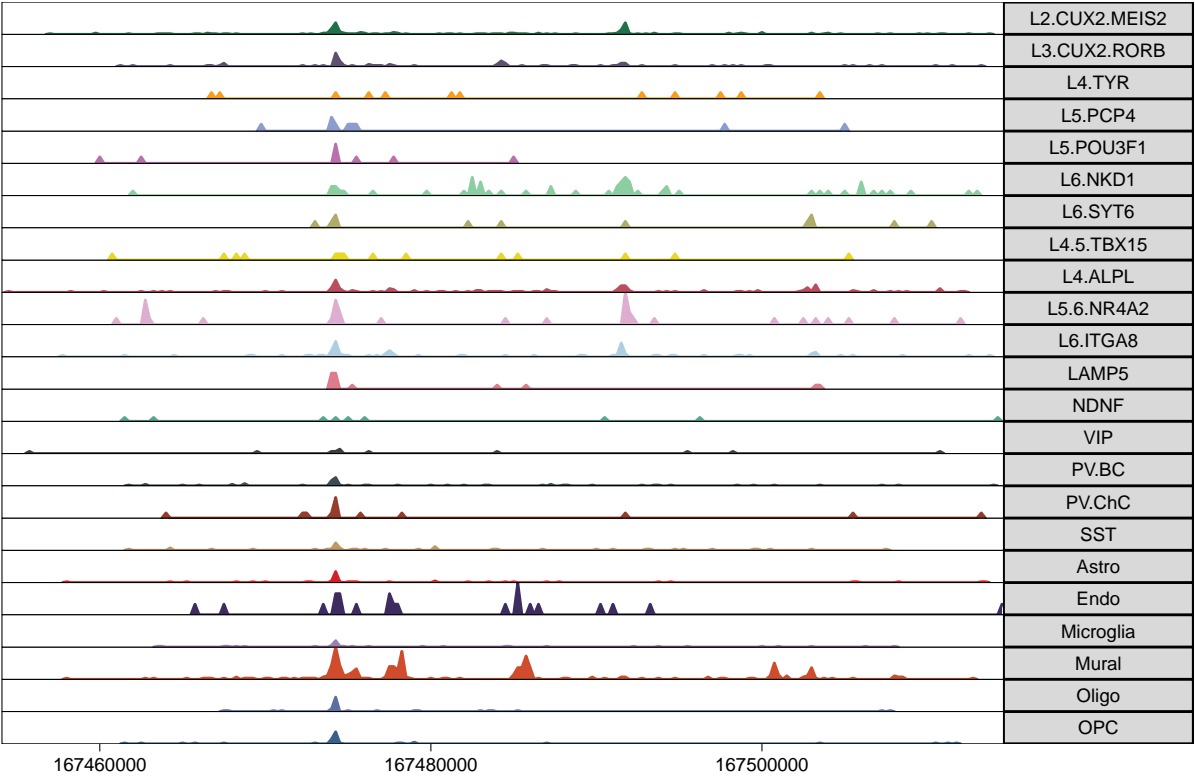
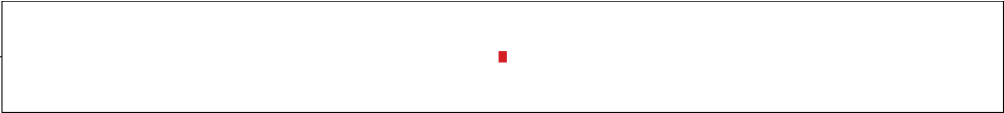


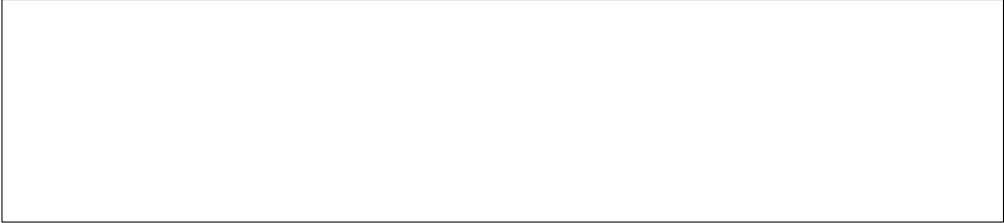
Coverage  
(Norm. ATAC Signal Range (0–1.19) by ReadsInTSS)



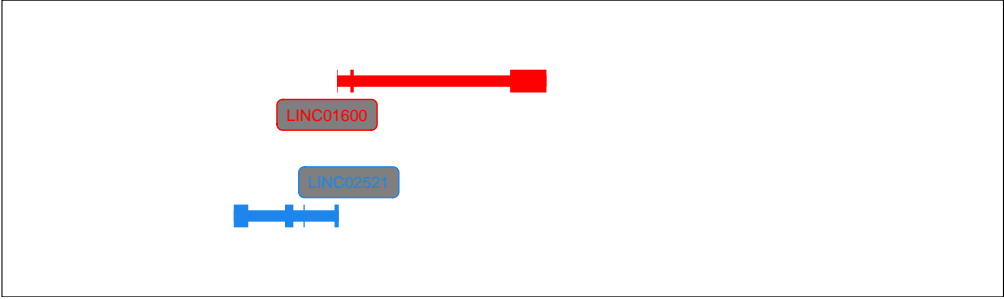
L3.CUX2.RORB



Peaks

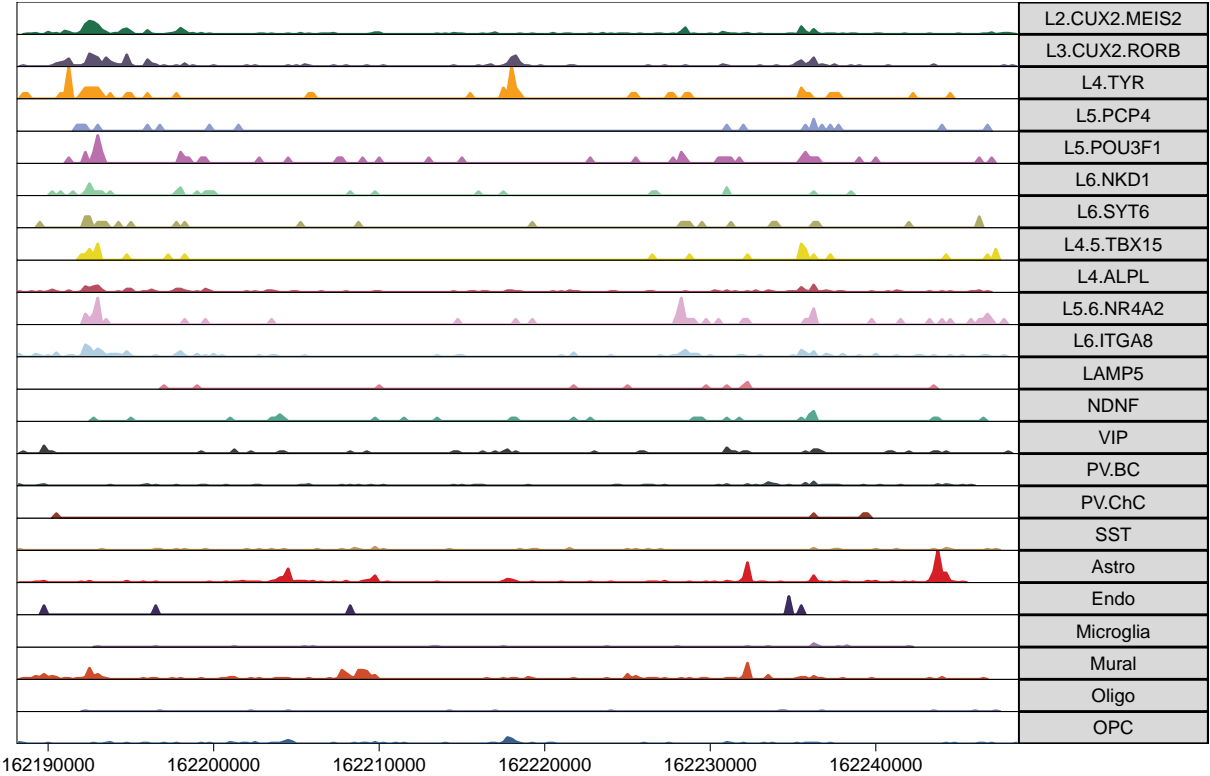


LoopTrack



Genes

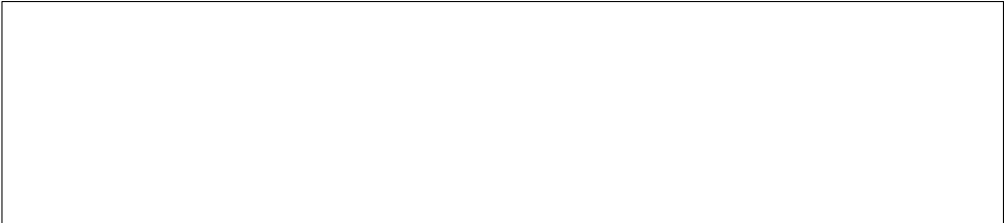
Coverage  
(Norm. ATAC Signal Range 0–1.37) by ReadsInTSS)



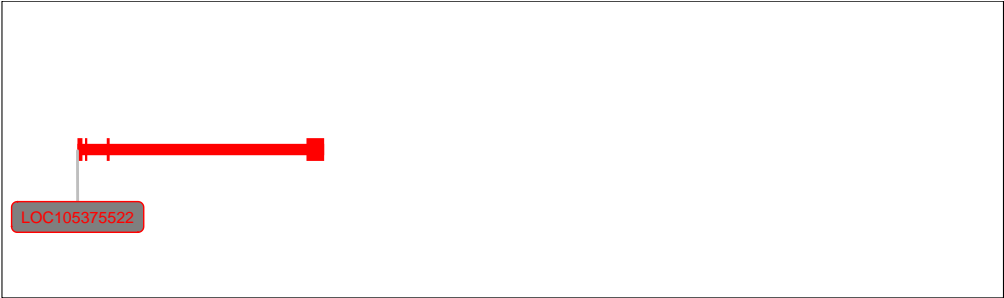
L3.CUX2.RORB



Peaks

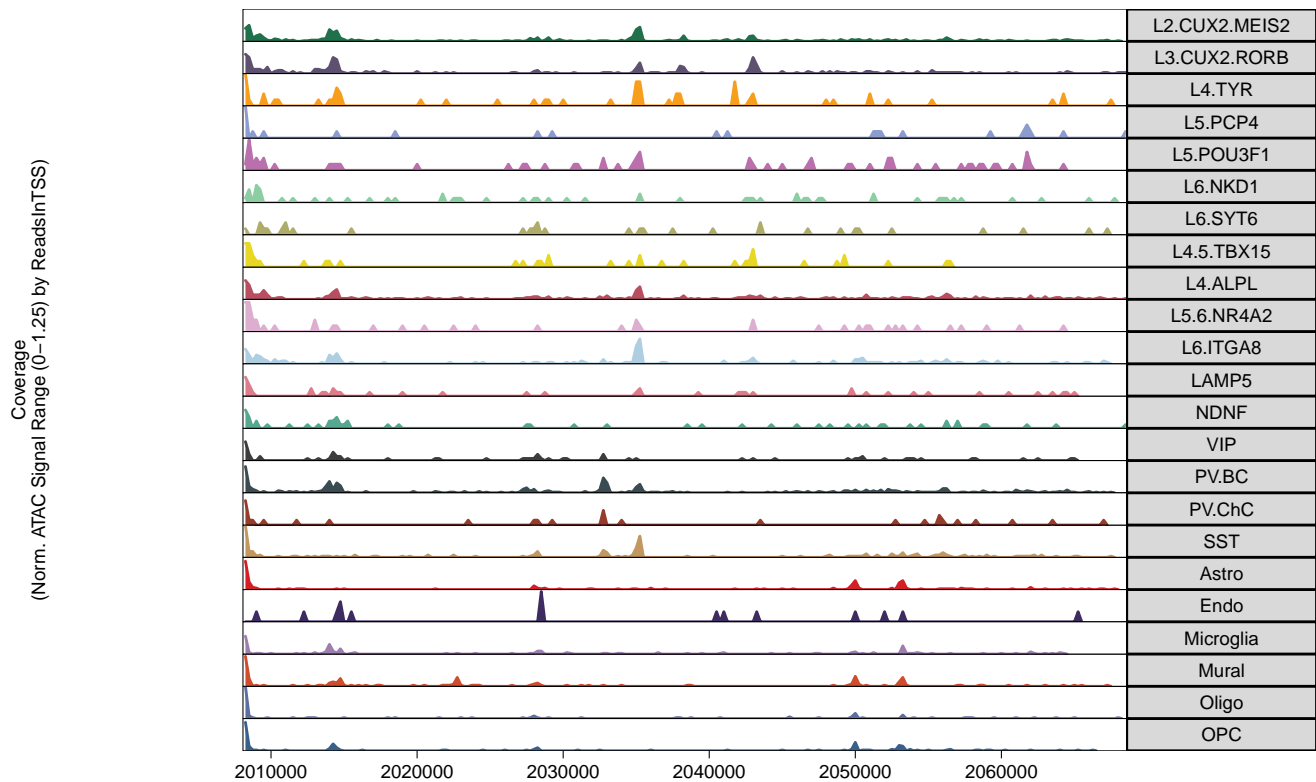


LoopTrack

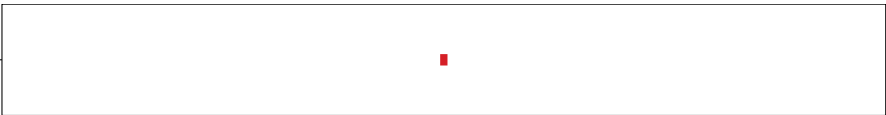


Genes

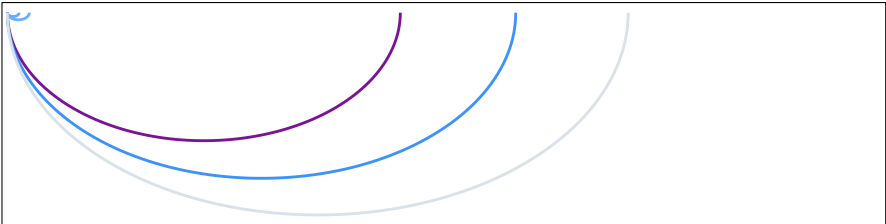
chr15:2008078–2068579 Marker Peak Tracks for: L3.CUX2.RORB



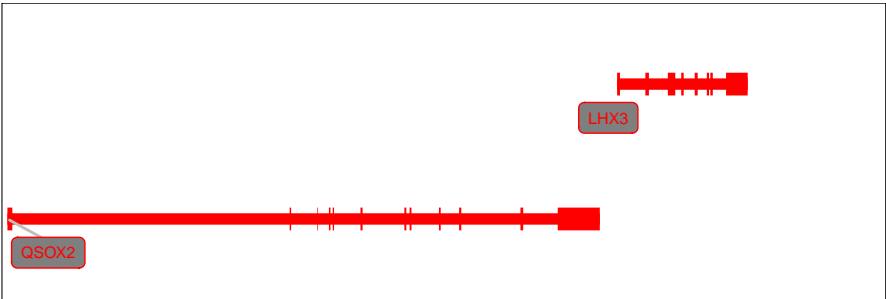
L3.CUX2.RORB



Peaks



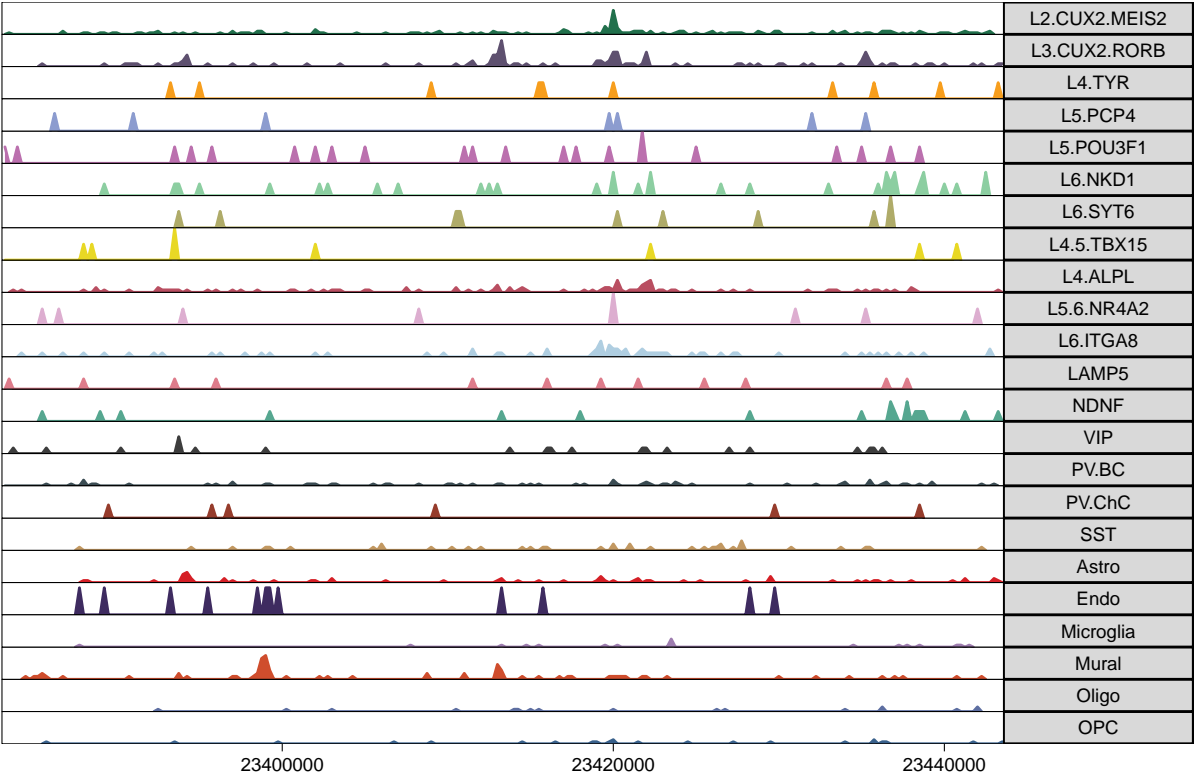
Peak2GeneLinks



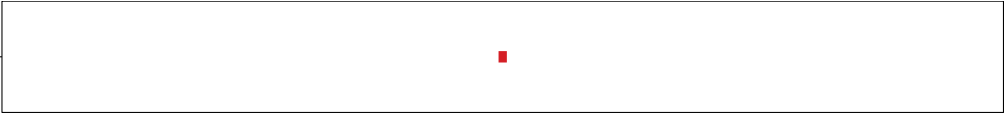
Genes

chr17:23383068–23443569 Marker Peak Tracks for: L3.CUX2.RORB

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



L3.CUX2.RORB



Peaks

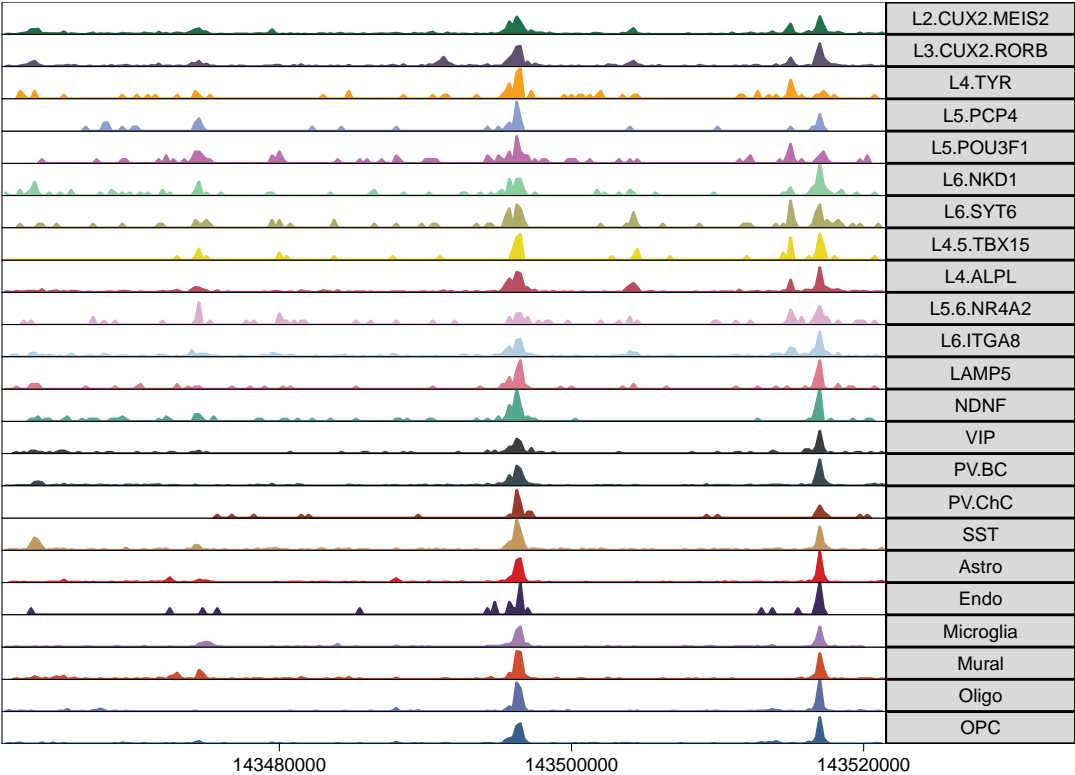


LoopTrack

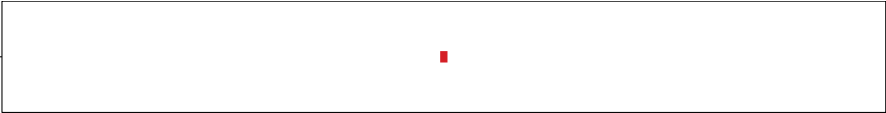


Genes

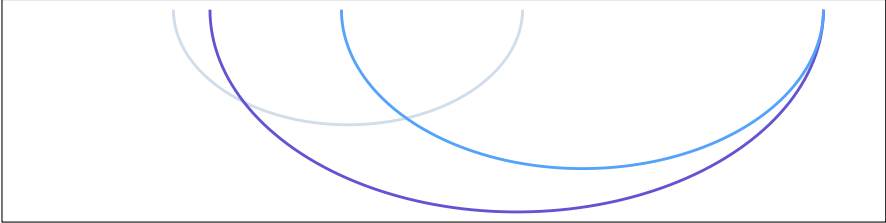
Coverage  
(Norm. ATAC Signal Range (0–1.99) by ReadsInTSS)



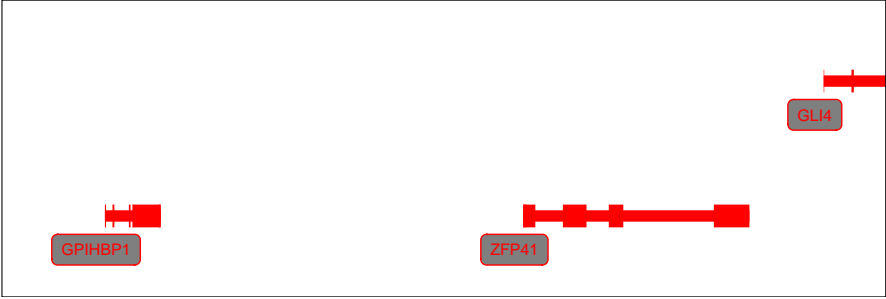
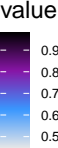
L3.CUX2.RORB



Peaks



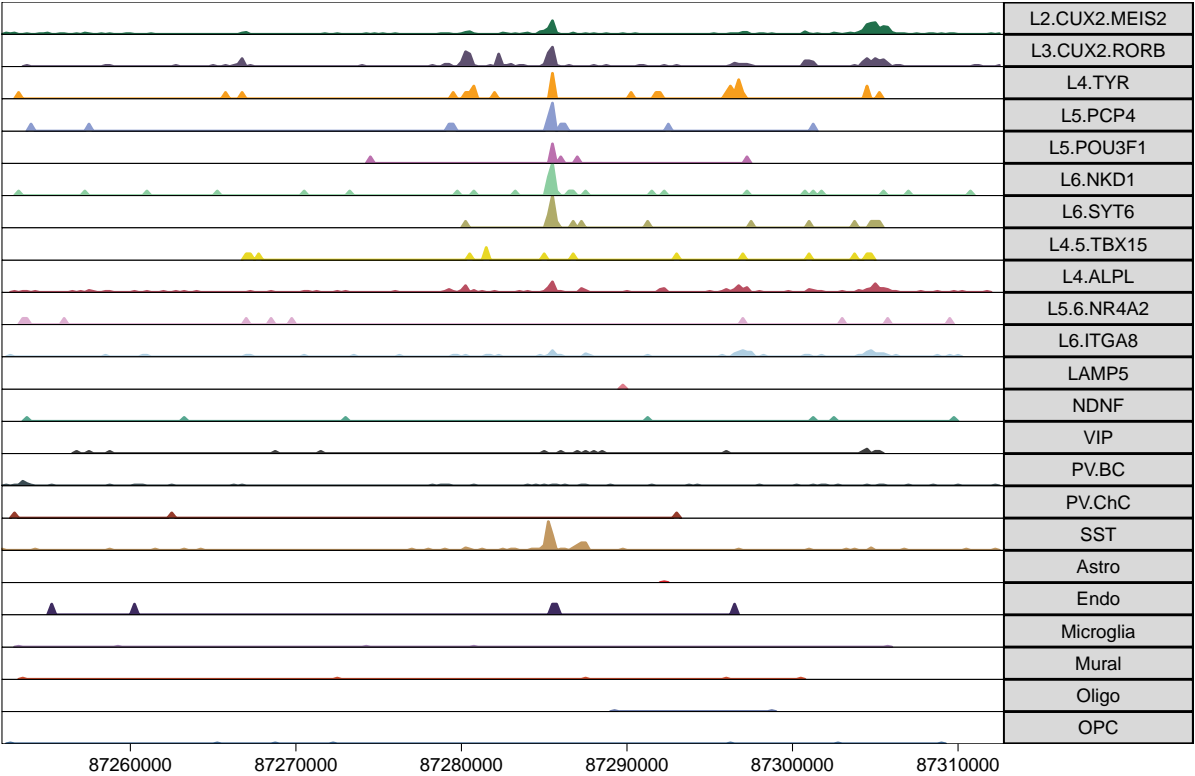
Peak2GeneLinks



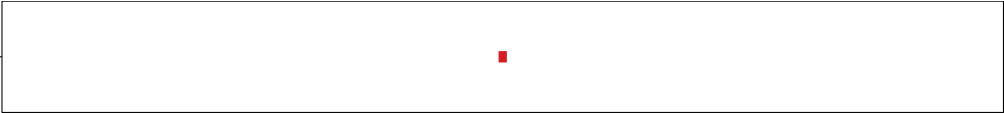
Genes

chr4:87252242–87312743 Marker Peak Tracks for: L3.CUX2.RORB

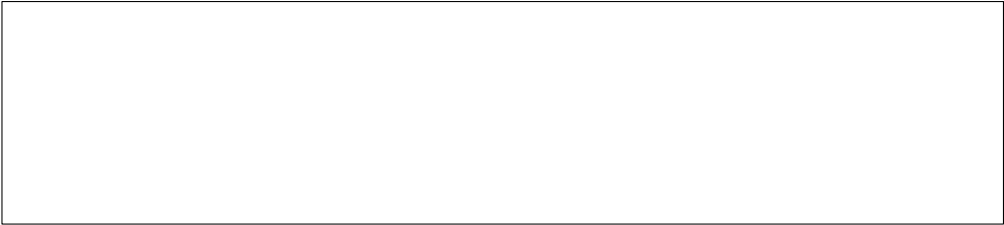
Coverage  
(Norm. ATAC Signal Range (0–1.18) by ReadsInTSS)



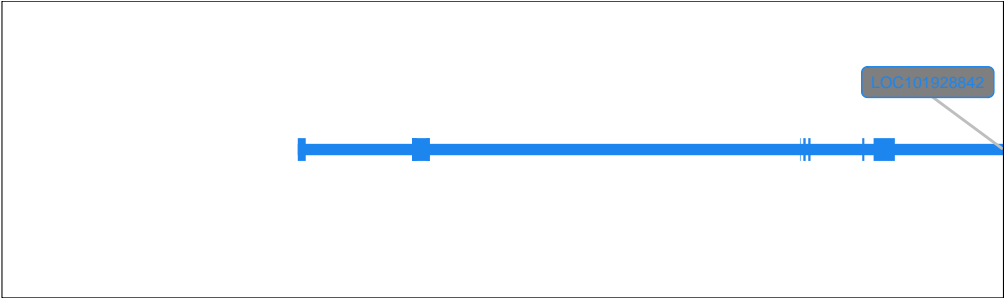
L3.CUX2.RORB



Peaks



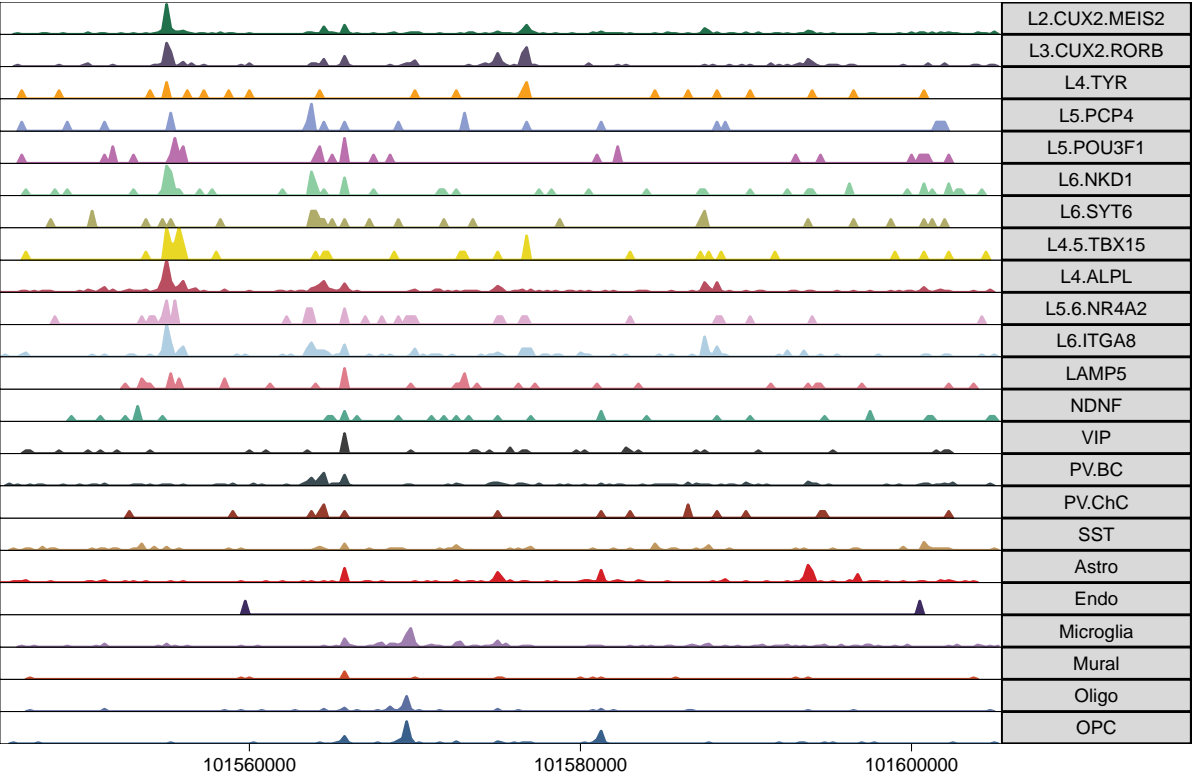
LoopTrack



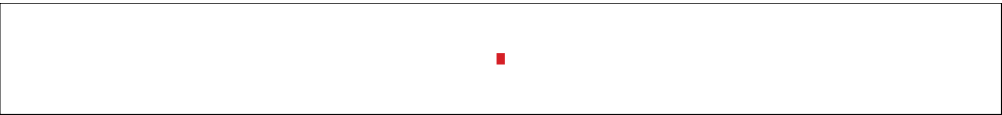
Genes

chr11:101544932–101605433 Marker Peak Tracks for: L3.CUX2.RORB

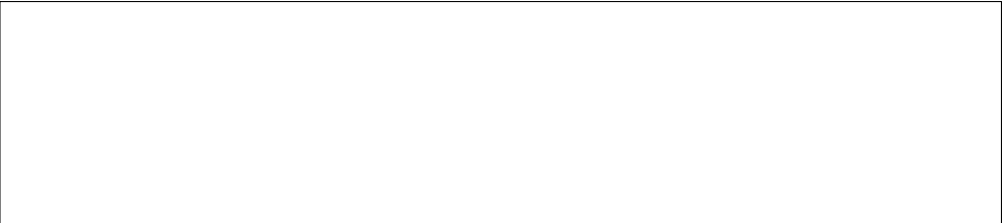
Coverage  
(Norm. ATAC Signal Range 0–0.92) by ReadsInTSS)



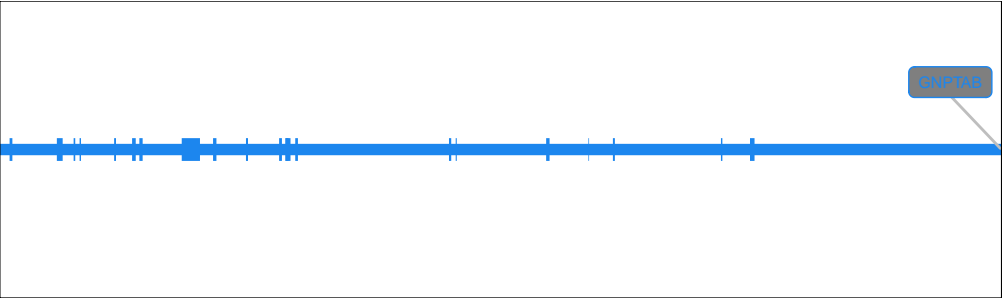
L3.CUX2.RORB



Peaks

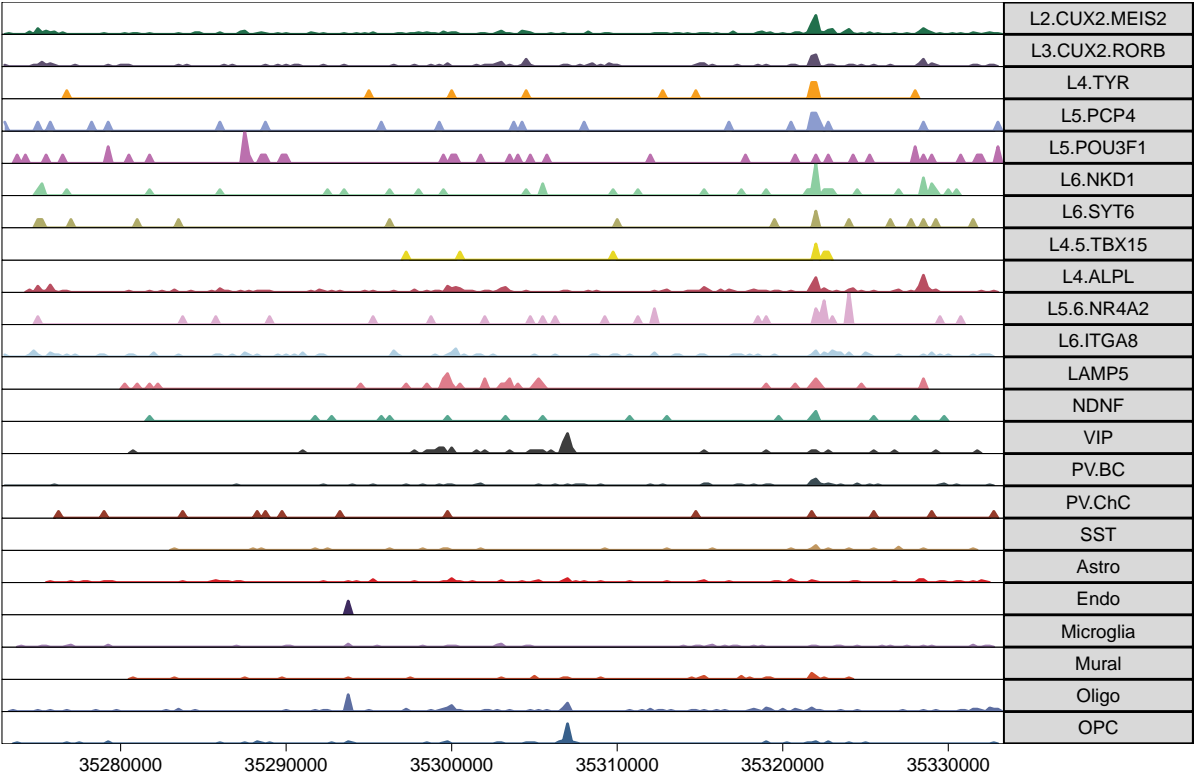


LoopTrack

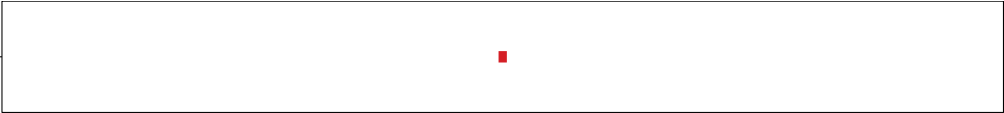


Genes

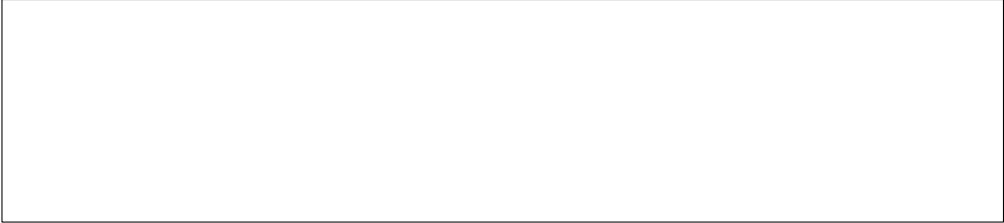
Coverage  
(Norm. ATAC Signal Range (0–0.92) by ReadsInTSS)



L3.CUX2.RORB



Peaks



LoopTrack

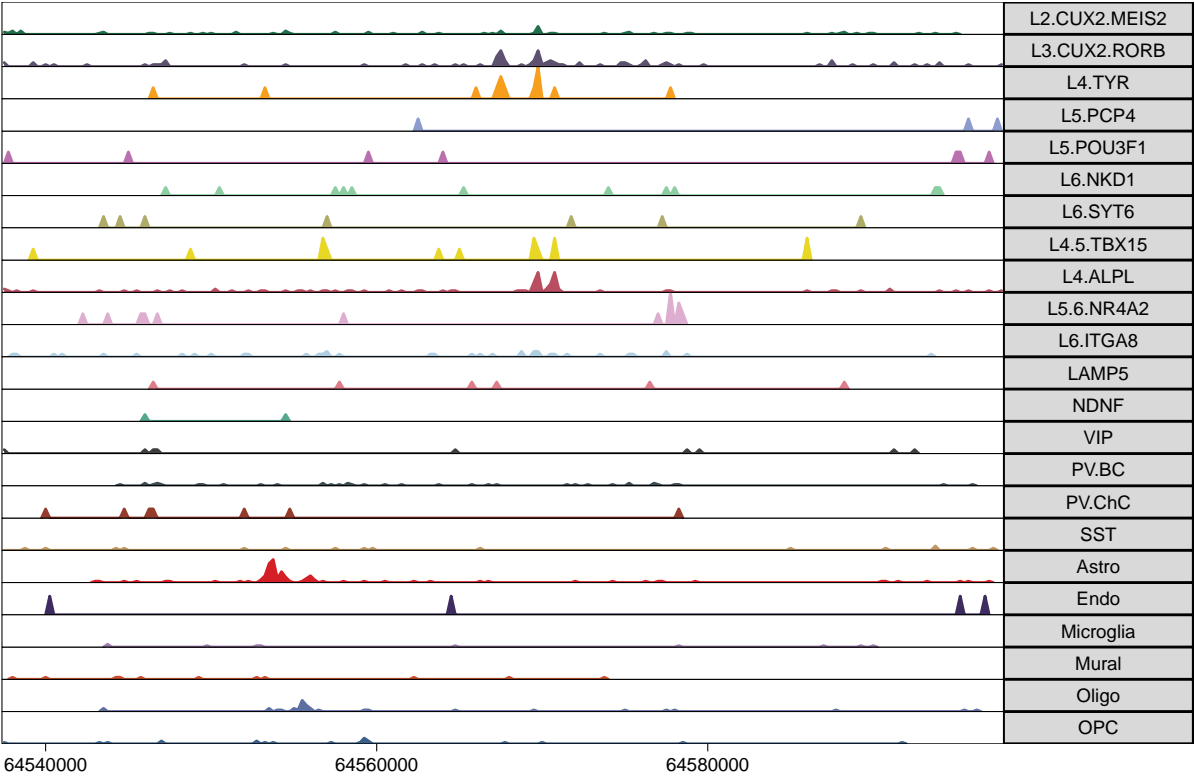


Genes

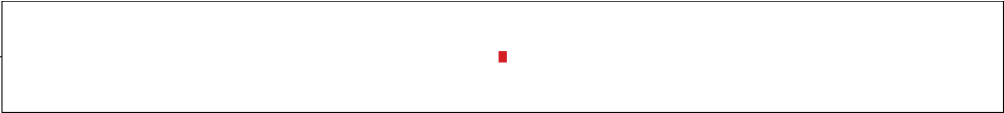


chr20:64537362–64597863 Marker Peak Tracks for: L3.CUX2.RORB

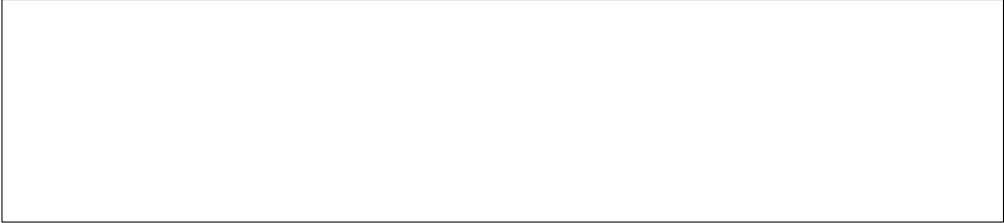
Coverage  
(Norm. ATAC Signal Range (0–0.68) by ReadsInTSS)



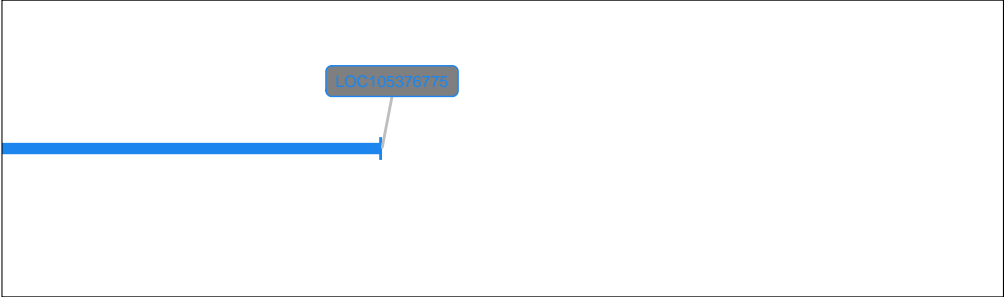
L3.CUX2.RORB



Peaks

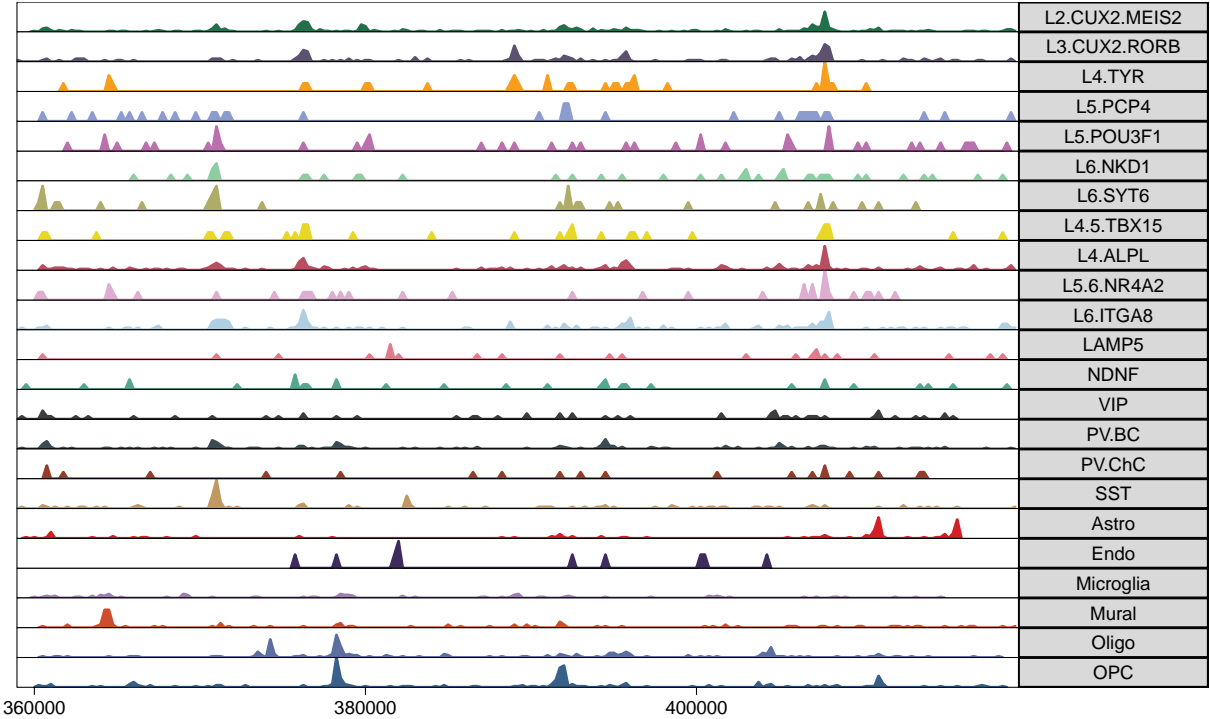


LoopTrack

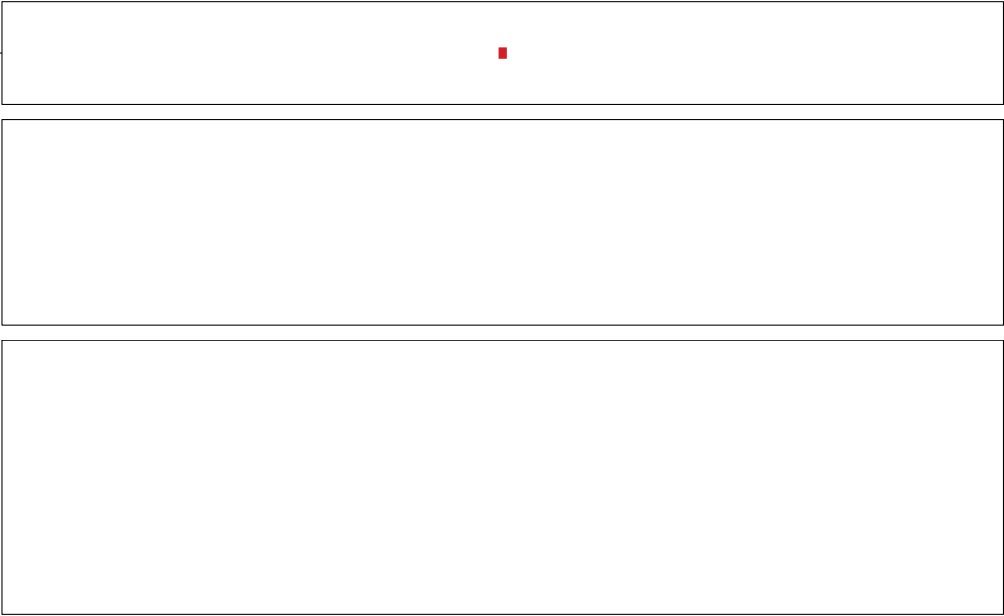


Genes

Coverage  
(Norm. ATAC Signal Range (0–0.87) by Reads in TSS)



L3.CUX2.RORB

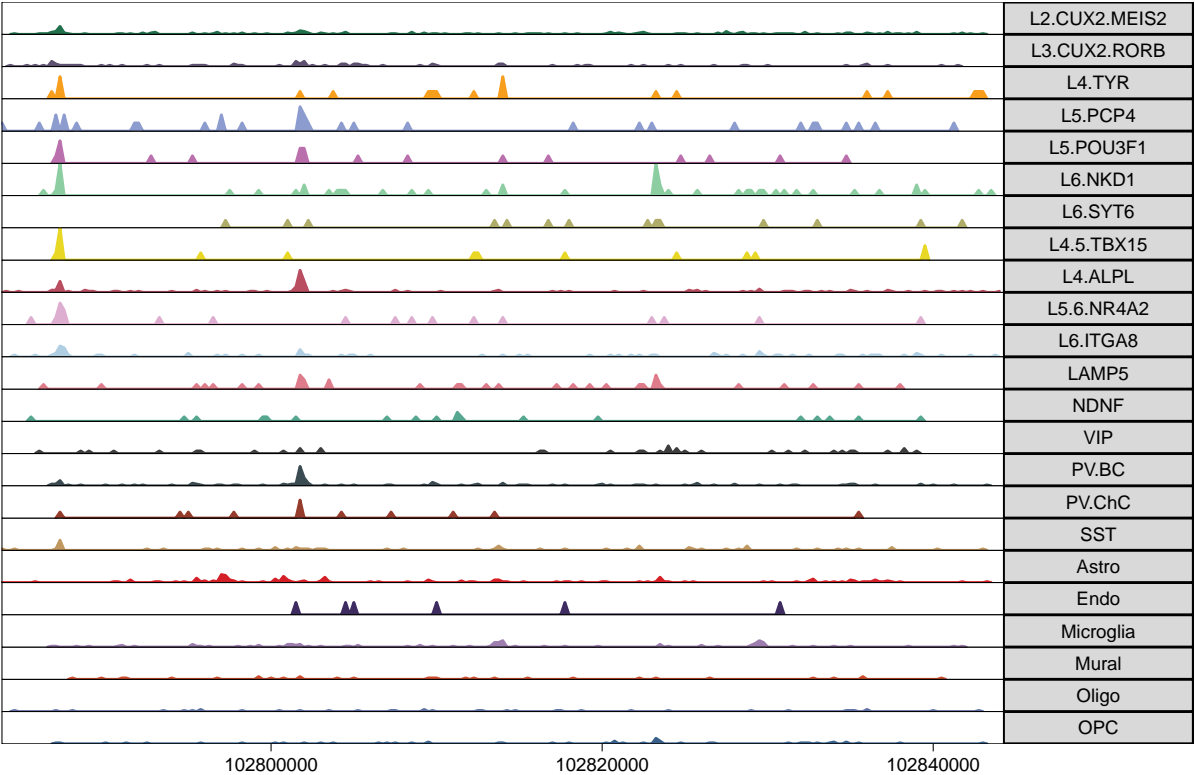


Peaks

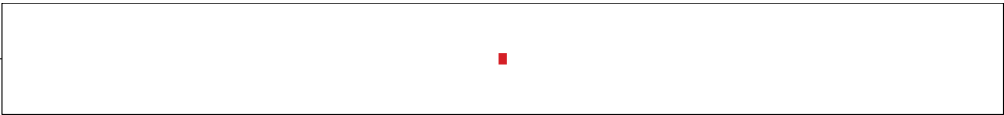
LoopTrack

GeneTrack

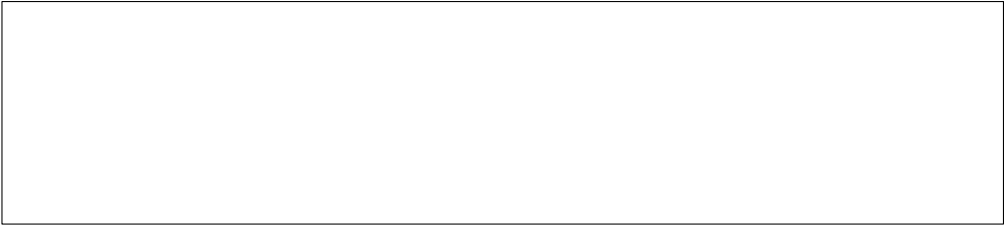
Coverage  
(Norm. ATAC Signal Range 0–1.02) by ReadsInTSS)



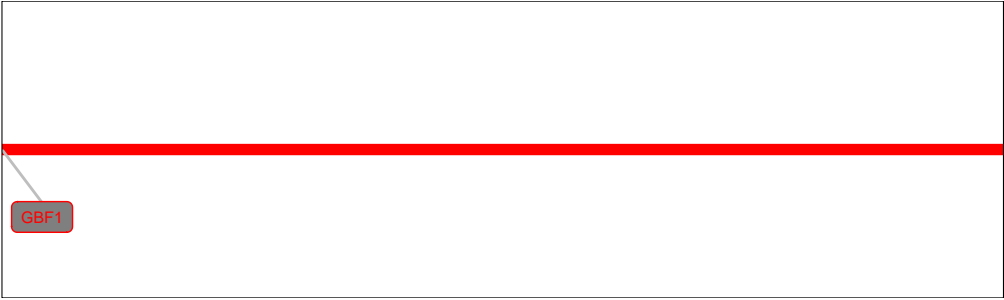
L3.CUX2.RORB



Peaks

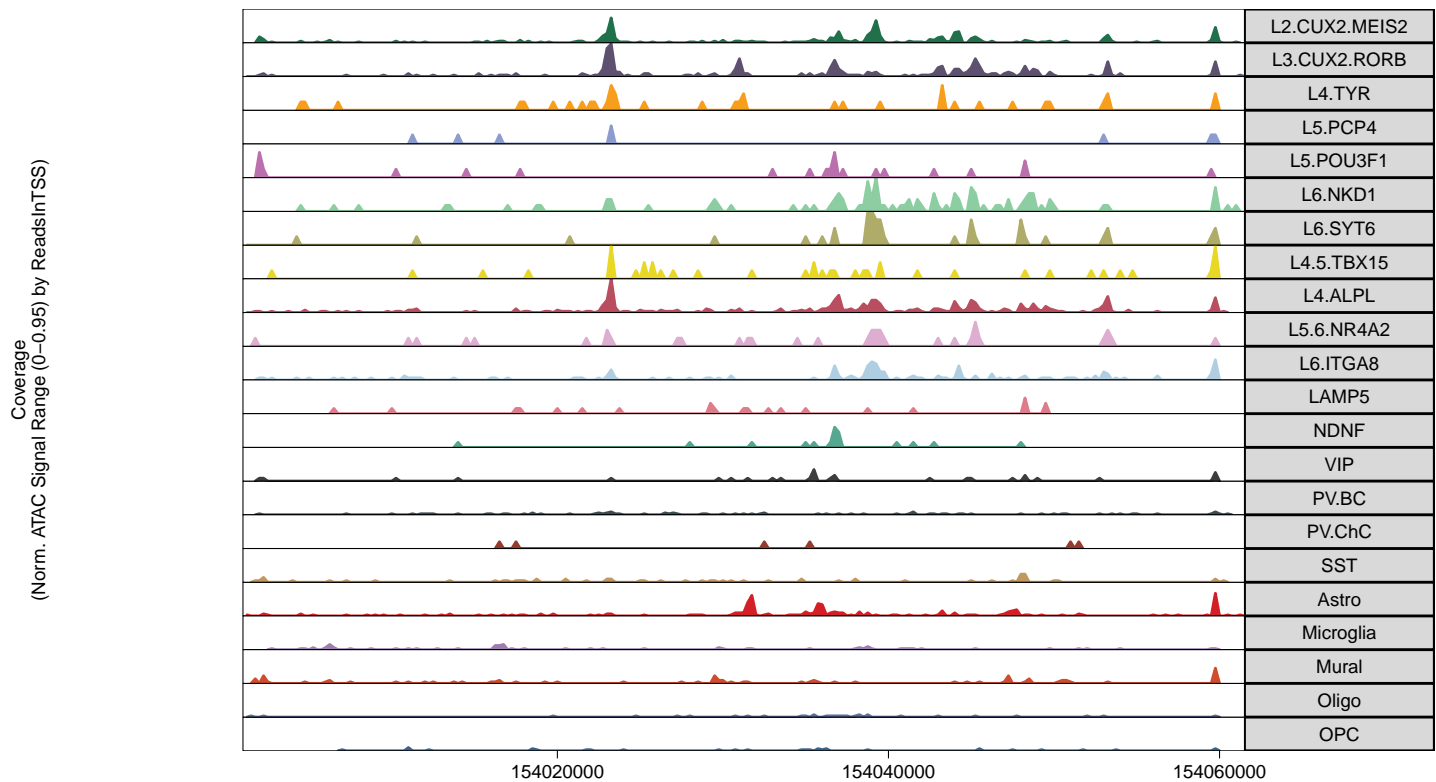


LoopTrack



Genes

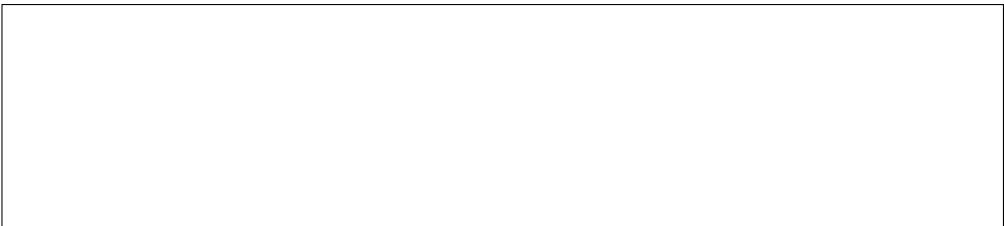
chr7:154001005–154061506 Marker Peak Tracks for: L3.CUX2.RORB



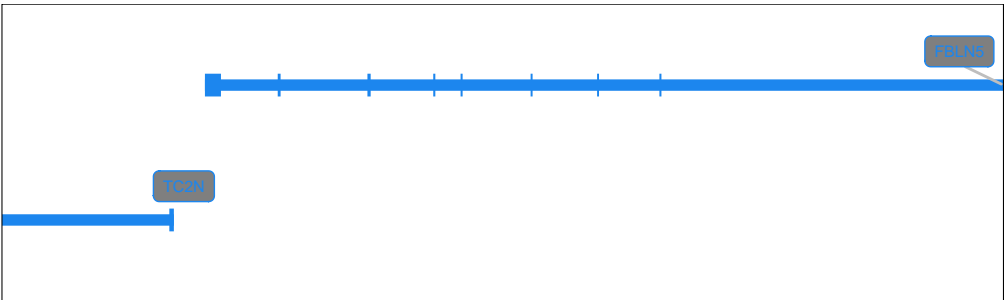
L3.CUX2.RORB



Peaks



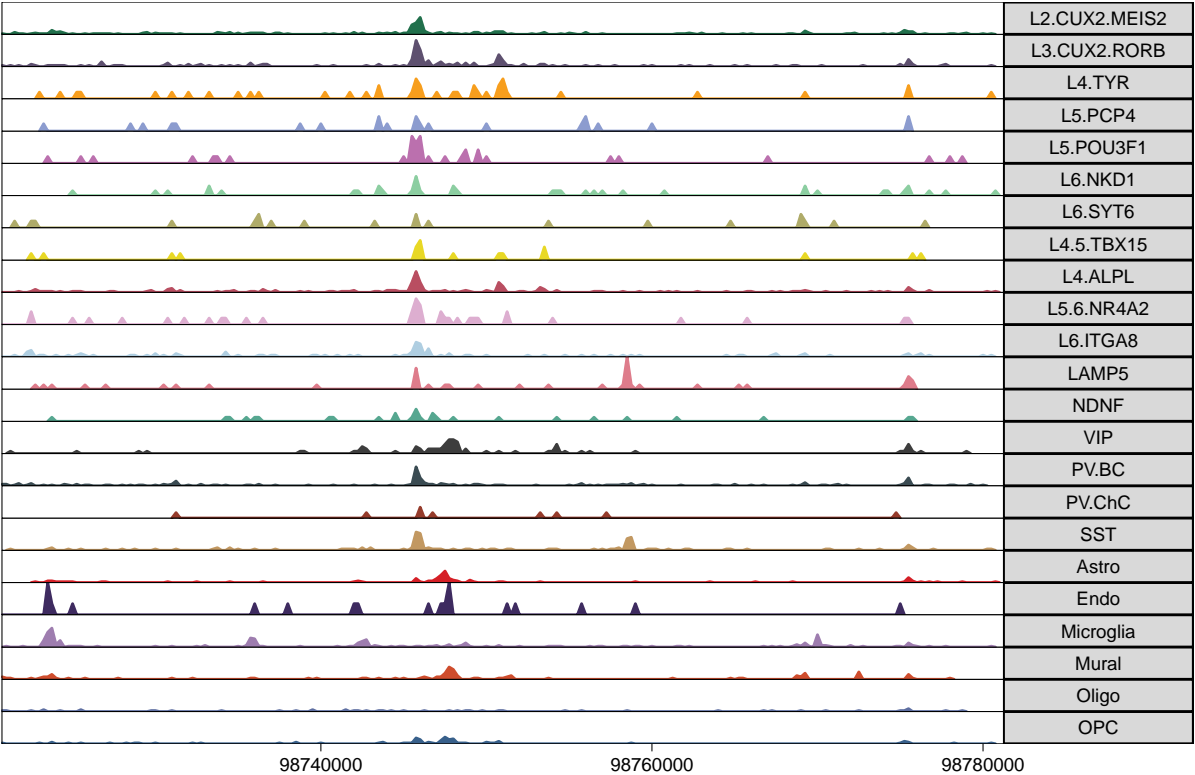
LoopTrack



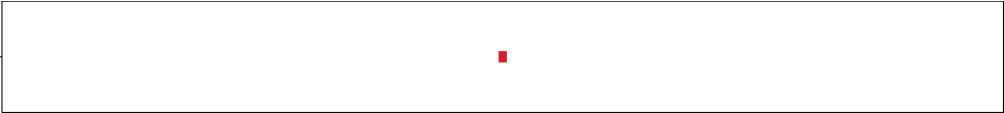
Genes

chr13:98720733–98781234 Marker Peak Tracks for: L3.CUX2.RORB

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



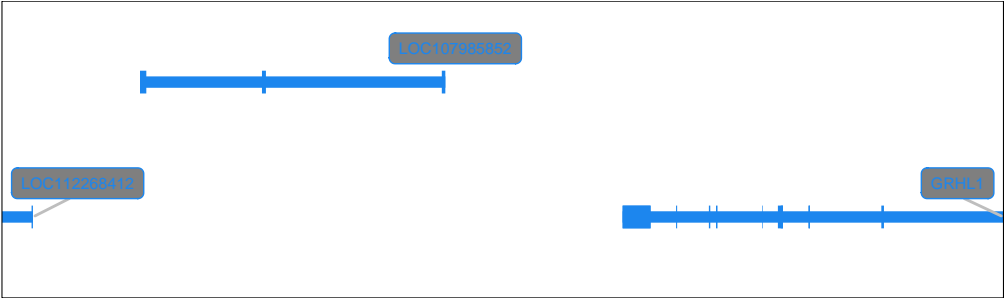
L3.CUX2.RORB



Peaks



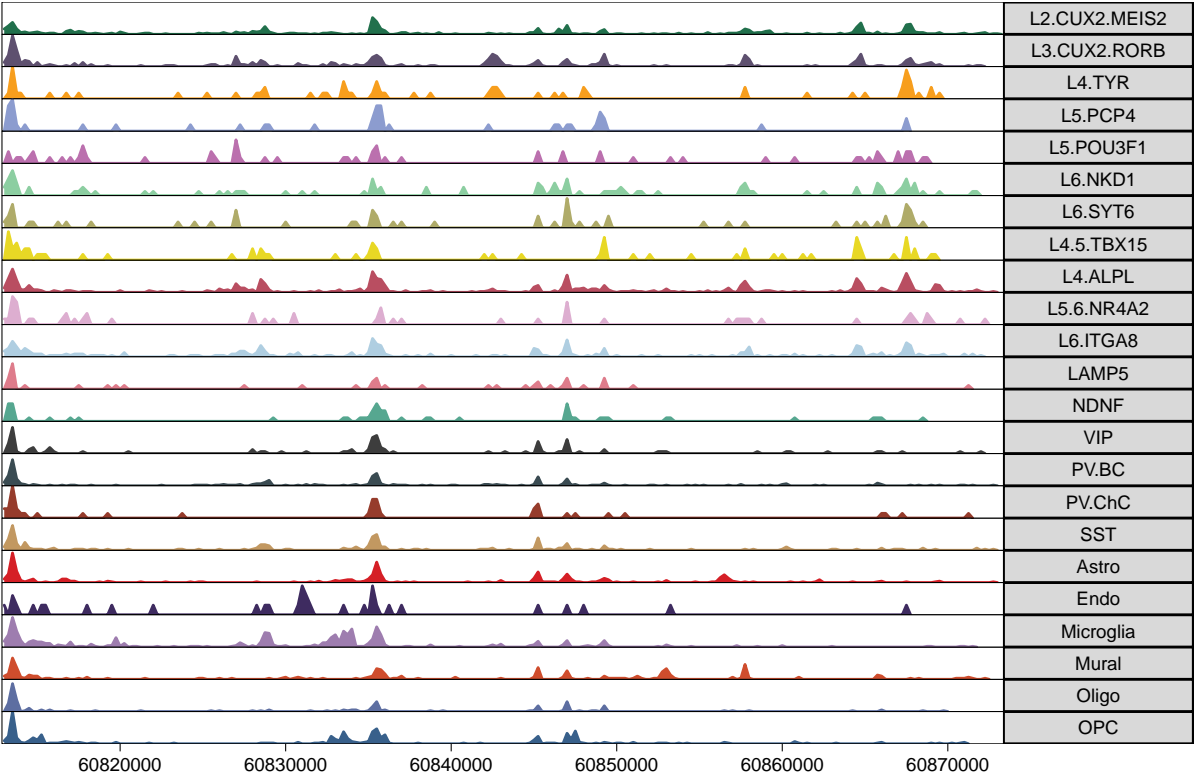
LoopTrack



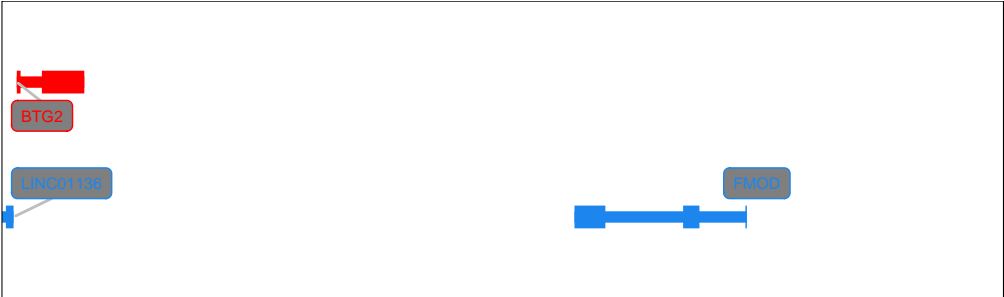
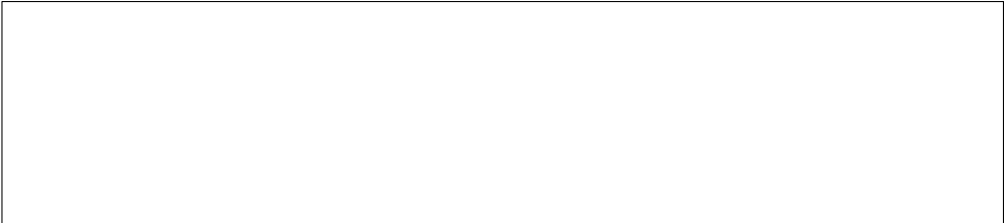
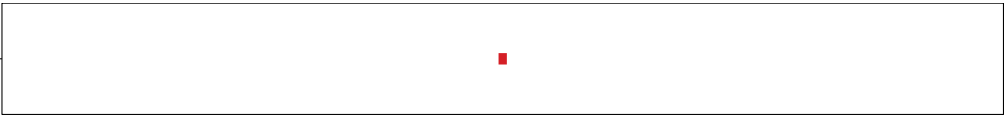
Genes

chr1:60812861–60873362 Marker Peak Tracks for: L3.CUX2.RORB

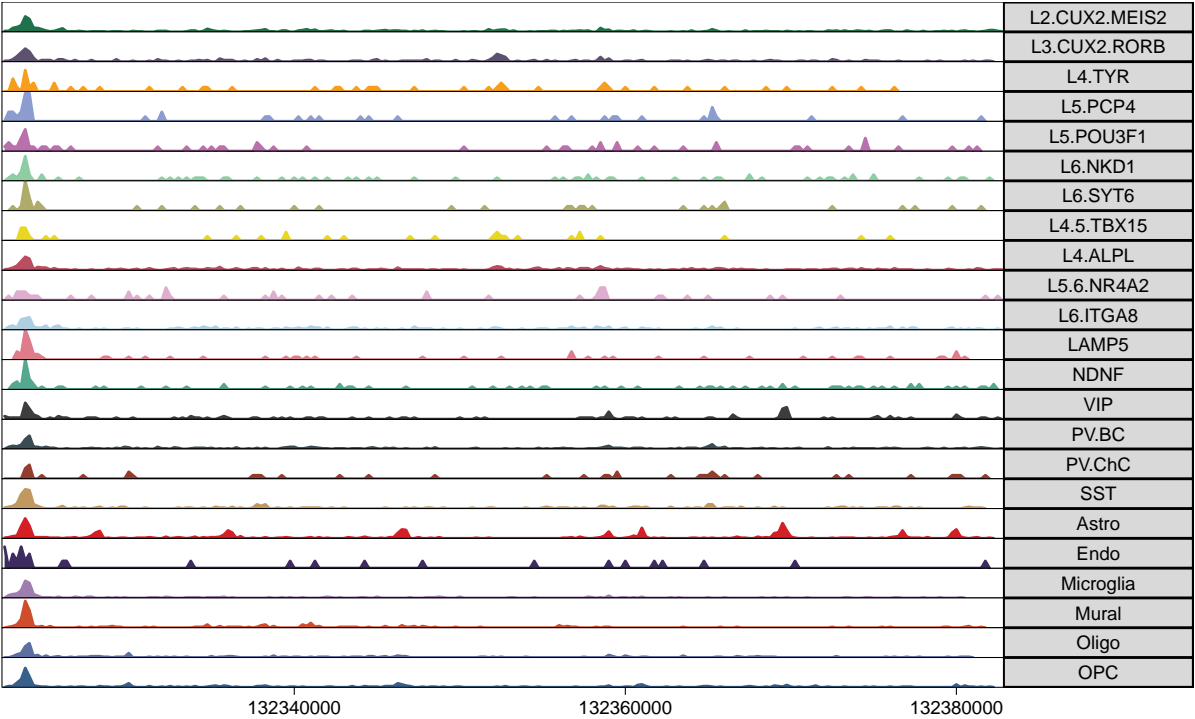
Coverage  
(Norm. ATAC Signal Range 0–1.31) by ReadsInTSS)



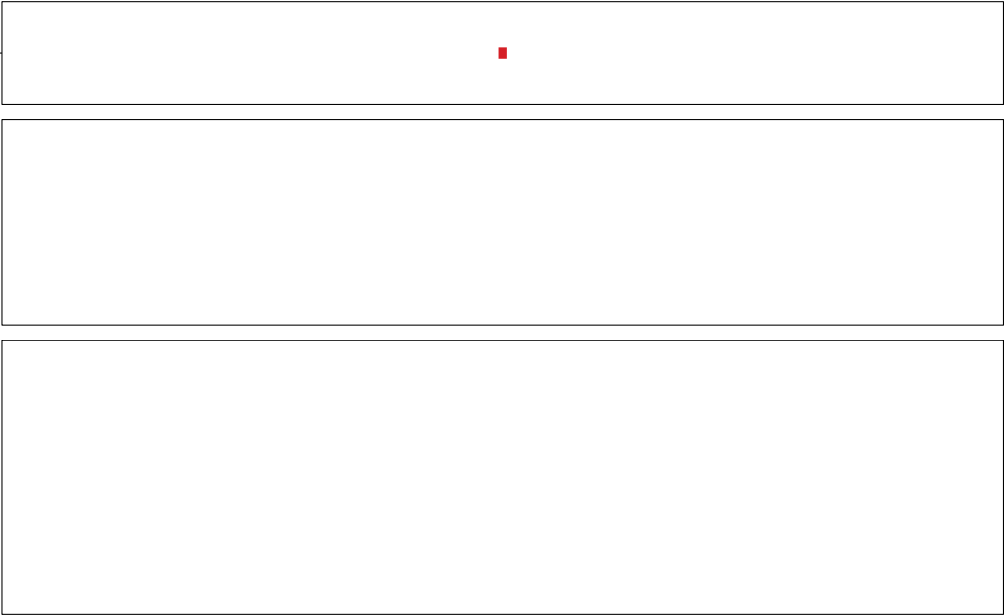
L3.CUX2.RORB



Coverage  
(Norm. ATAC Signal Range (0–1.63) by Reads in TSS)

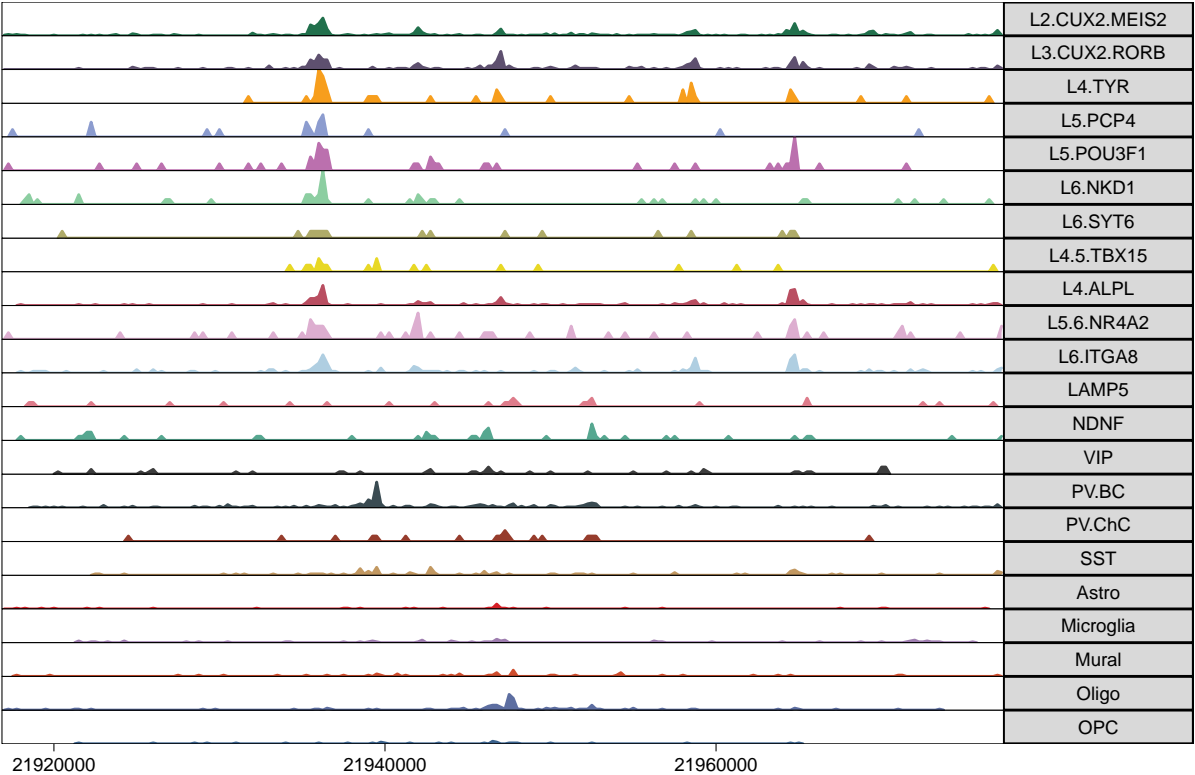


L3.CUX2.RORB

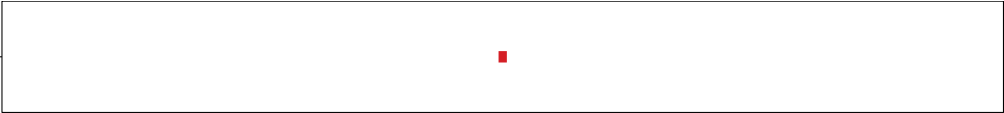


chr17:21916861–21977362 Marker Peak Tracks for: L3.CUX2.RORB

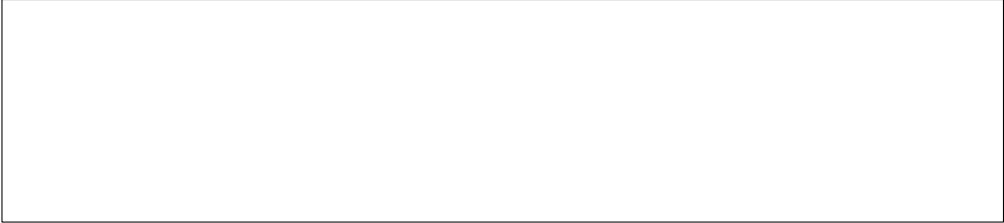
Coverage  
(Norm. ATAC Signal Range (0–1.17) by ReadsInTSS)



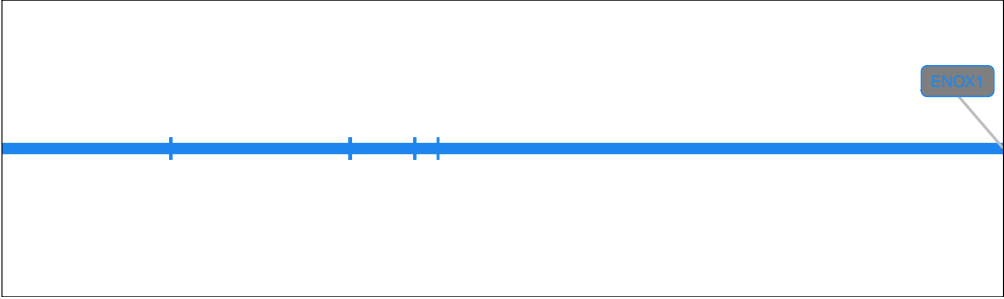
L3.CUX2.RORB



Peaks



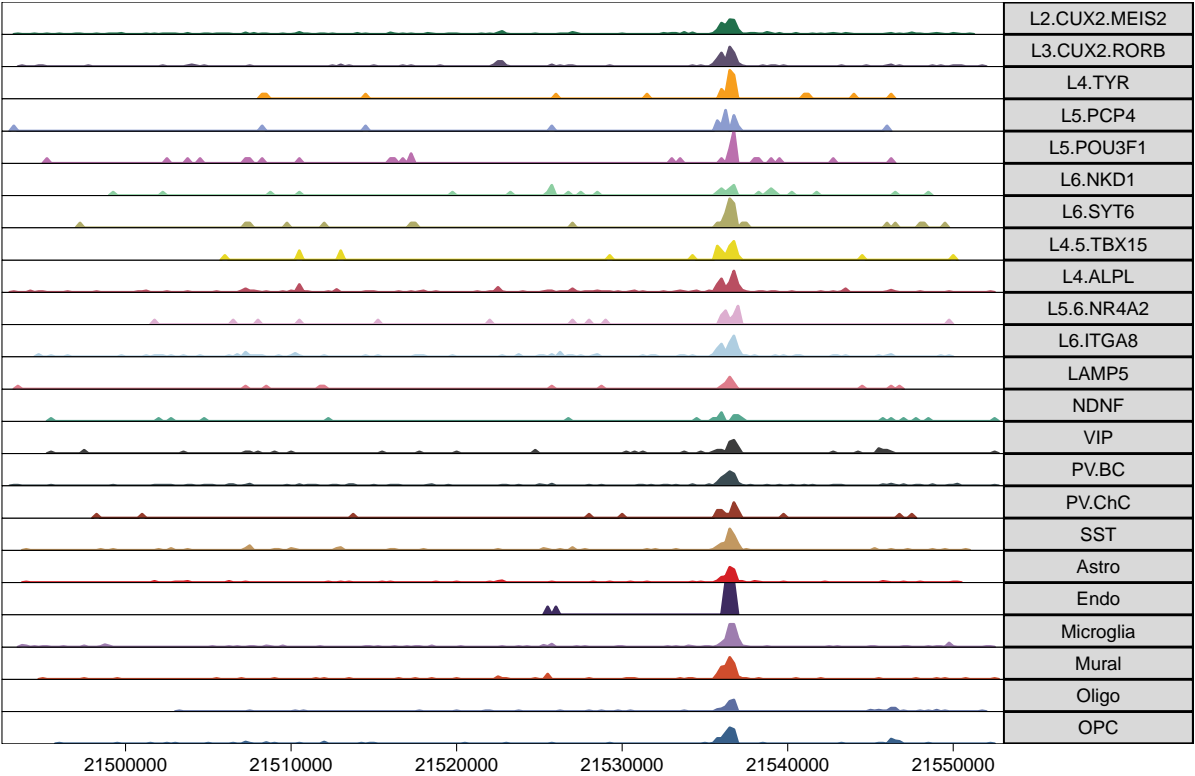
LoopTrack



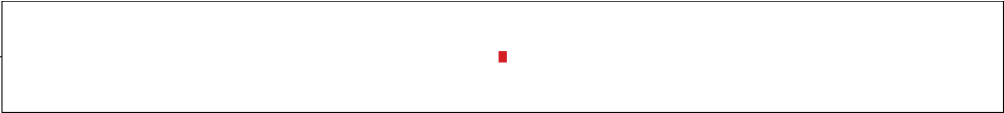
Genes



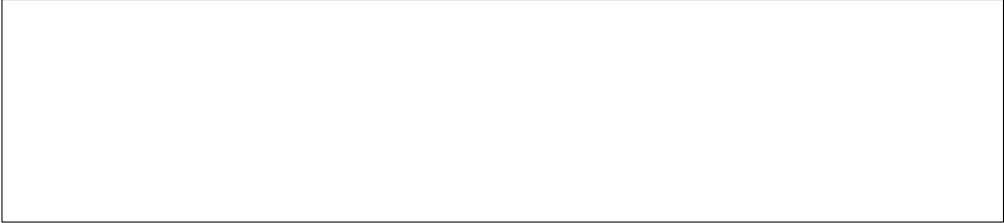
Coverage  
(Norm. ATAC Signal Range 0–1.59) by ReadsInTSS)



L3.CUX2.RORB



Peaks



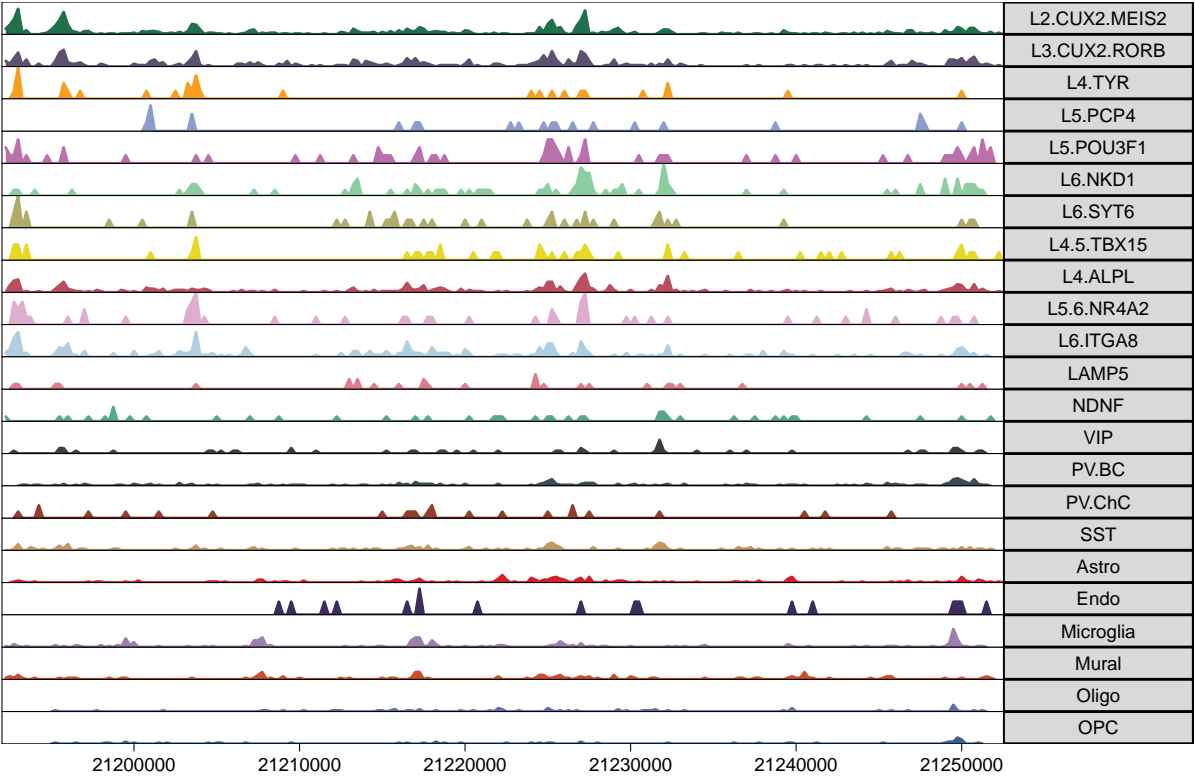
LoopTrack



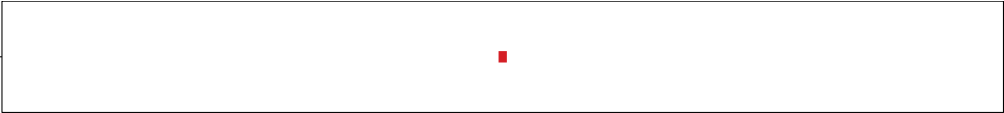
Genes

chr18:21192023–21252524 Marker Peak Tracks for: L3.CUX2.RORB

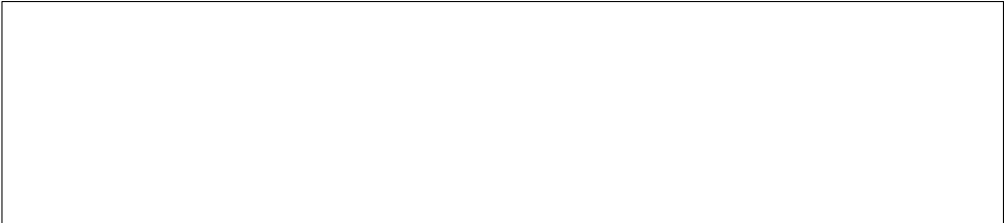
Coverage  
(Norm. ATAC Signal Range 0–0.97) by ReadsInTSS)



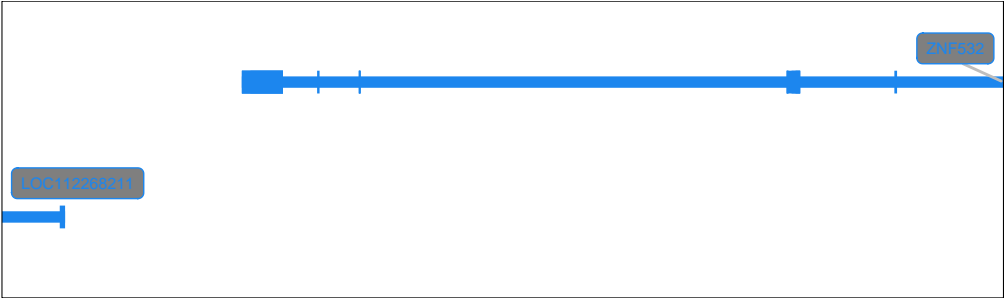
L3.CUX2.RORB



Peaks

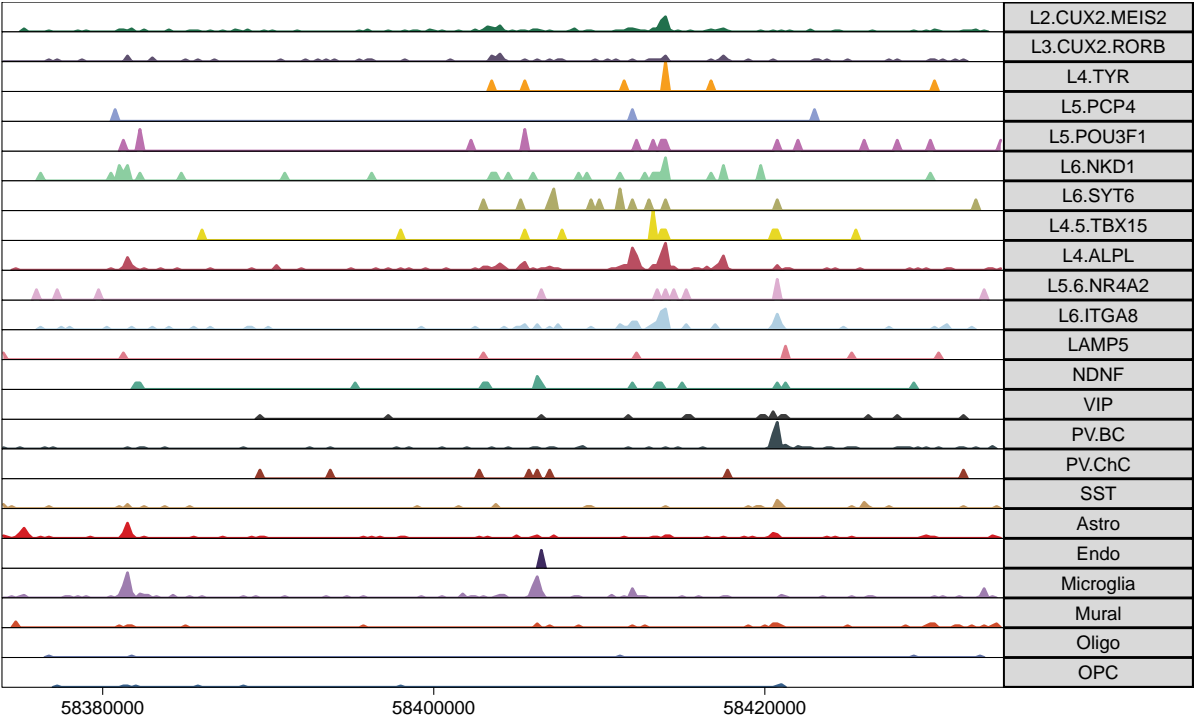


LoopTrack

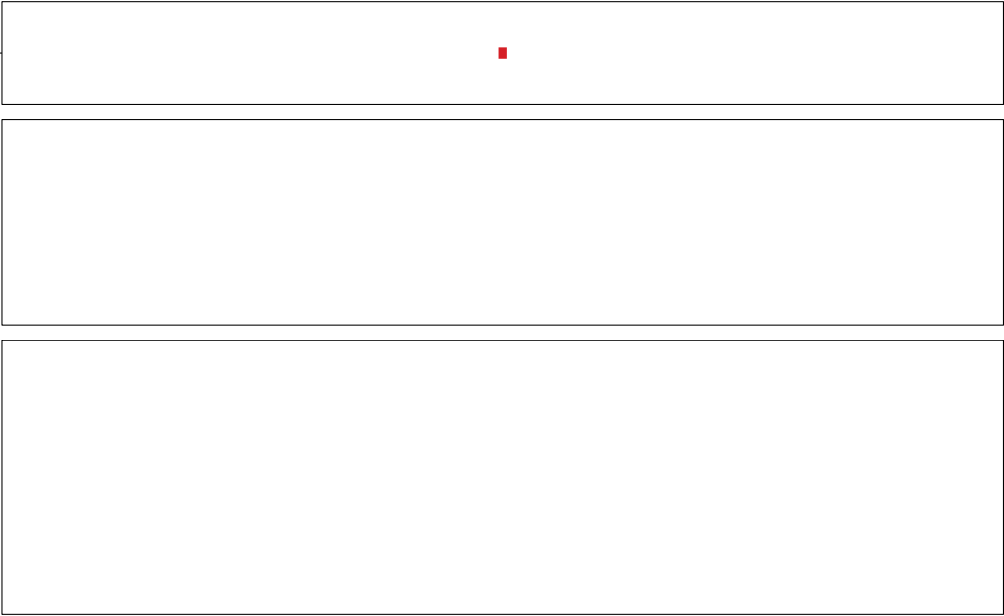


Genes

Coverage  
(Norm. ATAC Signal Range (0–0.65) by Reads in TSS)



L3.CUX2.RORB



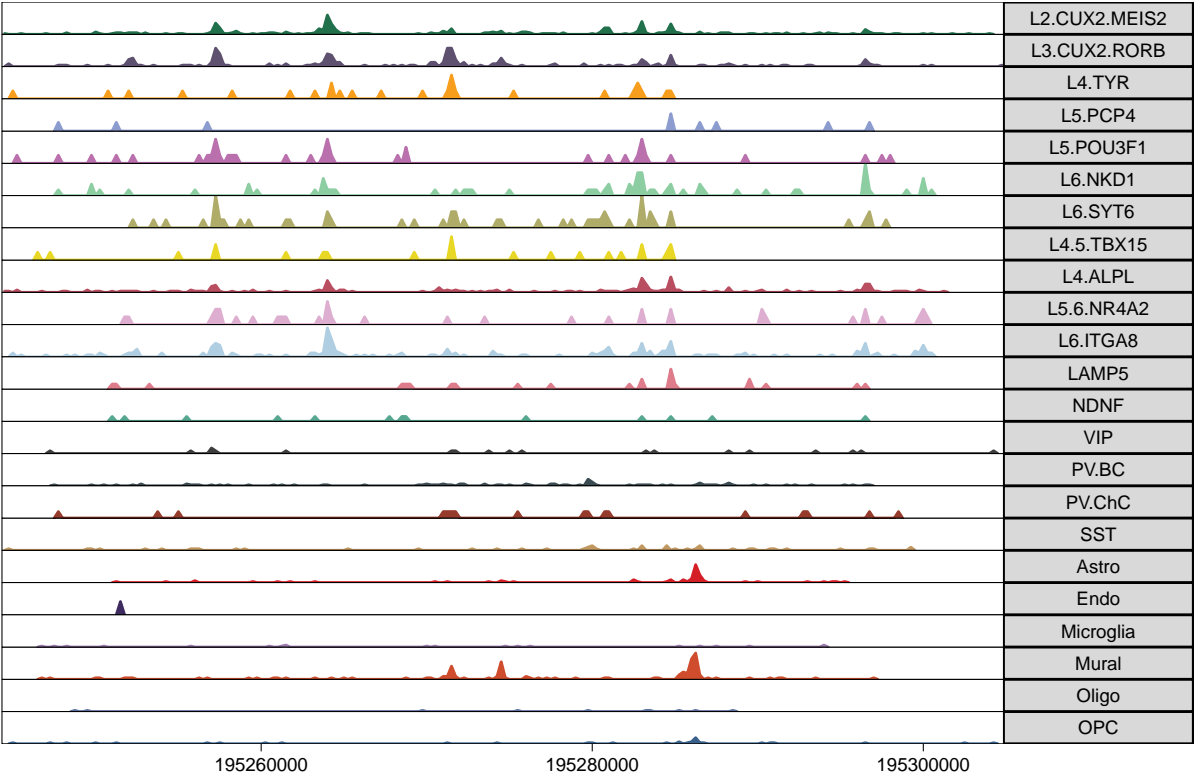
Peaks

LoopTrack

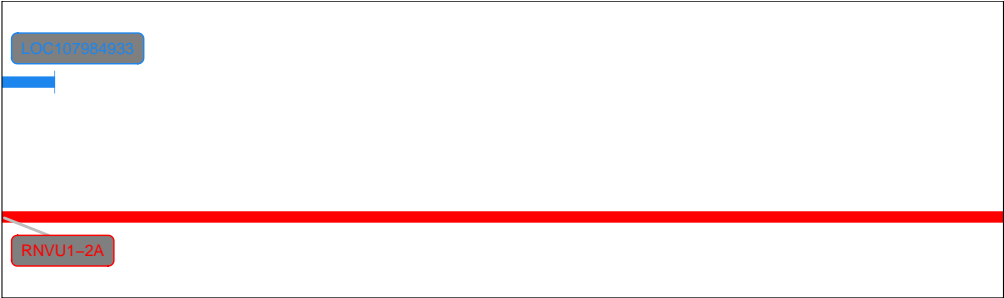
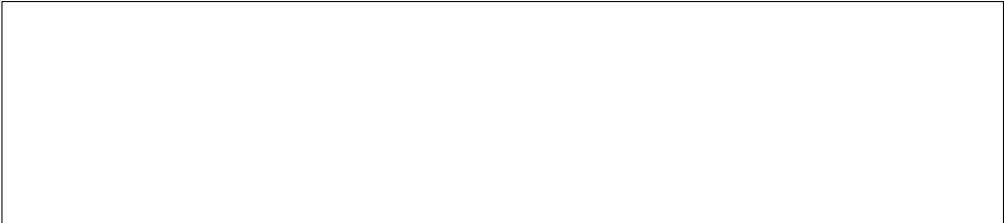
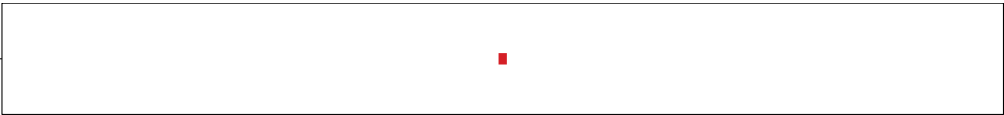
GeneTrack

chr1:195244341–195304842 Marker Peak Tracks for: L3.CUX2.RORB

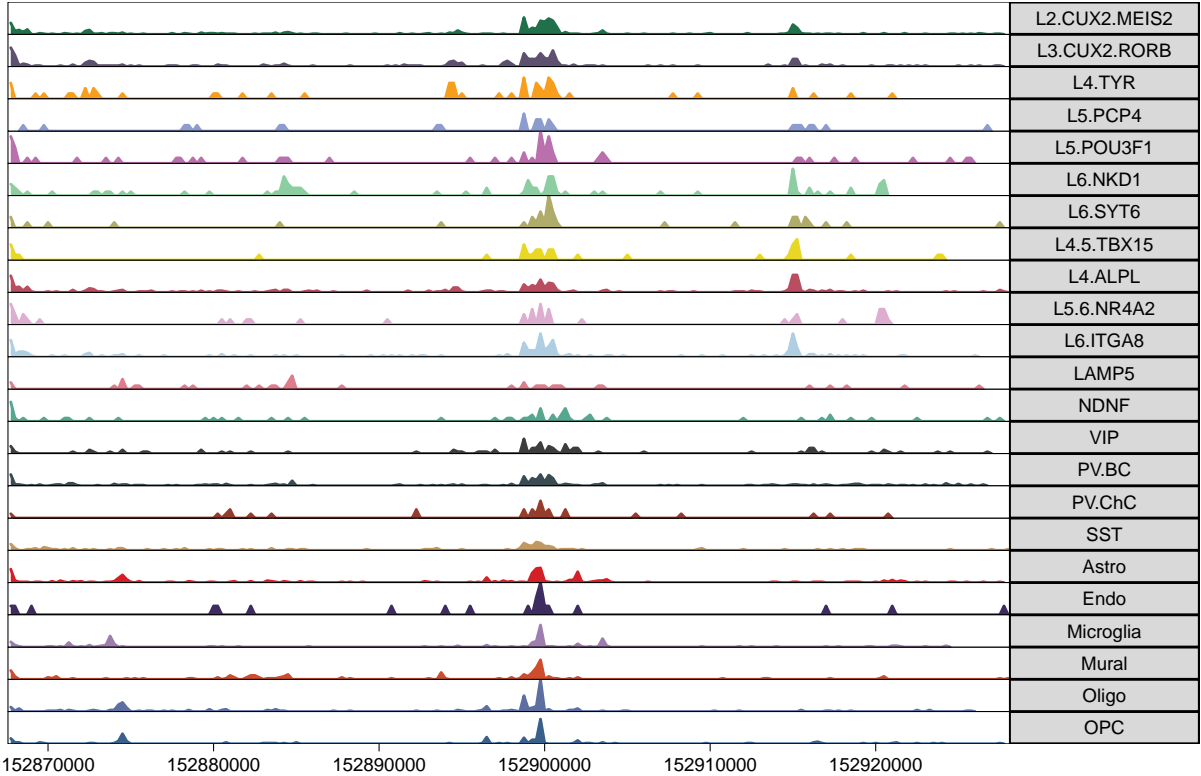
Coverage  
(Norm. ATAC Signal Range (0–0.95) by ReadsInTSS)



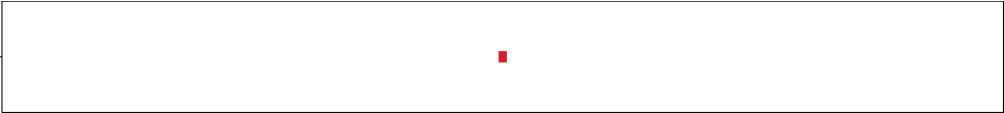
L3.CUX2.RORB



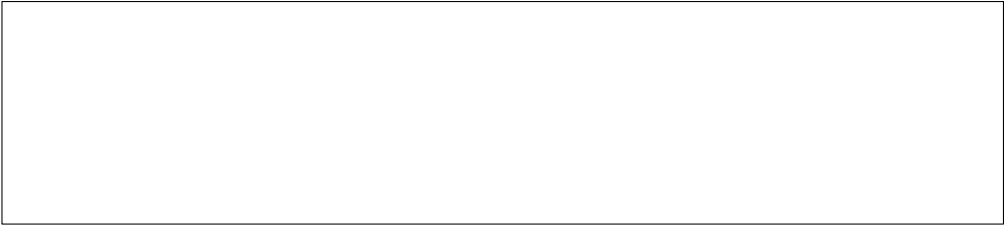
Coverage  
(Norm. ATAC Signal Range 0–1.43) by ReadsInTSS)



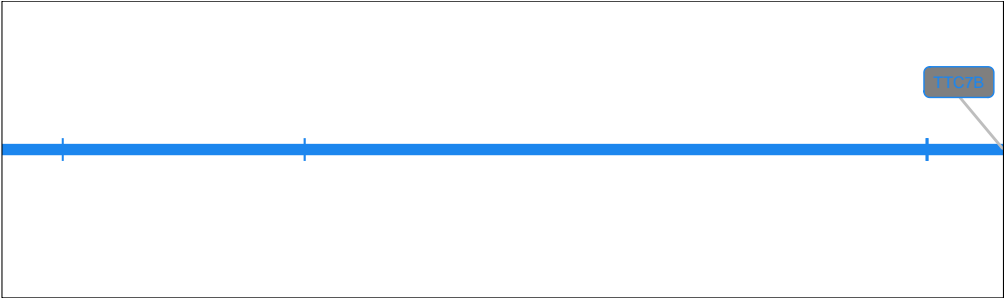
L3.CUX2.RORB



Peaks



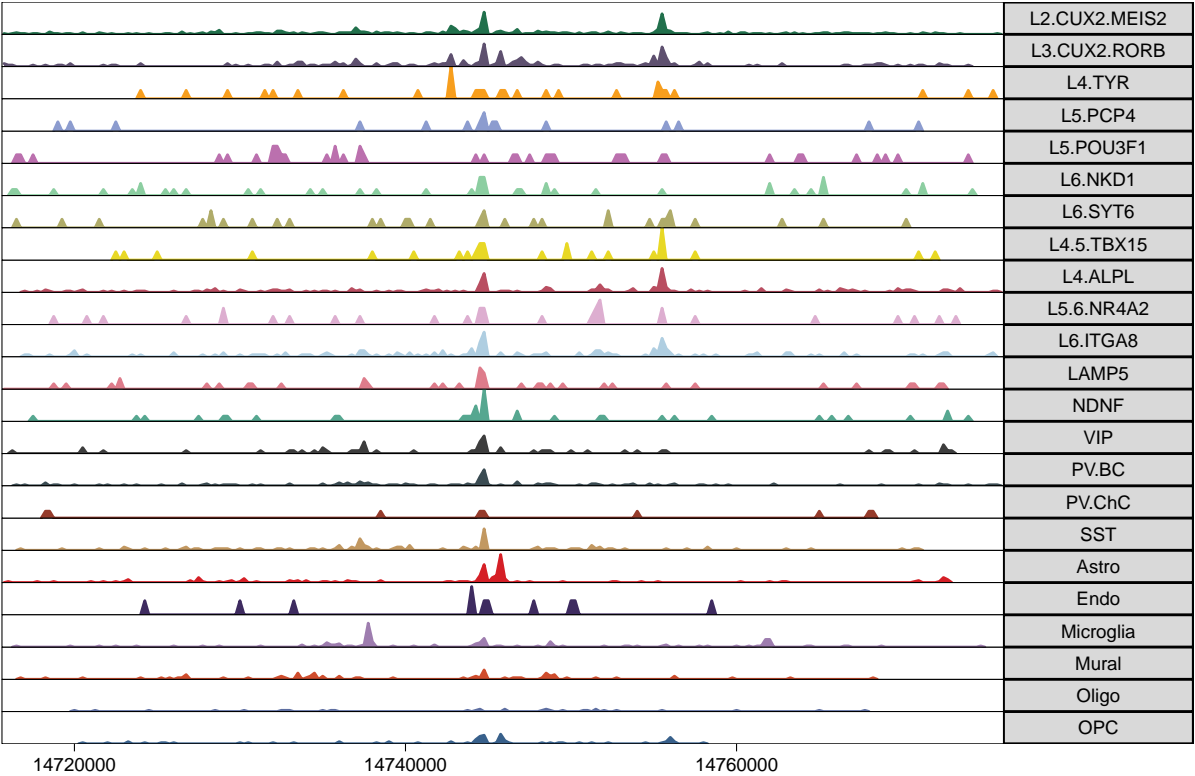
LoopTrack



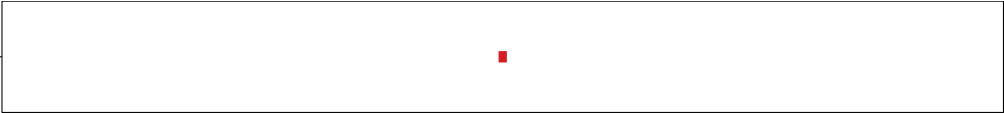
Genes

chr6:14715622–14776123 Marker Peak Tracks for: L3.CUX2.RORB

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



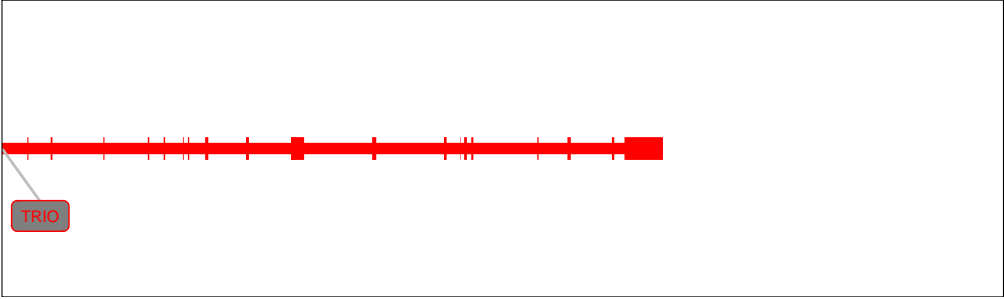
L3.CUX2.RORB



Peaks



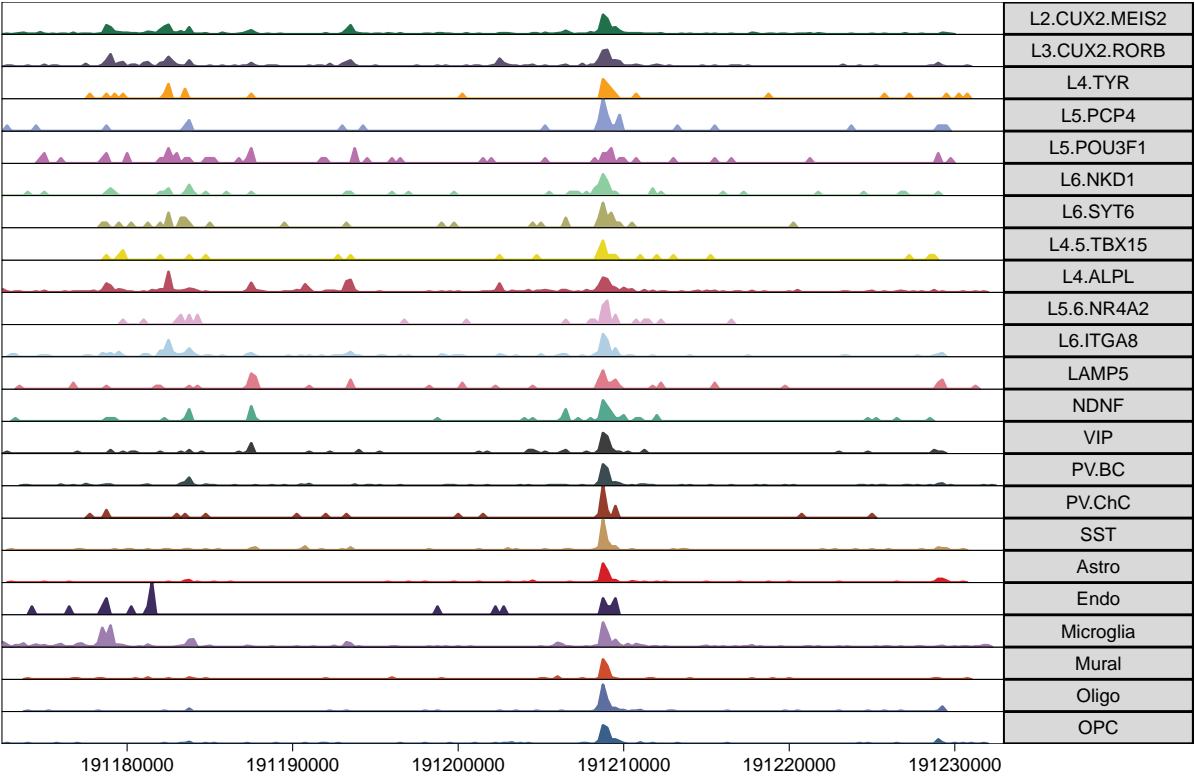
LoopTrack



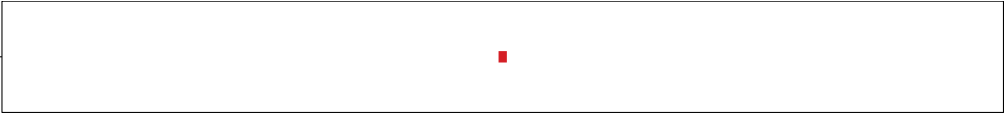
Genes

chr1:191172439–191232940 Marker Peak Tracks for: L3.CUX2.RORB

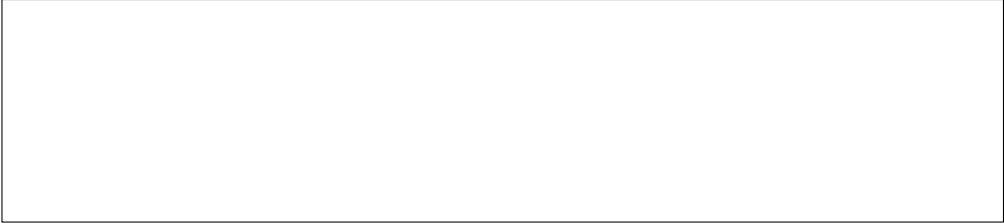
Coverage  
(Norm. ATAC Signal Range 0–1.54) by ReadsInTSS)



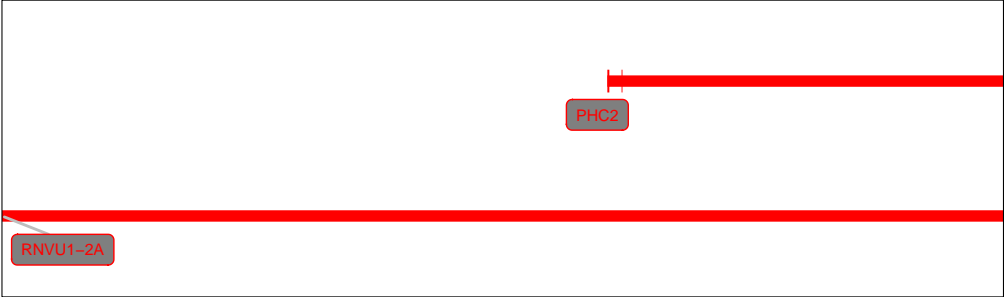
L3.CUX2.RORB



Peaks

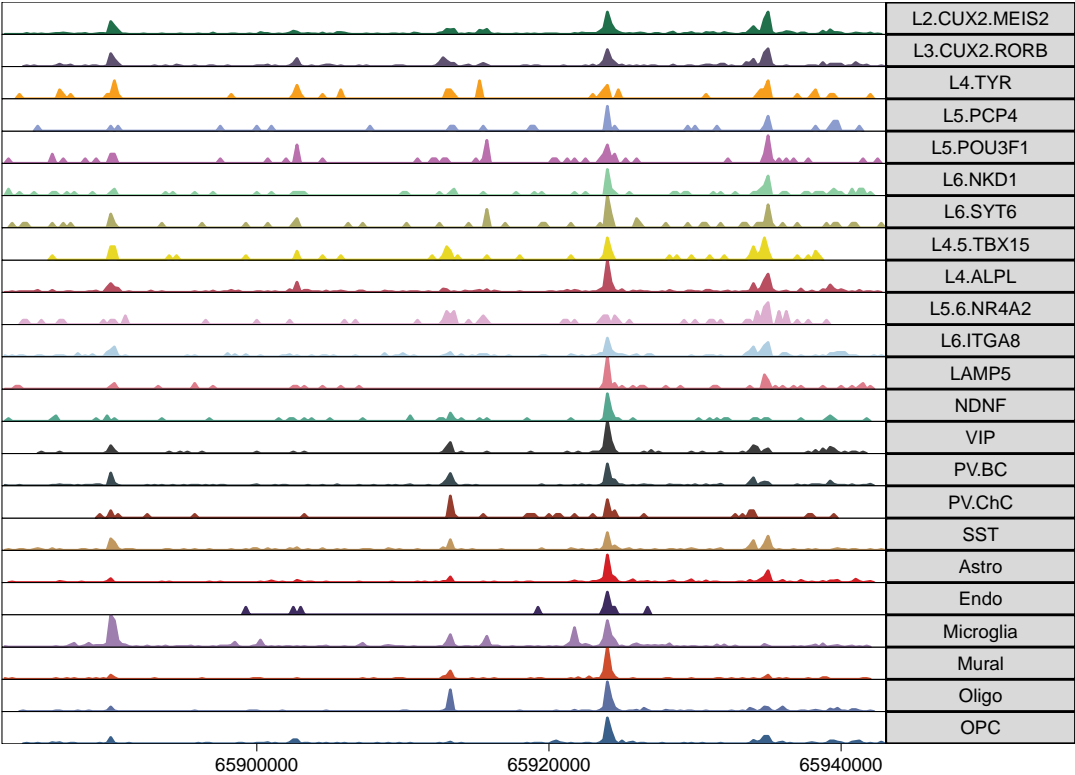


LoopTrack



Genes

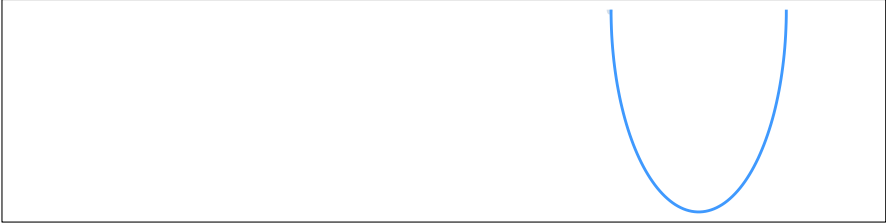
Coverage  
(Norm. ATAC Signal Range 0–1.67) by ReadsInTSS)



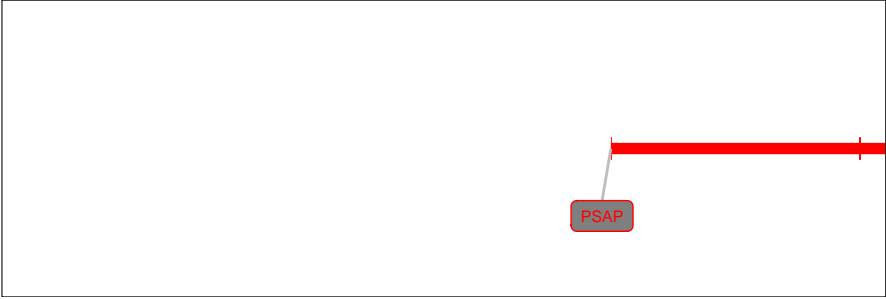
L3.CUX2.RORB



Peaks



Peak2GeneLinks



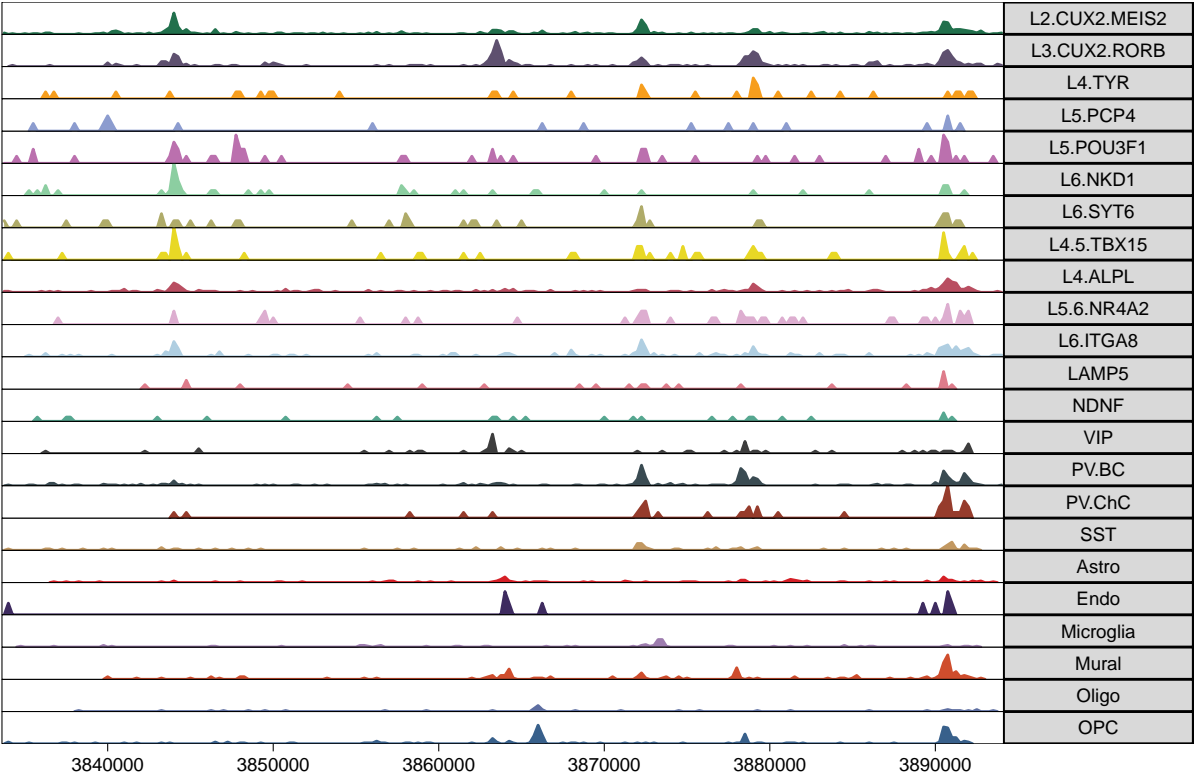
Genes



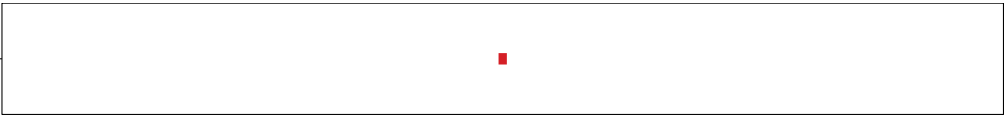


chr2:3833613–3894114 Marker Peak Tracks for: L3.CUX2.RORB

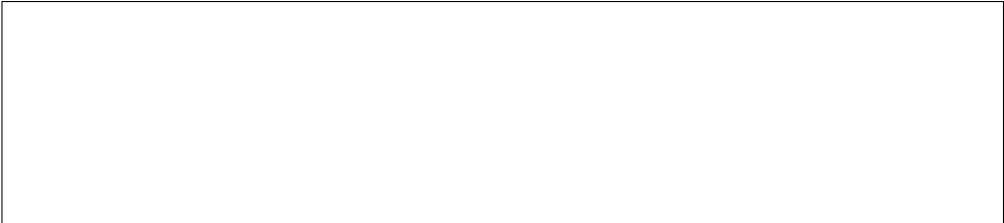
Coverage  
(Norm. ATAC Signal Range 0–1.08) by ReadsInTSS)



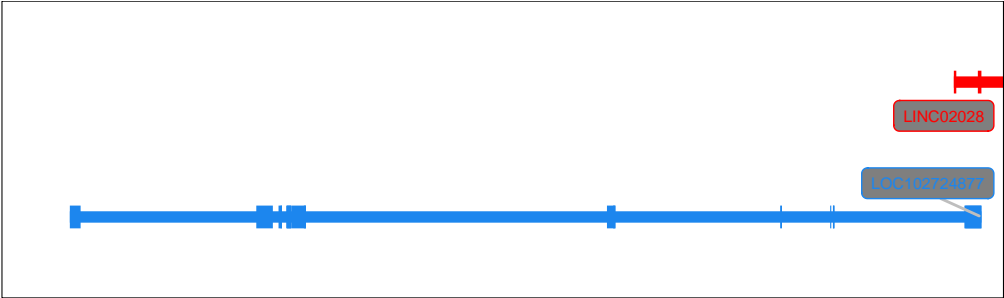
L3.CUX2.RORB



Peaks



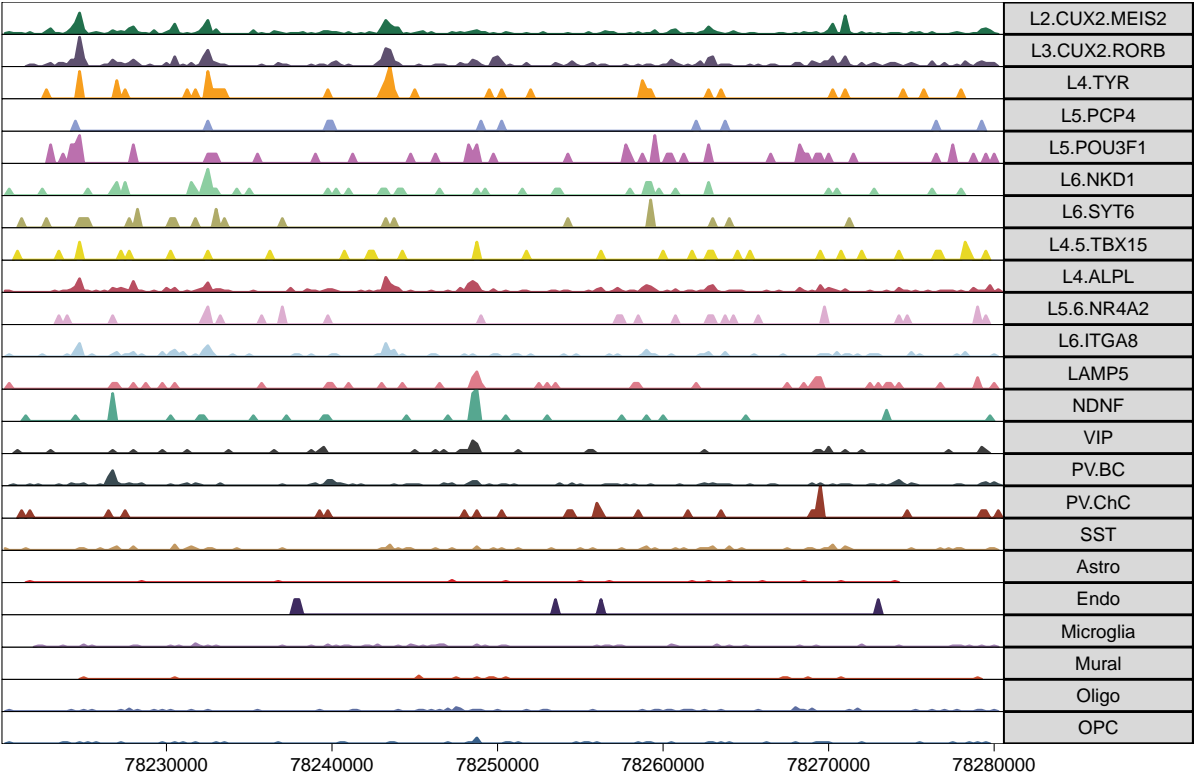
LoopTrack



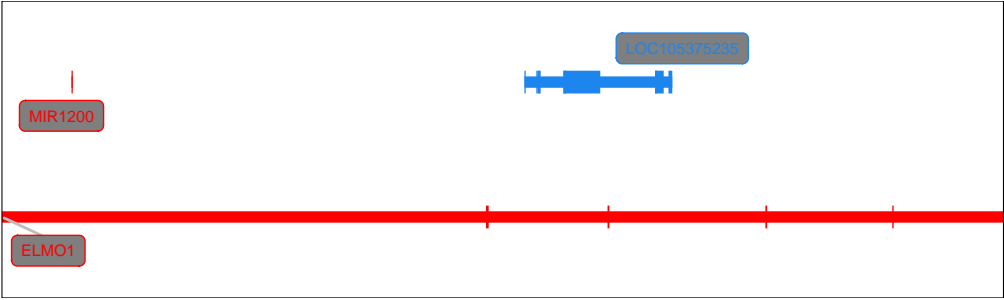
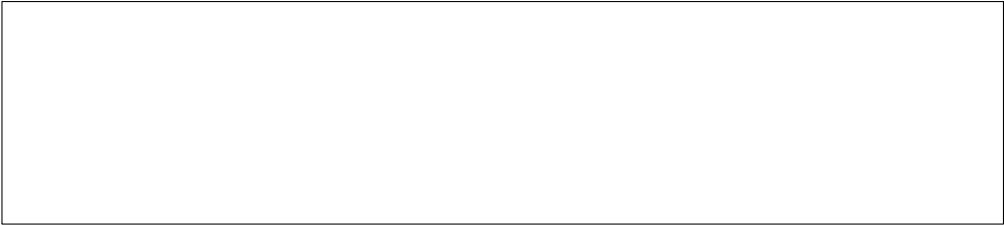
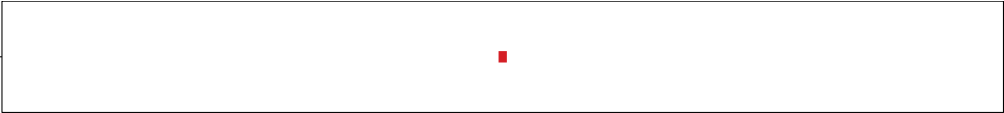
Genes

chr3:78220061–78280562 Marker Peak Tracks for: L3.CUX2.RORB

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

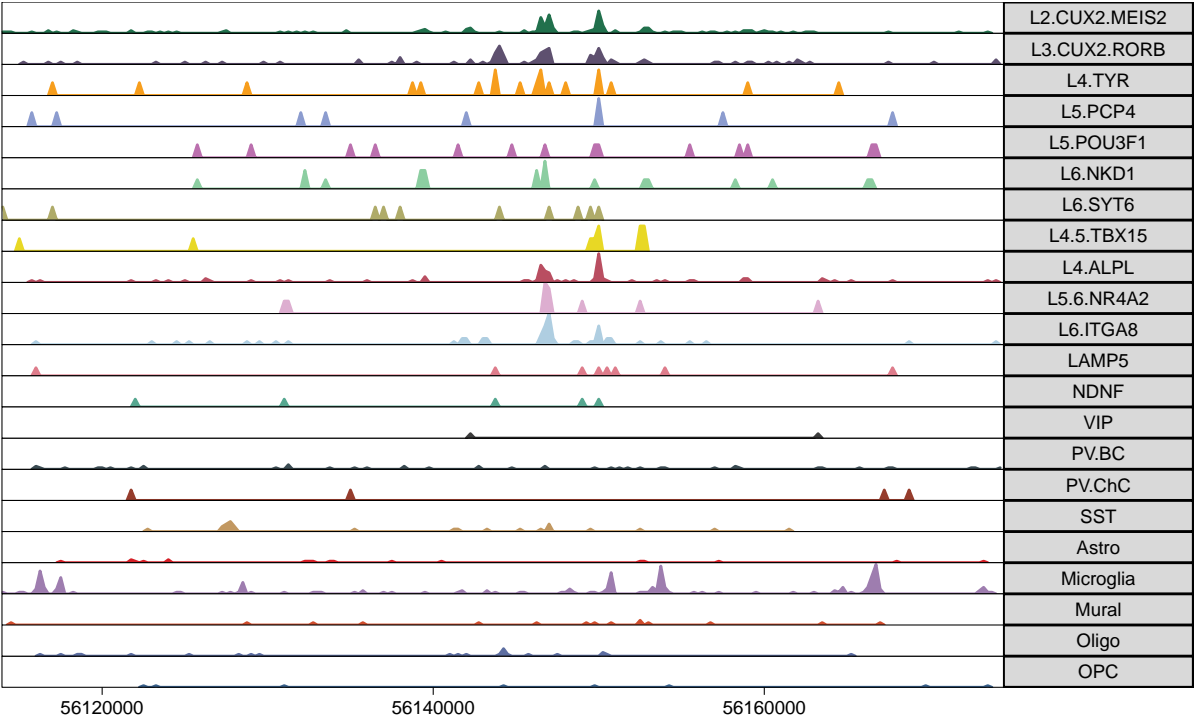


L3.CUX2.RORB

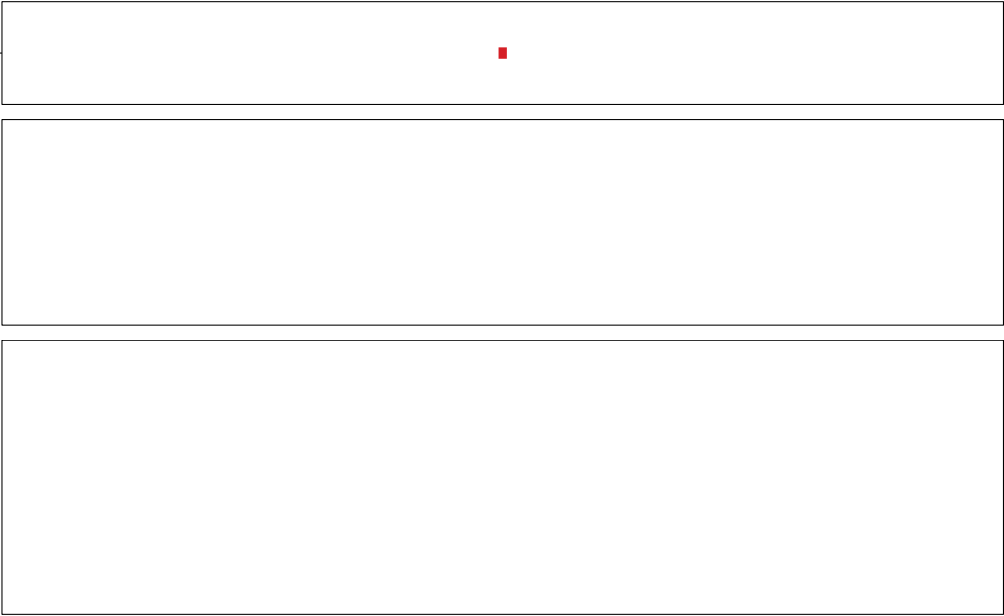


chr15:56113947–56174448 Marker Peak Tracks for: L3.CUX2.RORB

Coverage  
(Norm. ATAC Signal Range (0–0.57) by Reads in TSS)



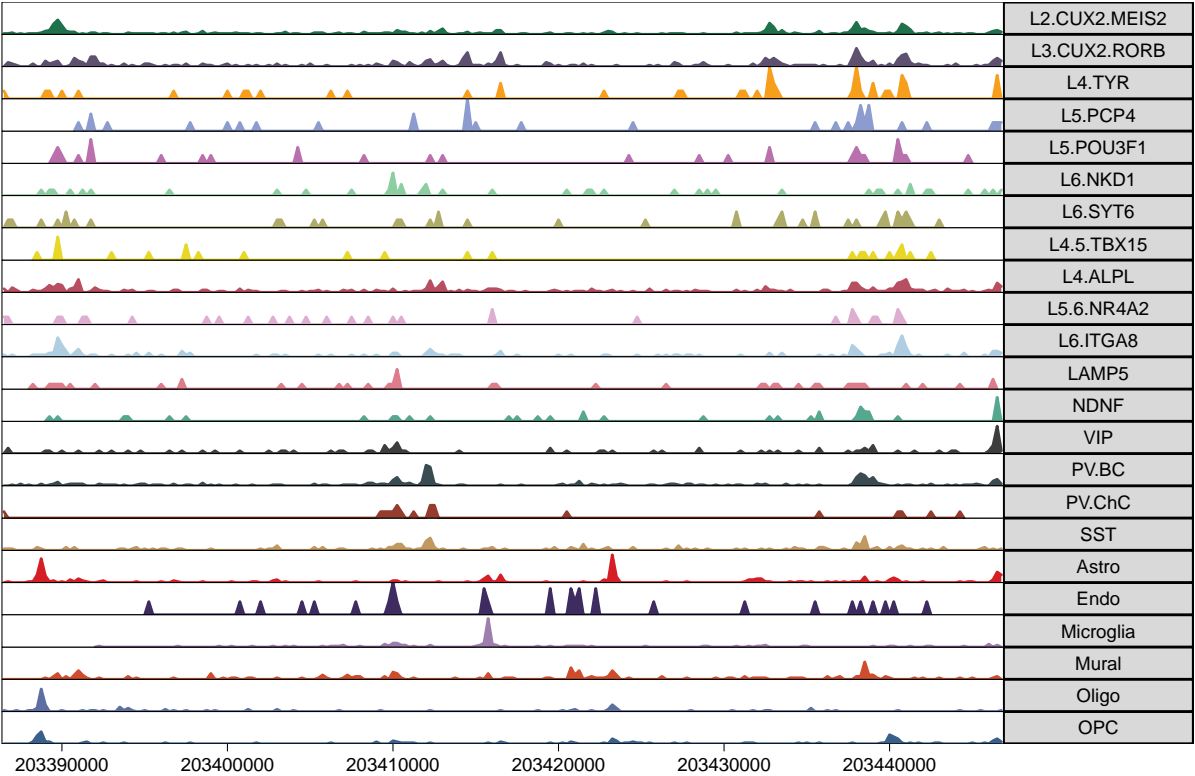
L3.CUX2.RORB



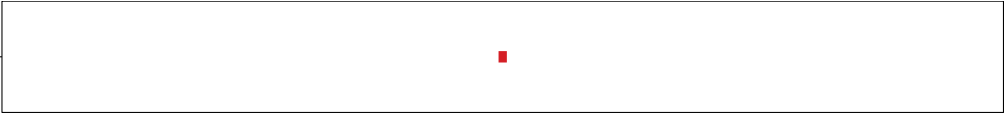
strand ● ·

chr1:203386378–203446879 Marker Peak Tracks for: L3.CUX2.RORB

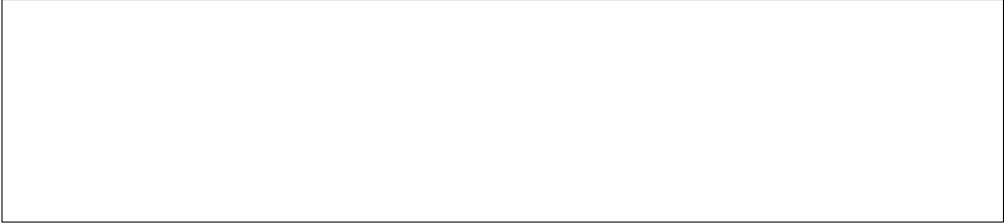
Coverage  
(Norm. ATAC Signal Range (0–0.96) by ReadsInTSS)



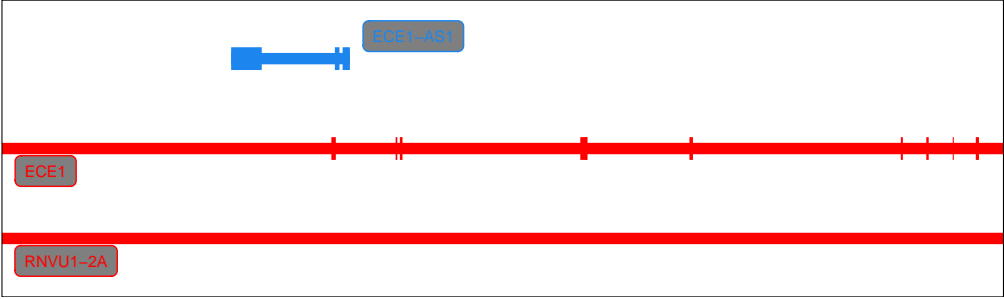
L3.CUX2.RORB



Peaks

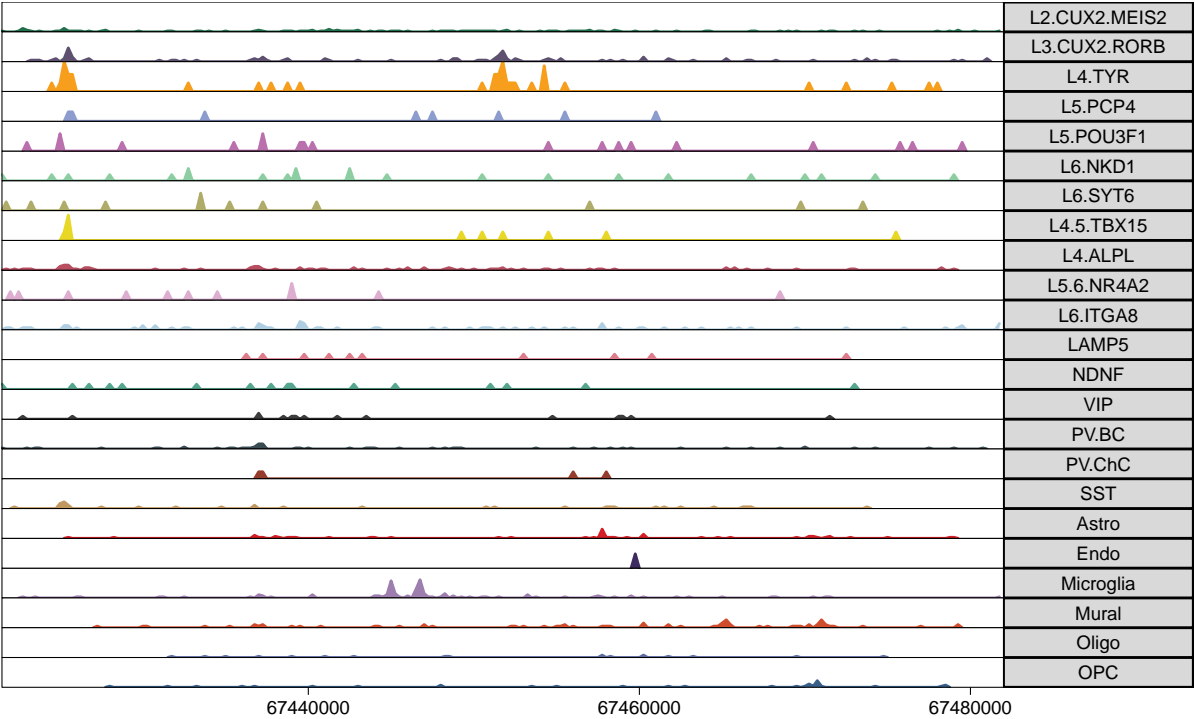


LoopTrack

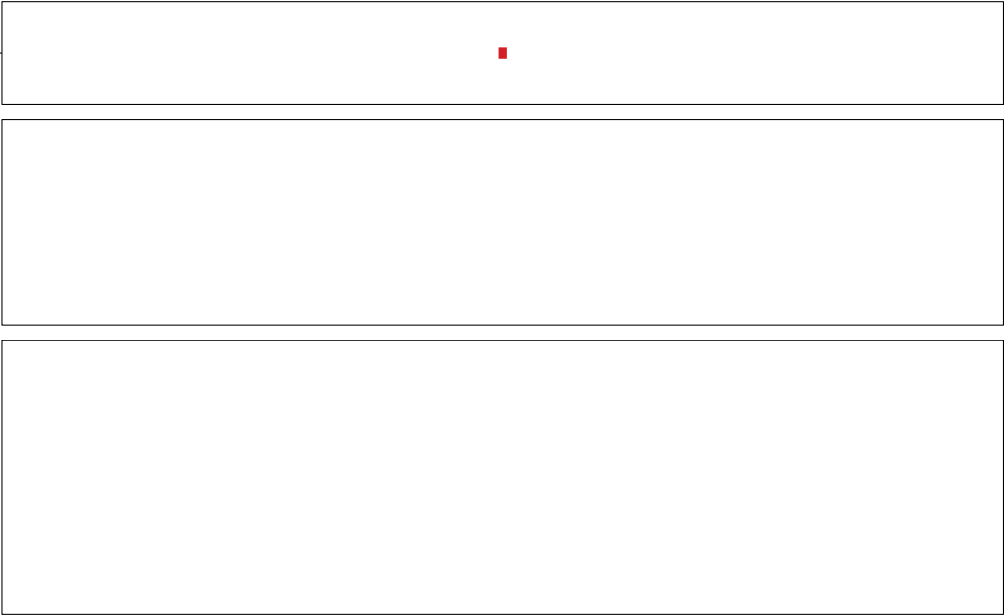


Genes

Coverage  
(Norm. ATAC Signal Range (0–0.81) by Reads in TSS)



L3.CUX2.RORB

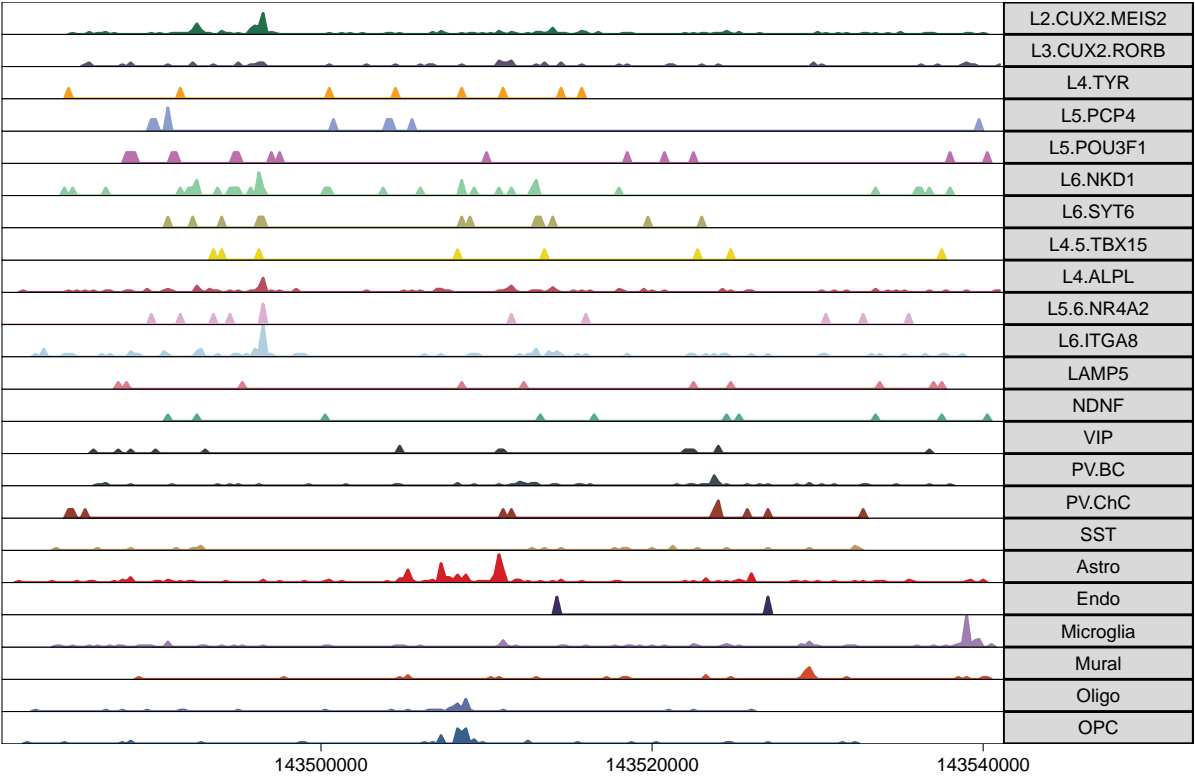


Peaks

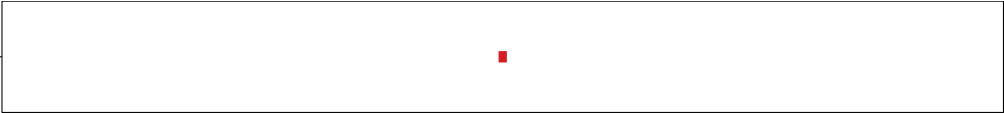
LoopTrack

GeneTrack

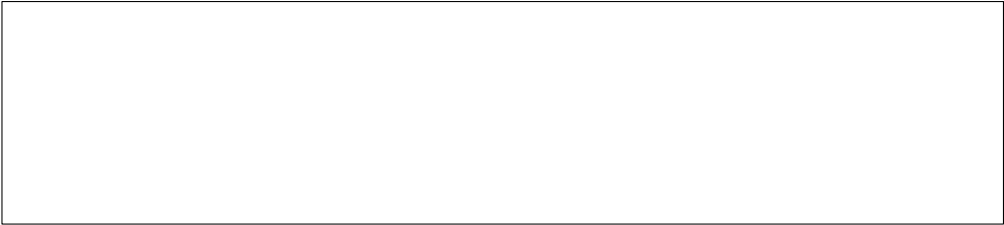
Coverage  
(Norm. ATAC Signal Range (0–0.71) by ReadsInTSS)



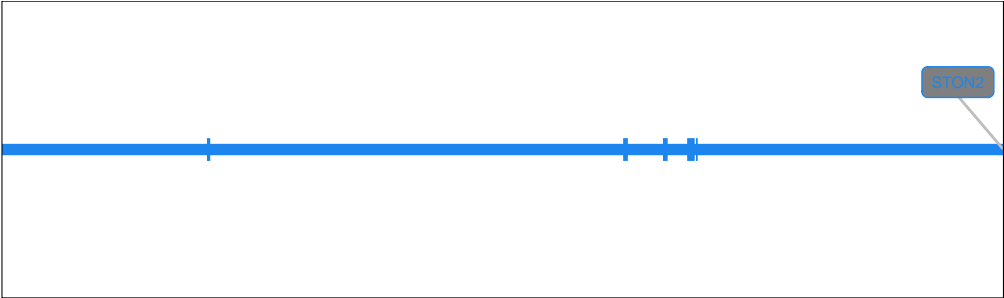
L3.CUX2.RORB



Peaks



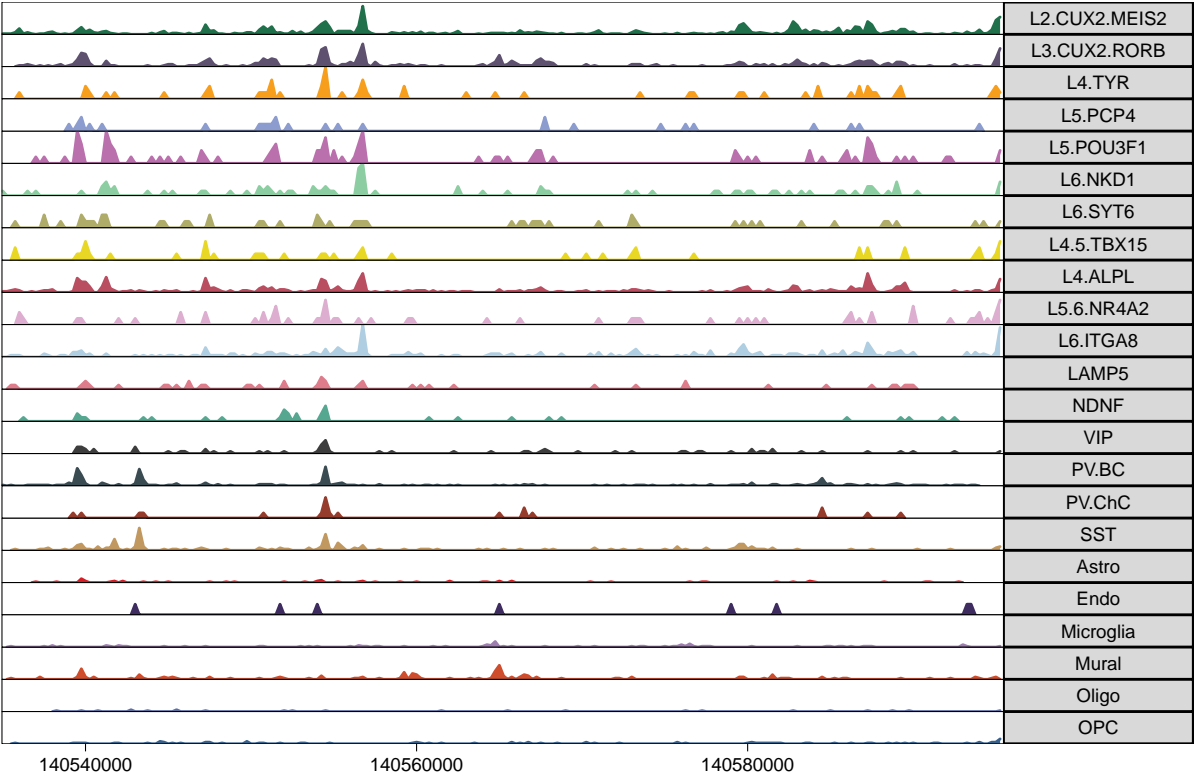
LoopTrack



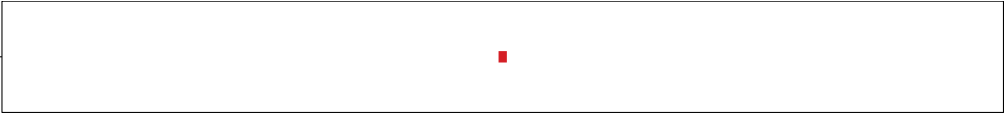
Genes

chr6:140534952–140595453 Marker Peak Tracks for: L3.CUX2.RORB

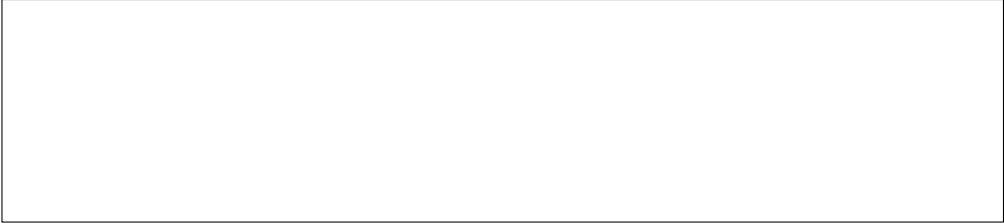
Coverage  
(Norm. ATAC Signal Range (0–1.19) by ReadsInTSS)



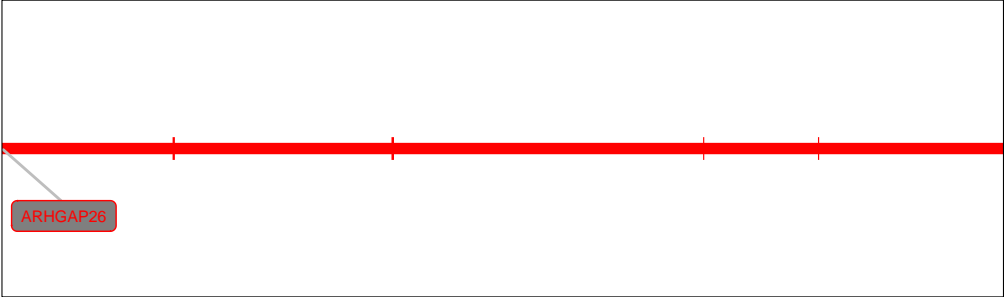
L3.CUX2.RORB



Peaks

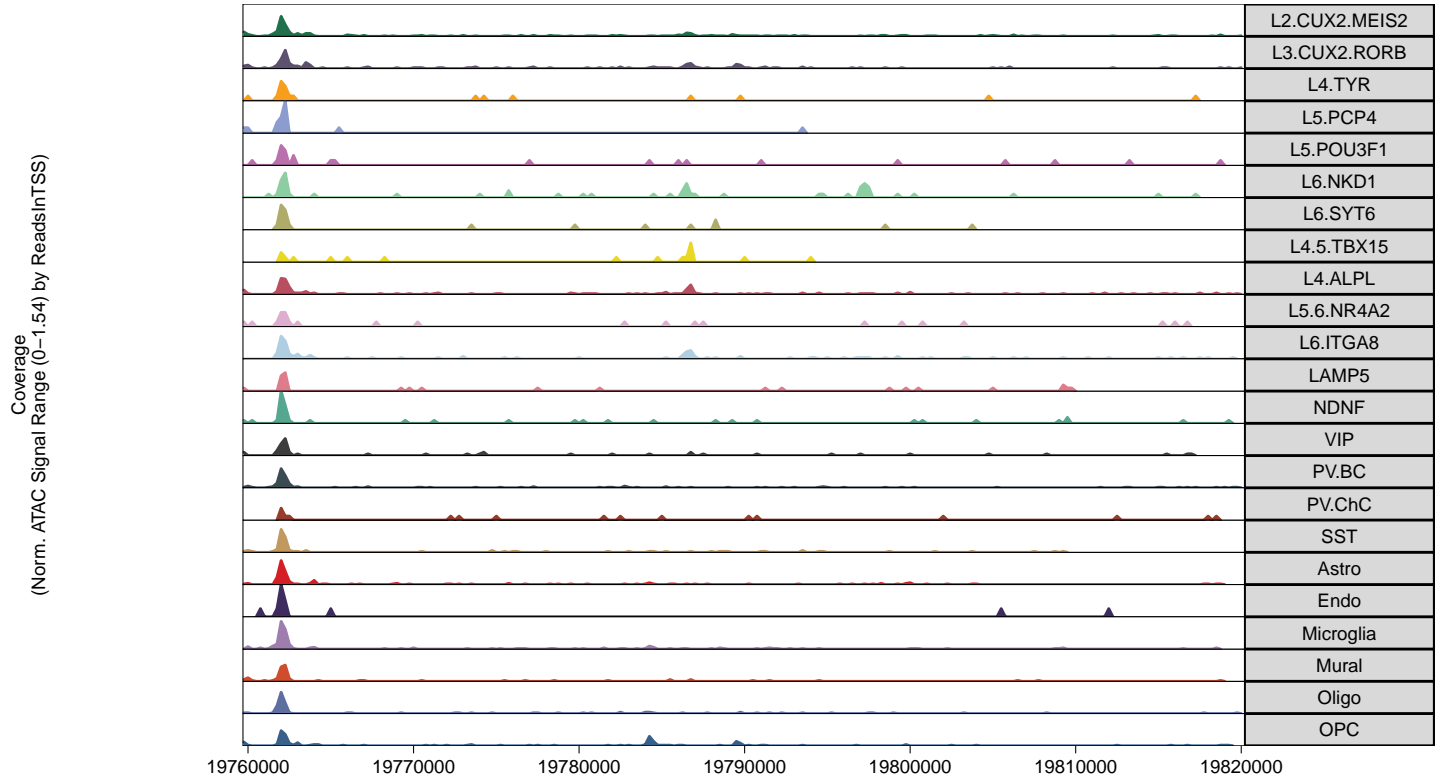


LoopTrack

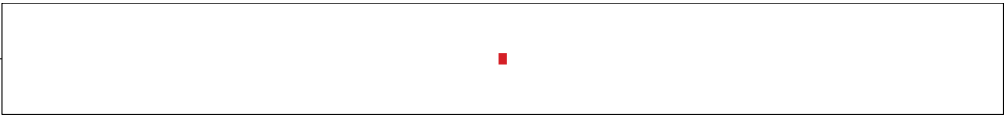


Genes

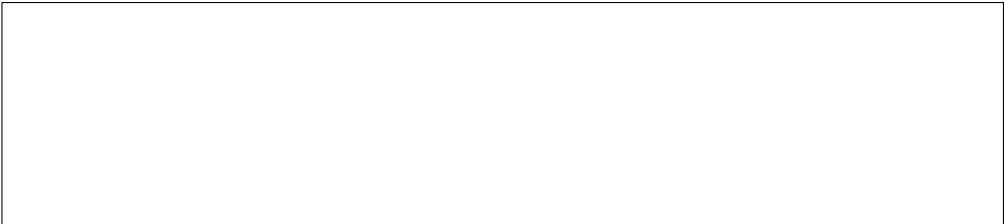
chr10:19759694–19820195 Marker Peak Tracks for: L3.CUX2.RORB



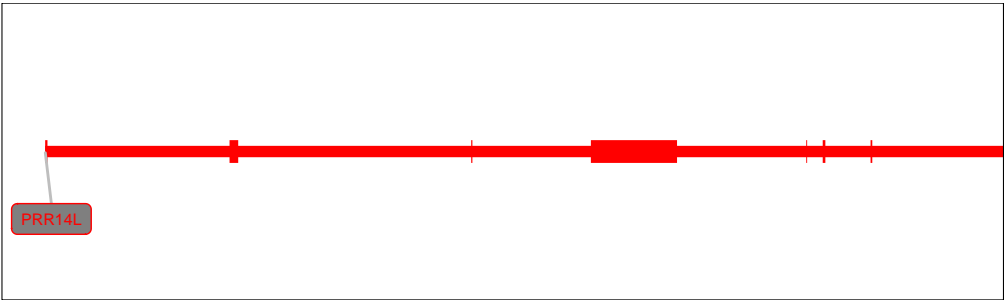
L3.CUX2.RORB



Peaks



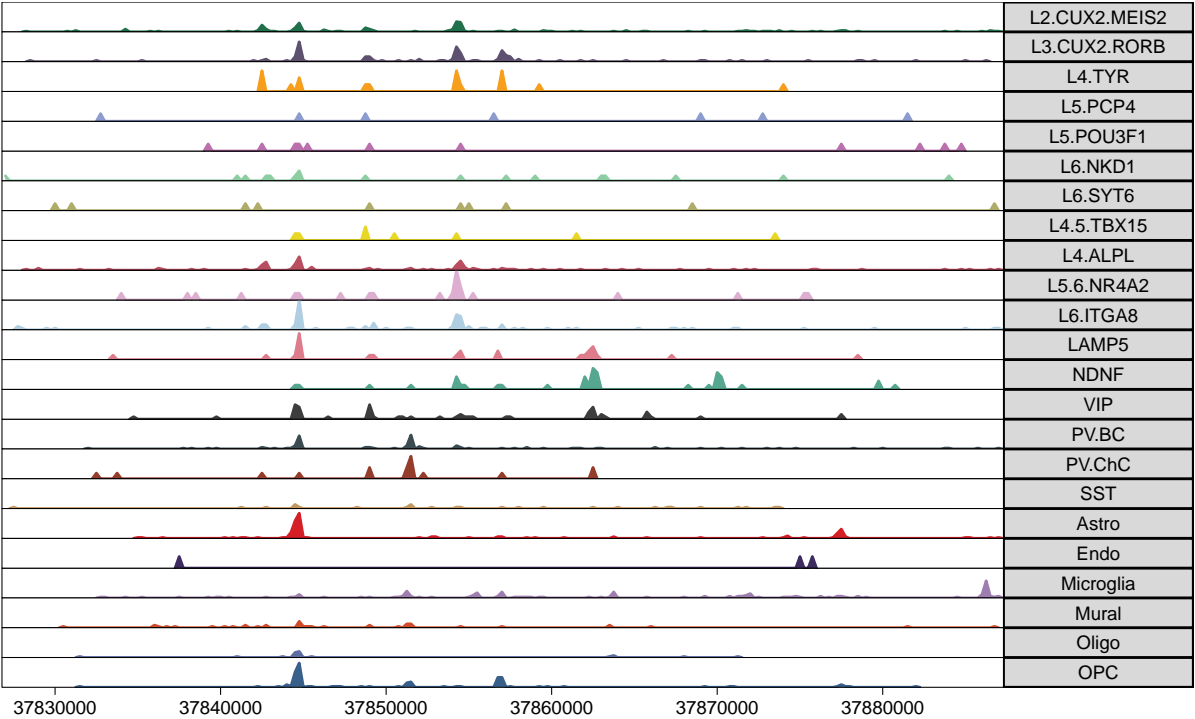
LoopTrack



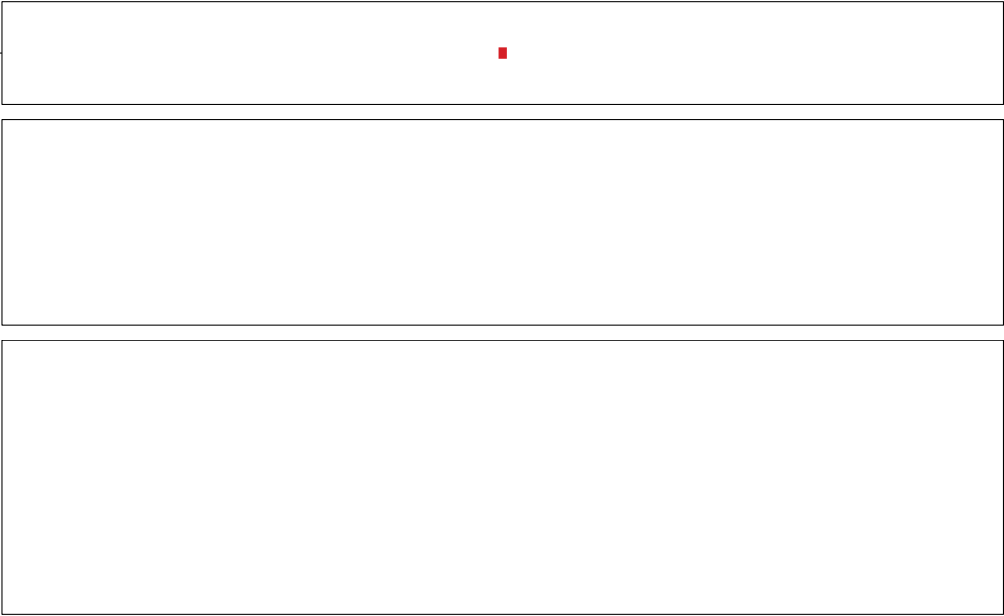
Genes



Coverage  
(Norm. ATAC Signal Range (0–1.01) by Reads in TSS)



L3.CUX2.RORB



Peaks

LoopTrack

GeneTrack