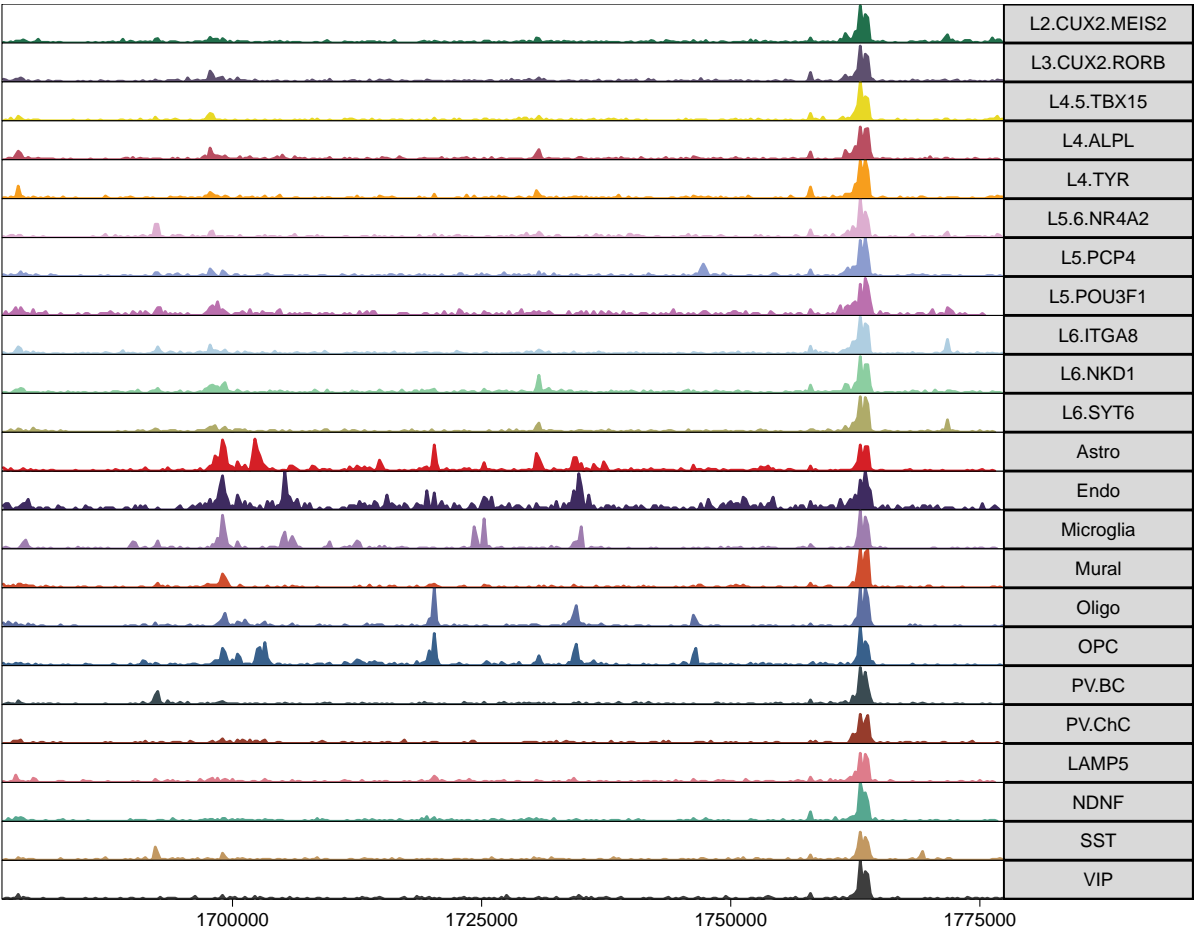
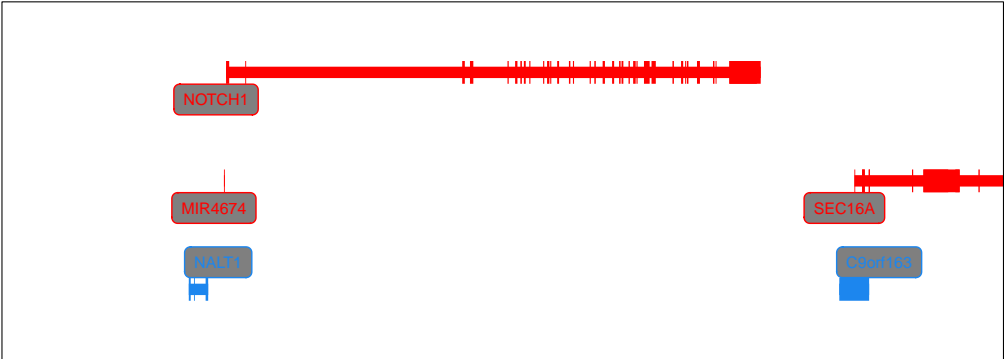
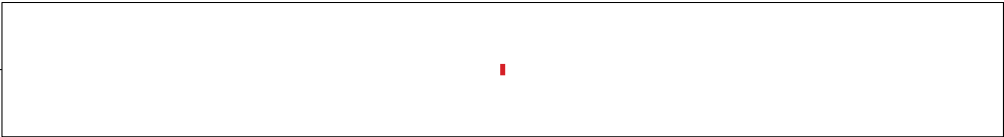


chr15:1676865–1777366 Marker Peak Tracks for: L3.CUX2.RORB

Coverage
(Norm. ATAC Signal Range (0–1.46) by ReadslnTSS)

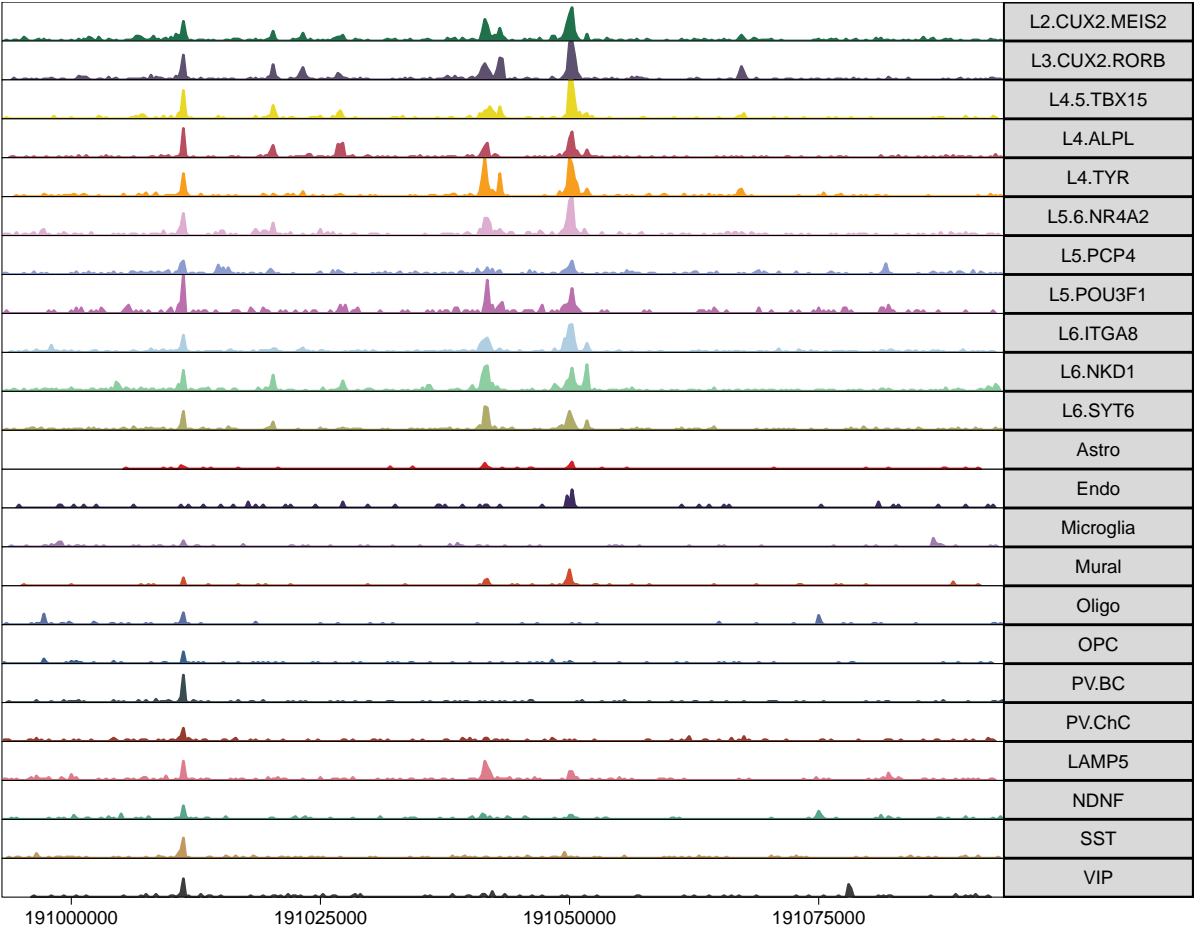


L3.CUX2.RORB

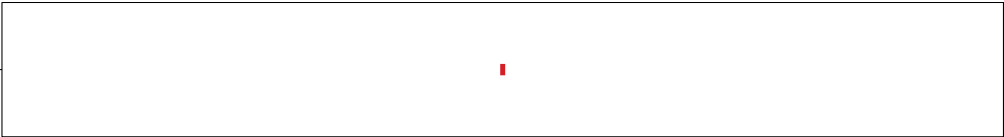


chr1:190993041–191093542 Marker Peak Tracks for: L3.CUX2.RORB

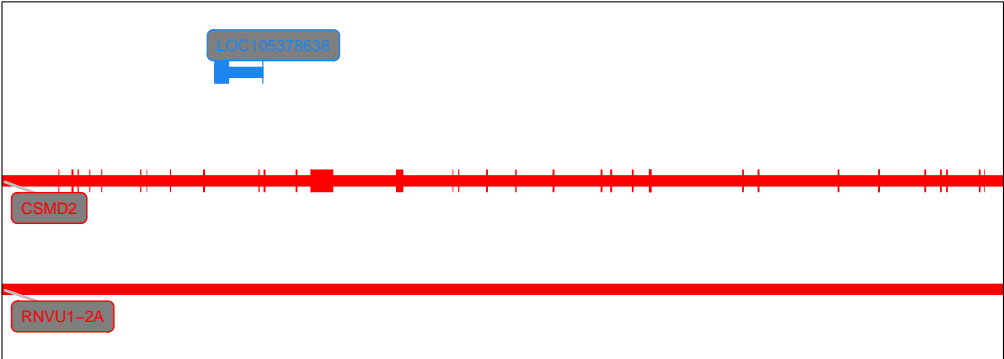
Coverage
(Norm. ATAC Signal Range (0–1.03) by ReadsInTSS)



L3.CUX2.RORB



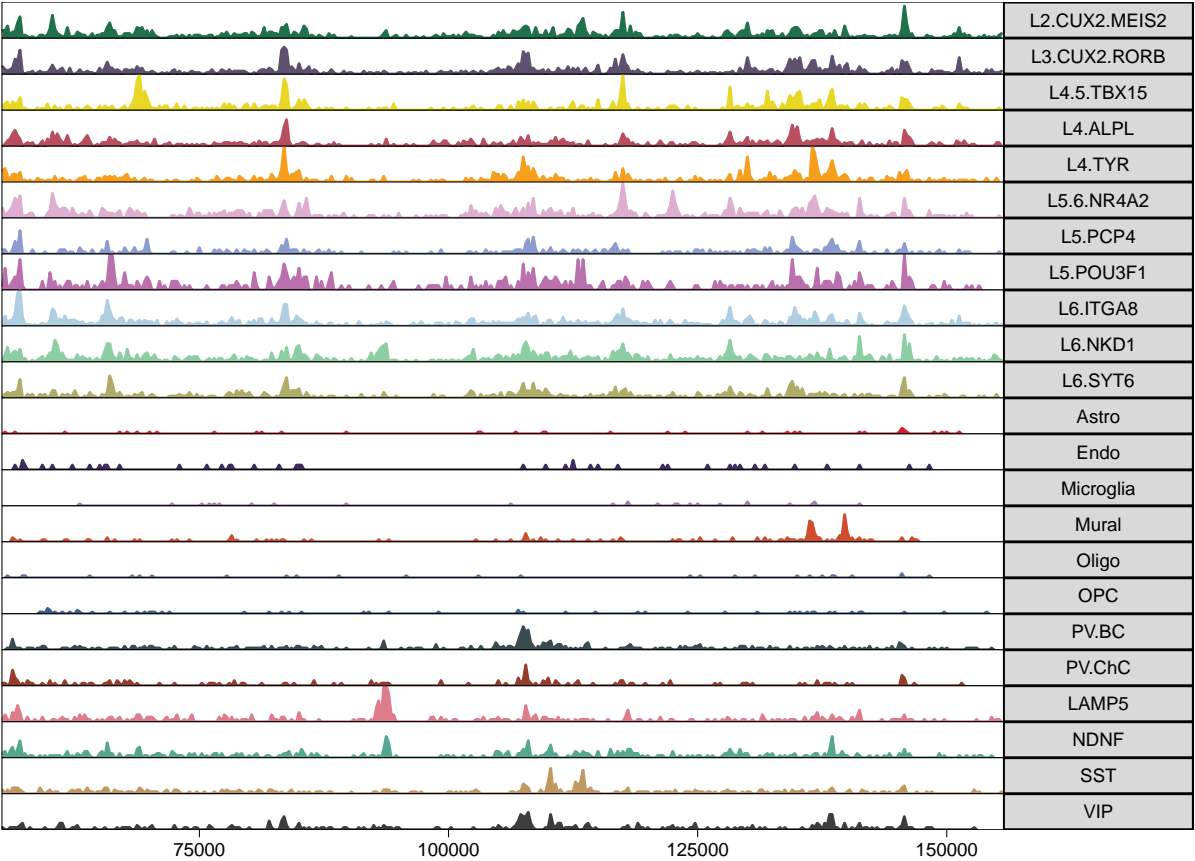
Peaks



Genes

chr15:55188–155689 Marker Peak Tracks for: L3.CUX2.RORB

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



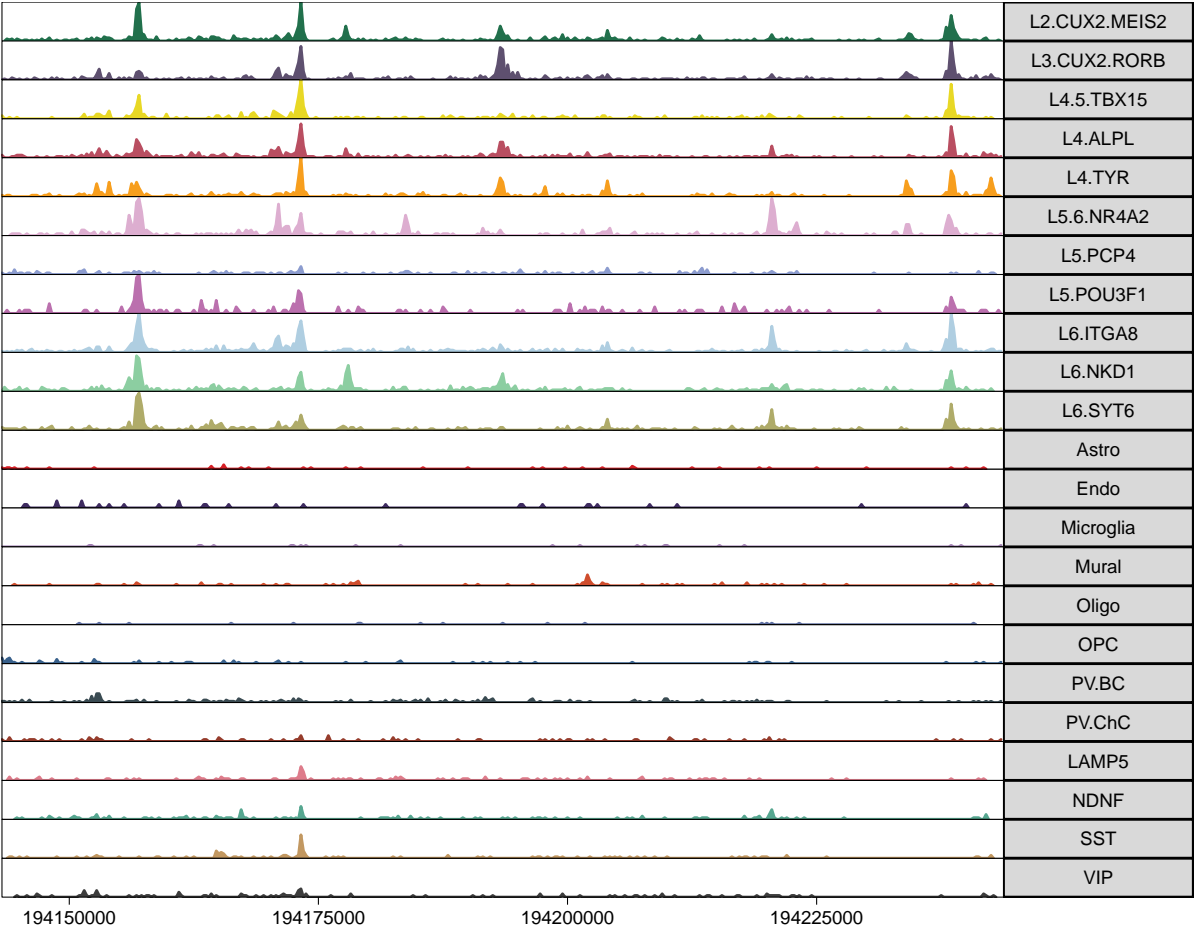
L3.CUX2.RORB



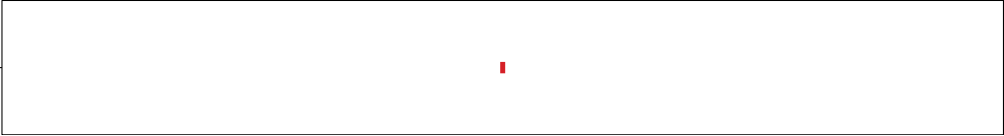
GeneTrack

strand ● ·

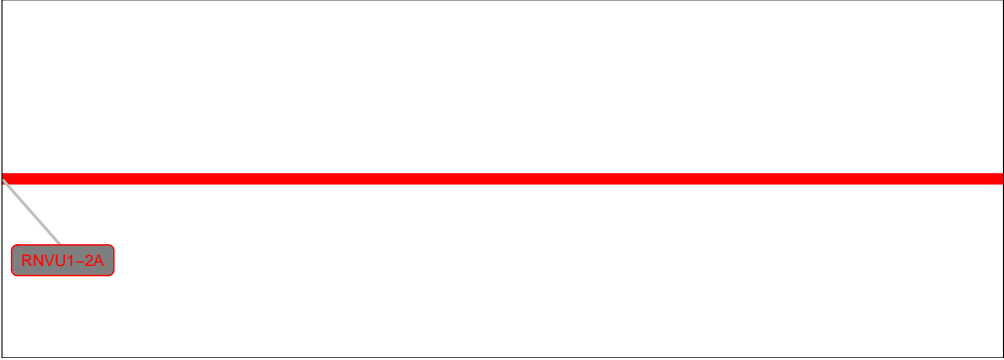
Coverage
(Norm. ATAC Signal Range (0–0.88) by ReadsInTSS)



L3.CUX2.RORB

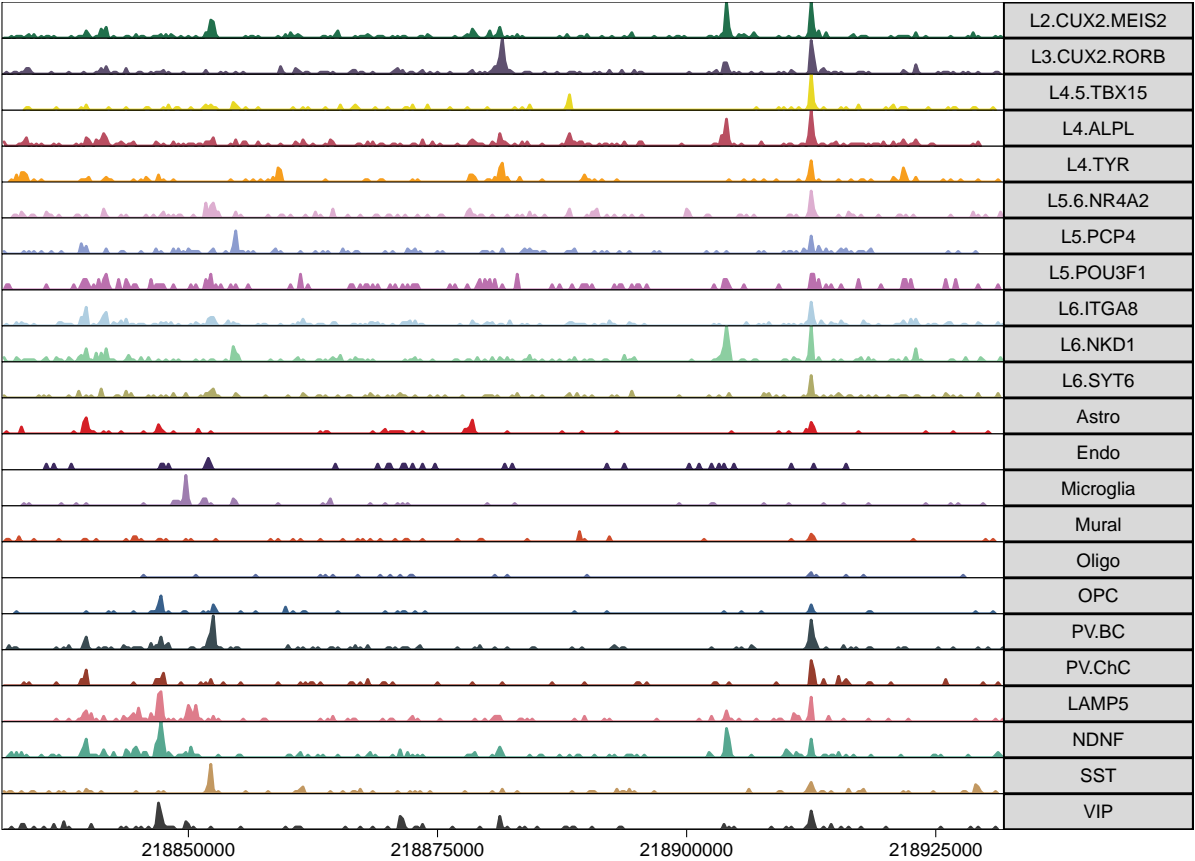


Peaks



Genes

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



L3.CUX2.RORB

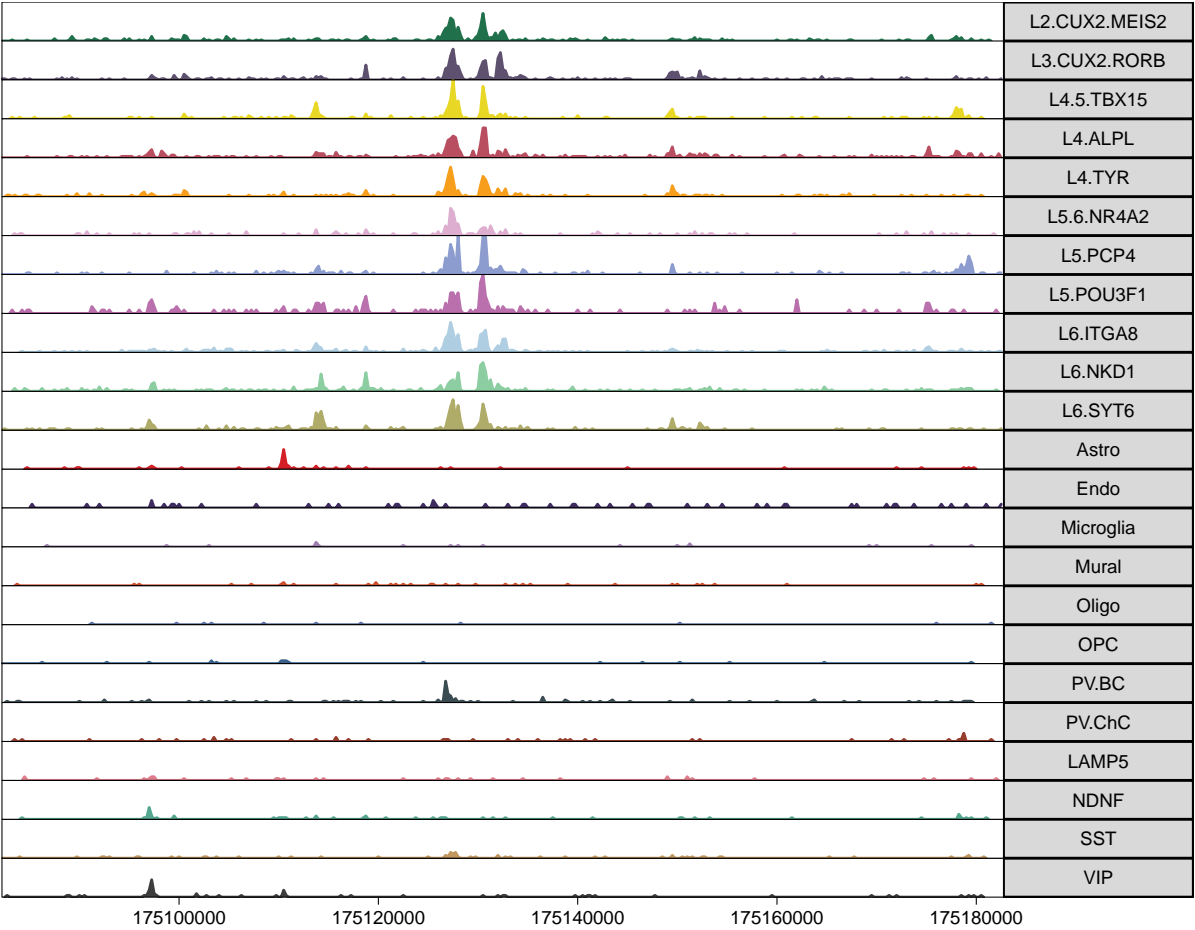


Peaks

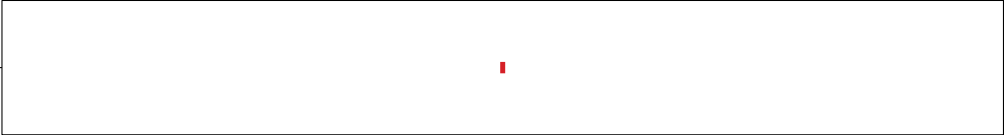
GeneTrack

strand ● ·

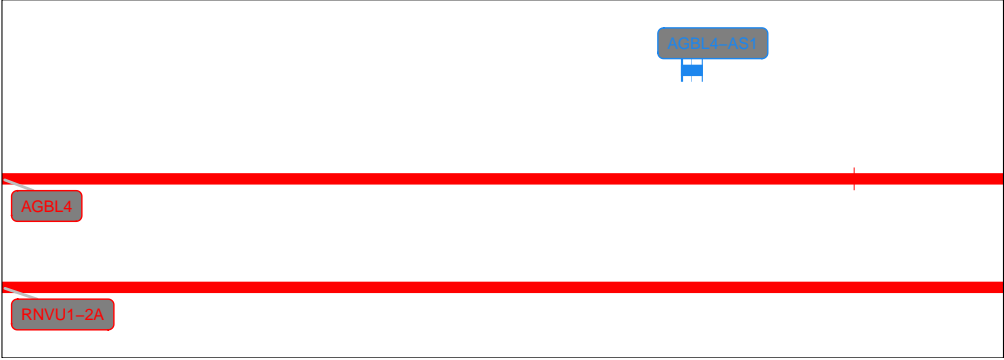
Coverage
(Norm. ATAC Signal Range (0–0.83) by ReadslnTSS)



L3.CUX2.RORB



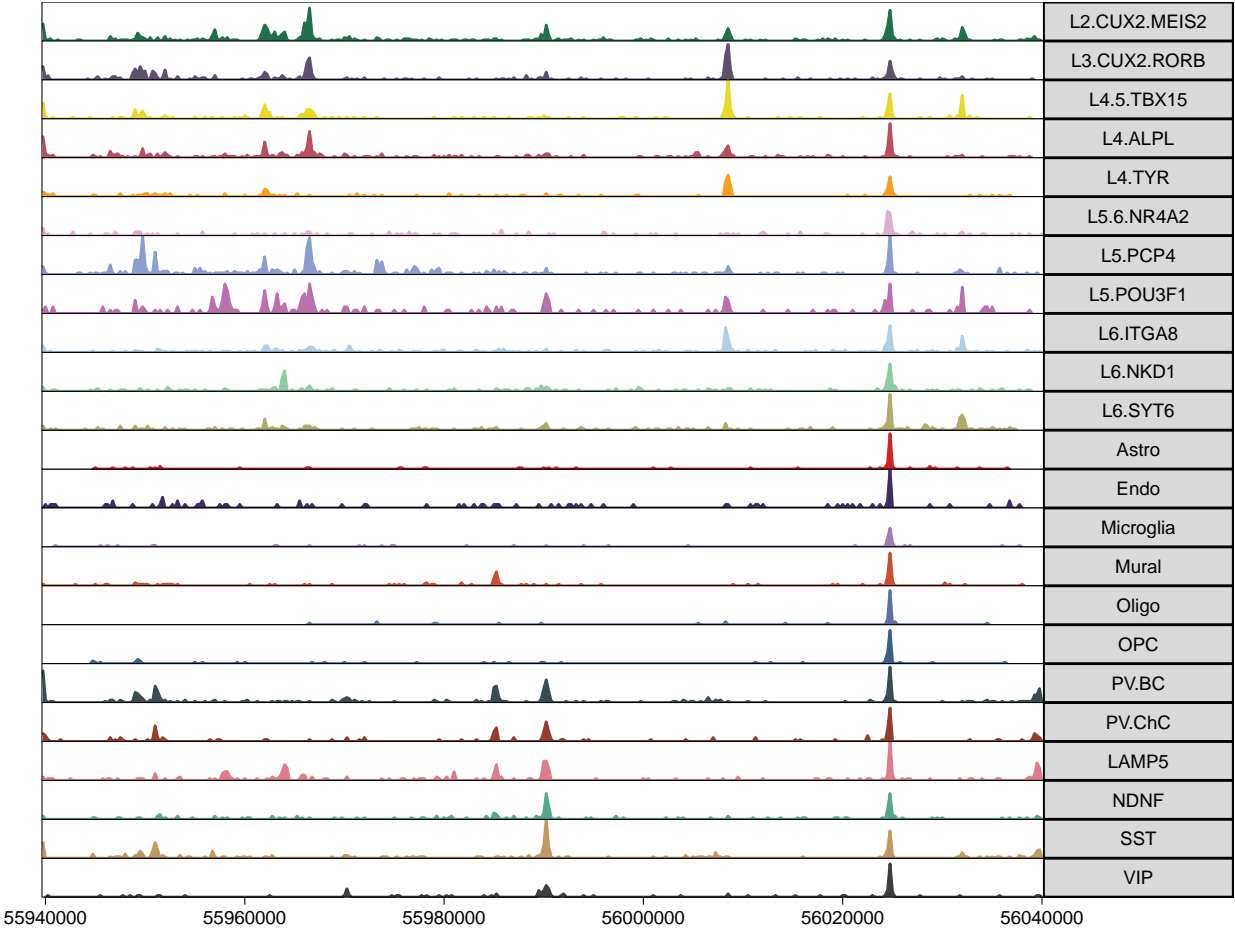
Peaks



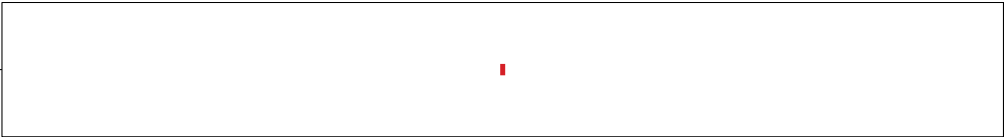
Genes

chr12:55939645–56040146 Marker Peak Tracks for: L3.CUX2.RORB

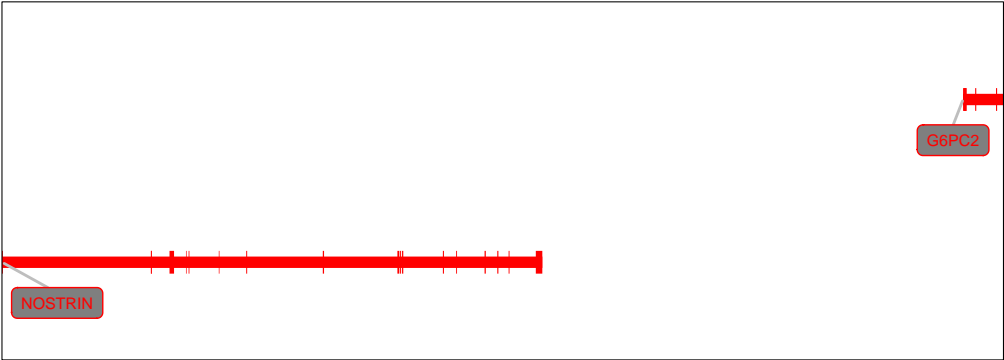
Coverage
(Norm. ATAC Signal Range (0–0.87) by ReadsInTSS)



L3.CUX2.RORB

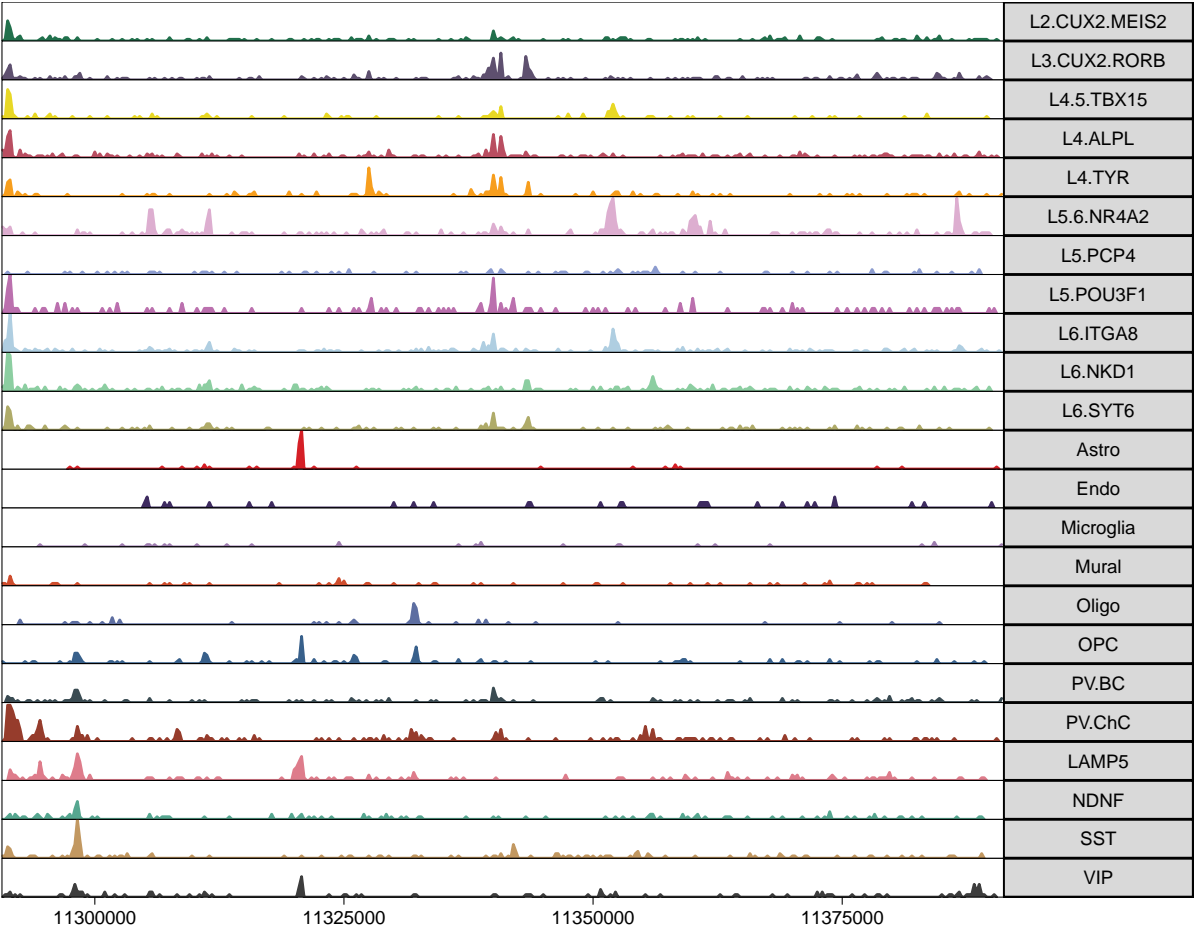


Peaks

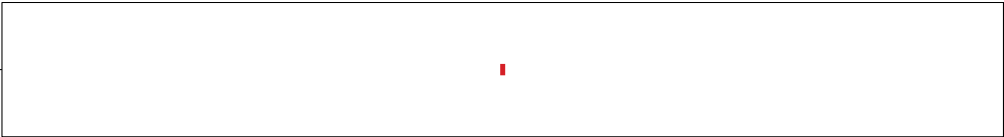


Genes

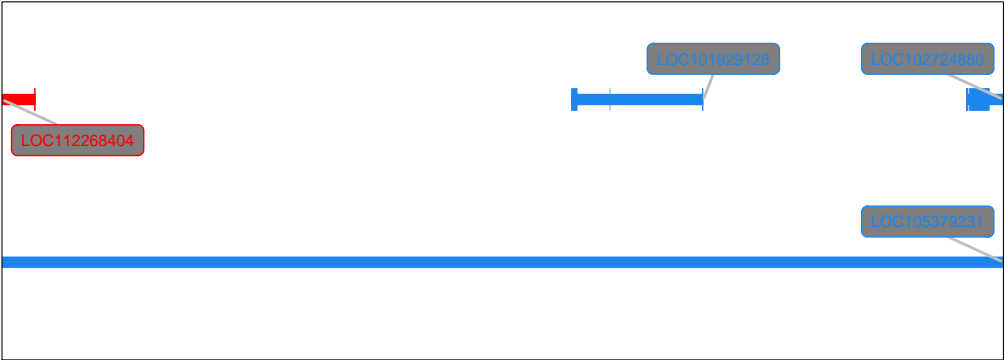
Coverage
(Norm. ATAC Signal Range (0–0.57) by ReadsInTSS)



L3.CUX2.RORB



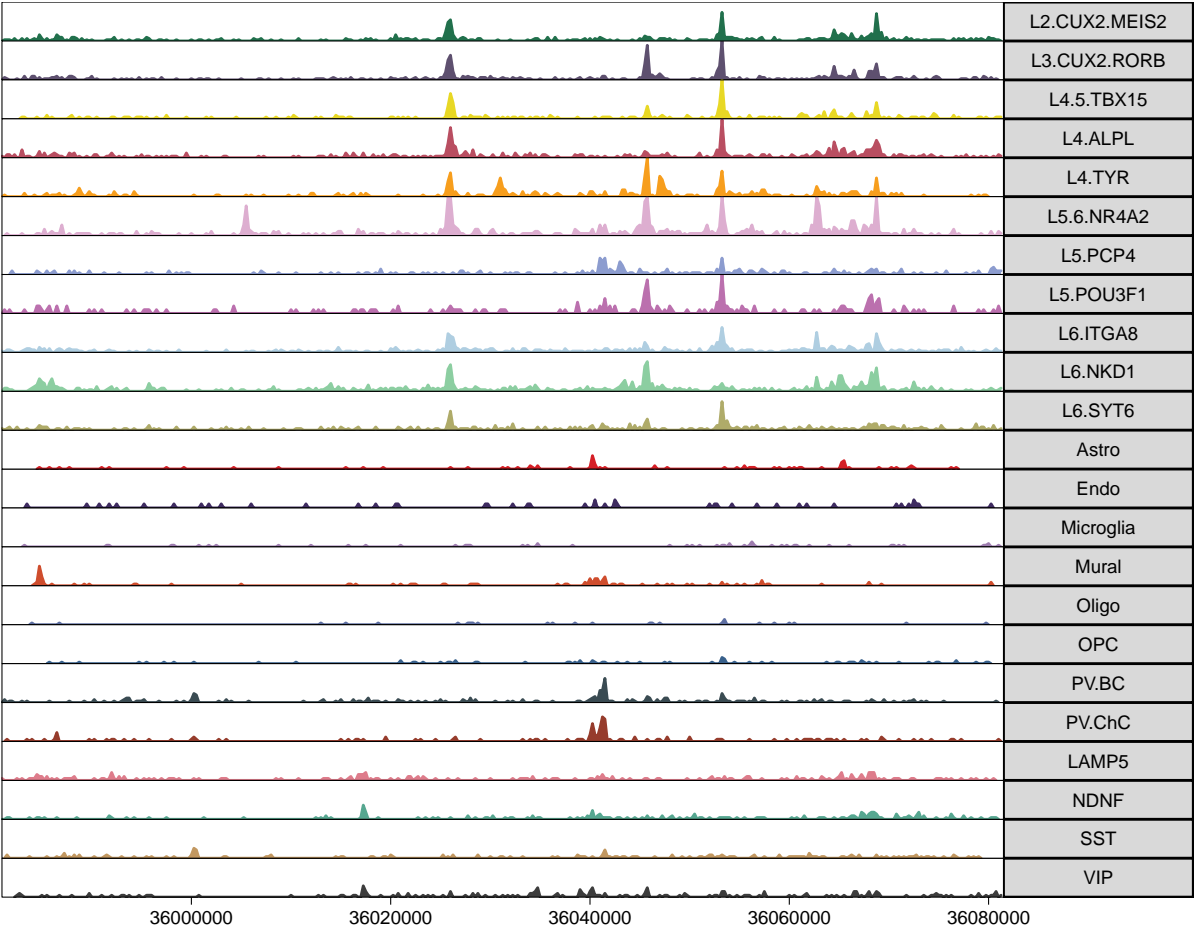
Peaks



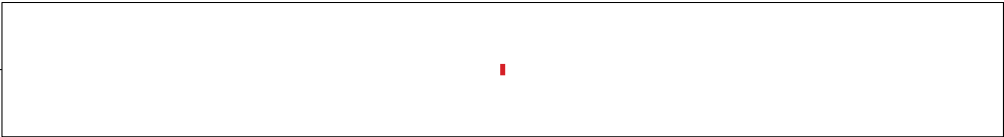
Genes

chr18:35980990–36081491 Marker Peak Tracks for: L3.CUX2.RORB

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



L3.CUX2.RORB

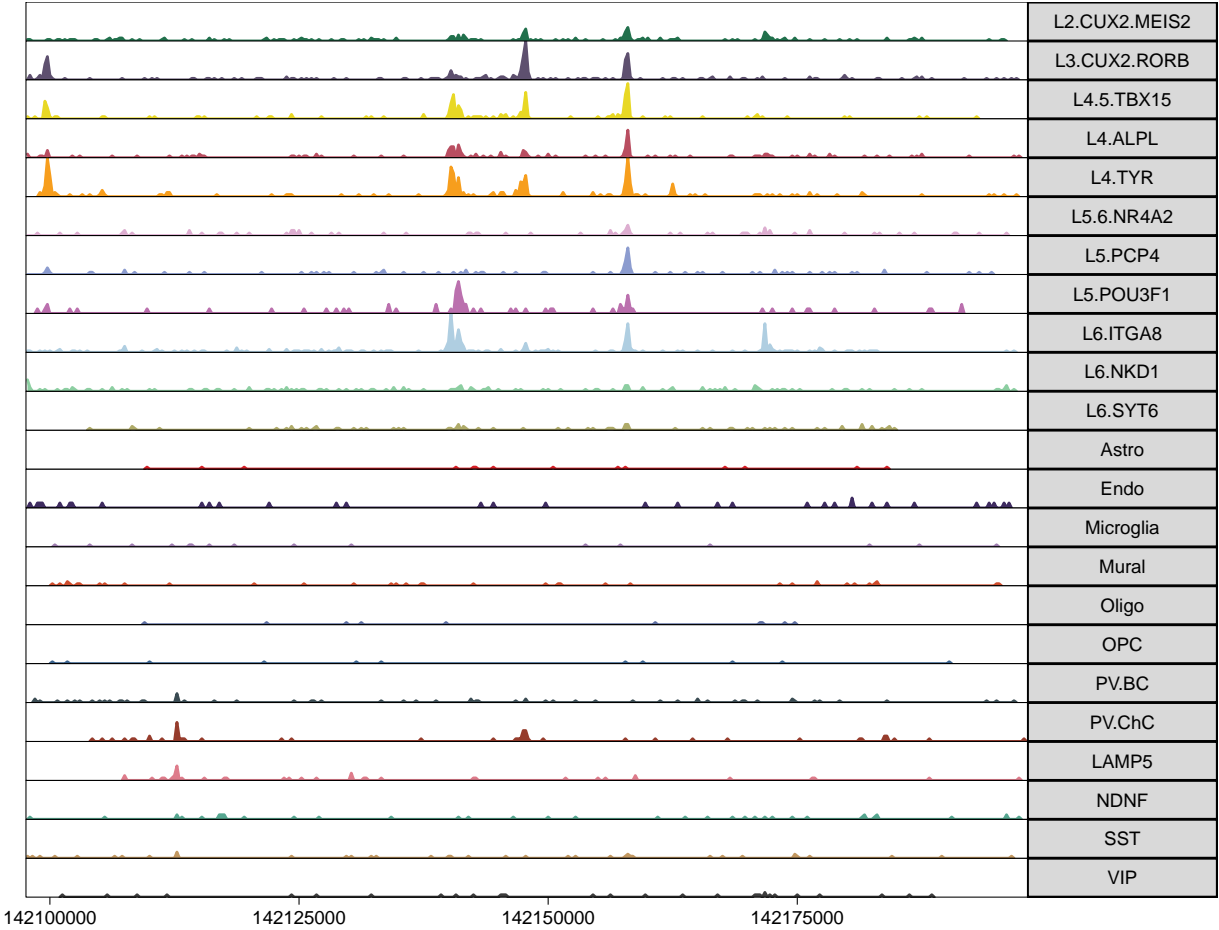


Peaks

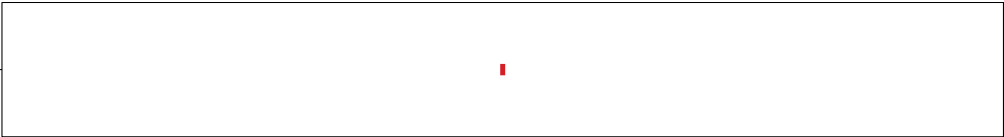


Genes

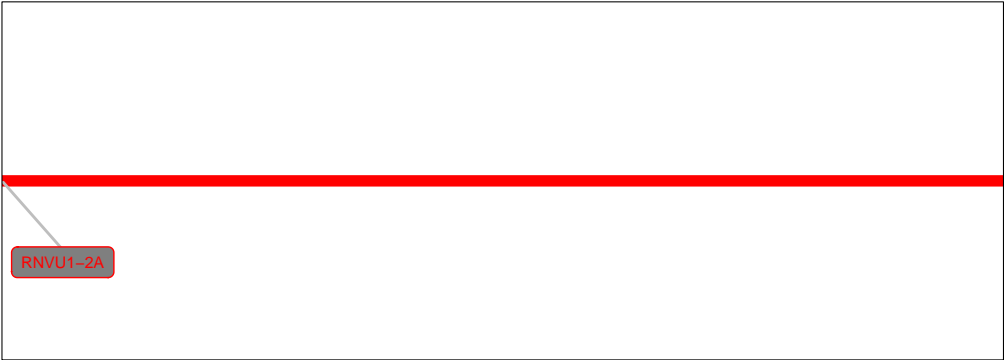
Coverage
(Norm. ATAC Signal Range (0–0.63) by ReadsInTSS)



L3.CUX2.RORB

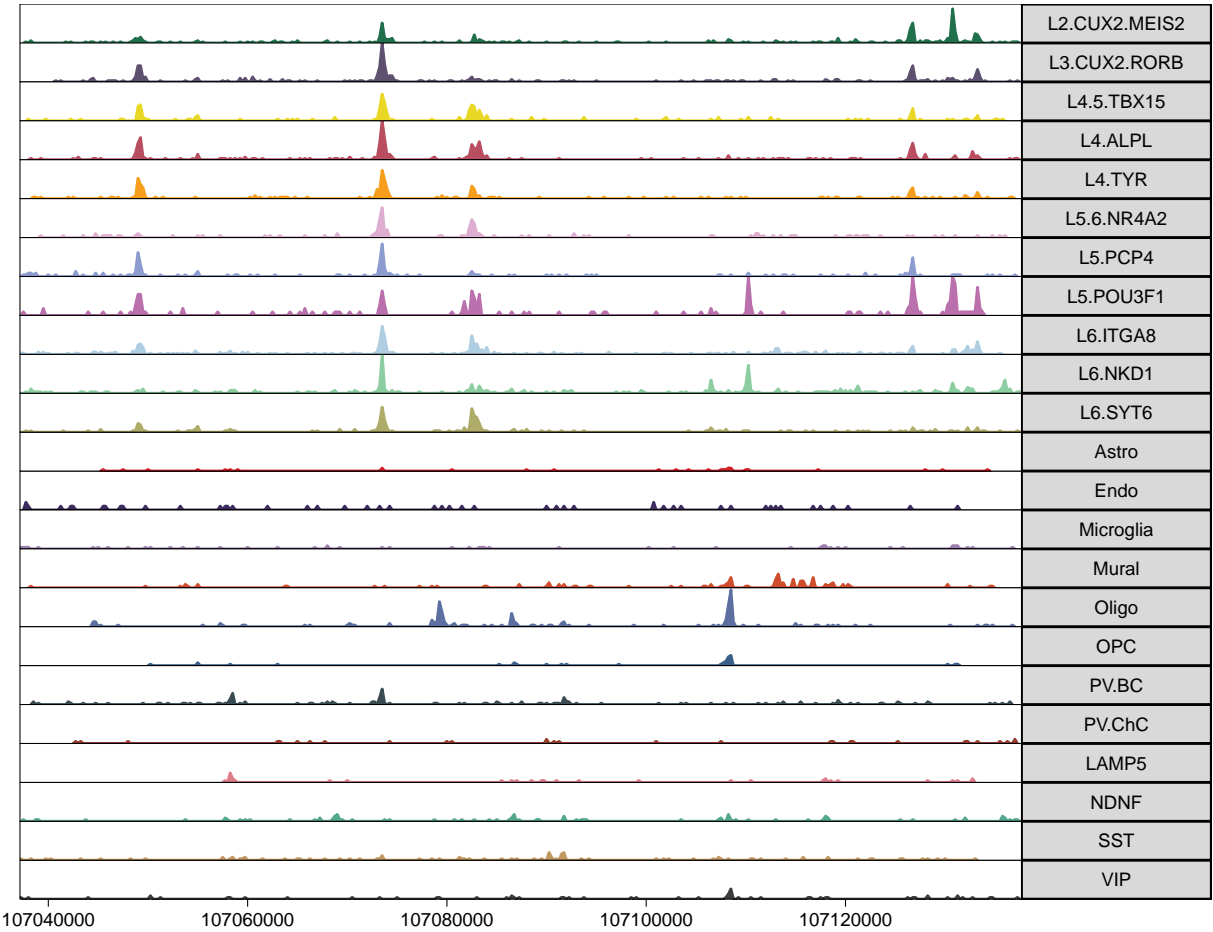


Peaks

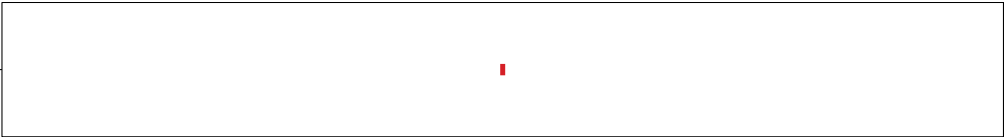


Genes

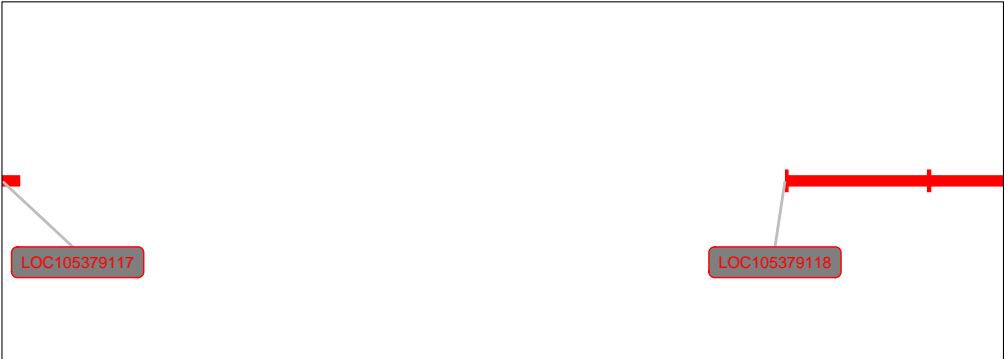
Coverage
(Norm. ATAC Signal Range (0–0.81) by ReadsInTSS)



L3.CUX2.RORB

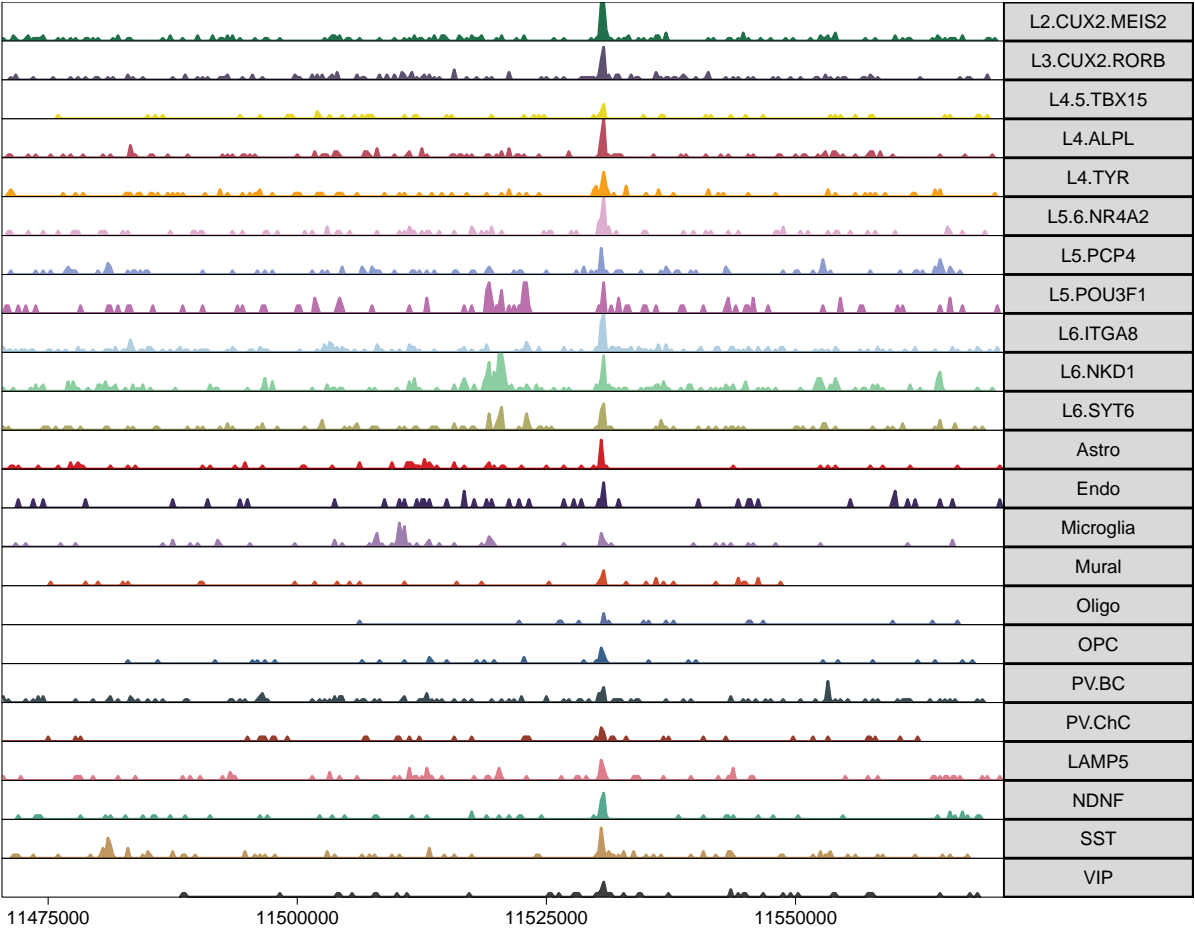


Peaks

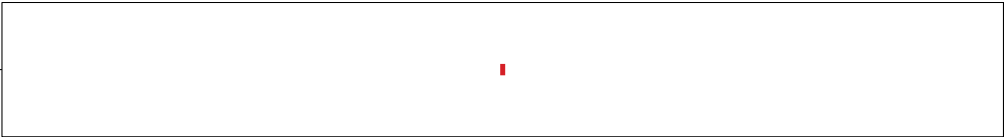


Genes

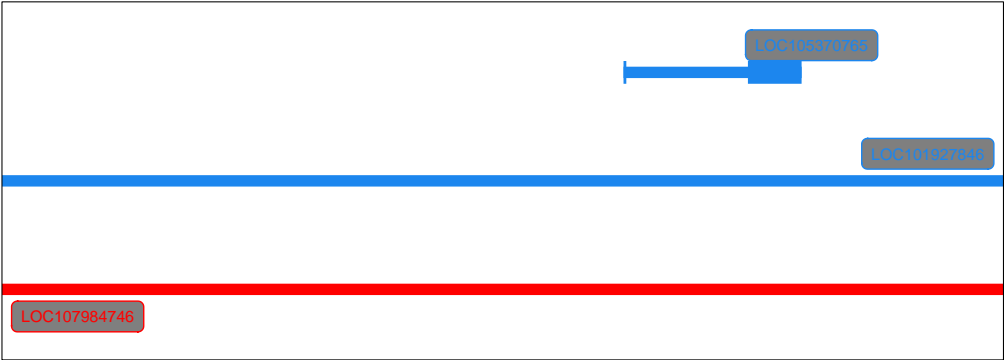
Coverage
(Norm. ATAC Signal Range (0–0.37) by ReadsInTSS)



L3.CUX2.RORB



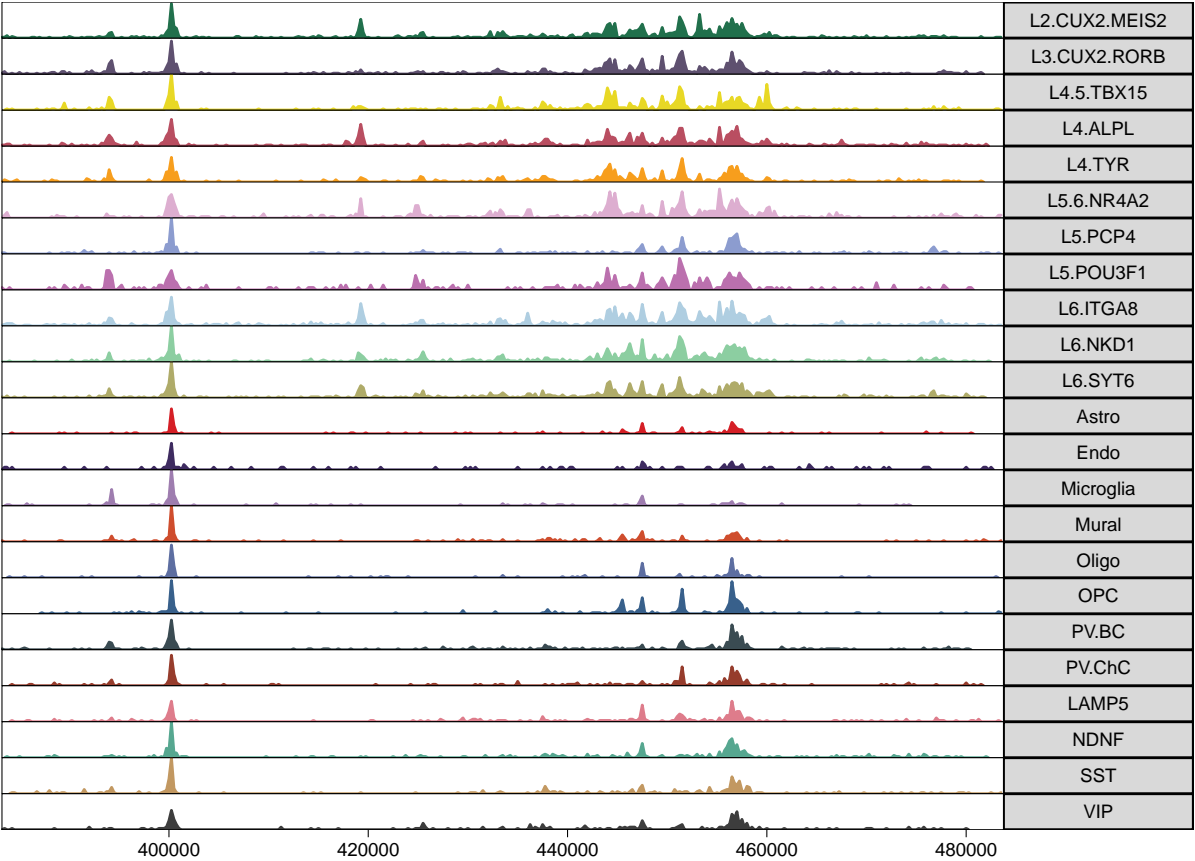
Peaks



Genes

chr17:383242–483743 Marker Peak Tracks for: L3.CUX2.RORB

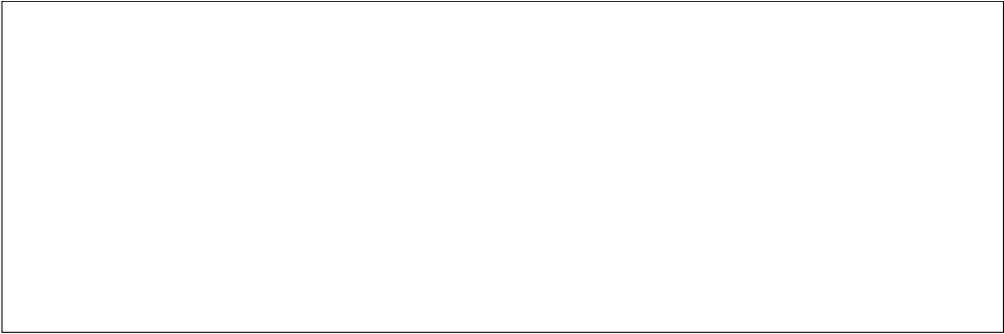
Coverage
(Norm. ATAC Signal Range (0–1.09) by ReadsInTSS)



L3.CUX2.RORB



Peaks

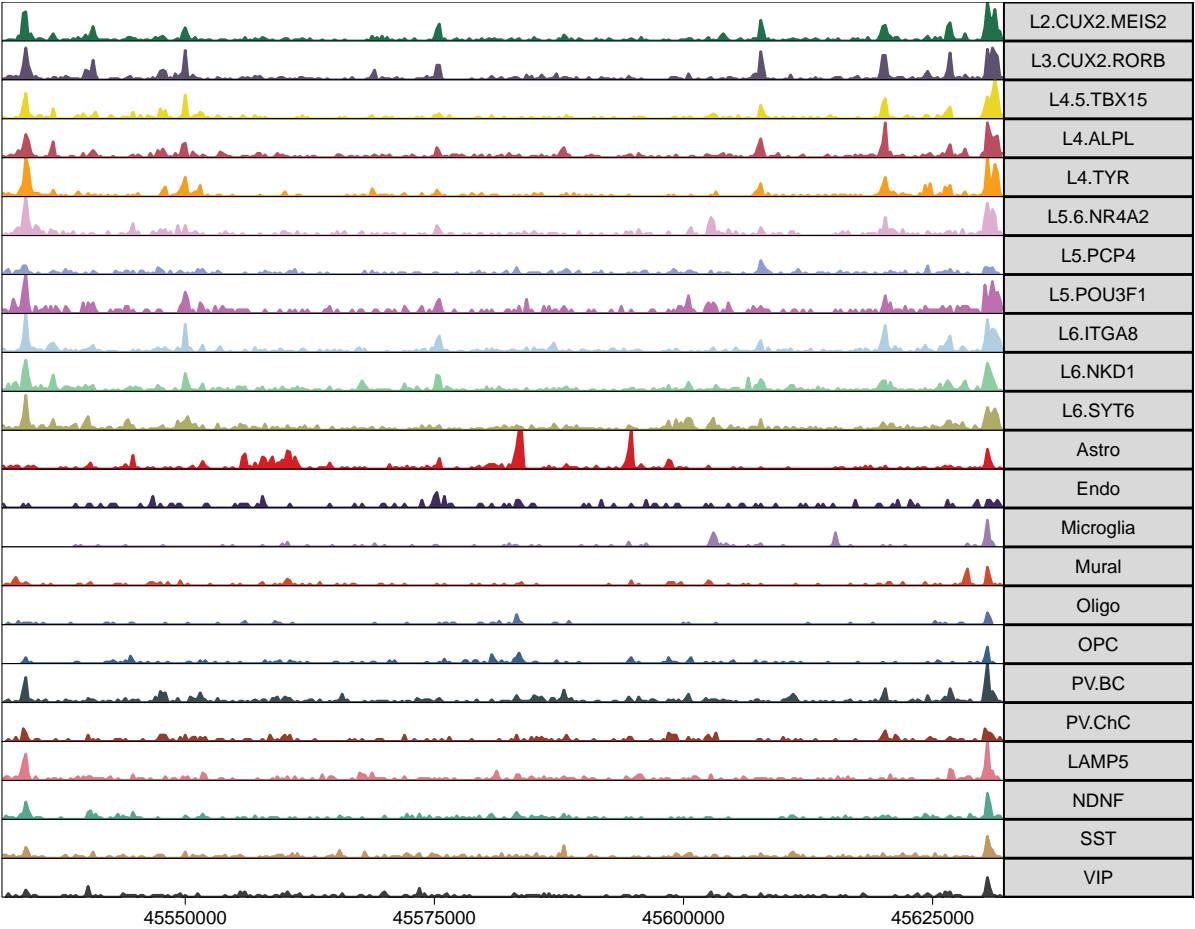


GeneTrack

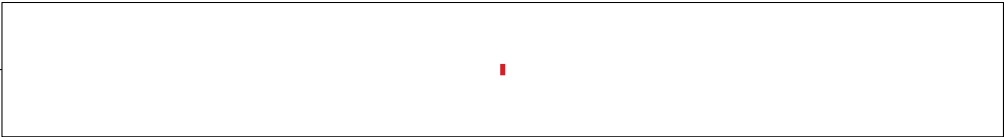
strand ● ·

chr14:45531605–45632106 Marker Peak Tracks for: L3.CUX2.RORB

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



L3.CUX2.RORB

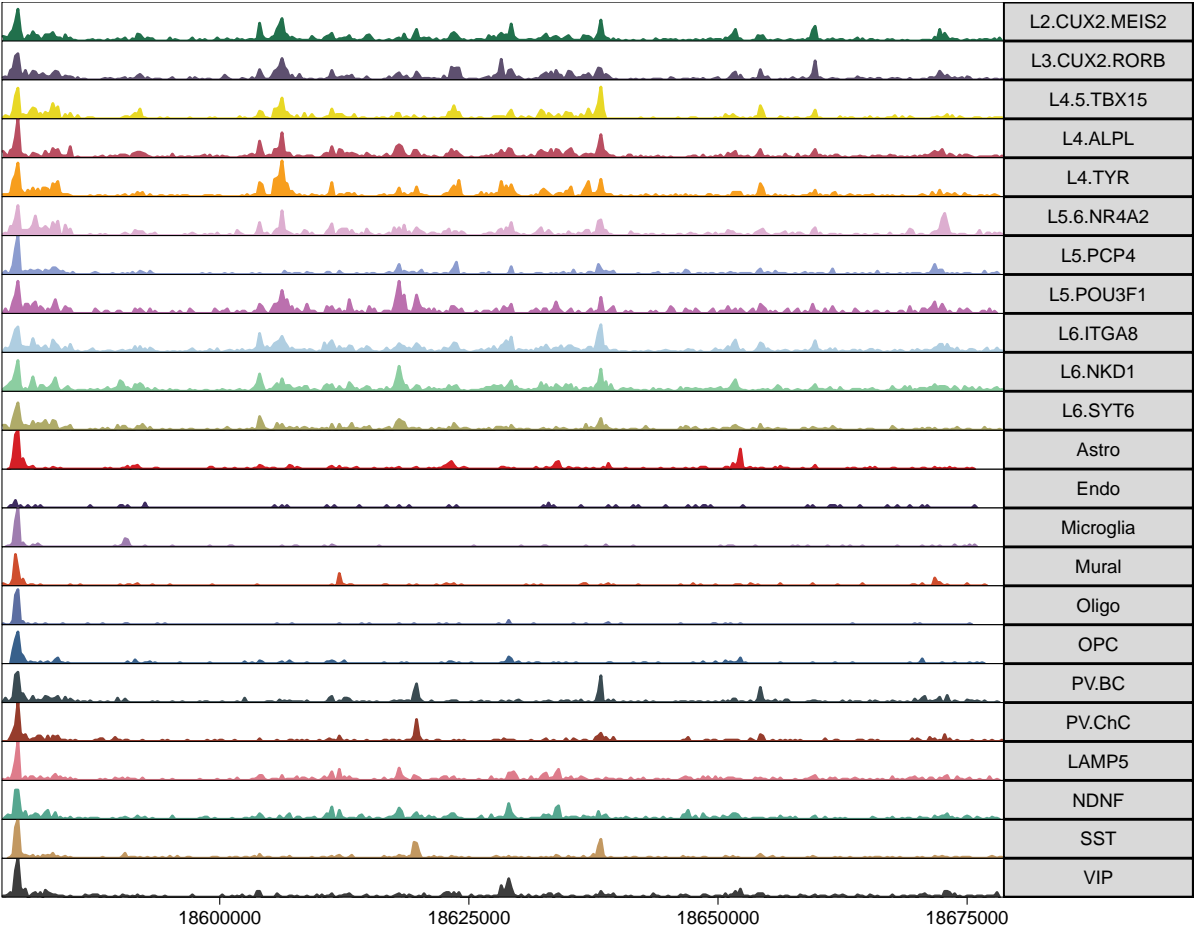


Peaks

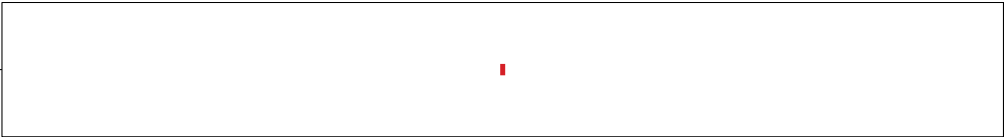


Genes

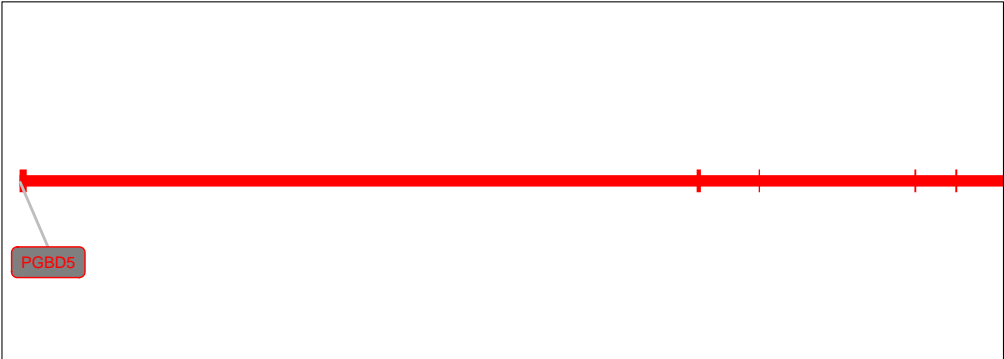
Coverage
(Norm. ATAC Signal Range (0–1.24) by ReadsInTSS)



L3.CUX2.RORB

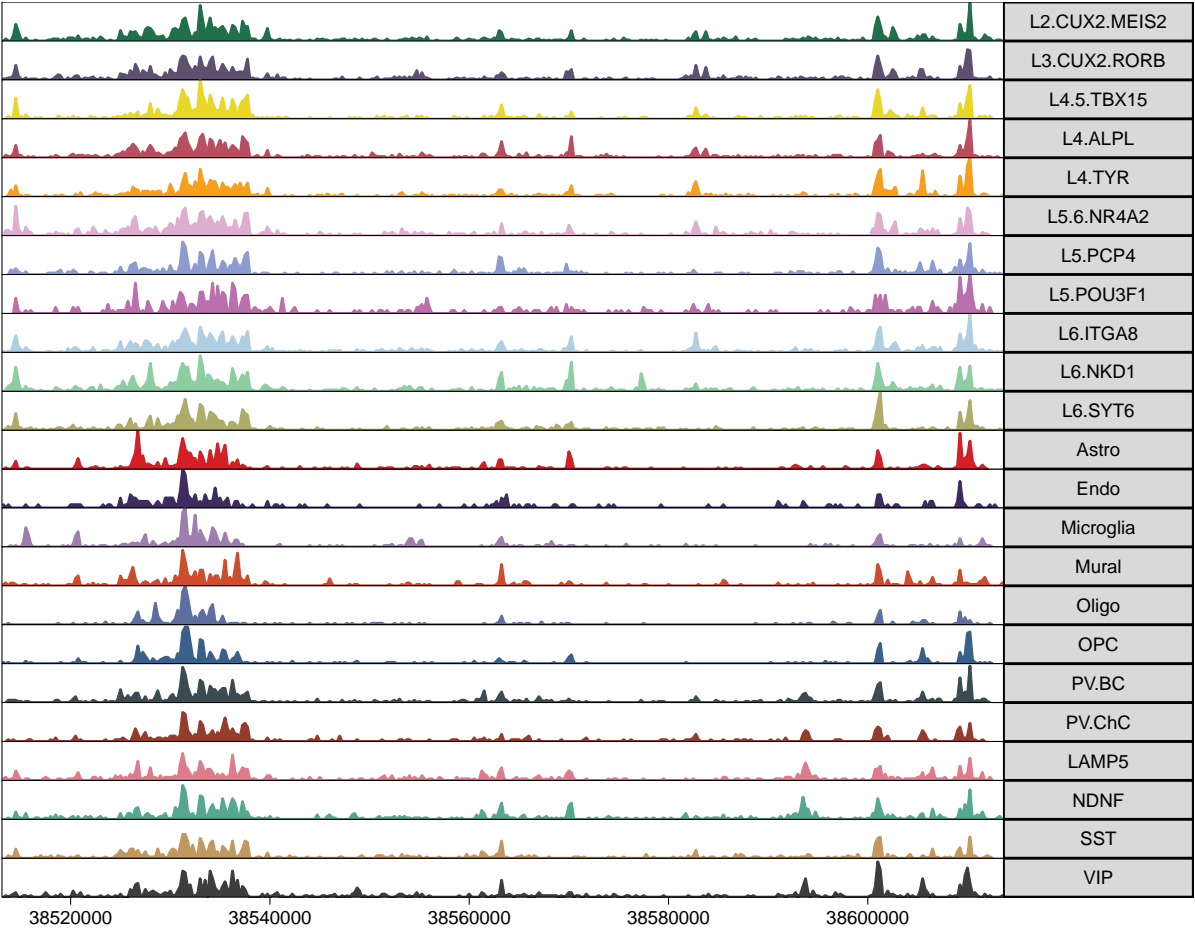


Peaks

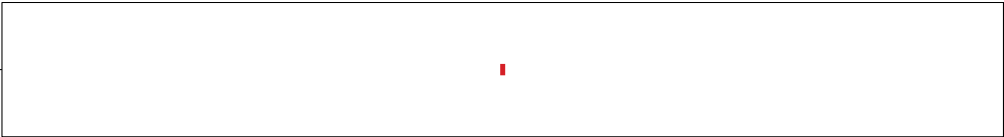


Genes

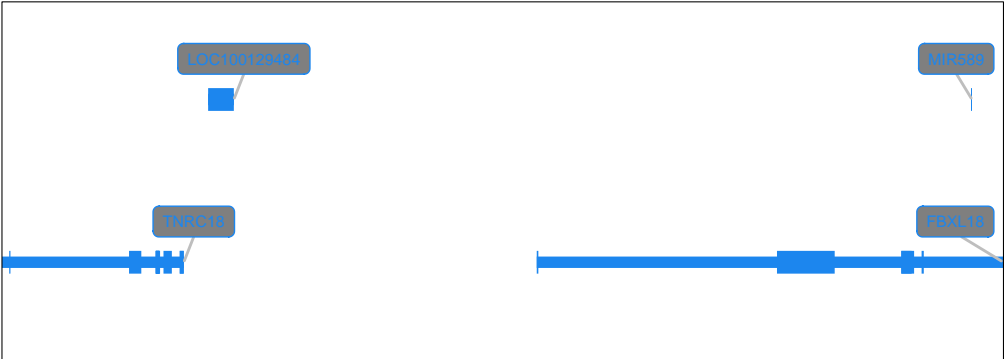
Coverage
(Norm. ATAC Signal Range (0–0.94) by ReadsInTSS)



L3.CUX2.RORB



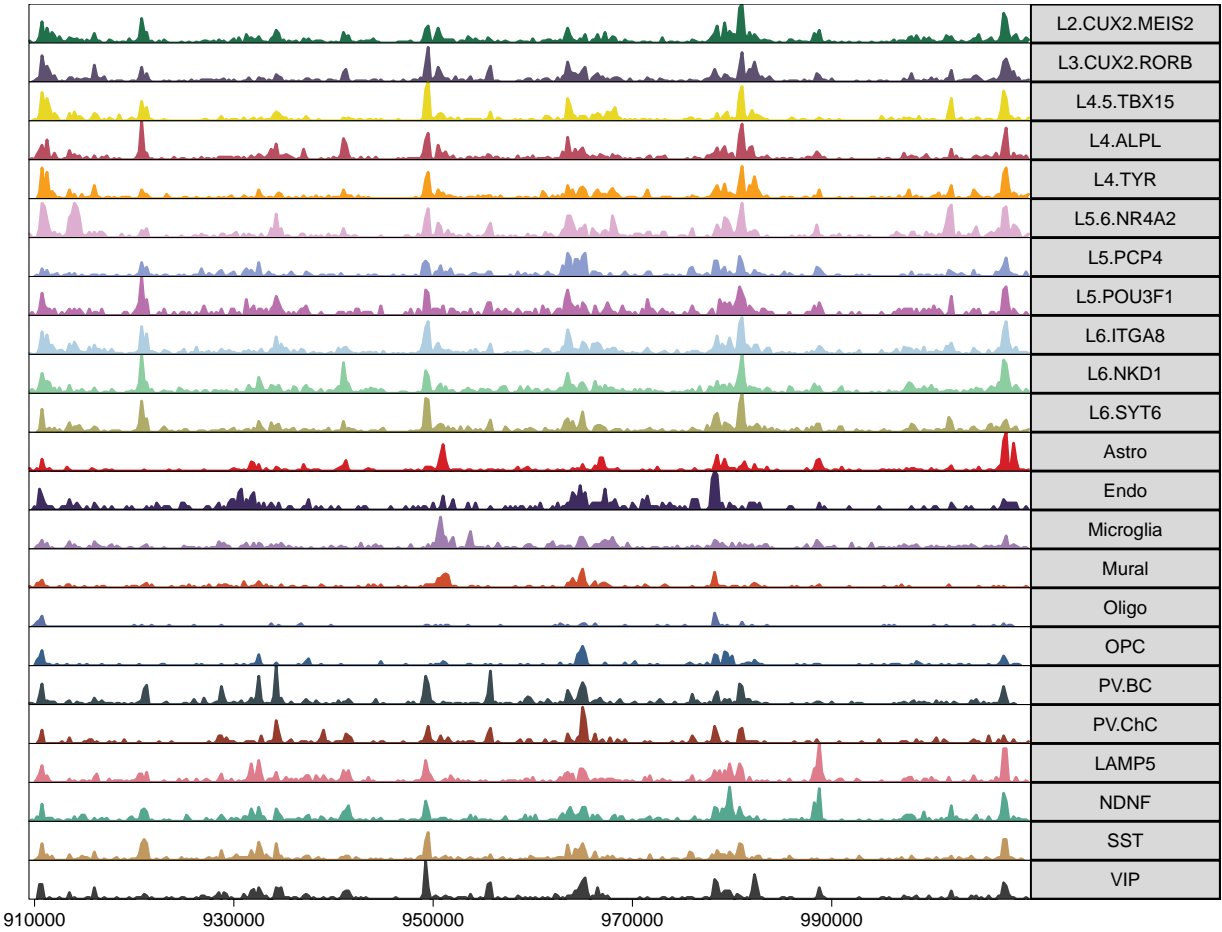
Peaks



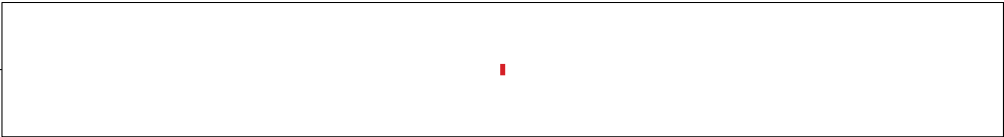
Genes

chr16:909439–1009940 Marker Peak Tracks for: L3.CUX2.RORB

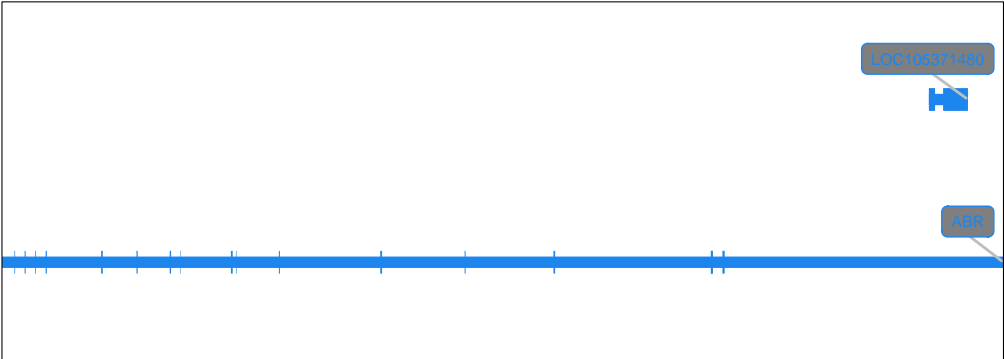
Coverage
(Norm. ATAC Signal Range (0–0.9) by ReadsInTSS)



L3.CUX2.RORB



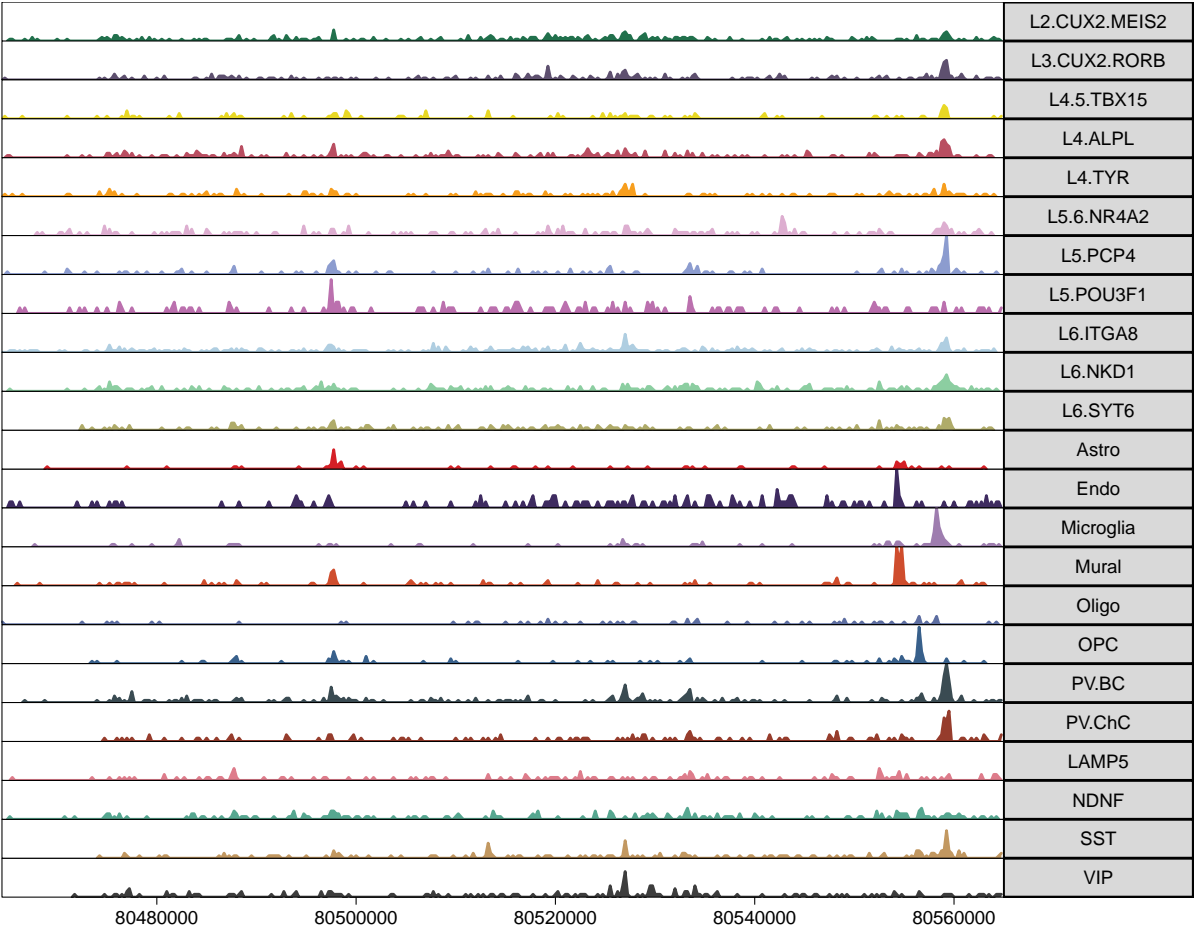
Peaks



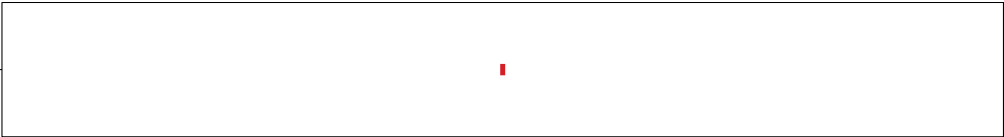
Genes

chr17:80464460–80564961 Marker Peak Tracks for: L3.CUX2.RORB

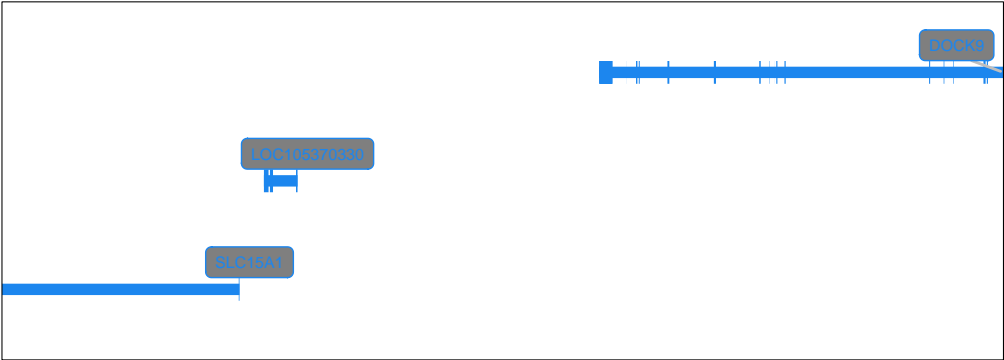
Coverage
(Norm. ATAC Signal Range (0–0.51) by ReadsInTSS)



L3.CUX2.RORB

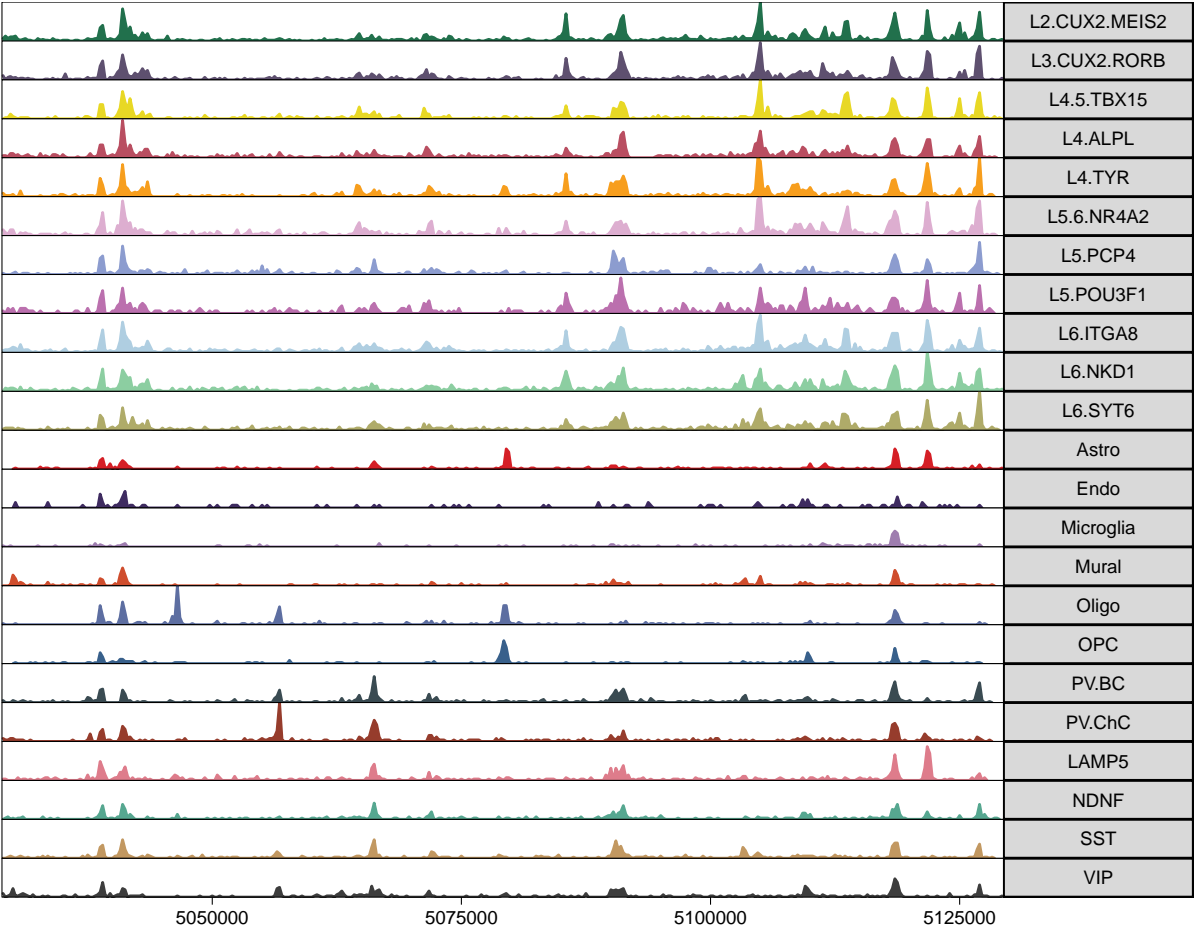


Peaks

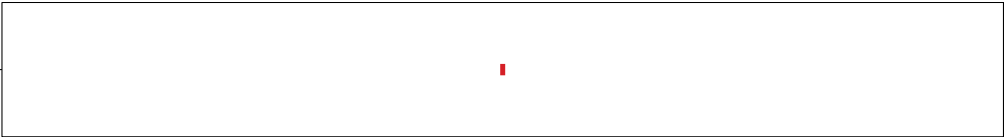


Genes

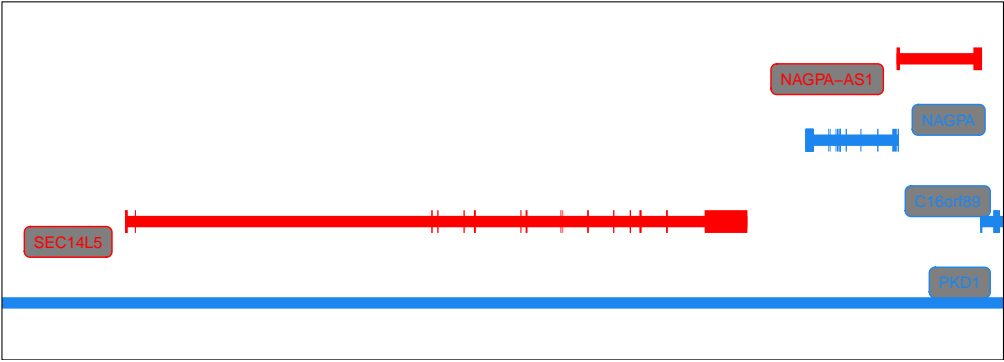
Coverage
(Norm. ATAC Signal Range (0–1.13) by ReadsInTSS)



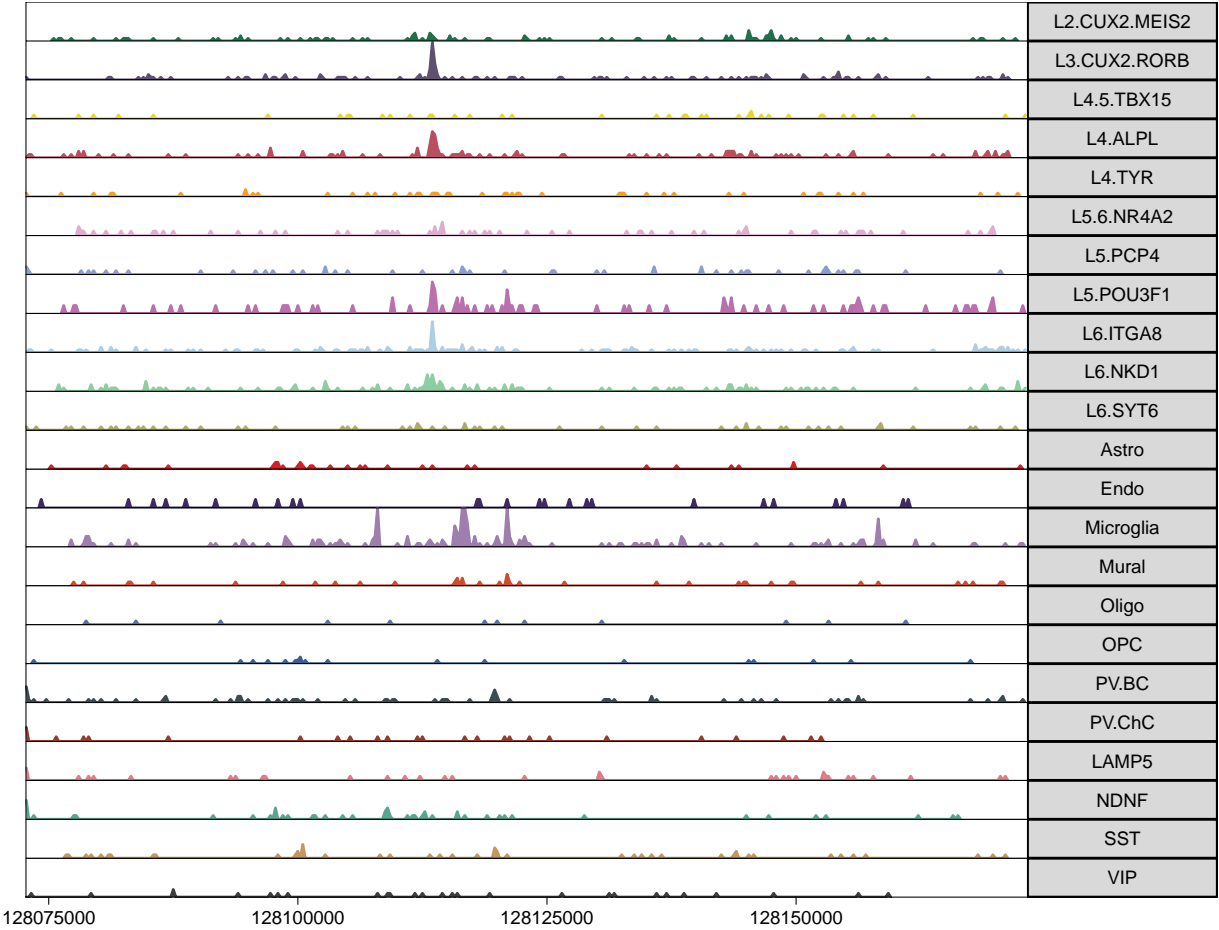
L3.CUX2.RORB



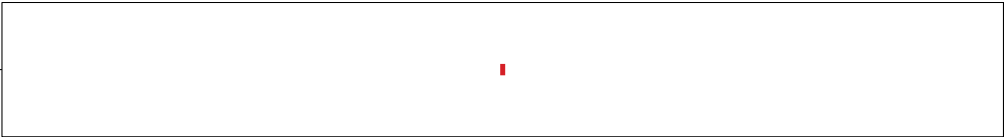
Peaks



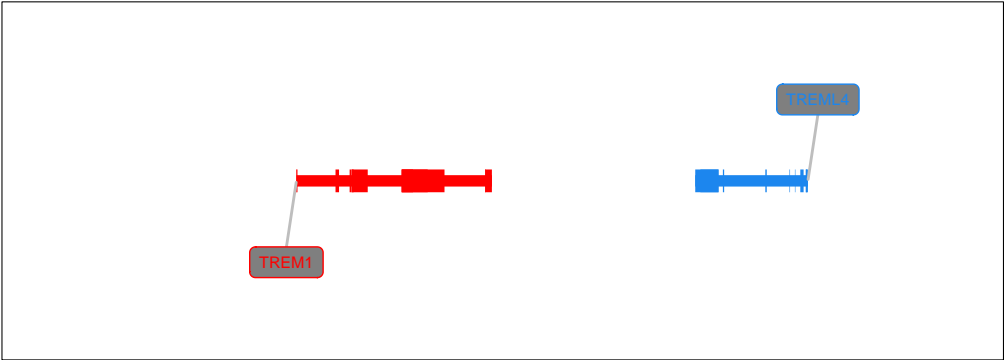
Coverage
(Norm. ATAC Signal Range (0–0.36) by ReadsInTSS)



L3.CUX2.RORB

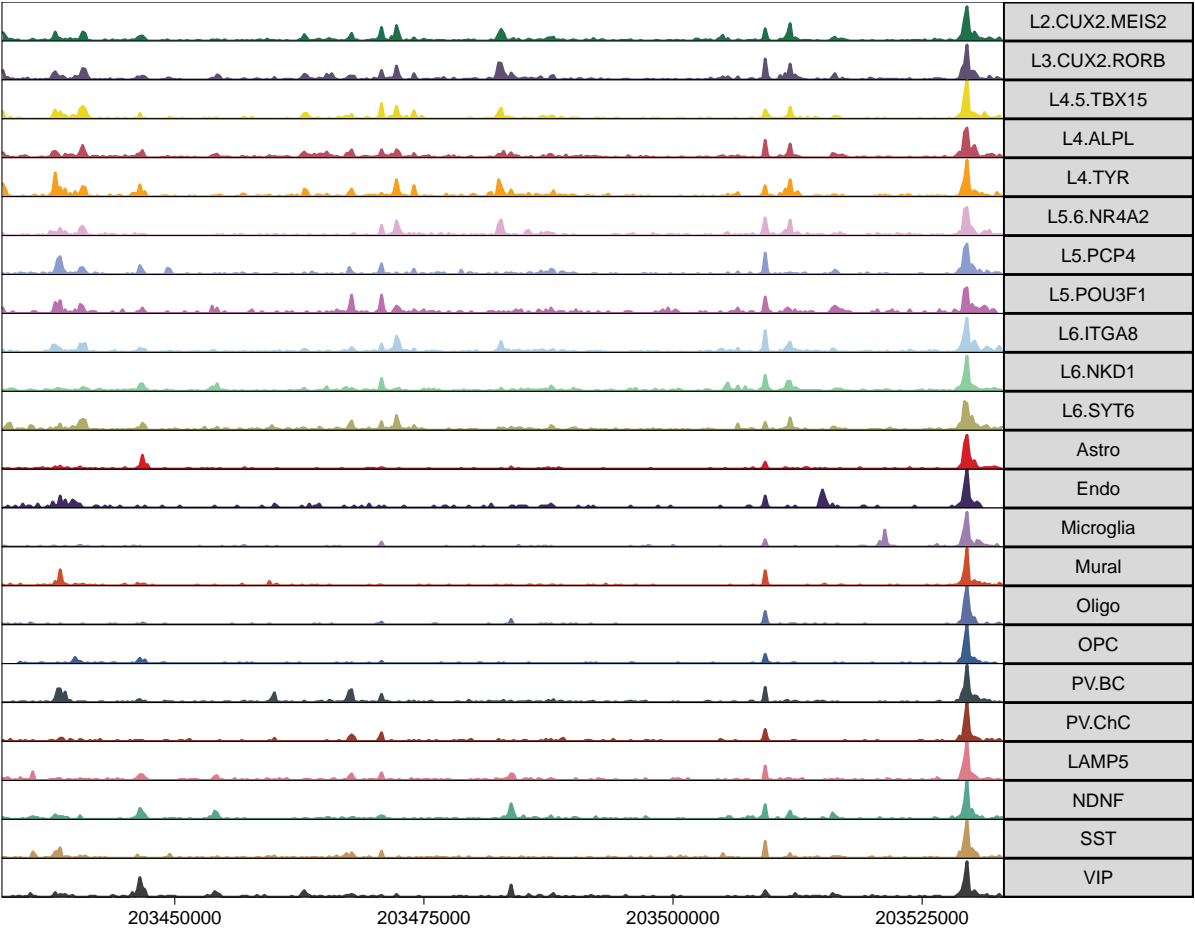


Peaks

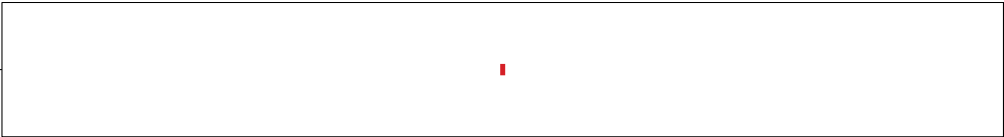


Genes

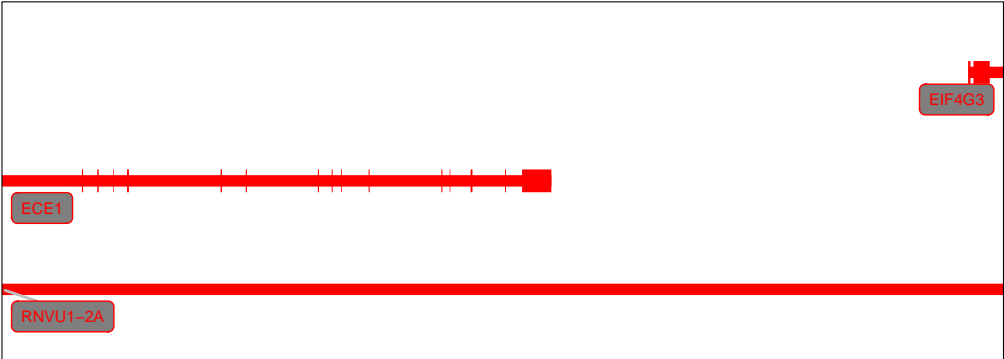
Coverage
(Norm. ATAC Signal Range (0–1.56) by ReadsInTSS)



L3.CUX2.RORB



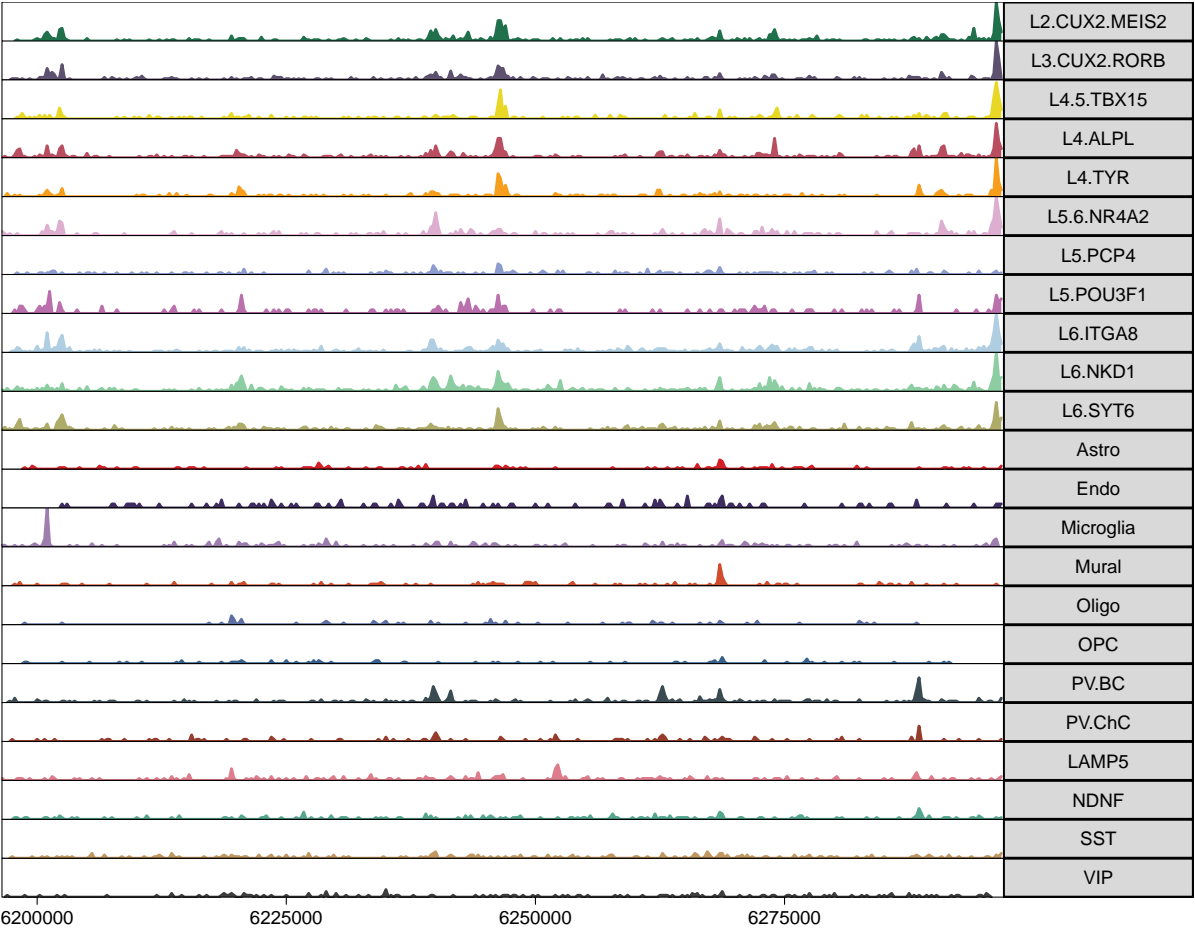
Peaks



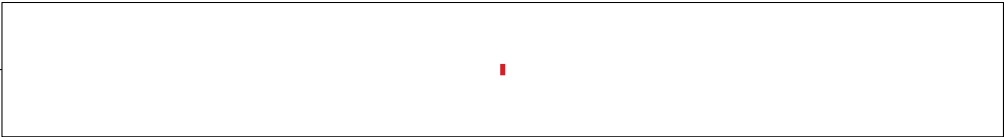
Genes

chr14:6196467–6296968 Marker Peak Tracks for: L3.CUX2.RORB

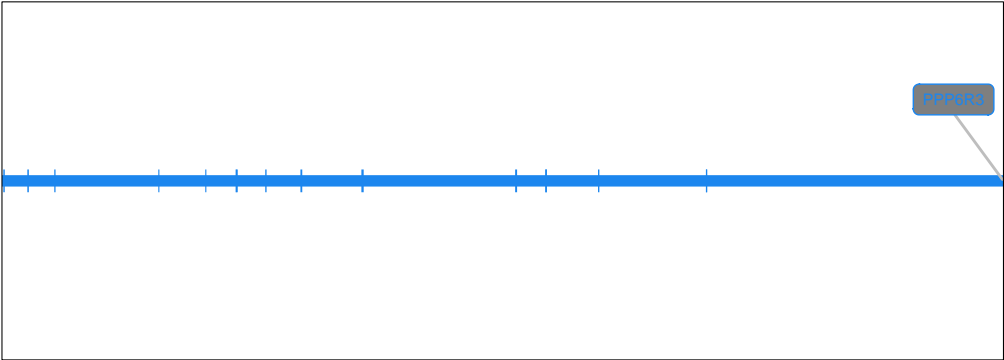
Coverage
(Norm. ATAC Signal Range (0–0.78) by ReadsInTSS)



L3.CUX2.RORB

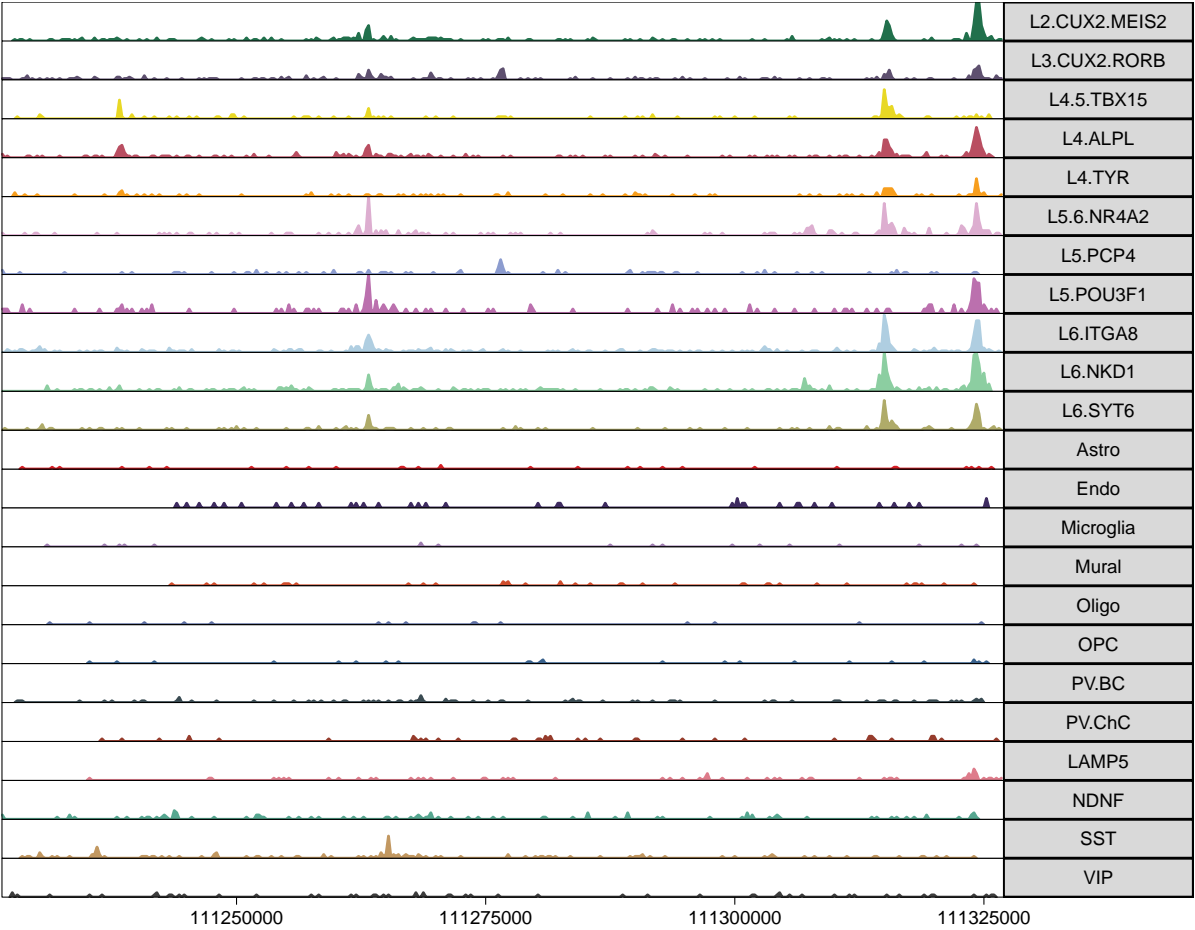


Peaks

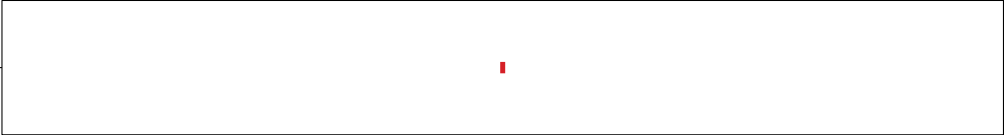


Genes

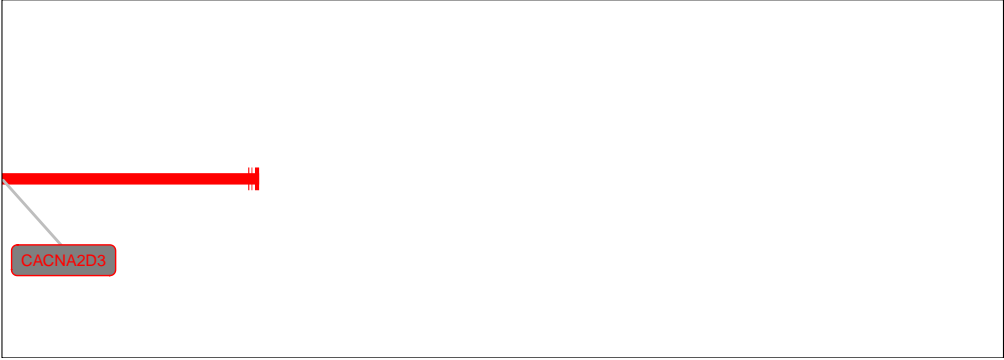
Coverage
(Norm. ATAC Signal Range (0–0.66) by ReadsInTSS)



L3.CUX2.RORB



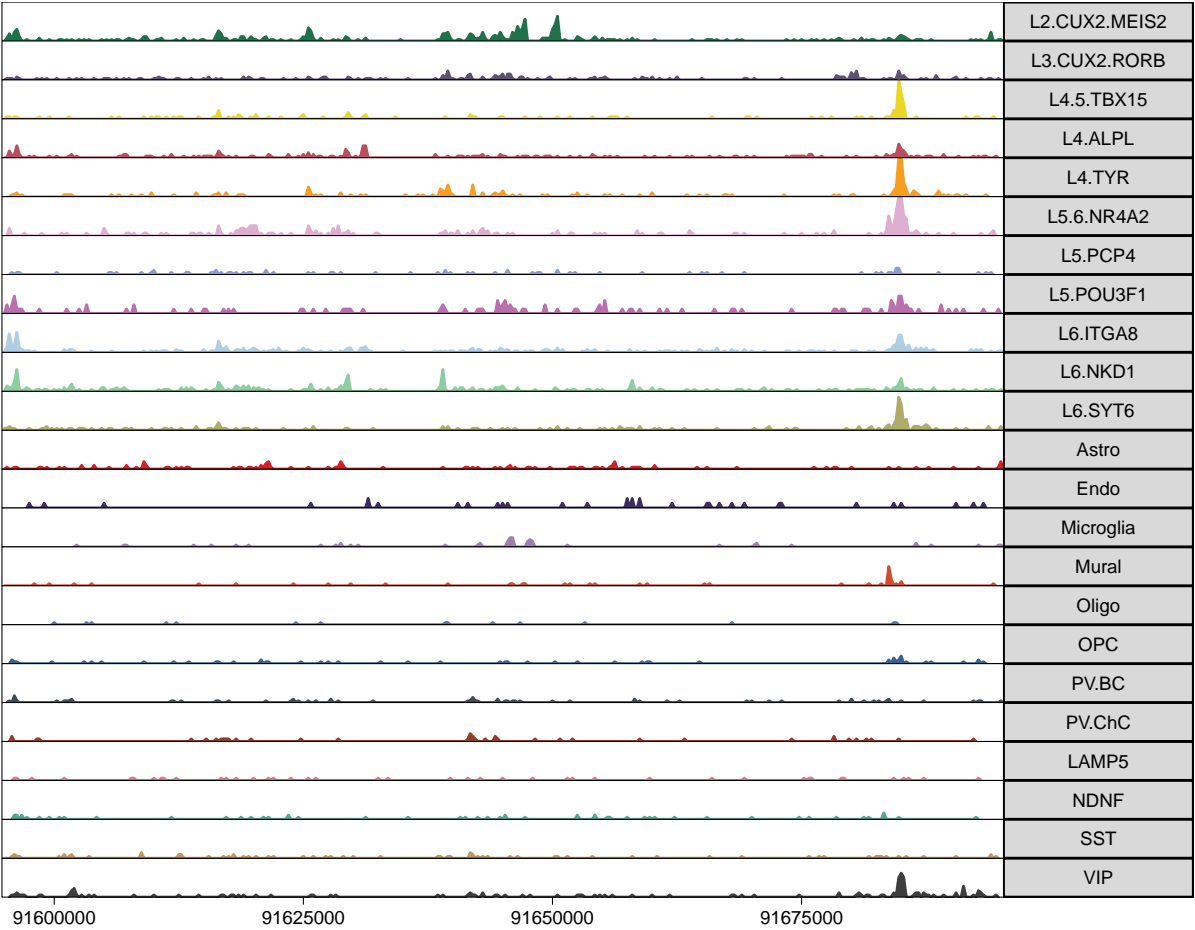
Peaks



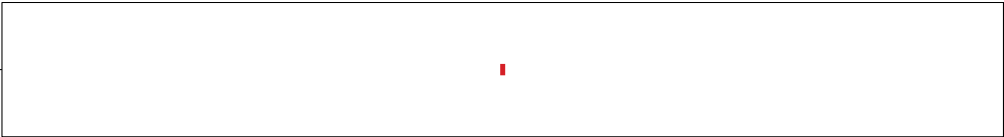
Genes

chr15:91594754–91695255 Marker Peak Tracks for: L3.CUX2.RORB

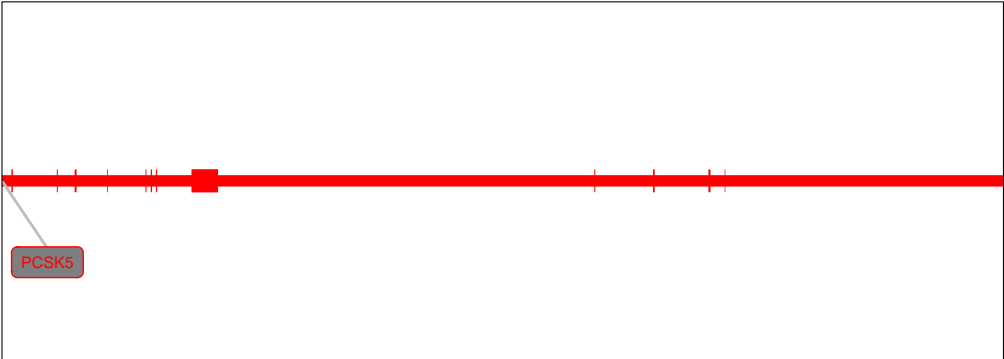
Coverage
(Norm. ATAC Signal Range (0–0.66) by ReadsInTSS)



L3.CUX2.RORB

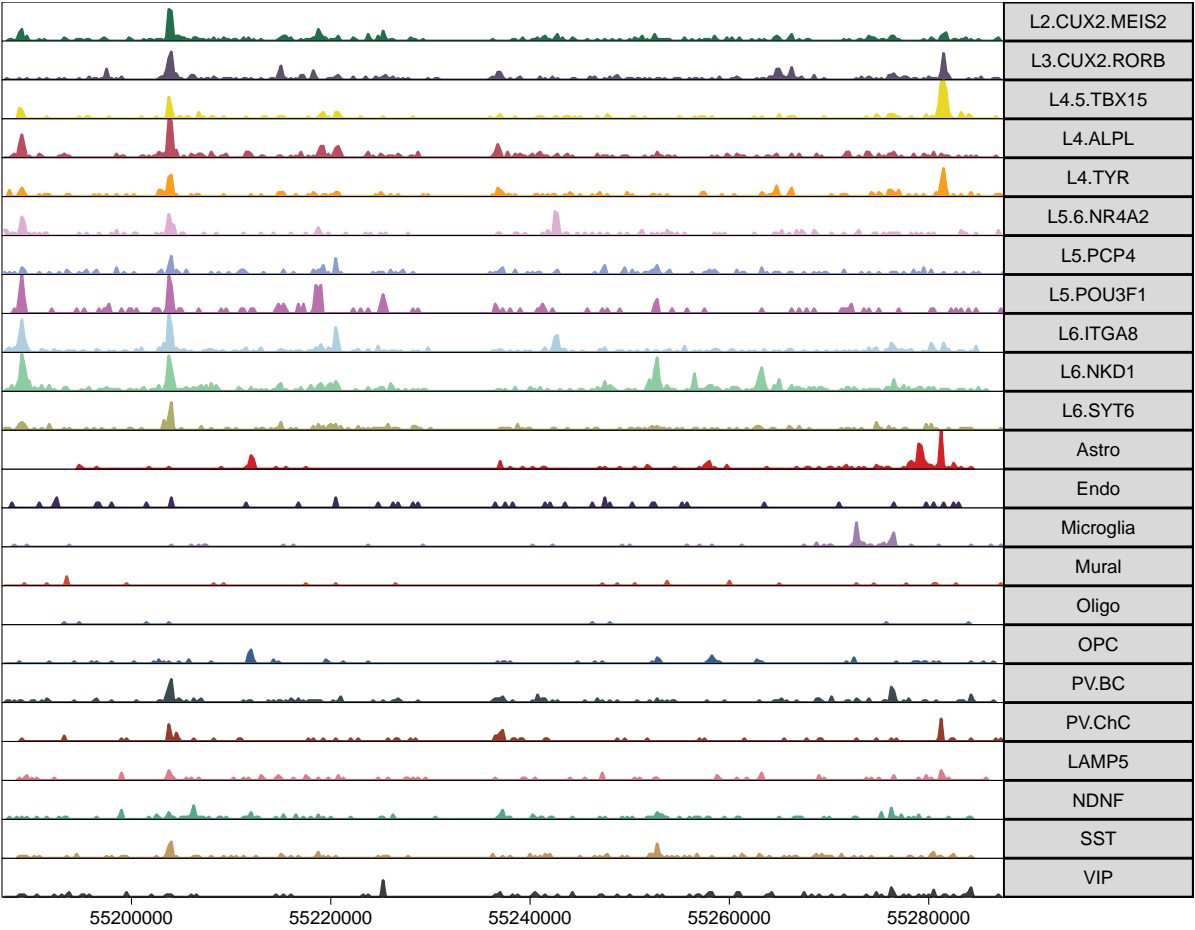


Peaks

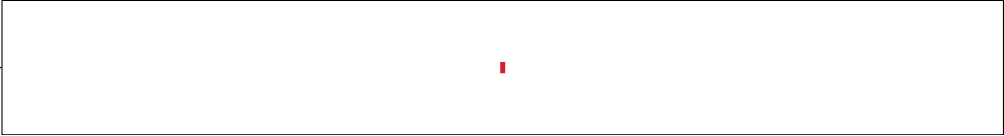


Genes

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



L3.CUX2.RORB

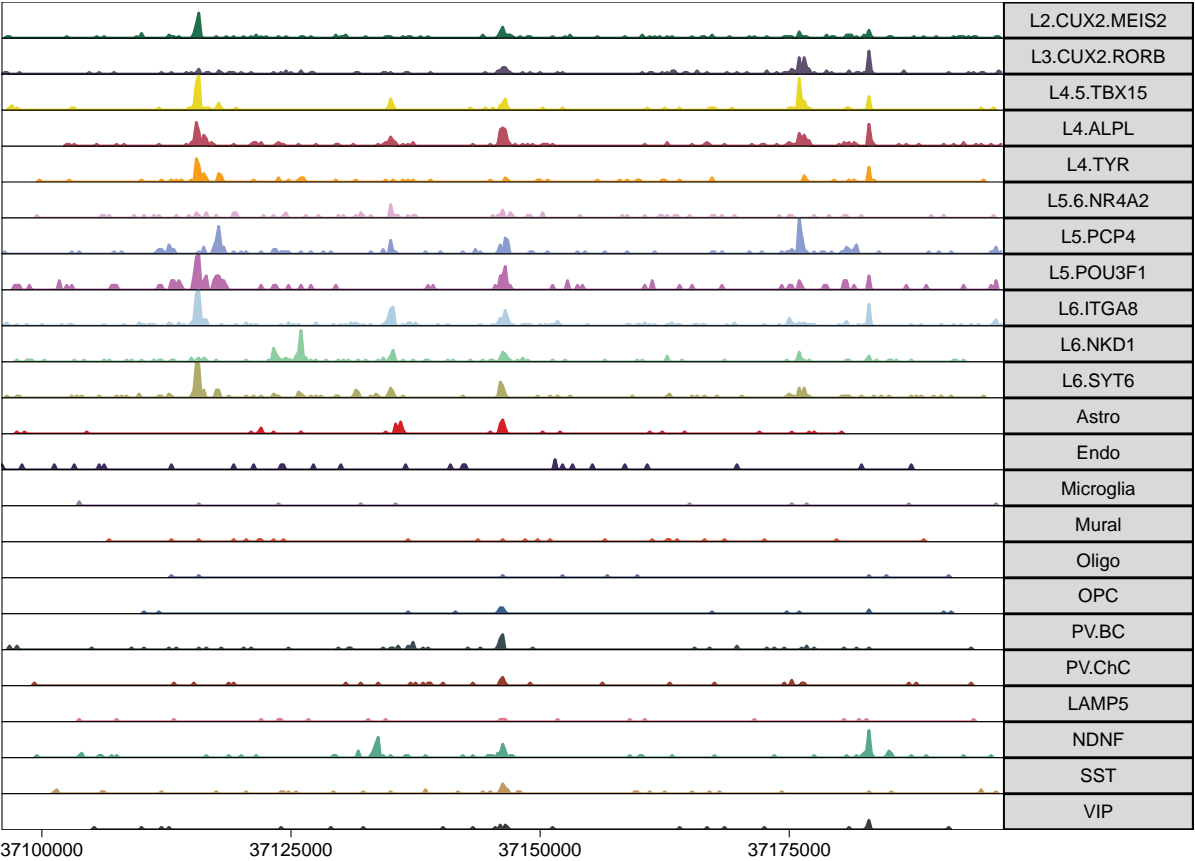


Peaks

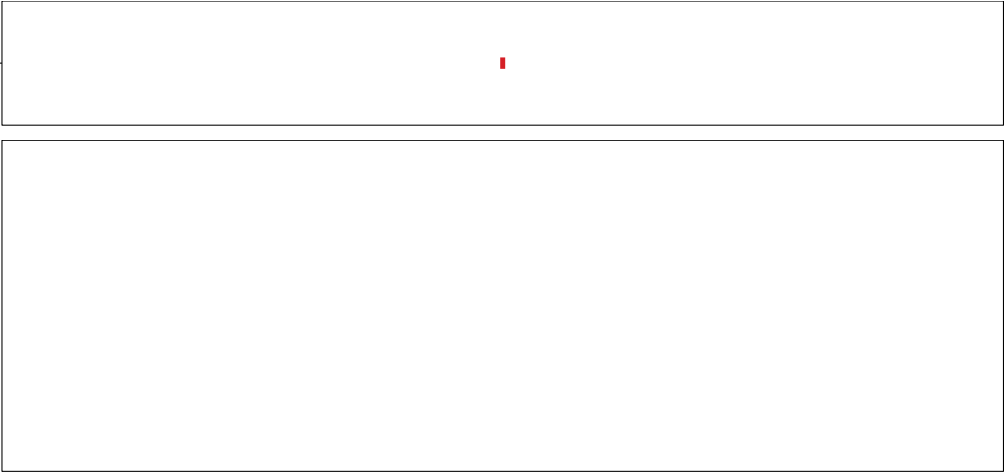


Genes

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



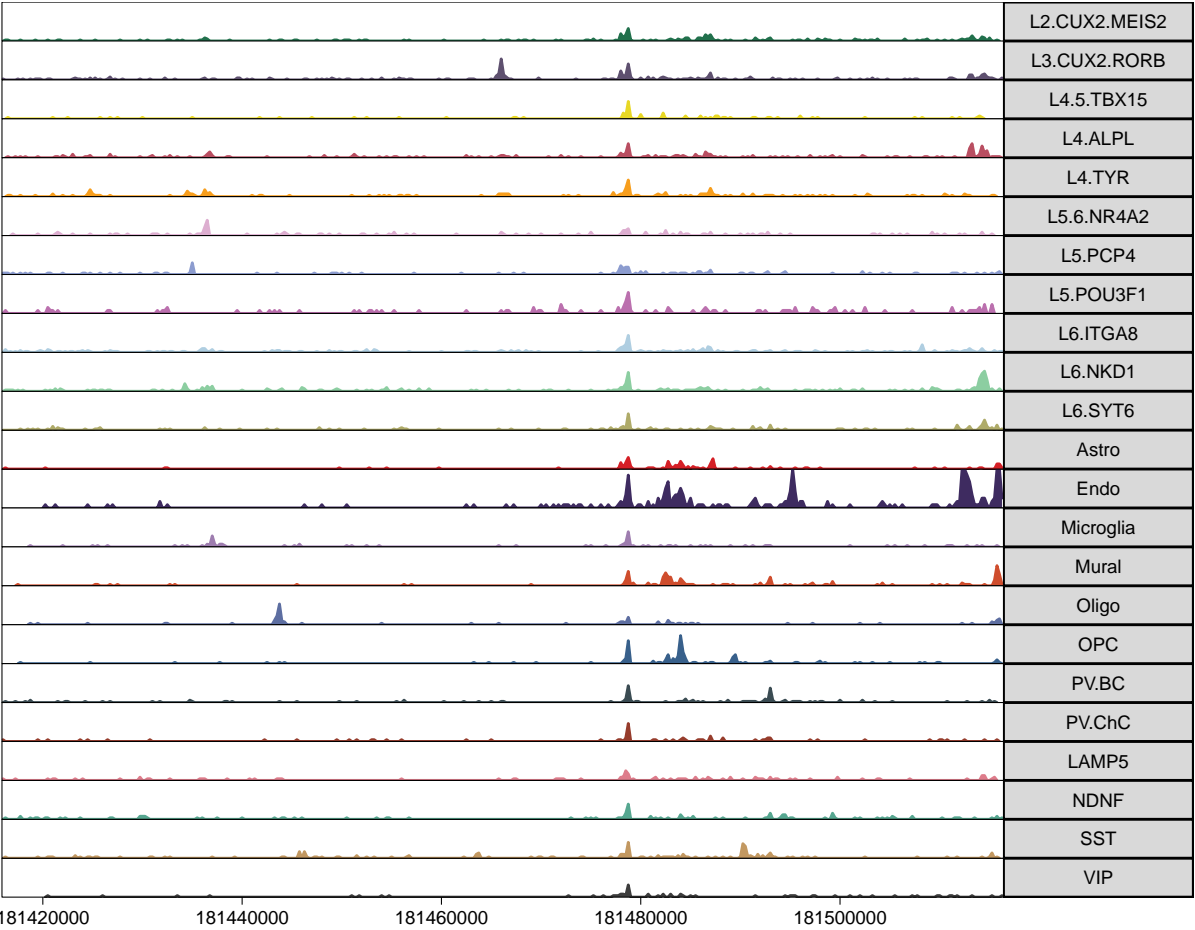
L3.CUX2.RORB



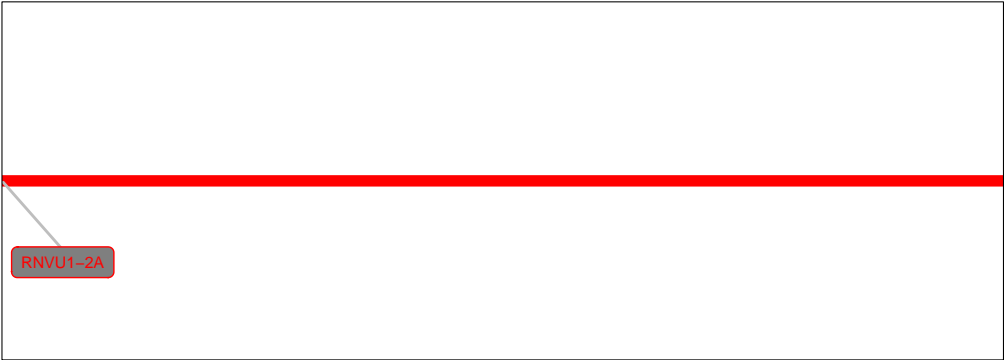
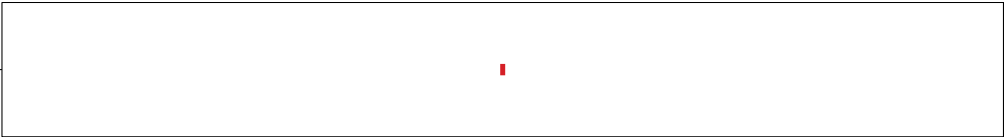
Peaks

GeneTrack

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

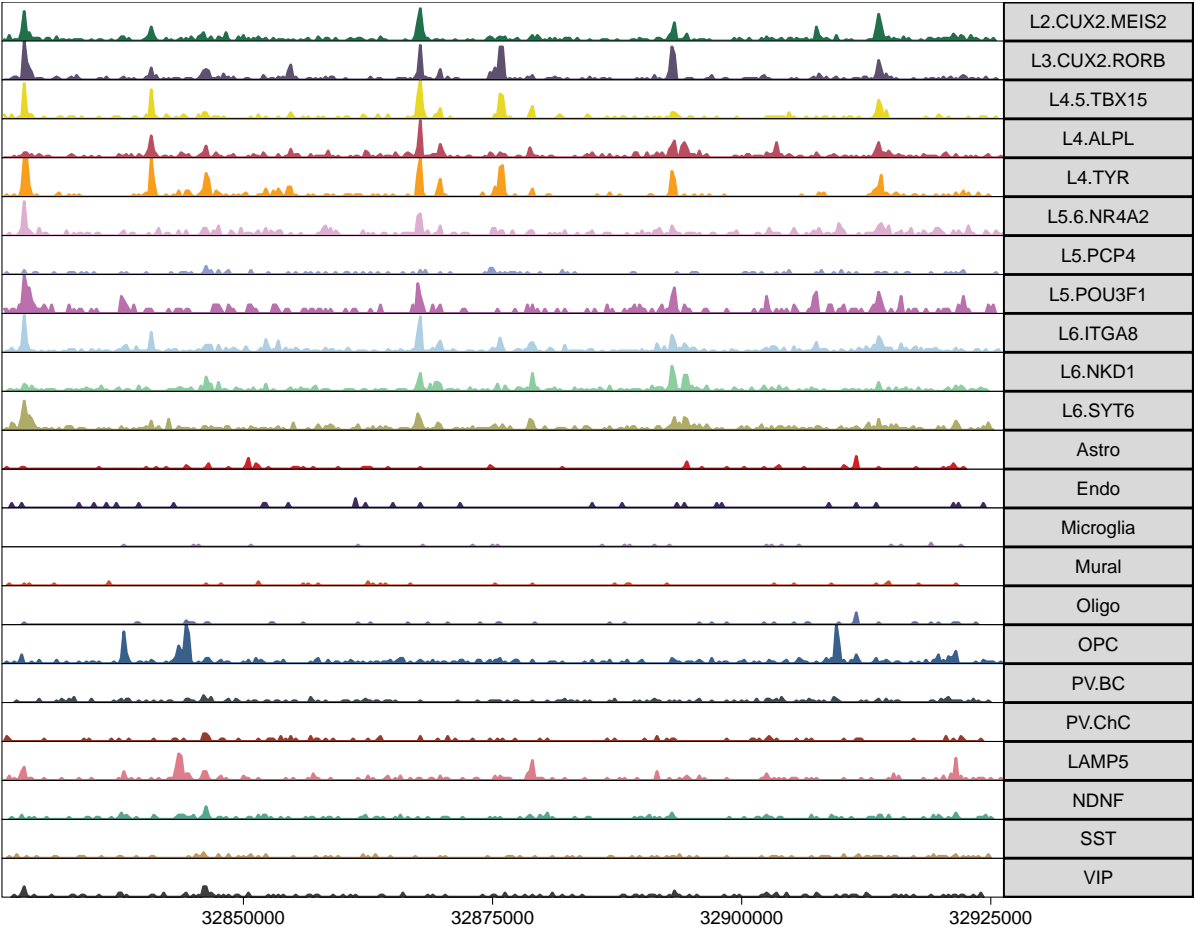


L3.CUX2.RORB

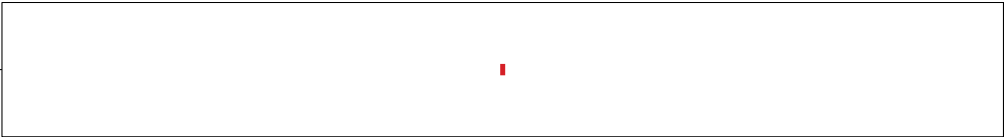


chr1:32825768–32926269 Marker Peak Tracks for: L3.CUX2.RORB

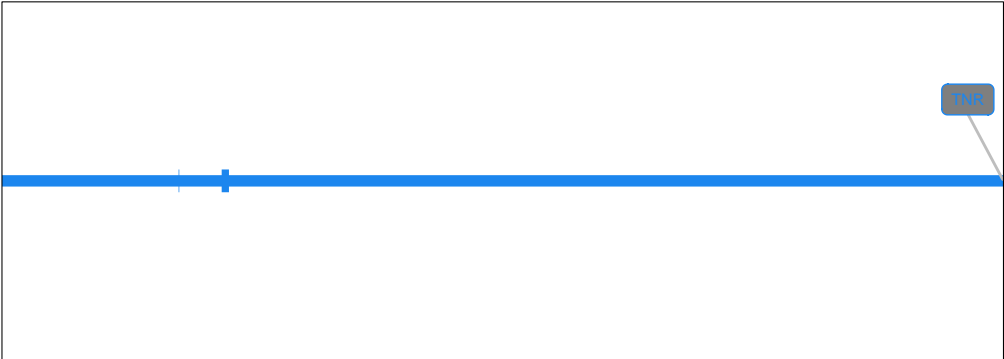
Coverage
(Norm. ATAC Signal Range (0–0.67) by ReadsInTSS)



L3.CUX2.RORB



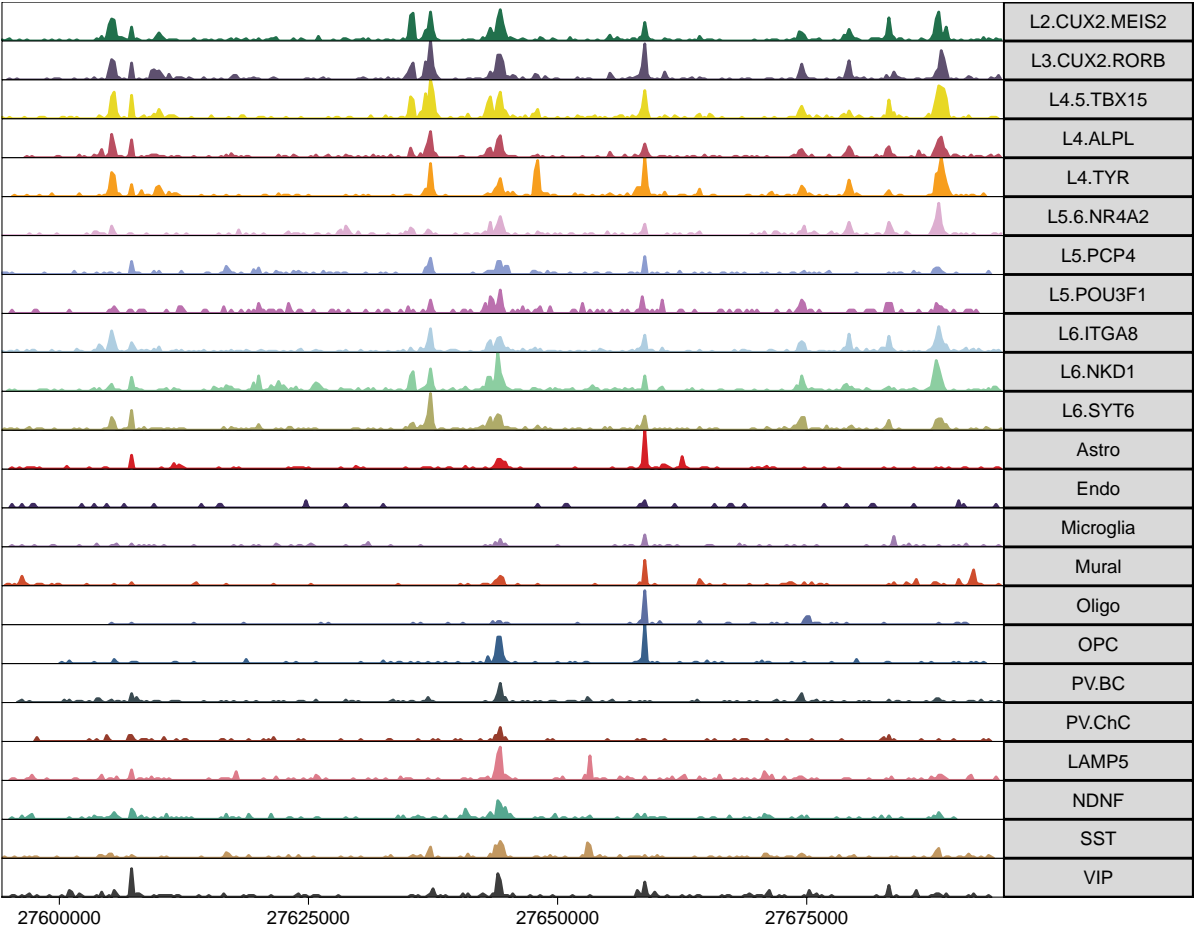
Peaks



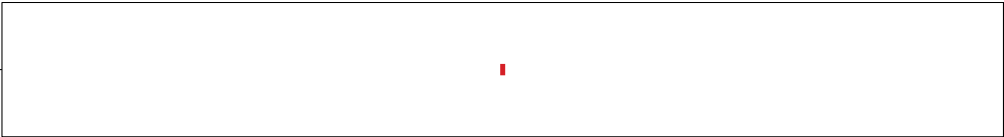
Genes

chr7:27594241–27694742 Marker Peak Tracks for: L3.CUX2.RORB

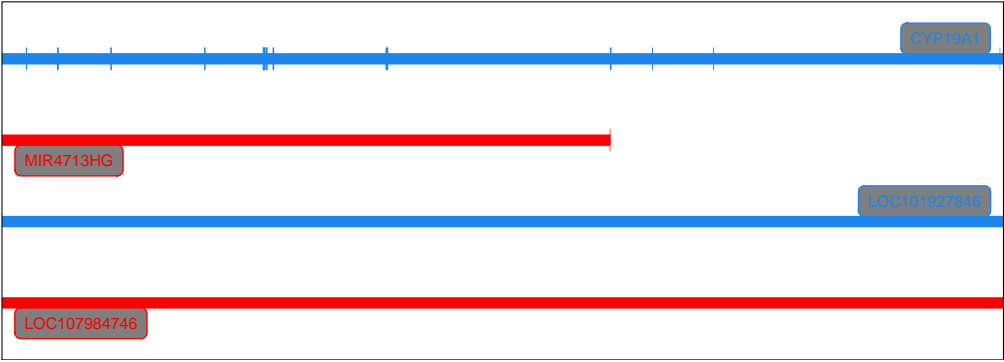
Coverage
(Norm. ATAC Signal Range (0–0.85) by ReadsInTSS)



L3.CUX2.RORB

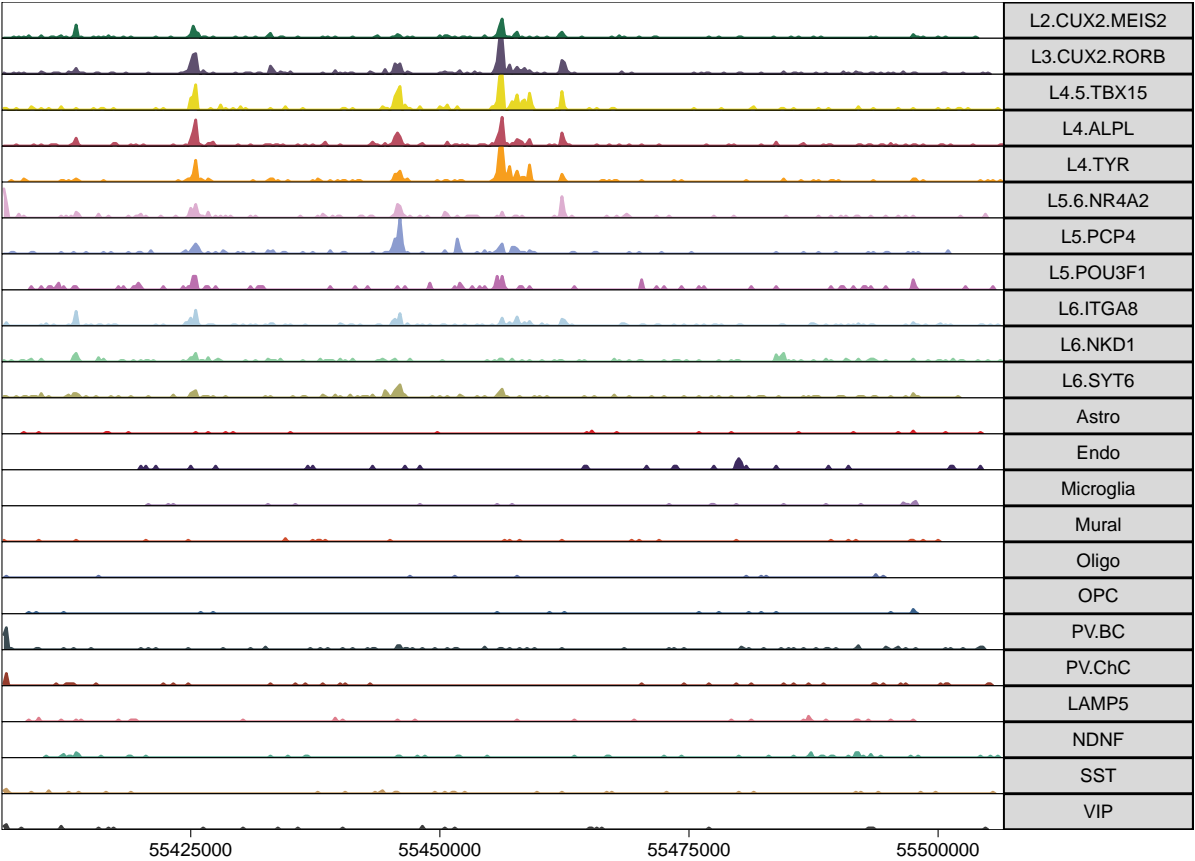


Peaks



Genes

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



L3.CUX2.RORB

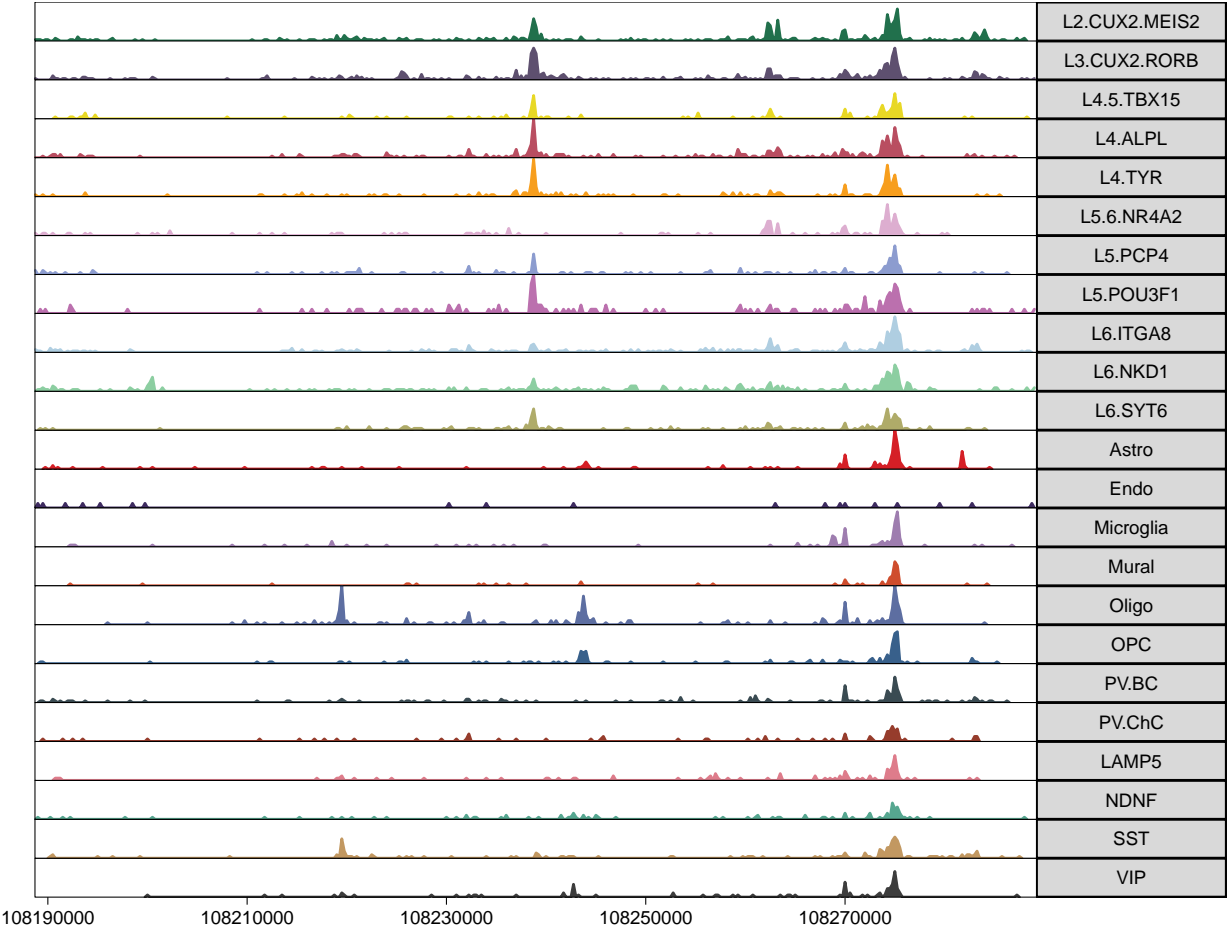


Peaks

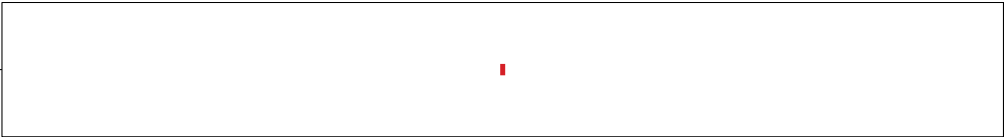
GeneTrack

chr1:108188707–108289208 Marker Peak Tracks for: L3.CUX2.RORB

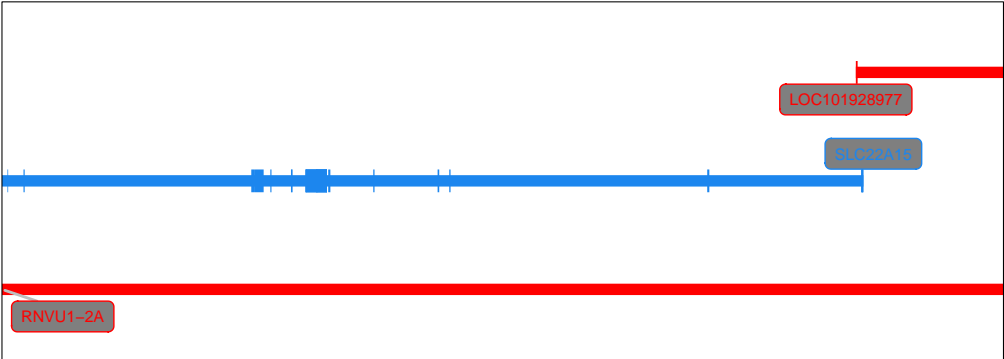
Coverage
(Norm. ATAC Signal Range (0–0.68) by ReadsInTSS)



L3.CUX2.RORB

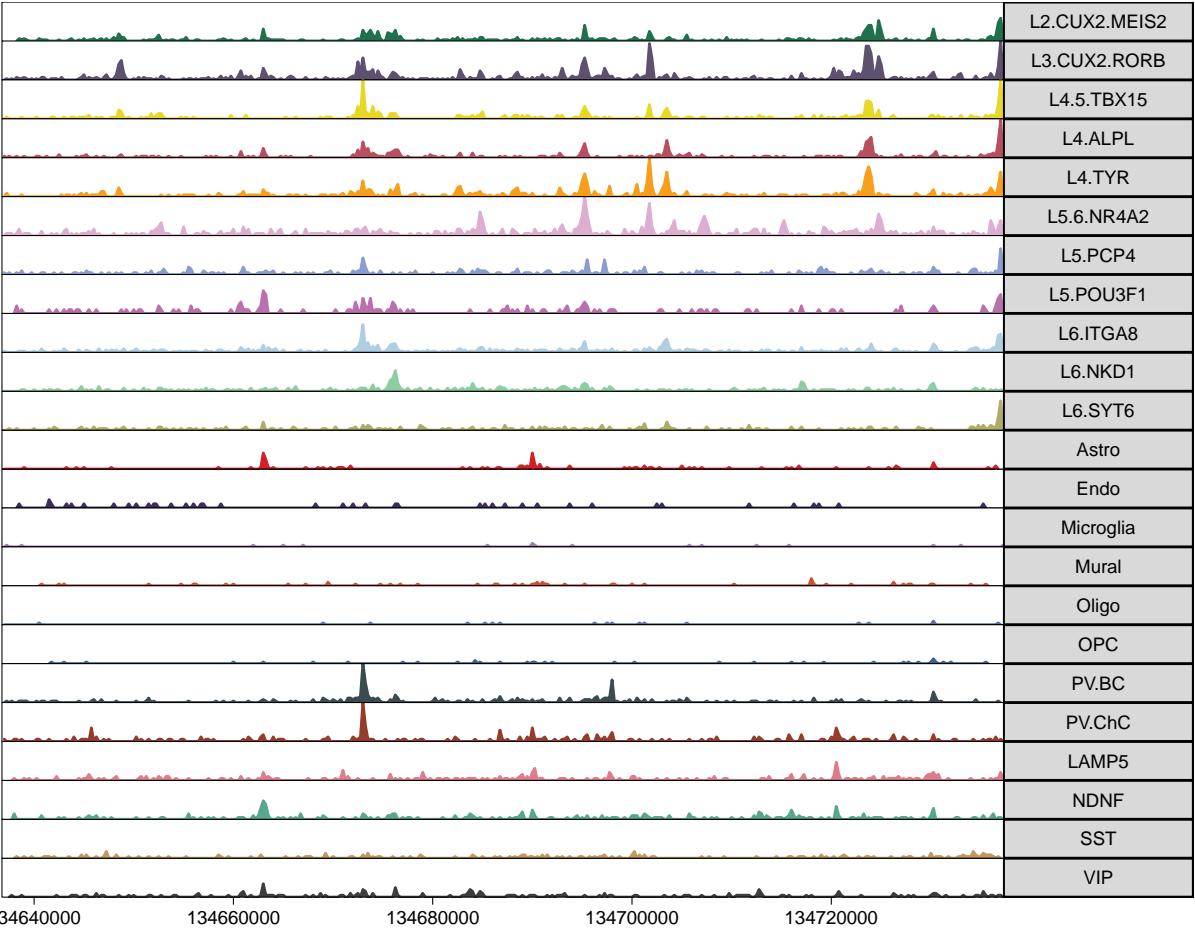


Peaks

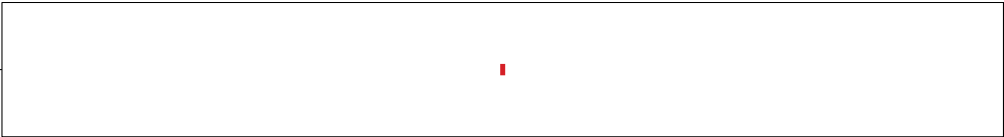


Genes

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

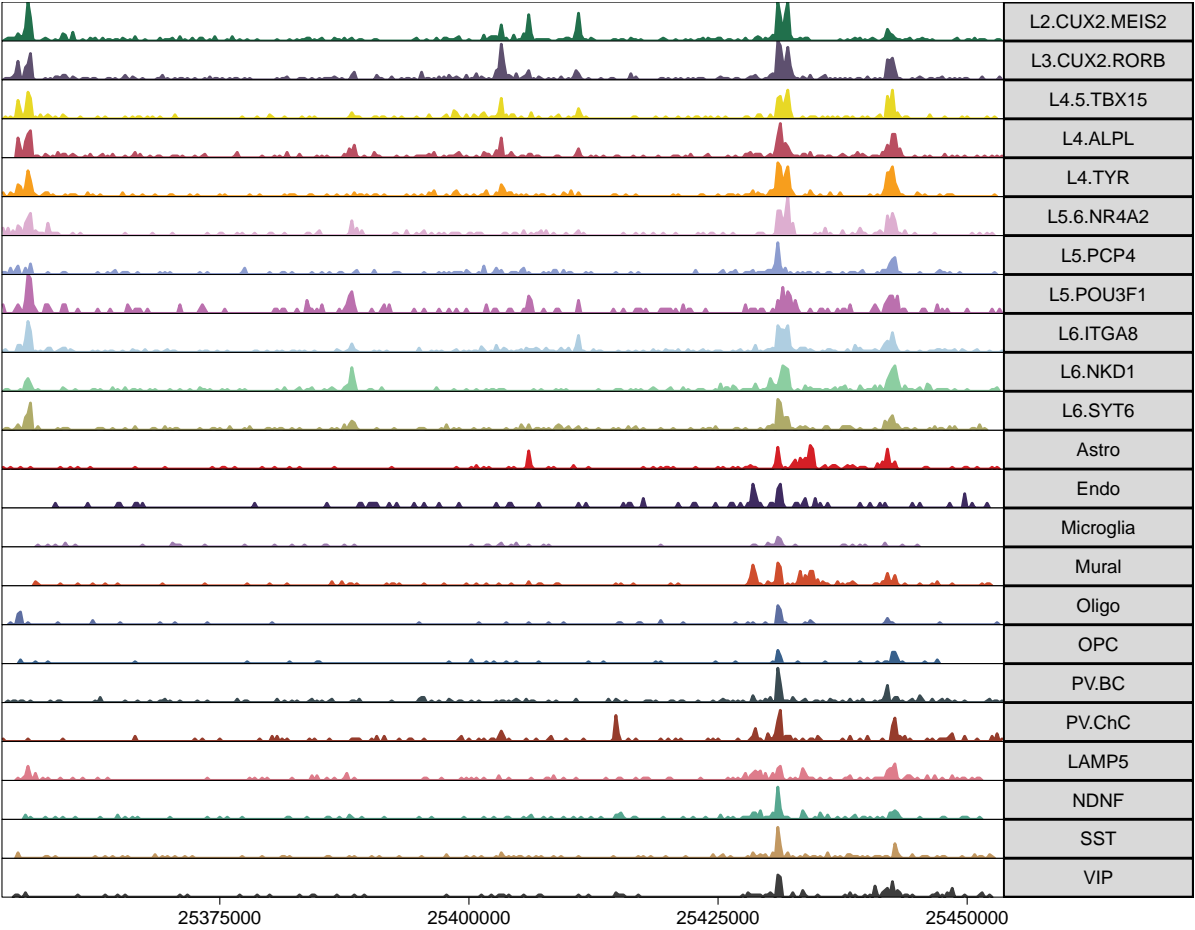


L3.CUX2.RORB

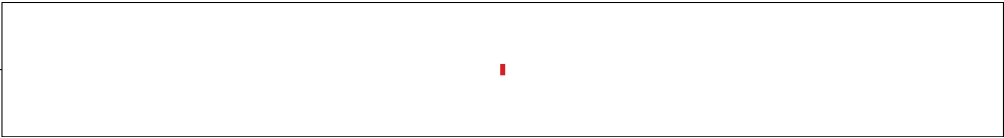


chr1:25353139–25453640 Marker Peak Tracks for: L3.CUX2.RORB

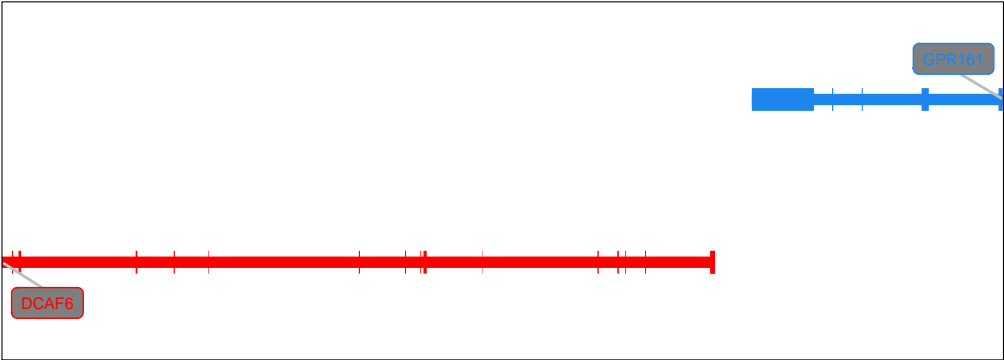
Coverage
(Norm. ATAC Signal Range (0–0.66) by ReadsInTSS)



L3.CUX2.RORB



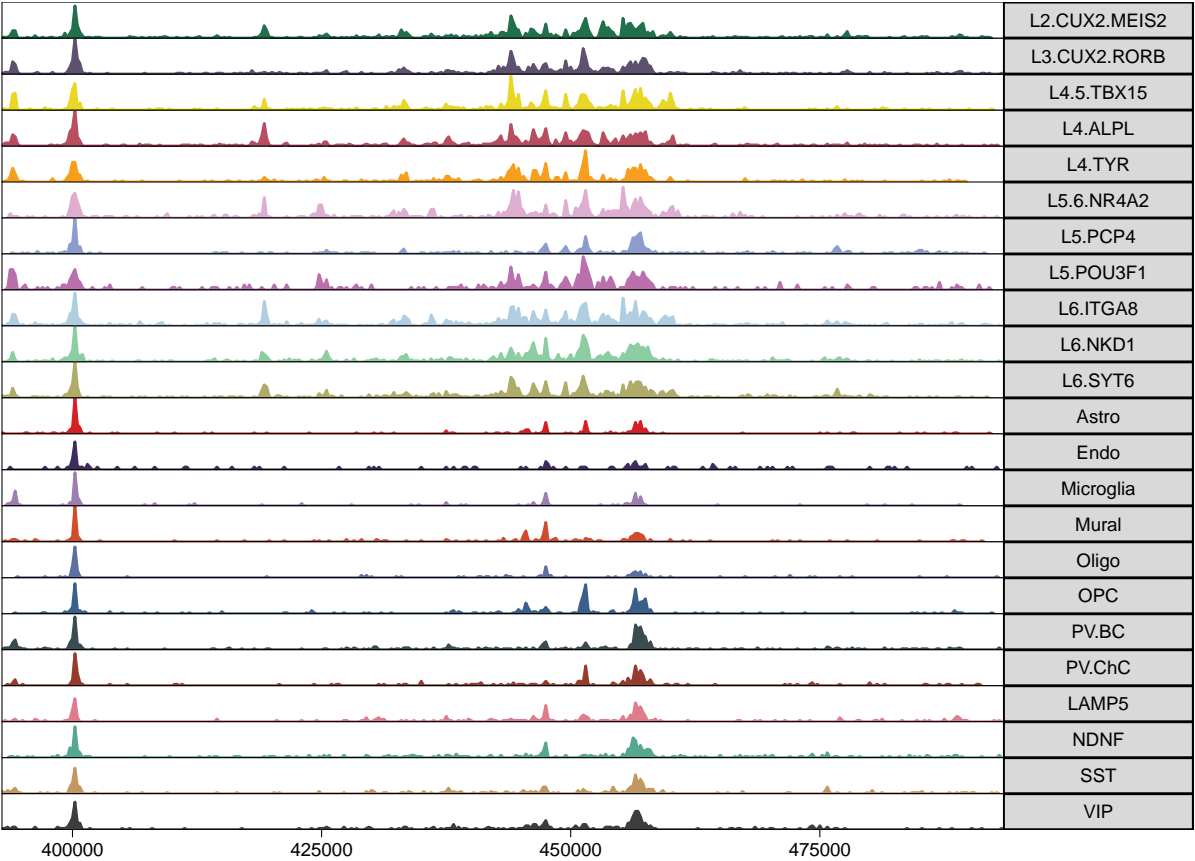
Peaks



Genes

chr17:392924–493425 Marker Peak Tracks for: L3.CUX2.RORB

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



L3.CUX2.RORB

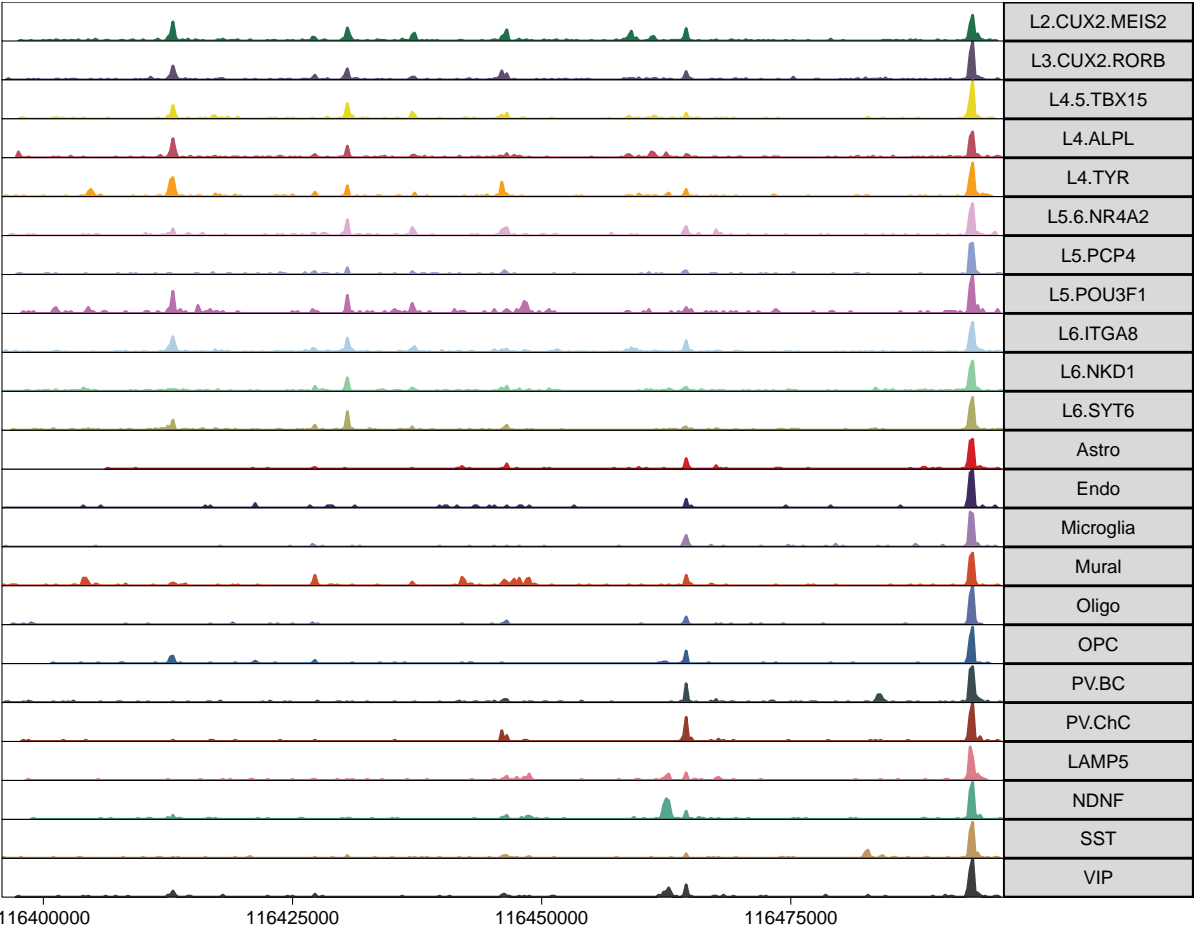


Peaks

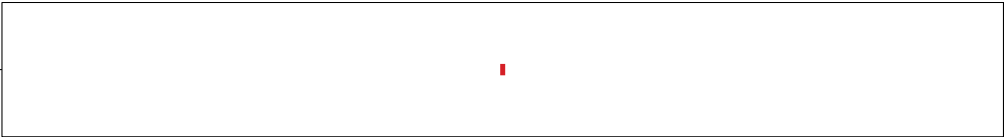
GeneTrack

strand ● ·

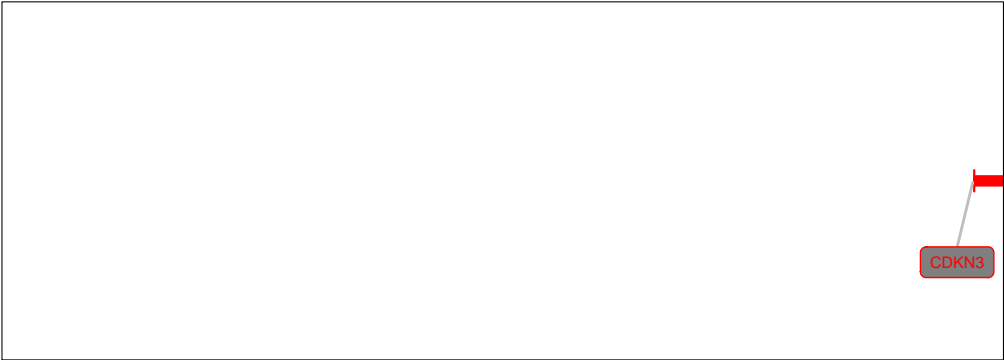
Coverage
(Norm. ATAC Signal Range (0–1.42) by ReadsInTSS)



L3.CUX2.RORB

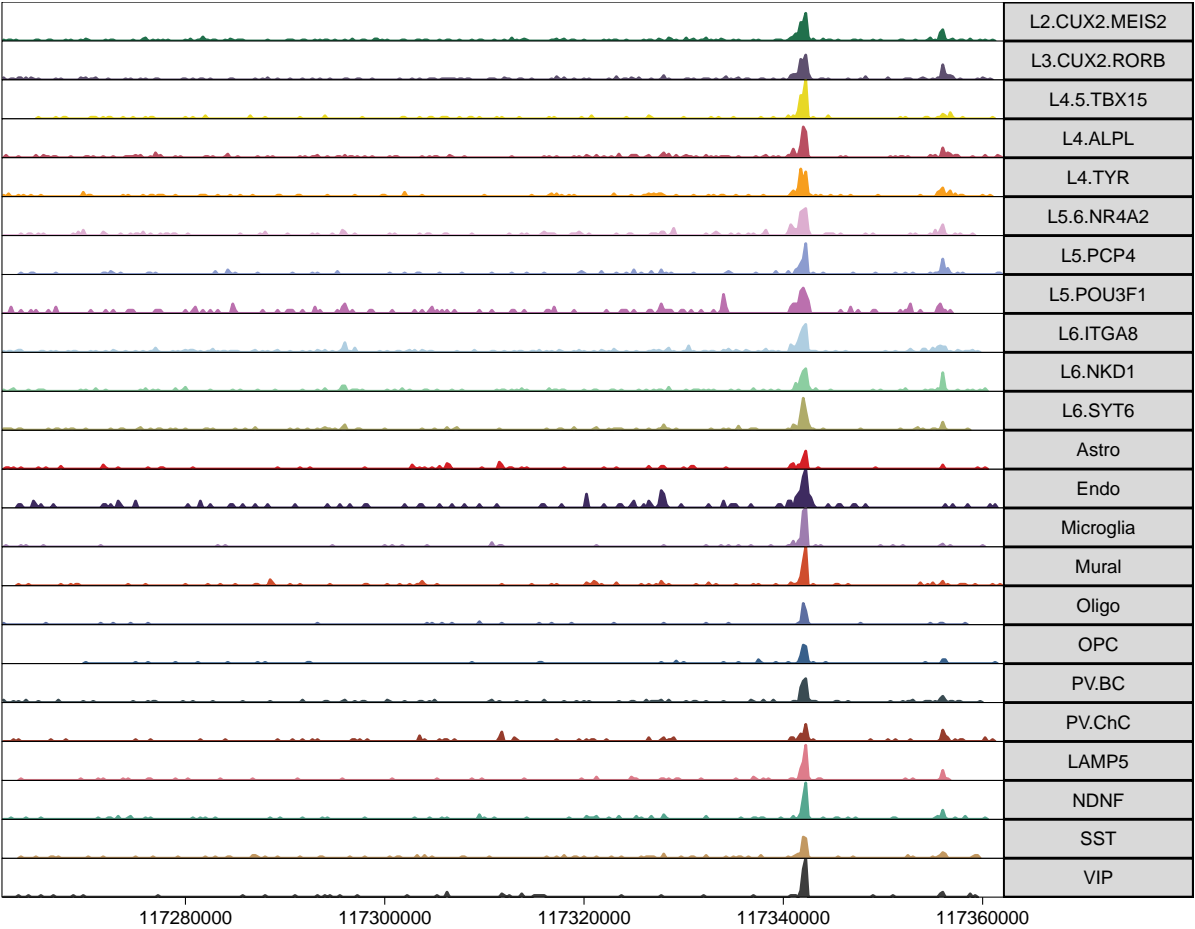


Peaks

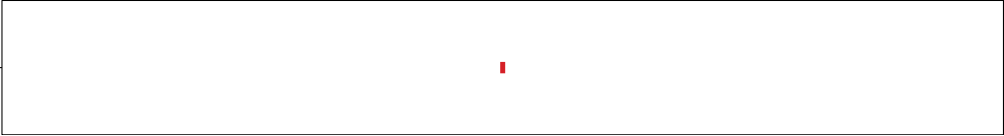


Genes

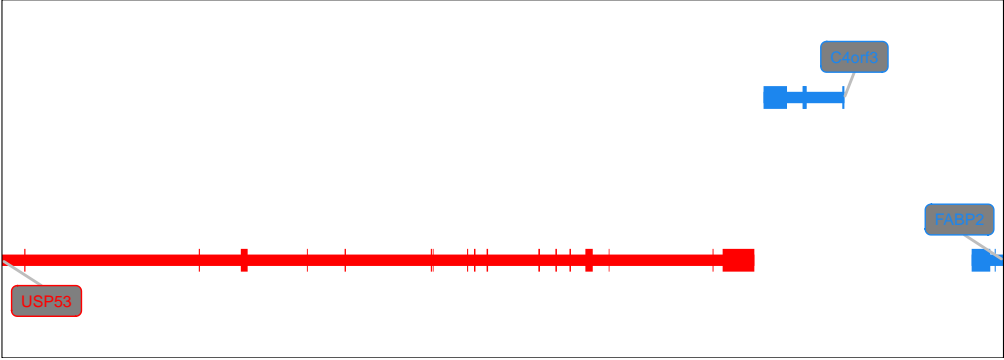
Coverage
(Norm. ATAC Signal Range (0–0.9) by ReadsInTSS)



L3.CUX2.RORB

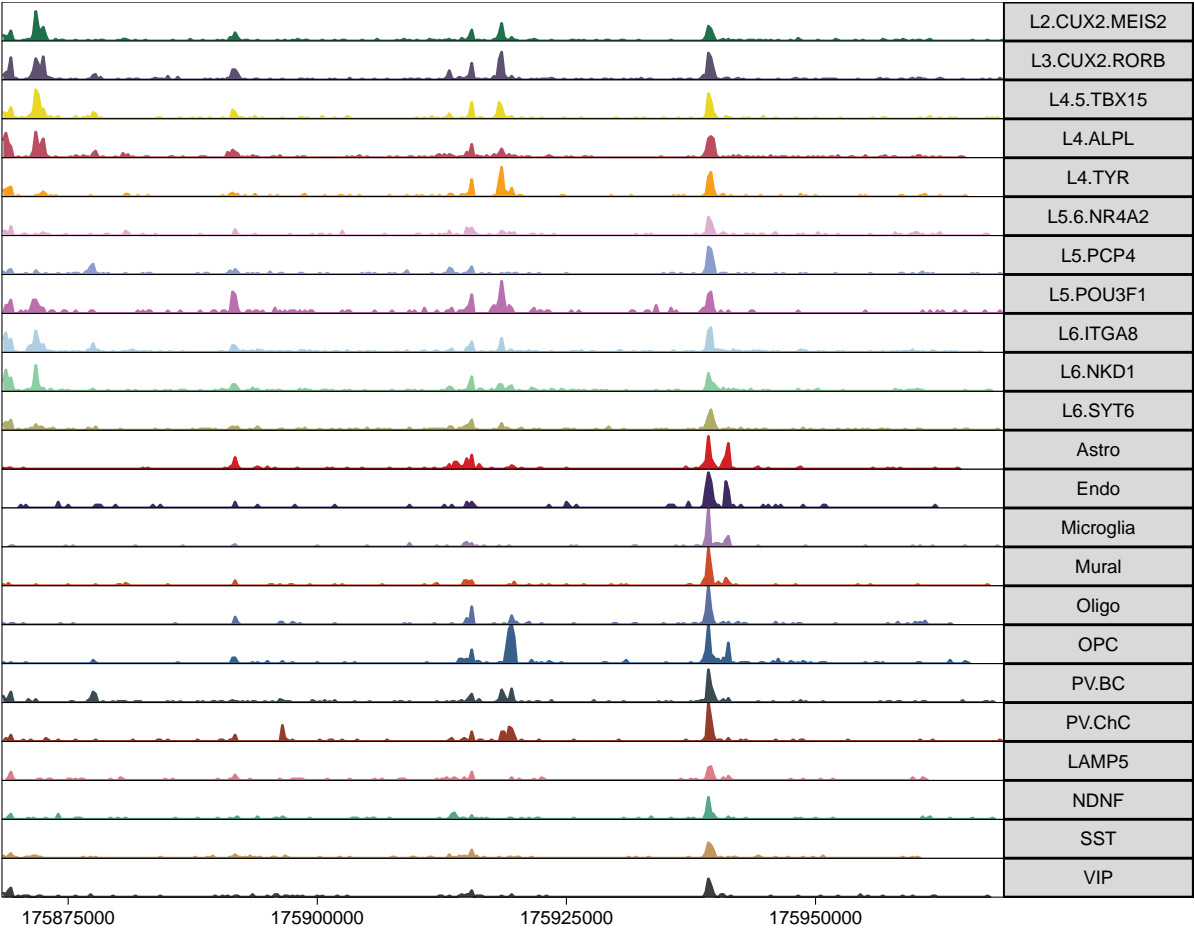


Peaks

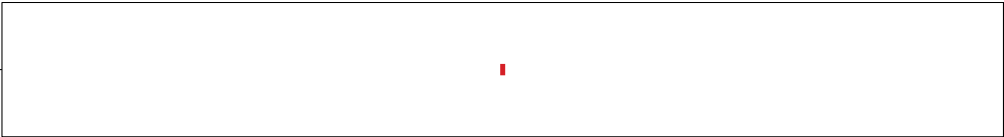


Genes

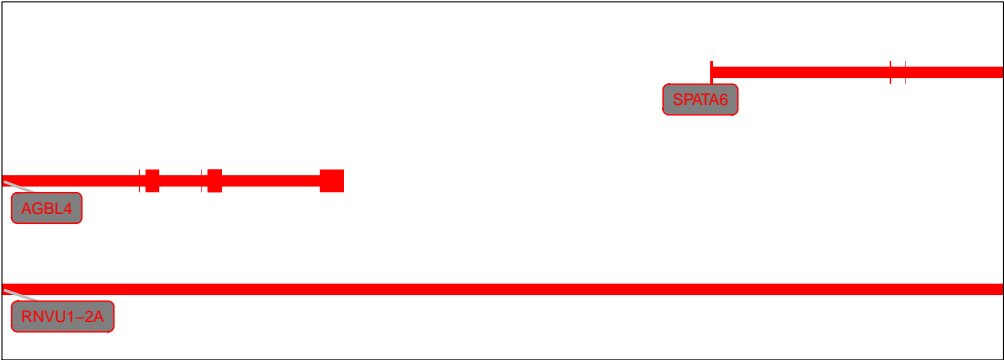
Coverage
(Norm. ATAC Signal Range (0–1.06) by ReadsInTSS)



L3.CUX2.RORB



Peaks



Genes