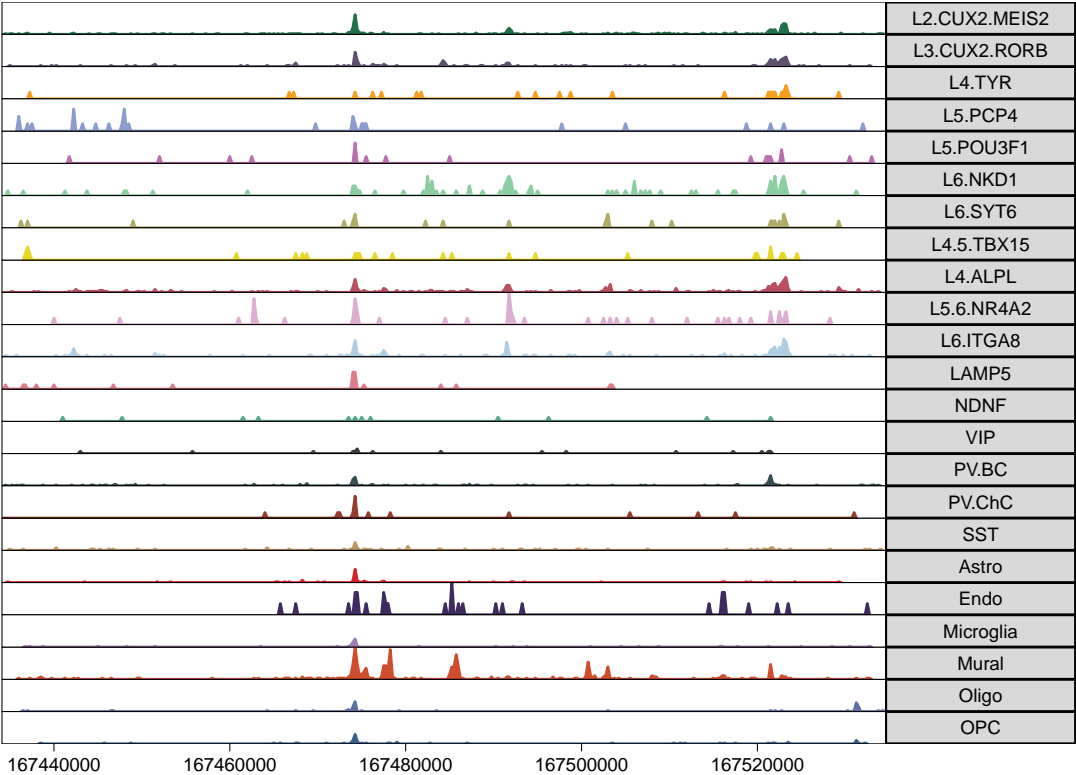


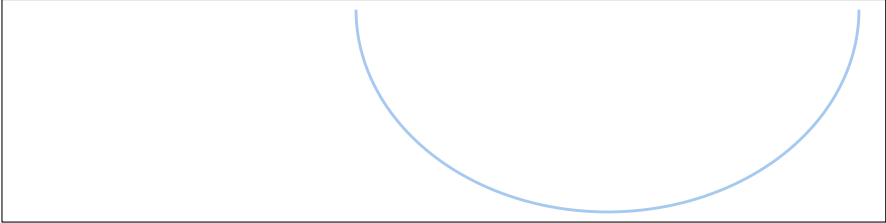
Coverage  
(Norm. ATAC Signal Range 0–1.12) by ReadsInTSS)



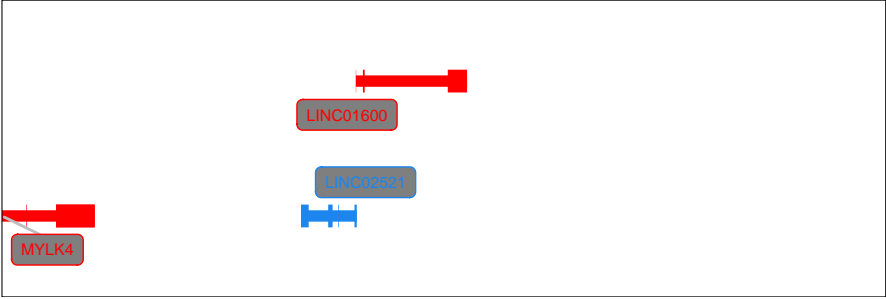
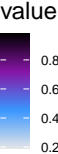
L3.CUX2.RORB



Peaks

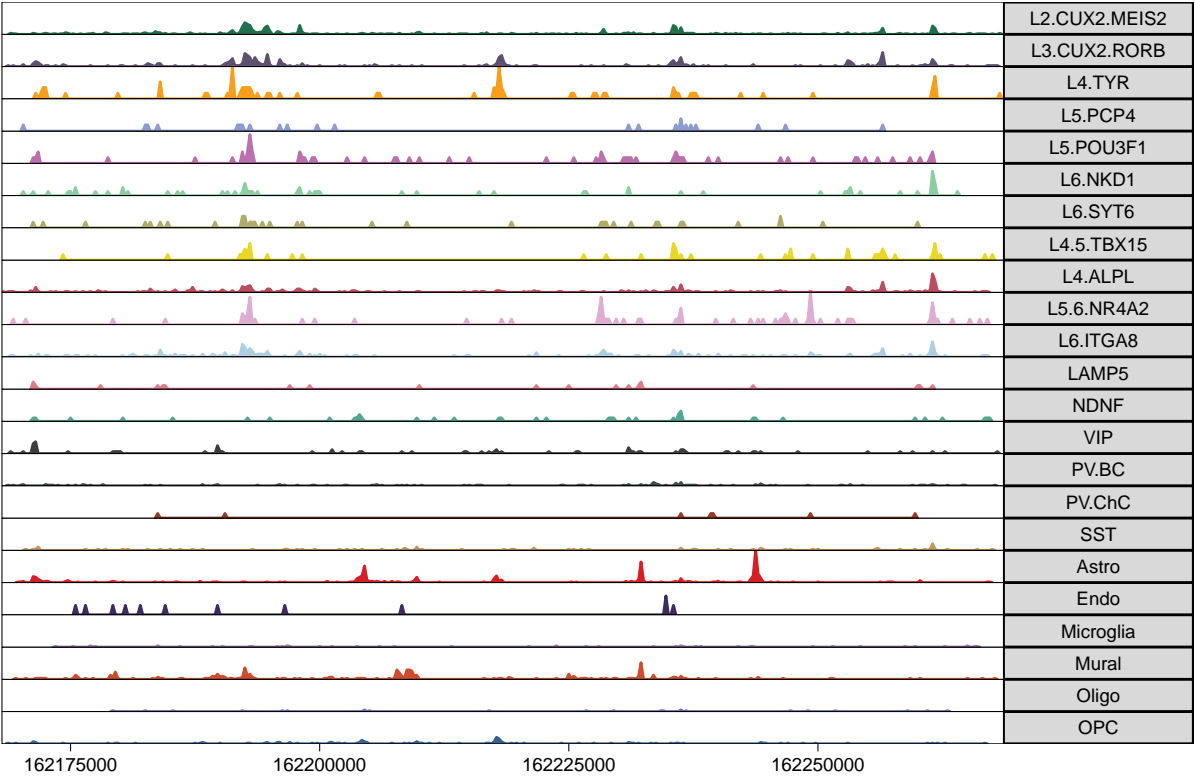


Peak2GeneLinks

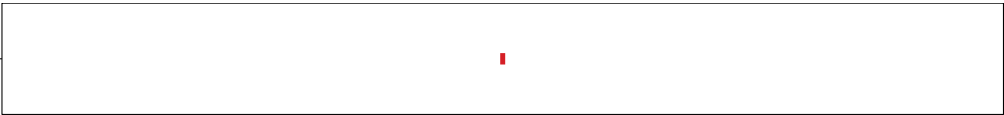


Genes

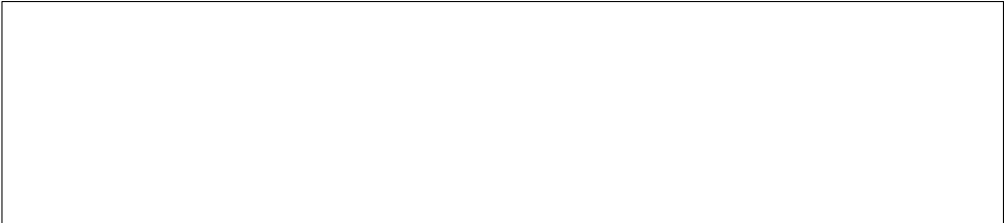
Coverage  
(Norm. ATAC Signal Range 0–1.33) by ReadsInTSS)



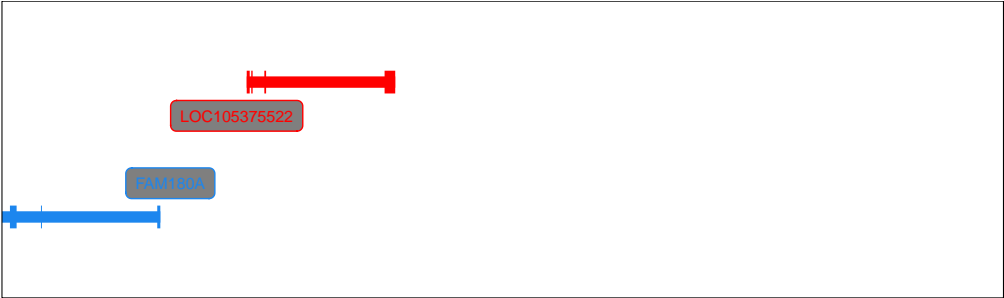
L3.CUX2.RORB



Peaks

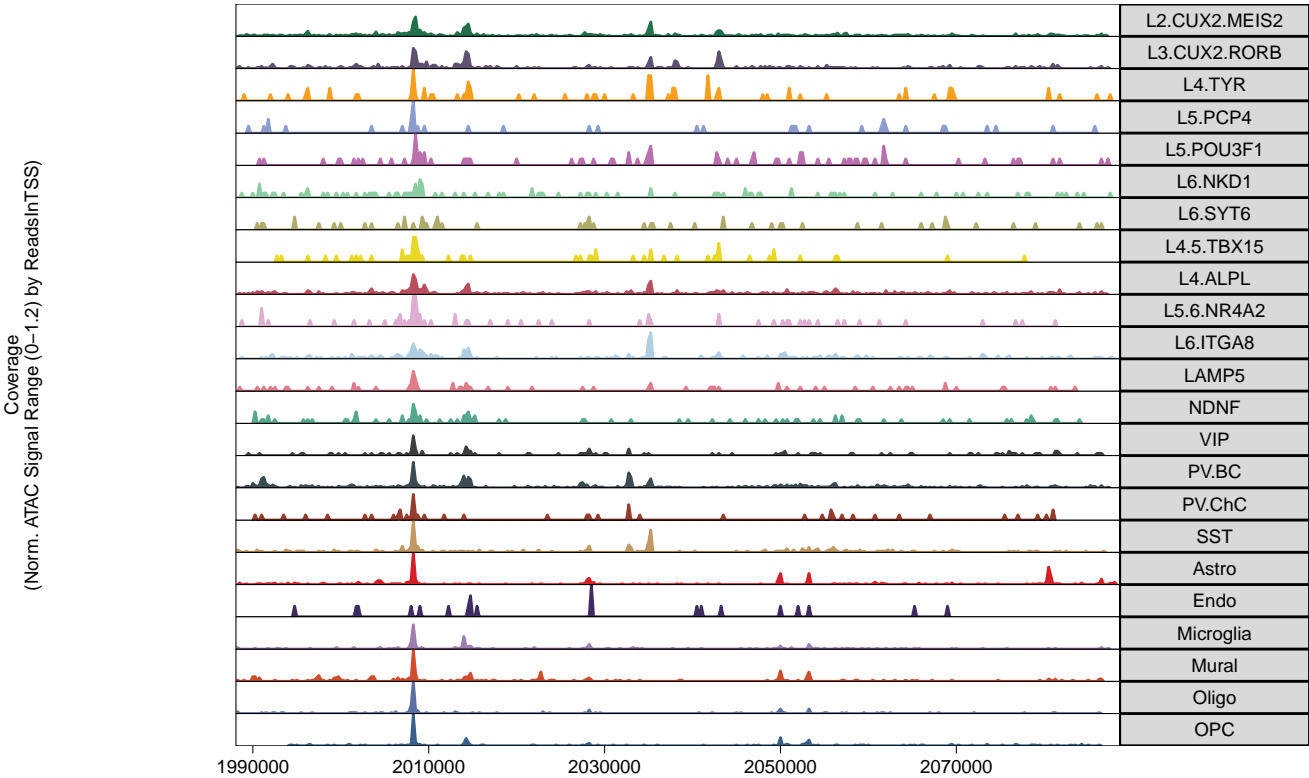


LoopTrack



Genes

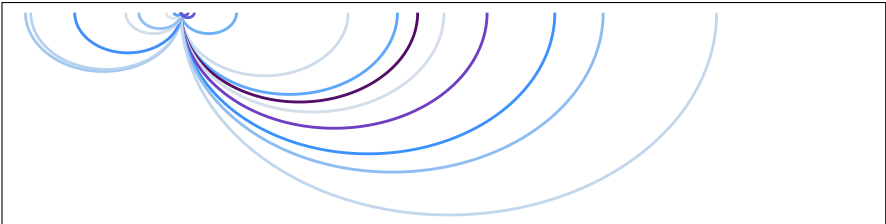
chr15:1988078–2088579 Marker Peak Tracks for: L3.CUX2.RORB



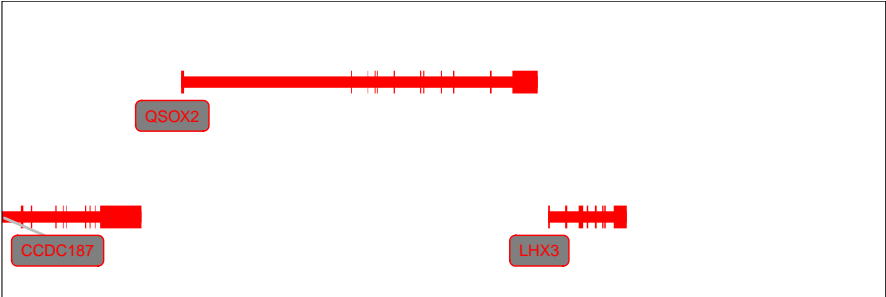
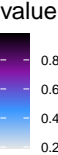
L3.CUX2.RORB



Peaks



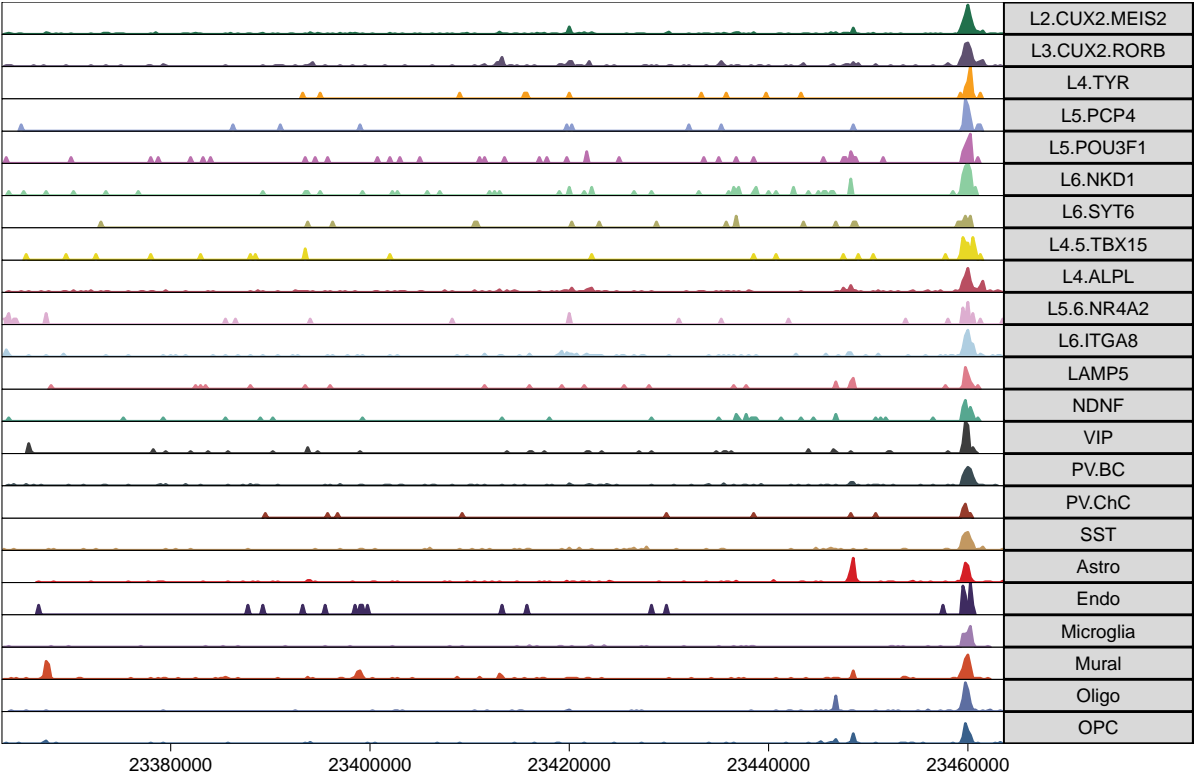
Peak2GeneLinks



Genes

chr17:23363068–23463569 Marker Peak Tracks for: L3.CUX2.RORB

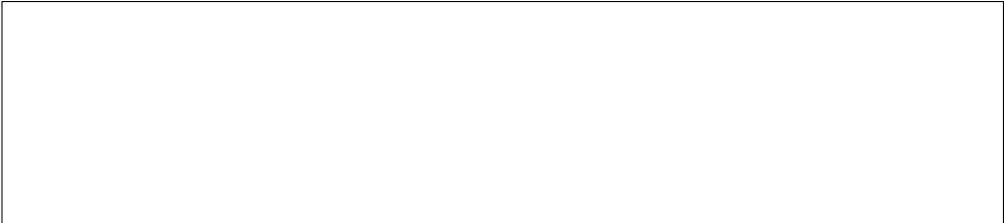
Coverage  
(Norm. ATAC Signal Range (0–1.31) by ReadsInTSS)



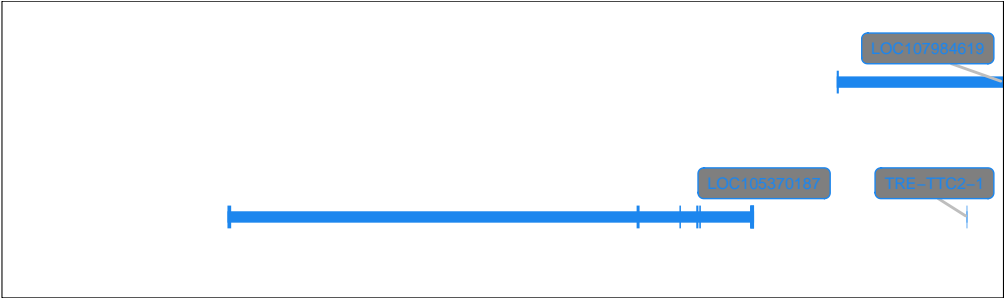
L3.CUX2.RORB



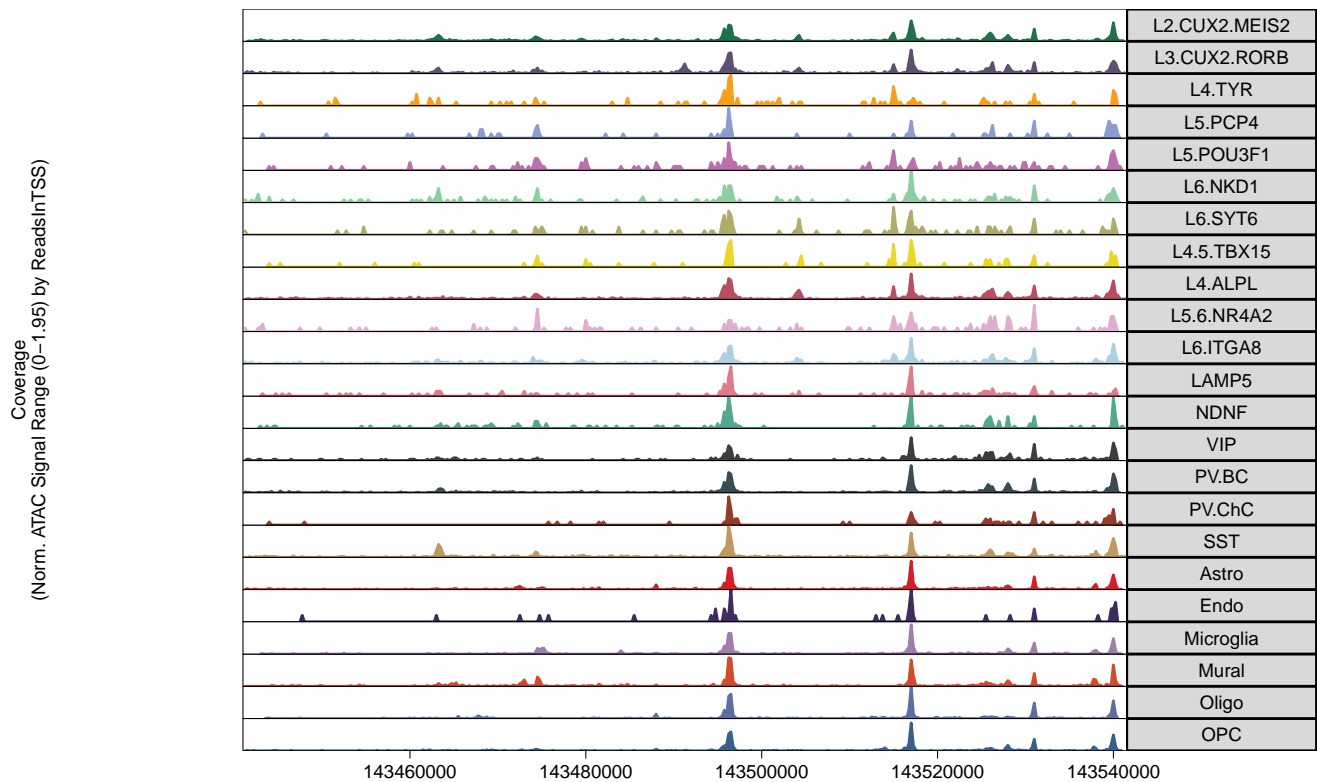
Peaks



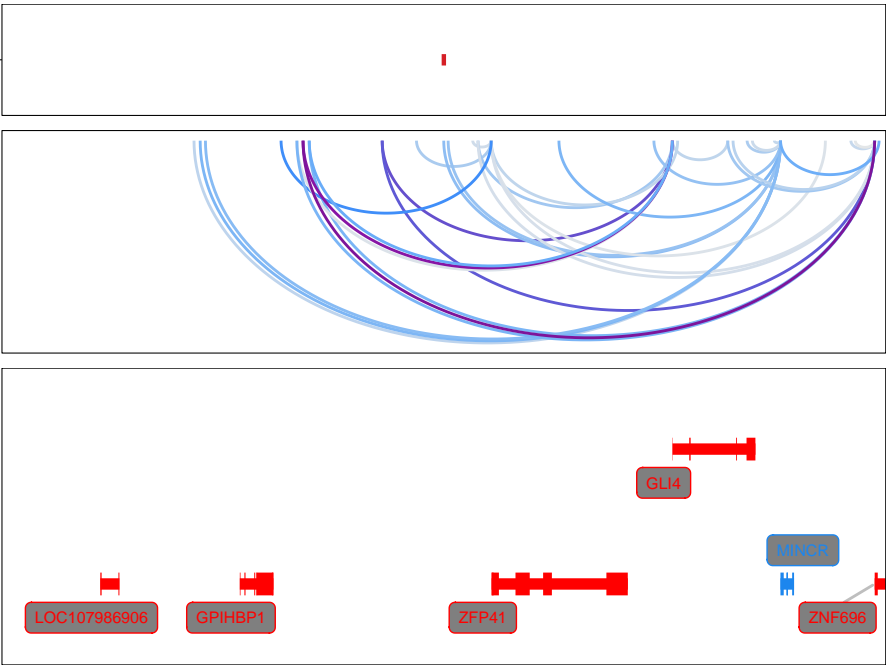
LoopTrack



Genes

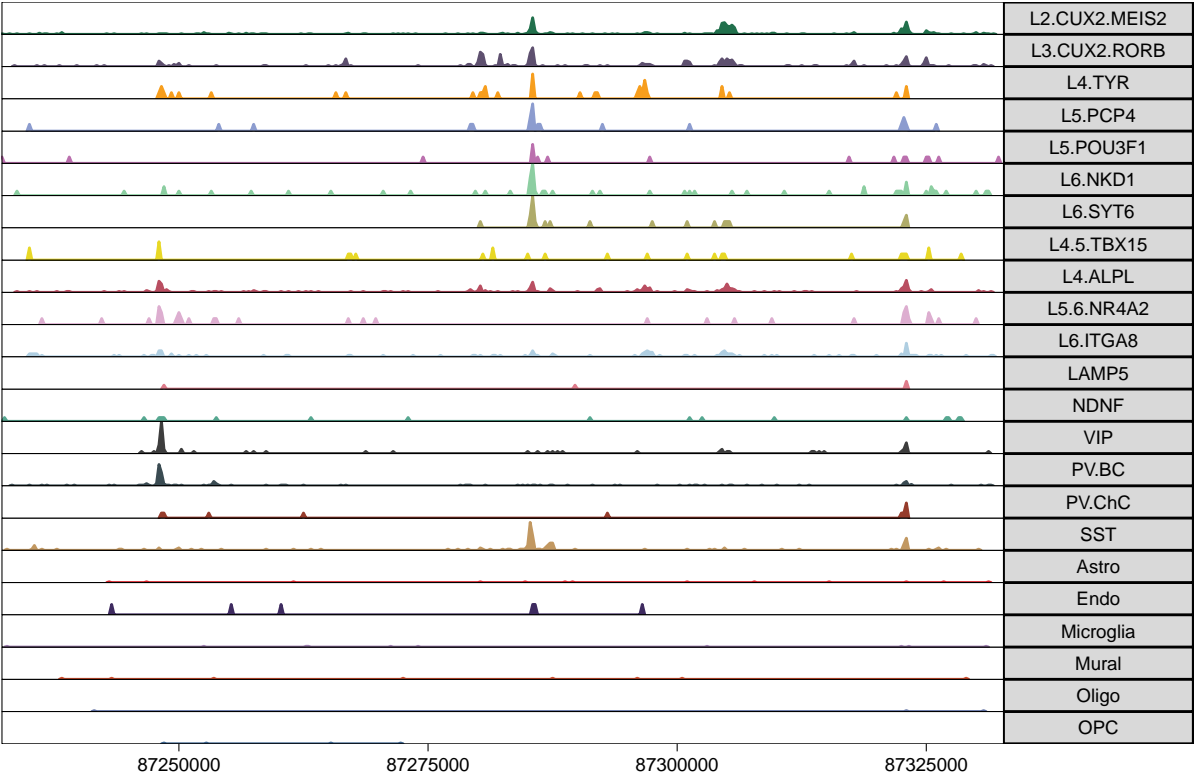


L3.CUX2.RORB



chr4:87232242–87332743 Marker Peak Tracks for: L3.CUX2.RORB

Coverage  
(Norm. ATAC Signal Range 0–1.21) by ReadsInTSS)



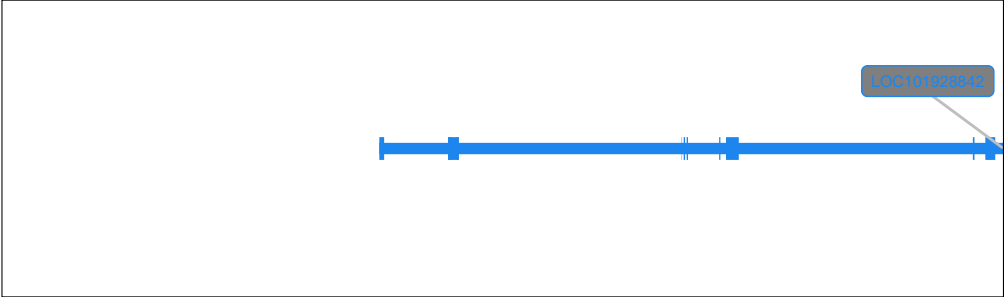
L3.CUX2.RORB



Peaks



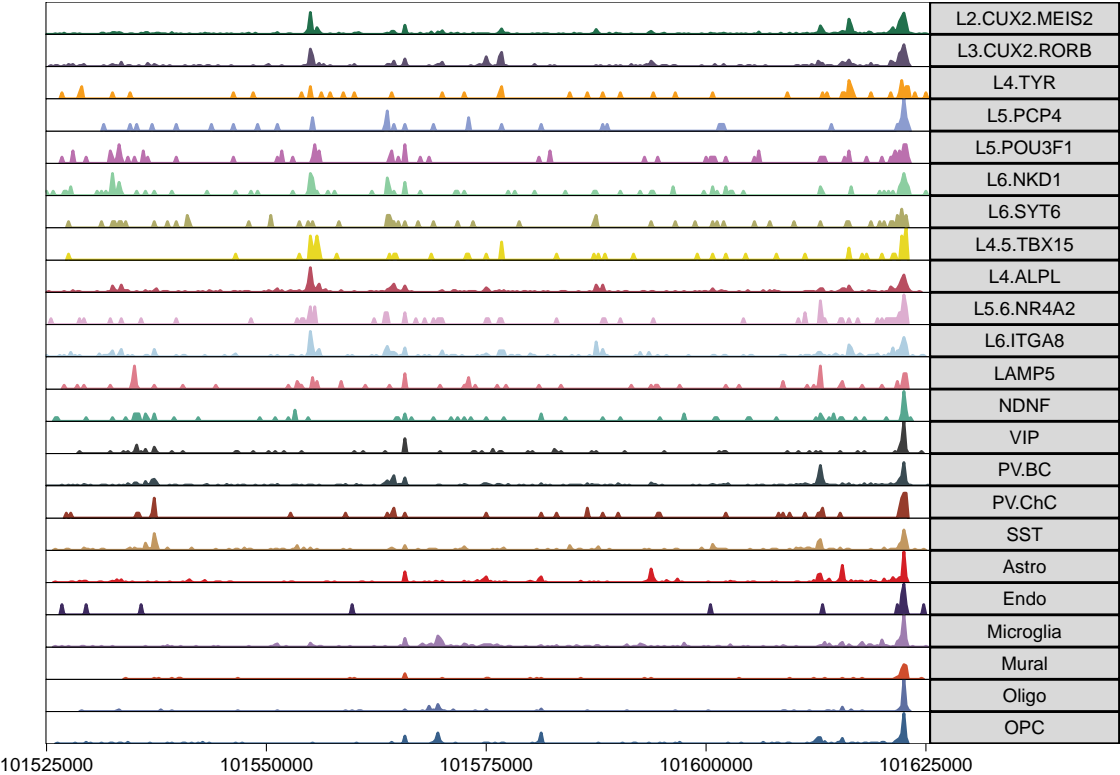
LoopTrack



Genes

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

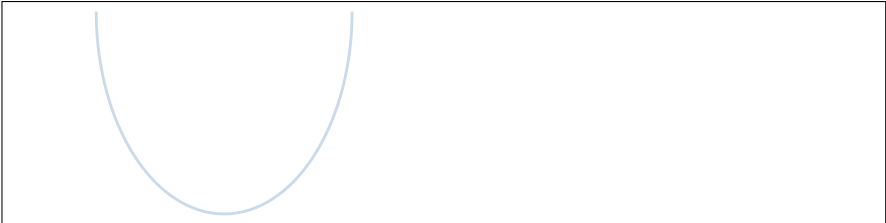
Coverage  
(Norm. ATAC Signal Range 0–1.23) by ReadsInTSS)



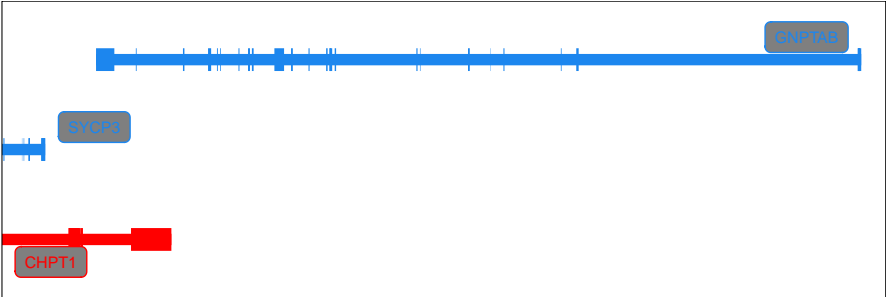
L3.CUX2.RORB



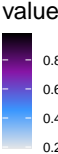
Peaks



Peak2GeneLinks

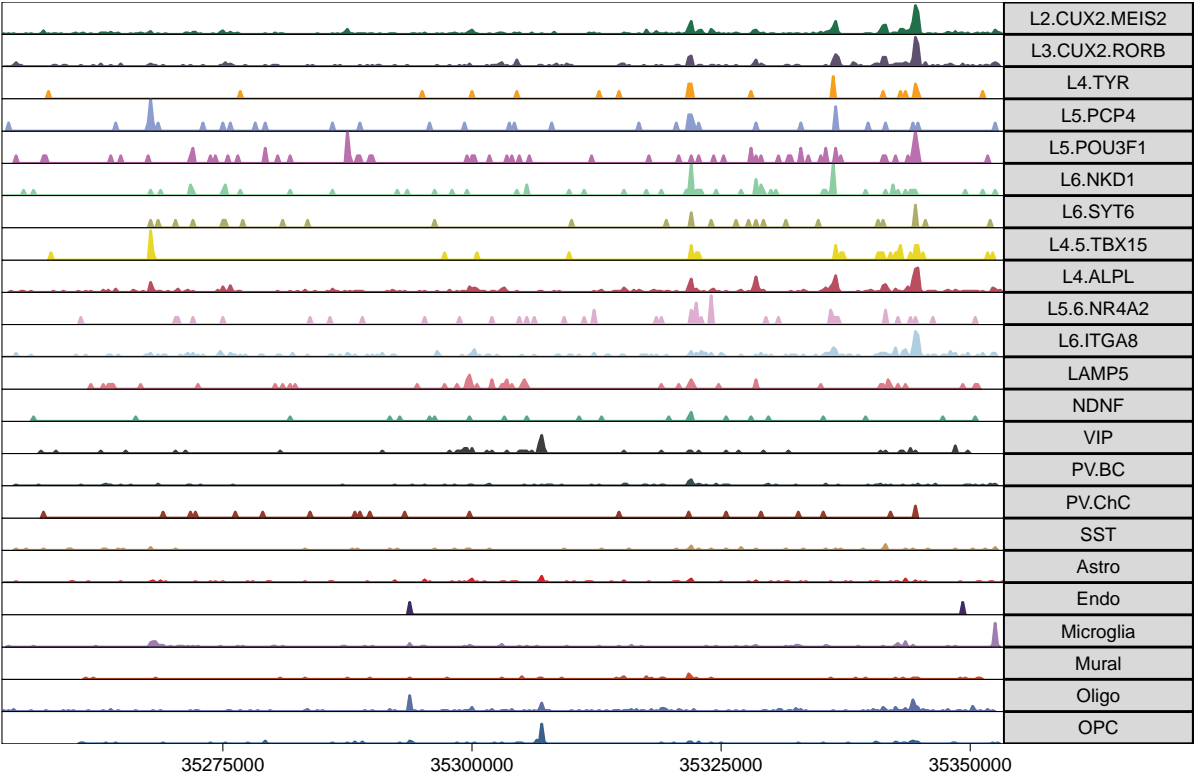


Genes



chr5:35252832–35353333 Marker Peak Tracks for: L3.CUX2.RORB

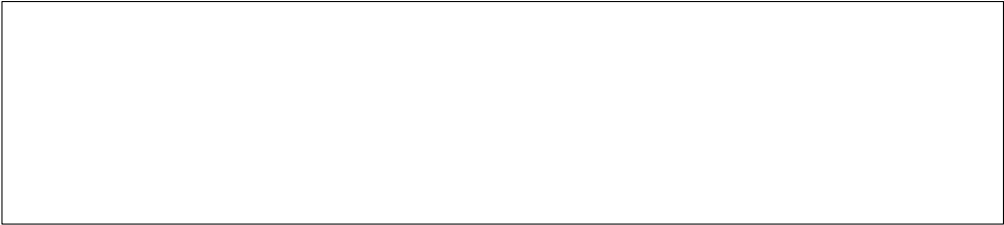
Coverage  
(Norm. ATAC Signal Range 0–1.01) by ReadsInTSS)



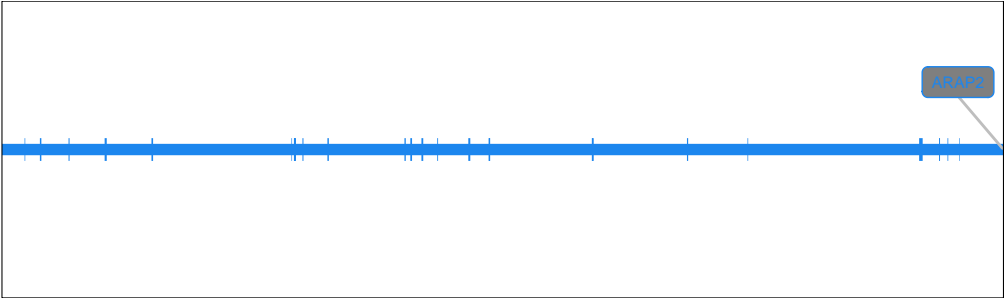
L3.CUX2.RORB



Peaks



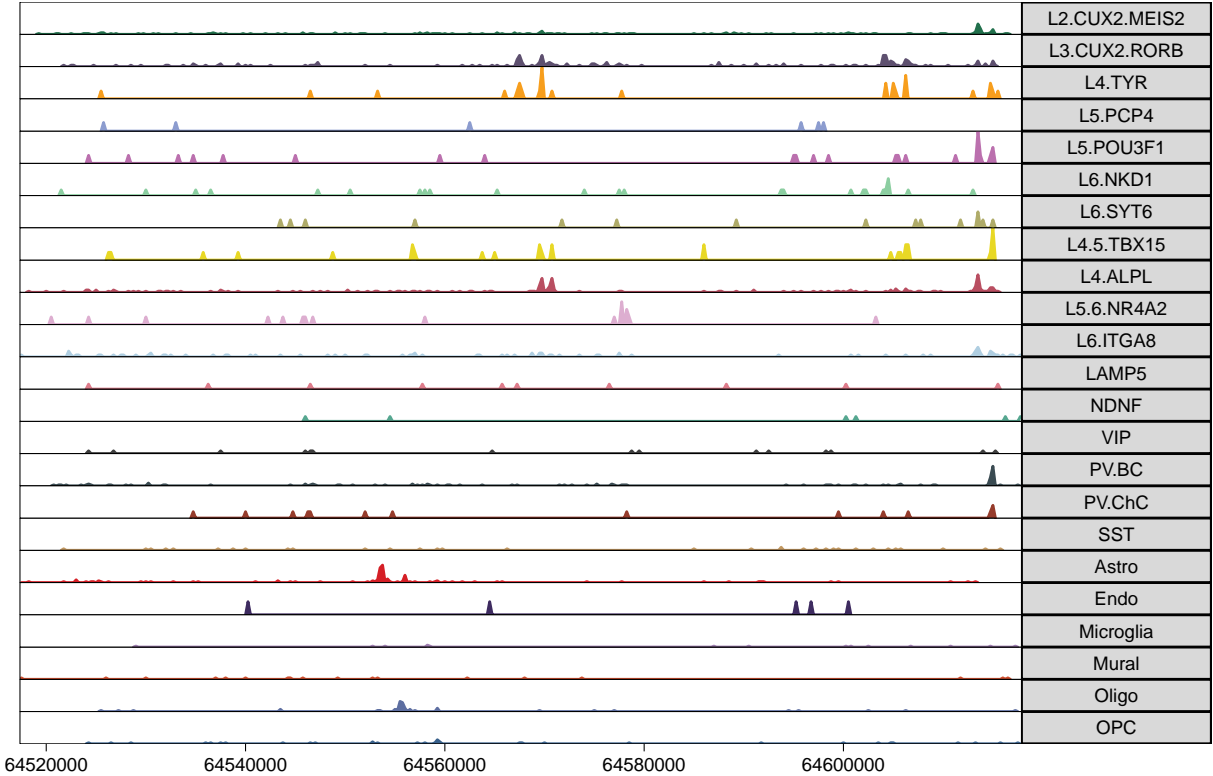
LoopTrack



Genes



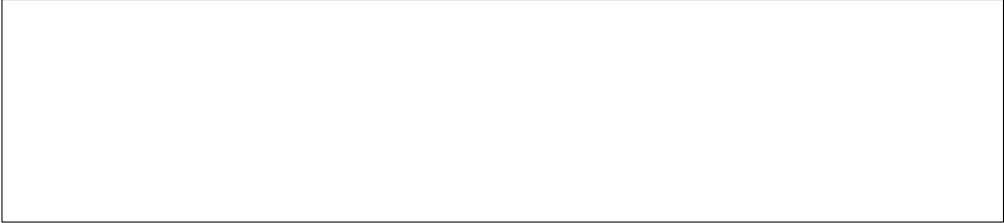
Coverage  
(Norm. ATAC Signal Range (0–0.96) by ReadsInTSS)



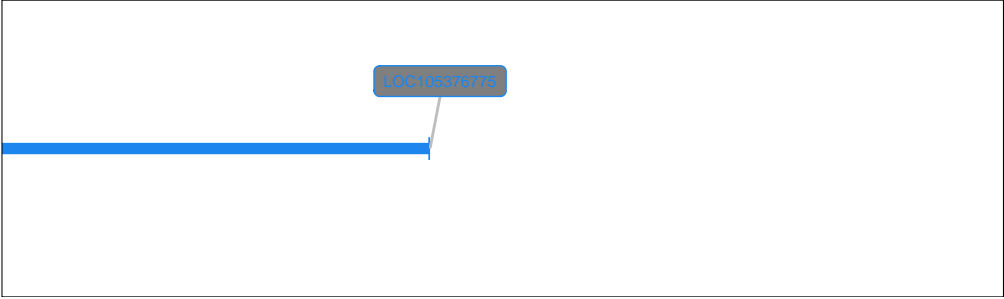
L3.CUX2.RORB



Peaks

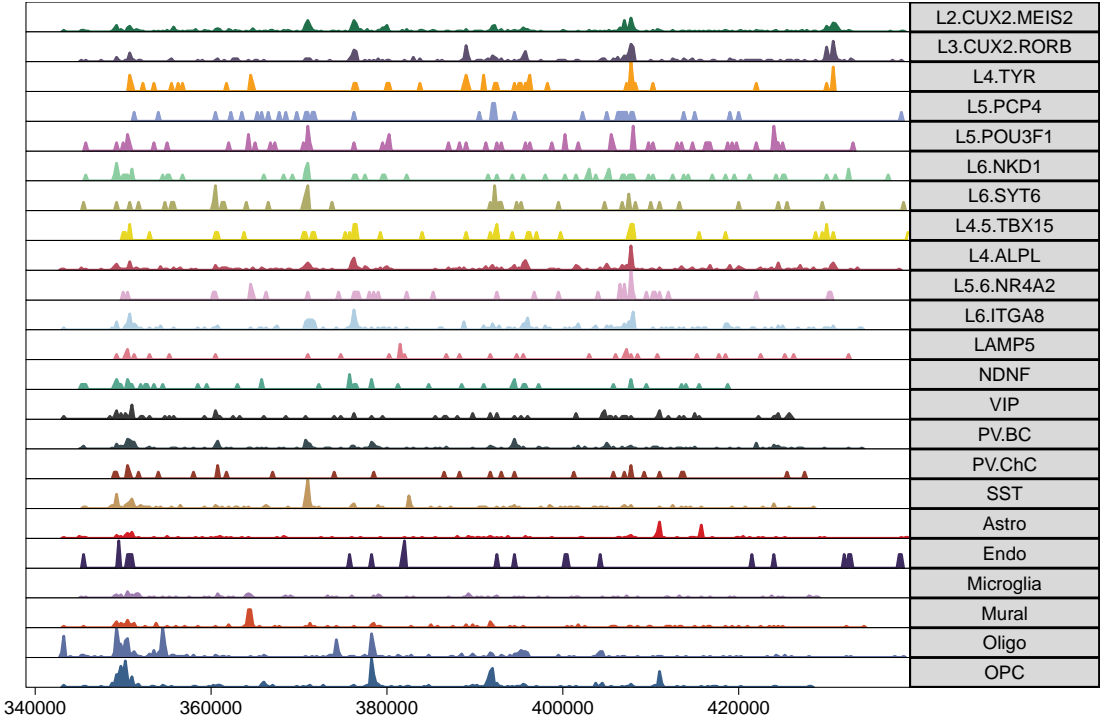


LoopTrack



Genes

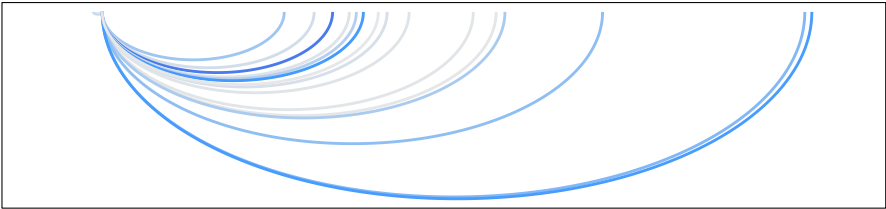
Coverage  
(Norm. ATAC Signal Range (0–0.85) by ReadsInTSS)



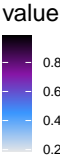
L3.CUX2.RORB



Peaks



Peak2GeneLinks

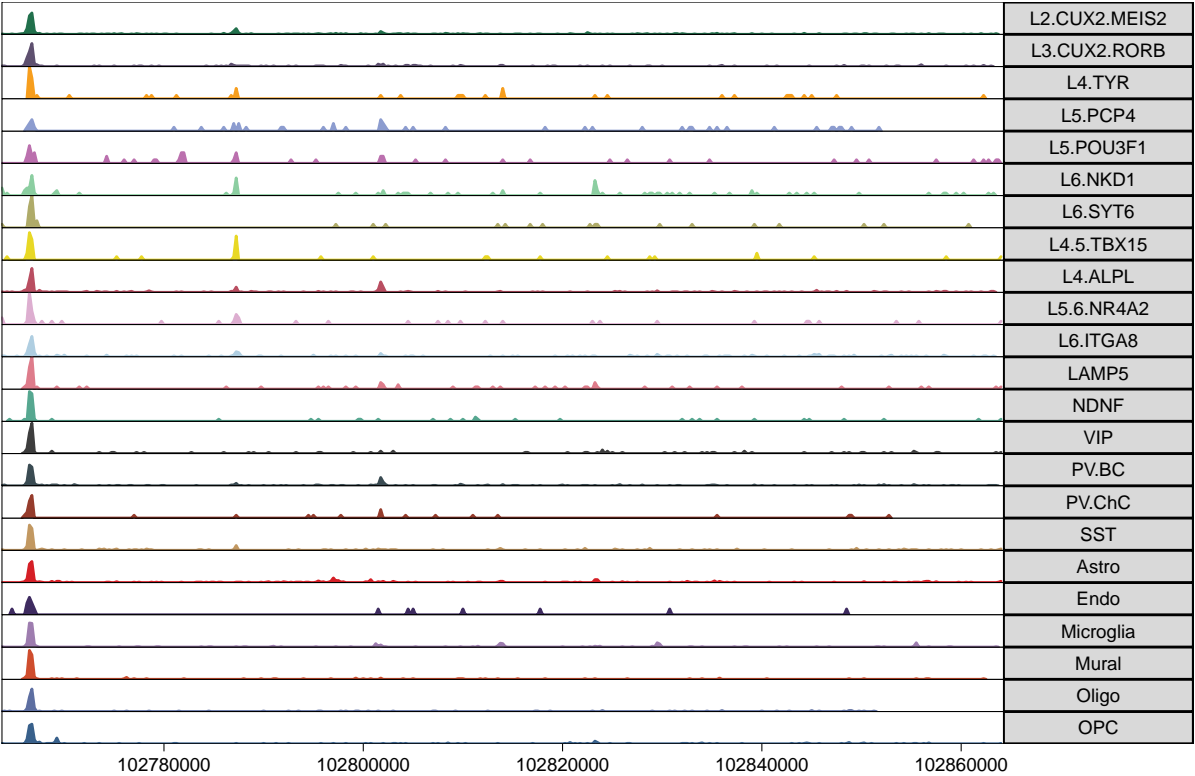


GeneTrack

strand    ●    ·

chr9:102763742–102864243 Marker Peak Tracks for: L3.CUX2.RORB

Coverage  
(Norm. ATAC Signal Range 0–2.16) by ReadsInTSS)



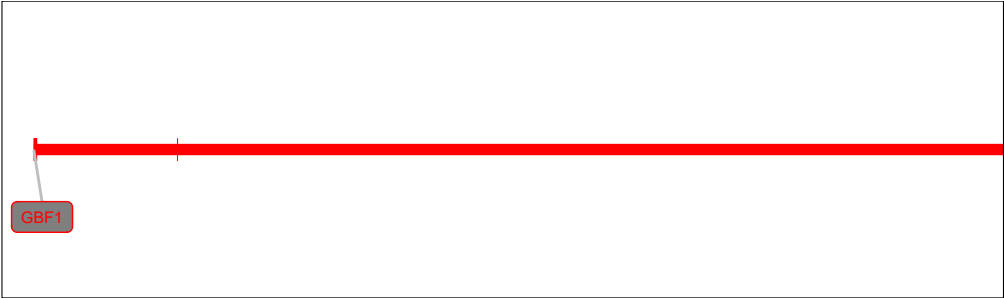
L3.CUX2.RORB



Peaks

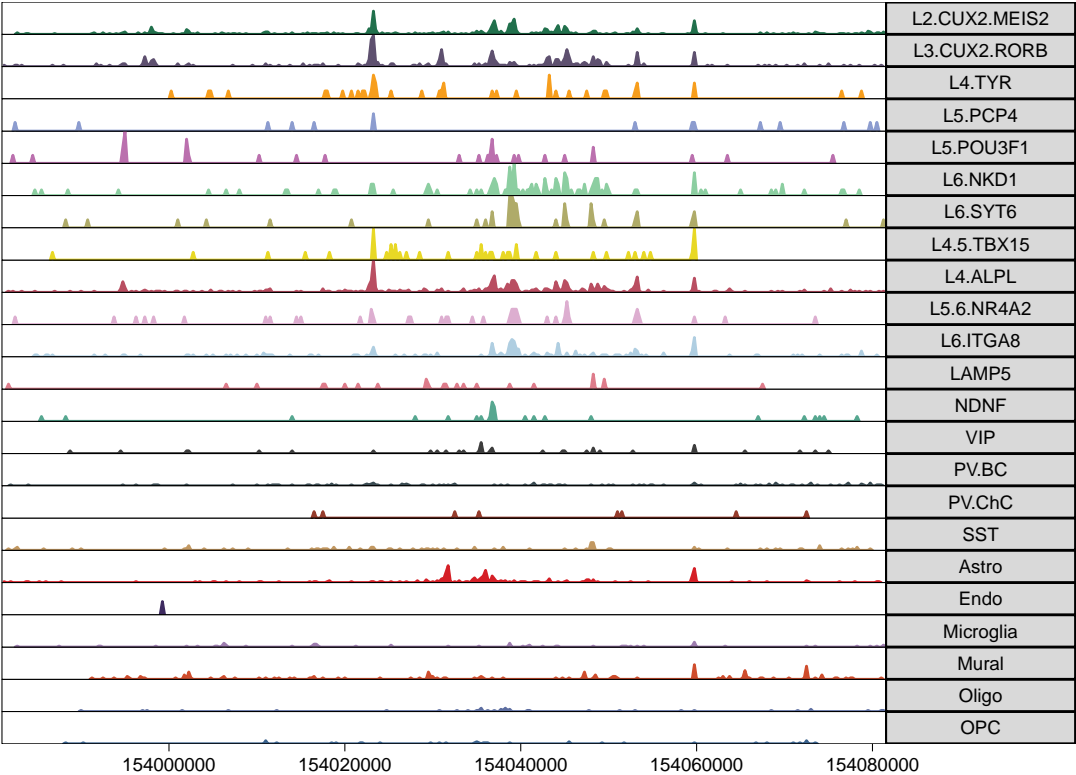


LoopTrack

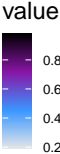
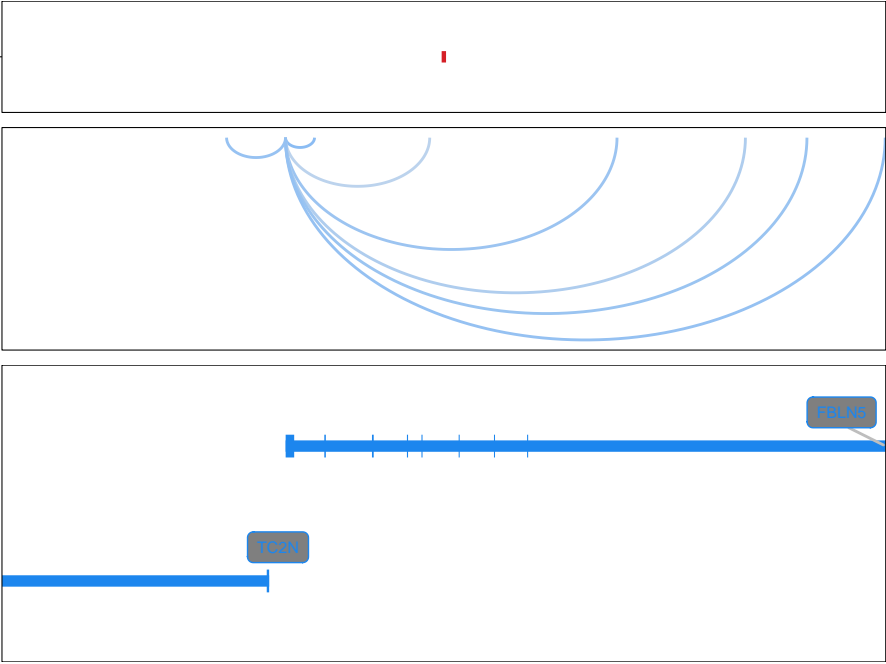


Genes

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

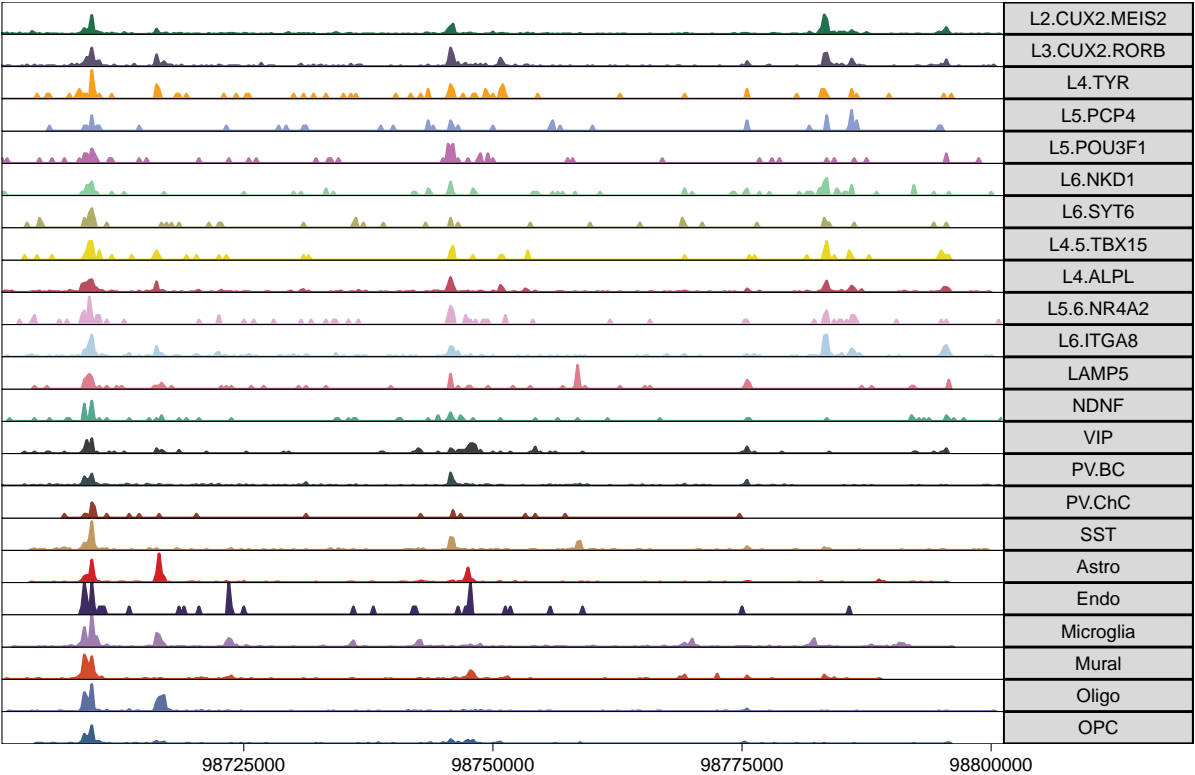


L3.CUX2.RORB



chr13:98700733–98801234 Marker Peak Tracks for: L3.CUX2.RORB

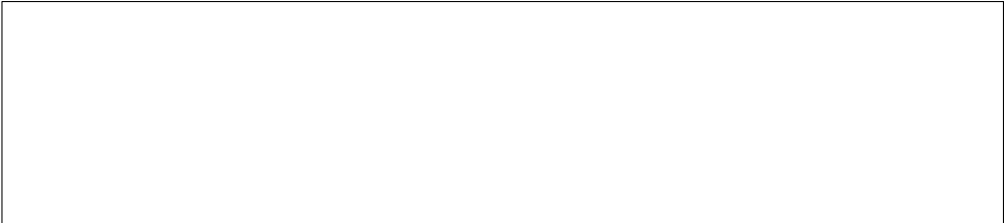
Coverage  
(Norm. ATAC Signal Range 0–1.59) by ReadsInTSS)



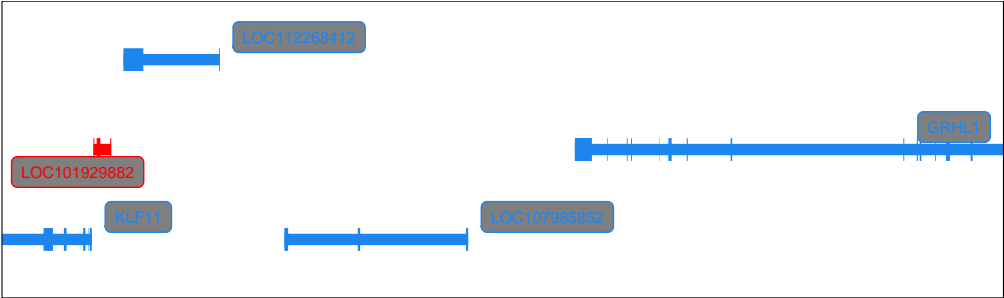
L3.CUX2.RORB



Peaks

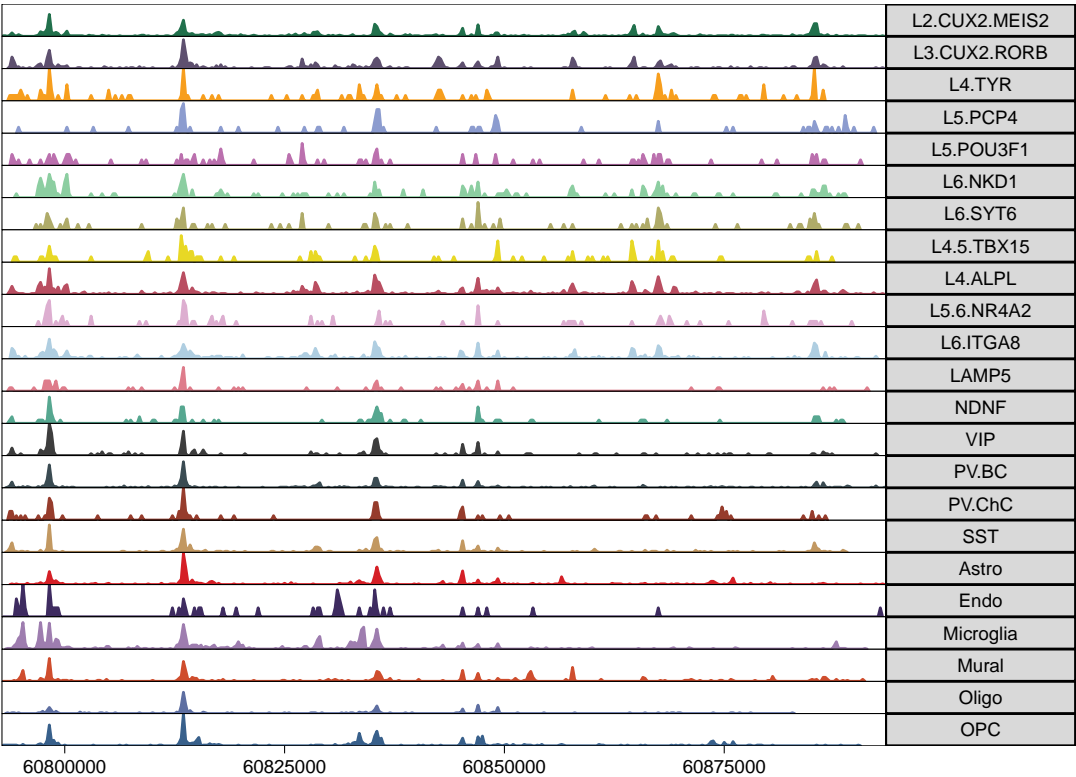


LoopTrack



Genes

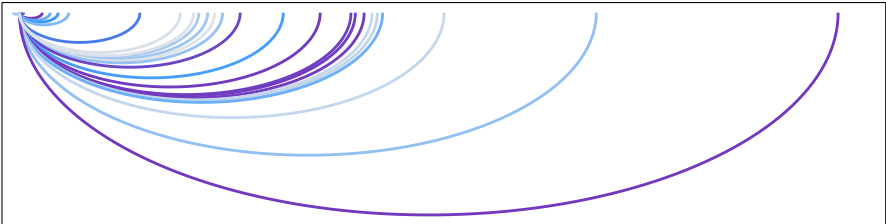
Coverage  
(Norm. ATAC Signal Range (0–1.4) by ReadsInTSS)



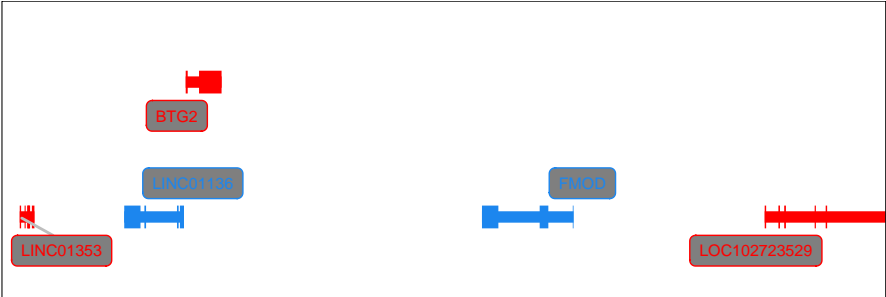
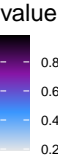
L3.CUX2.RORB



Peaks

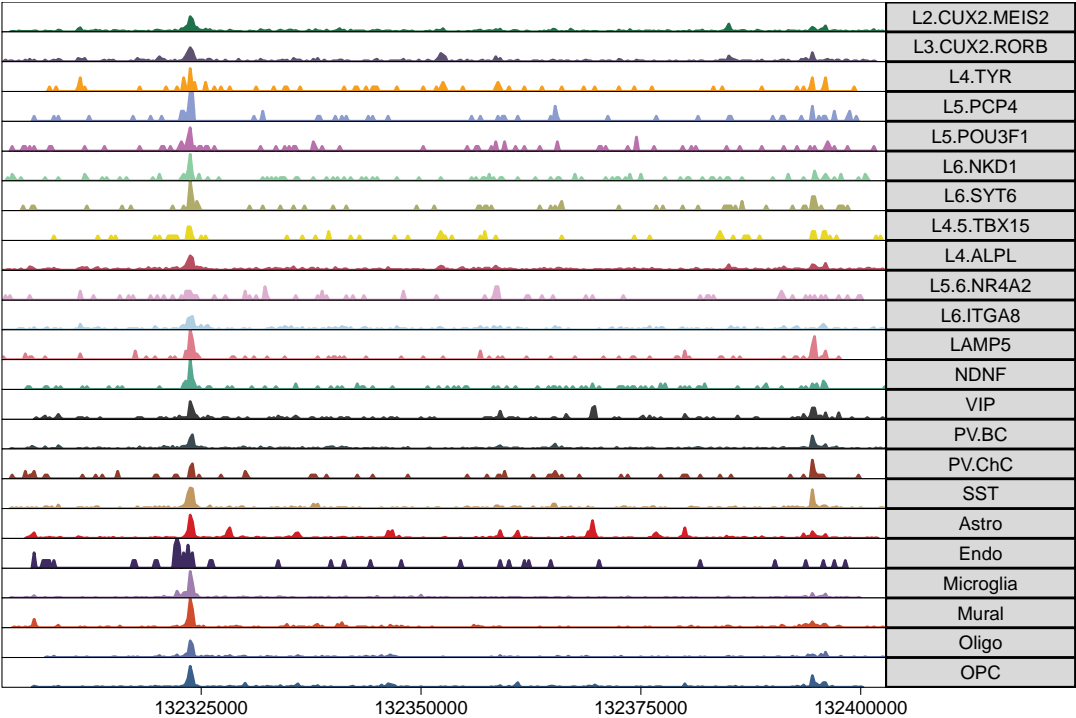


Peak2GeneLinks



Genes

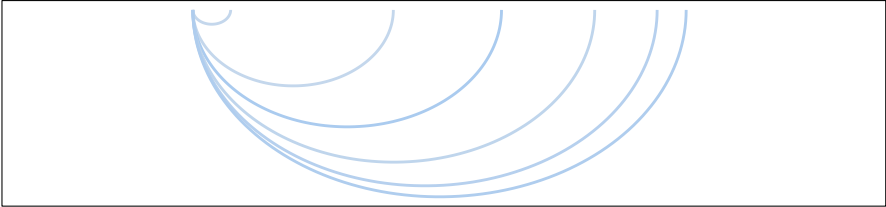
Coverage  
(Norm. ATAC Signal Range (0–1.52) by ReadslnTSS)



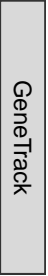
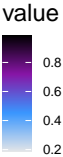
L3.CUX2.RORB



Peaks



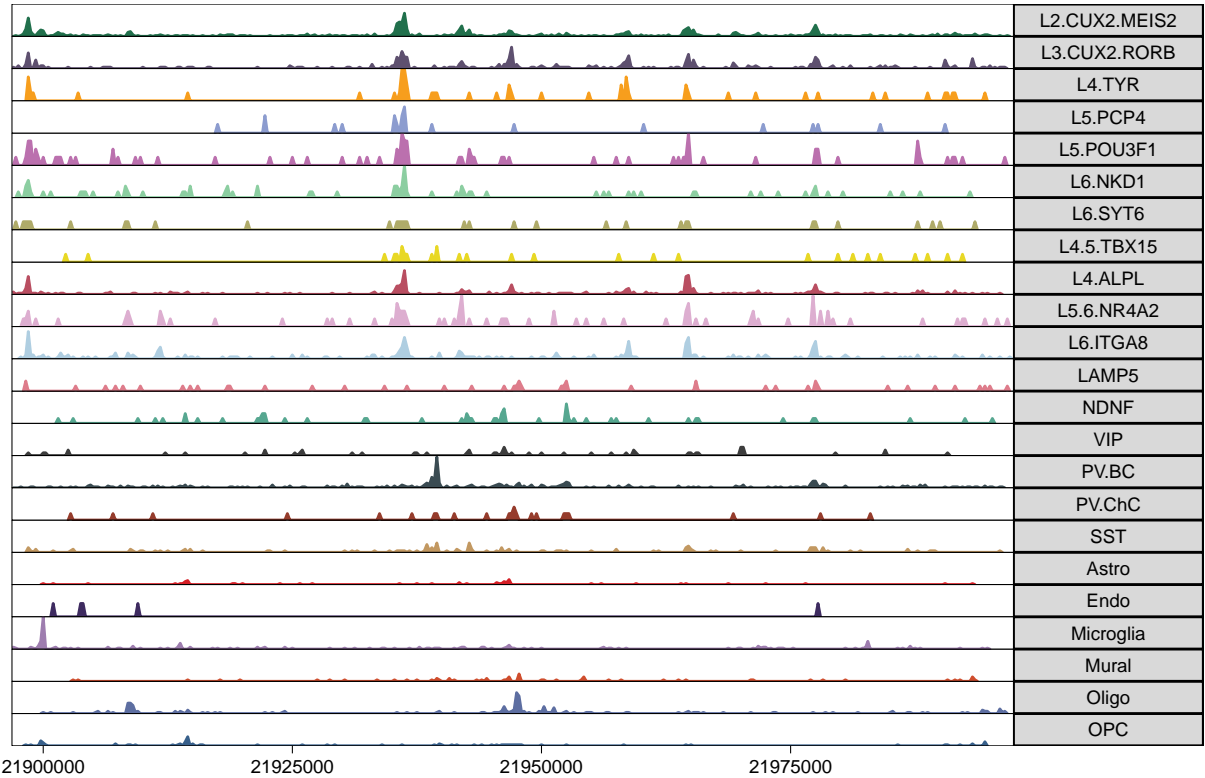
Peak2GeneLinks



strand ● ·

chr17:21896861–21997362 Marker Peak Tracks for: L3.CUX2.RORB

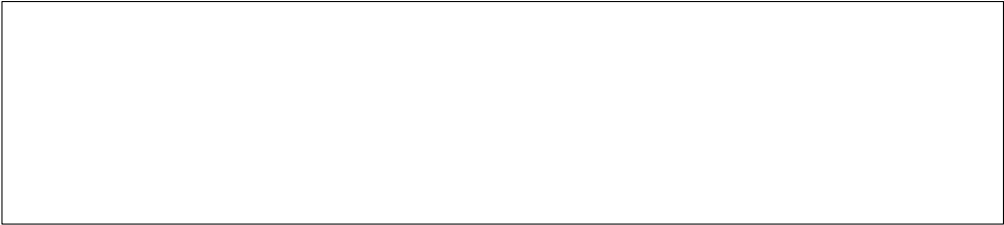
Coverage  
(Norm. ATAC Signal Range (0–0.95) by ReadsInTSS)



L3.CUX2.RORB



Peaks



LoopTrack

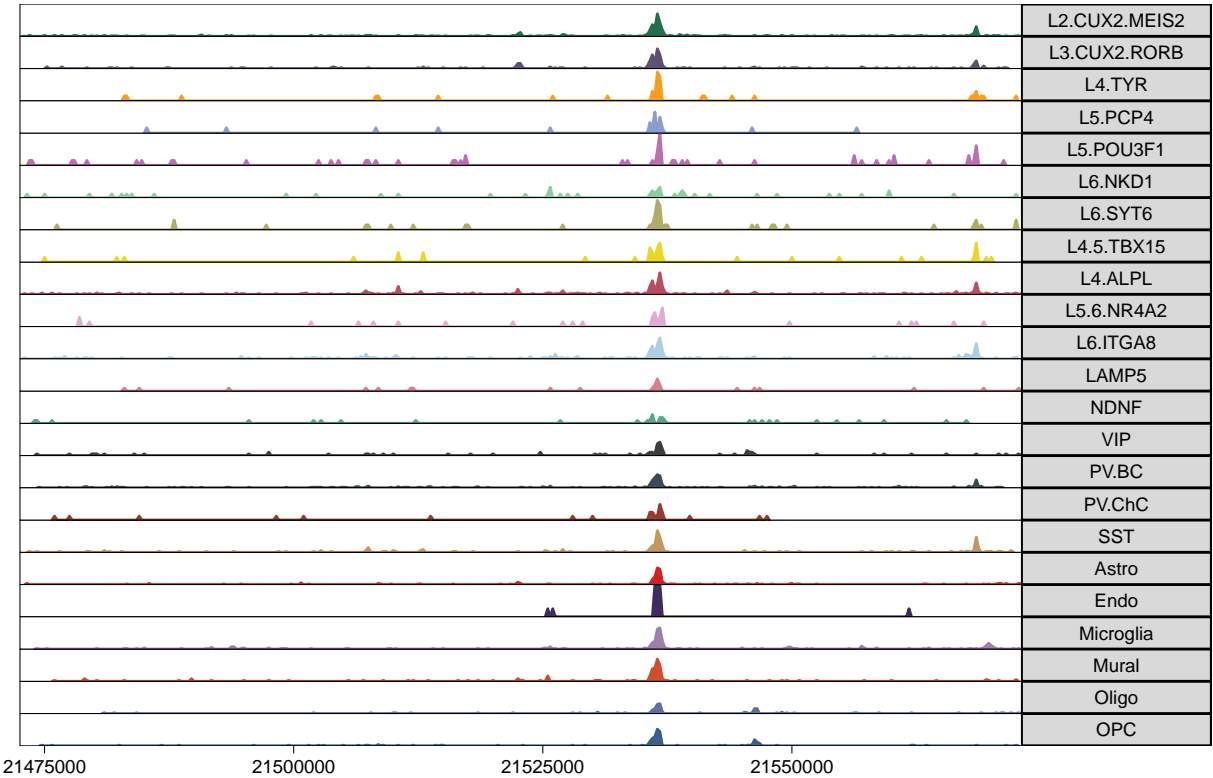


Genes

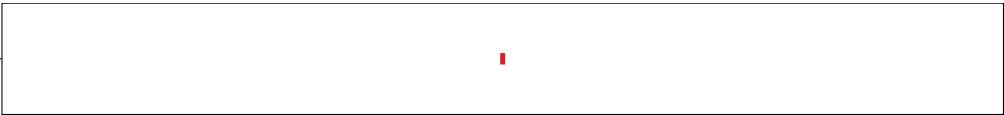


chr18:21472533–21573034 Marker Peak Tracks for: L3.CUX2.RORB

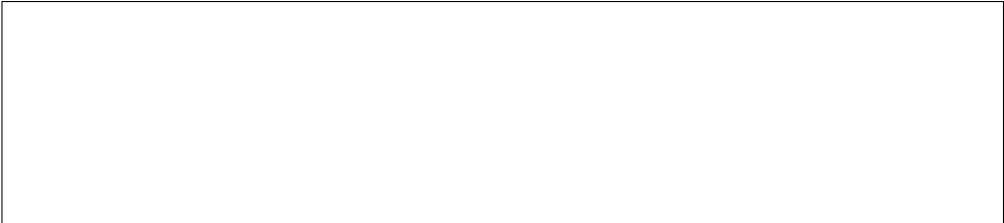
Coverage  
(Norm. ATAC Signal Range 0–1.56) by ReadsInTSS)



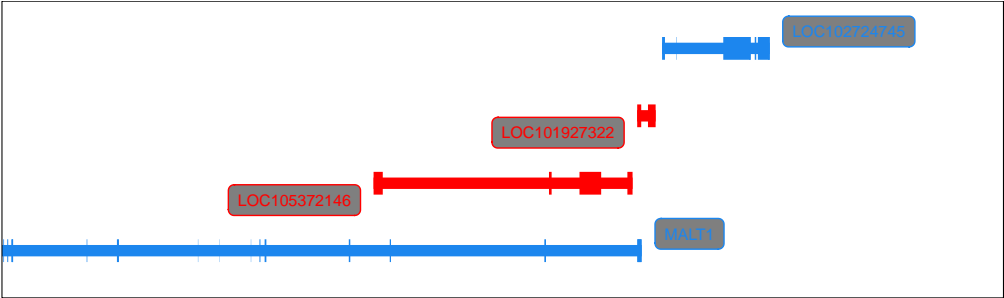
L3.CUX2.RORB



Peaks



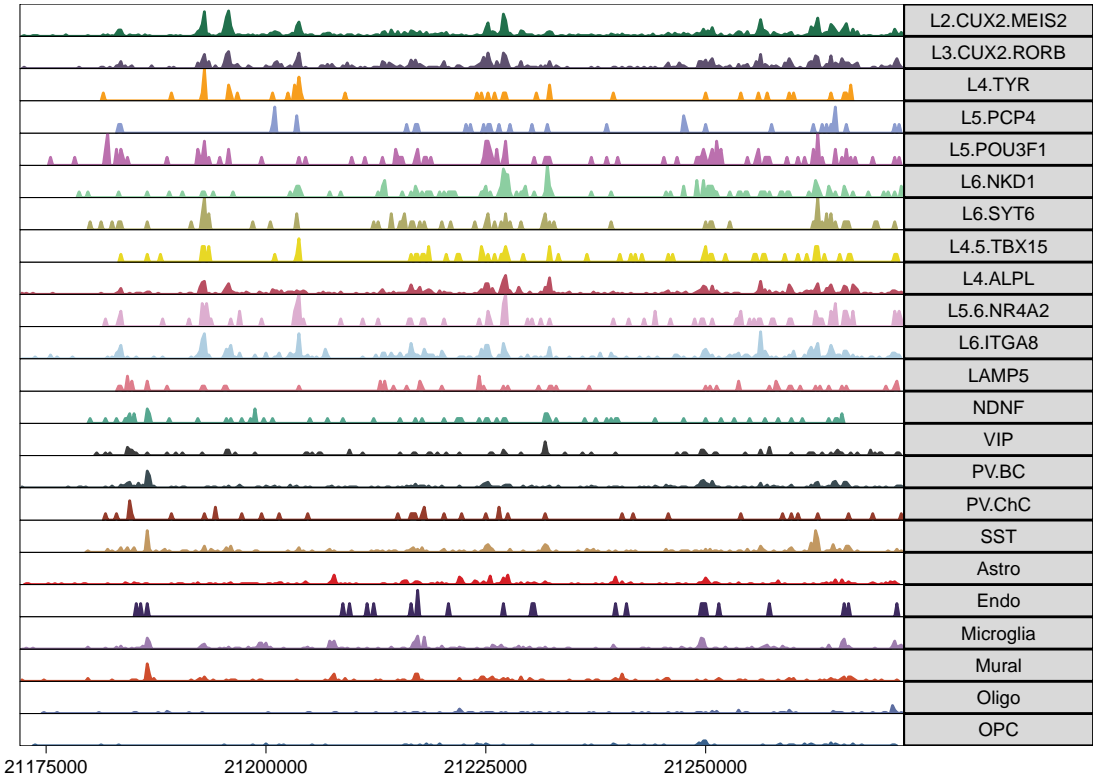
LoopTrack



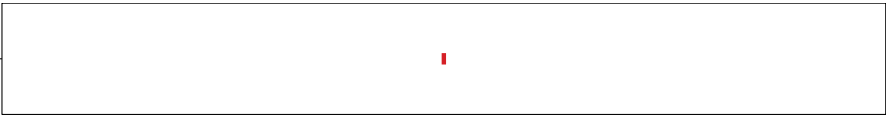
Genes

chr18:21172023–21272524 Marker Peak Tracks for: L3.CUX2.RORB

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



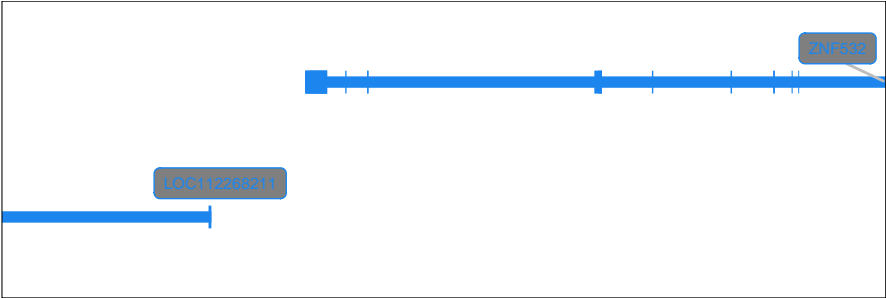
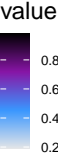
L3.CUX2.RORB



Peaks



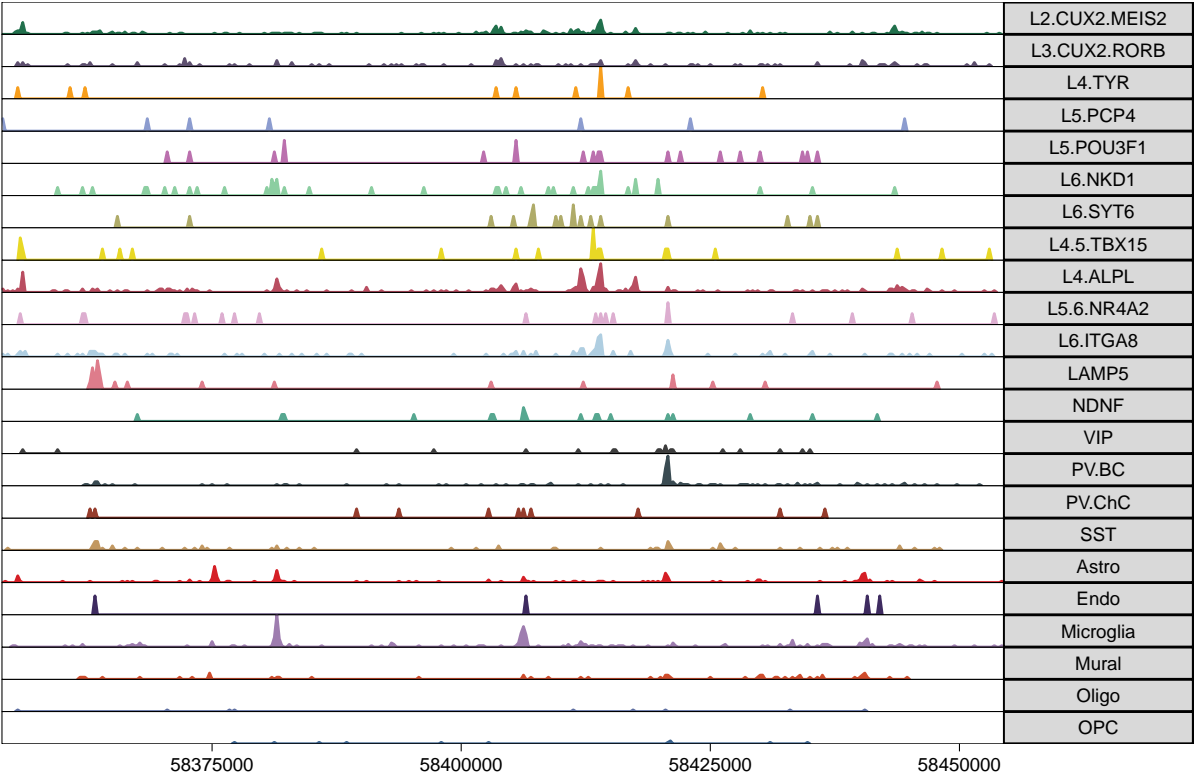
Peak2GeneLinks



Genes

chr17:58353914–58454415 Marker Peak Tracks for: L3.CUX2.RORB

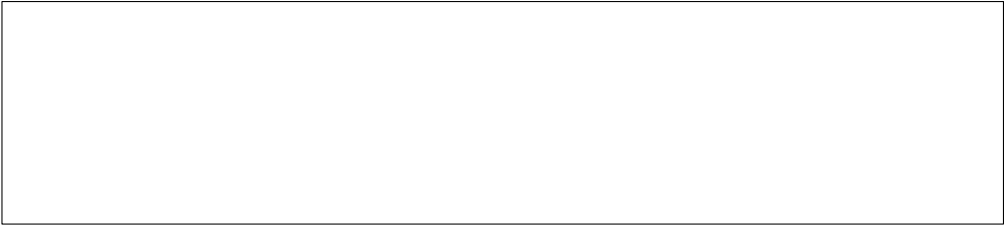
Coverage  
(Norm. ATAC Signal Range (0–0.67) by ReadsInTSS)



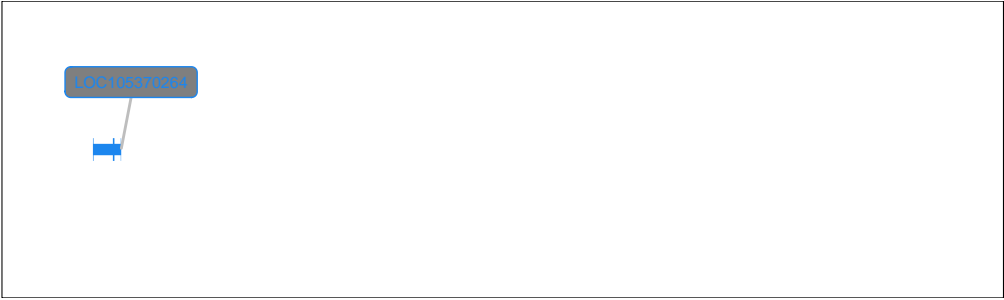
L3.CUX2.RORB



Peaks

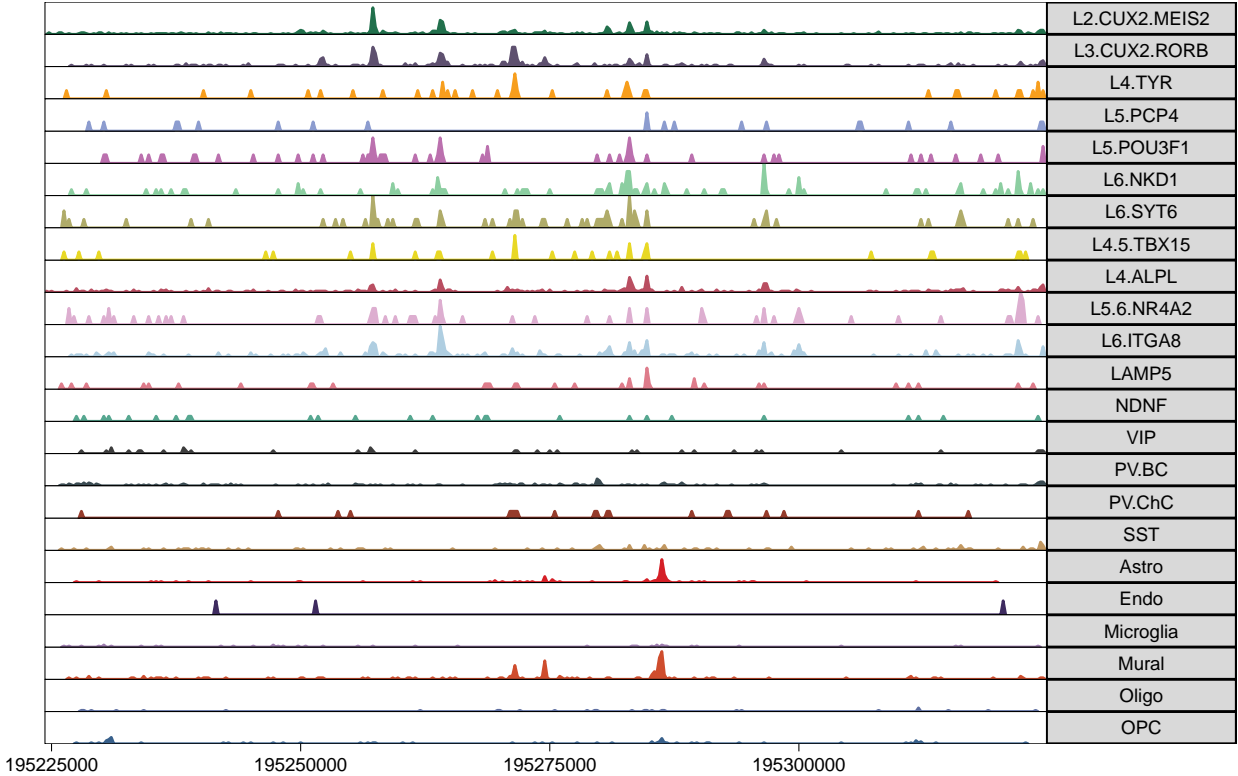


LoopTrack

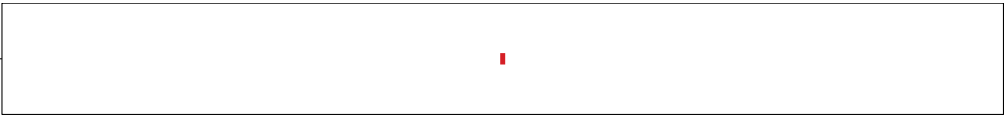


Genes

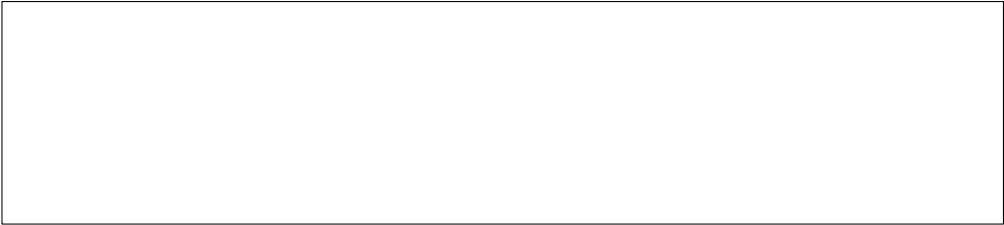
Coverage  
(Norm. ATAC Signal Range (0–0.9) by Reads in TSS)



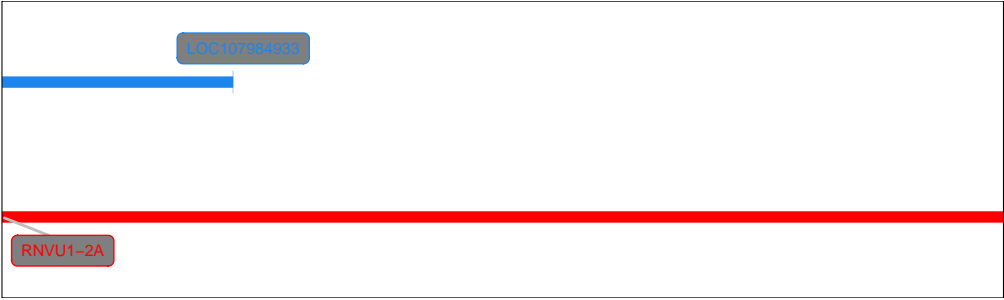
L3.CUX2.RORB



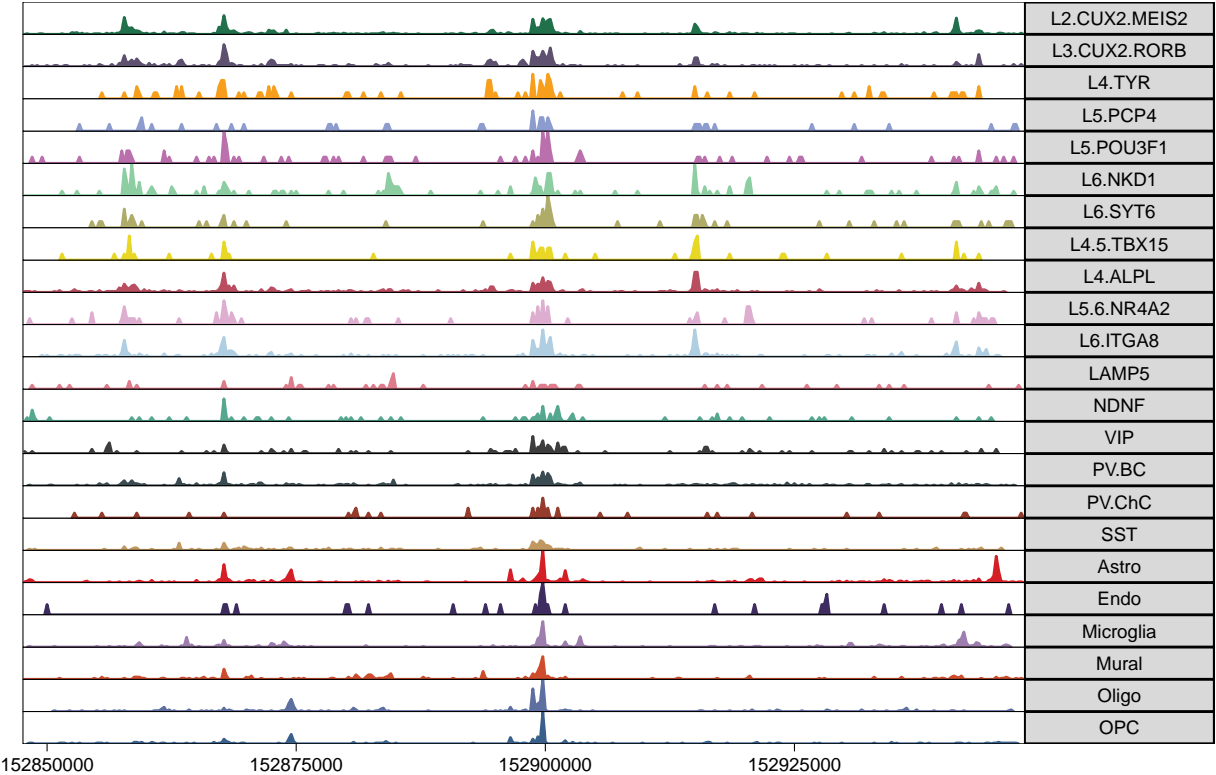
Peaks



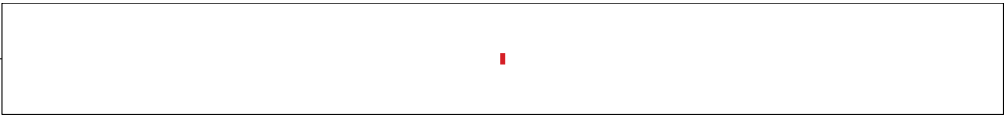
LoopTrack



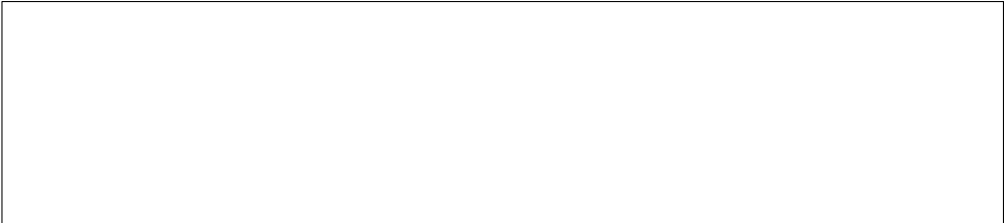
Coverage  
(Norm. ATAC Signal Range 0–1.23) by ReadsInTSS)



L3.CUX2.RORB



Peaks



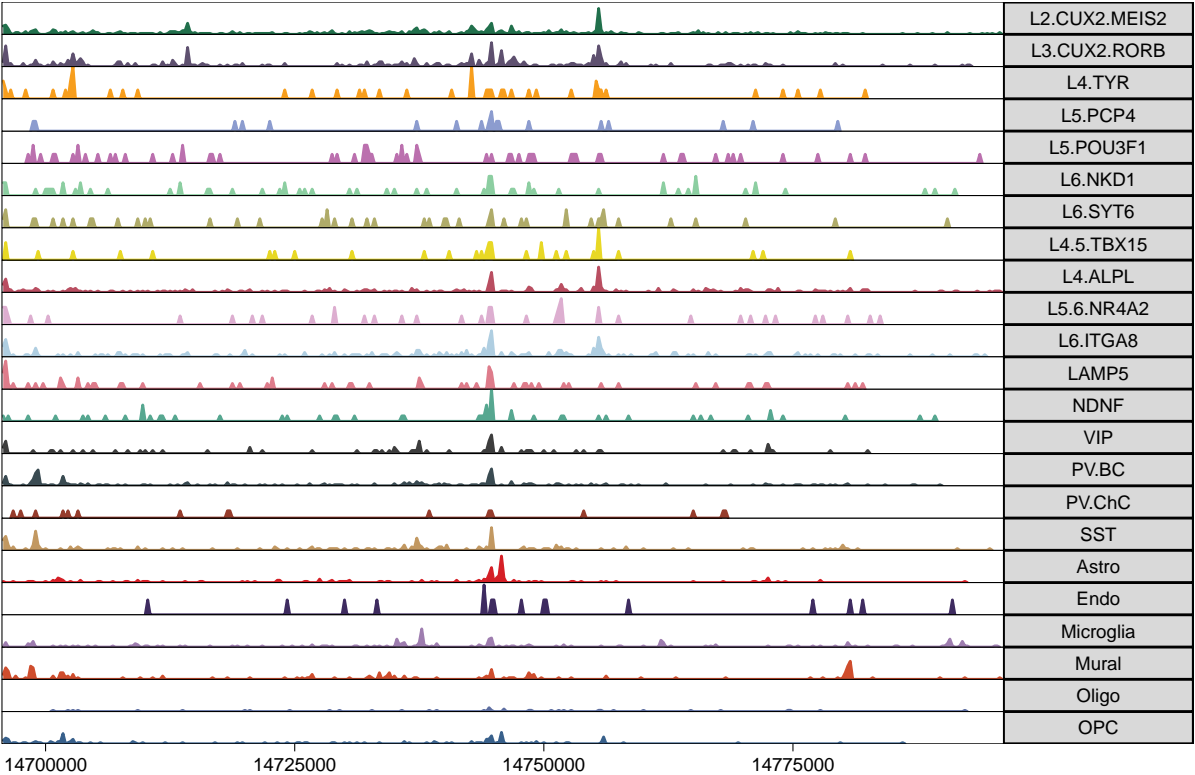
LoopTrack



Genes

chr6:14695622–14796123 Marker Peak Tracks for: L3.CUX2.RORB

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



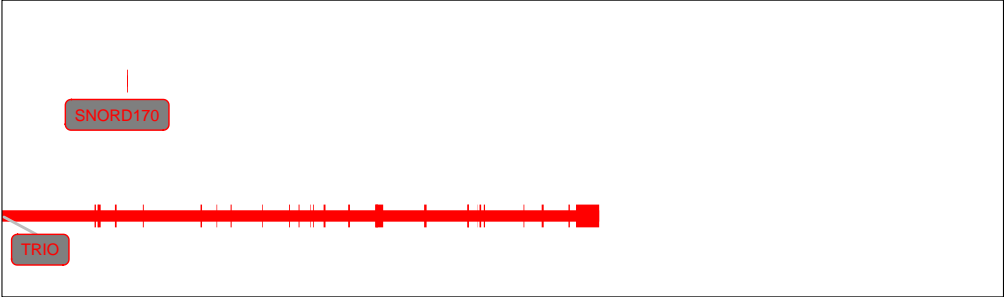
L3.CUX2.RORB



Peaks



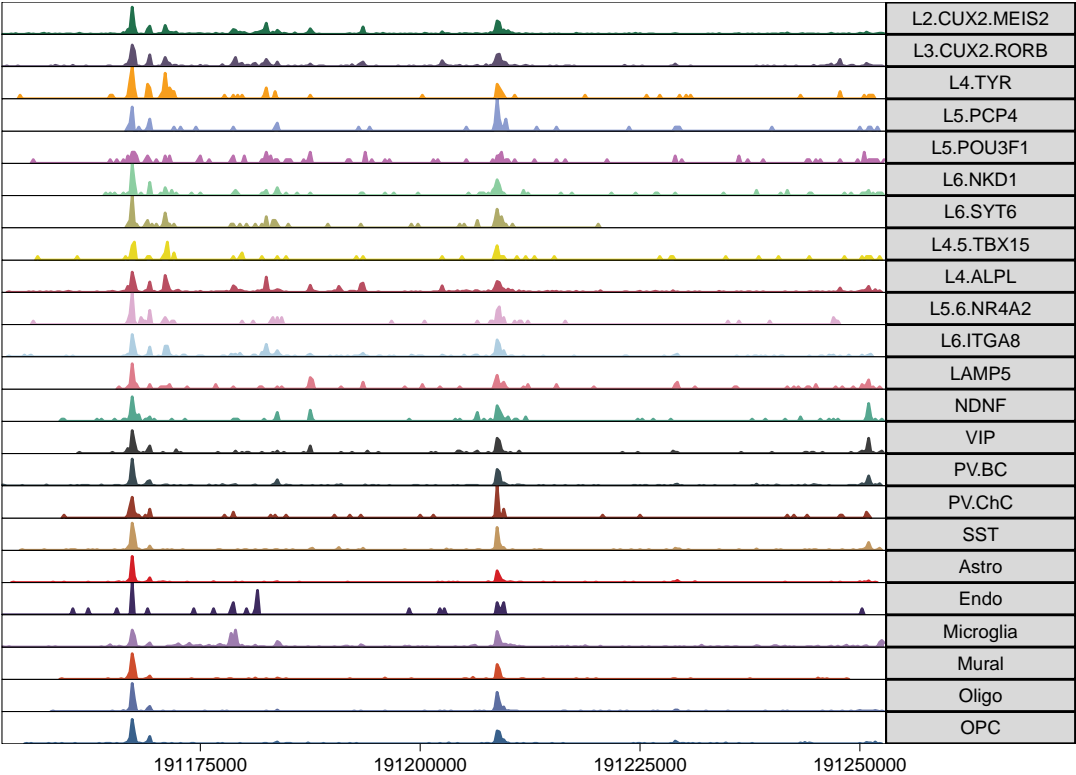
LoopTrack



Genes

chr1:191152439–191252940 Marker Peak Tracks for: L3.CUX2.RORB

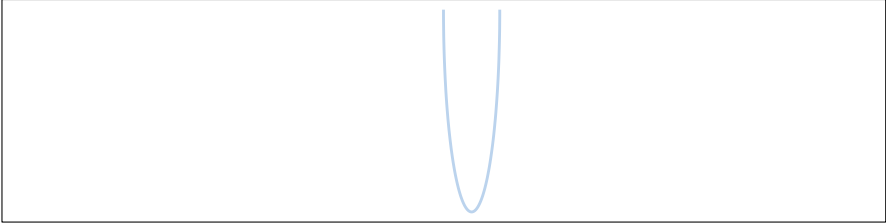
Coverage  
(Norm. ATAC Signal Range 0–2.06) by ReadsInTSS)



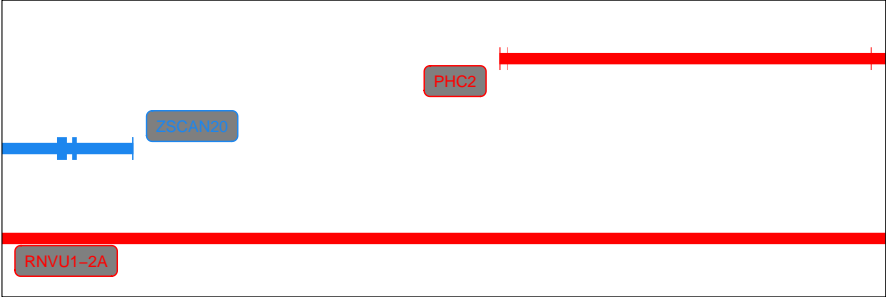
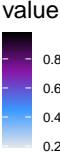
L3.CUX2.RORB



Peaks



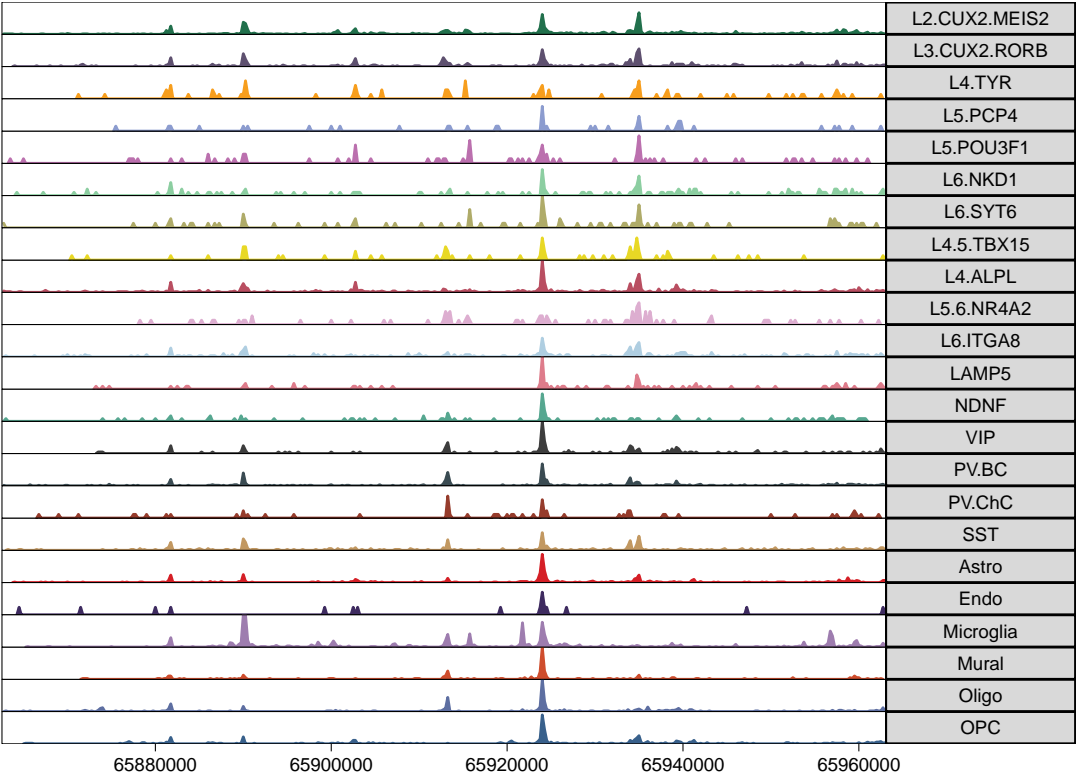
Peak2GeneLinks



Genes

chr9:65862553–65963054 Marker Peak Tracks for: L3.CUX2.RORB

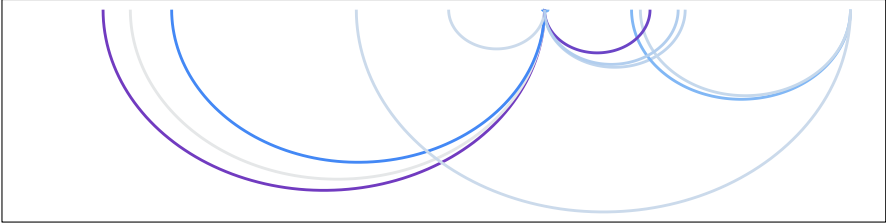
Coverage  
(Norm. ATAC Signal Range 0–1.67) by ReadsInTSS)



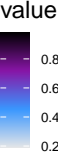
L3.CUX2.RORB



Peaks



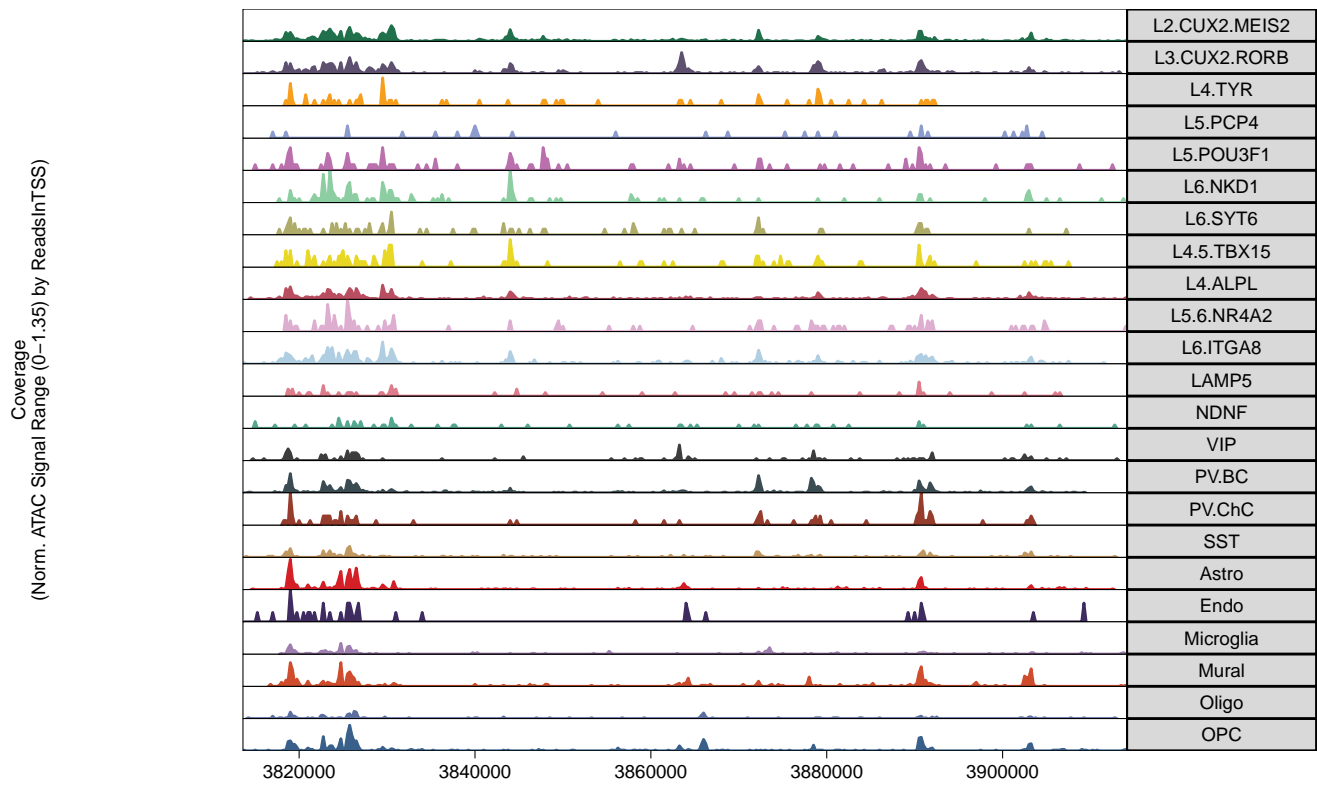
Peak2GeneLinks



Genes



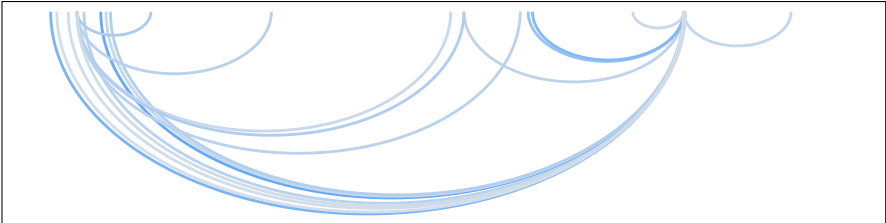
chr2:3813613–3914114 Marker Peak Tracks for: L3.CUX2.RORB



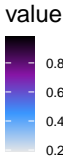
L3.CUX2.RORB



Peaks

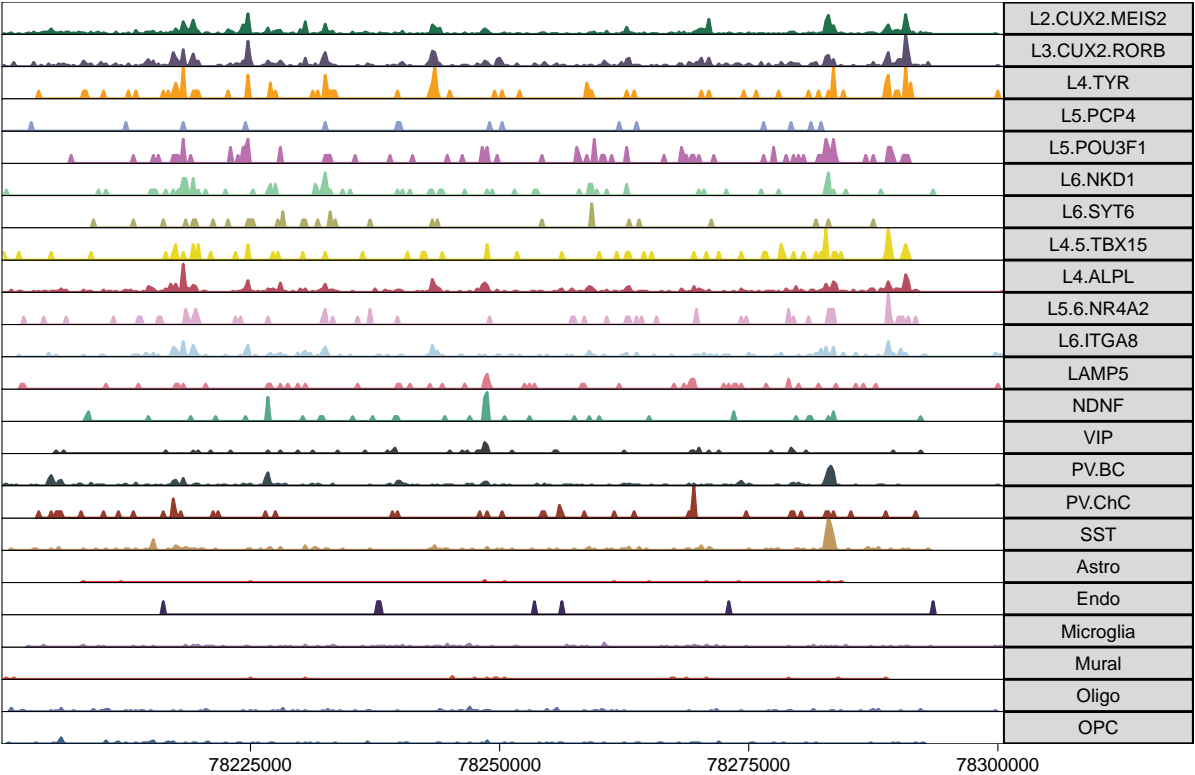


Peak2GeneLinks



Genes

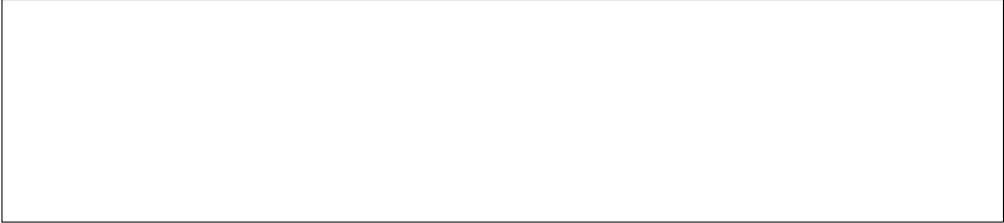
Coverage  
(Norm. ATAC Signal Range (0–0.95) by ReadsInTSS)



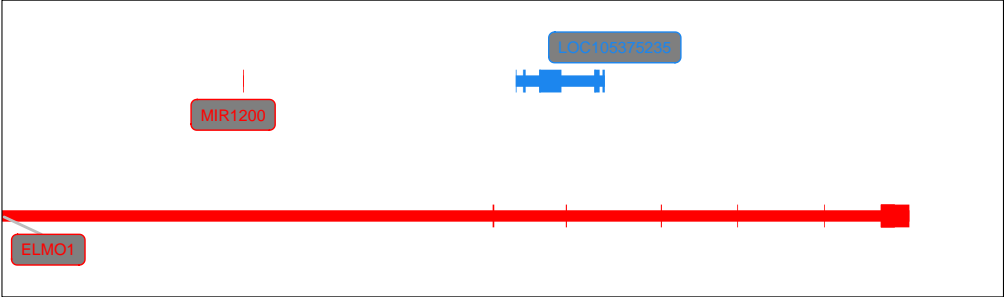
L3.CUX2.RORB



Peaks



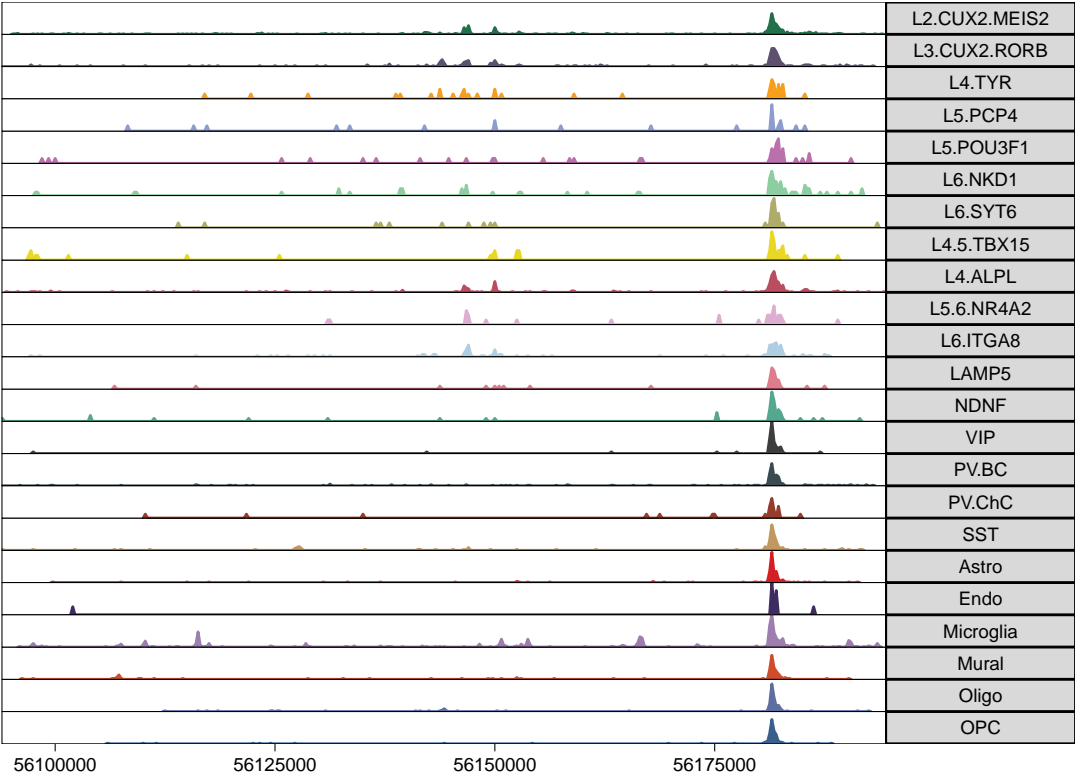
LoopTrack



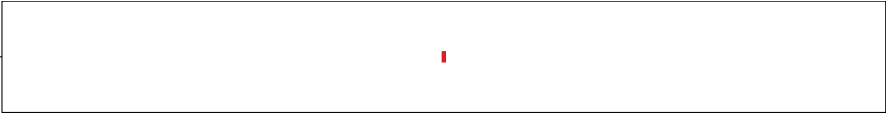
Genes

chr15:56093947–56194448 Marker Peak Tracks for: L3.CUX2.RORB

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



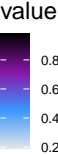
L3.CUX2.RORB



Peaks



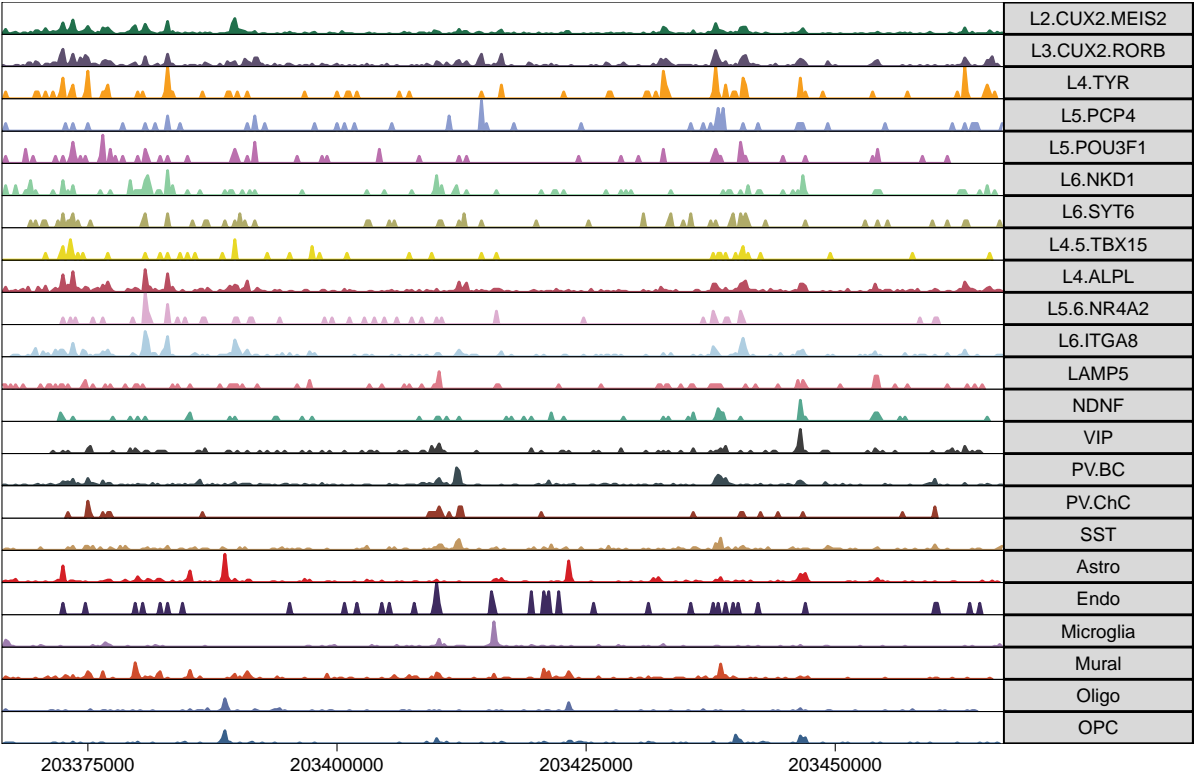
Peak2GeneLinks



Genes

chr1:203366378–203466879 Marker Peak Tracks for: L3.CUX2.RORB

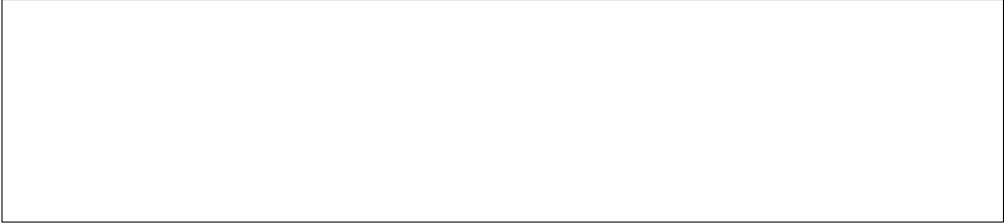
Coverage  
(Norm. ATAC Signal Range 0–1.09) by ReadsInTSS)



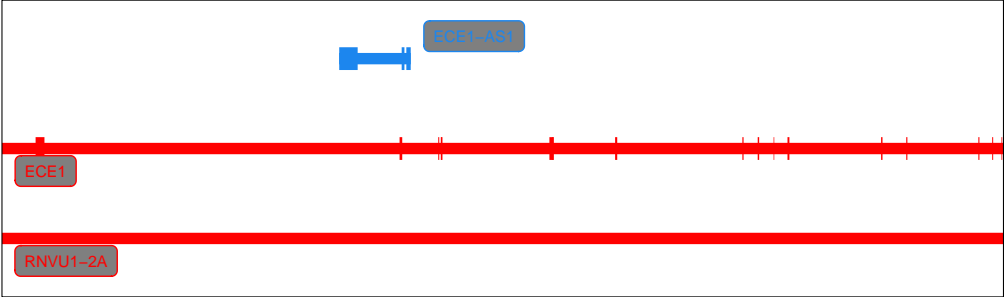
L3.CUX2.RORB



Peaks

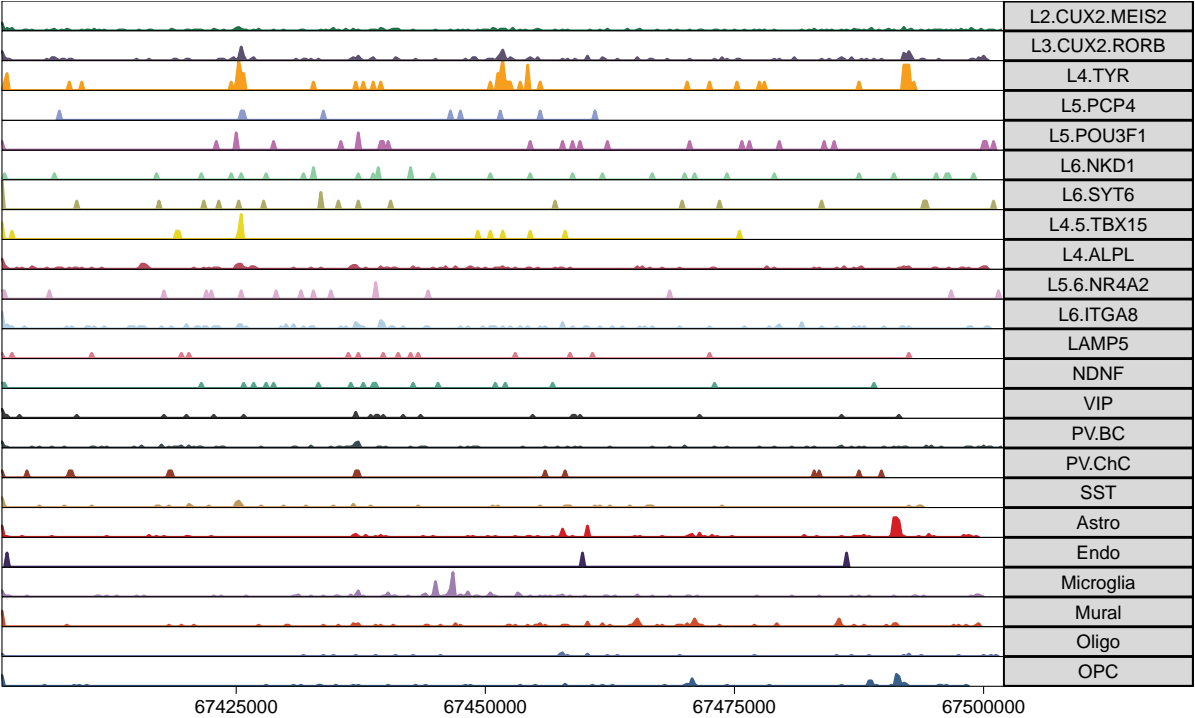


LoopTrack

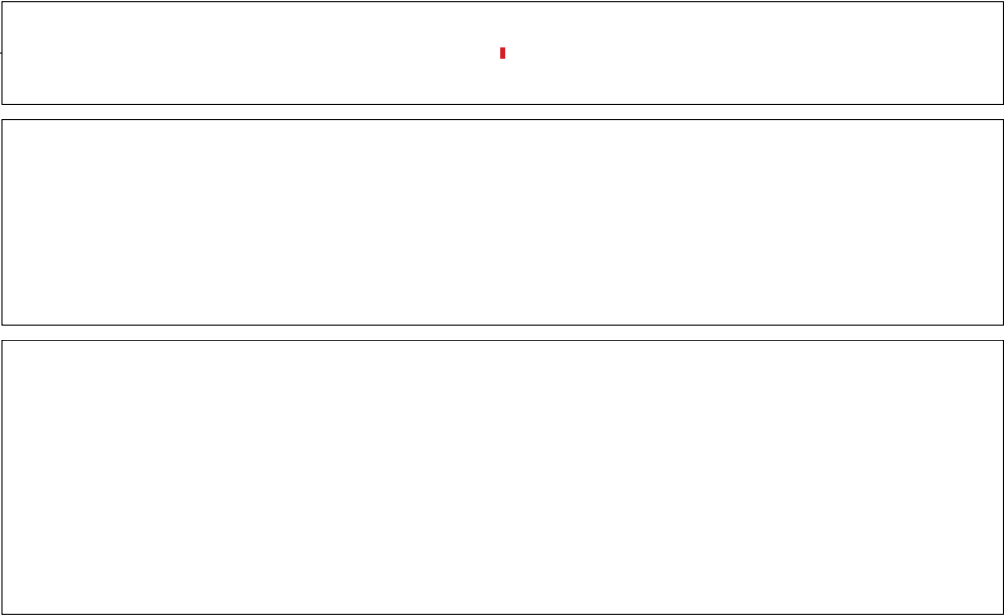


Genes

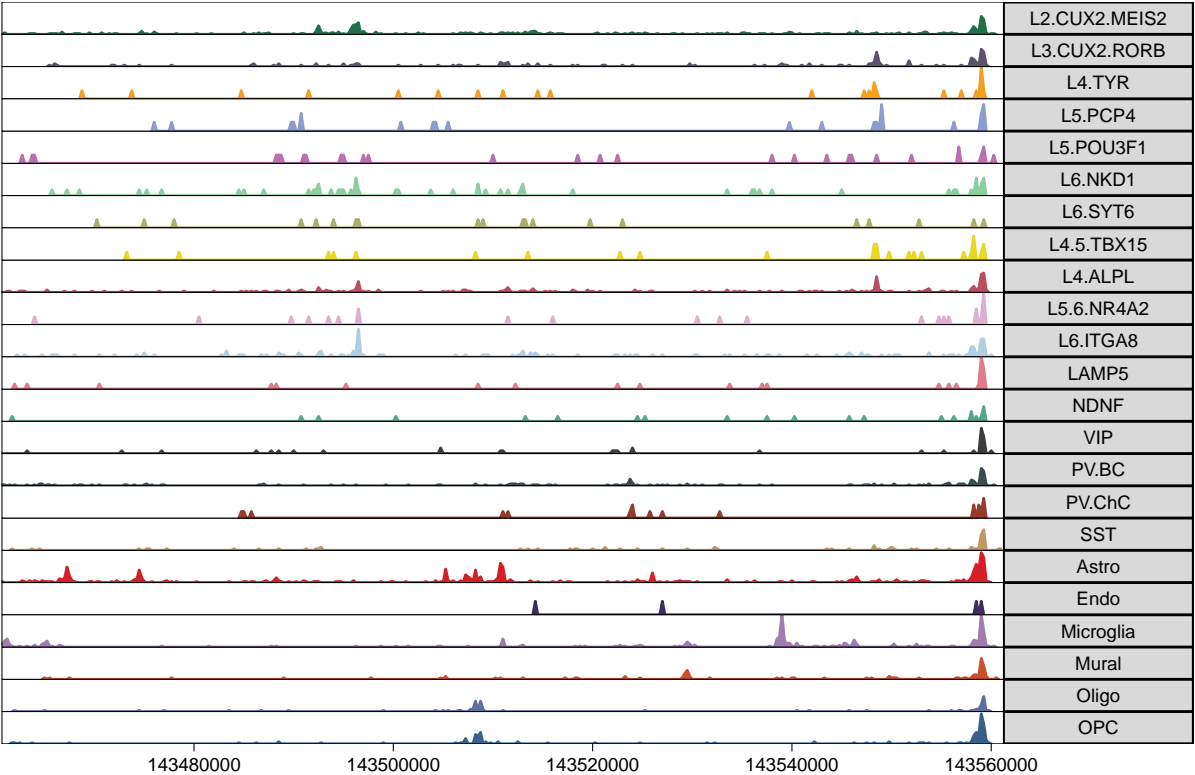
Coverage  
(Norm. ATAC Signal Range (0–0.82) by Reads in TSS)



L3.CUX2.RORB



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



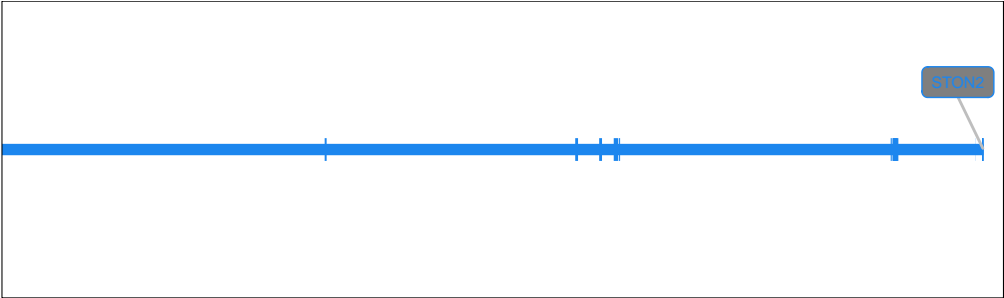
L3.CUX2.RORB



Peaks

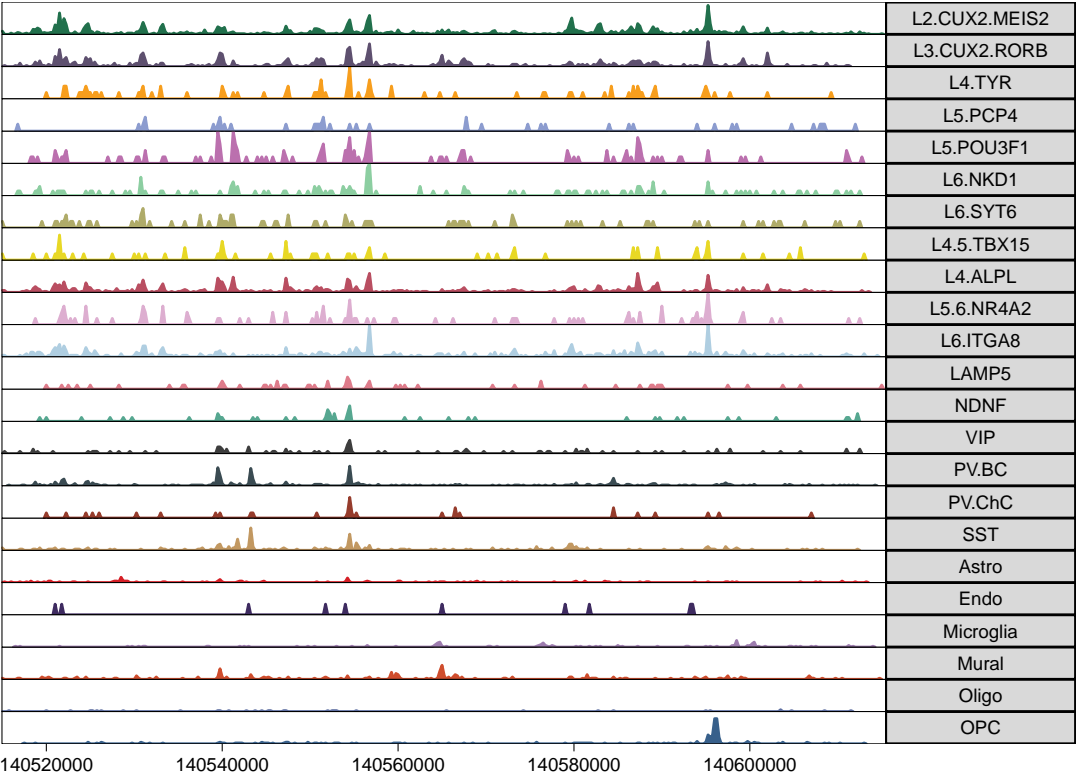


LoopTrack

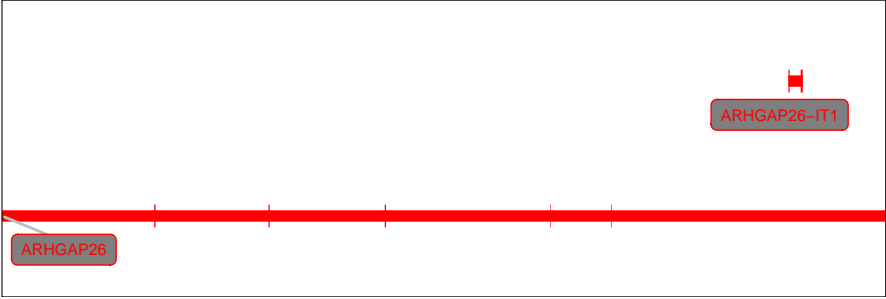
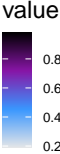
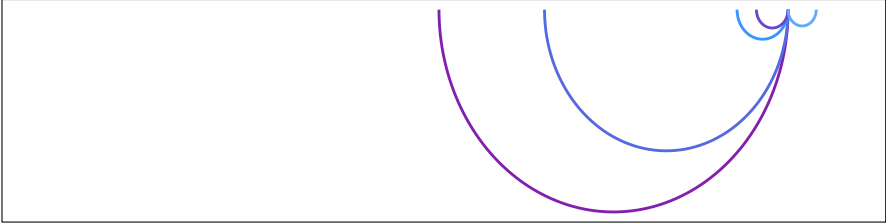


Genes

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

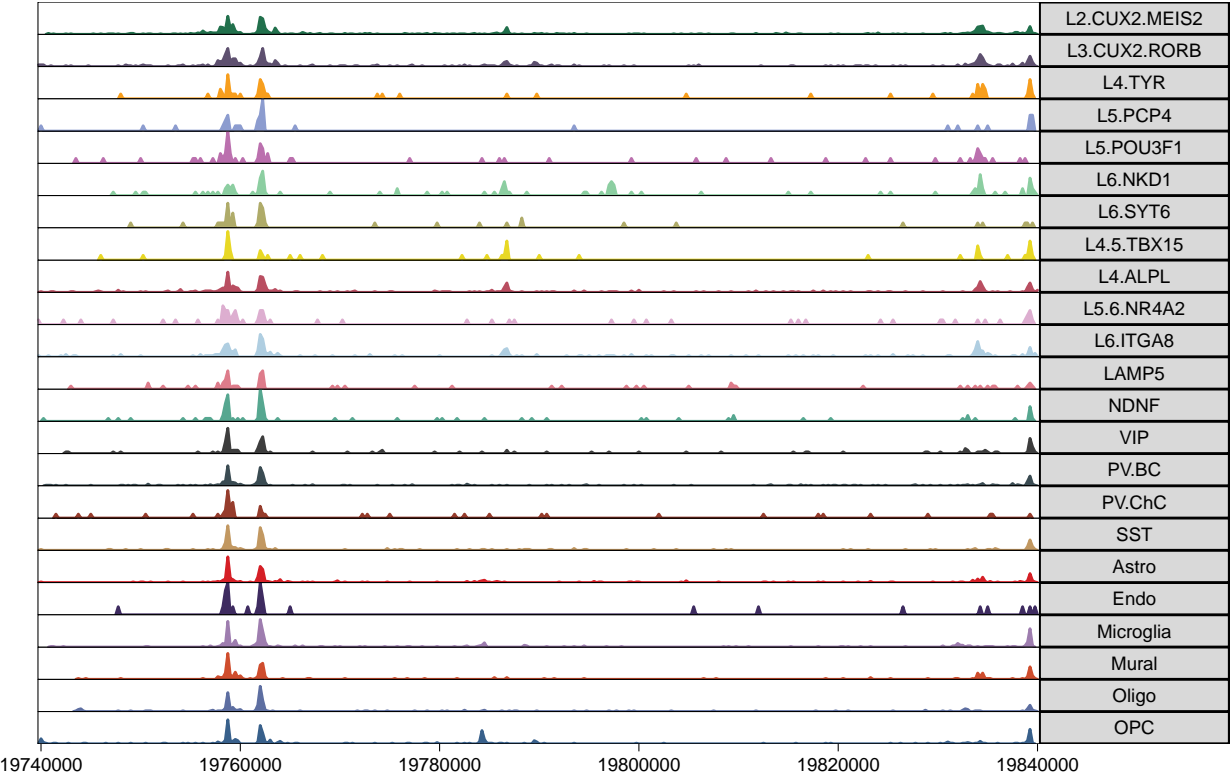


L3.CUX2.RORB



chr10:19739694–19840195 Marker Peak Tracks for: L3.CUX2.RORB

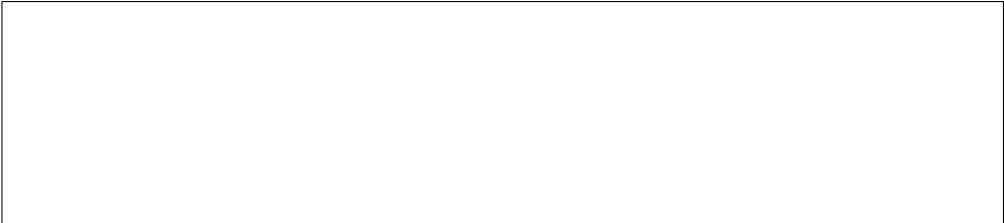
Coverage  
(Norm. ATAC Signal Range 0–1.56) by ReadsInTSS)



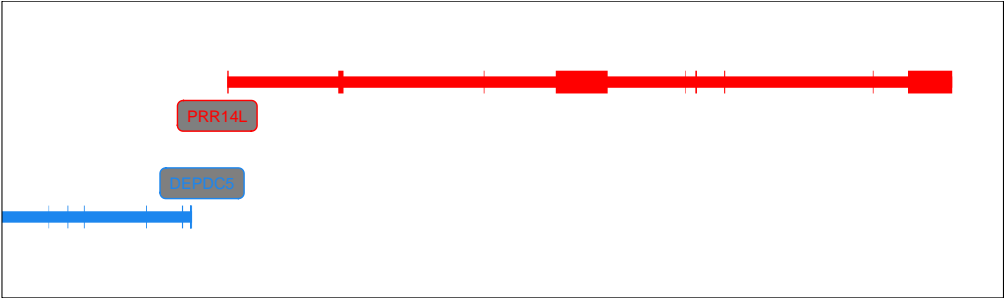
L3.CUX2.RORB



Peaks



LoopTrack

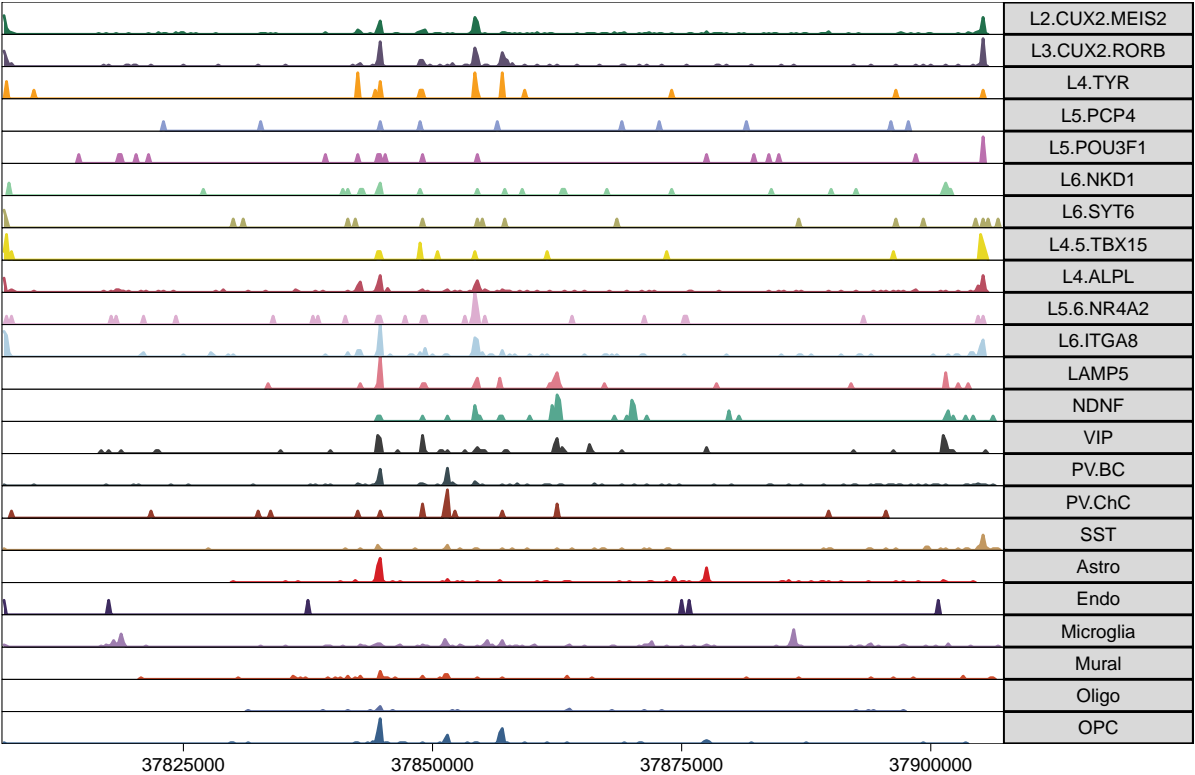


Genes



chr18:37806792–37907293 Marker Peak Tracks for: L3.CUX2.RORB

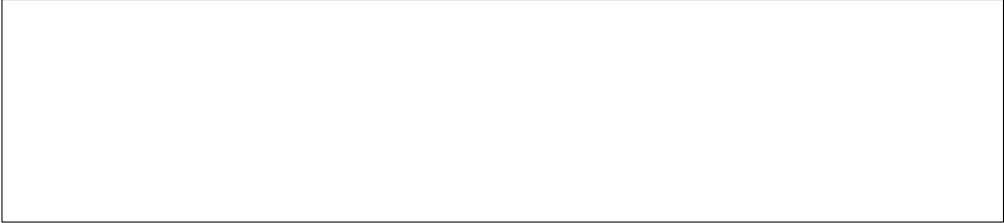
Coverage  
(Norm. ATAC Signal Range (0–0.87) by ReadsInTSS)



L3.CUX2.RORB



Peaks



LoopTrack



Genes