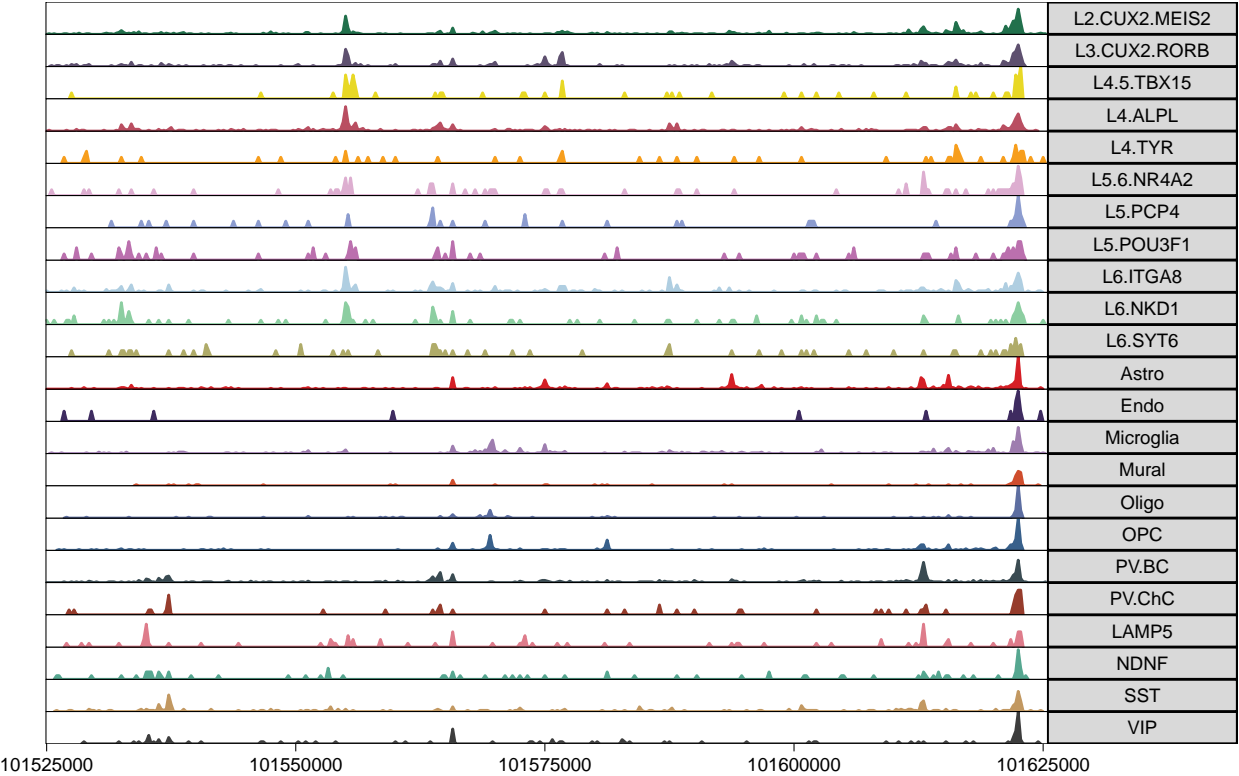


chr11:101524932–101625433 Marker Peak Tracks for: L3.CUX2.RORB

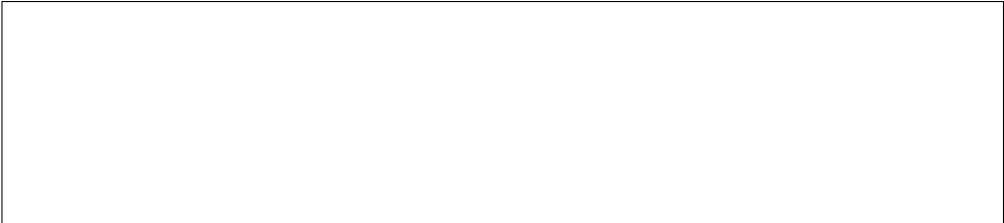
Coverage  
(Norm. ATAC Signal Range 0–1.24) by ReadsInTSS)



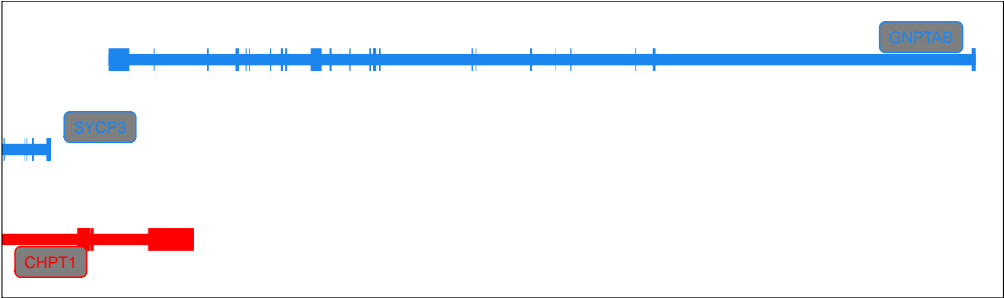
L3.CUX2.RORB



Peaks

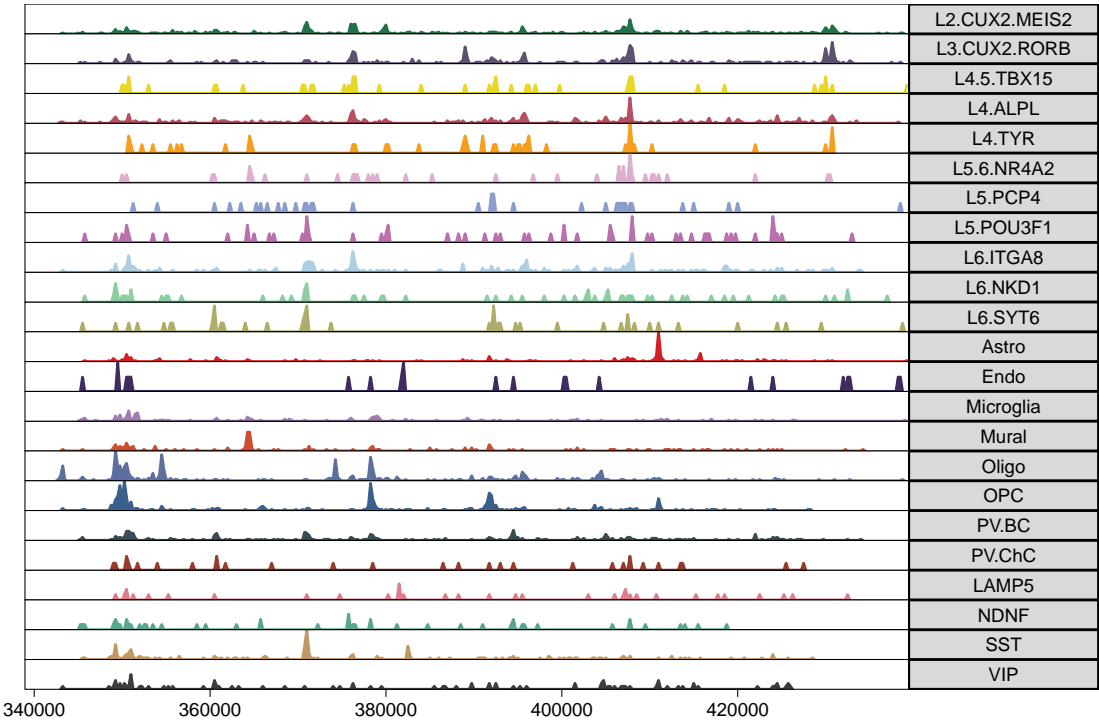


LoopTrack

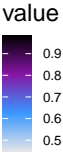
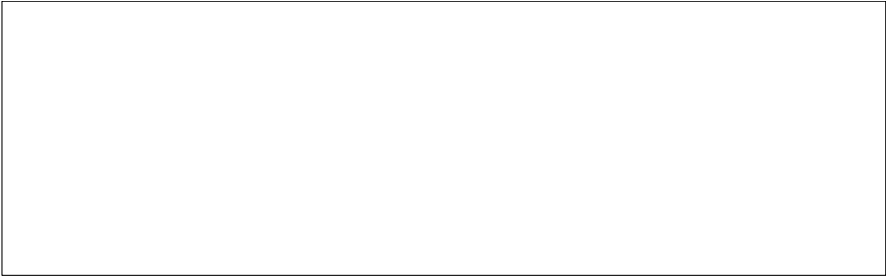


Genes

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

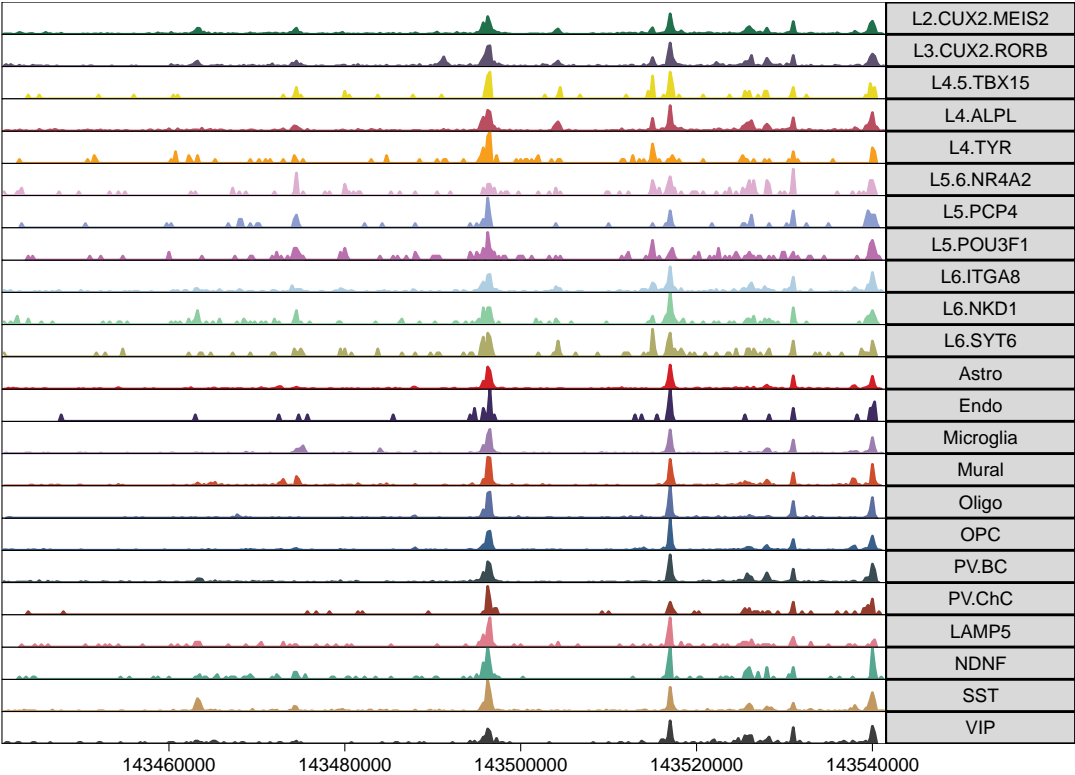


L3.CUX2.RORB

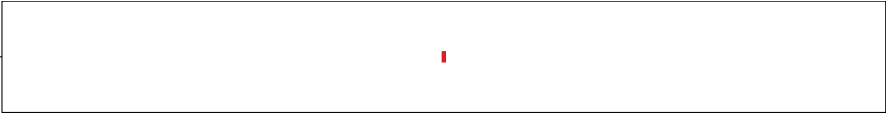


strand    ●    ·

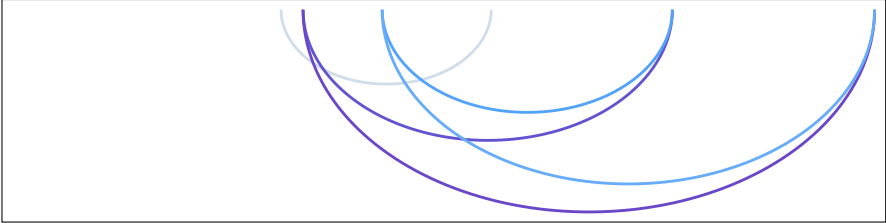
Coverage  
(Norm. ATAC Signal Range 0–1.94) by ReadsInTSS)



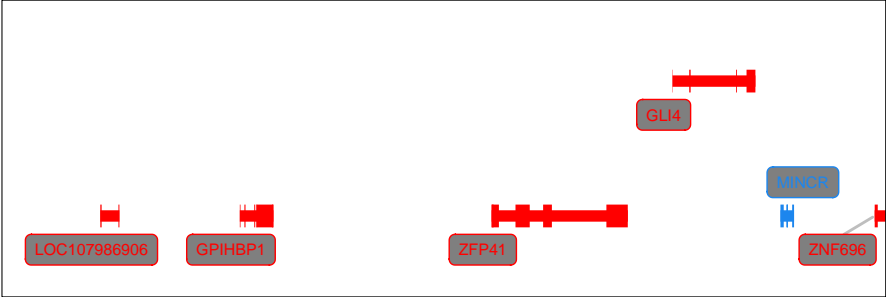
L3.CUX2.RORB



Peaks

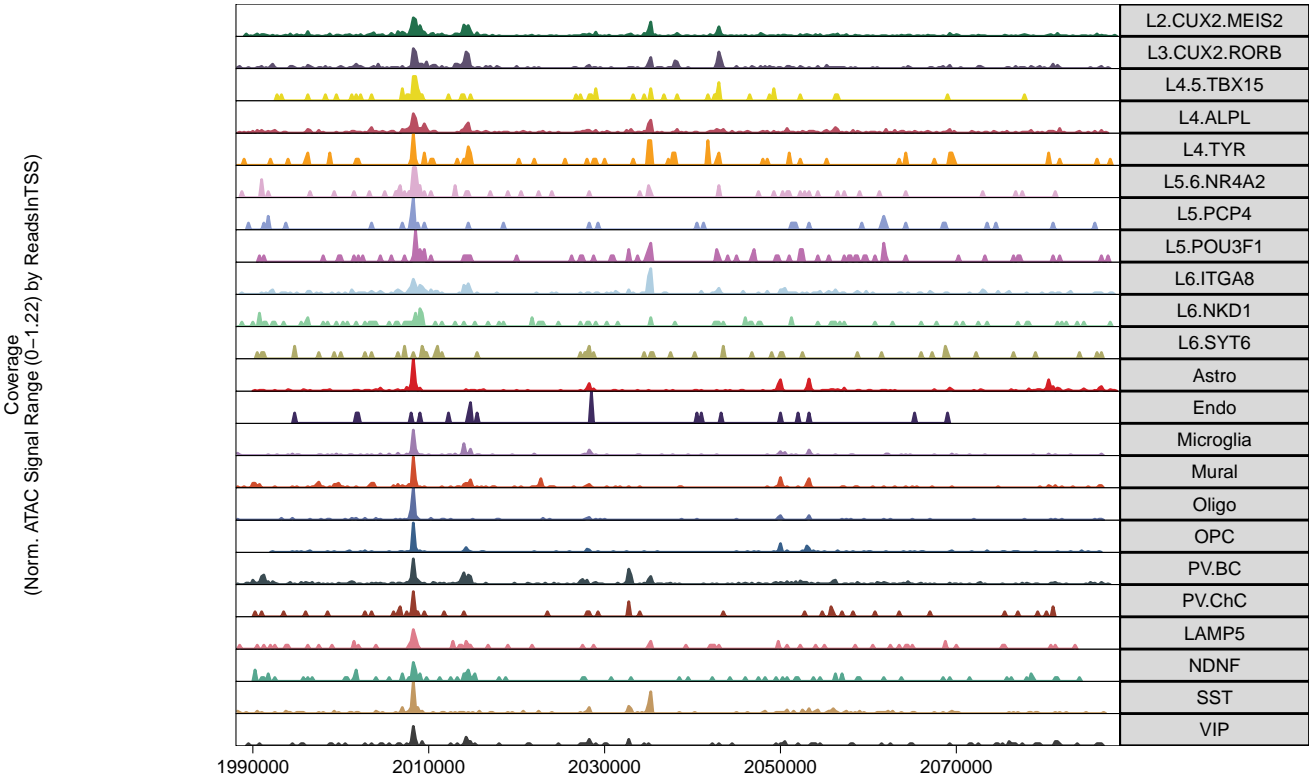


Peak2GeneLinks

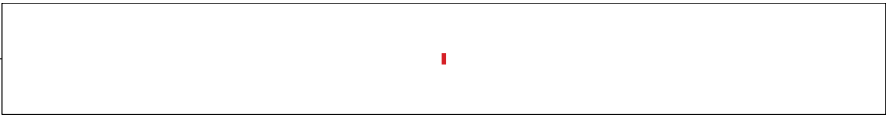


Genes

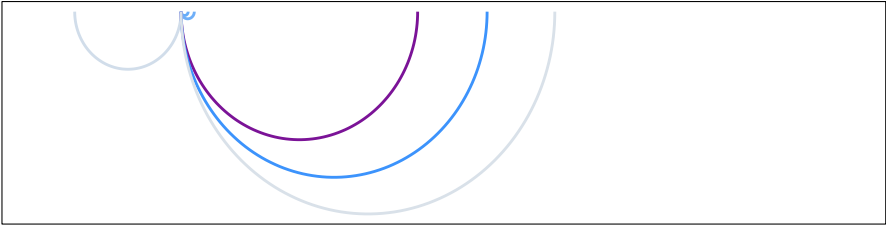
chr15:1988078–2088579 Marker Peak Tracks for: L3.CUX2.RORB



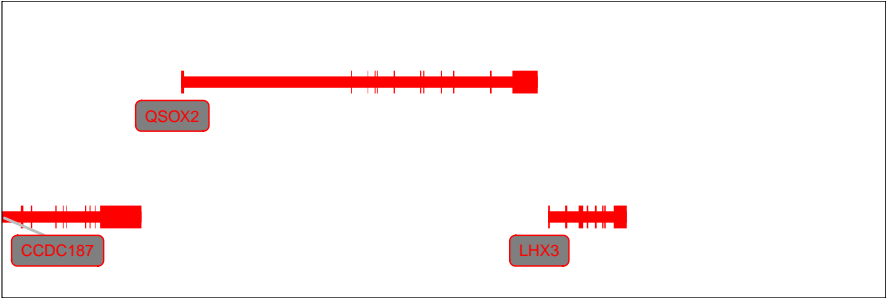
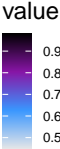
L3.CUX2.RORB



Peaks

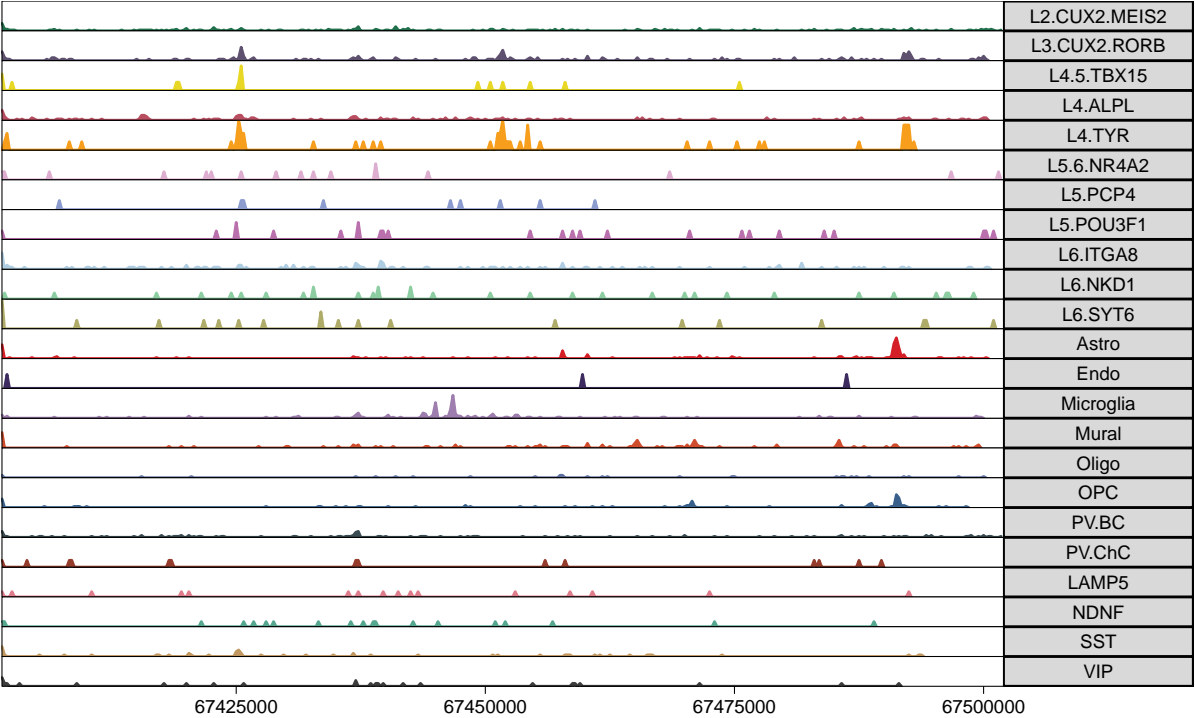


Peak2GeneLinks



Genes

Coverage  
(Norm. ATAC Signal Range (0–0.83) by Reads in TSS)



L3.CUX2.RORB

