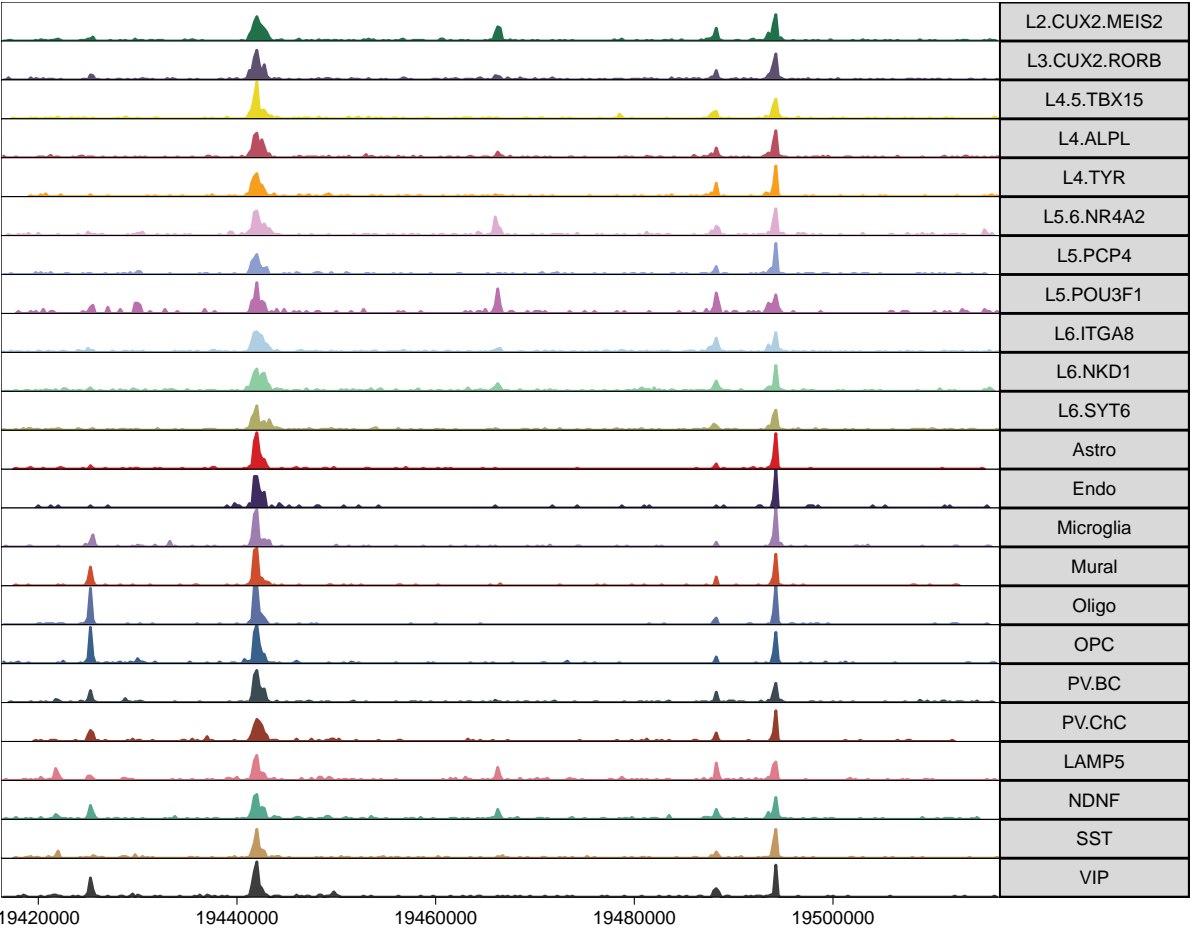
Peaks



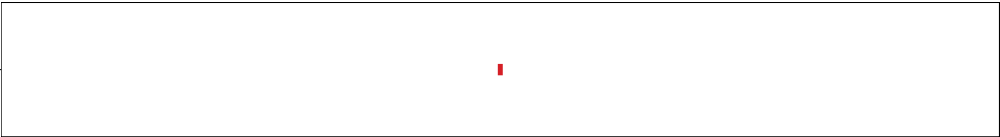
Genes

chr1:19416257-19516758 Marker Peak Tracks for: L2.CUX2.MEIS2

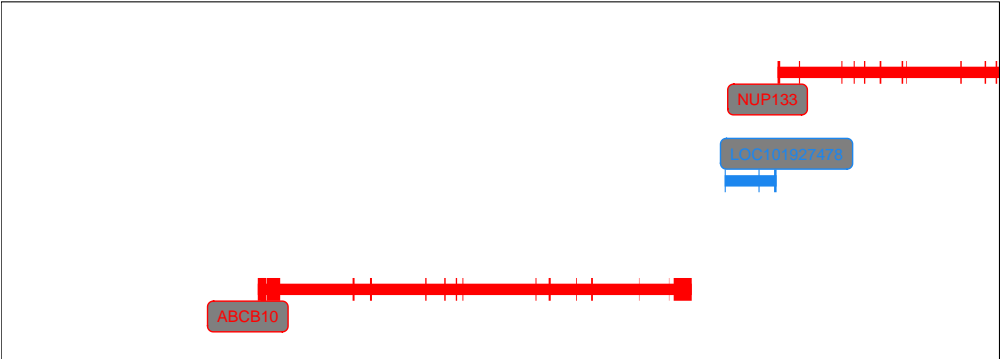
Coverage
(Norm. ATAC Signal Range (0-1.39) by ReadsInTSS)



L2.CUX2.MEIS2



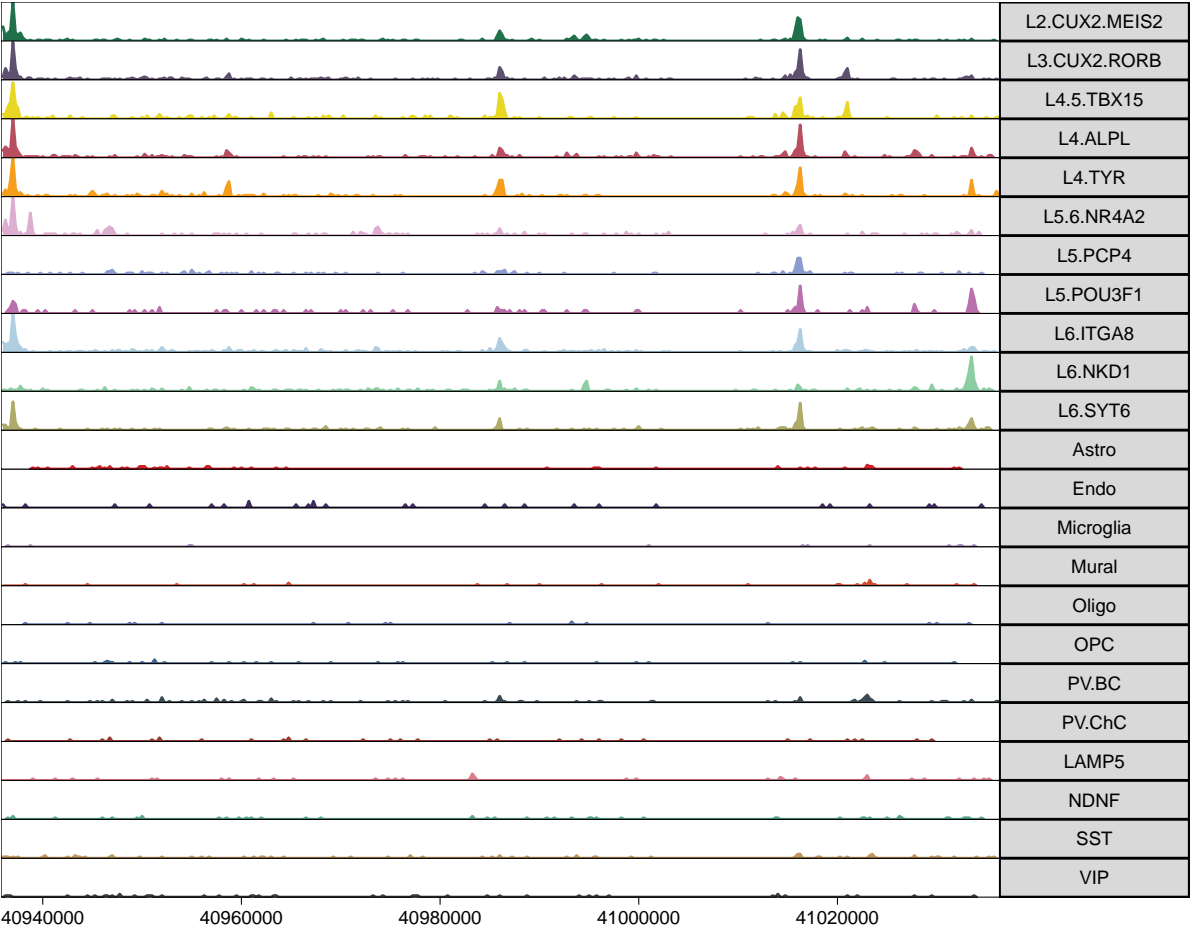
Peaks



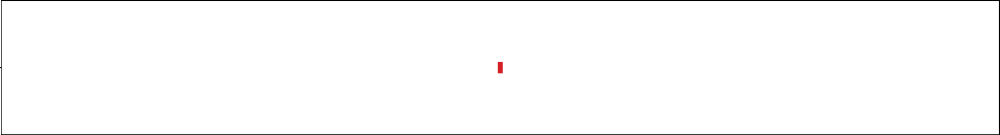
Genes

chr11:40935810–41036311 Marker Peak Tracks for: L2.CUX2.MEIS2

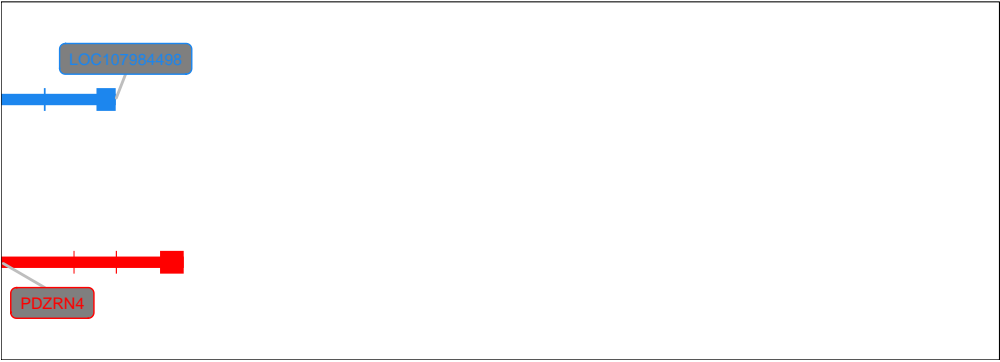
Coverage
(Norm. ATAC Signal Range (0–0.93) by ReadsInTSS)



L2.CUX2.MEIS2

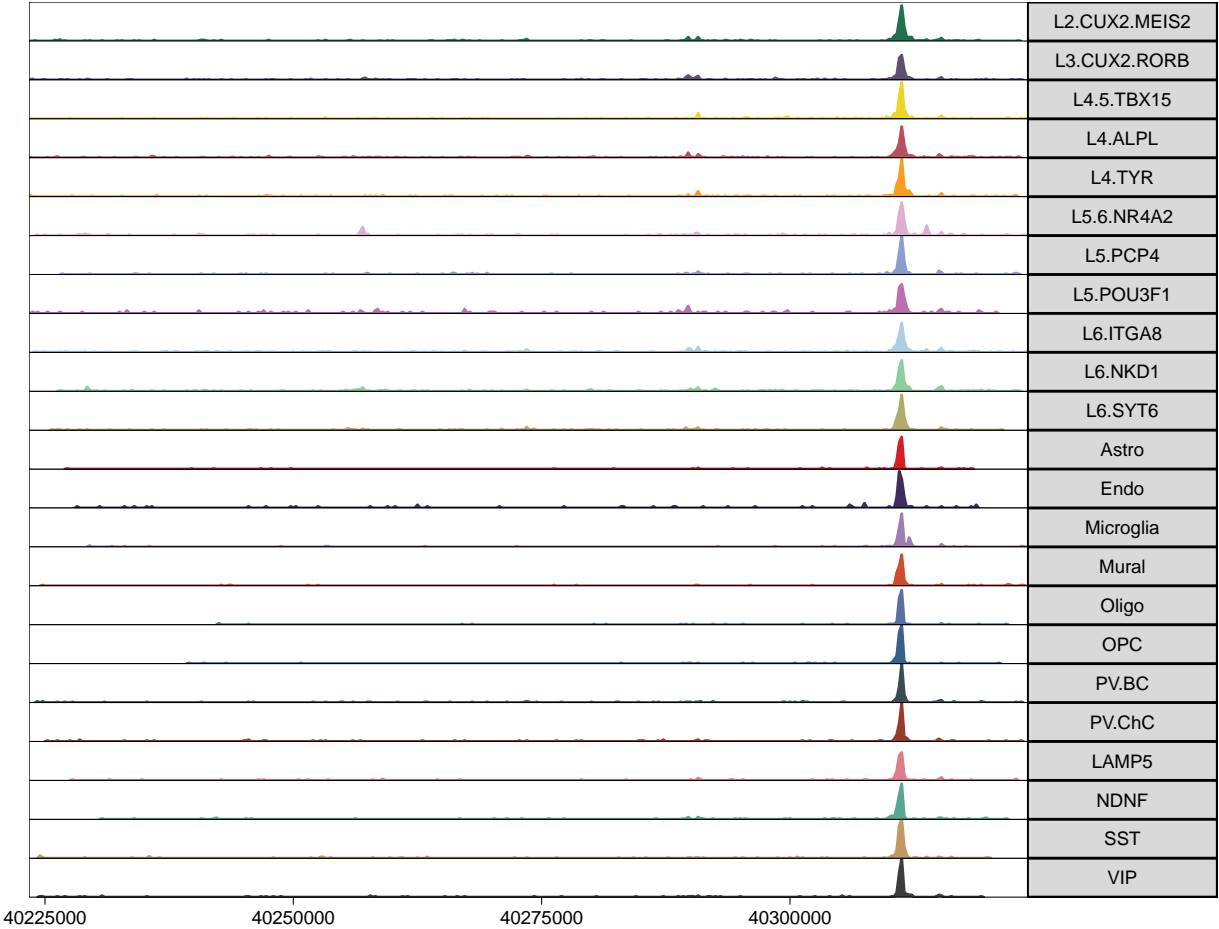


Peaks

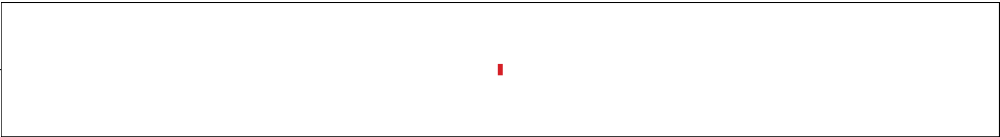


Genes

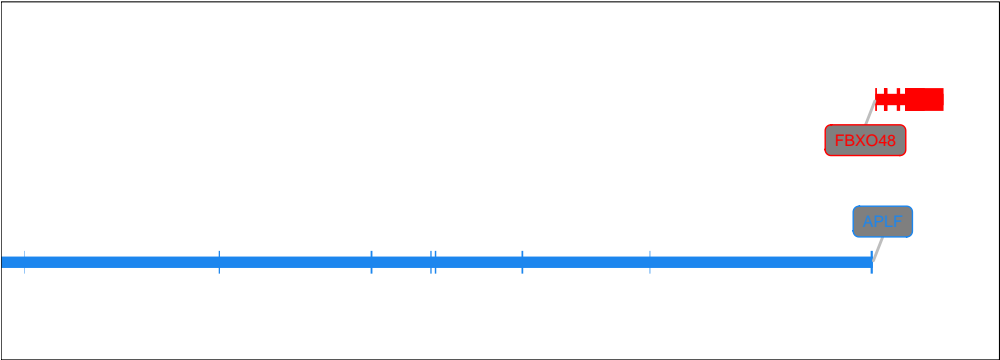
Coverage
(Norm. ATAC Signal Range (0–1.97) by ReadsInTSS)



L2.CUX2.MEIS2



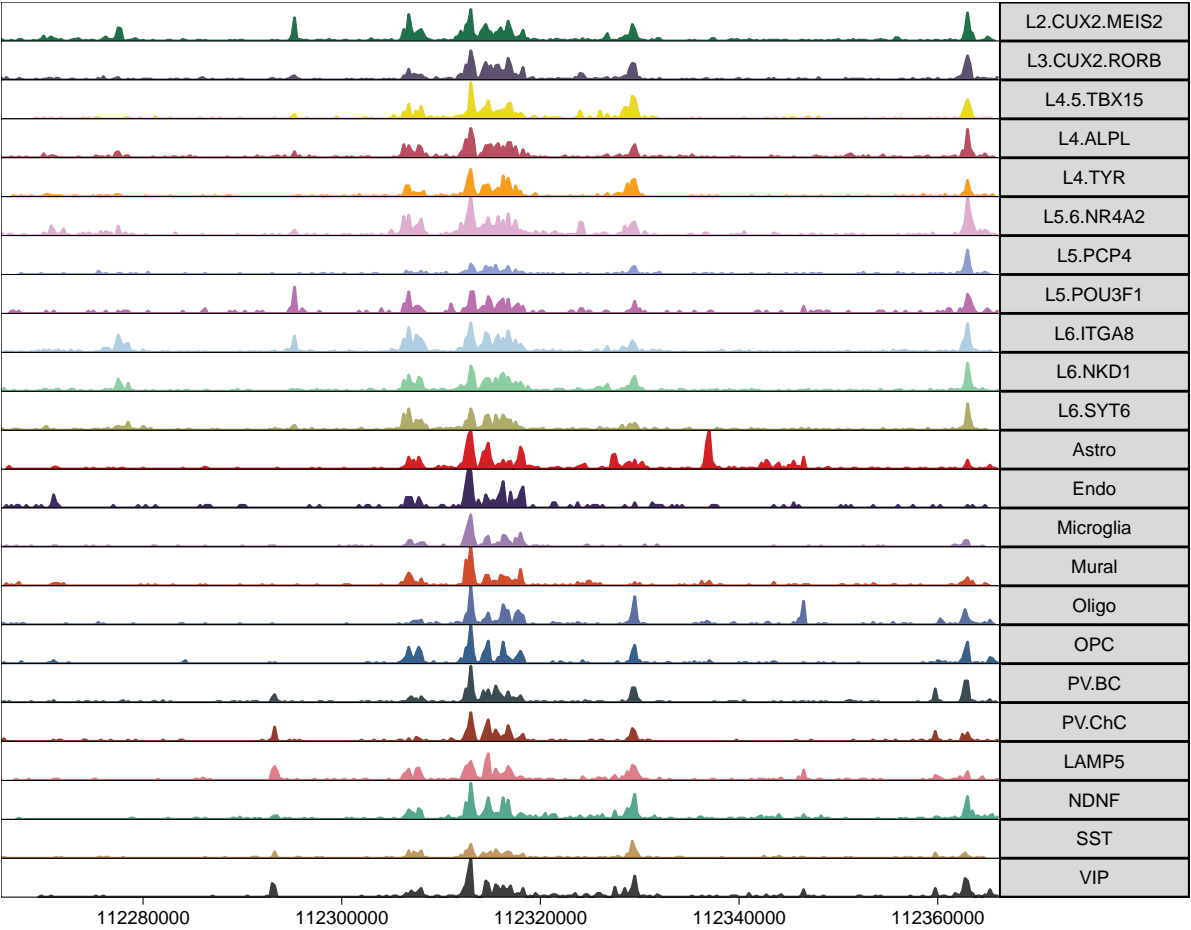
Peaks



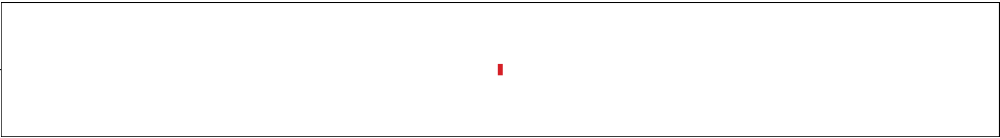
Genes

chr15:112265712–112366213 Marker Peak Tracks for: L2.CUX2.MEIS2

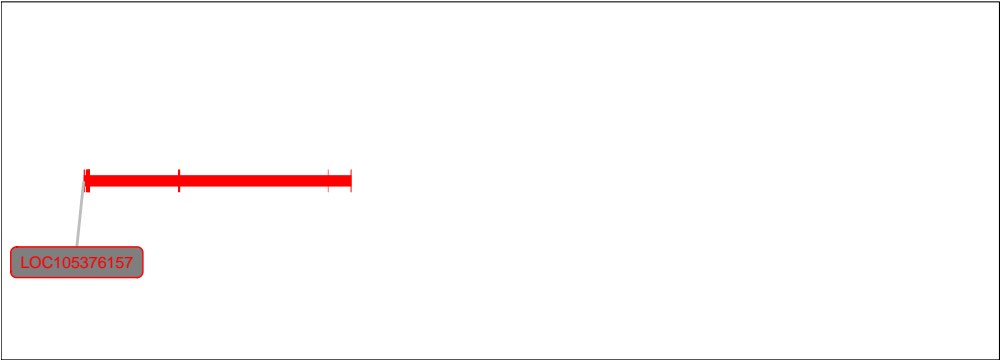
Coverage
(Norm. ATAC Signal Range (0–1.19) by ReadsInTSS)



L2.CUX2.MEIS2

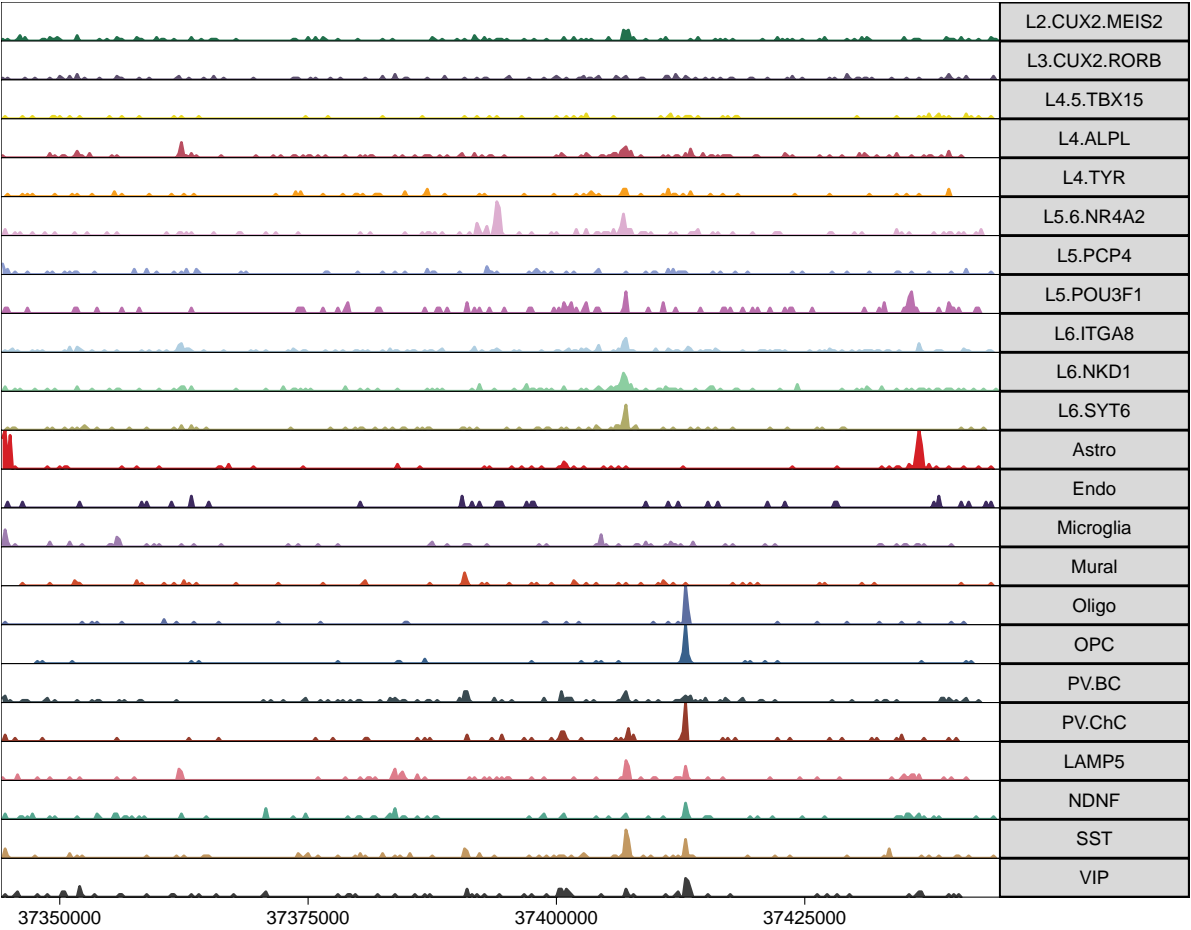


Peaks

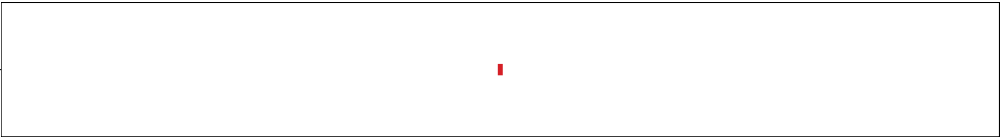


Genes

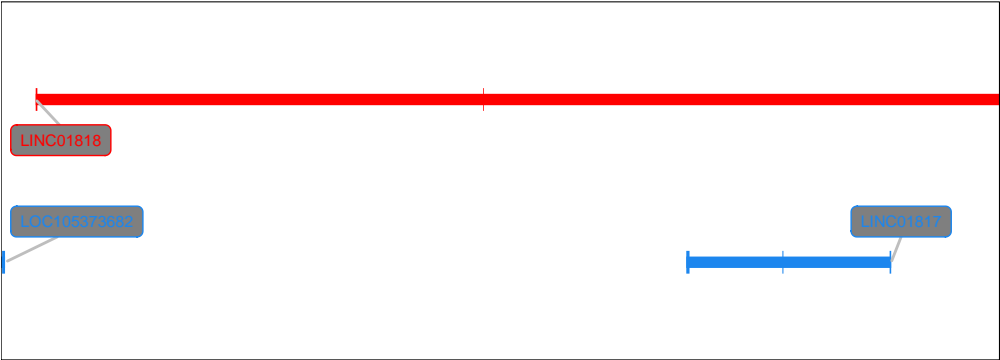
Coverage
(Norm. ATAC Signal Range (0–0.53) by ReadsInTSS)



L2.CUX2.MEIS2



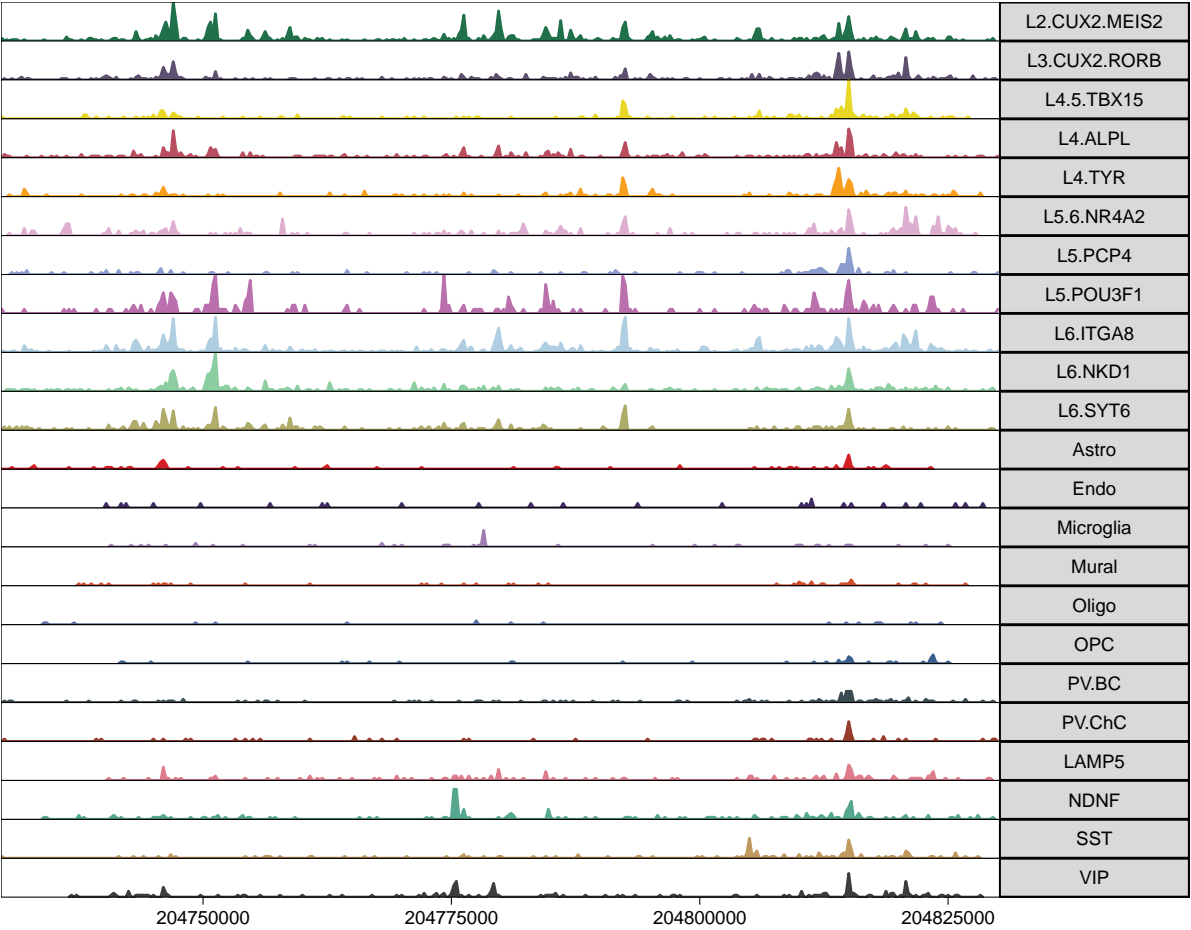
Peaks



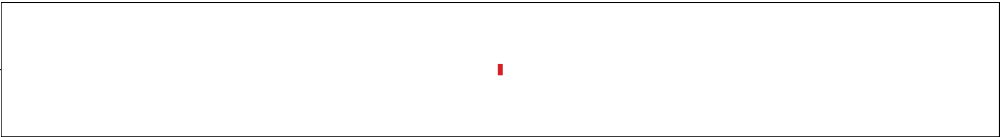
Genes

chr1:204729670–204830171 Marker Peak Tracks for: L2.CUX2.MEIS2

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



L2.CUX2.MEIS2

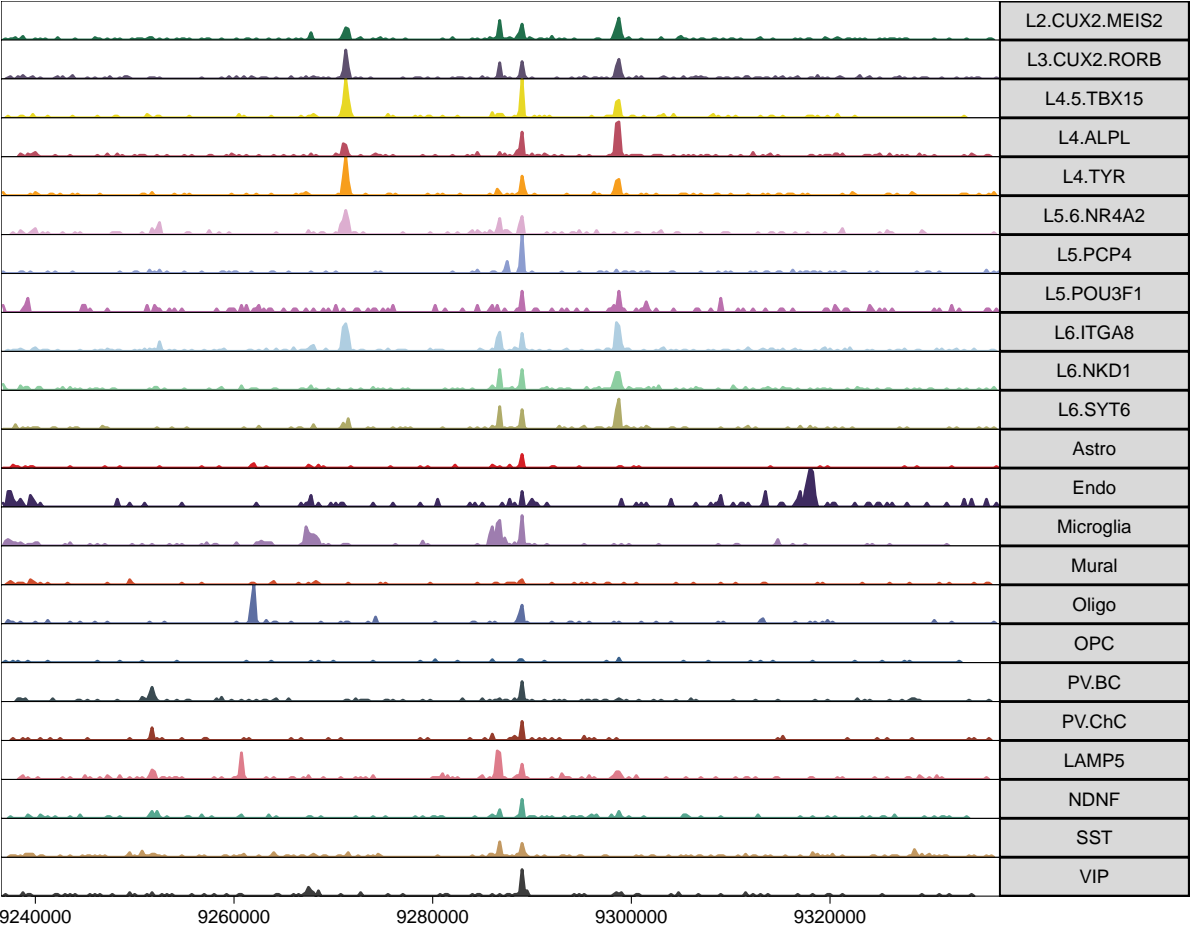


Peaks

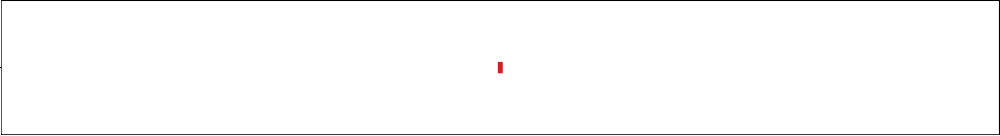


chr15:9236555-9337056 Marker Peak Tracks for: L2.CUX2.MEIS2

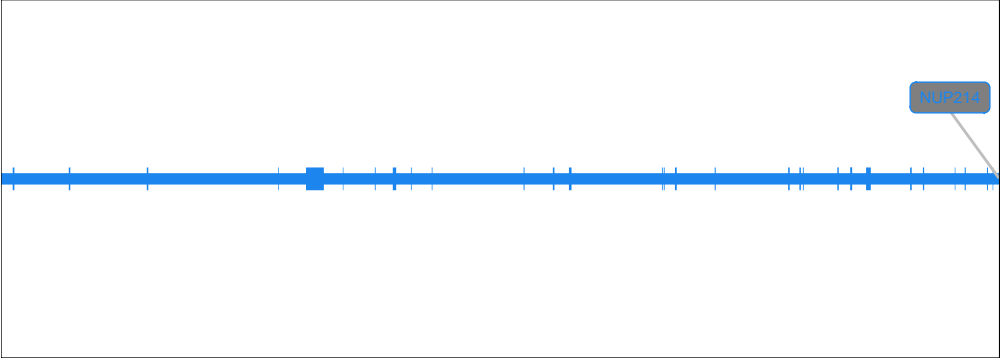
Coverage
(Norm. ATAC Signal Range (0-0.81) by ReadsInTSS)



L2.CUX2.MEIS2



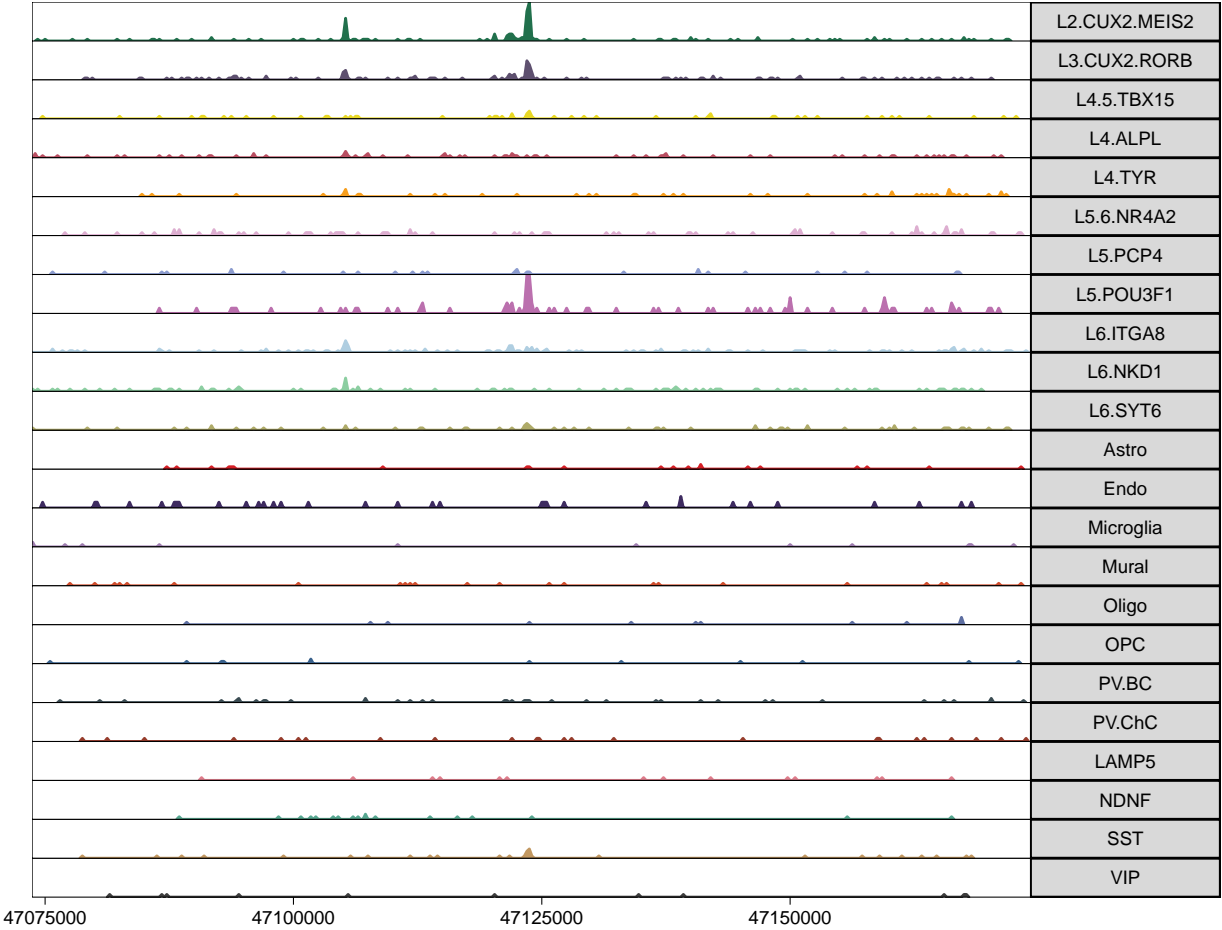
Peaks



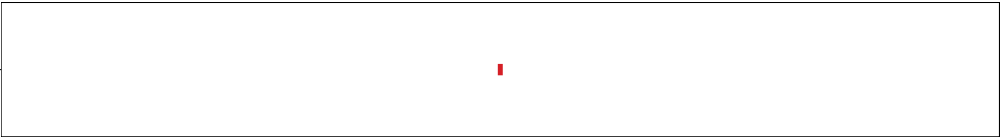
Genes

chr1:47073692–47174193 Marker Peak Tracks for: L2.CUX2.MEIS2

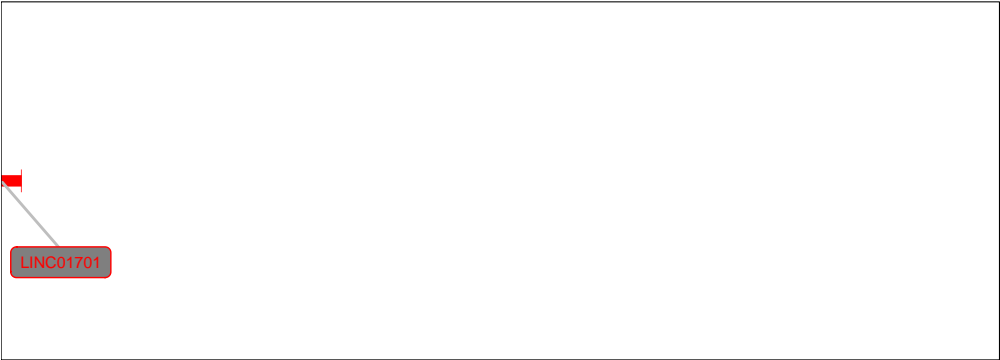
Coverage
(Norm. ATAC Signal Range (0–0.54) by ReadsInTSS)



L2.CUX2.MEIS2

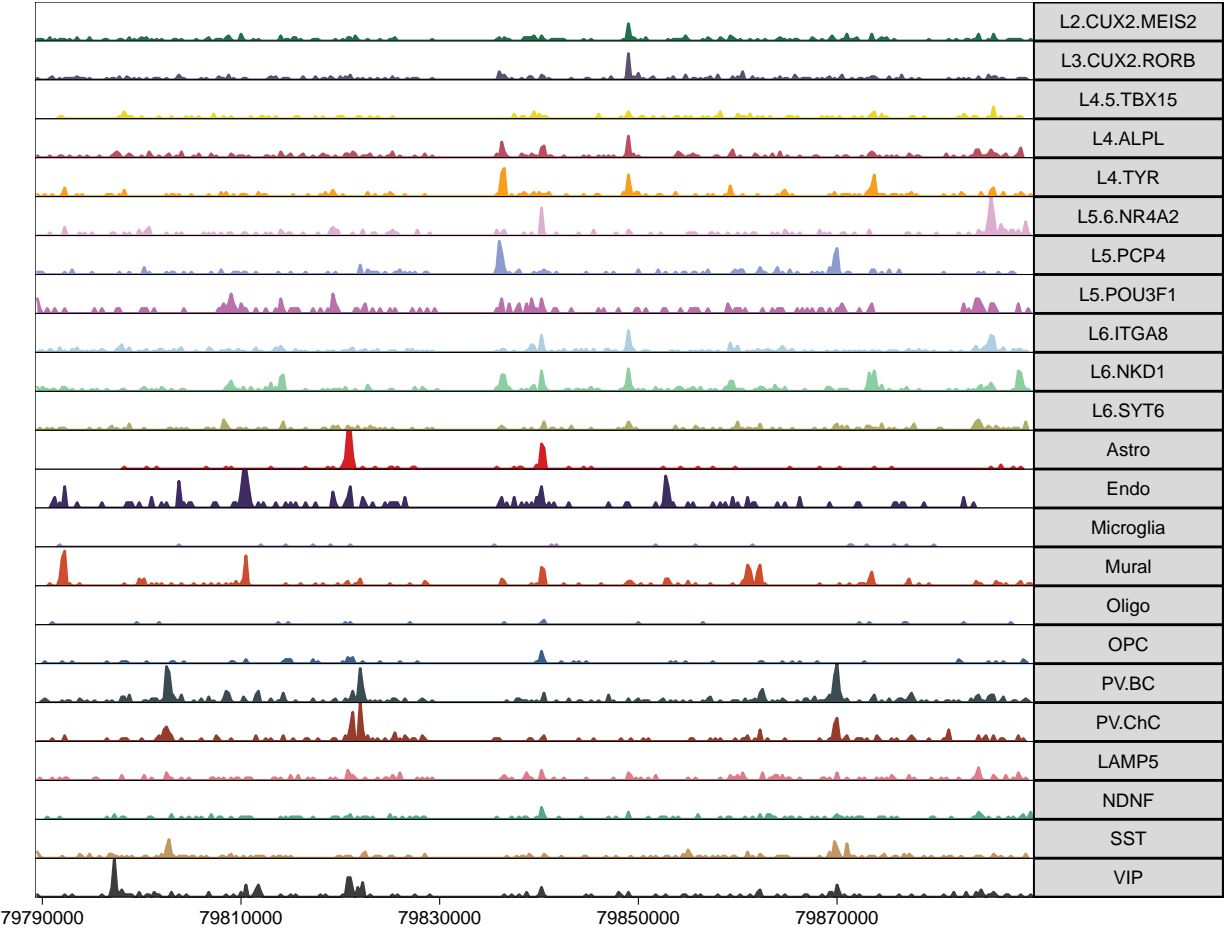


Peaks

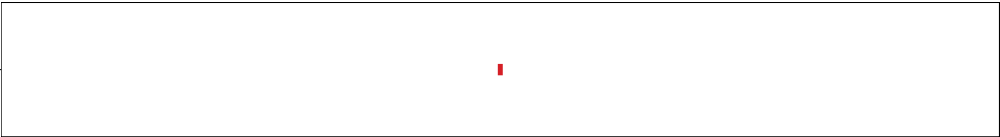


Genes

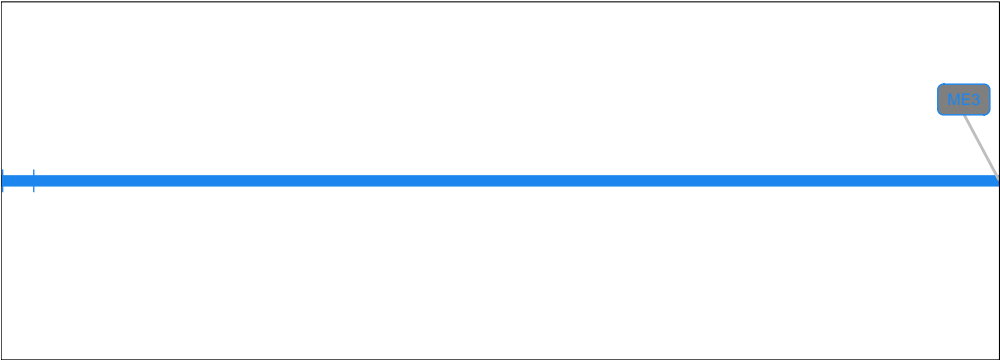
Coverage
(Norm. ATAC Signal Range (0–0.59) by ReadsInTSS)



L2.CUX2.MEIS2

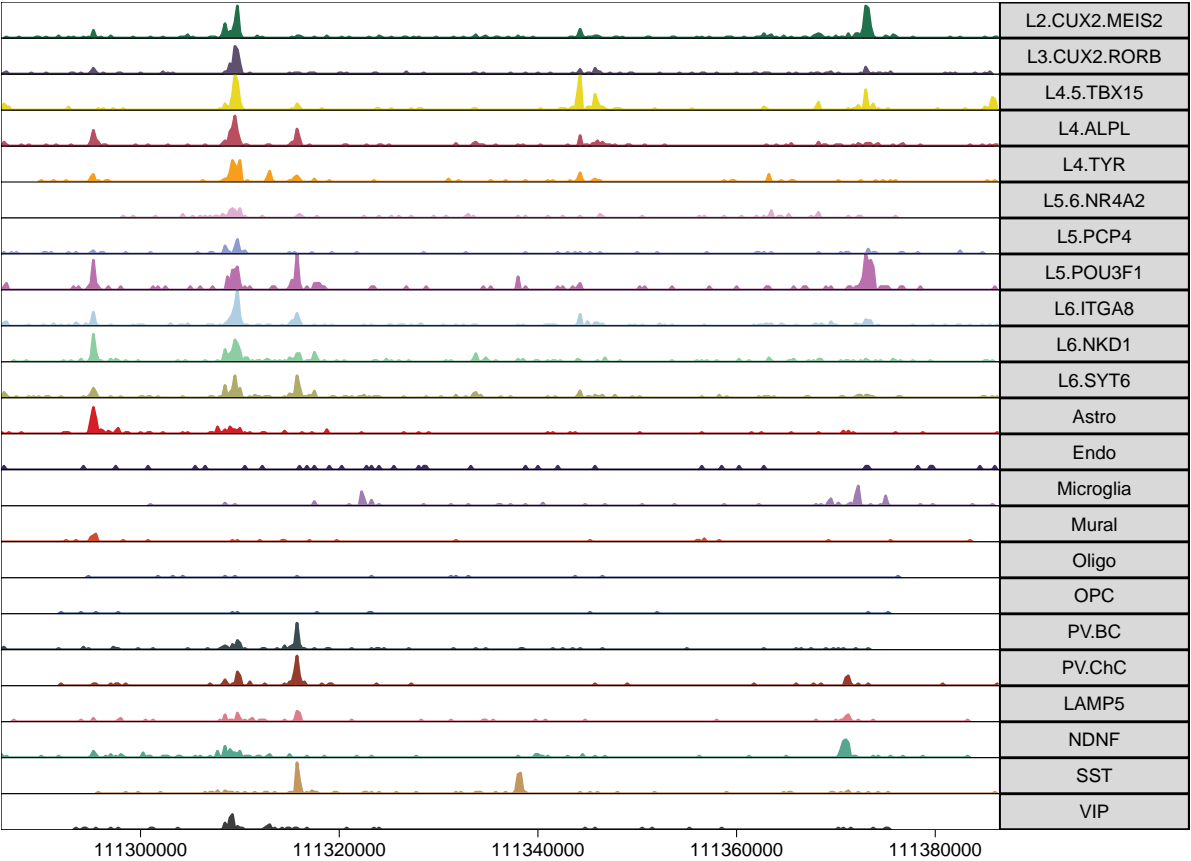


Peaks

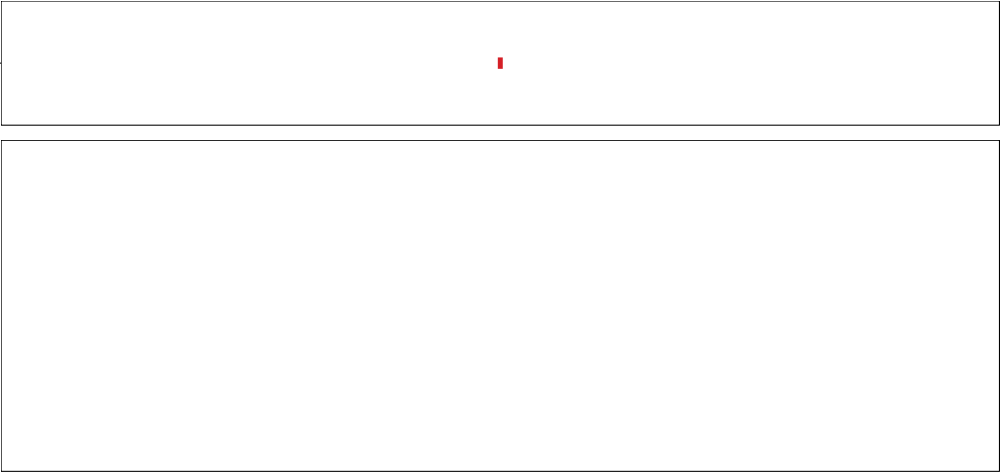


Genes

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



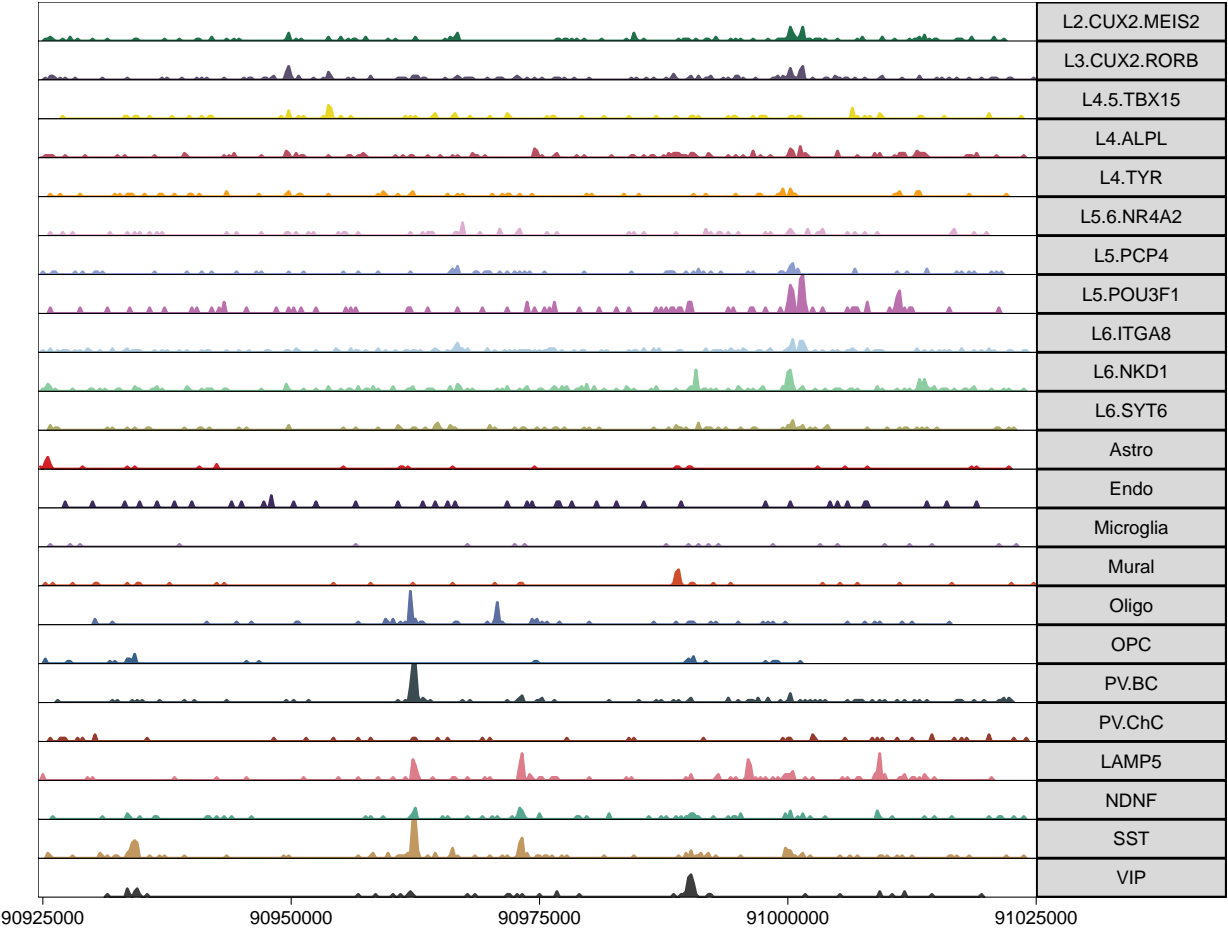
L2.CUX2.MEIS2



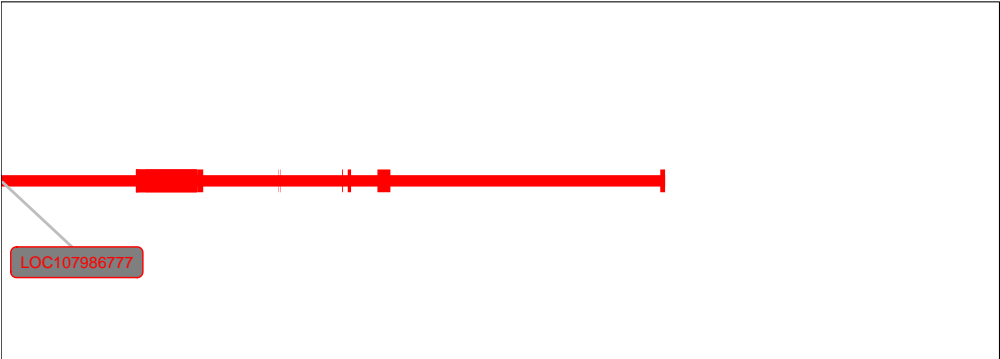
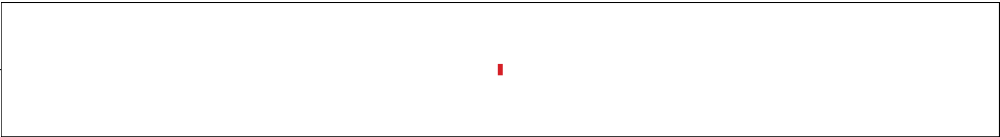
Peaks

GeneTrack

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

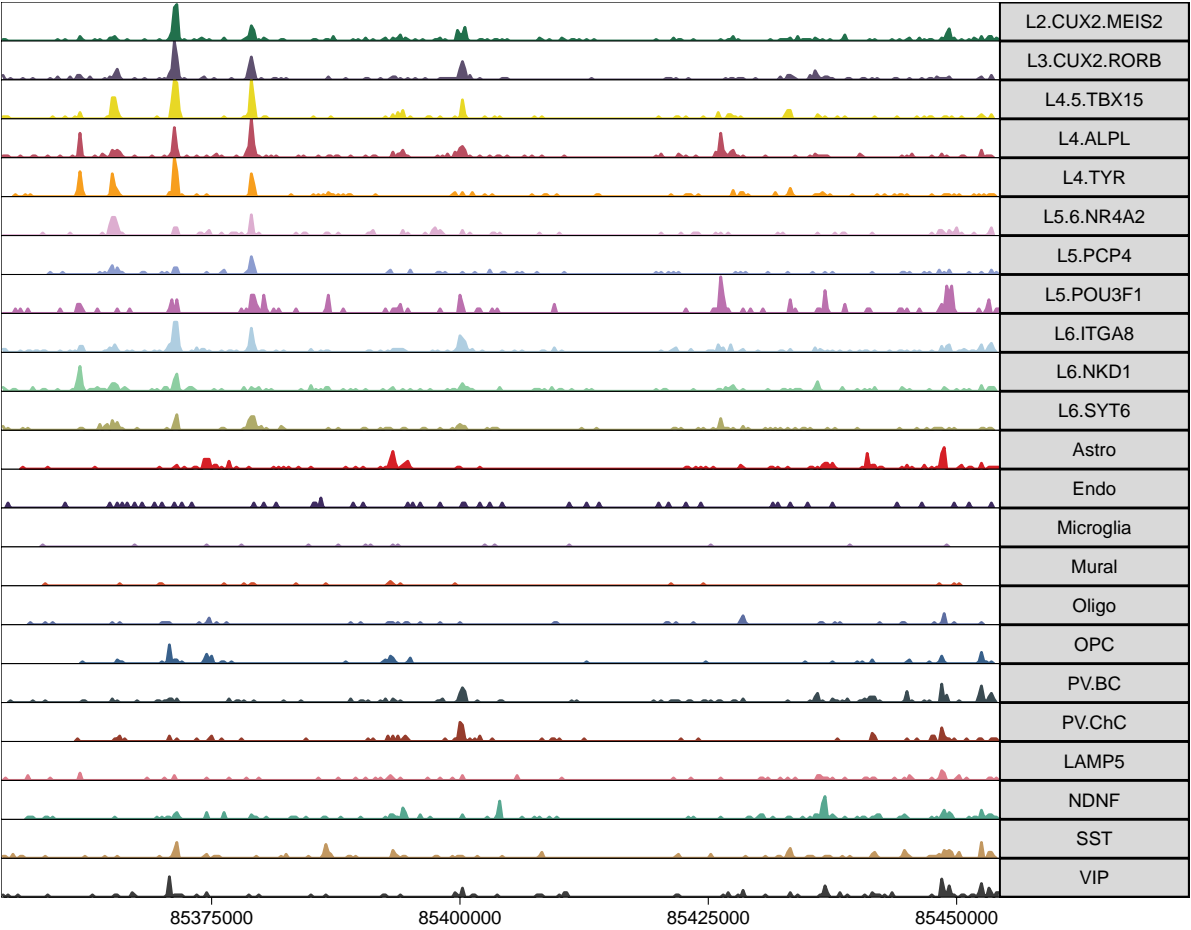


L2.CUX2.MEIS2

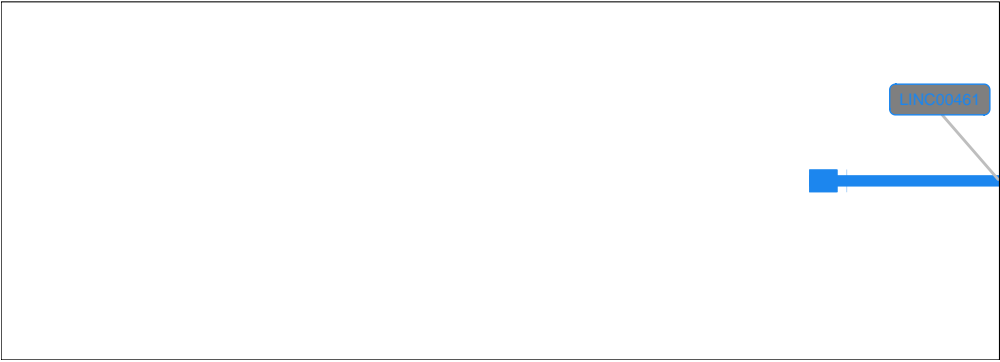
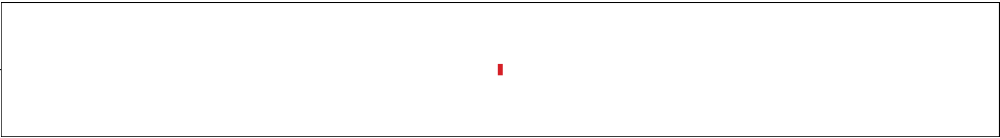


chr6:85353809–85454310 Marker Peak Tracks for: L2.CUX2.MEIS2

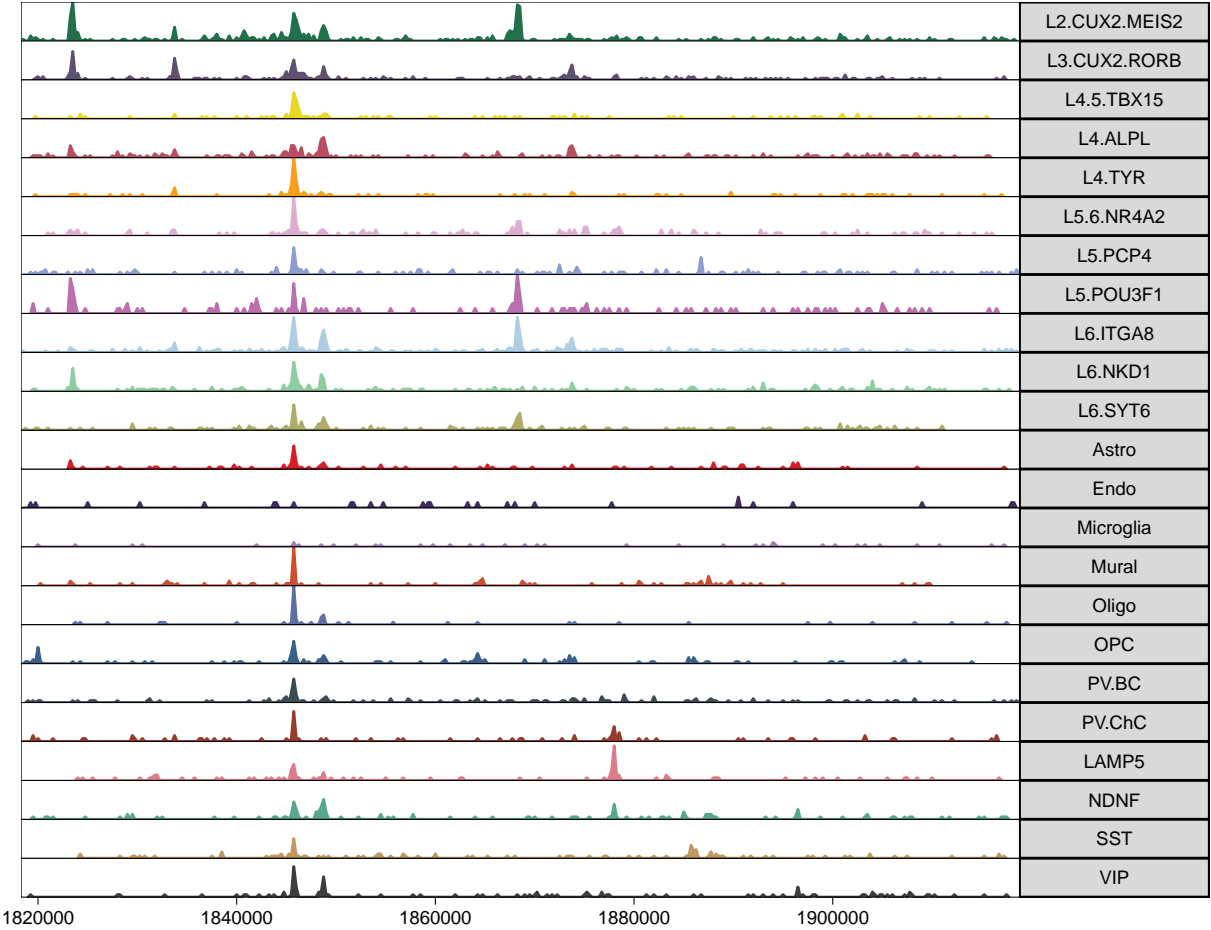
Coverage
(Norm. ATAC Signal Range (0–0.63) by ReadsInTSS)



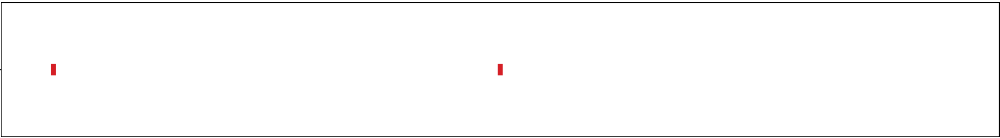
L2.CUX2.MEIS2



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



L2.CUX2.MEIS2

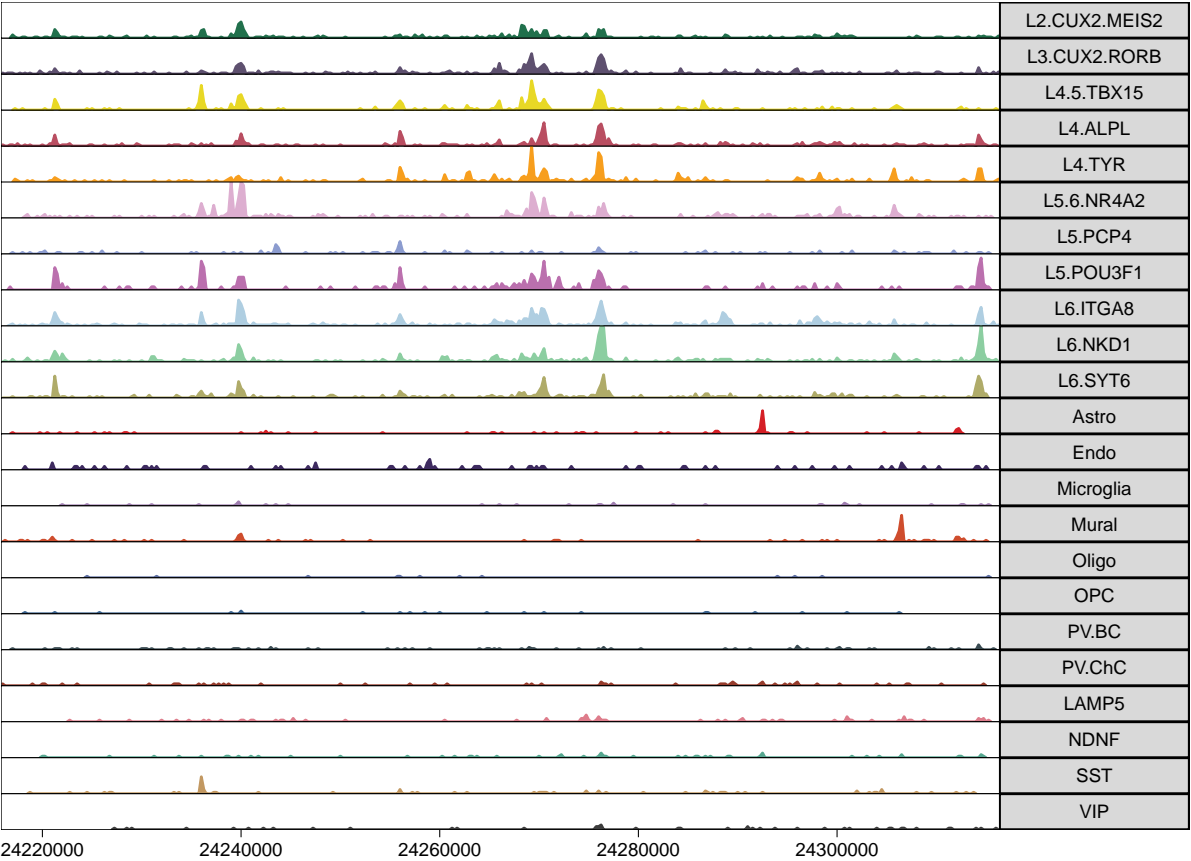


Peaks

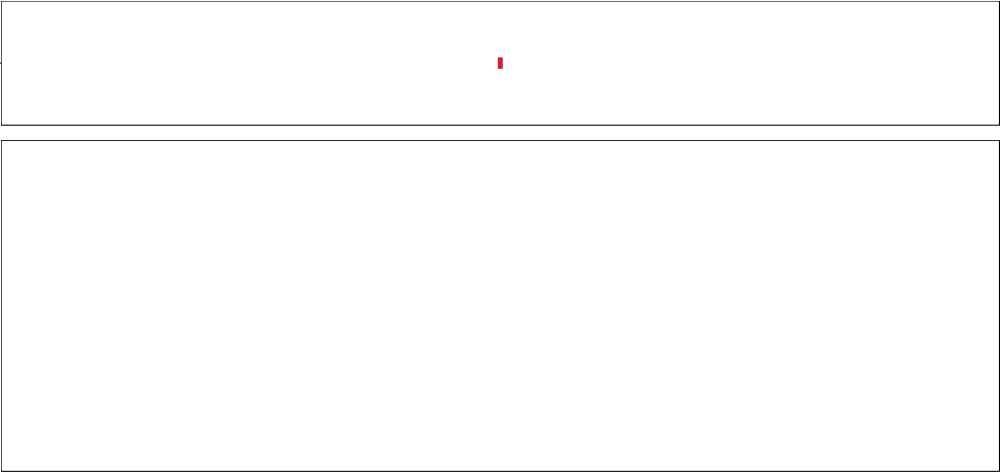


Genes

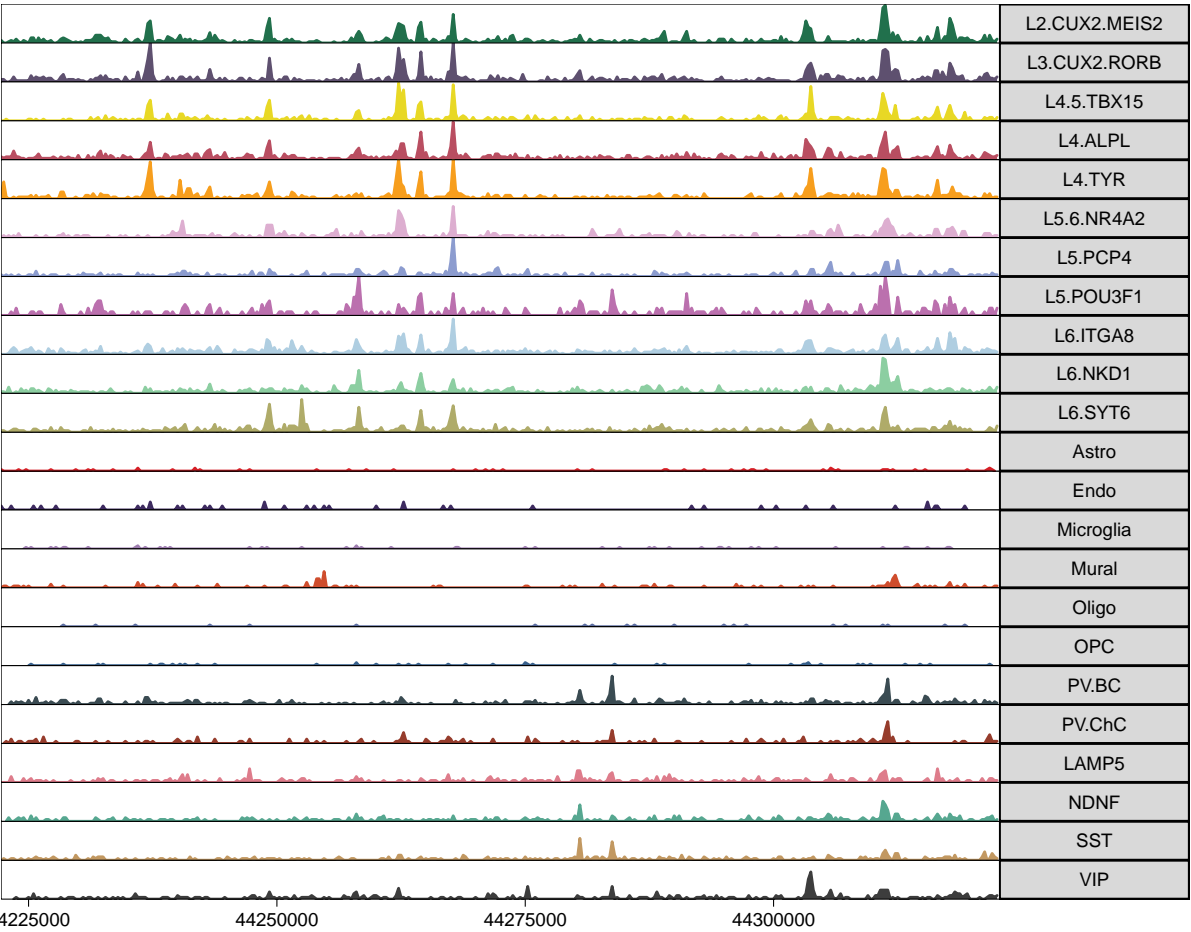
Coverage
(Norm. ATAC Signal Range (0–0.83) by ReadsInTSS)



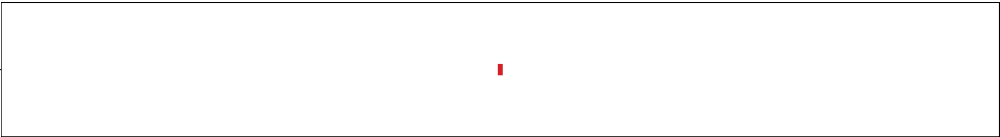
L2.CUX2.MEIS2



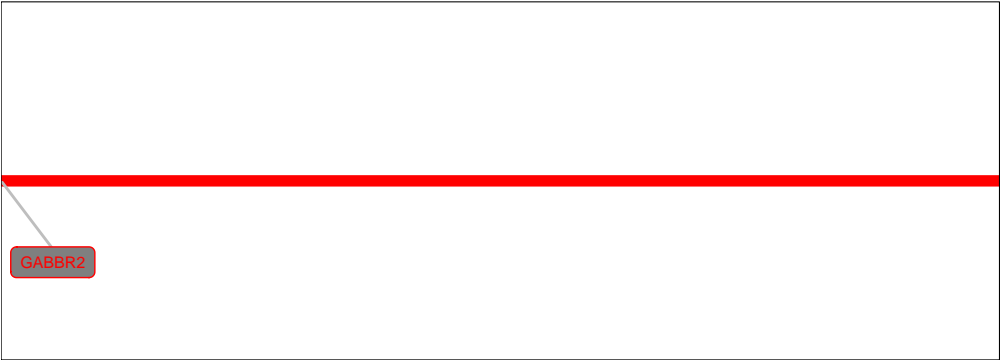
Coverage
(Norm. ATAC Signal Range (0-0.78) by ReadsInTSS)



L2.CUX2.MEIS2

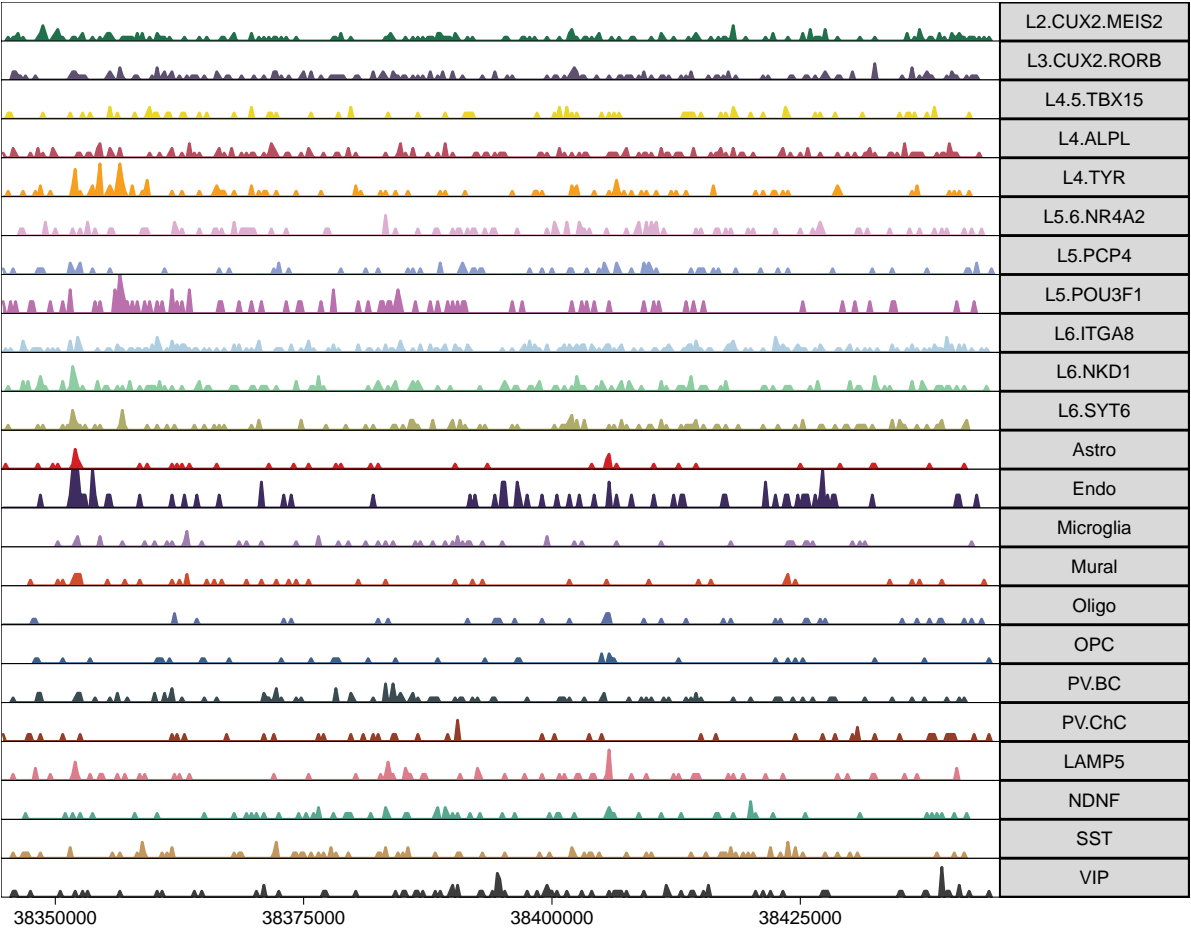


Peaks

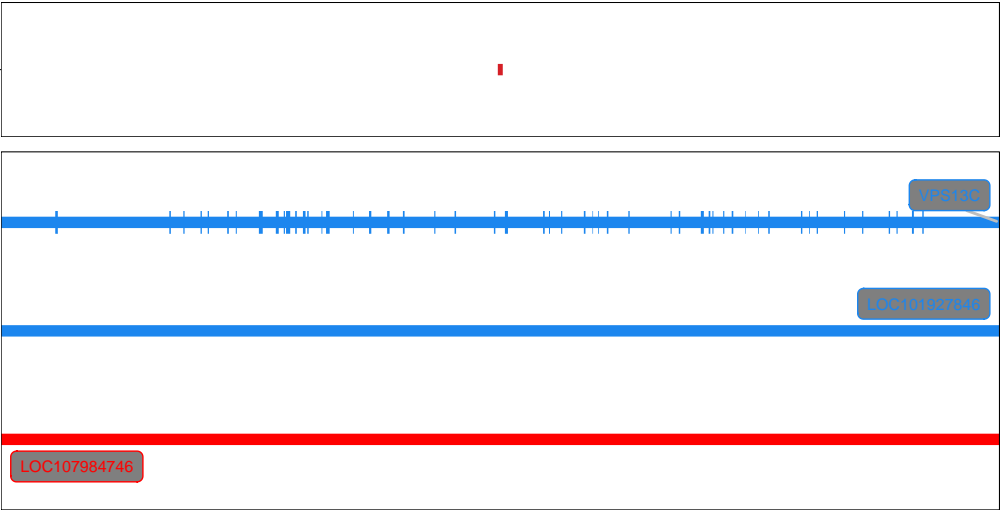


Genes

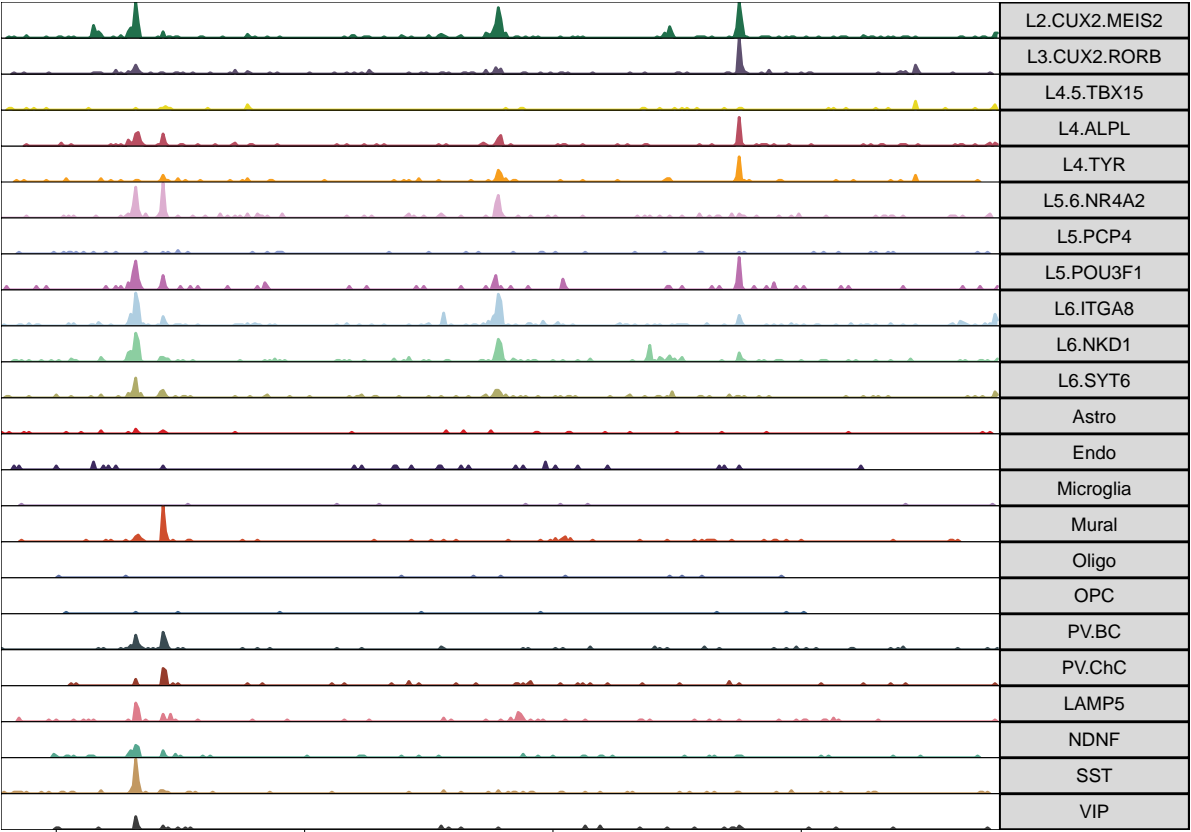
Coverage
(Norm. ATAC Signal Range (0-0.24) by ReadsInTSS)



L2.CUX2.MEIS2



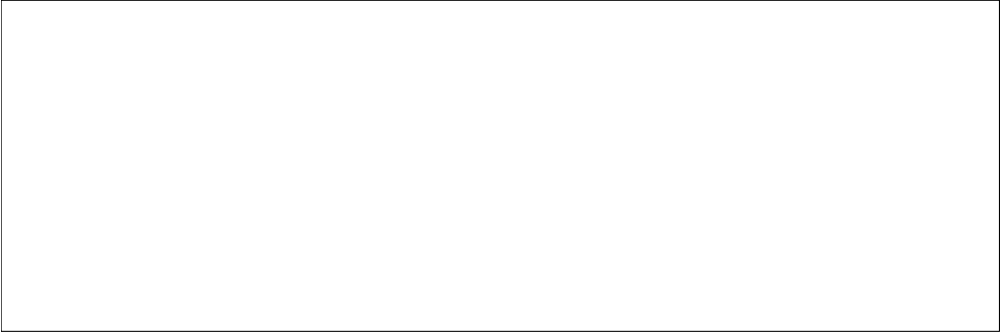
Coverage
(Norm. ATAC Signal Range (0-0.73) by ReadsInTSS)



L2.CUX2.MEIS2



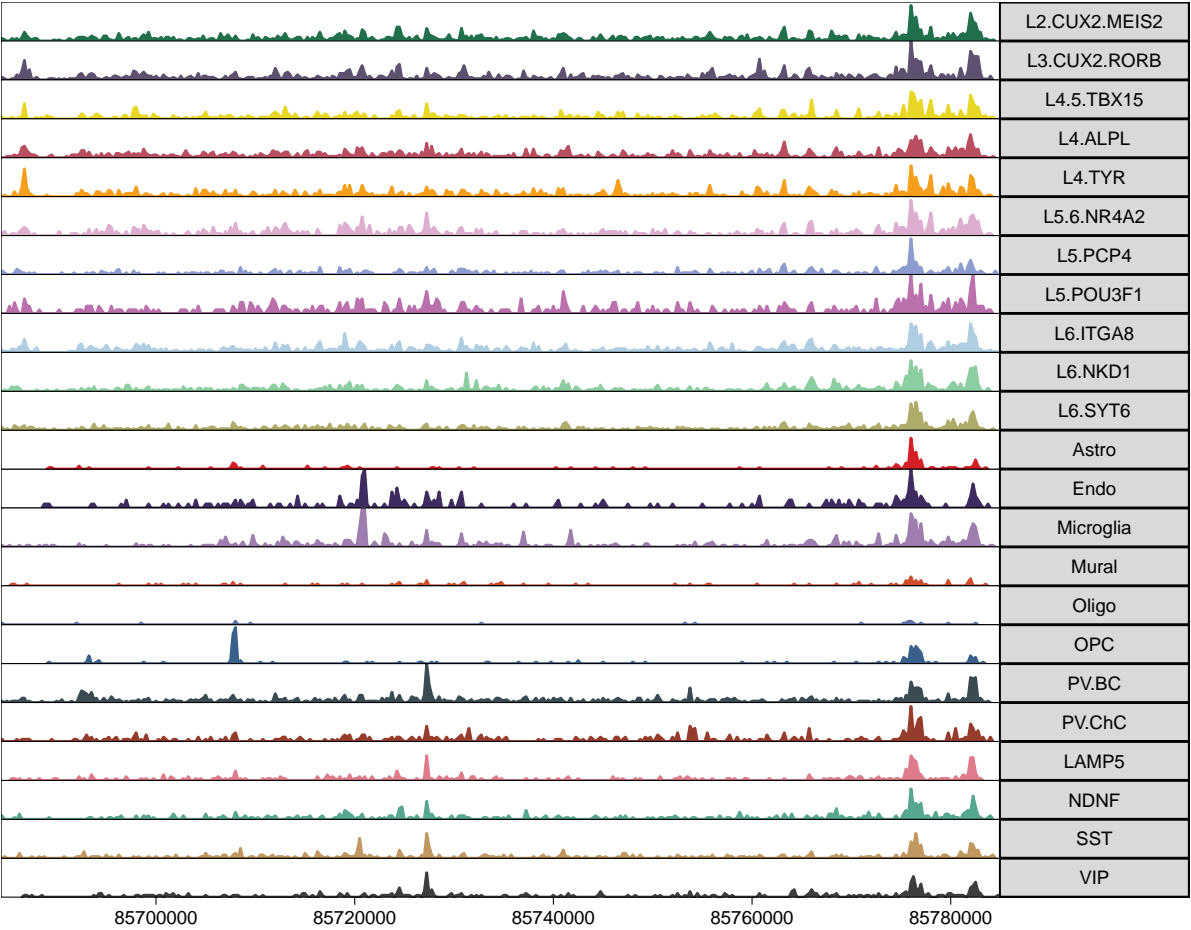
Peaks



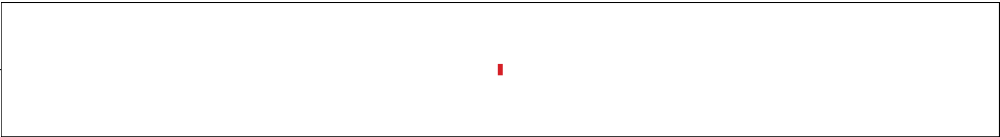
GeneTrack

chr6:85684405–85784906 Marker Peak Tracks for: L2.CUX2.MEIS2

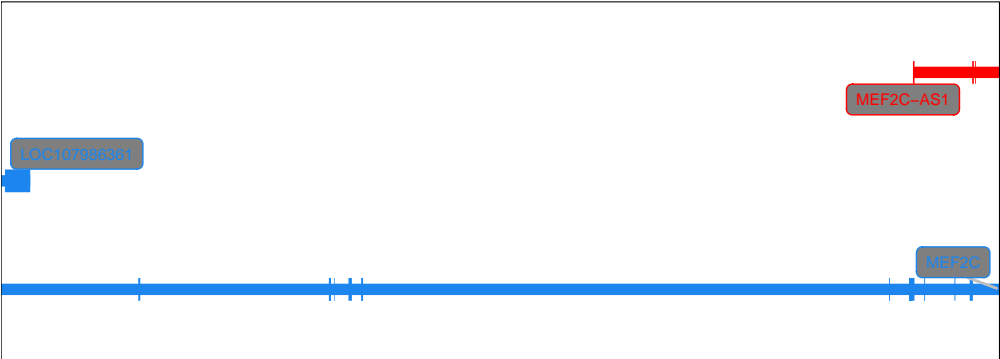
Coverage
(Norm. ATAC Signal Range (0–0.78) by ReadsInTSS)



L2.CUX2.MEIS2

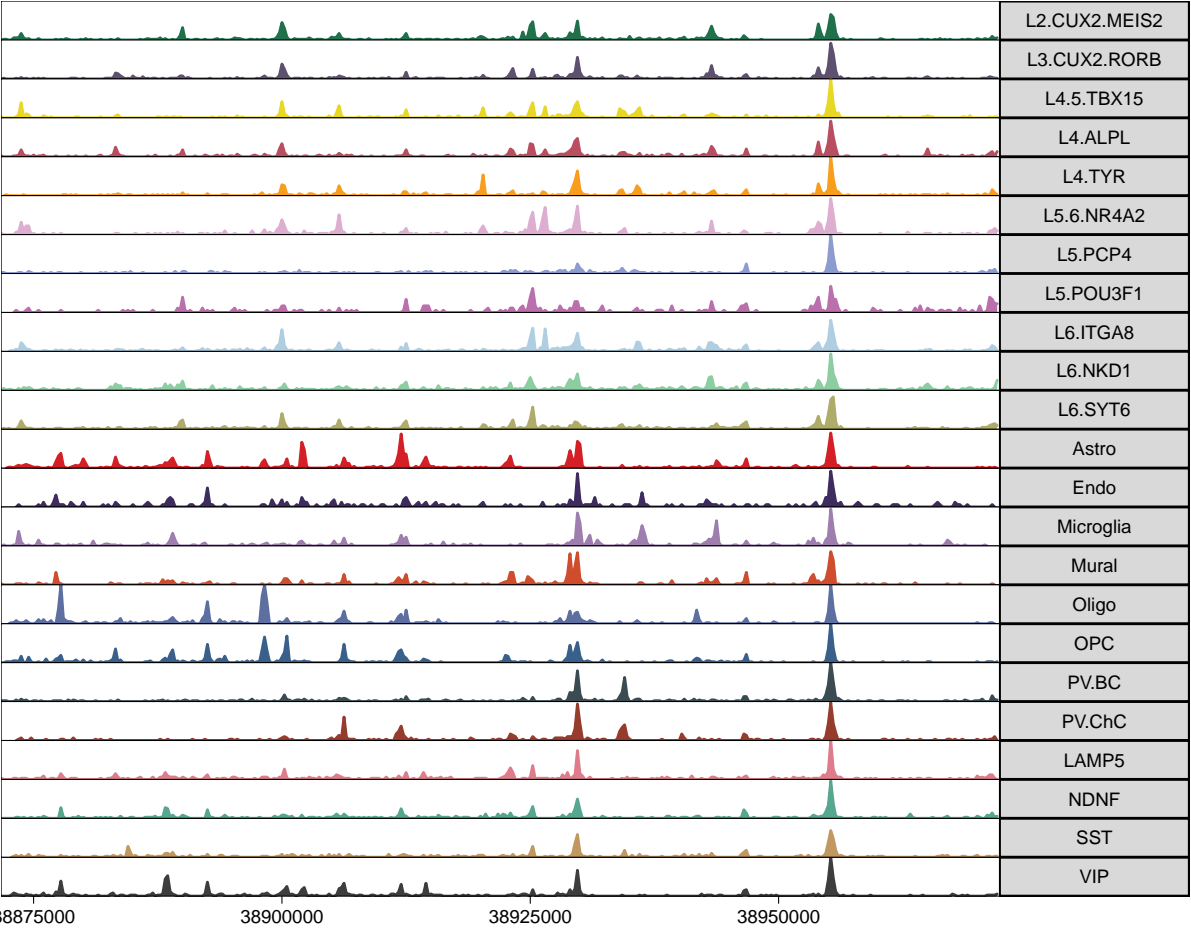


Peaks

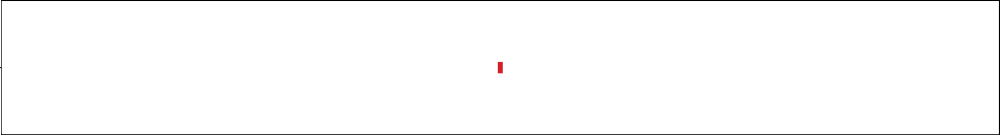


Genes

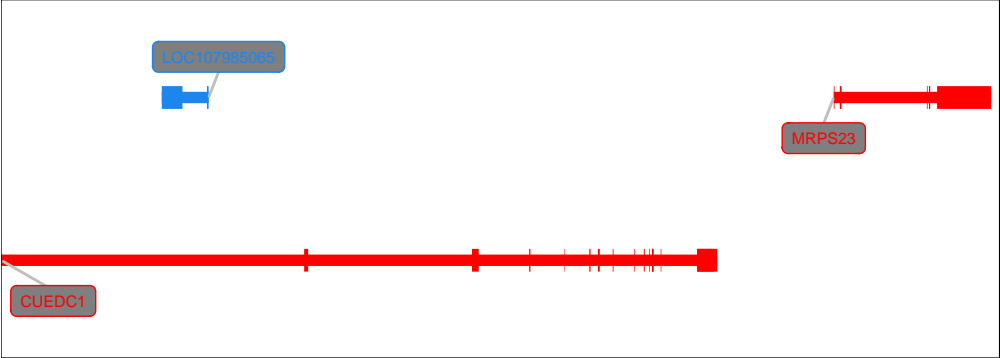
Coverage
(Norm. ATAC Signal Range (0–1.31) by ReadsInTSS)



L2.CUX2.MEIS2



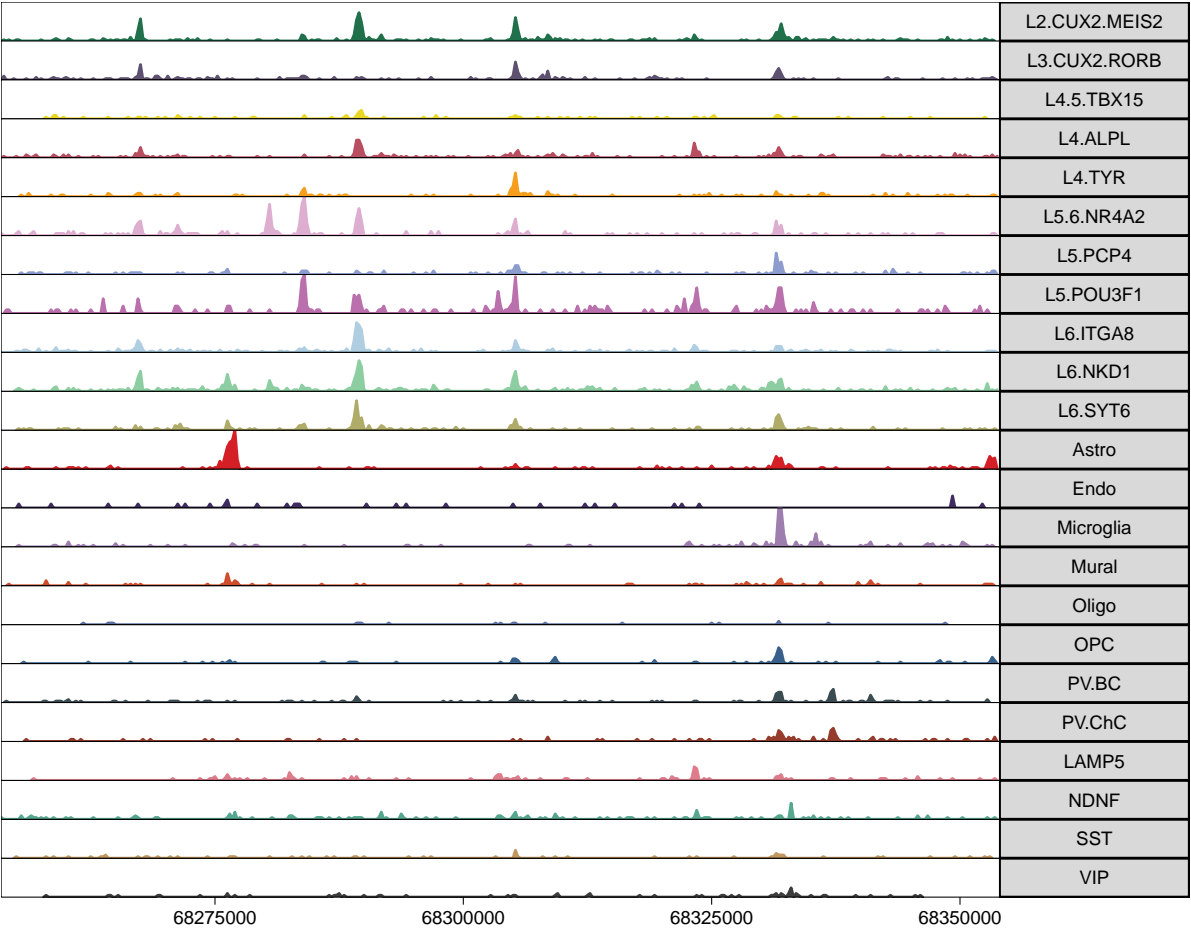
Peaks



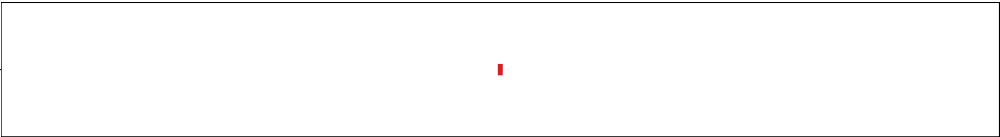
Genes

chr16:68253472–68353973 Marker Peak Tracks for: L2.CUX2.MEIS2

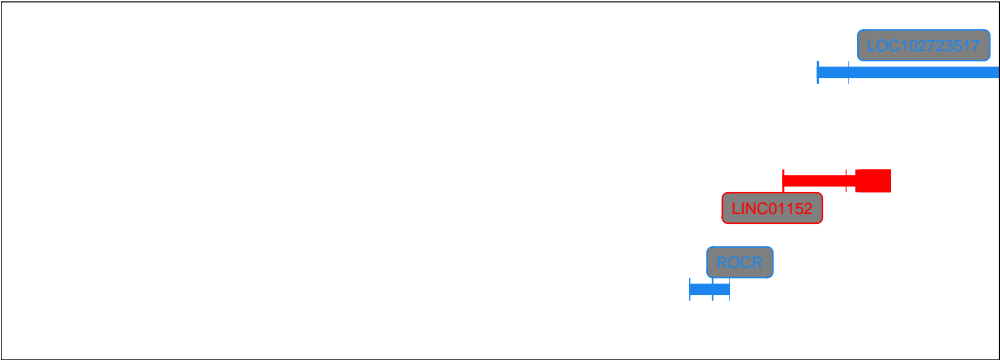
Coverage
(Norm. ATAC Signal Range (0–0.78) by ReadsInTSS)



L2.CUX2.MEIS2



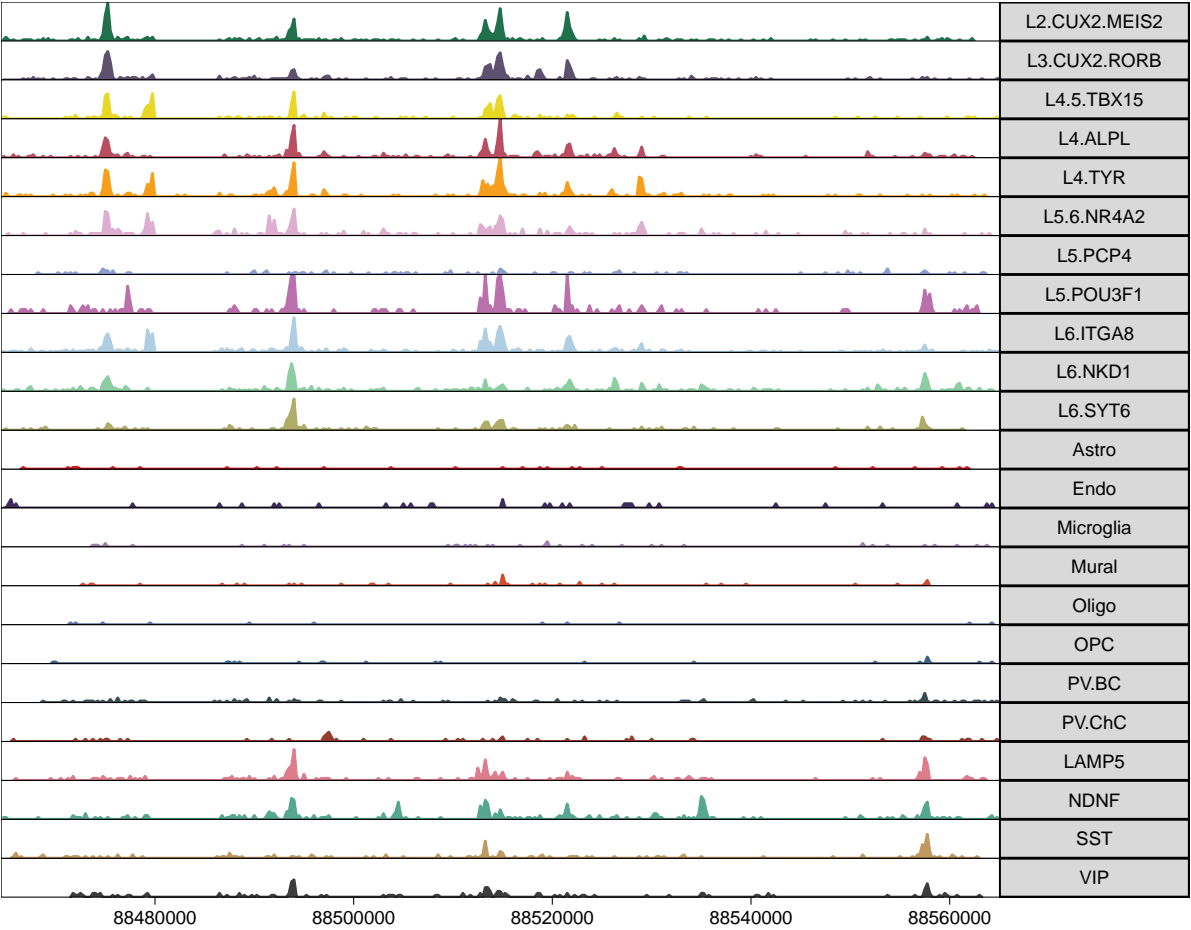
Peaks



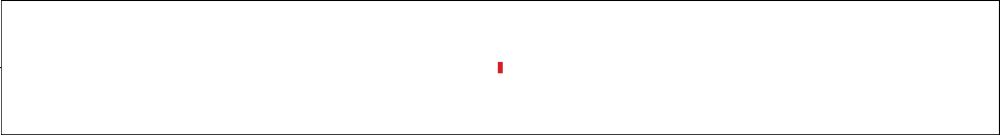
Genes

chr3:88464511–88565012 Marker Peak Tracks for: L2.CUX2.MEIS2

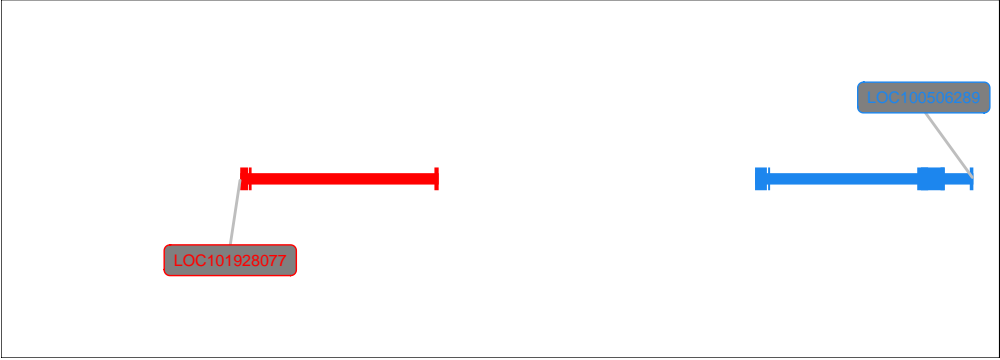
Coverage
(Norm. ATAC Signal Range (0–0.73) by ReadsInTSS)



L2.CUX2.MEIS2

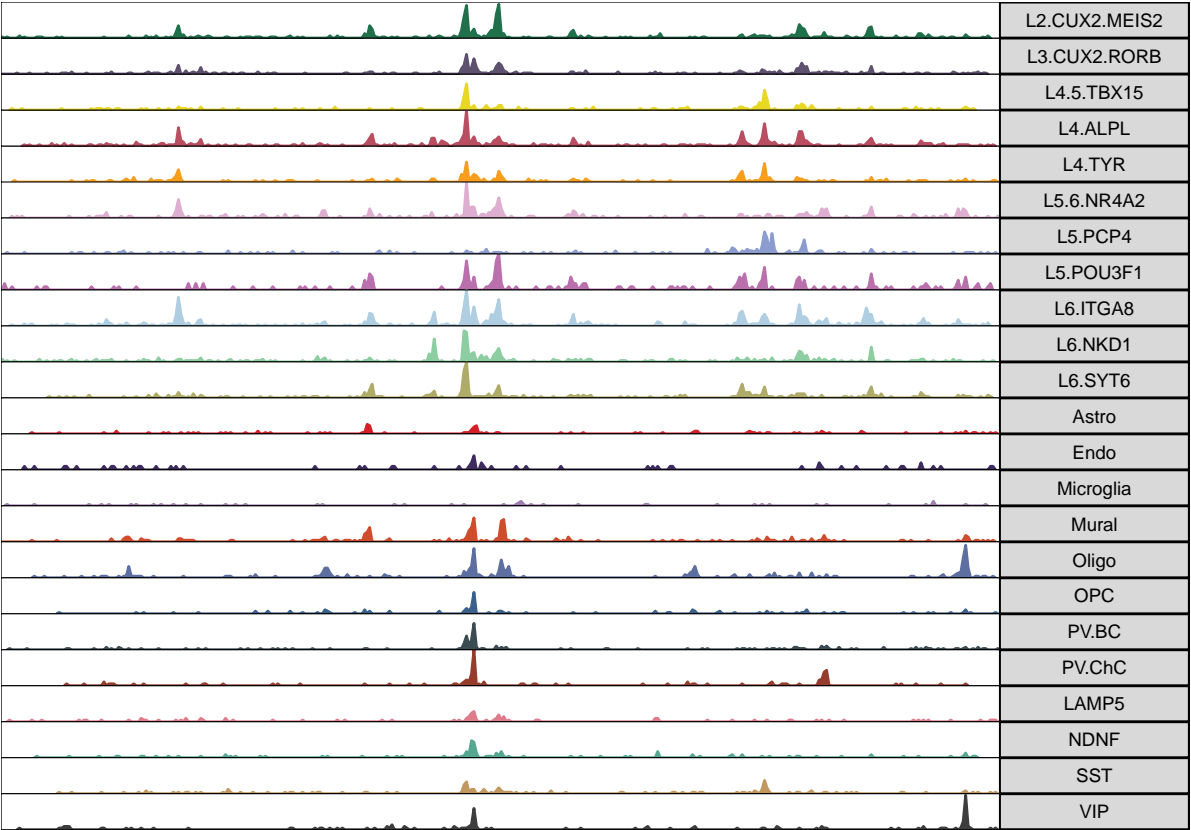


Peaks

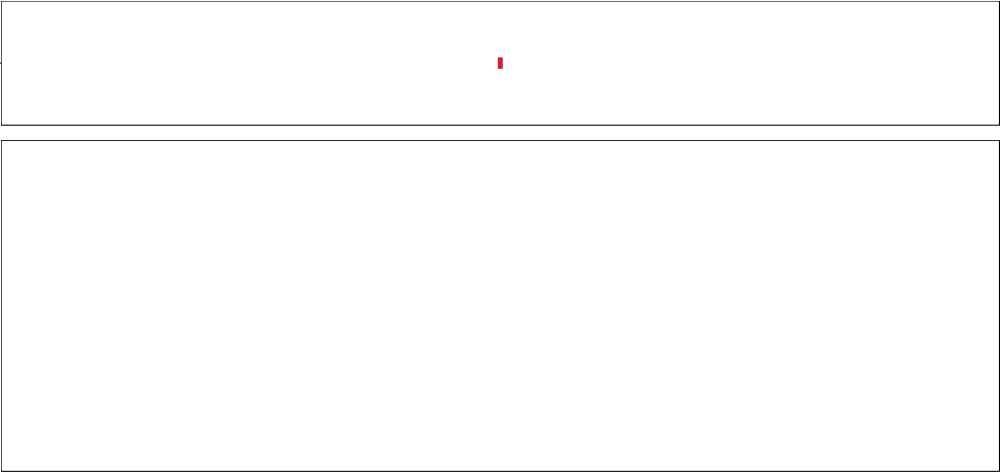


Genes

Coverage
(Norm. ATAC Signal Range (0–0.82) by ReadsInTSS)



L2.CUX2.MEIS2

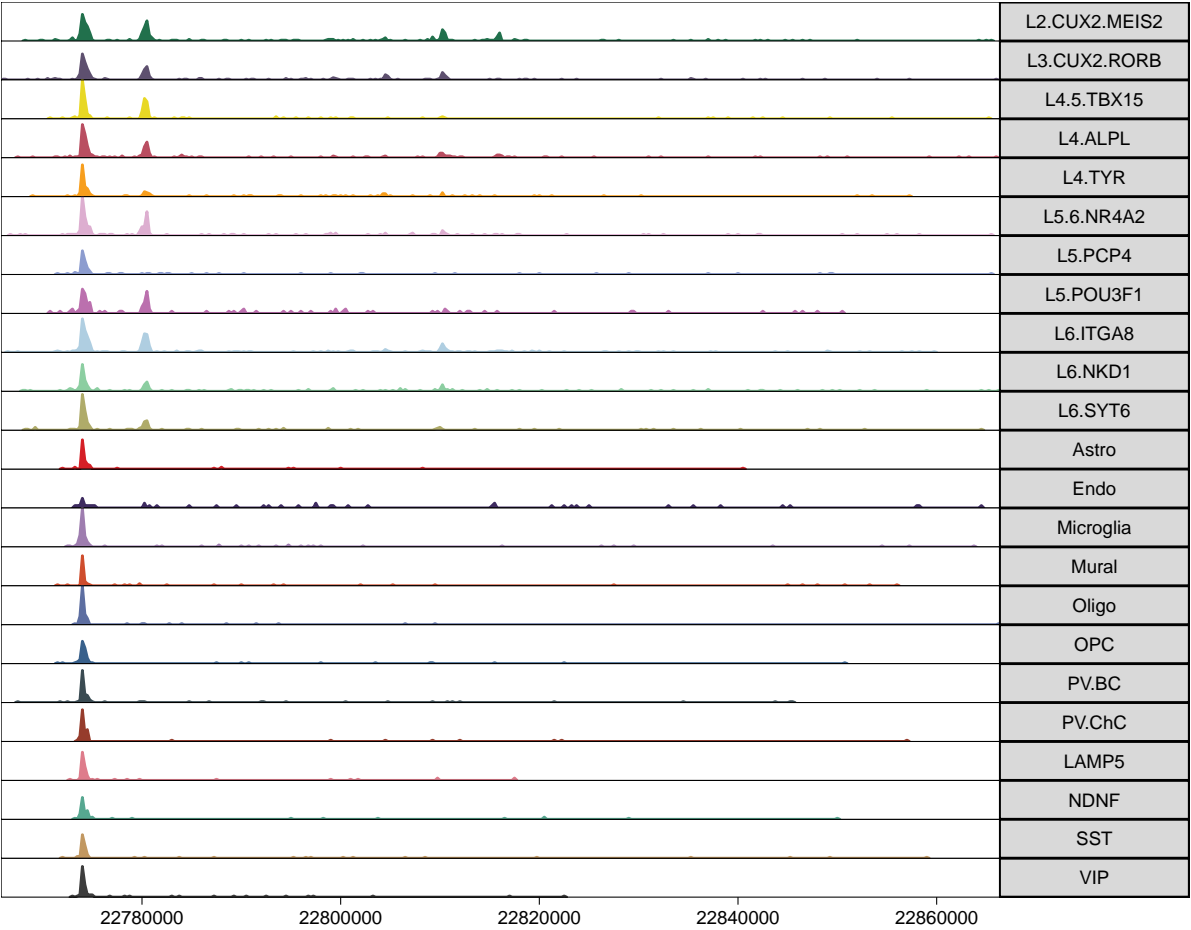


Peaks

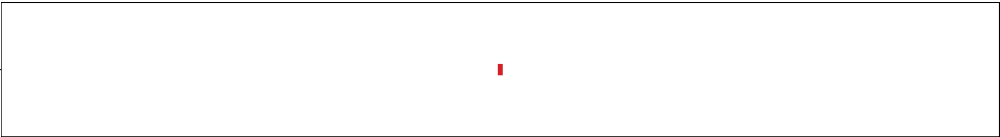
GeneTrack

chr19:22765819–22866320 Marker Peak Tracks for: L2.CUX2.MEIS2

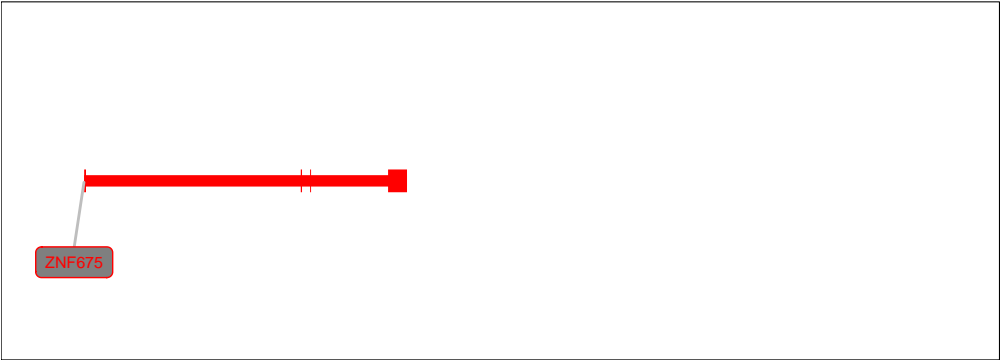
Coverage
(Norm. ATAC Signal Range (0–1.29) by ReadsInTSS)



L2.CUX2.MEIS2



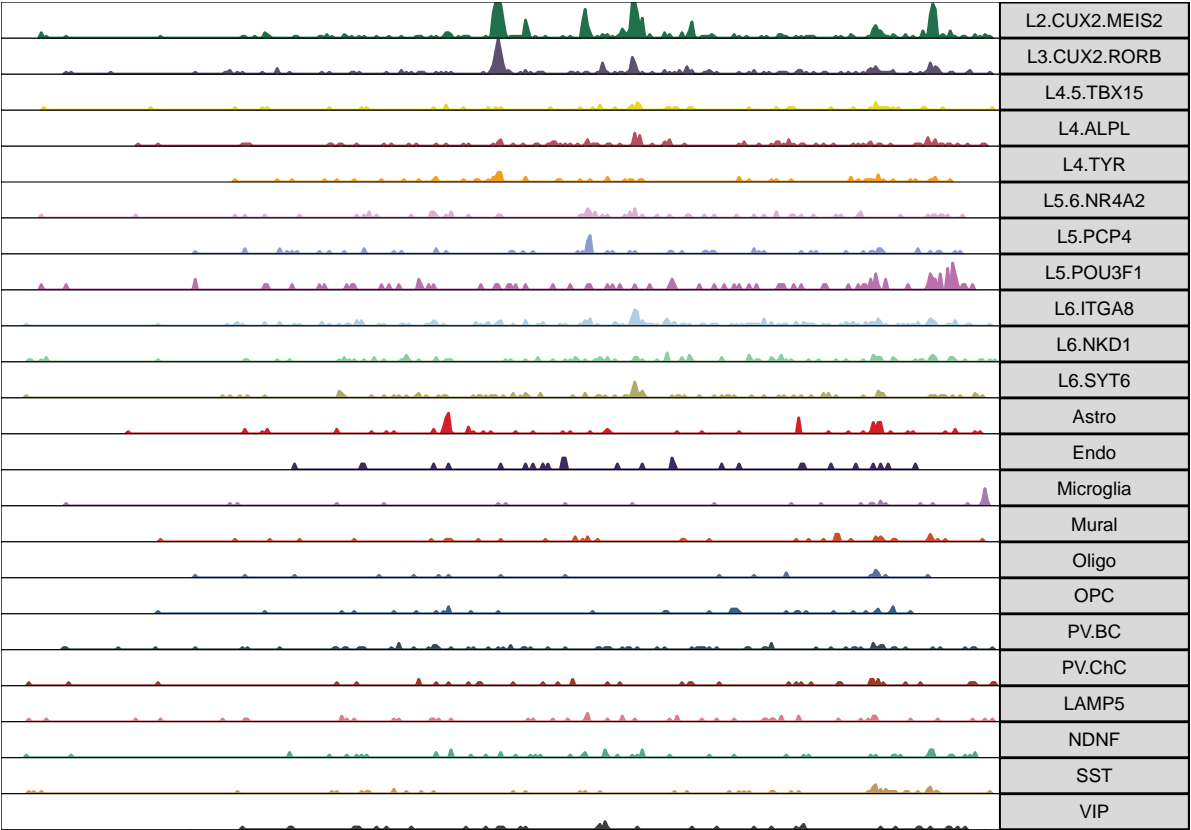
Peaks



Genes

chrX:1640959–1741460 Marker Peak Tracks for: L2.CUX2.MEIS2

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

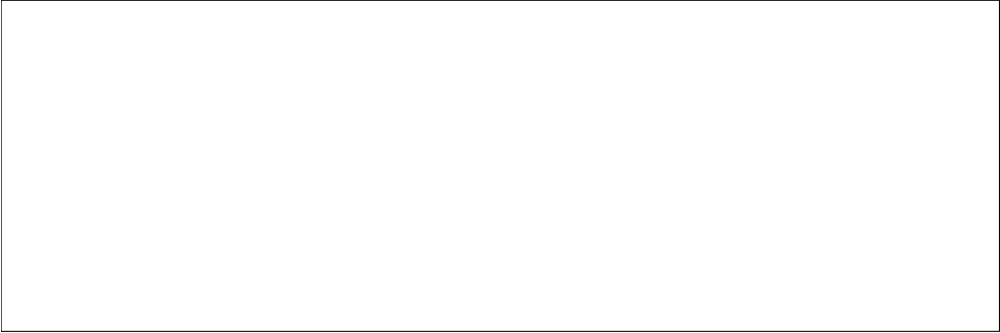


1660000 1680000 1700000 1720000 1740000

L2.CUX2.MEIS2



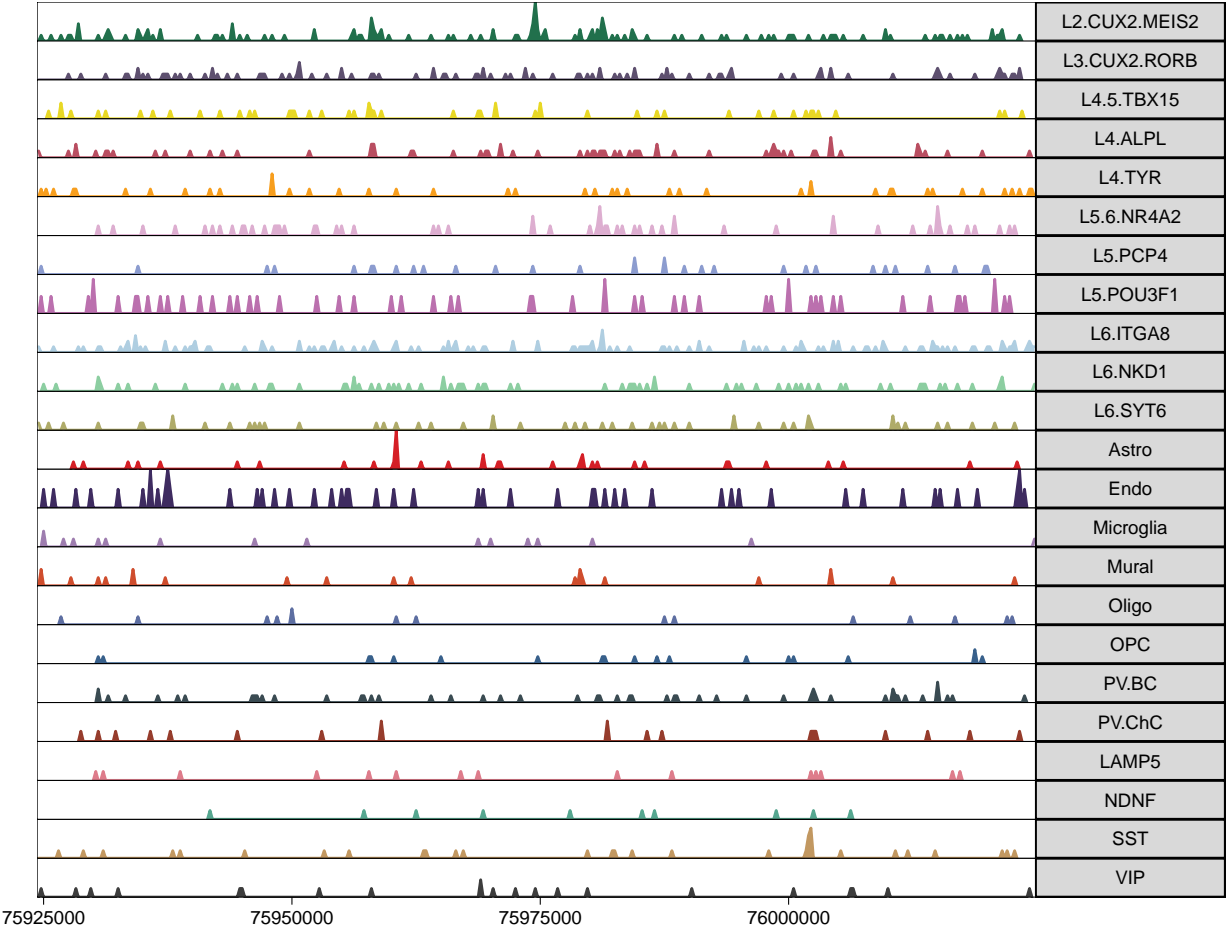
Peaks



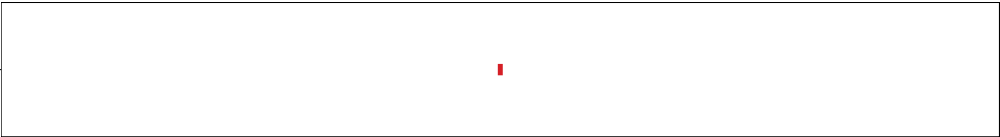
GeneTrack

strand ● ·

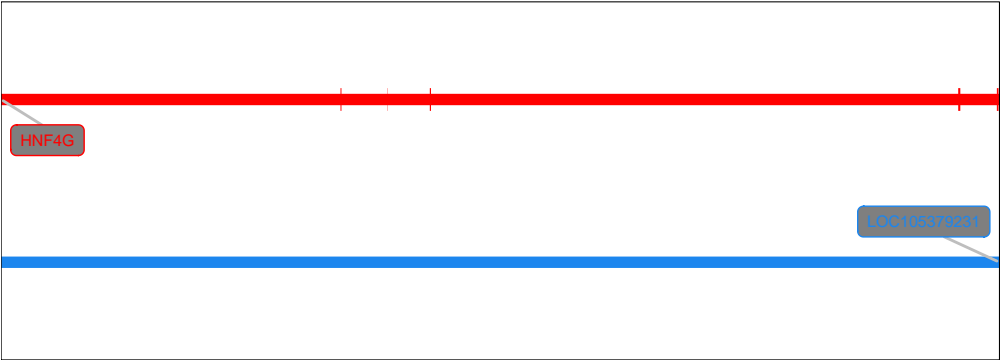
Coverage
(Norm. ATAC Signal Range (0–0.17) by ReadsInTSS)



L2.CUX2.MEIS2

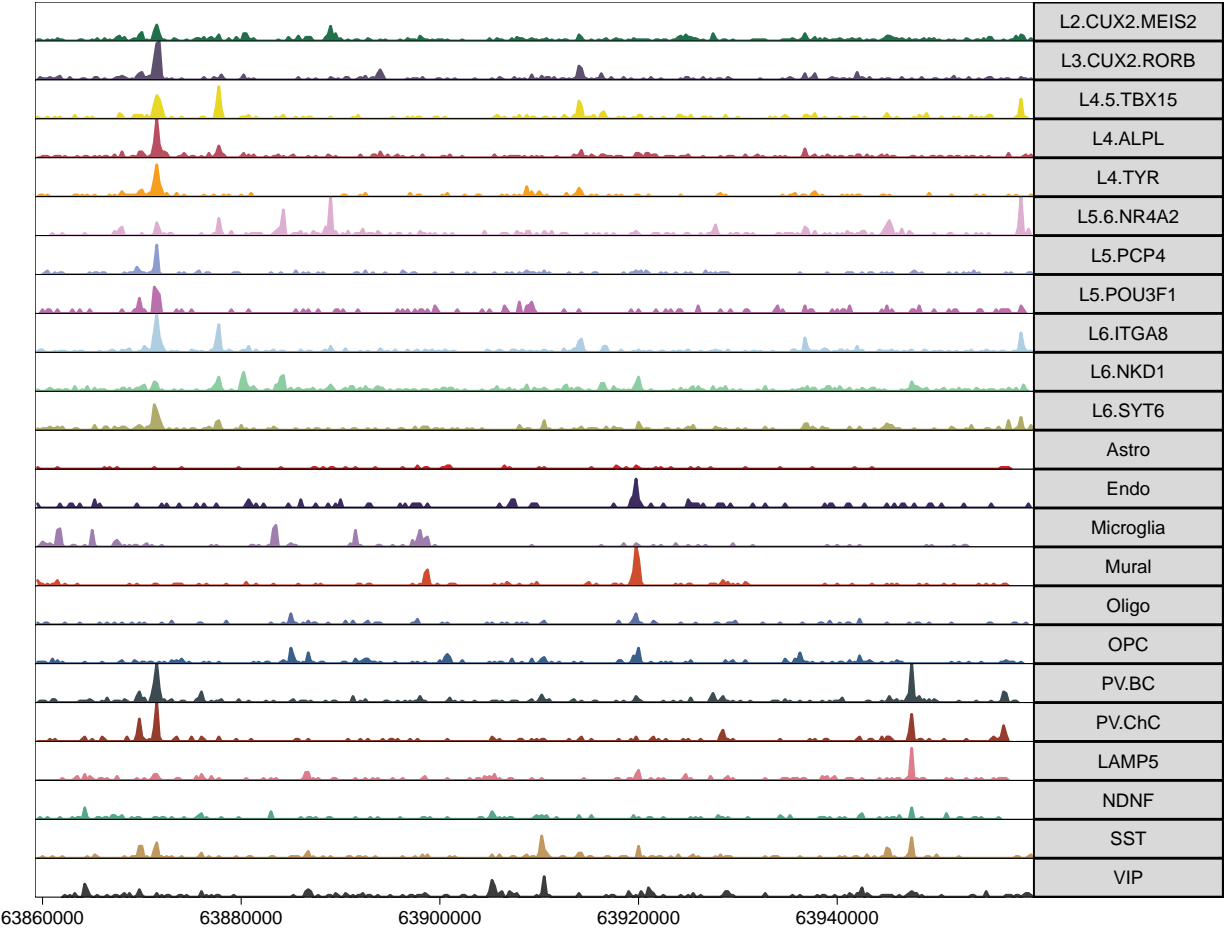


Peaks

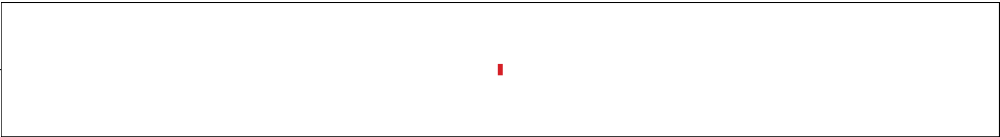


Genes

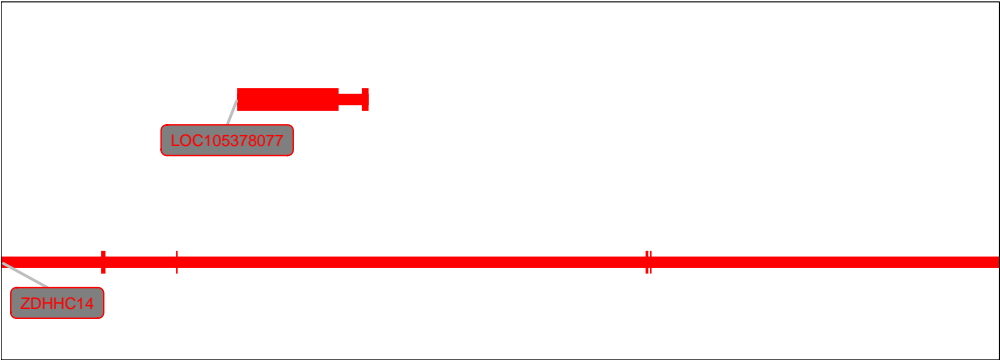
Coverage
(Norm. ATAC Signal Range (0–0.76) by ReadsInTSS)



L2.CUX2.MEIS2

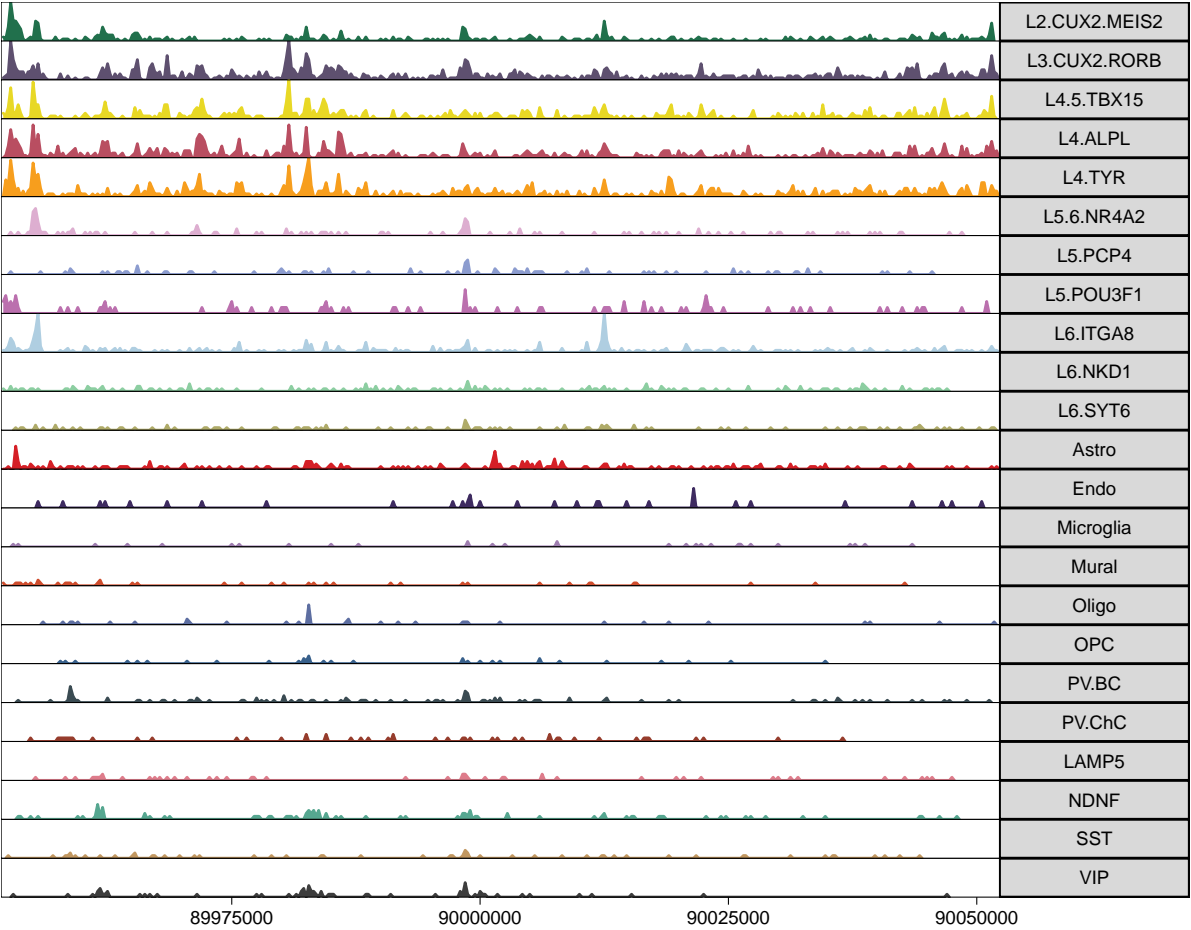


Peaks

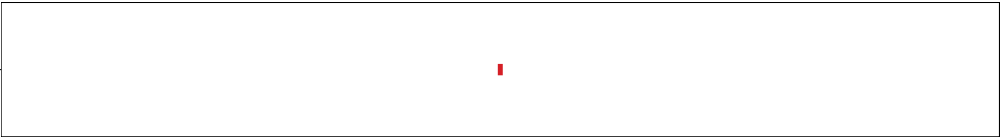


Genes

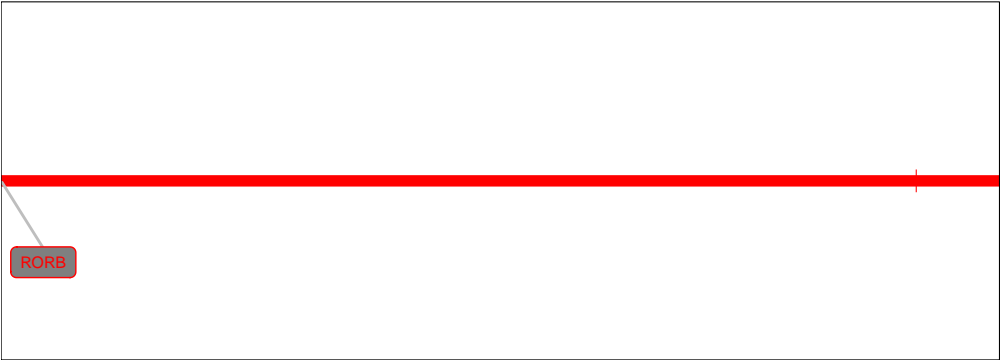
Coverage
(Norm. ATAC Signal Range (0–0.48) by ReadsInTSS)



L2.CUX2.MEIS2

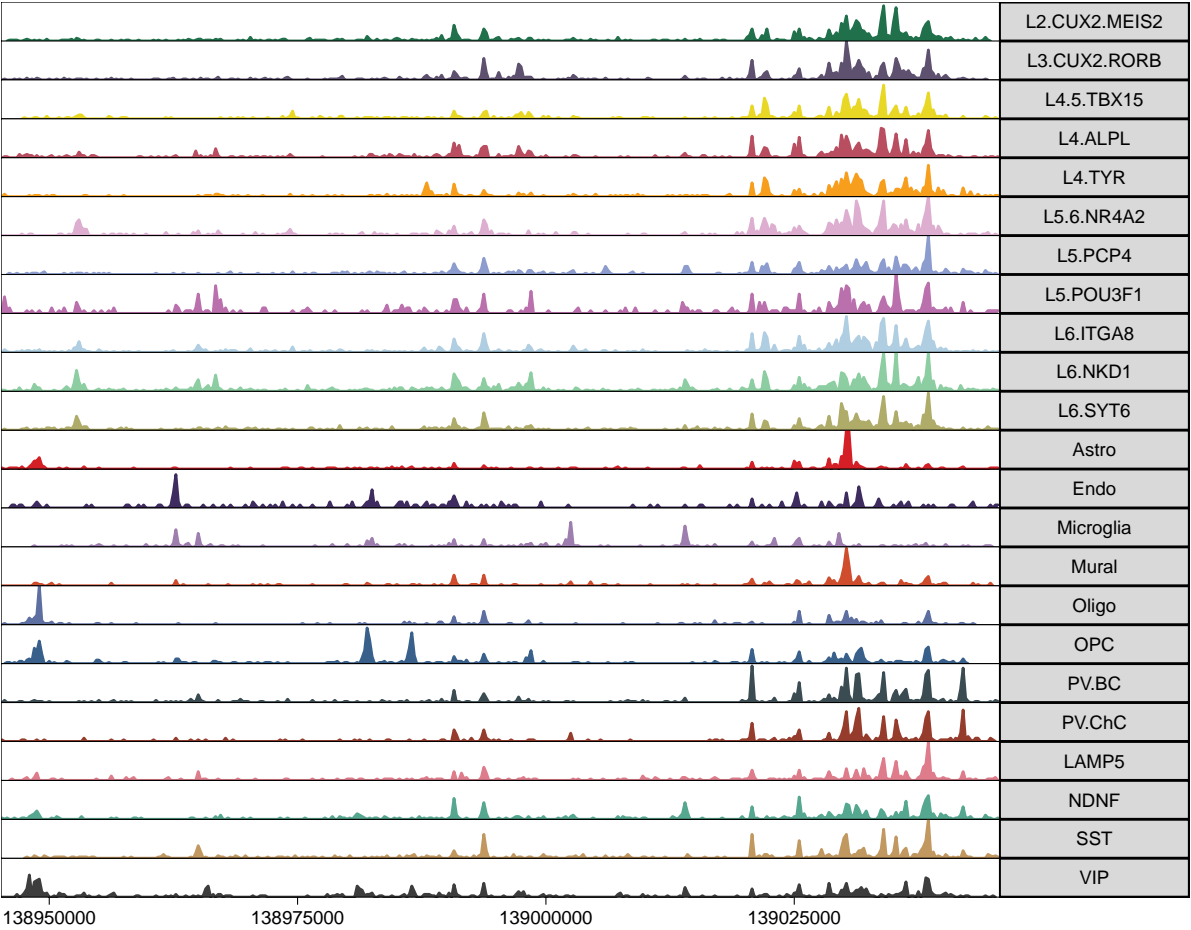


Peaks

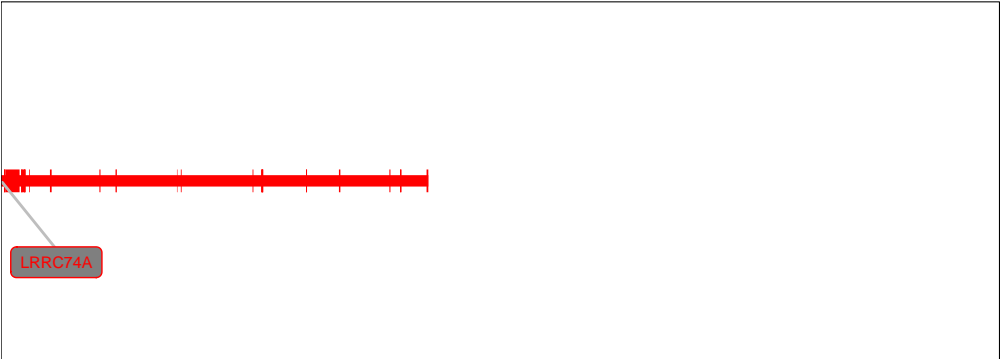
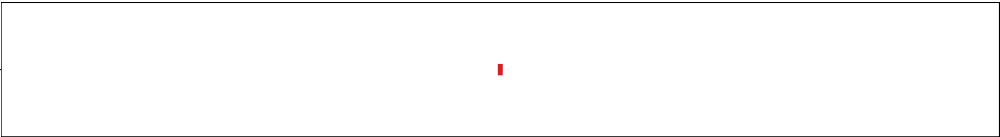


Genes

Coverage
(Norm. ATAC Signal Range (0–1.03) by ReadsInTSS)

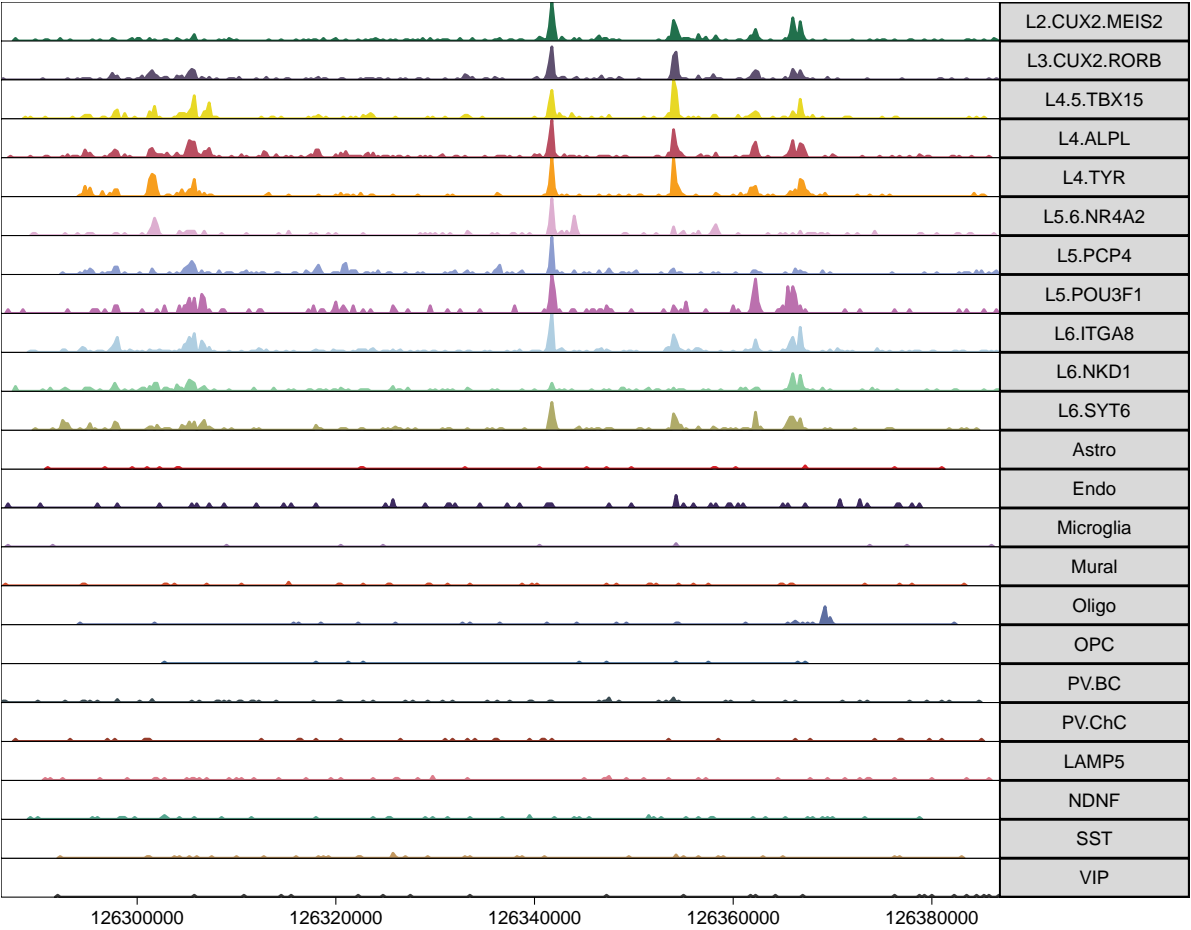


L2.CUX2.MEIS2

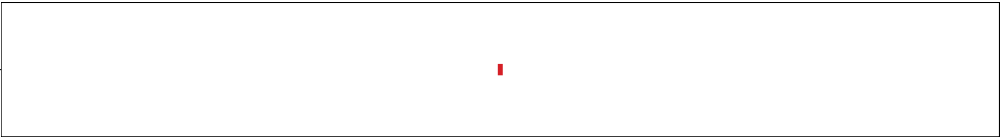


chr2:126286304–126386805 Marker Peak Tracks for: L2.CUX2.MEIS2

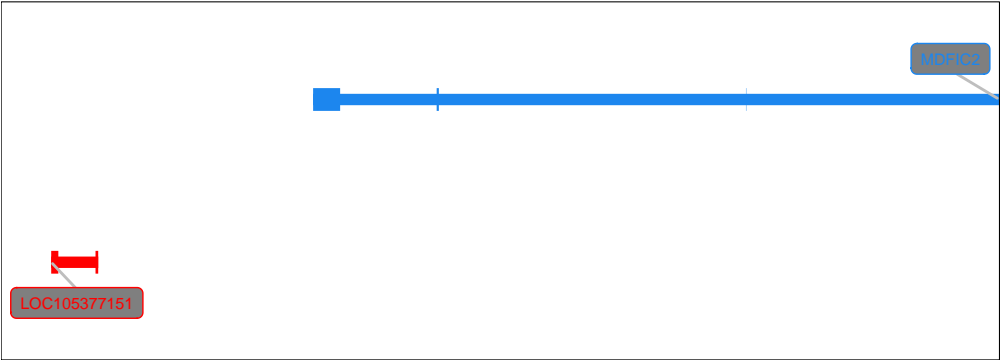
Coverage
(Norm. ATAC Signal Range (0–0.75) by ReadsInTSS)



L2.CUX2.MEIS2

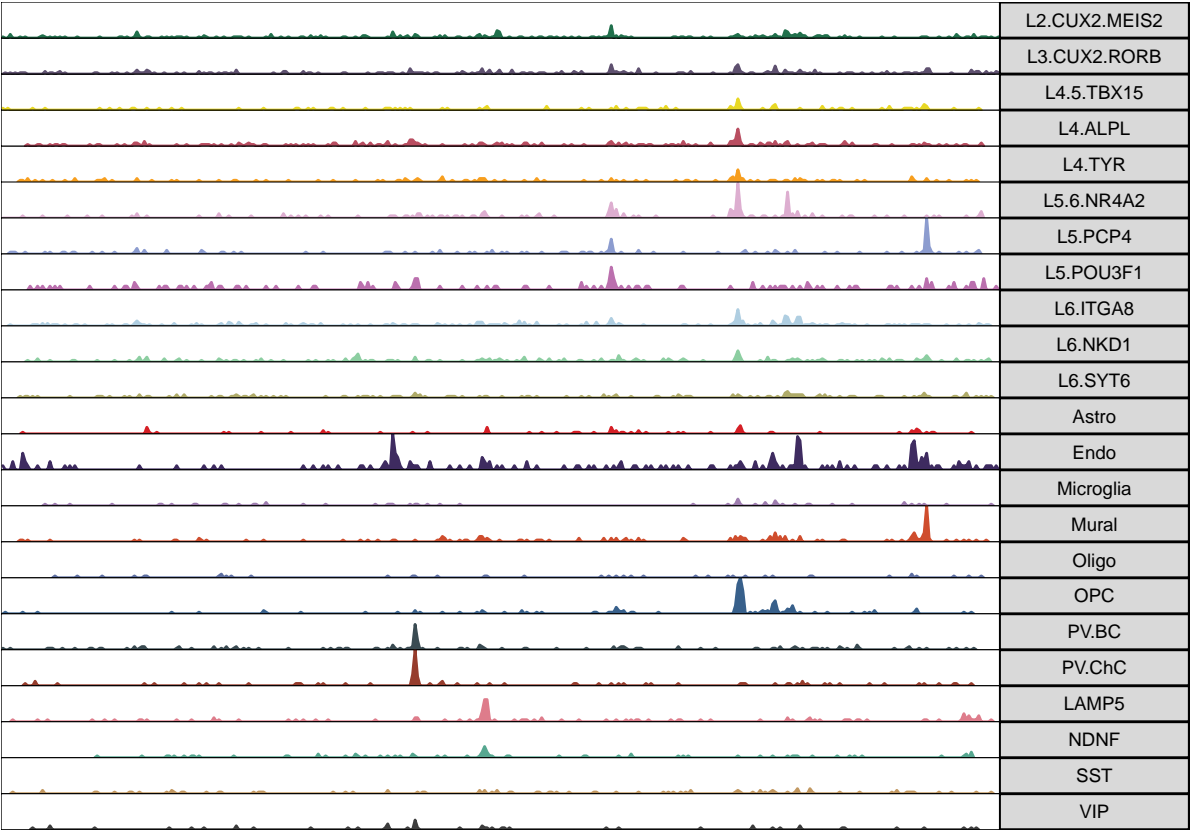


Peaks



Genes

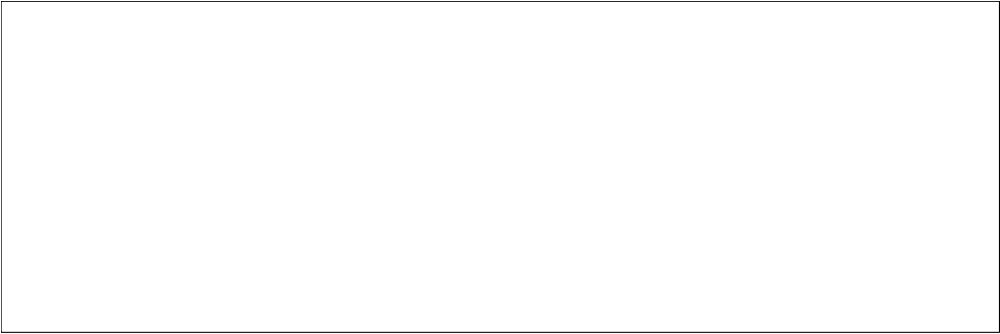
Coverage
(Norm. ATAC Signal Range (0-0.69) by ReadsInTSS)



L2.CUX2.MEIS2

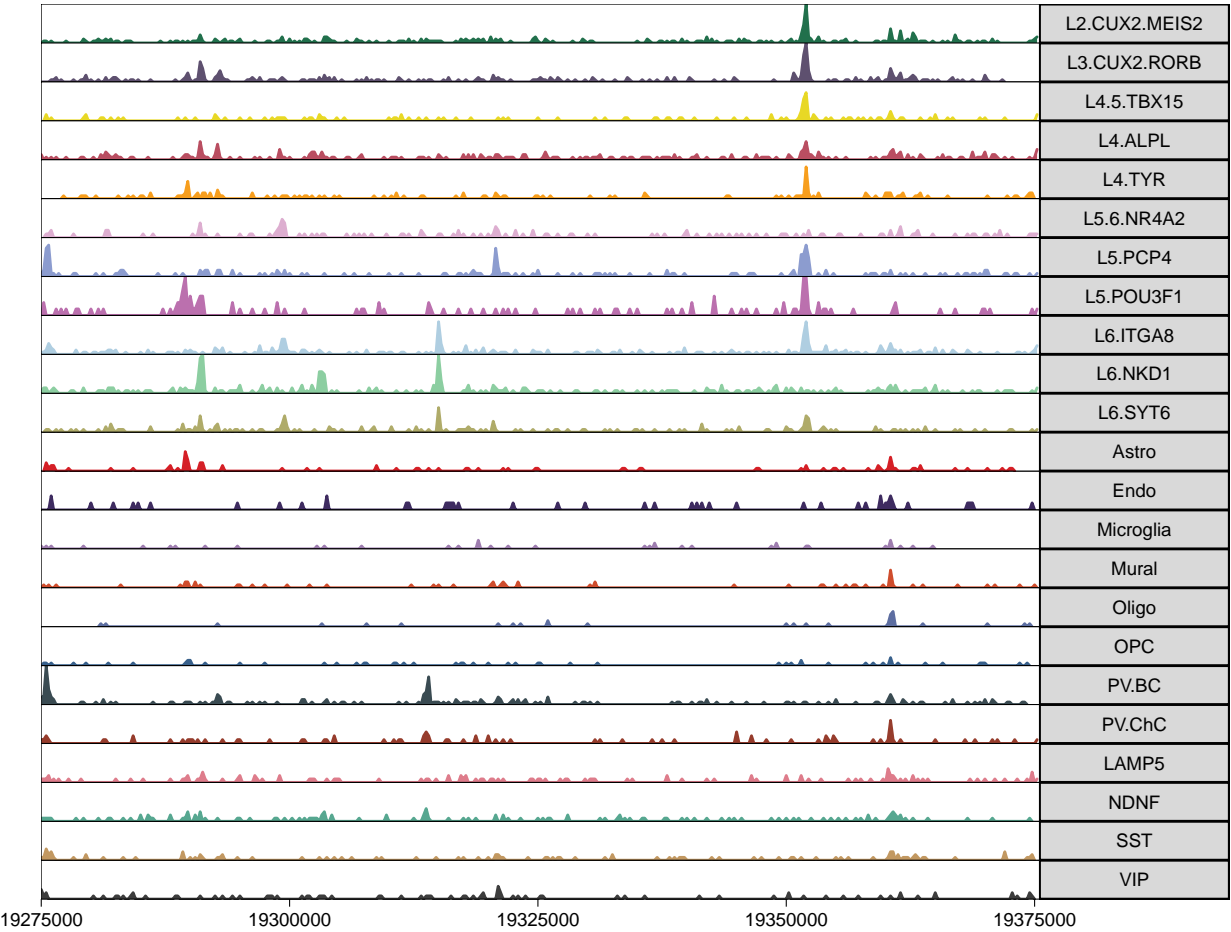


Peaks

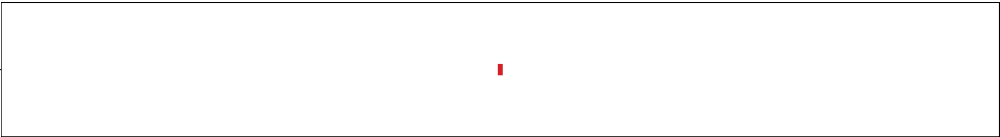


GeneTrack

Coverage
(Norm. ATAC Signal Range (0-0.44) by ReadsInTSS)



L2.CUX2.MEIS2



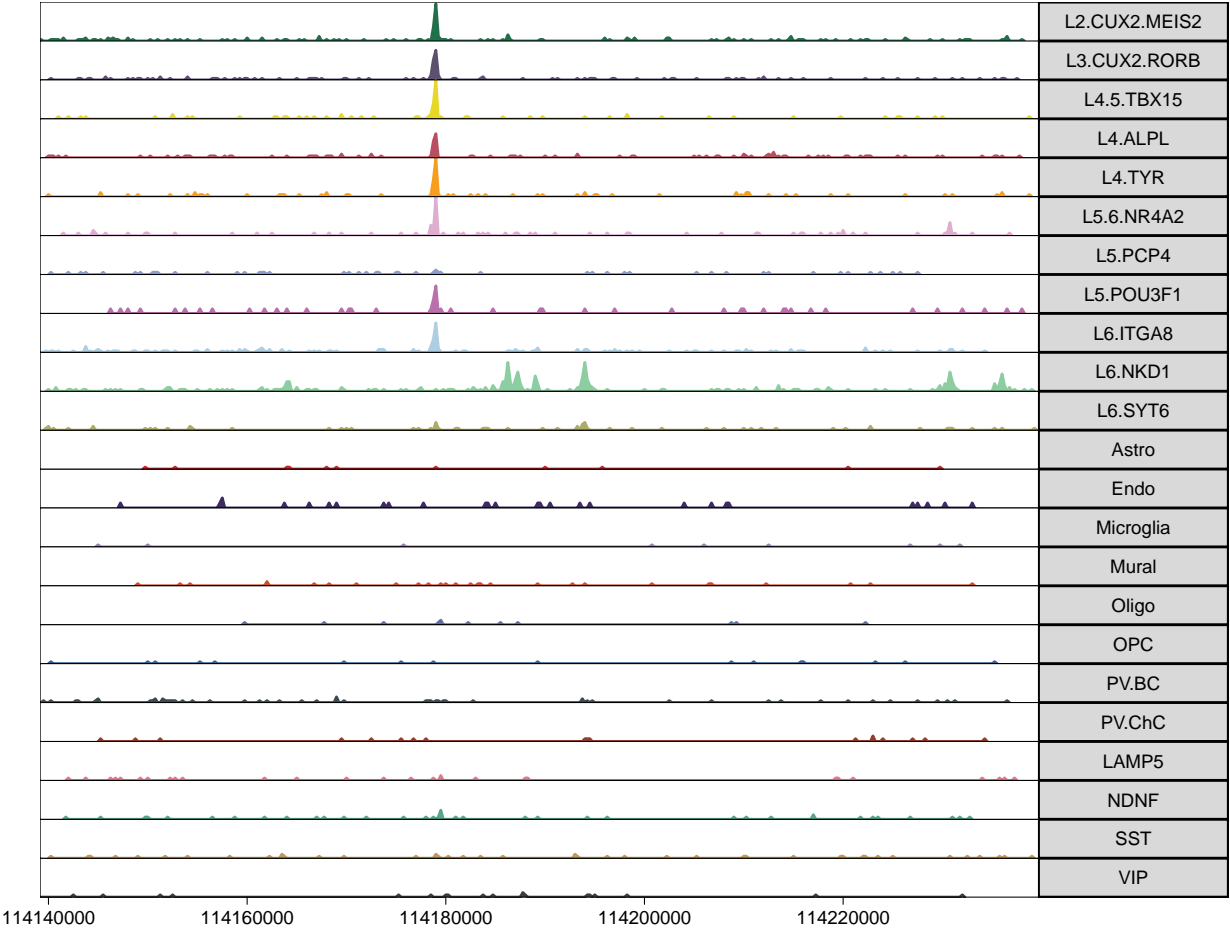
Peaks



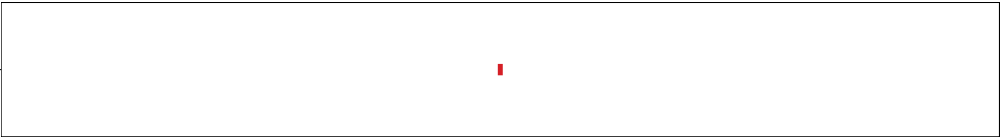
Genes

chr4:114139161–114239662 Marker Peak Tracks for: L2.CUX2.MEIS2

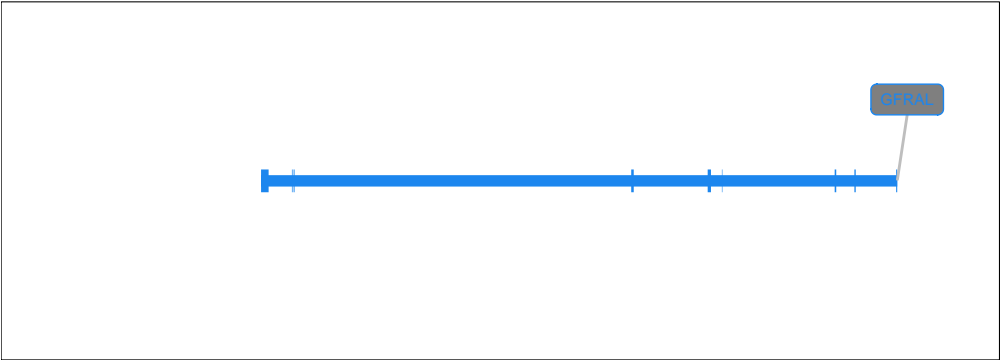
Coverage
(Norm. ATAC Signal Range (0–0.62) by ReadsInTSS)



L2.CUX2.MEIS2

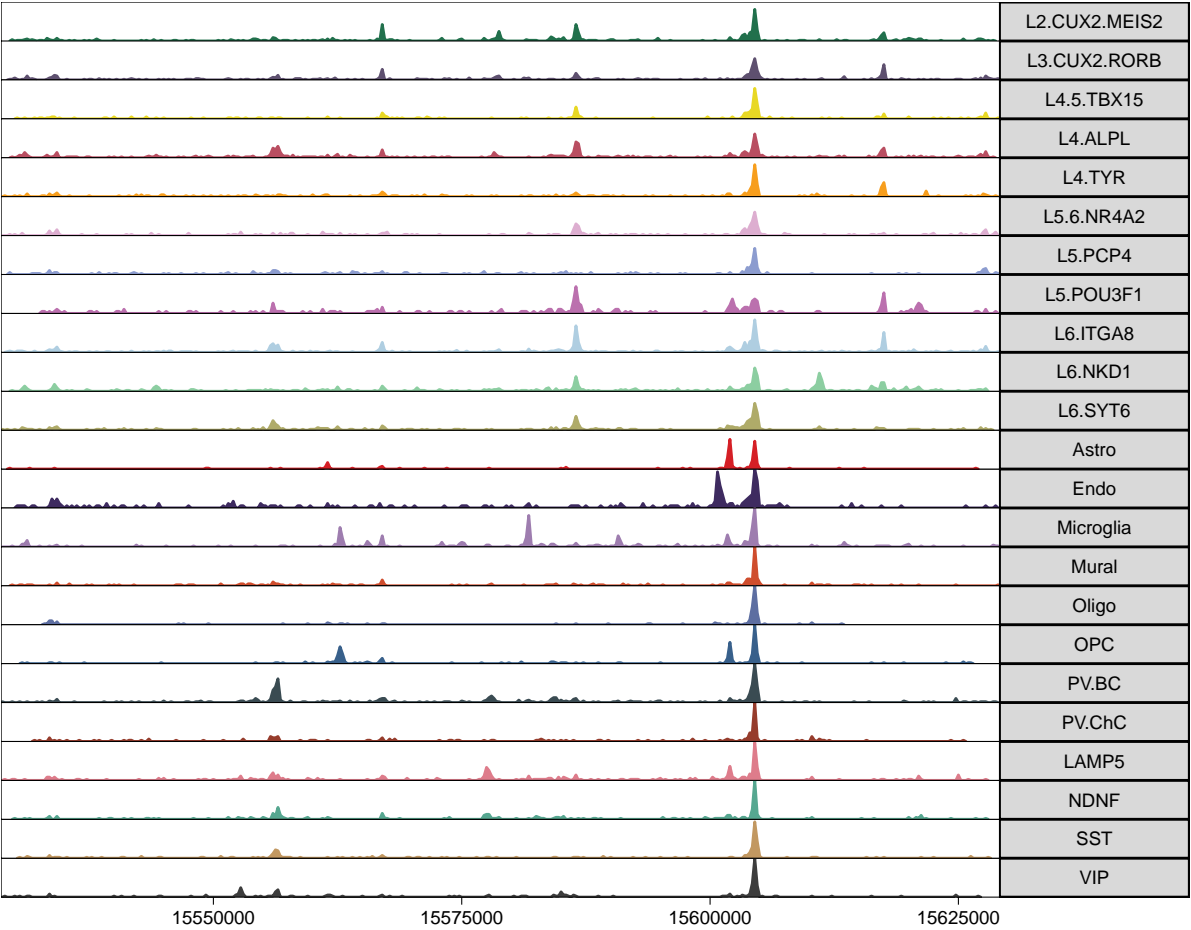


Peaks

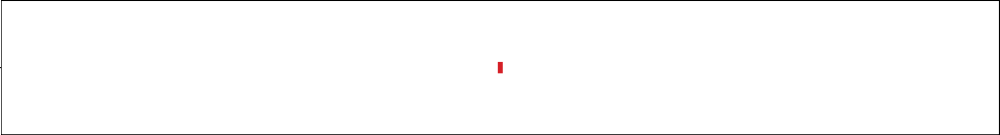


Genes

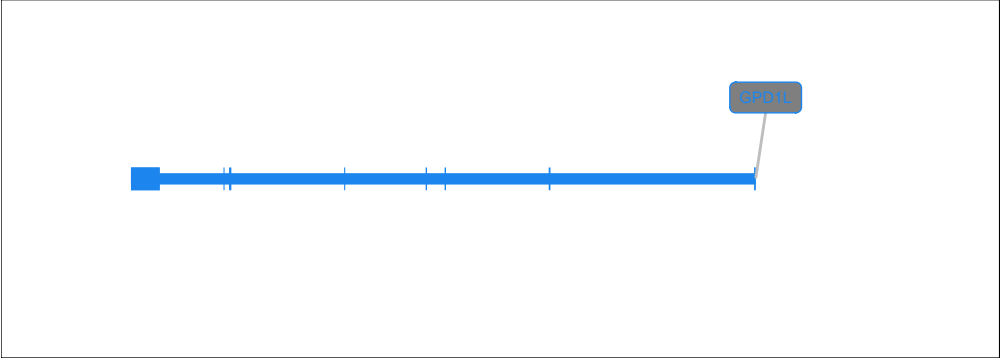
Coverage
(Norm. ATAC Signal Range (0–1.39) by ReadsInTSS)



L2.CUX2.MEIS2

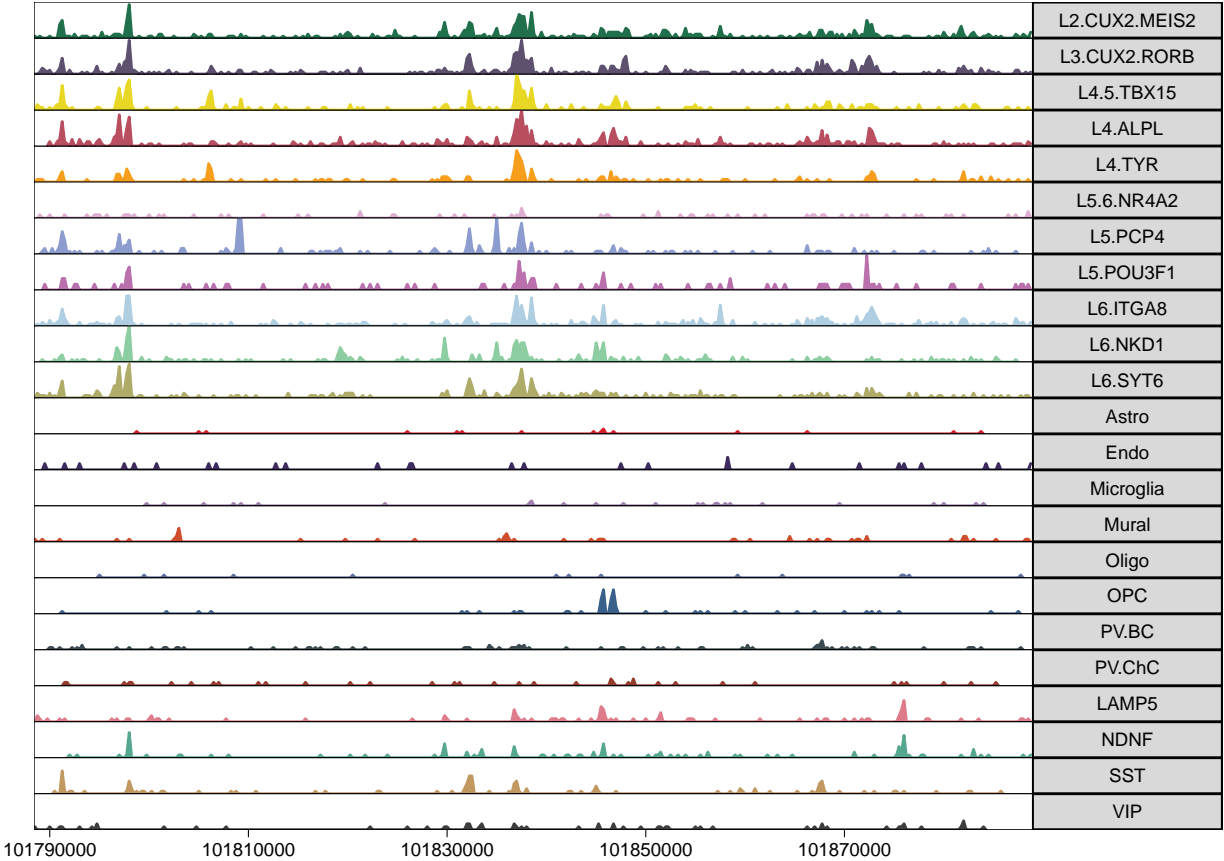


Peaks



Genes

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



L2.CUX2.MEIS2

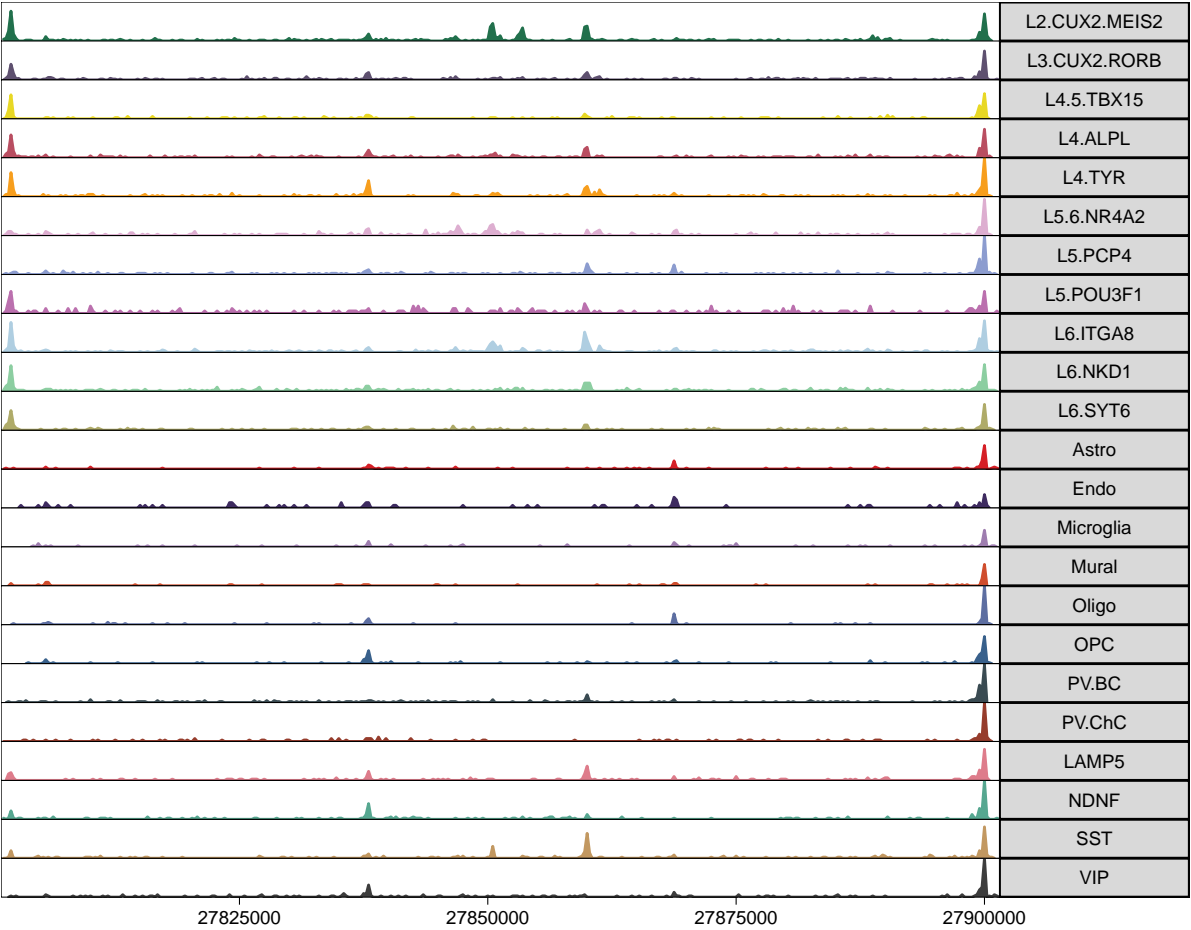


Peaks

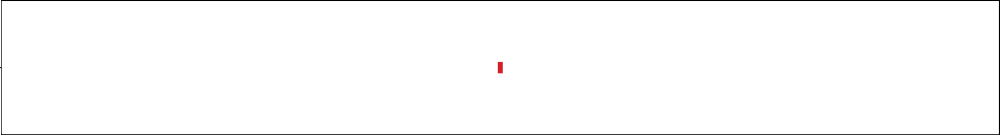
GeneTrack

chr10:27801007–27901508 Marker Peak Tracks for: L2.CUX2.MEIS2

Coverage
(Norm. ATAC Signal Range (0–1.17) by ReadsInTSS)



L2.CUX2.MEIS2

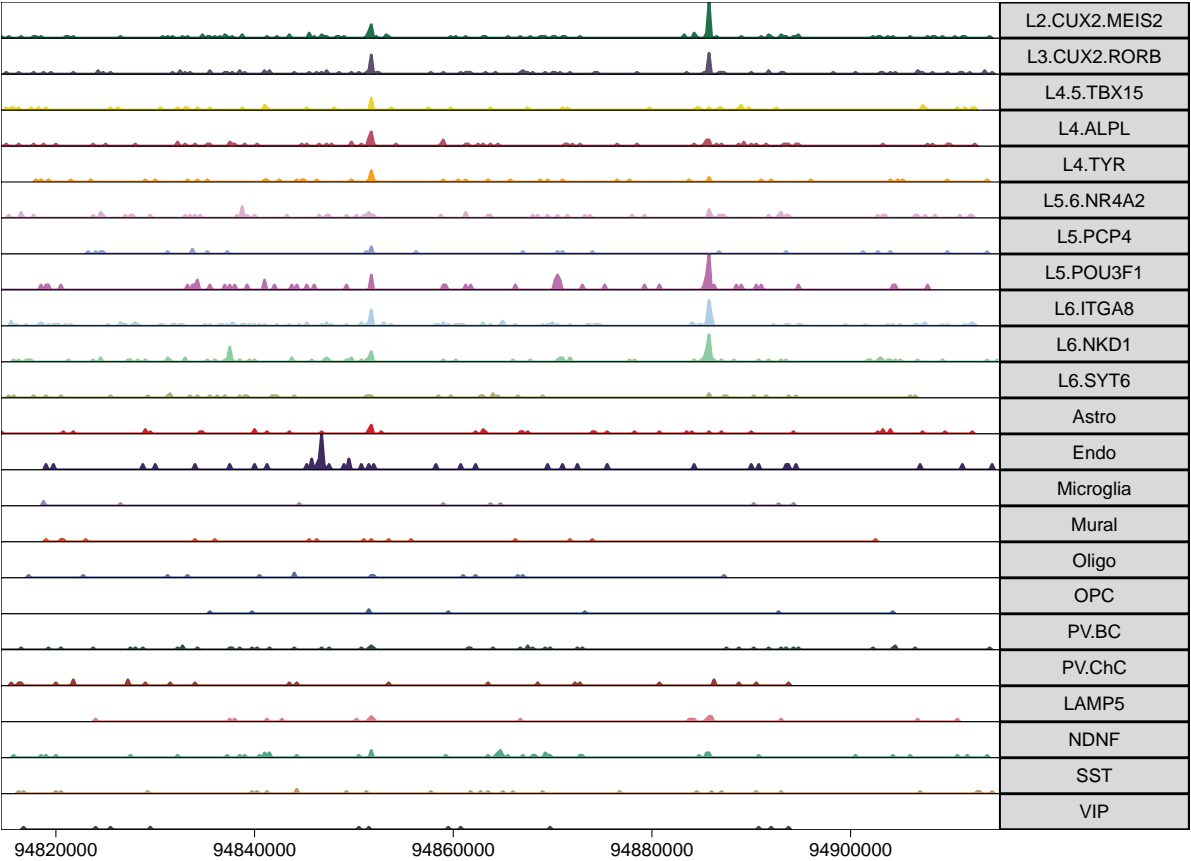


Peaks

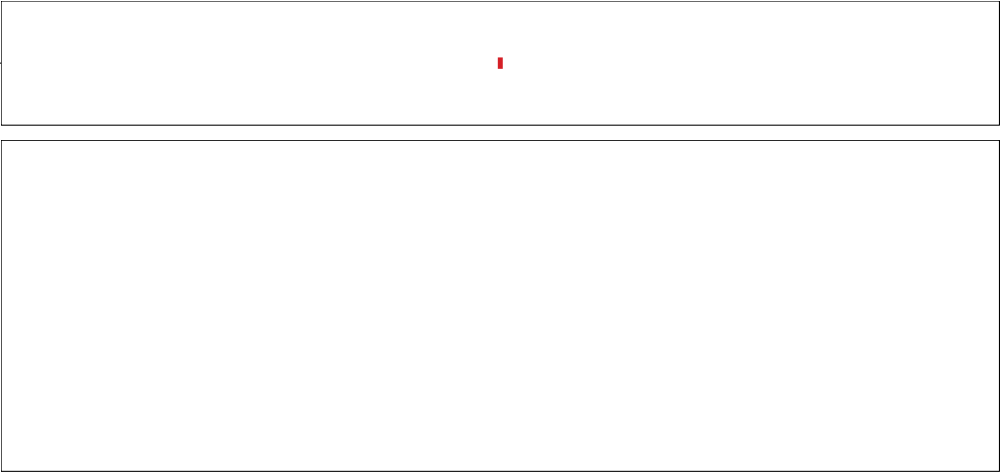


Genes

Coverage
(Norm. ATAC Signal Range (0-0.53) by ReadsInTSS)



L2.CUX2.MEIS2

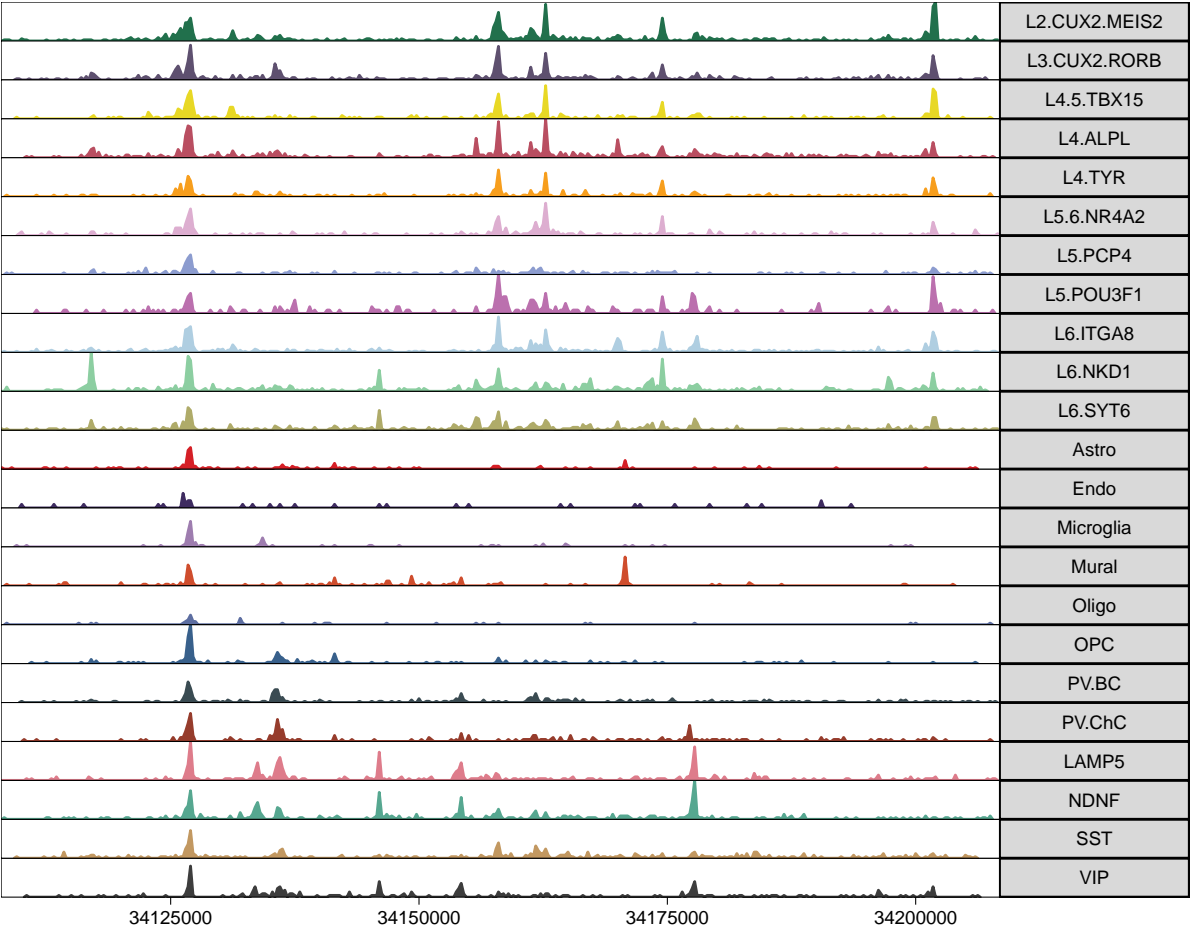


Peaks

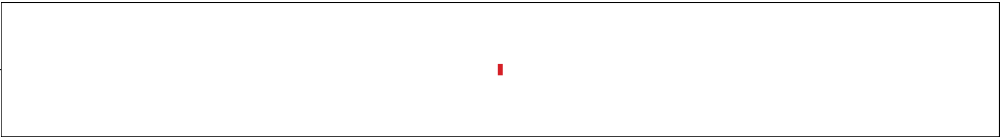
GeneTrack

chr20:34107932–34208433 Marker Peak Tracks for: L2.CUX2.MEIS2

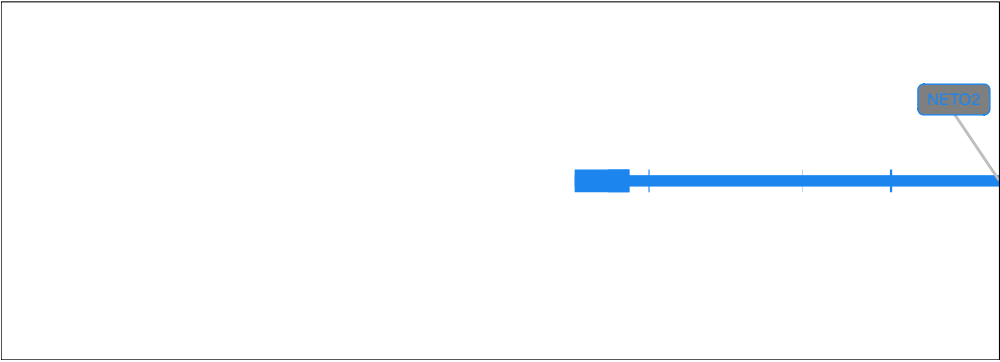
Coverage
(Norm. ATAC Signal Range (0–0.85) by ReadsInTSS)



L2.CUX2.MEIS2



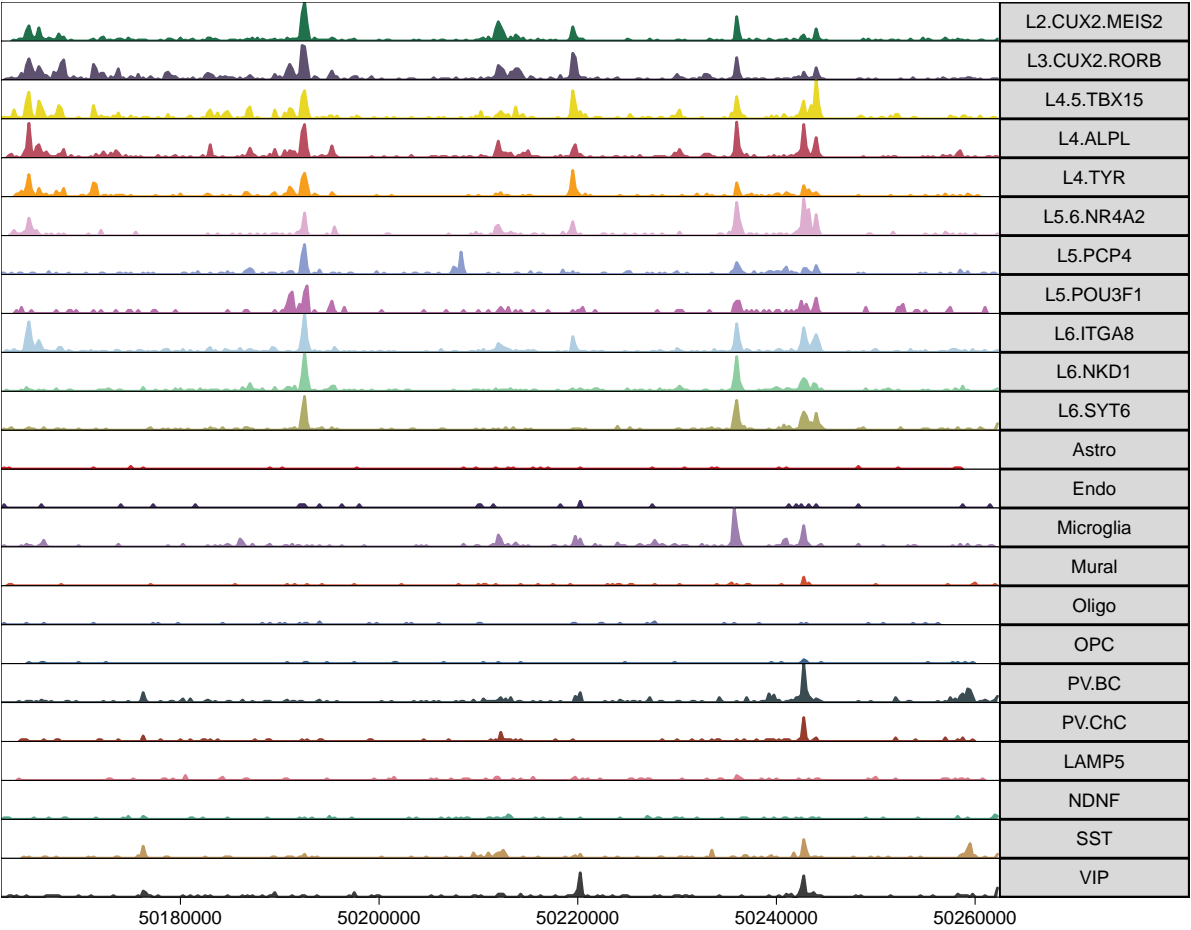
Peaks



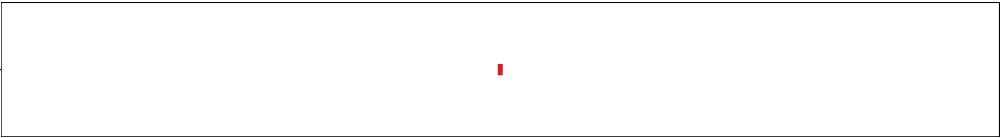
Genes

chr10:50161950–50262451 Marker Peak Tracks for: L2.CUX2.MEIS2

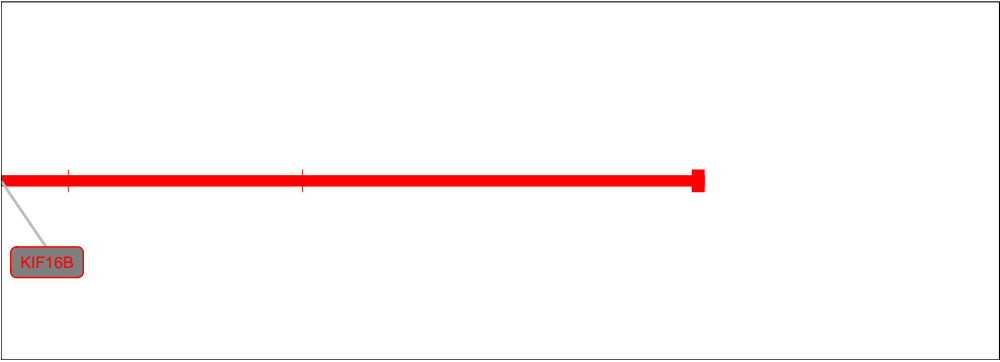
Coverage
(Norm. ATAC Signal Range (0–0.93) by ReadsInTSS)



L2.CUX2.MEIS2



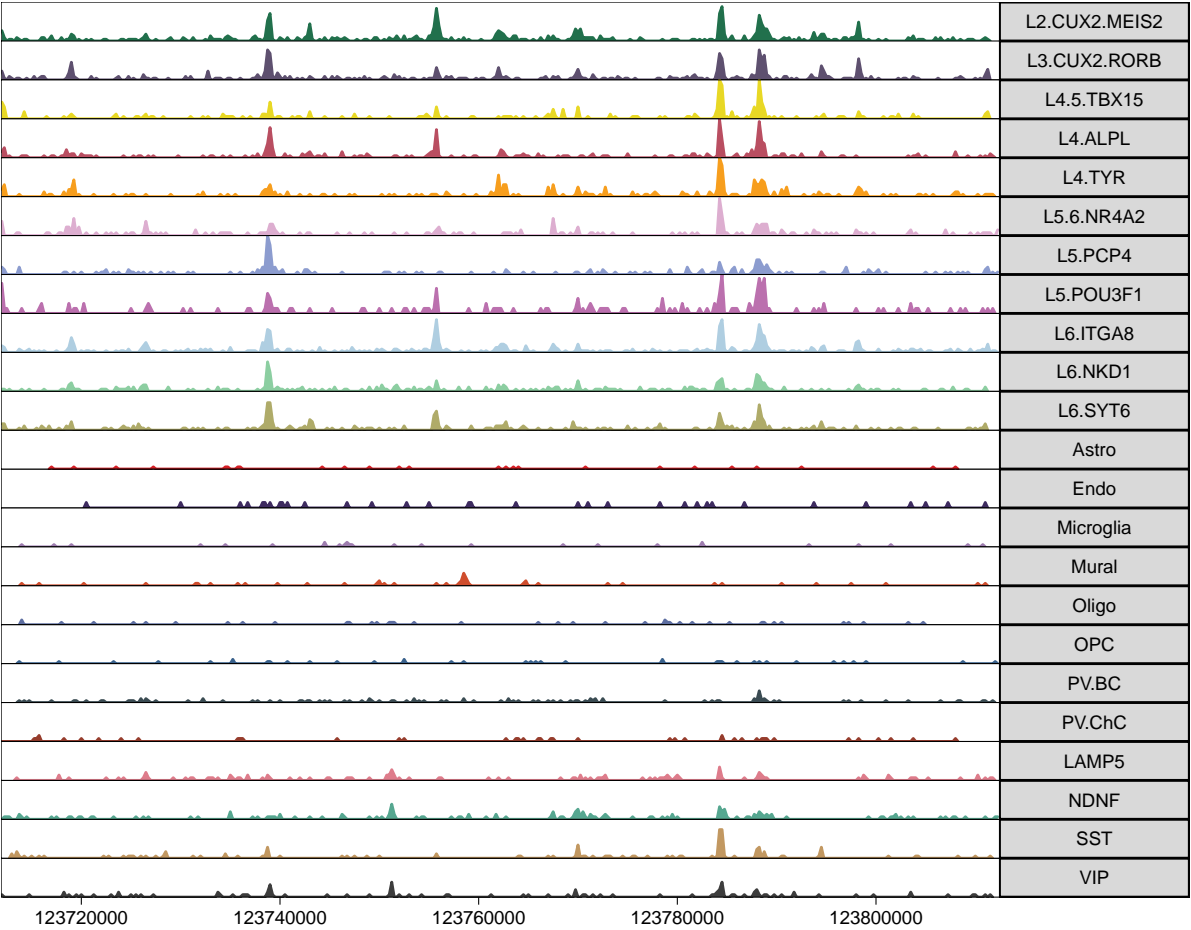
Peaks



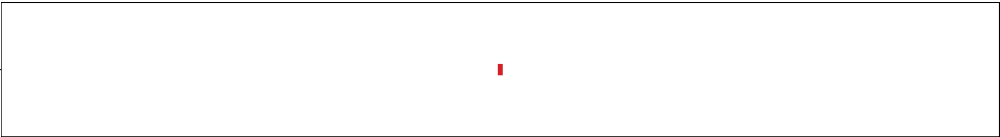
Genes

chr3:123711926–123812427 Marker Peak Tracks for: L2.CUX2.MEIS2

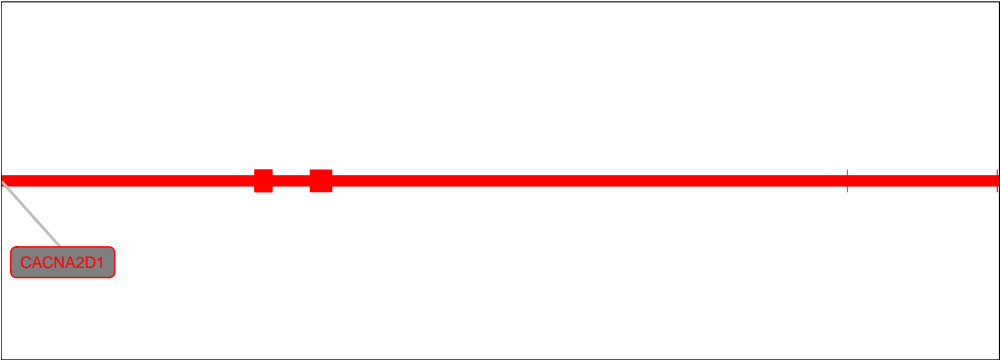
Coverage
(Norm. ATAC Signal Range (0–0.57) by ReadsInTSS)



L2.CUX2.MEIS2



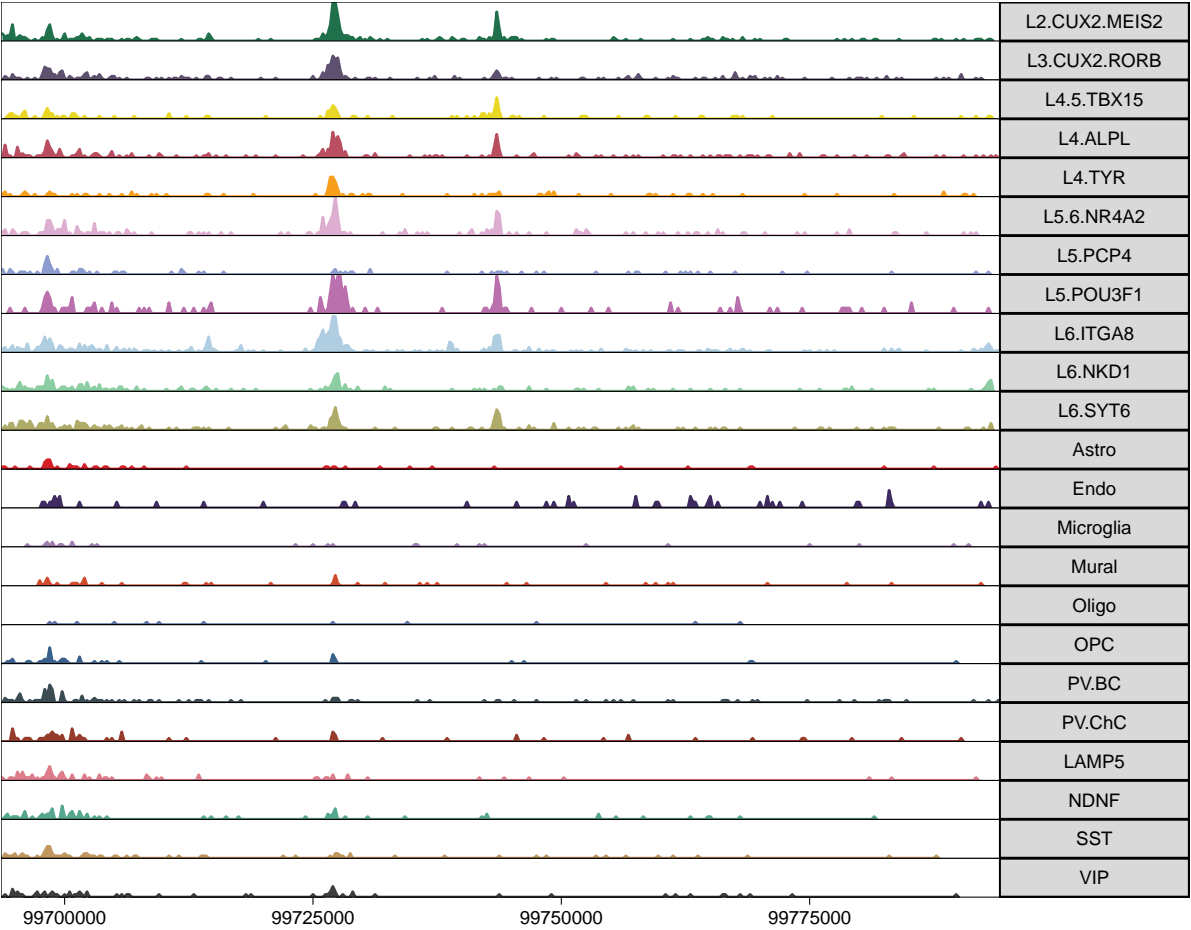
Peaks



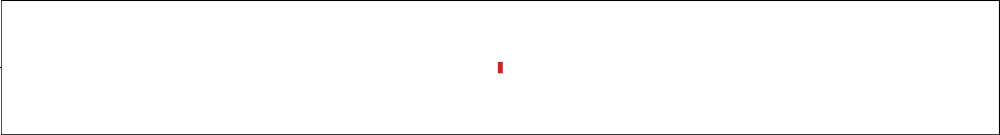
Genes

chr7:99693605-99794106 Marker Peak Tracks for: L2.CUX2.MEIS2

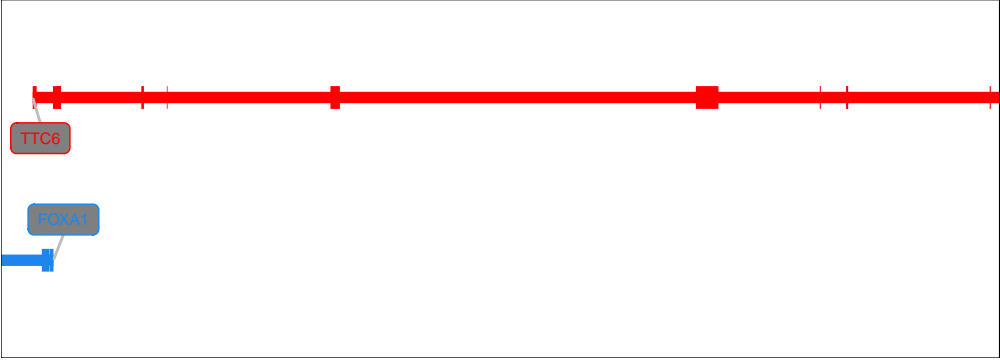
Coverage
(Norm. ATAC Signal Range (0-0.53) by ReadsInTSS)



L2.CUX2.MEIS2

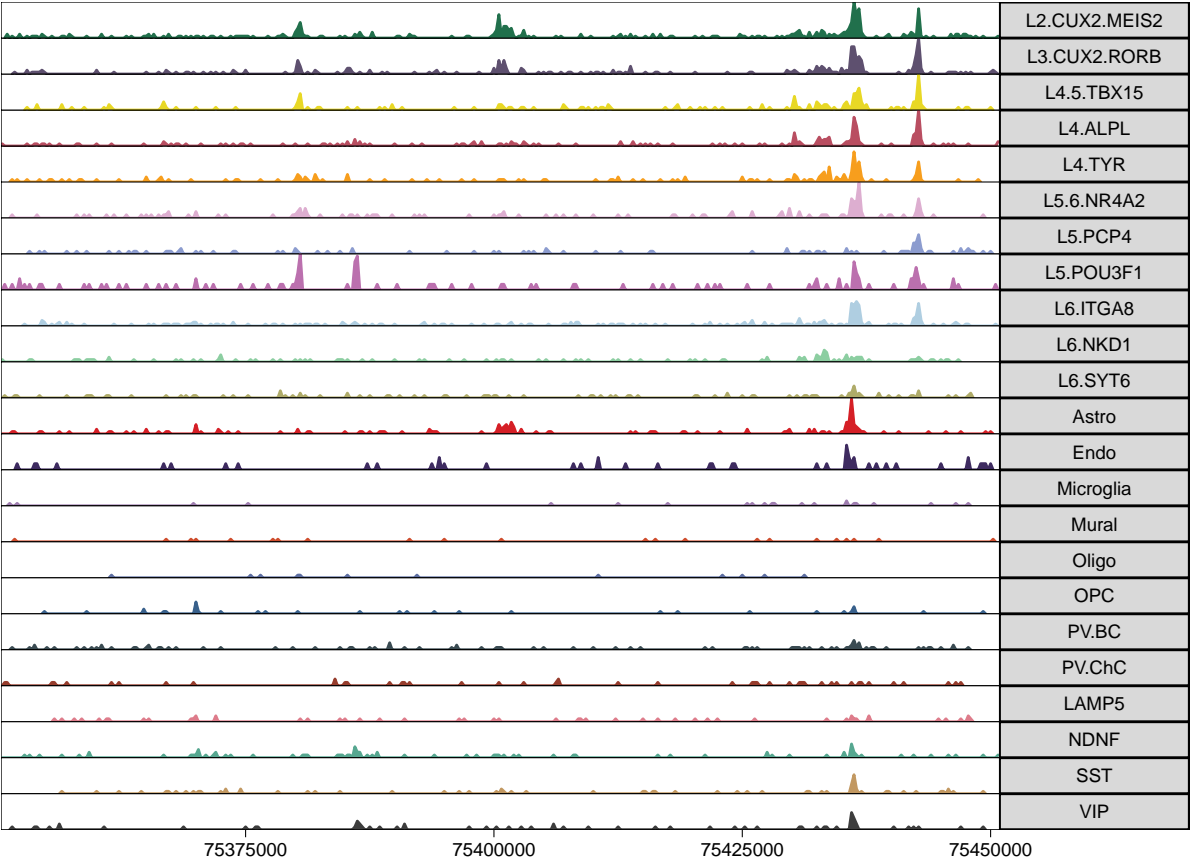


Peaks



Genes

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



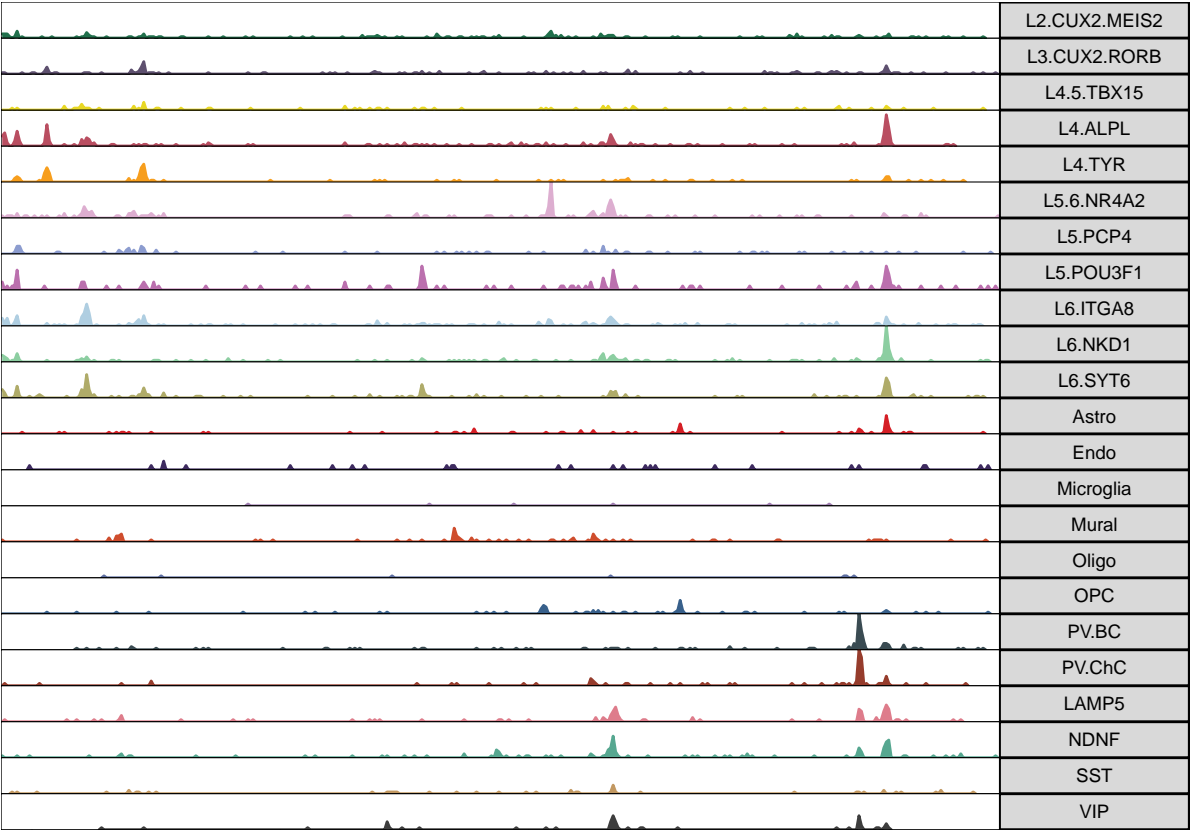
L2.CUX2.MEIS2



Peaks

GeneTrack

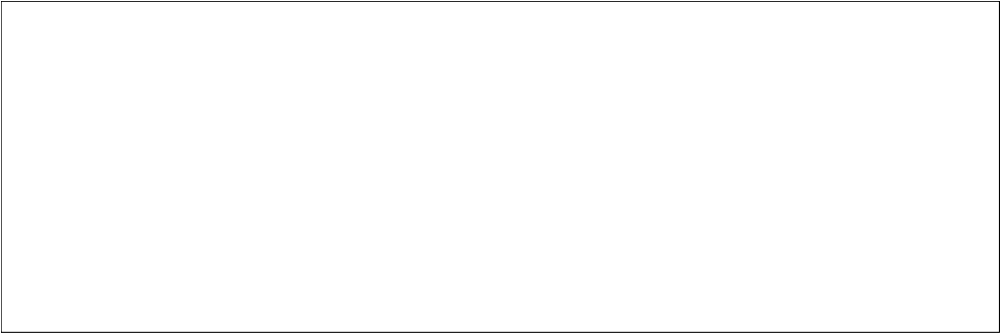
Coverage
(Norm. ATAC Signal Range (0-0.66) by ReadsInTSS)



L2.CUX2.MEIS2

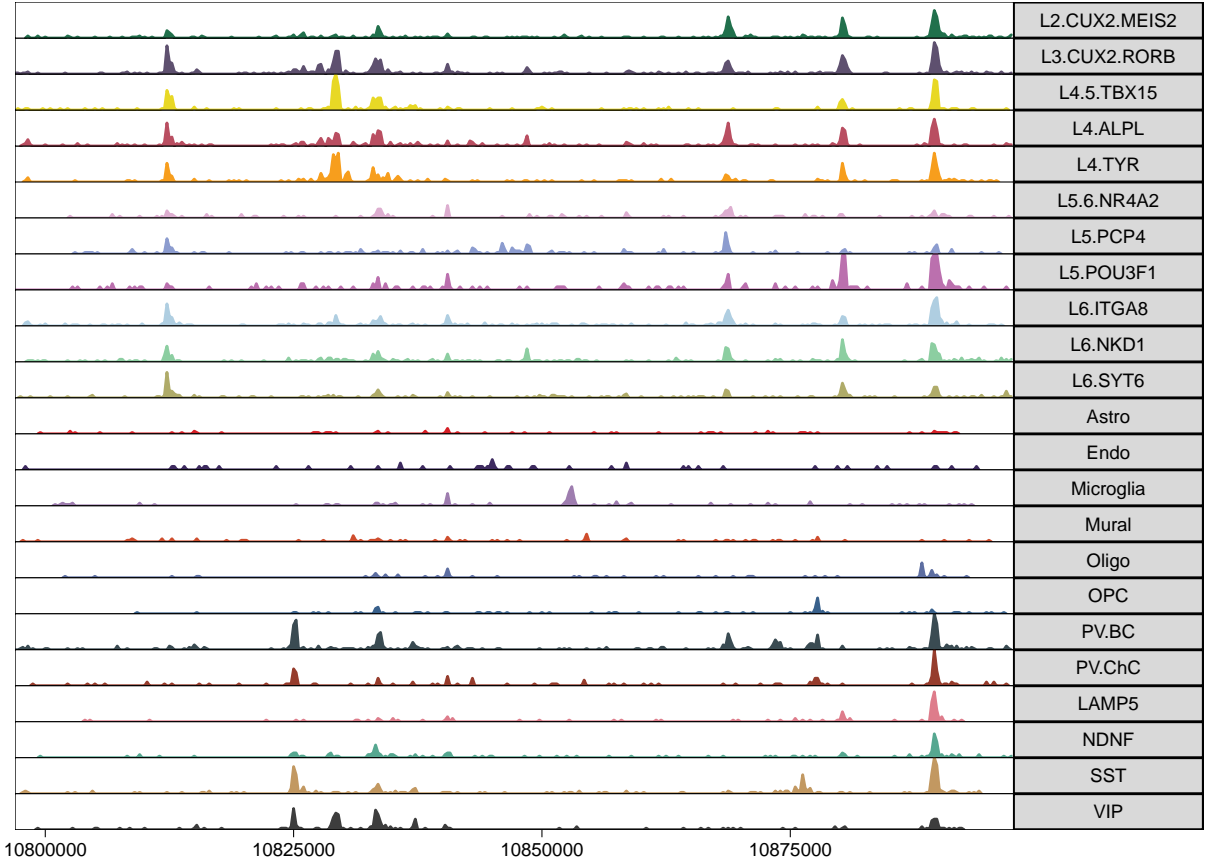


Peaks



GeneTrack

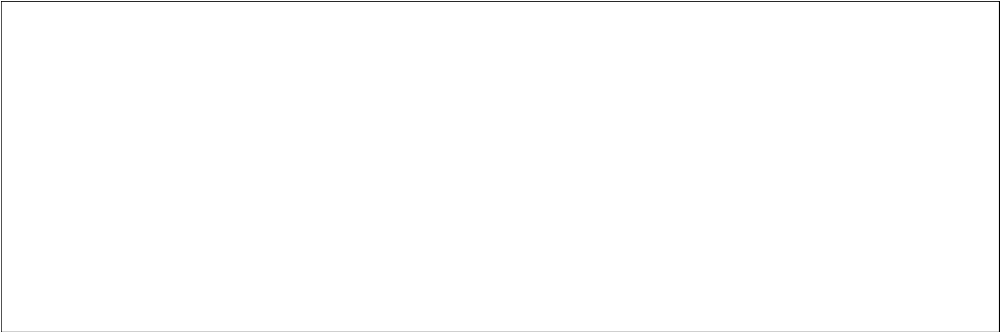
Coverage
(Norm. ATAC Signal Range (0–0.84) by ReadsInTSS)



L2.CUX2.MEIS2

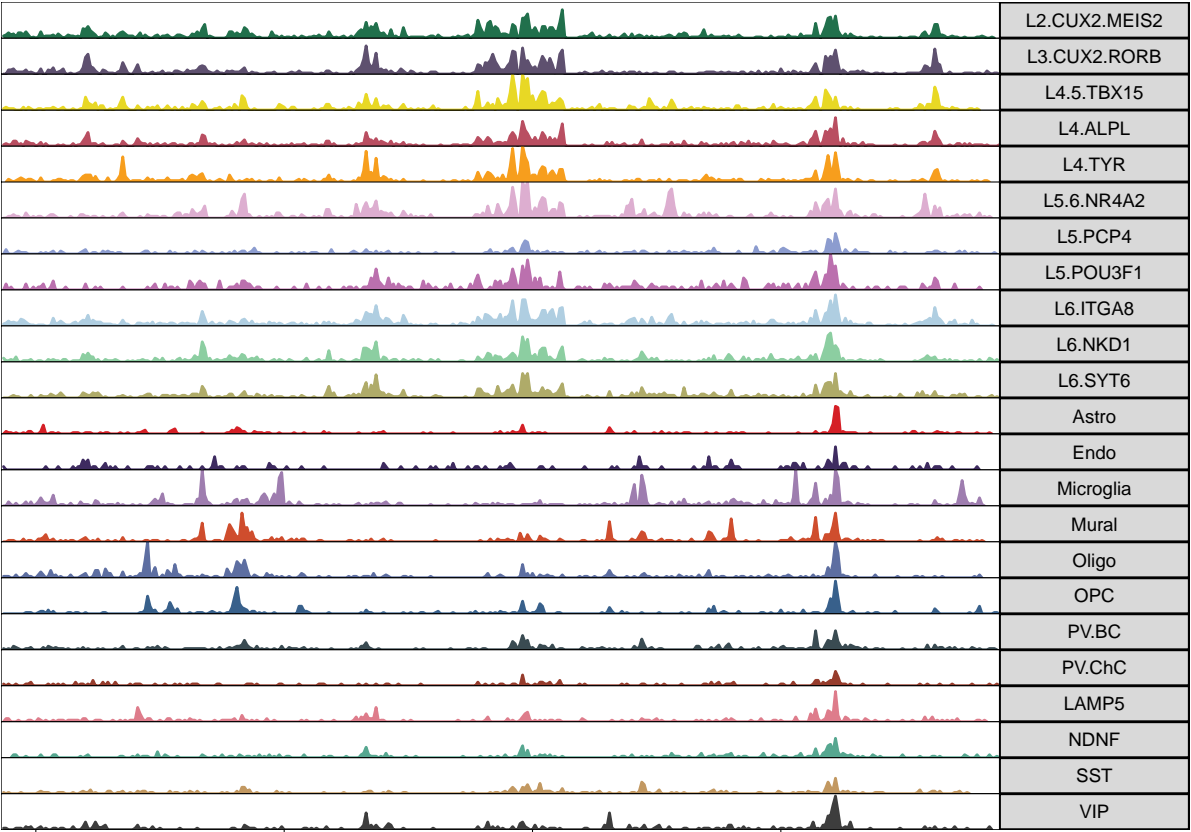


Peaks

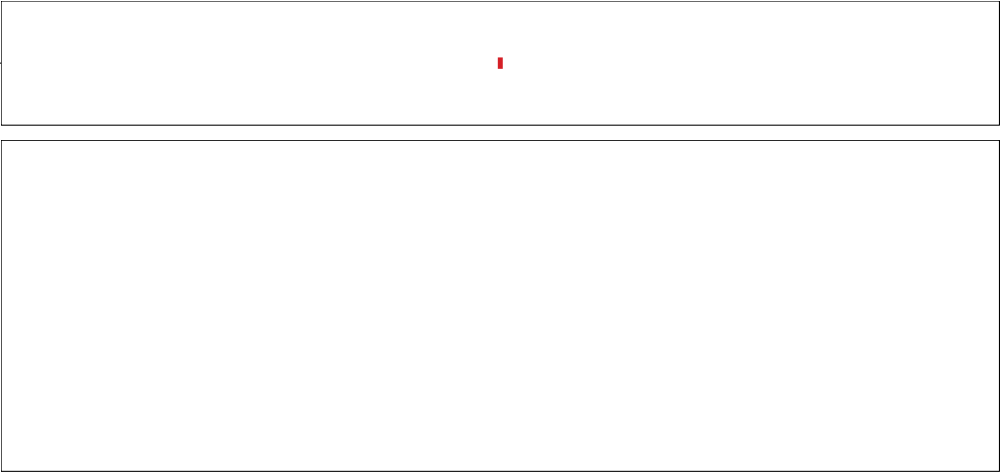


GeneTrack

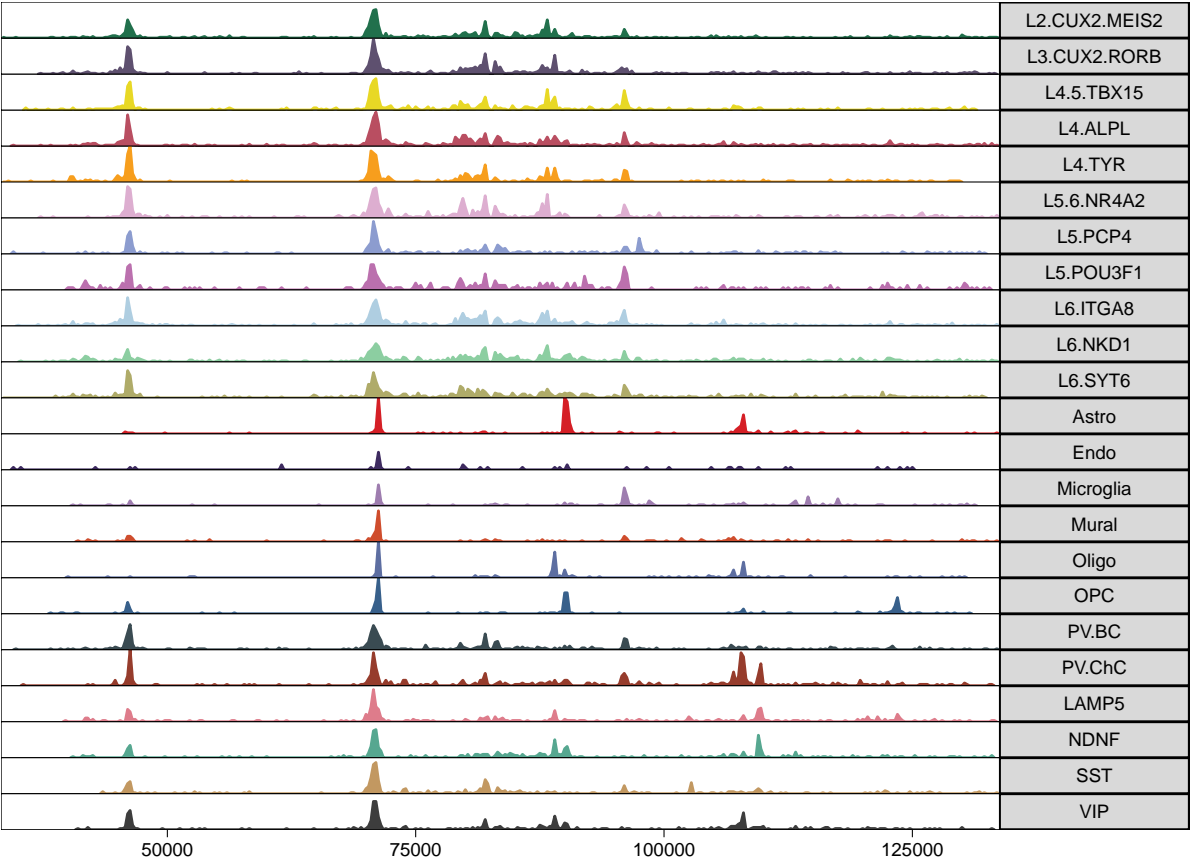
Coverage
(Norm. ATAC Signal Range (0–0.88) by ReadsInTSS)



L2.CUX2.MEIS2



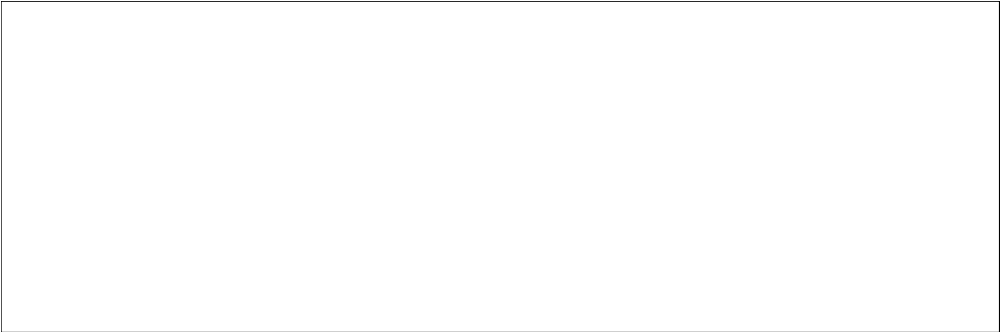
Coverage
(Norm. ATAC Signal Range (0–1.15) by ReadsInTSS)



L2.CUX2.MEIS2



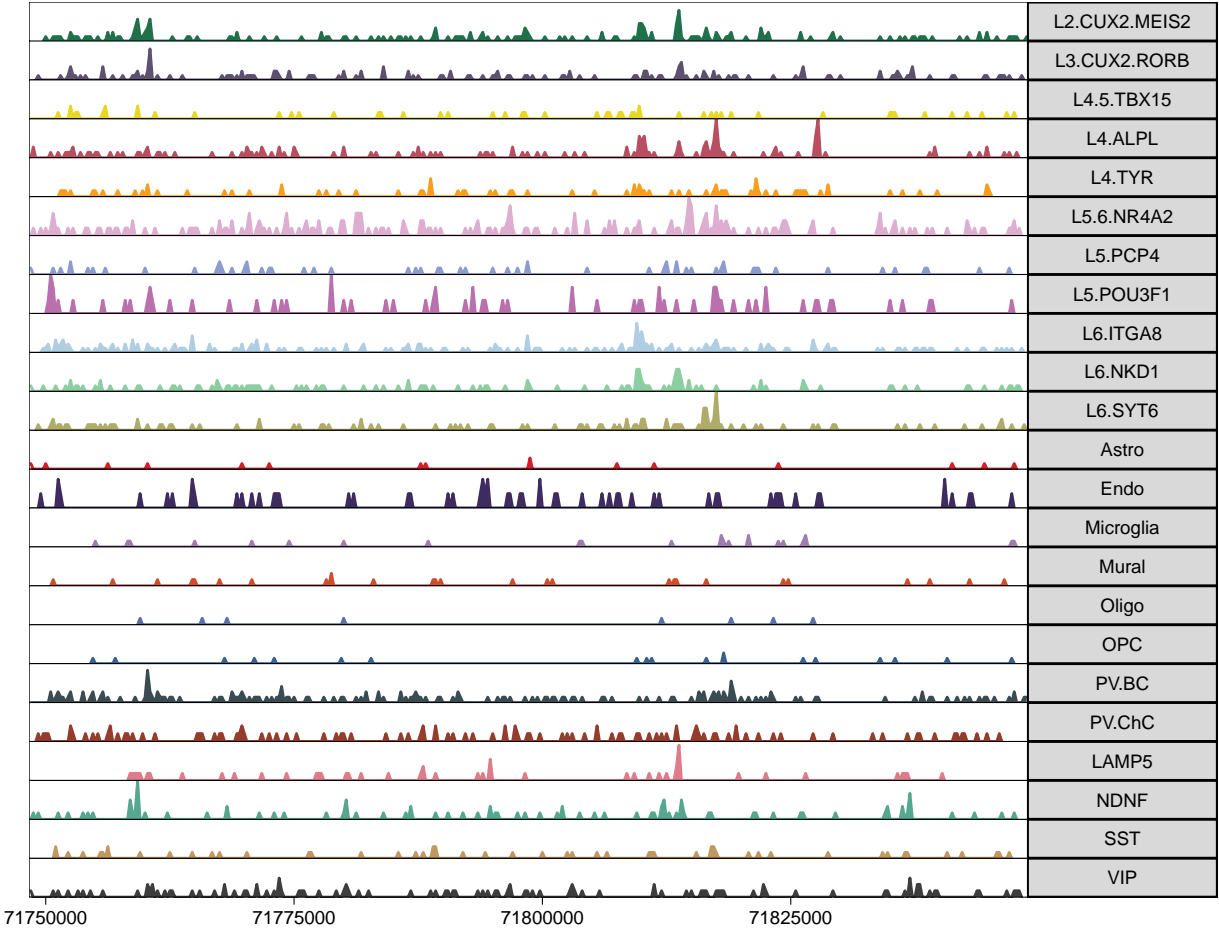
Peaks



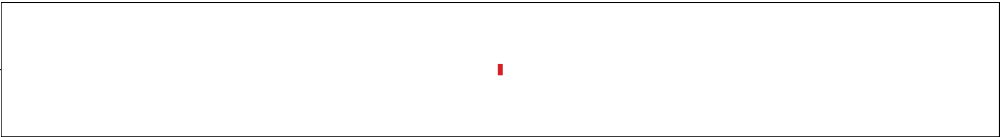
GeneTrack

chr12:71748329–71848830 Marker Peak Tracks for: L2.CUX2.MEIS2

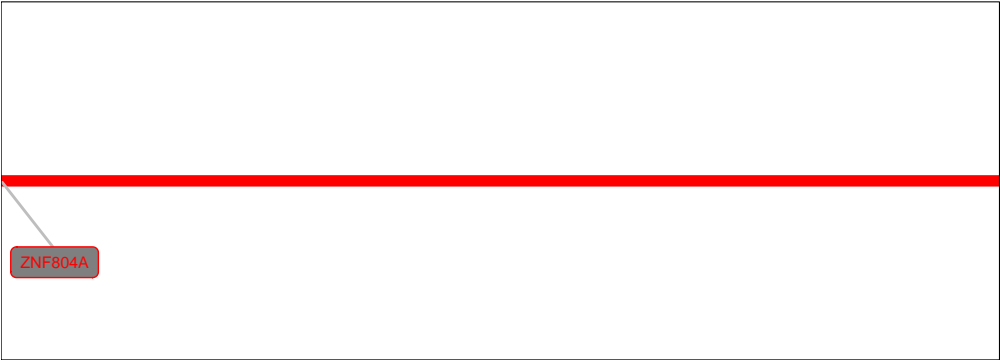
Coverage
(Norm. ATAC Signal Range (0–0.22) by ReadsInTSS)



L2.CUX2.MEIS2

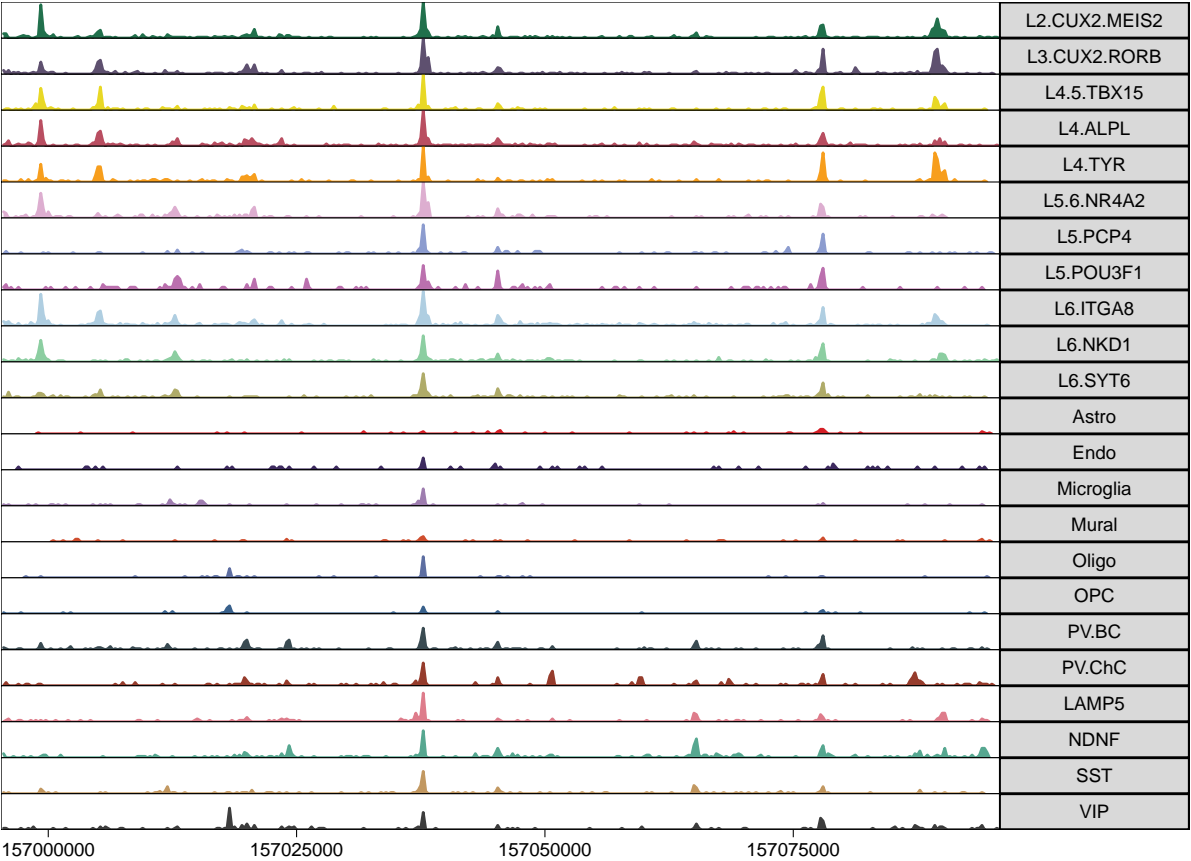


Peaks

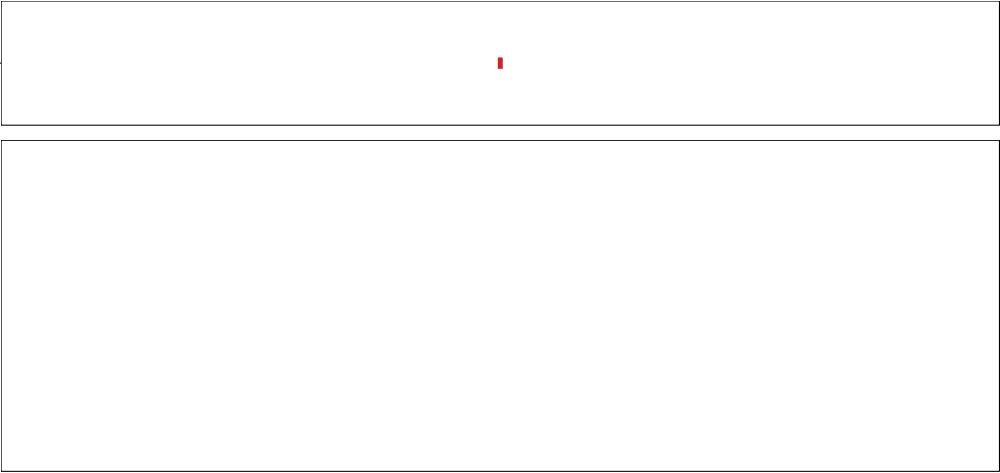


Genes

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



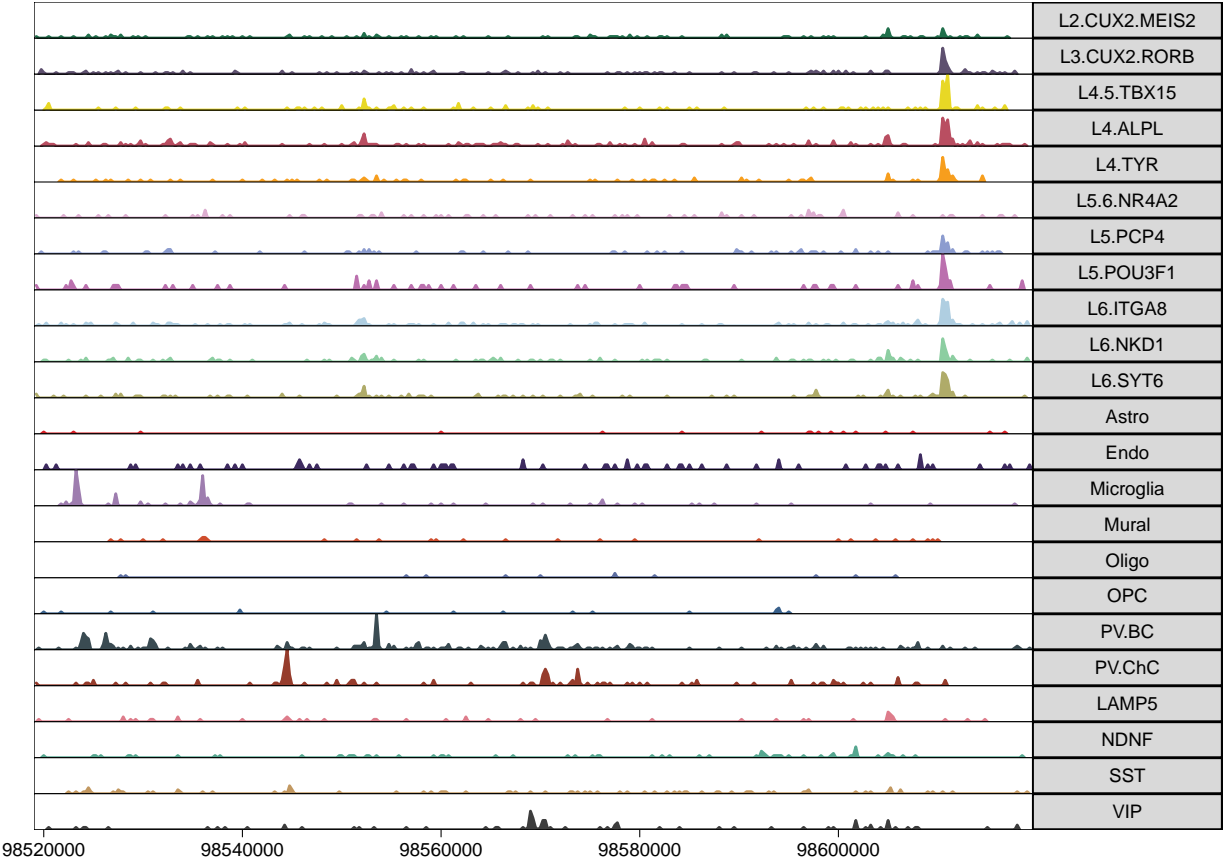
L2.CUX2.MEIS2



Peaks

GeneTrack

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



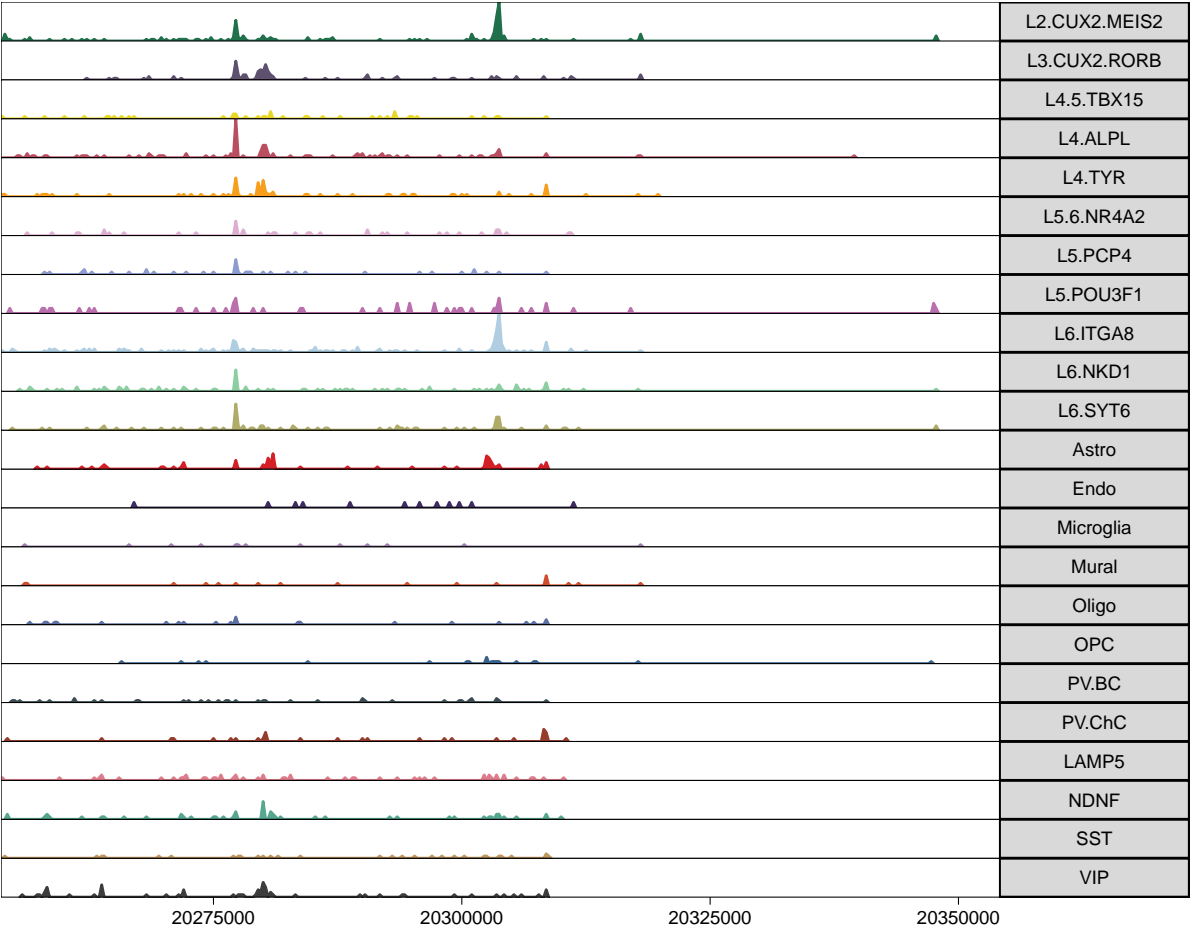
L2.CUX2.MEIS2

Peaks

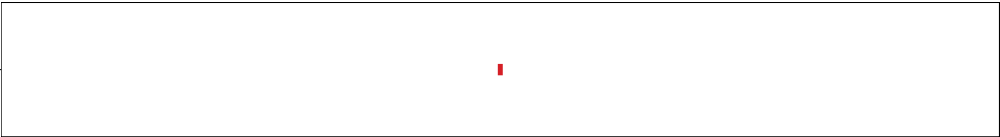
GeneTrack

strand ● ·

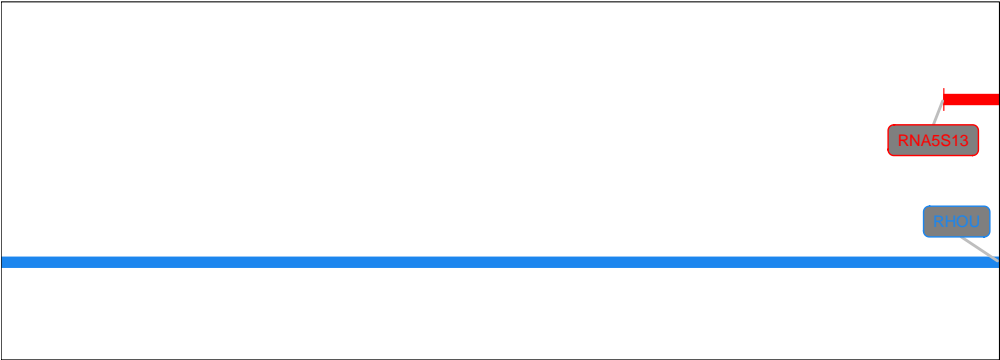
Coverage
(Norm. ATAC Signal Range (0-0.56) by ReadsInTSS)



L2.CUX2.MEIS2

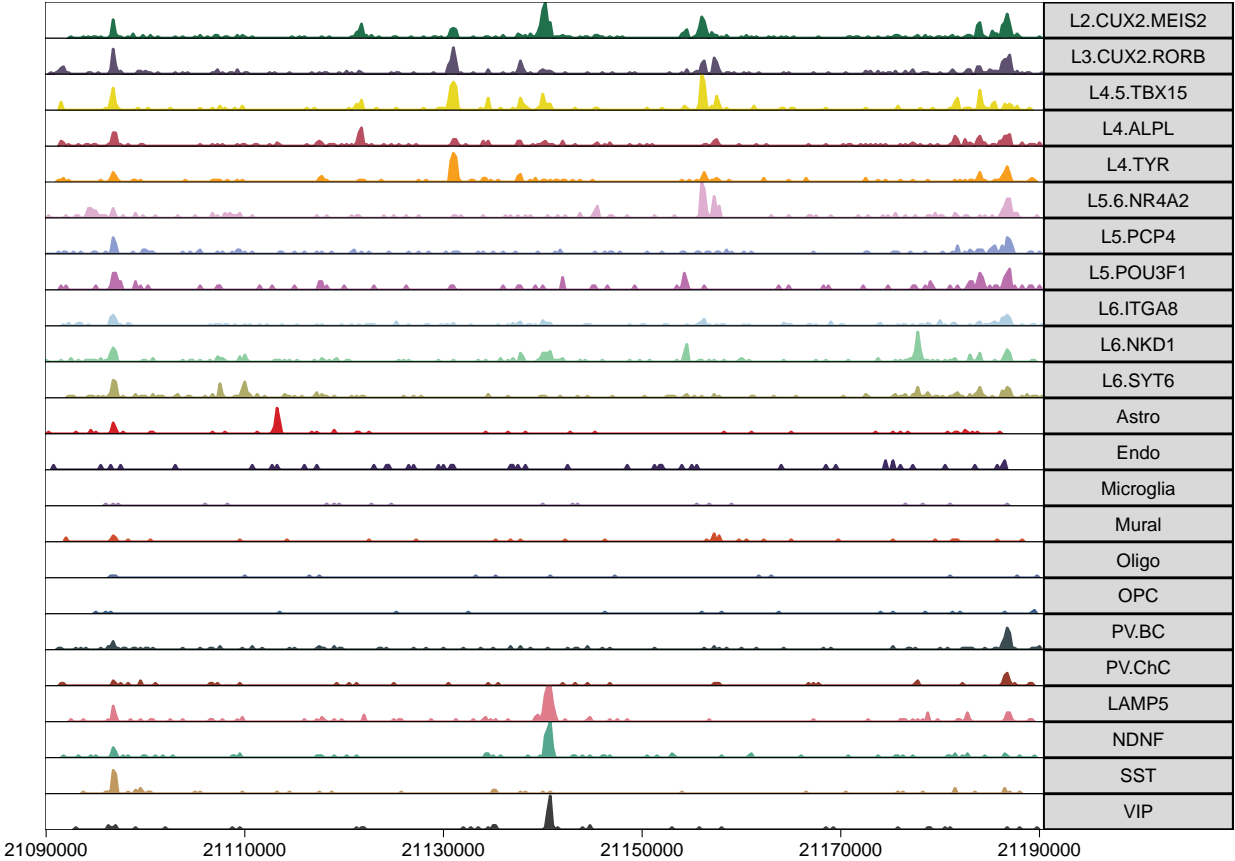


Peaks

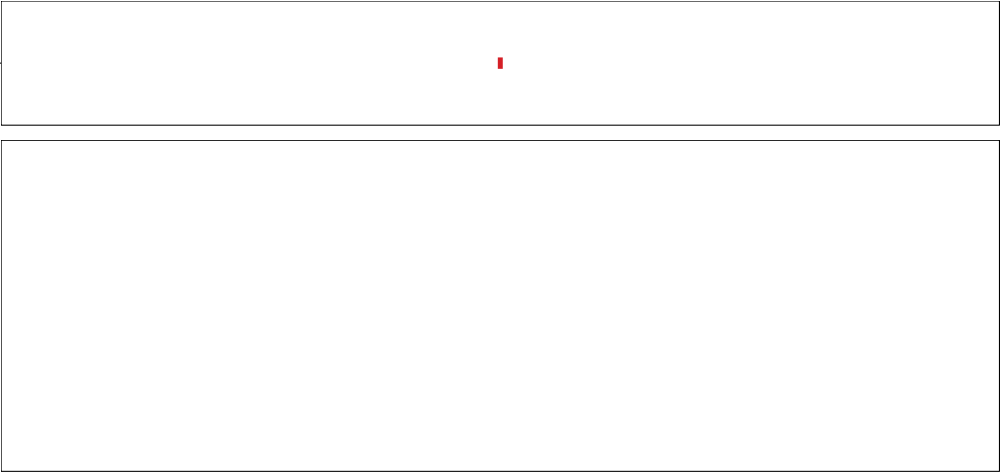


Genes

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



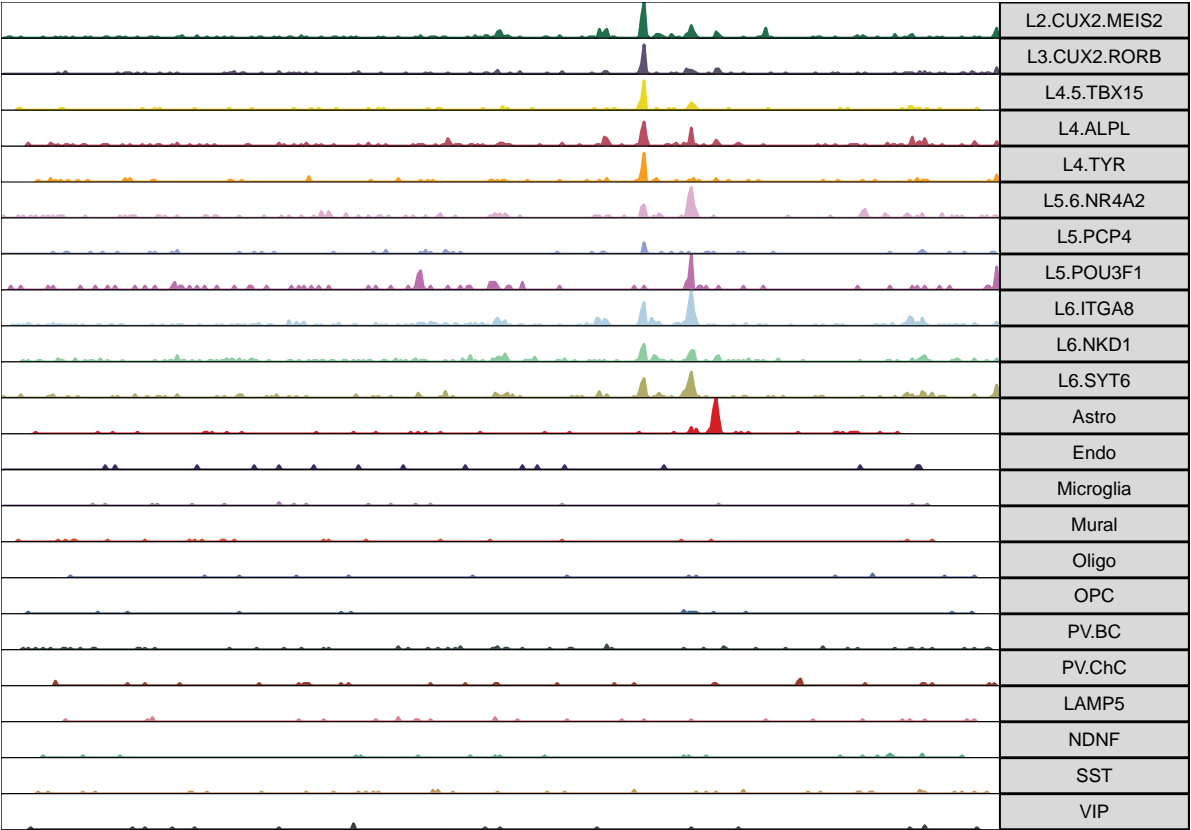
L2.CUX2.MEIS2



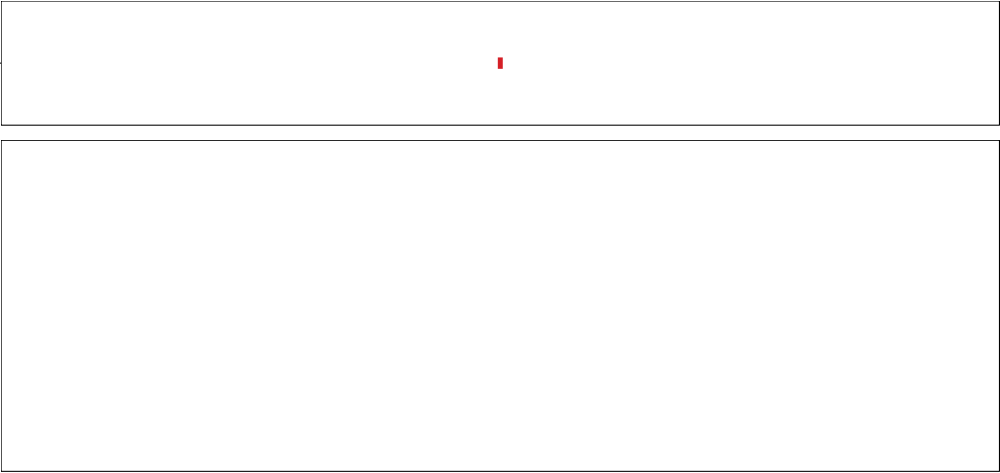
Peaks

GeneTrack

Coverage
(Norm. ATAC Signal Range (0-0.69) by ReadslnTSS)



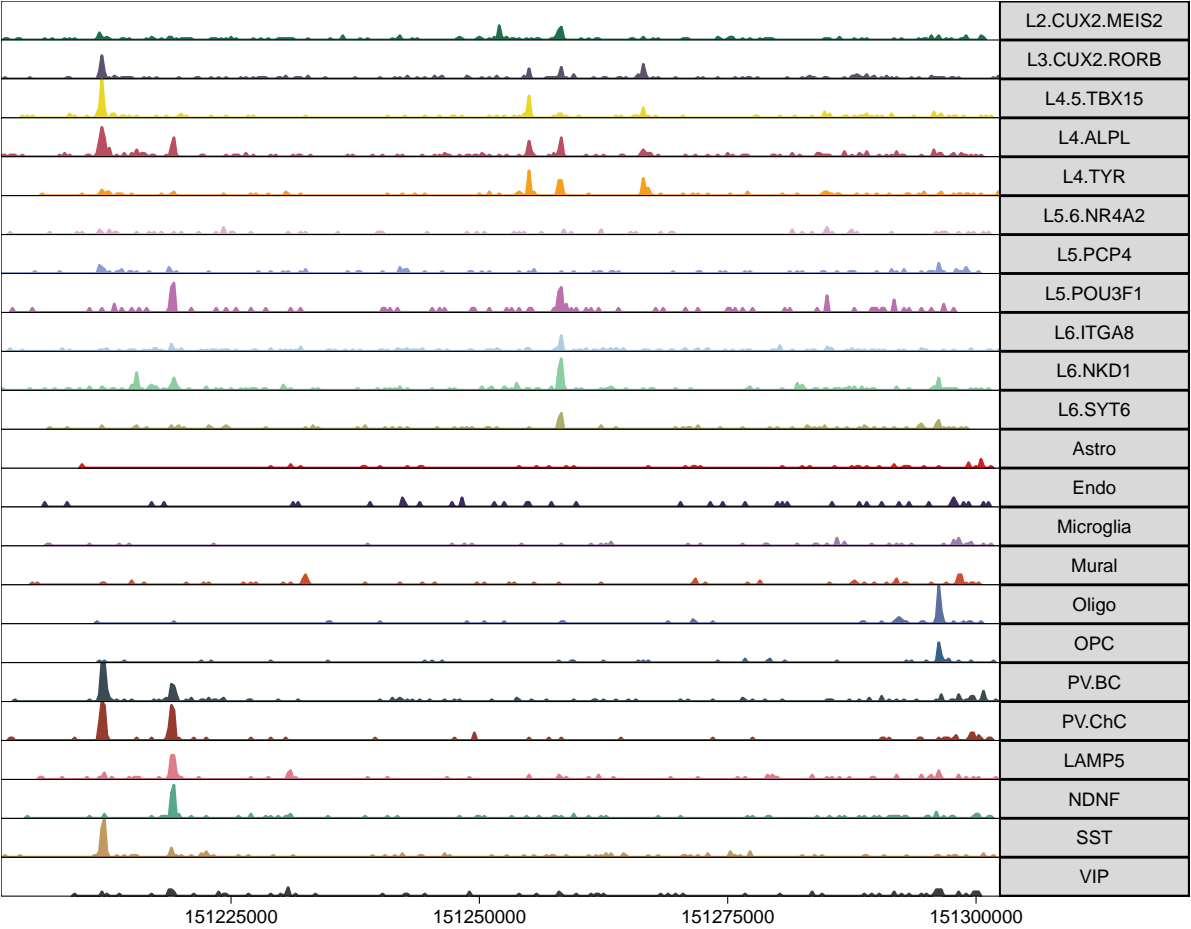
L2.CUX2.MEIS2



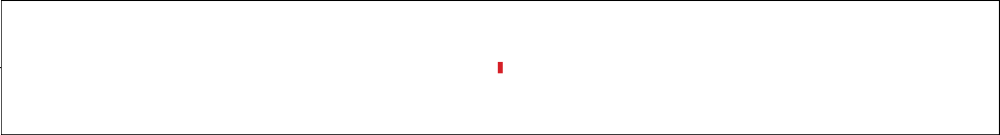
Peaks

GeneTrack

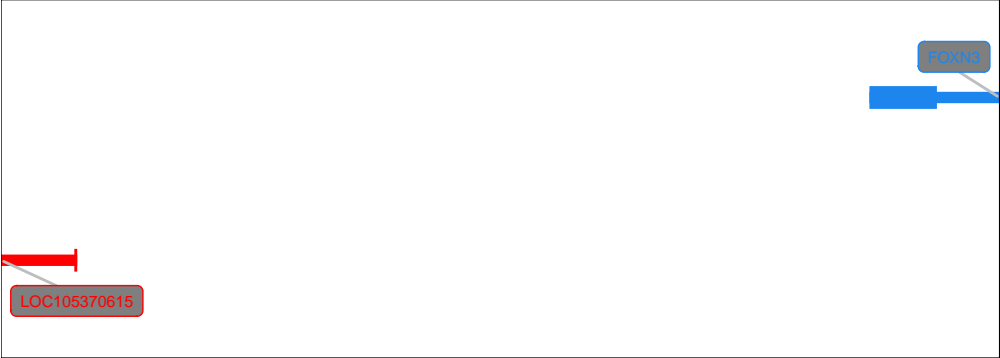
Coverage
(Norm. ATAC Signal Range (0–0.68) by ReadsInTSS)



L2.CUX2.MEIS2

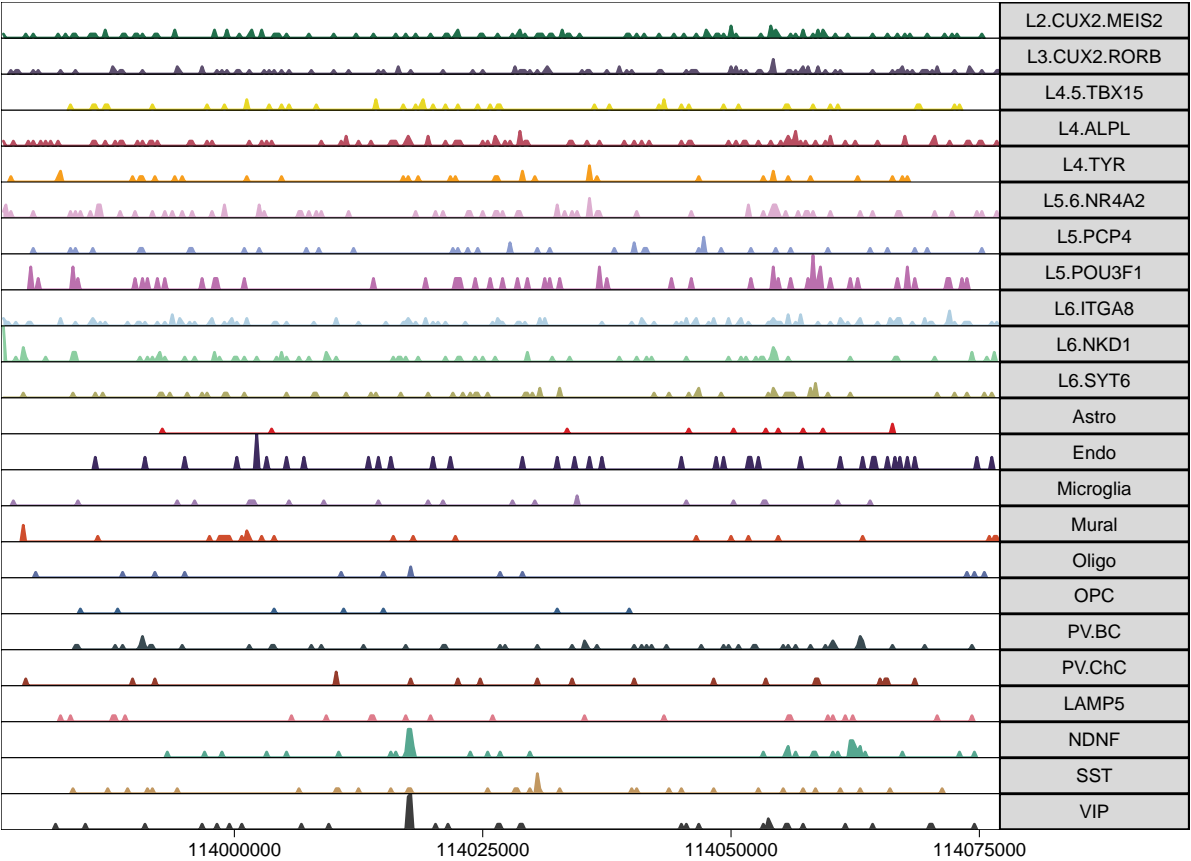


Peaks

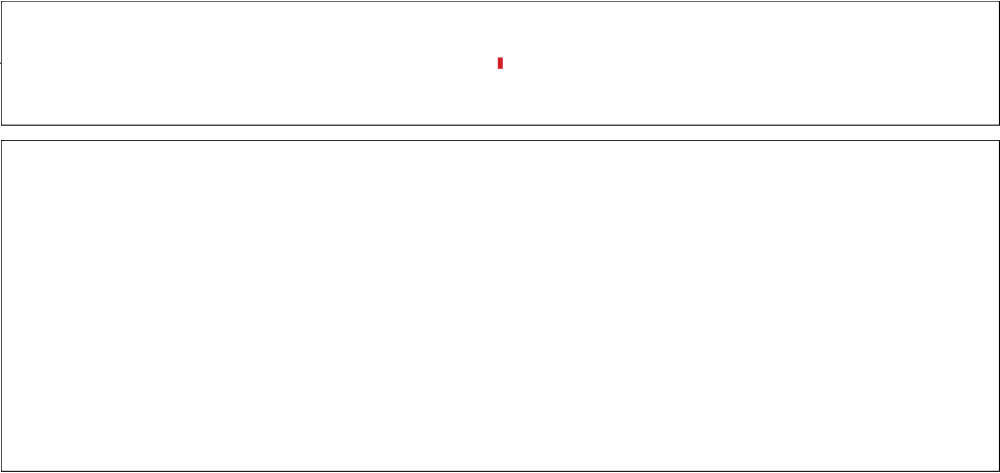


Genes

Coverage
(Norm. ATAC Signal Range (0–0.23) by ReadsInTSS)

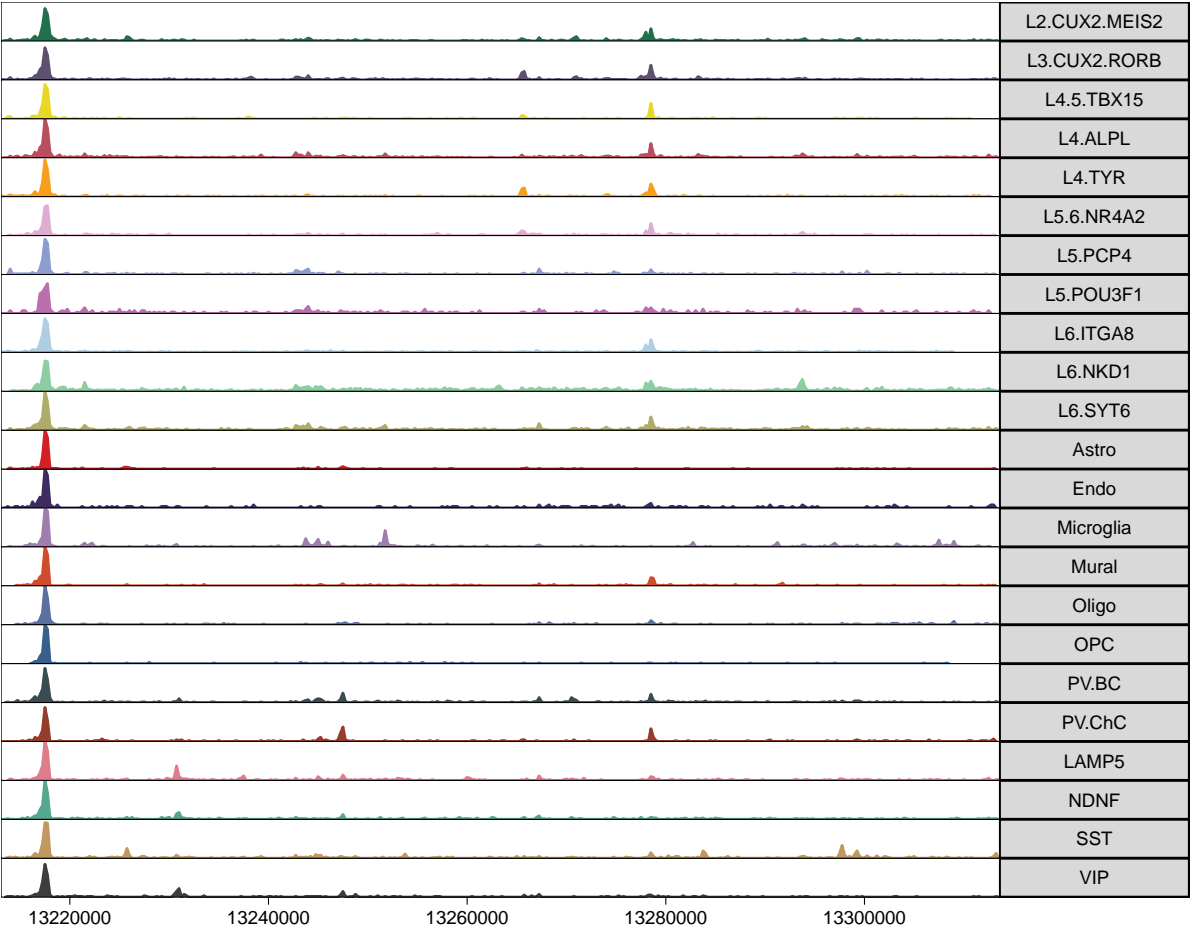


L2.CUX2.MEIS2

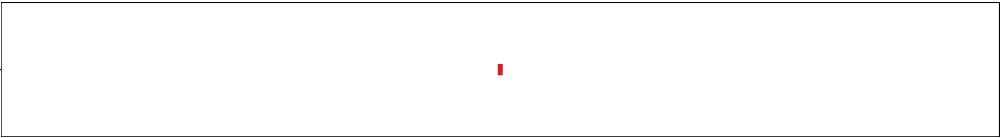


chr15:13213086–13313587 Marker Peak Tracks for: L2.CUX2.MEIS2

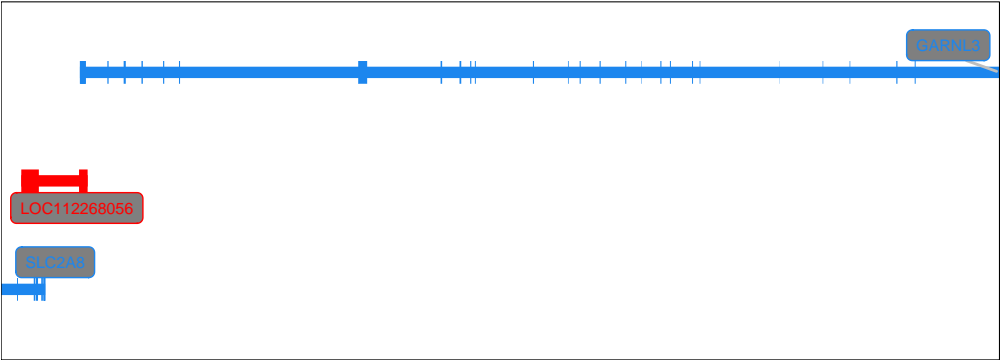
Coverage
(Norm. ATAC Signal Range (0–2.01) by ReadsInTSS)



L2.CUX2.MEIS2



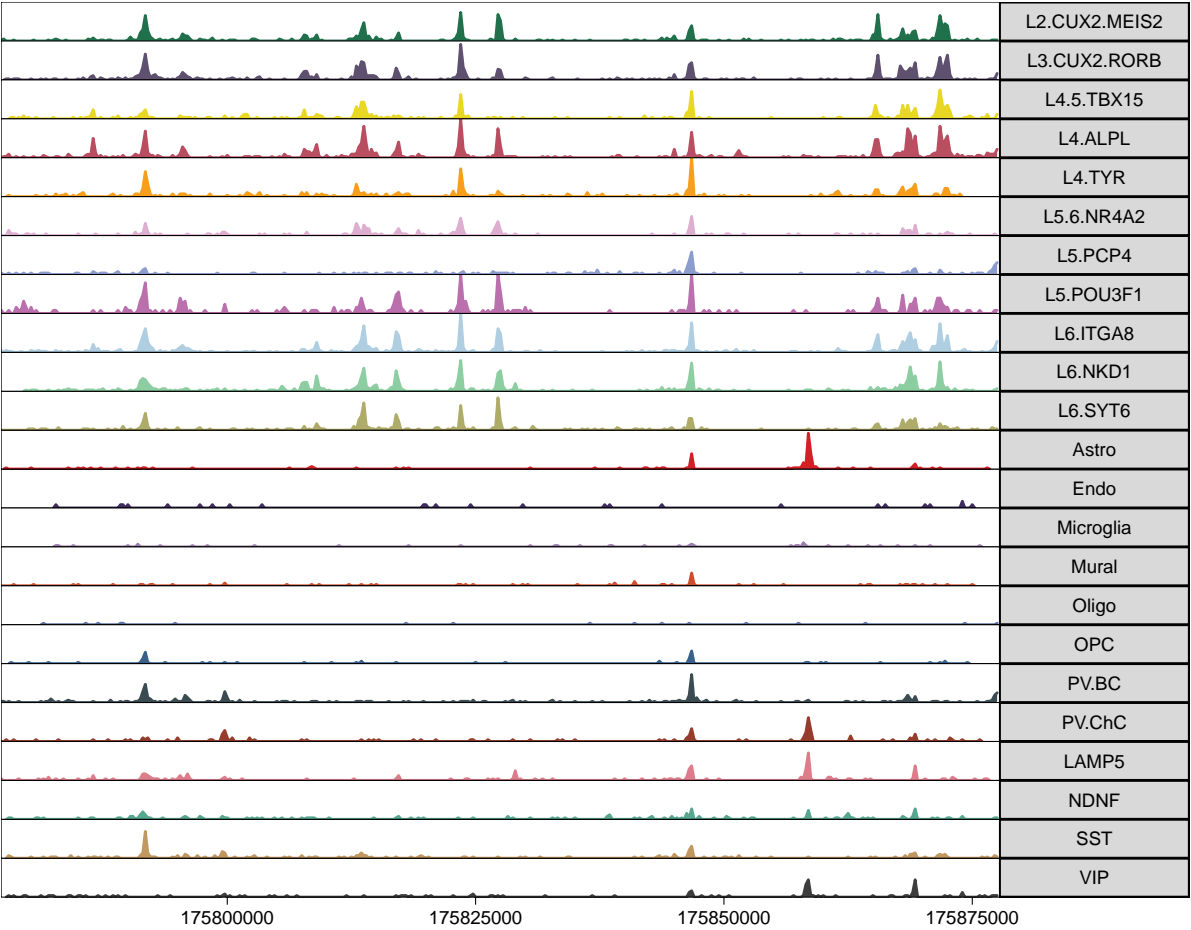
Peaks



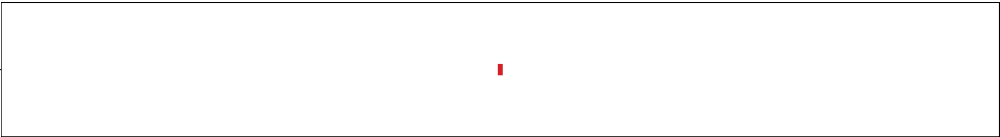
Genes

chr1:175777234–175877735 Marker Peak Tracks for: L2.CUX2.MEIS2

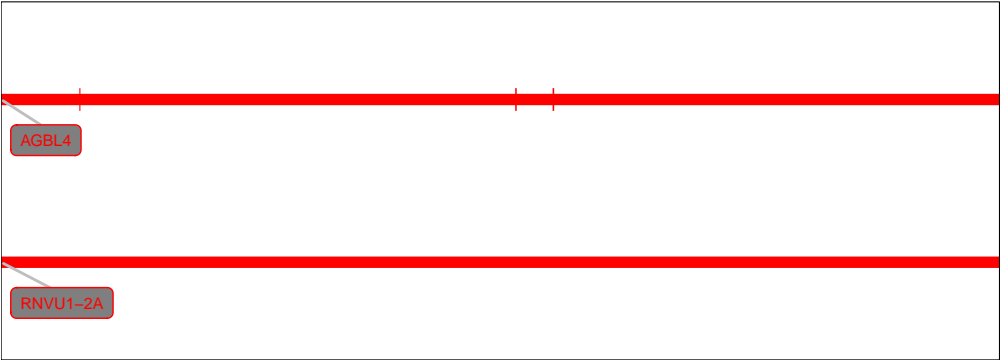
Coverage
(Norm. ATAC Signal Range (0–0.93) by ReadsInTSS)



L2.CUX2.MEIS2

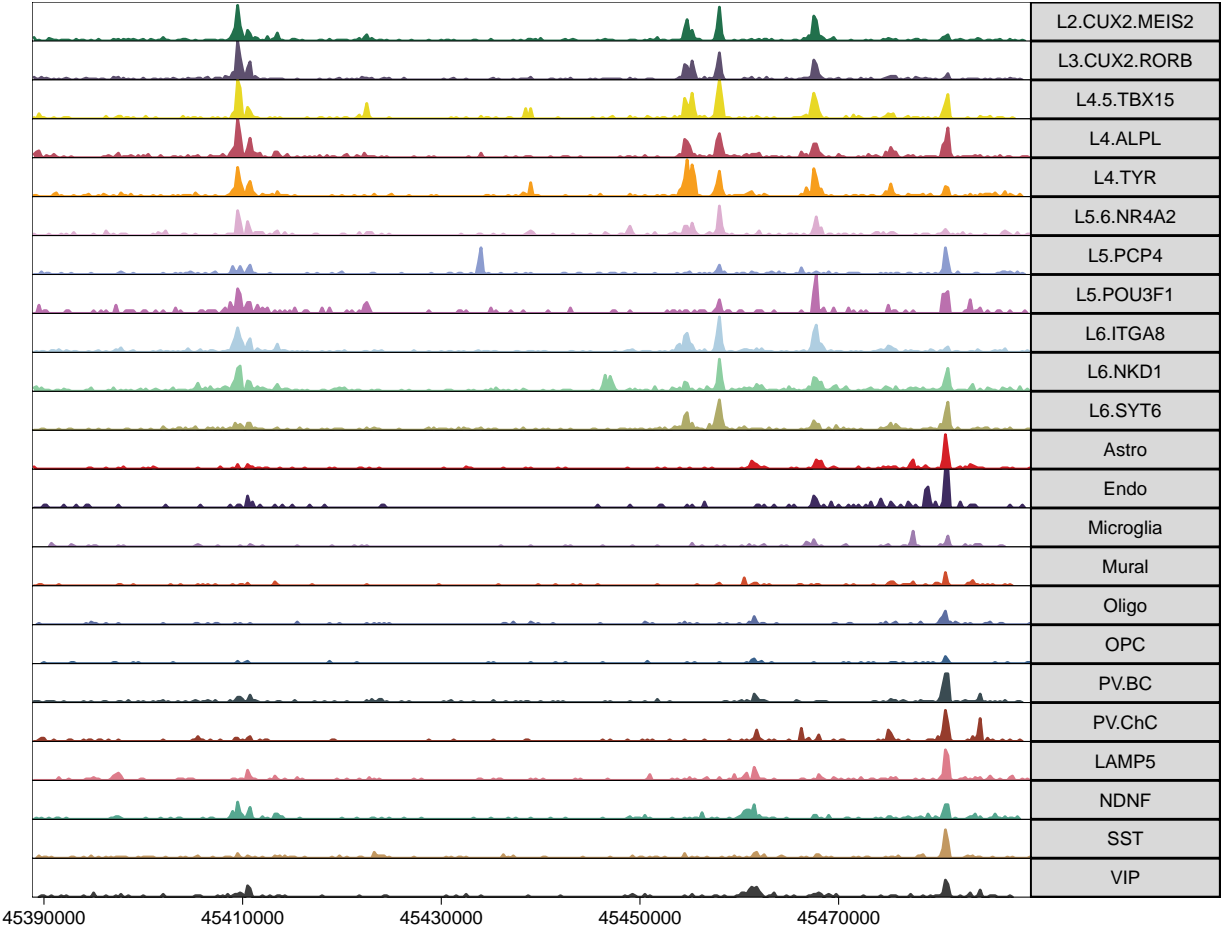


Peaks

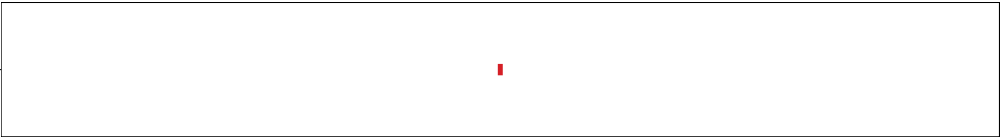


Genes

Coverage
(Norm. ATAC Signal Range (0–1.04) by ReadsInTSS)



L2.CUX2.MEIS2



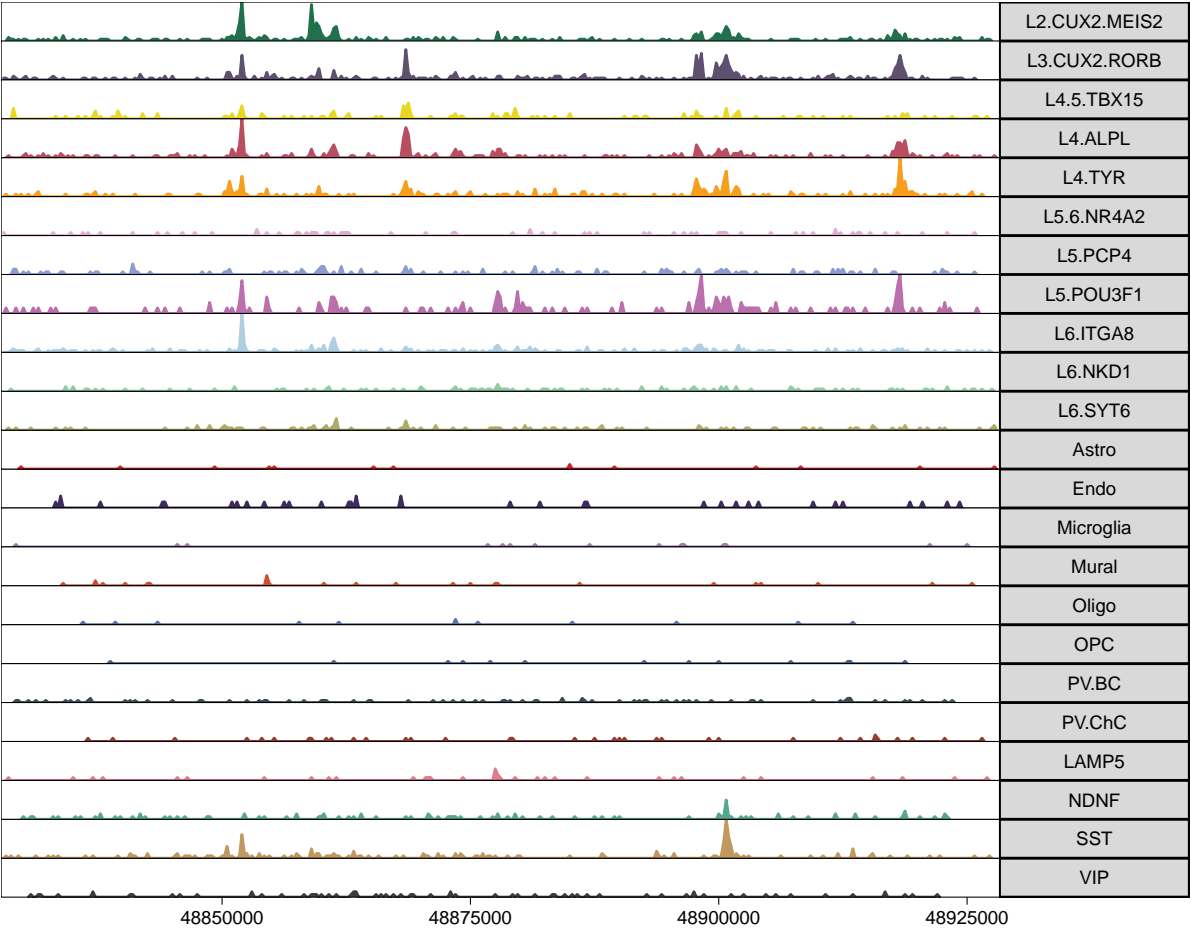
Peaks



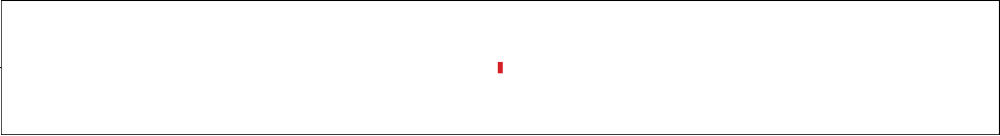
Genes

chr20:48827757-48928258 Marker Peak Tracks for: L2.CUX2.MEIS2

Coverage
(Norm. ATAC Signal Range (0-0.52) by ReadsInTSS)



L2.CUX2.MEIS2



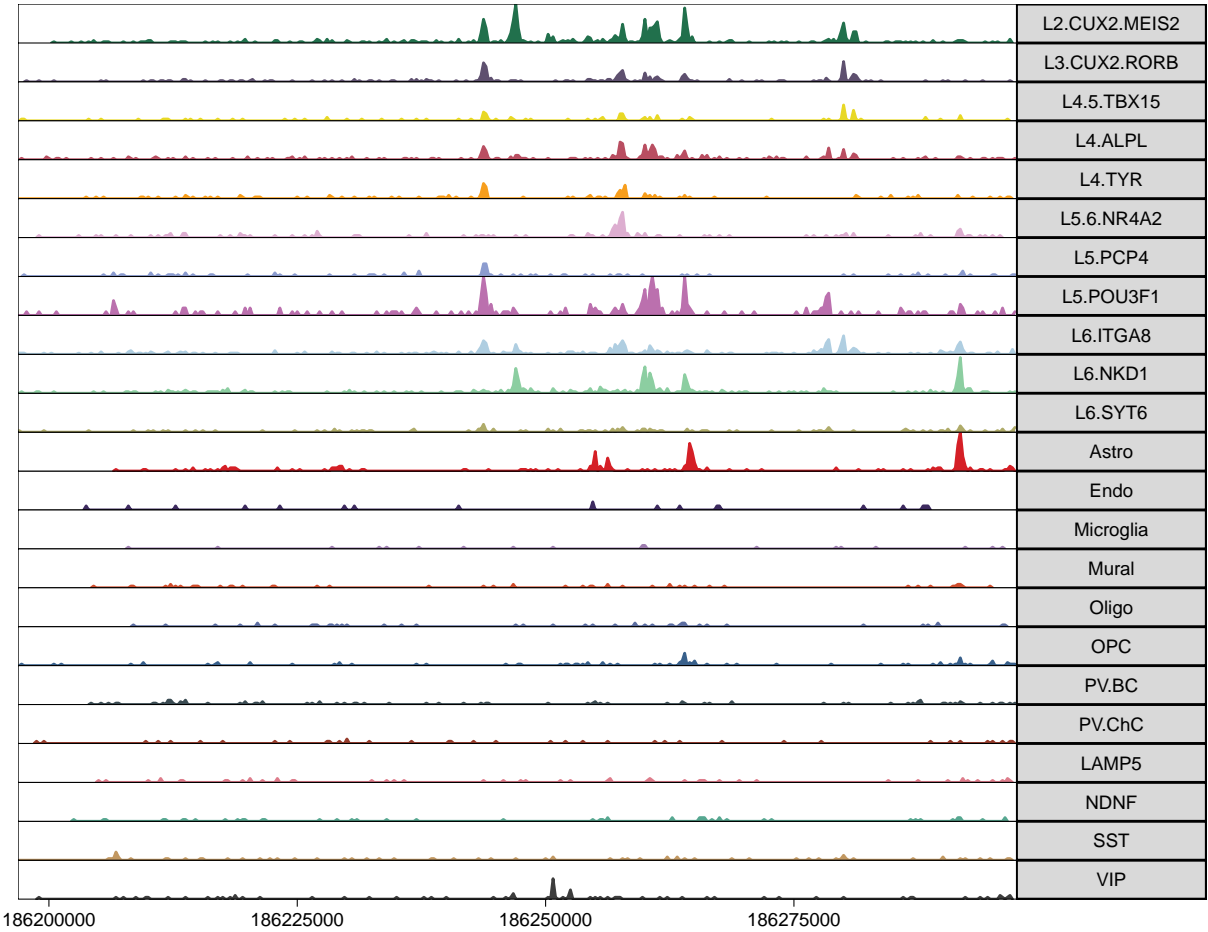
Peaks



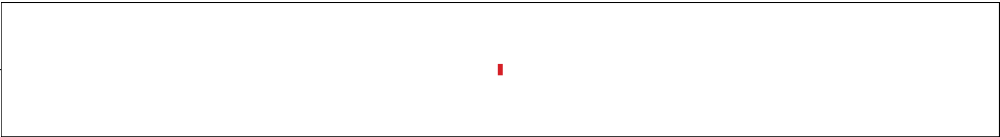
Genes

chr1:186196899–186297400 Marker Peak Tracks for: L2.CUX2.MEIS2

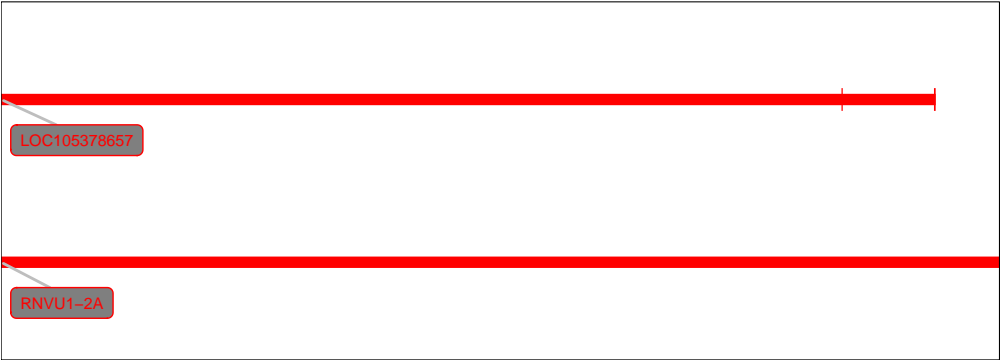
Coverage
(Norm. ATAC Signal Range (0–0.77) by ReadsInTSS)



L2.CUX2.MEIS2

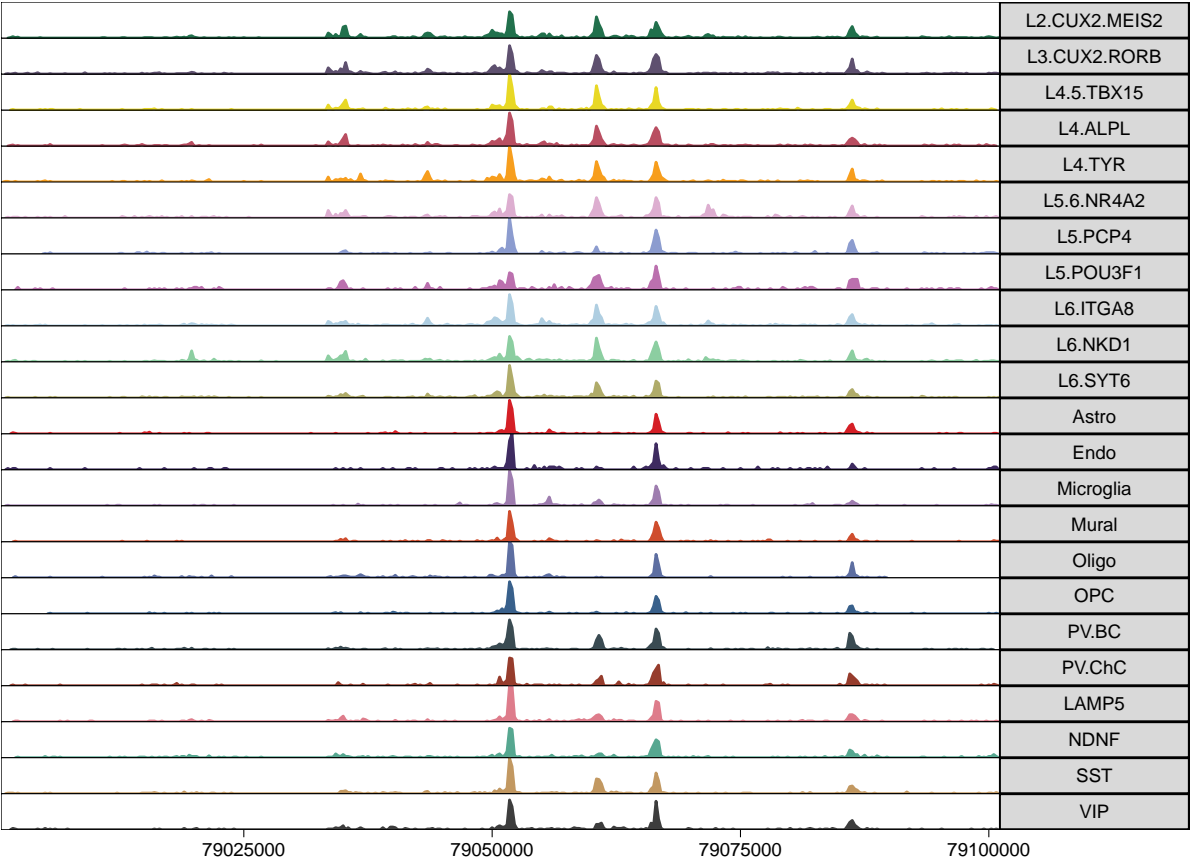


Peaks

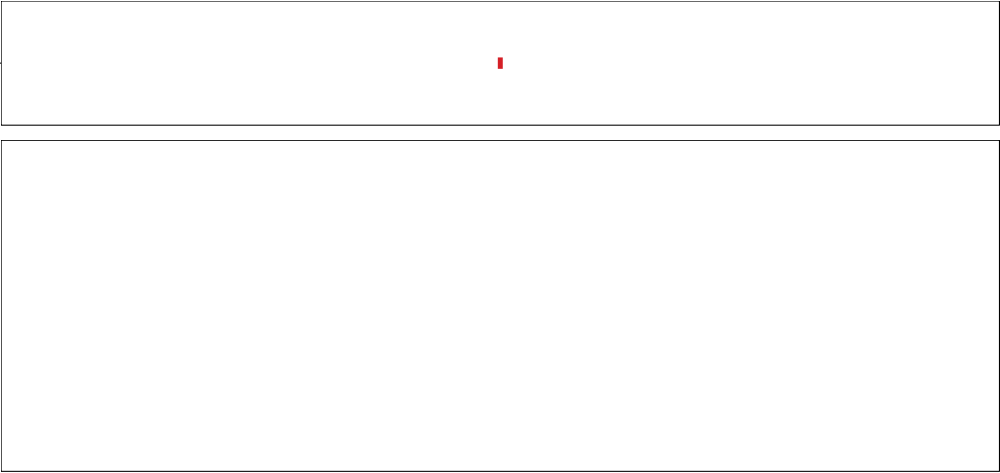


Genes

Coverage
(Norm. ATAC Signal Range (0–1.99) by ReadsInTSS)



L2.CUX2.MEIS2

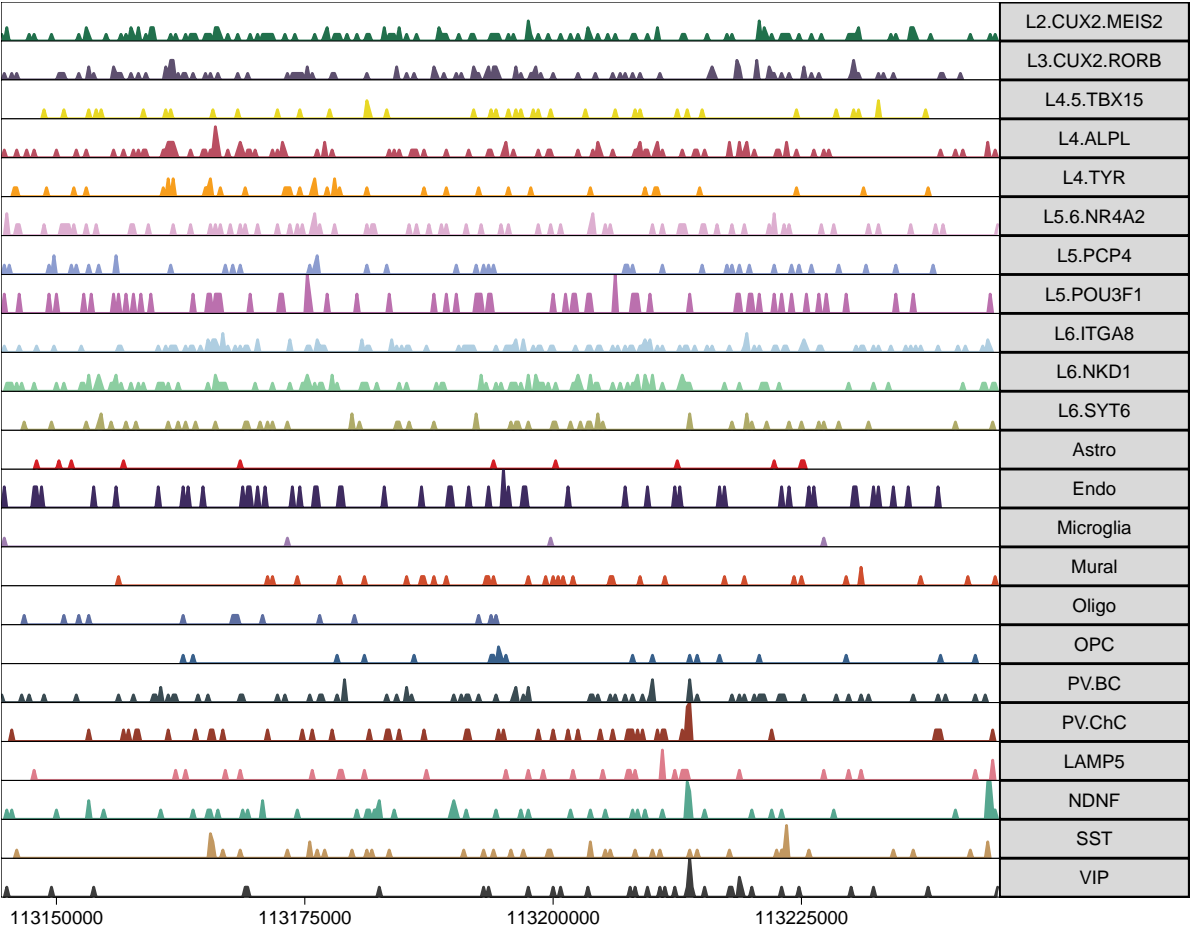


Peaks

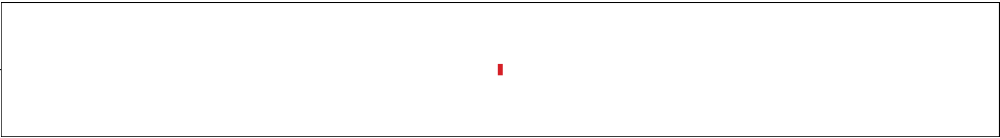
GeneTrack

chr5:113144430–113244931 Marker Peak Tracks for: L2.CUX2.MEIS2

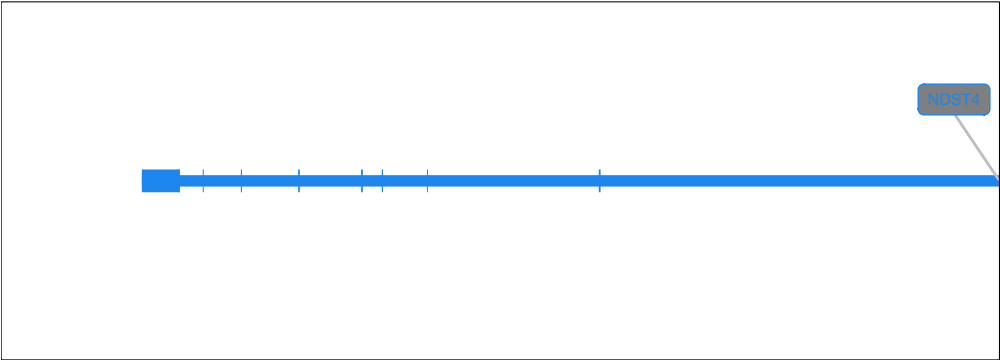
Coverage
(Norm. ATAC Signal Range (0–0.15) by ReadsInTSS)



L2.CUX2.MEIS2



Peaks



Genes