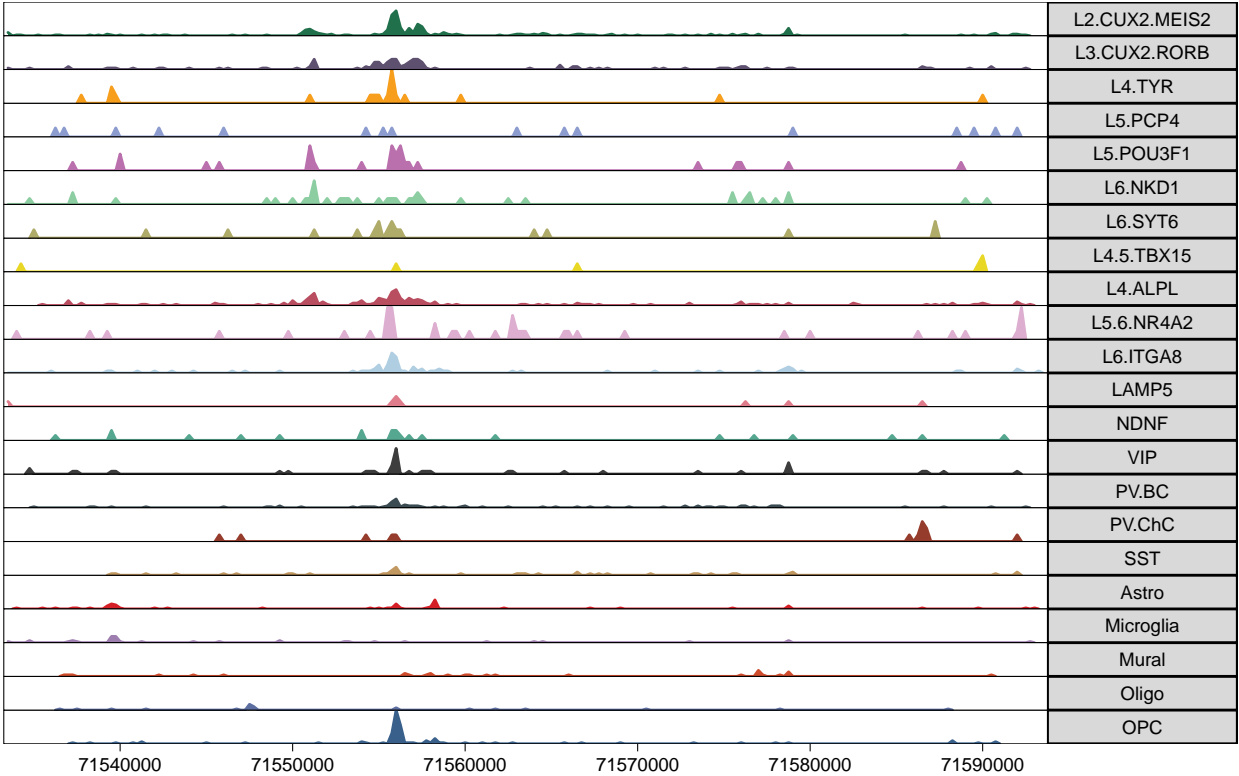
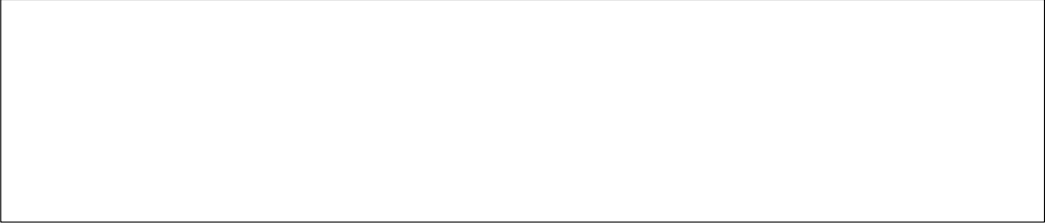
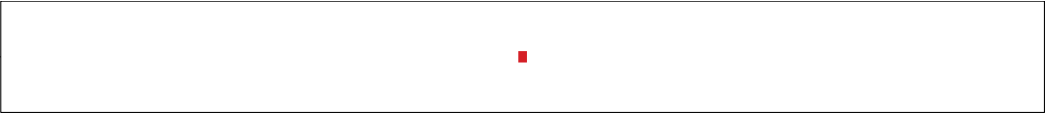


chr7:71533255–71593756 Marker Peak Tracks for: L5.6.NR4A2

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

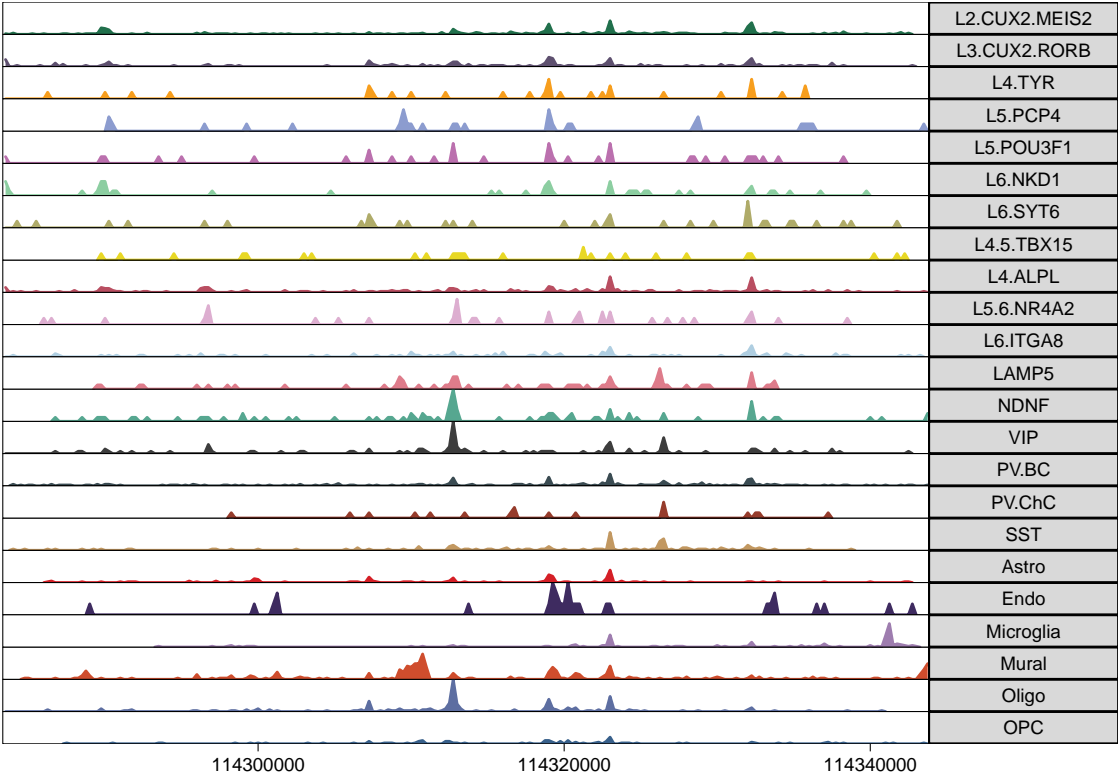


L5.6.NR4A2

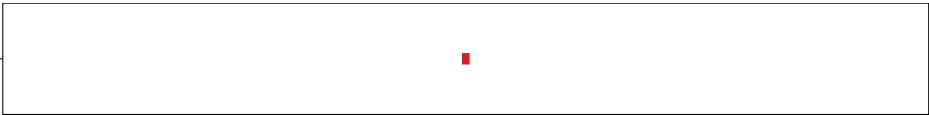


chr2:114283318-114343819 Marker Peak Tracks for: L5.6.NR4A2

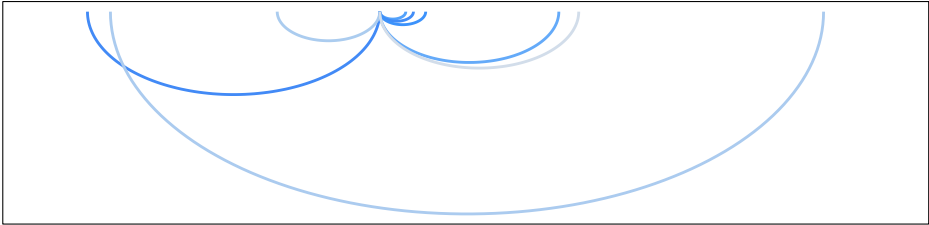
Coverage
(Norm. ATAC Signal Range 0-1.15) by ReadsInTSS)



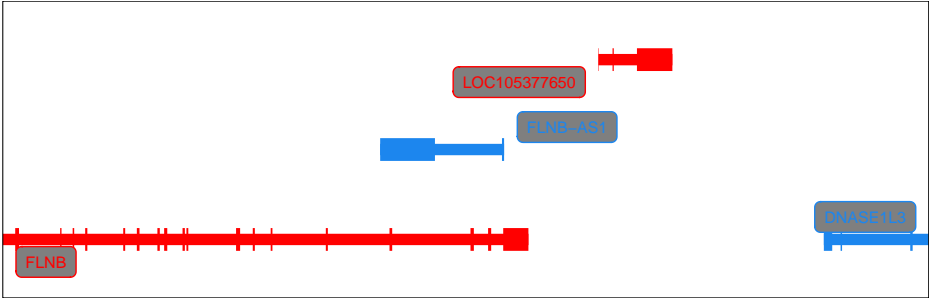
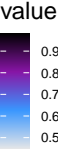
L5.6.NR4A2



Peaks

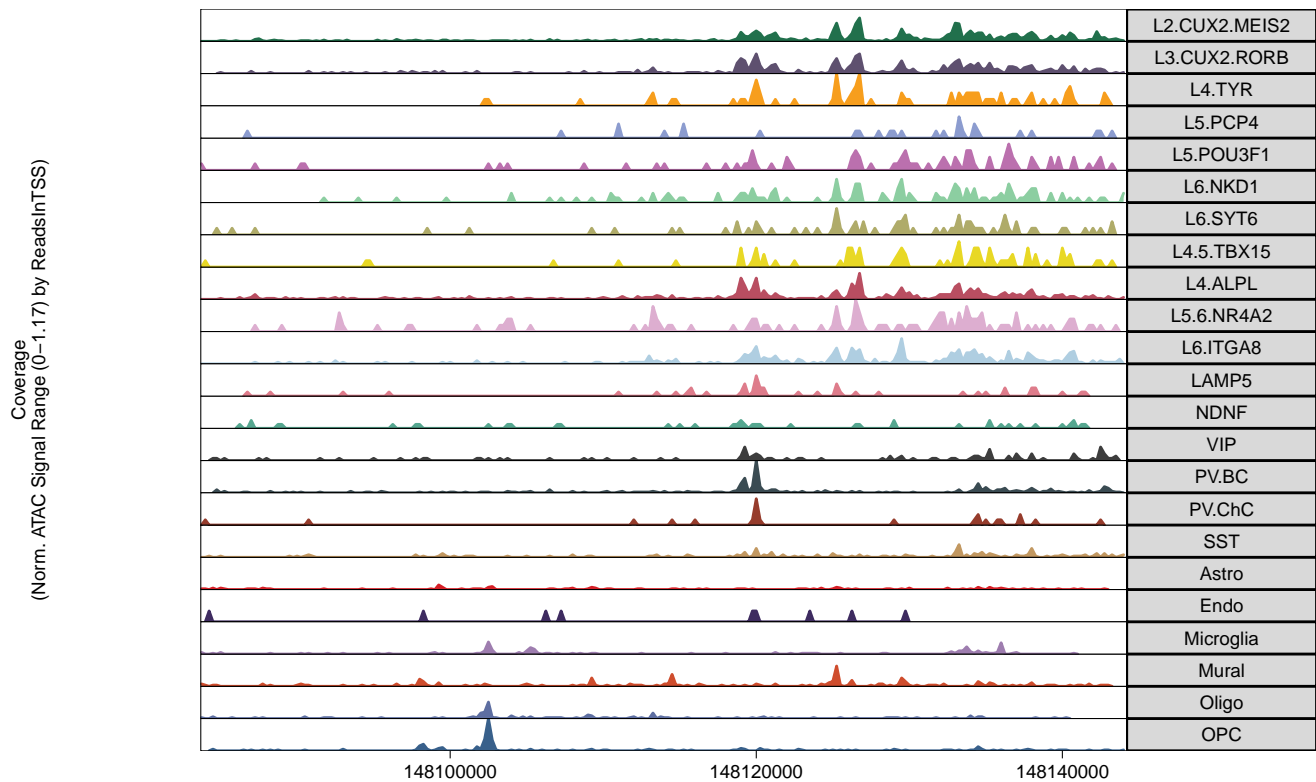


Peak2GeneLinks



Genes

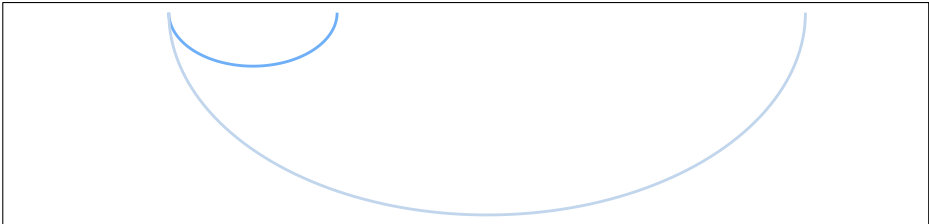
chr6:148083704–148144205 Marker Peak Tracks for: L5.6.NR4A2



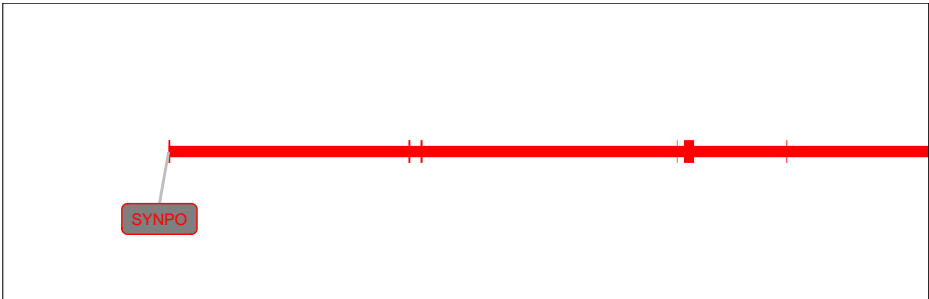
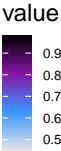
L5.6.NR4A2



Peaks

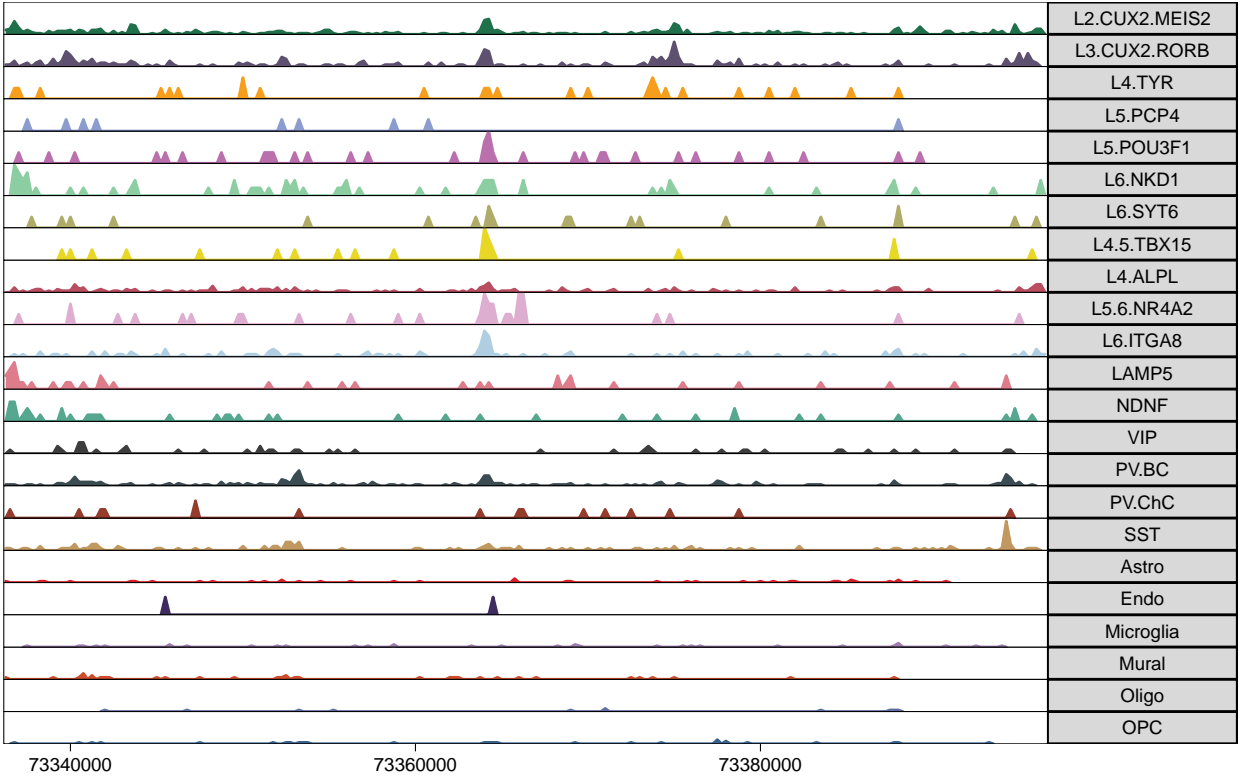


Peak2GeneLinks

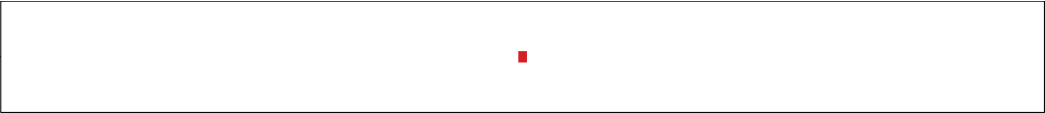


Genes

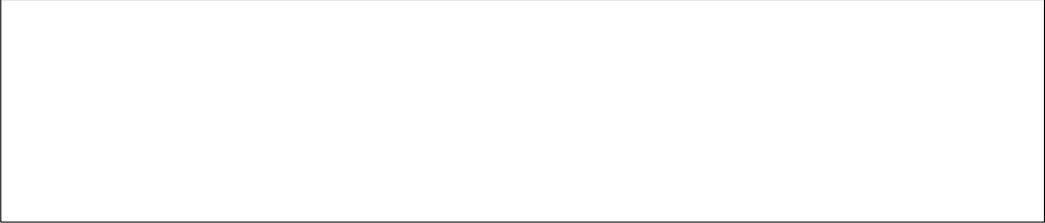
Coverage
(Norm. ATAC Signal Range (0–0.71) by ReadsInTSS)



L5.6.NR4A2



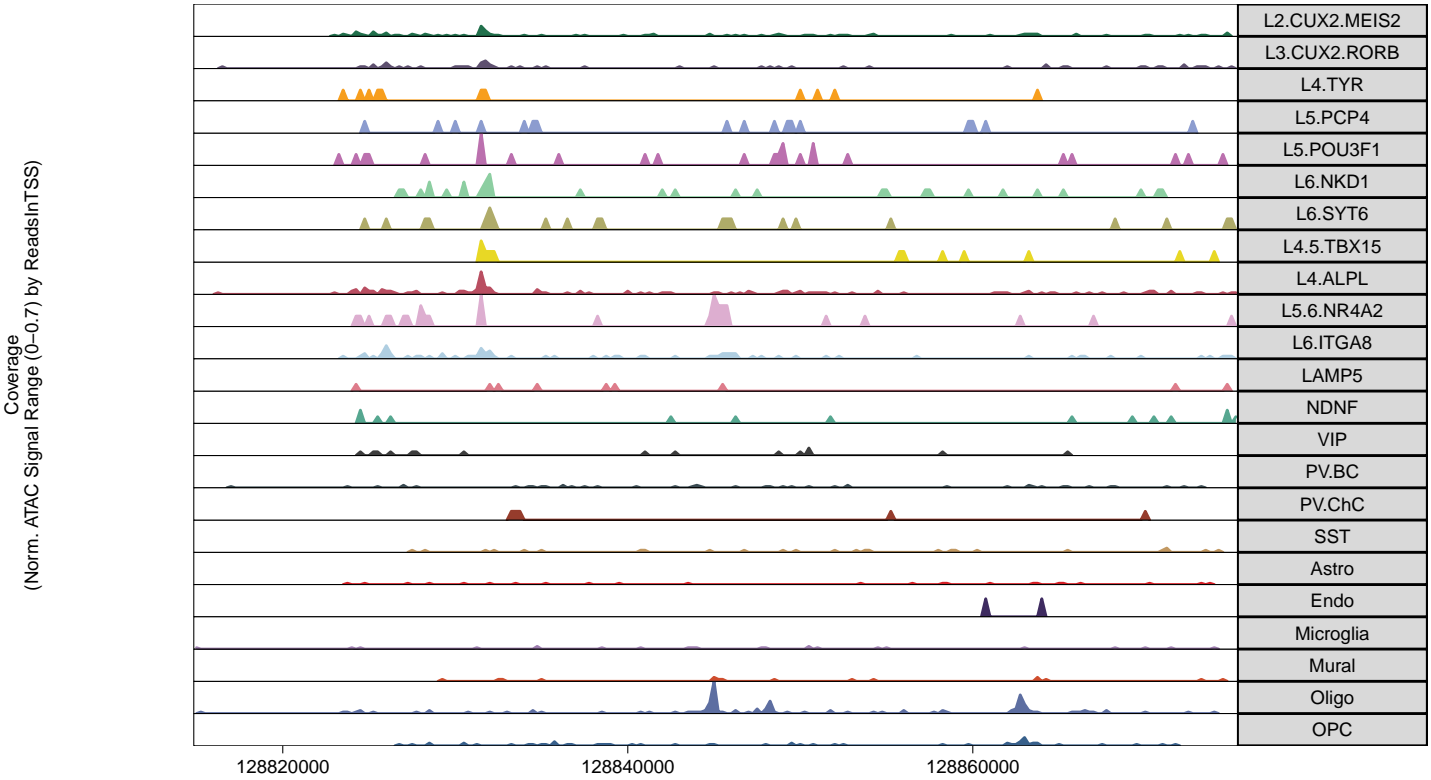
Peaks



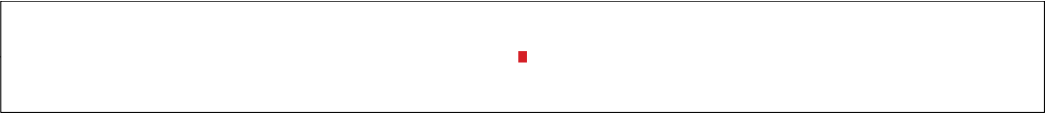
LoopTrack



Genes



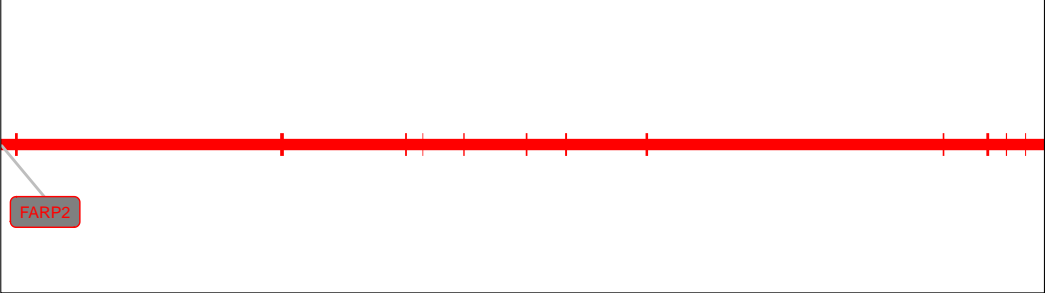
L5.6.NR4A2



Peaks

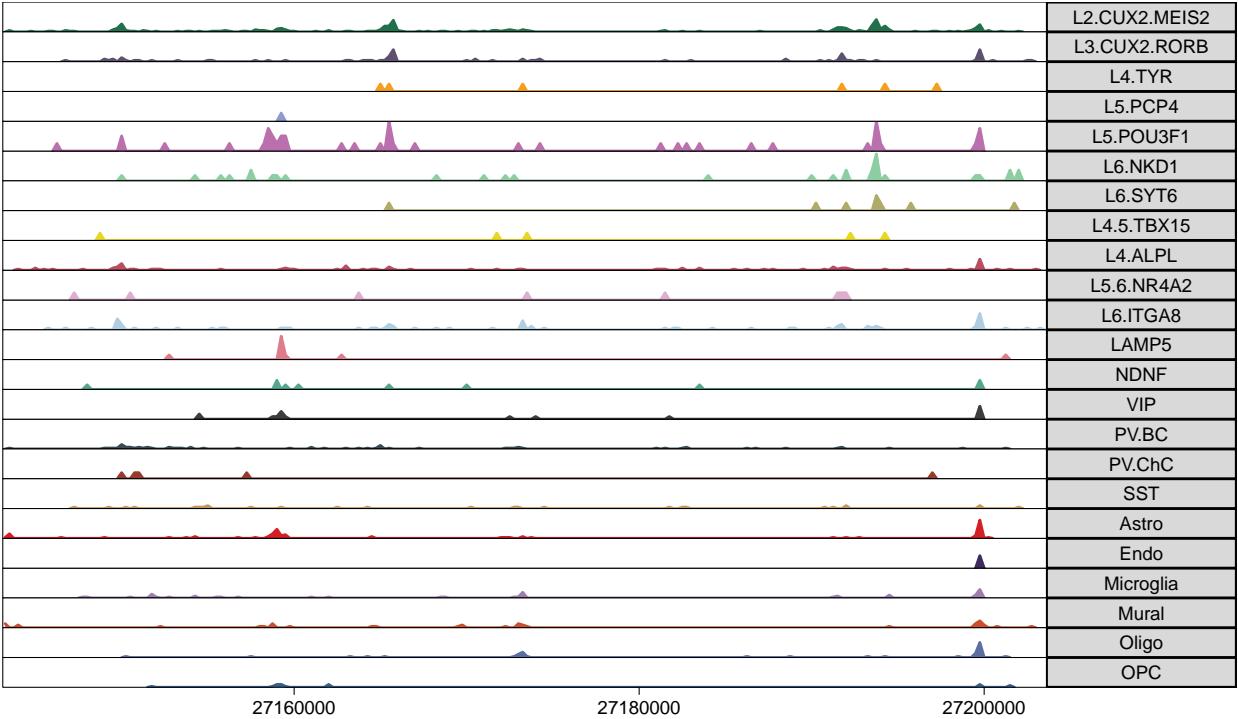


LoopTrack

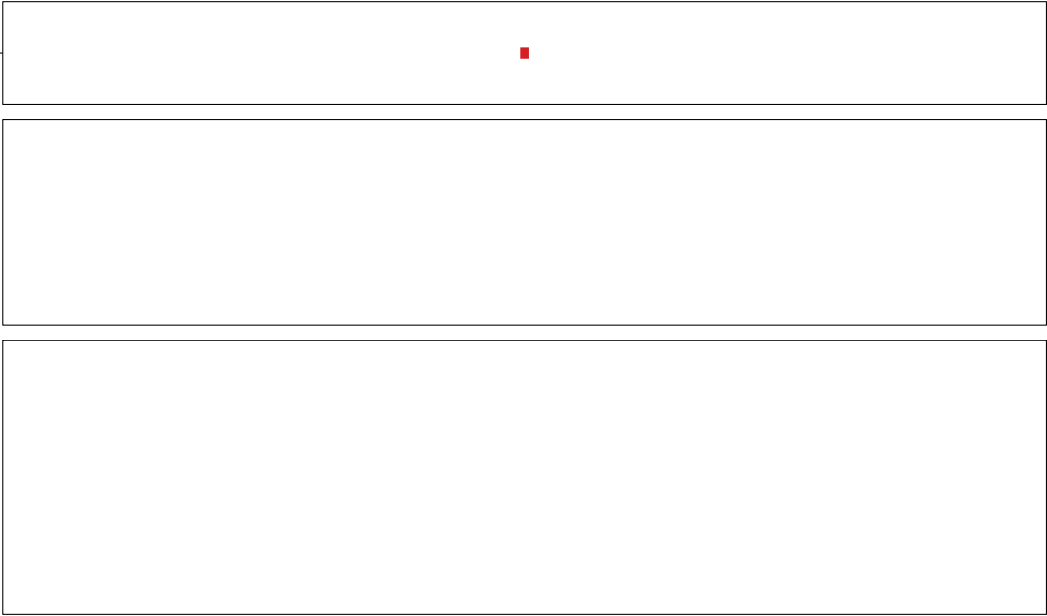


Genes

Coverage
(Norm. ATAC Signal Range (0–0.94) by Reads in TSS)



L5.6.NR4A2

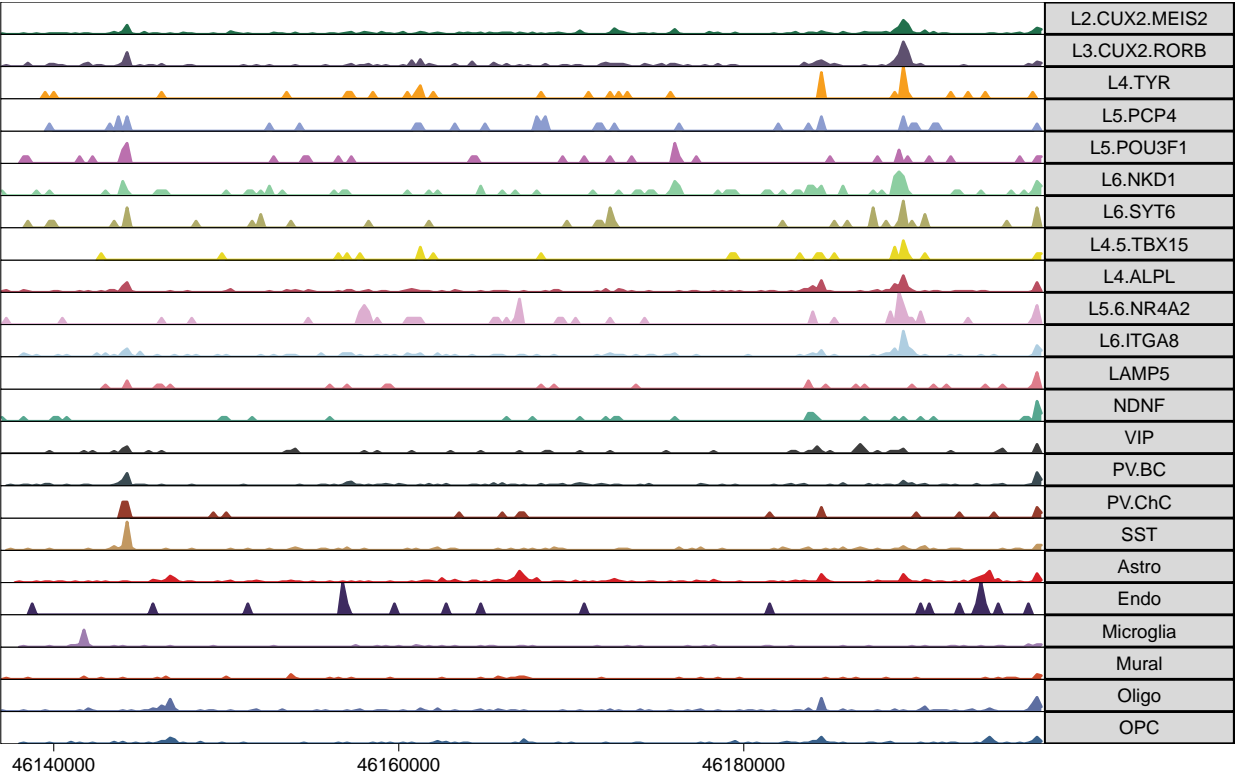


Peaks

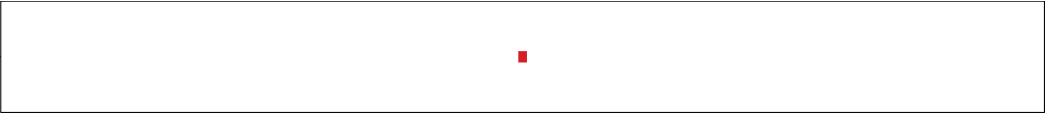
LoopTrack

GeneTrack

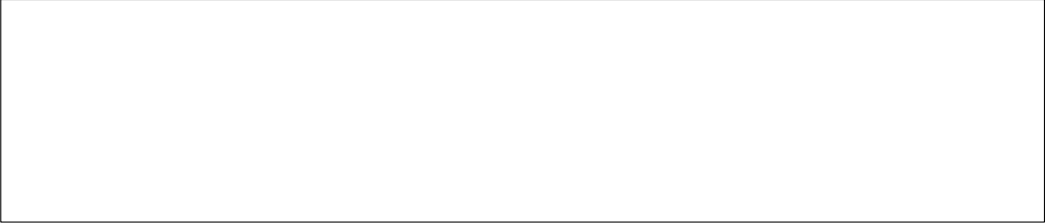
Coverage
(Norm. ATAC Signal Range 0-1.15) by ReadsInTSS)



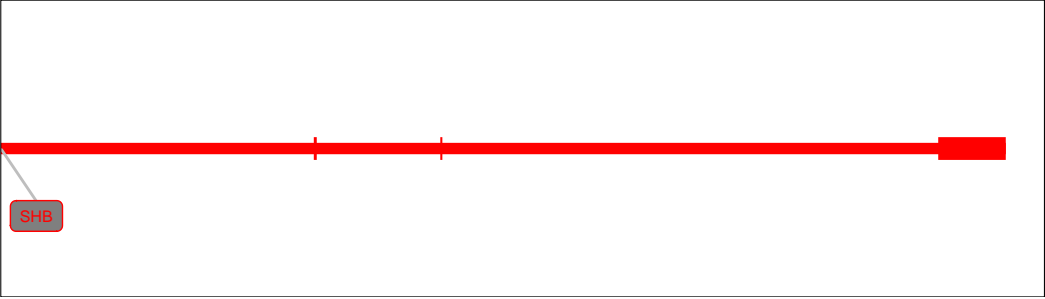
L5.6.NR4A2



Peaks



LoopTrack

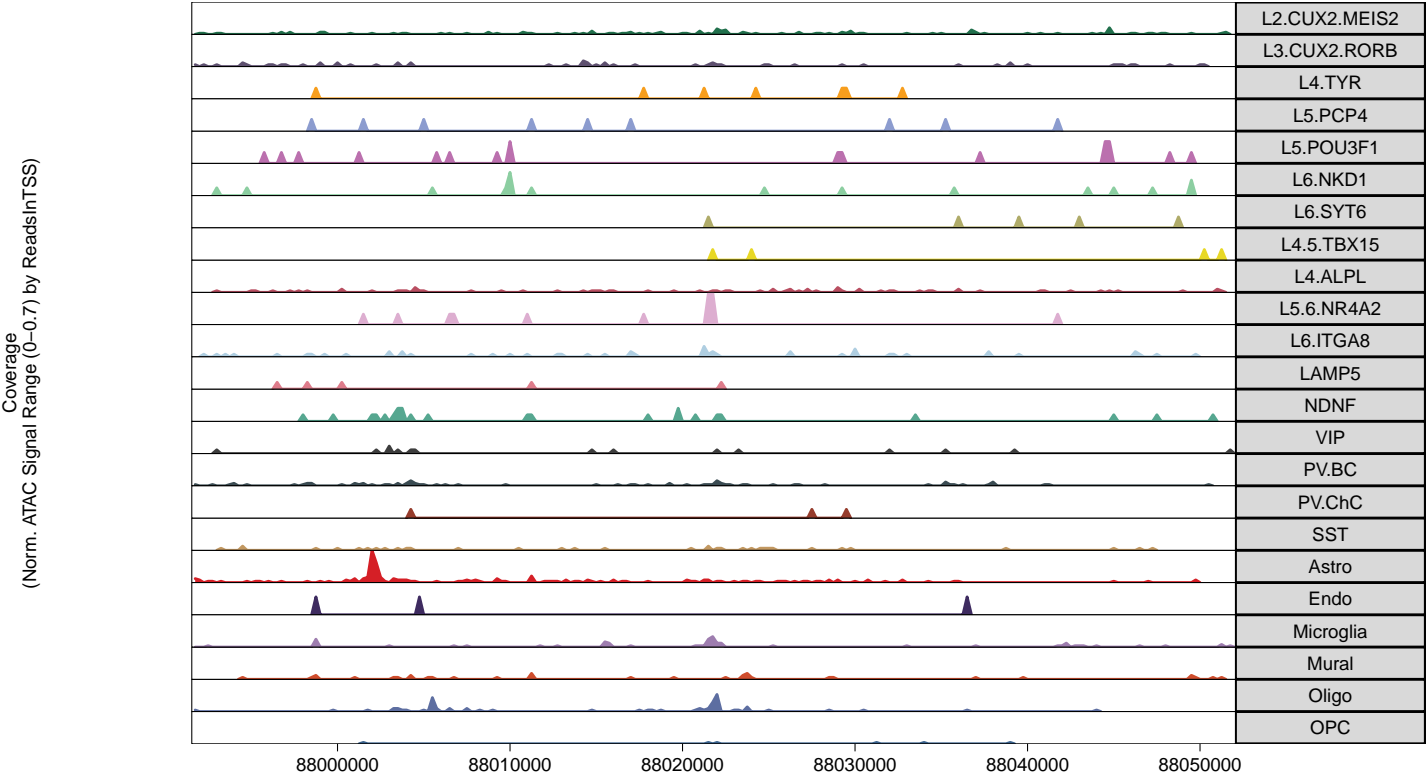


Genes

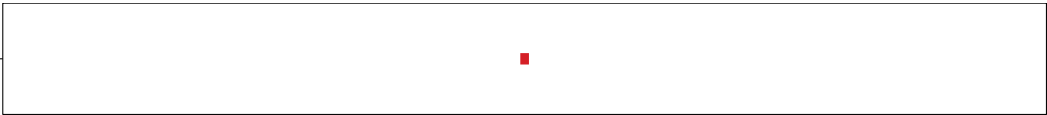
L5.6.NR4A2

LoopTrack

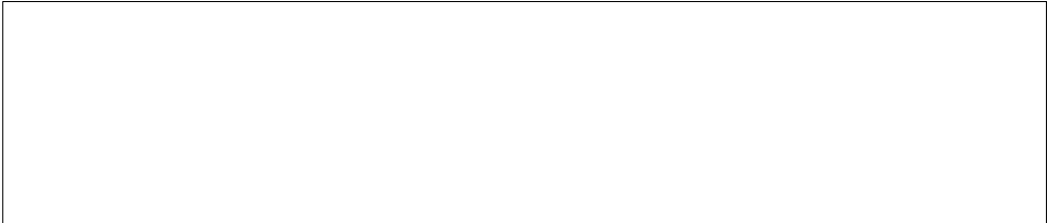
The diagram illustrates a horizontal bar with a blue segment on the left and a grey segment on the right. The blue segment is labeled 'H' and 'LOCALIZATION'. The grey segment is labeled 'CHPLM' and 'FROM'.



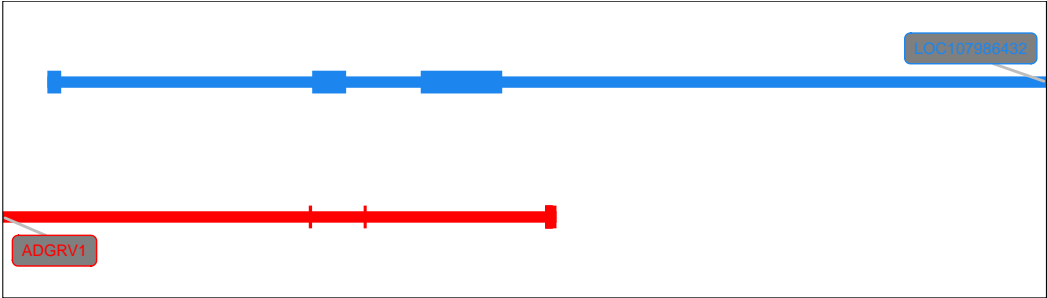
L5.6.NR4A2



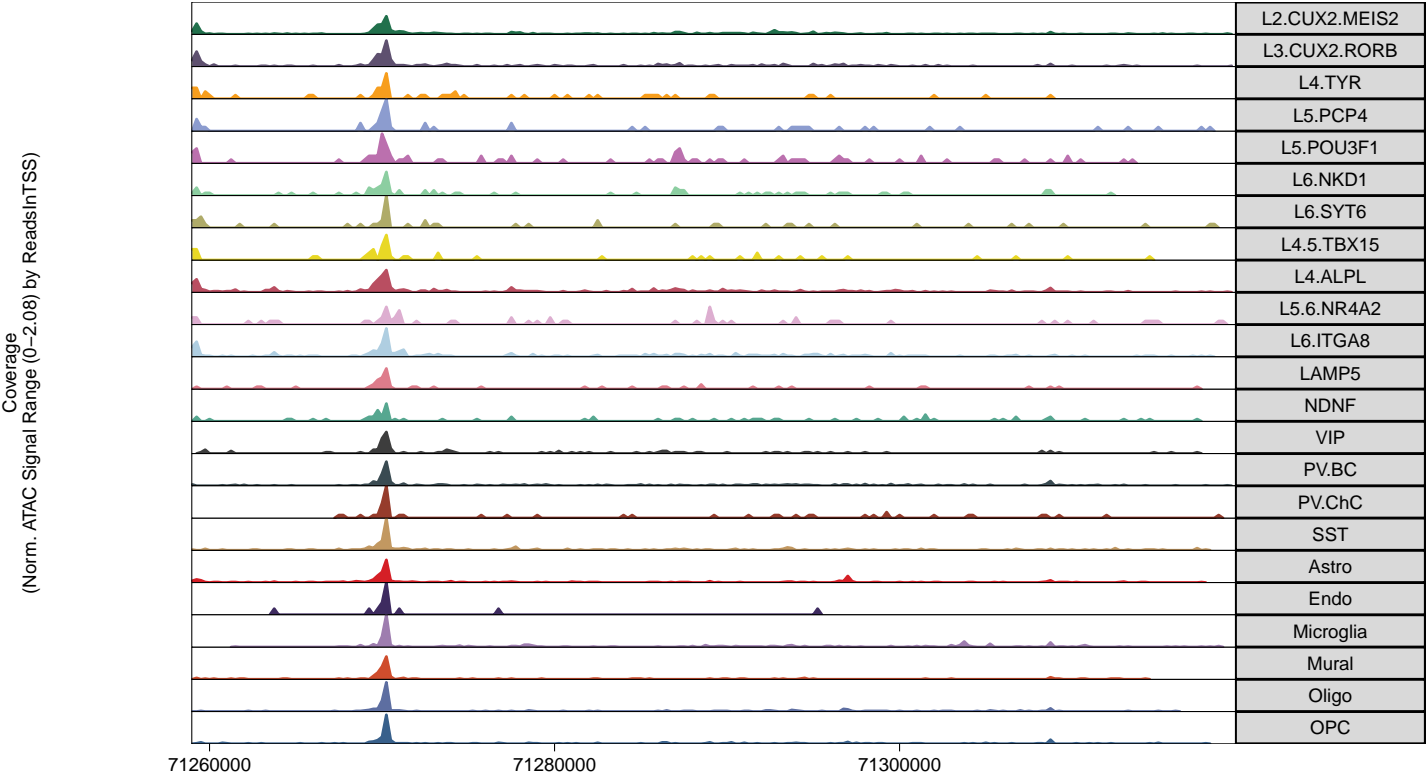
Peaks



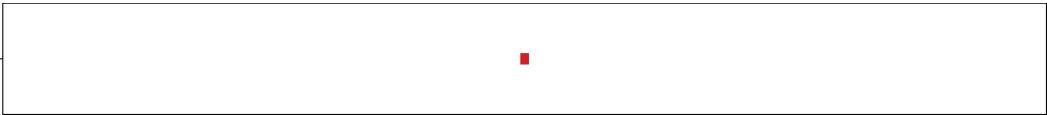
LoopTrack



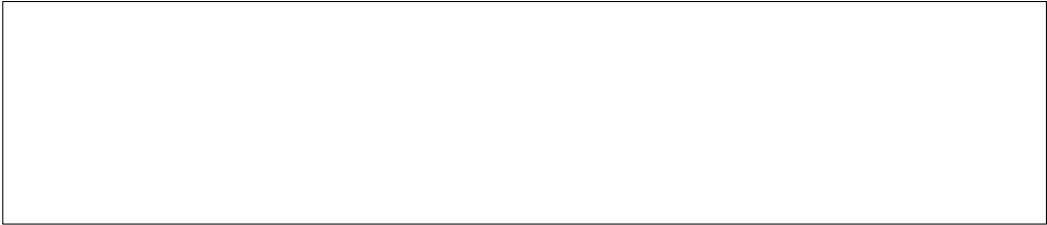
Genes



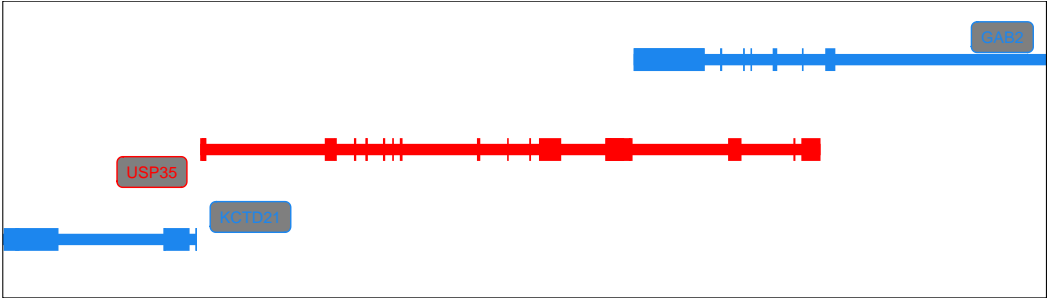
L5.6.NR4A2



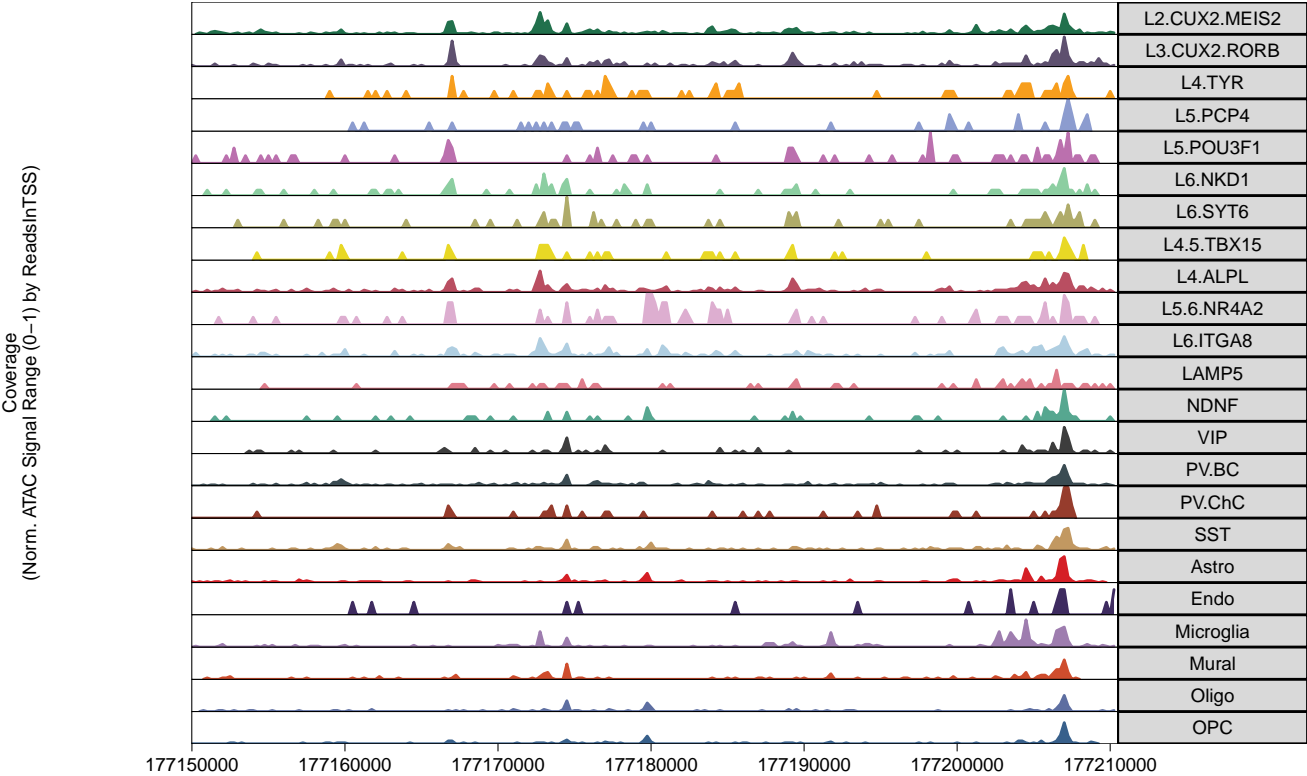
Peaks



LoopTrack



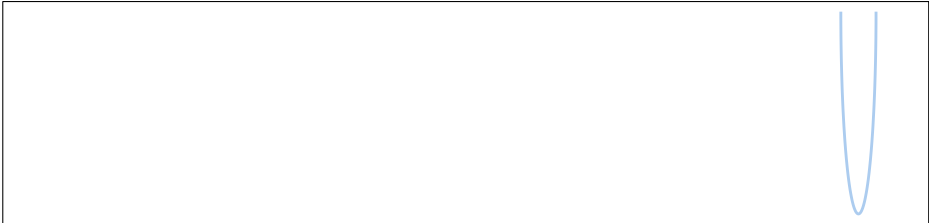
Genes



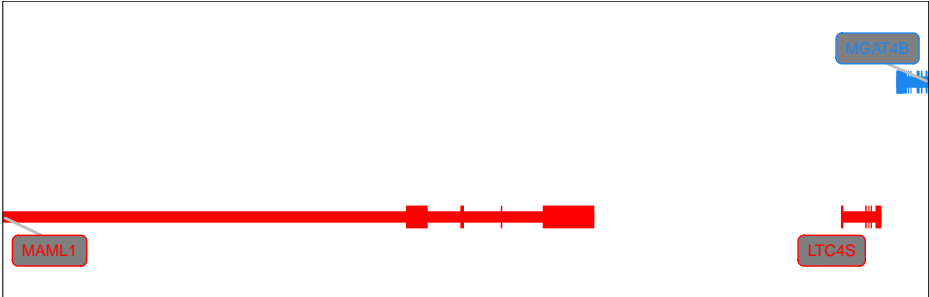
L5.6.NR4A2



Peaks



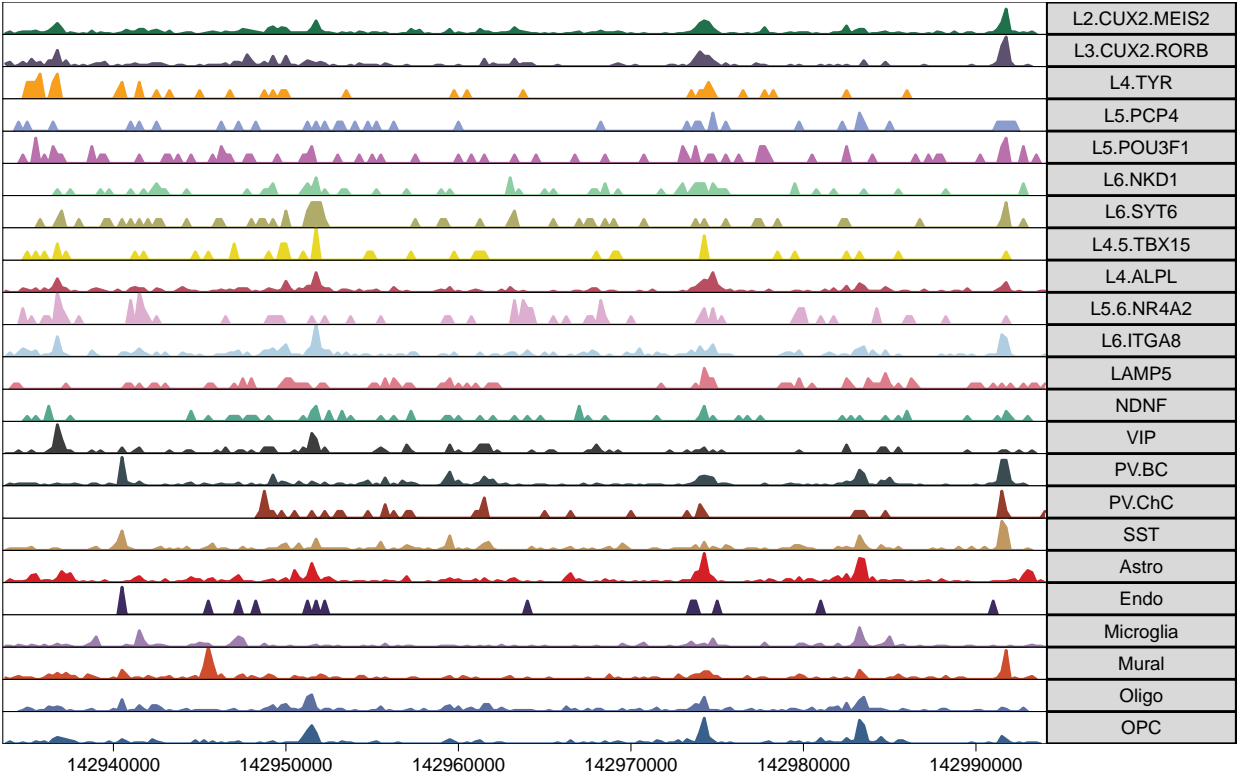
Peak2GeneLinks



Genes

chr2:142933586–142994087 Marker Peak Tracks for: L5.6.NR4A2

Coverage
(Norm. ATAC Signal Range (0–0.91) by ReadsInTSS)



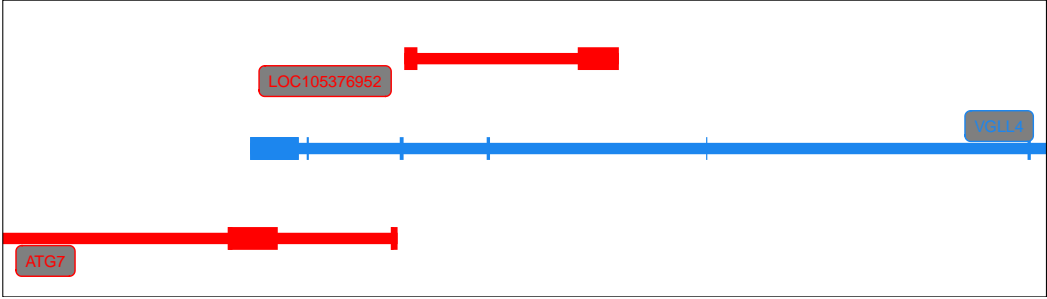
L5.6.NR4A2



Peaks



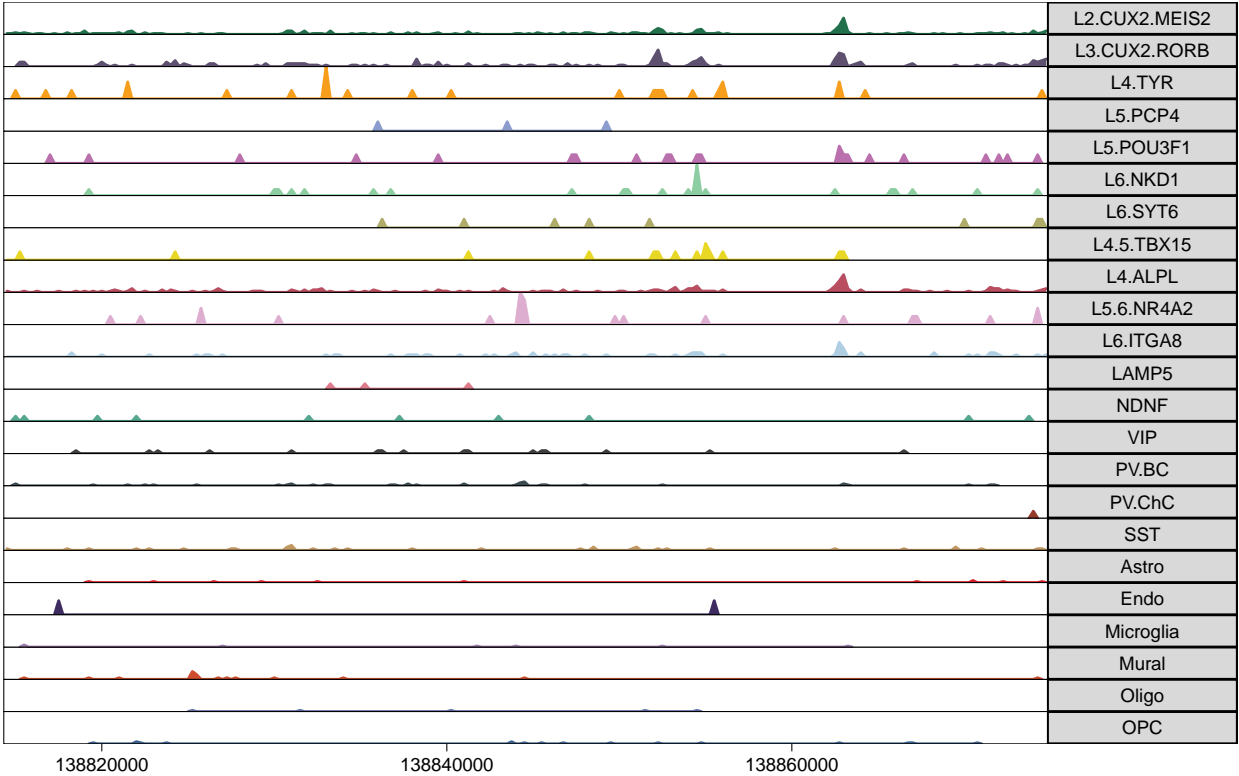
LoopTrack



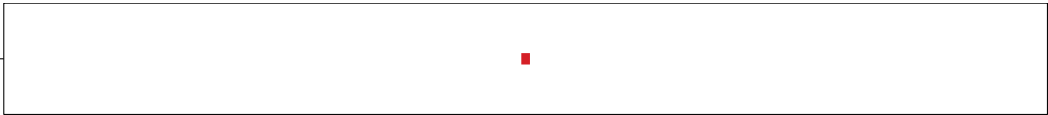
Genes

chr2:138814317-138874818 Marker Peak Tracks for: L5.6.NR4A2

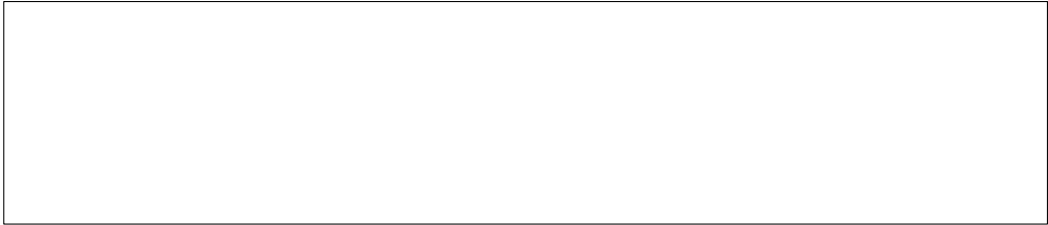
Coverage
(Norm. ATAC Signal Range (0-0.89) by ReadsInTSS)



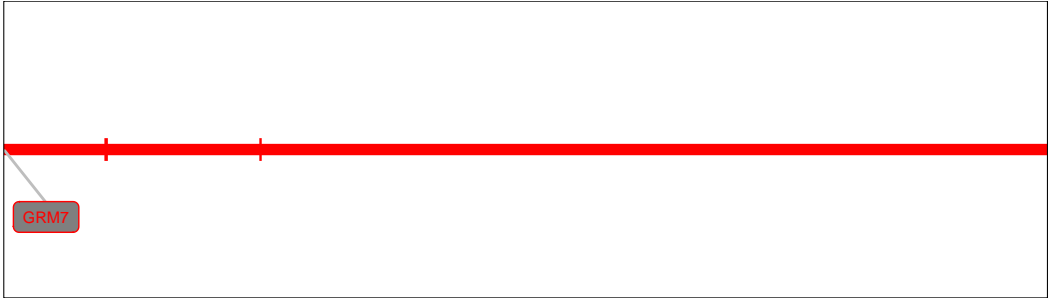
L5.6.NR4A2



Peaks

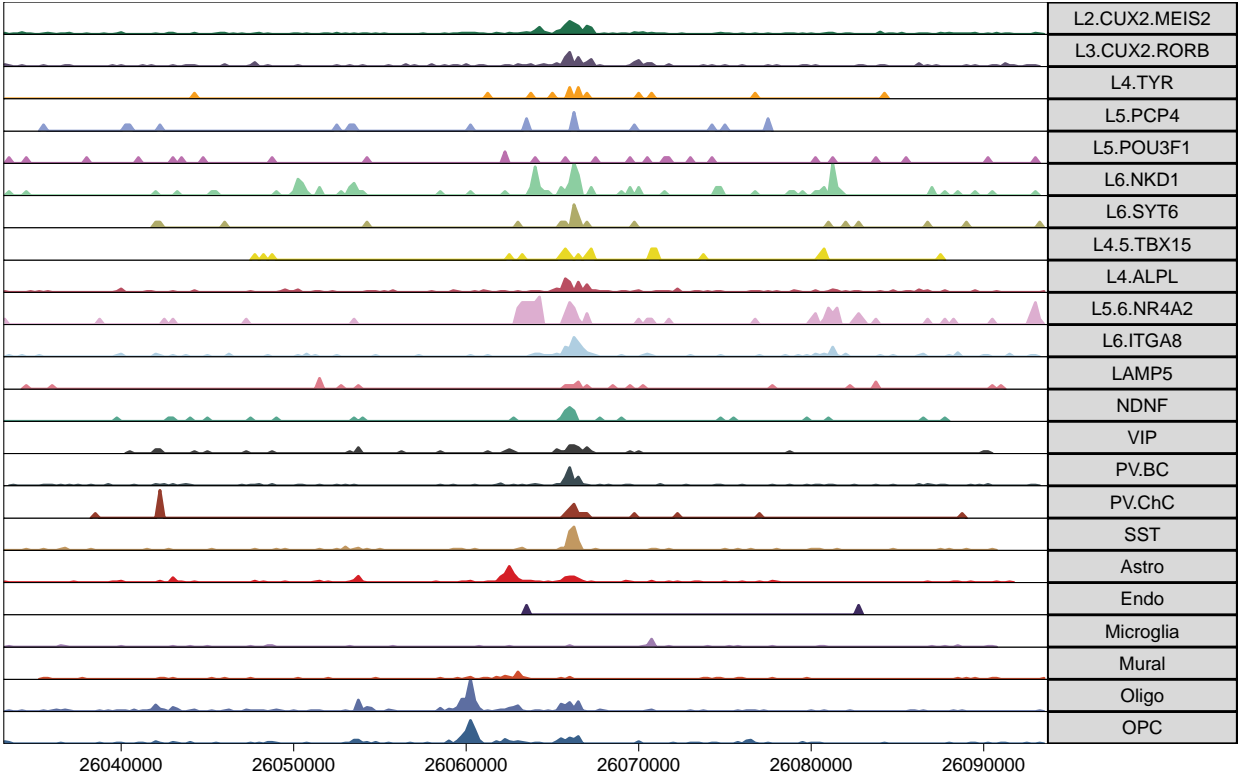


LoopTrack

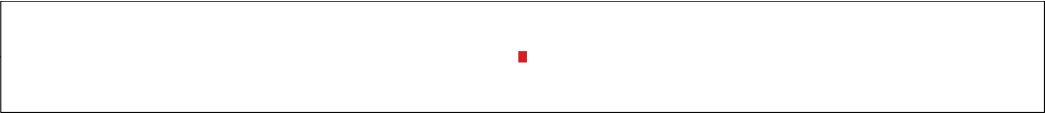


Genes

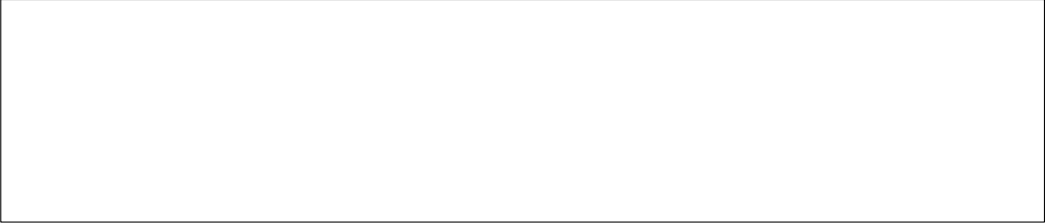
Coverage
(Norm. ATAC Signal Range 0–1.33) by ReadsInTSS)



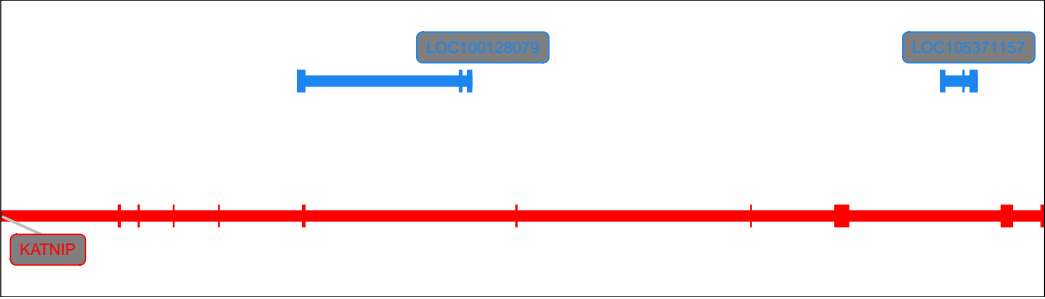
L5.6.NR4A2



Peaks

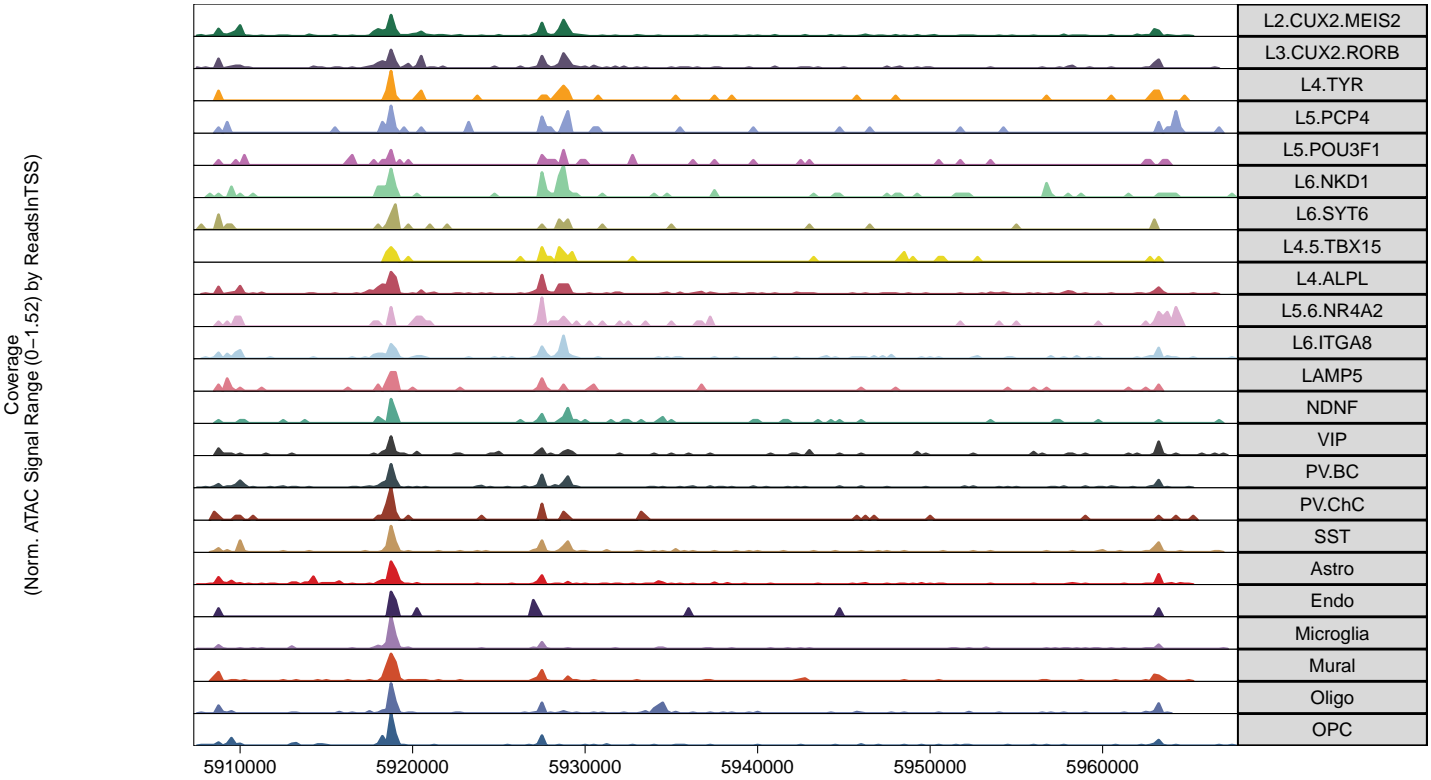


LoopTrack

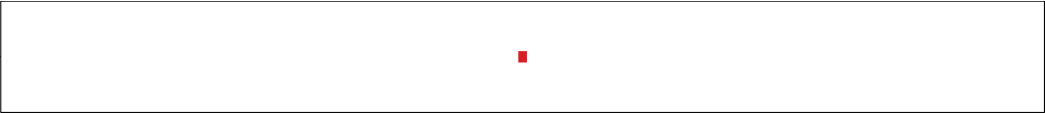


Genes

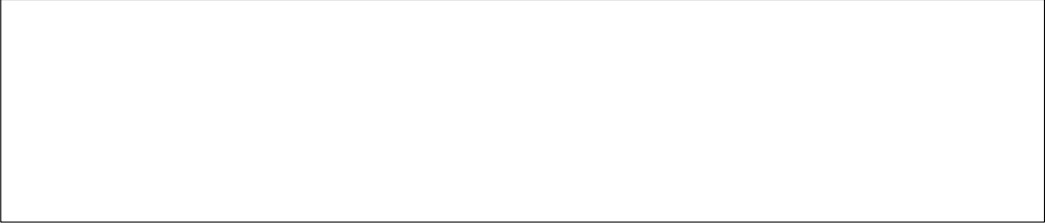
chr19:5907315–5967816 Marker Peak Tracks for: L5.6.NR4A2



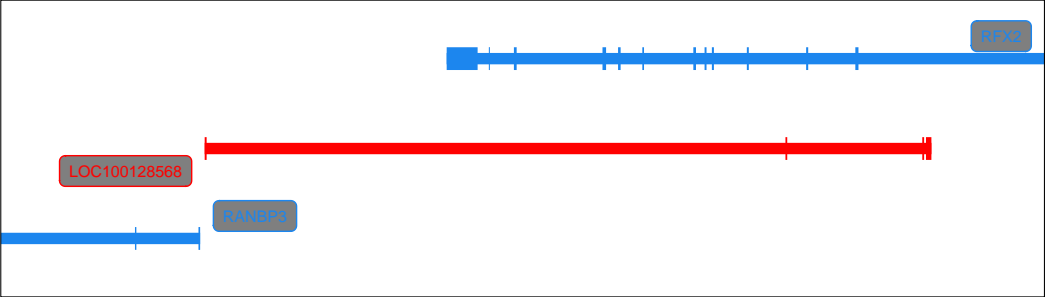
L5.6.NR4A2



Peaks

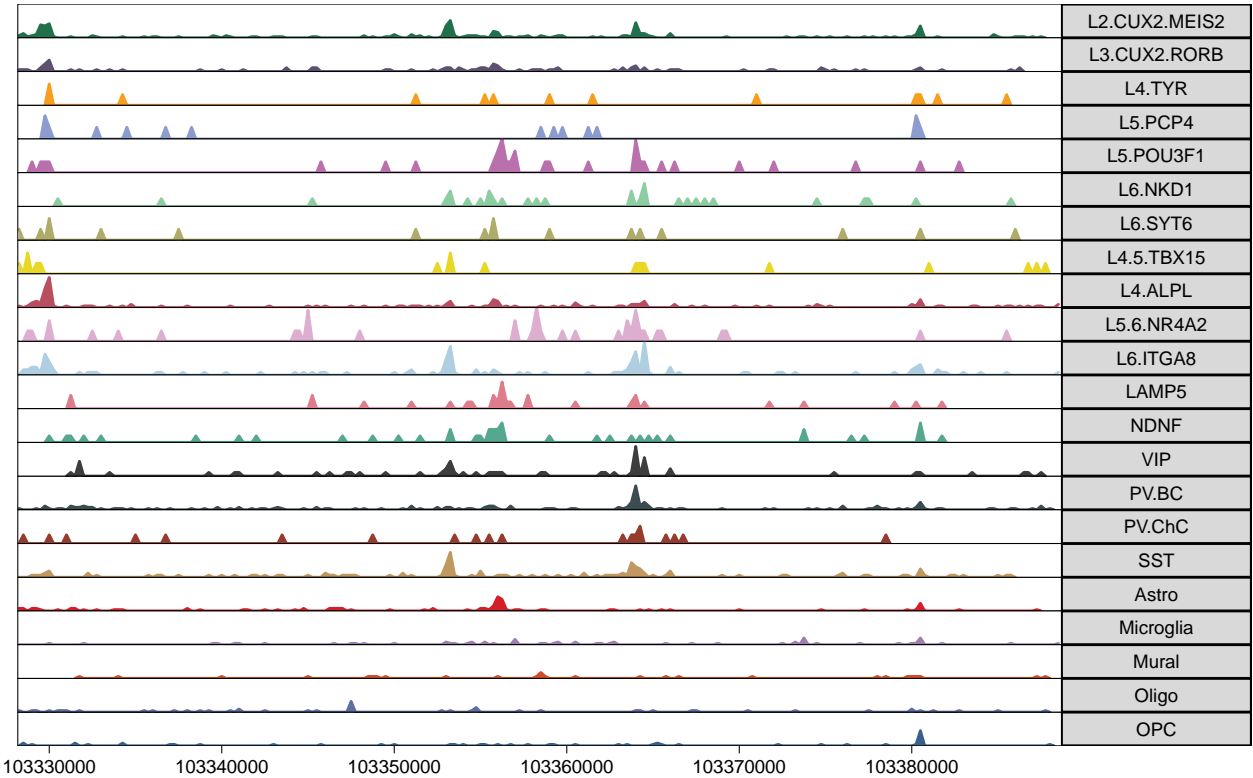


LoopTrack

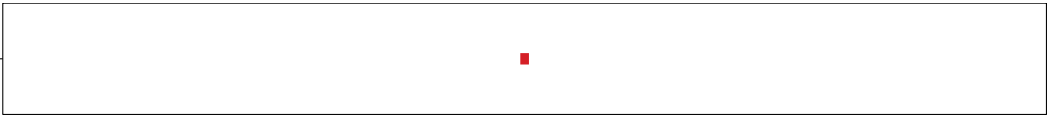


Genes

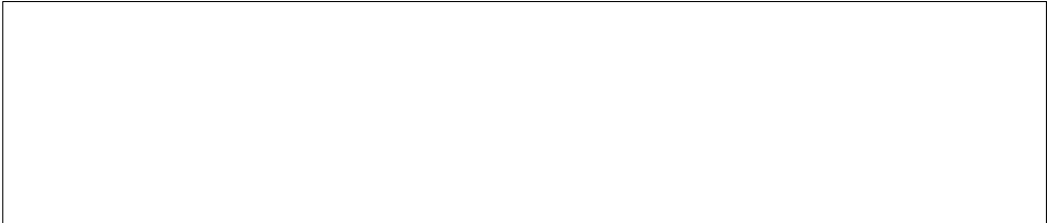
Coverage
(Norm. ATAC Signal Range 0–0.74) by ReadsInTSS)



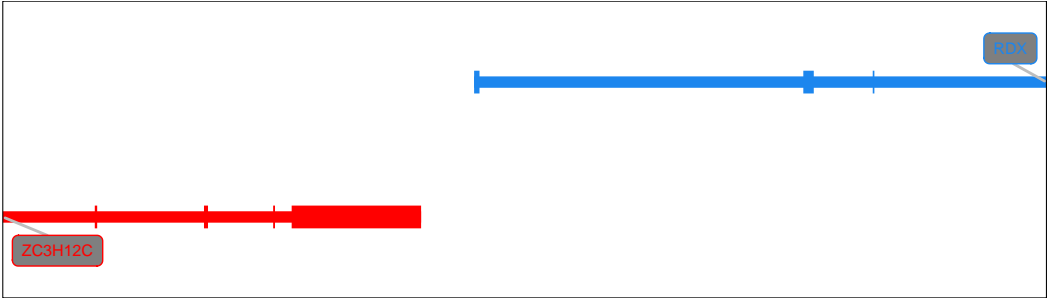
L5.6.NR4A2



Peaks



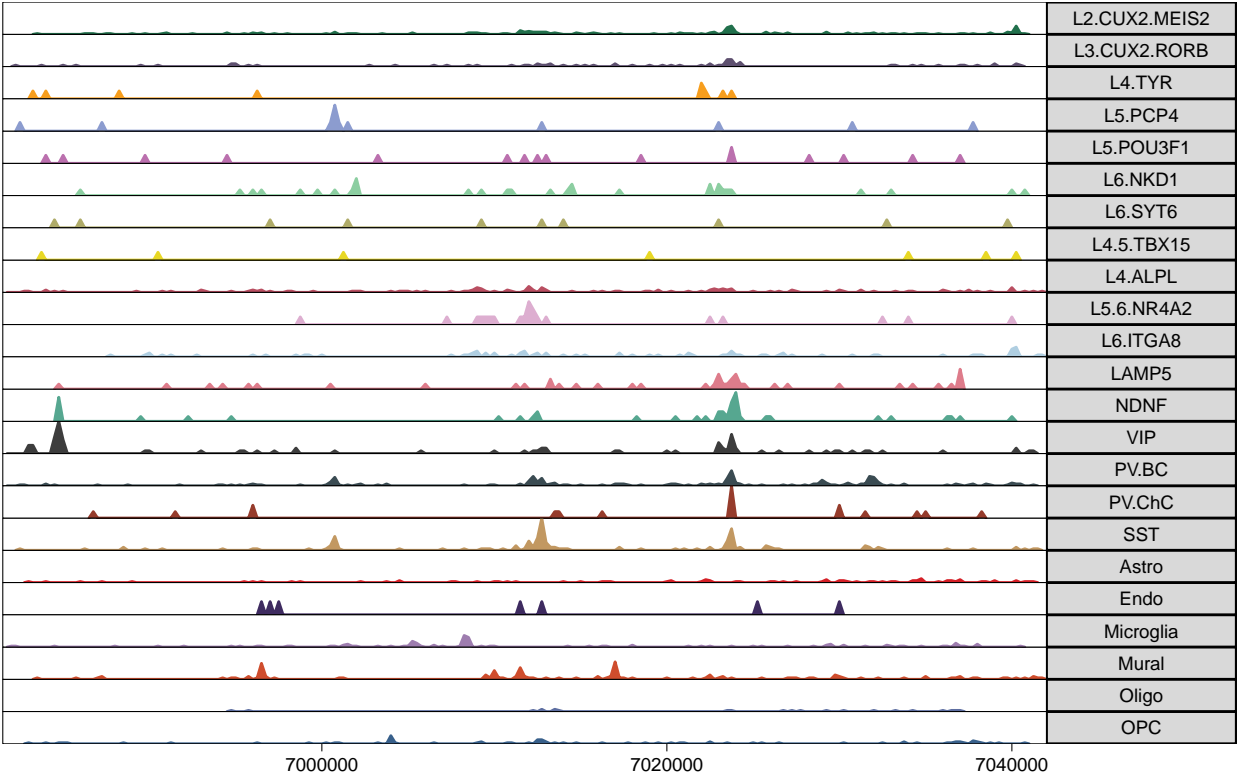
LoopTrack



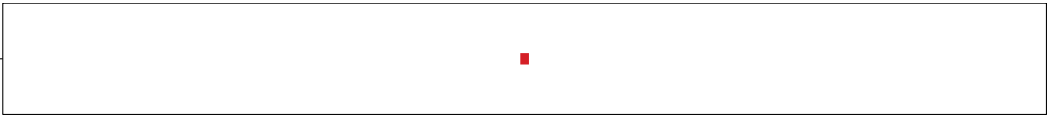
Genes

chr4:6981503–7042004 Marker Peak Tracks for: L5.6.NR4A2

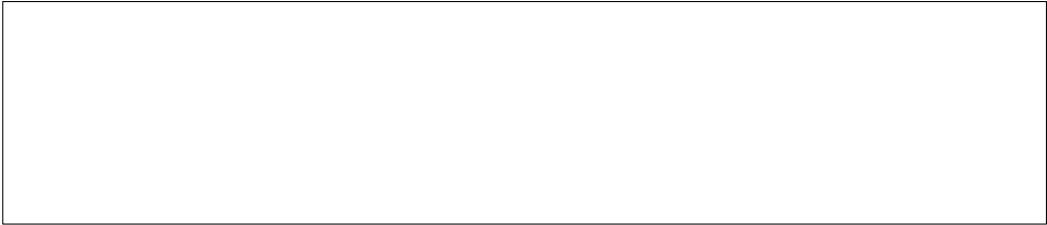
Coverage
(Norm. ATAC Signal Range (0–0.96) by ReadsInTSS)



L5.6.NR4A2



Peaks



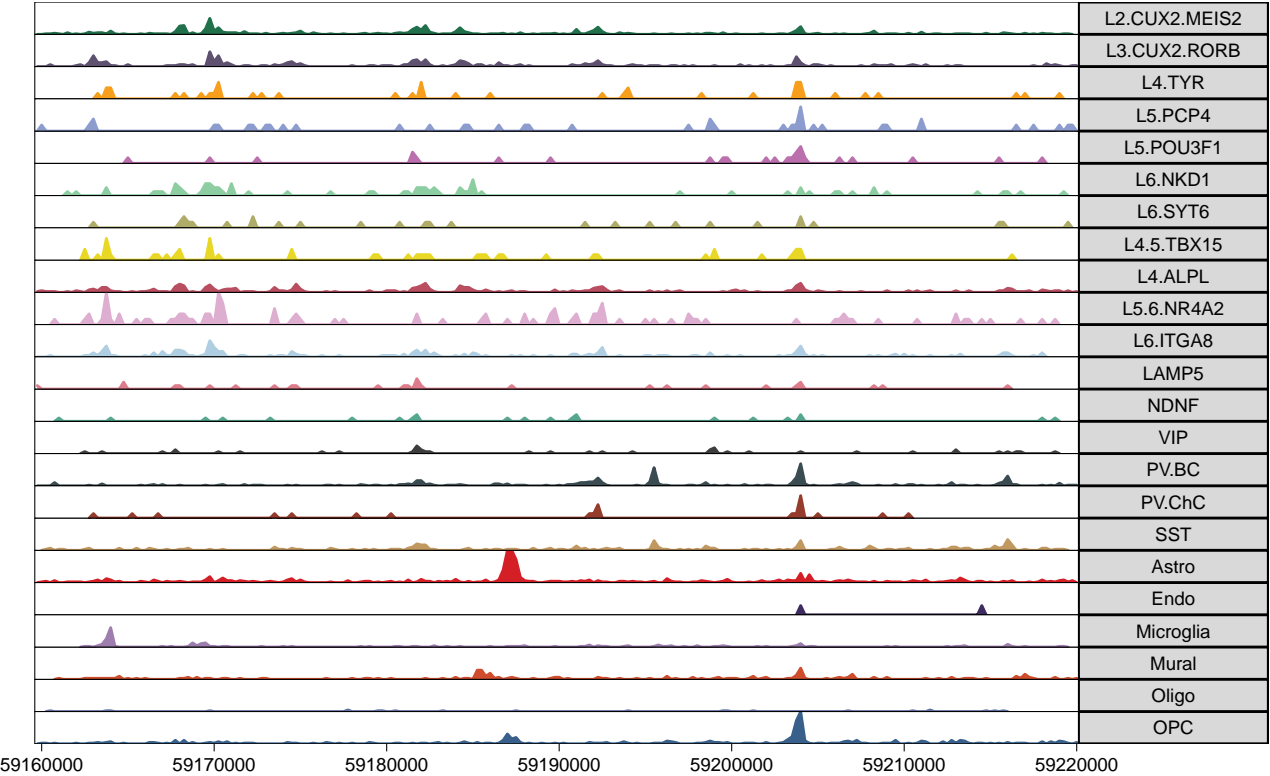
LoopTrack



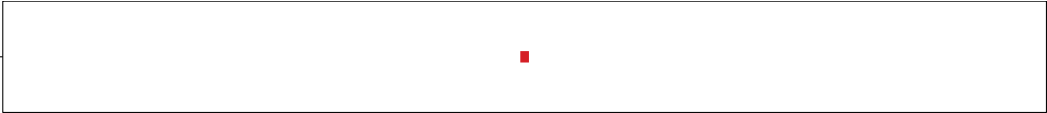
Genes

chr18:59159601–59220102 Marker Peak Tracks for: L5.6.NR4A2

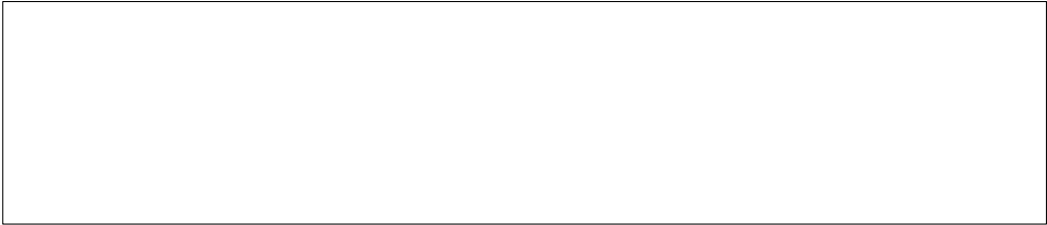
Coverage
(Norm. ATAC Signal Range 0–1.37) by ReadsInTSS)



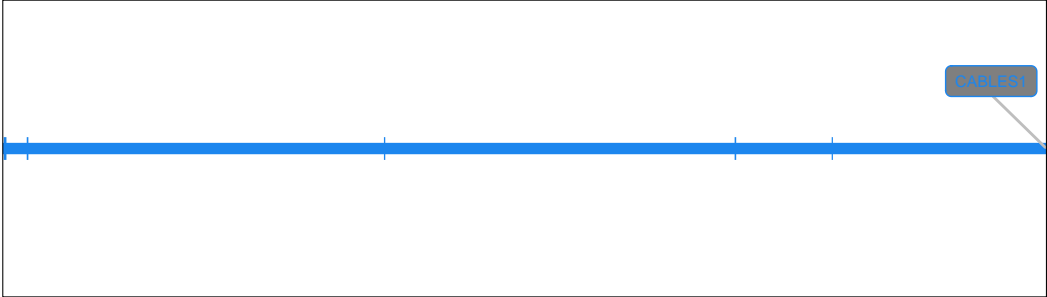
L5.6.NR4A2



Peaks

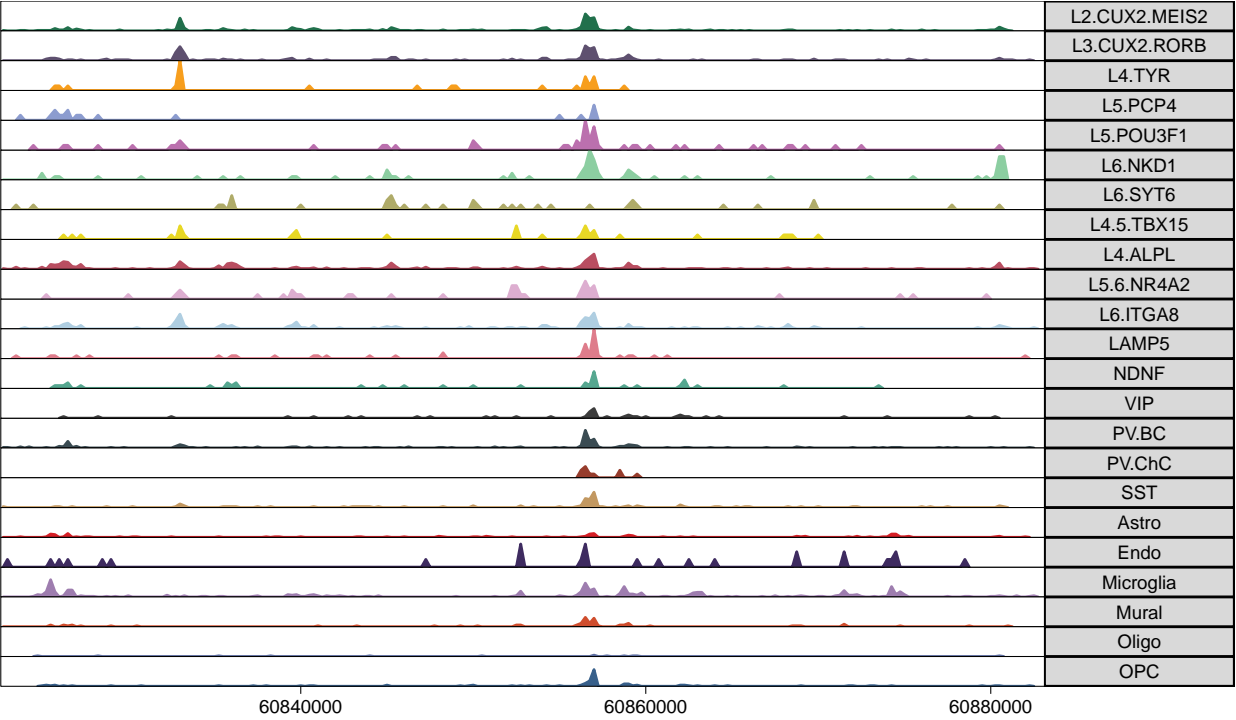


LoopTrack

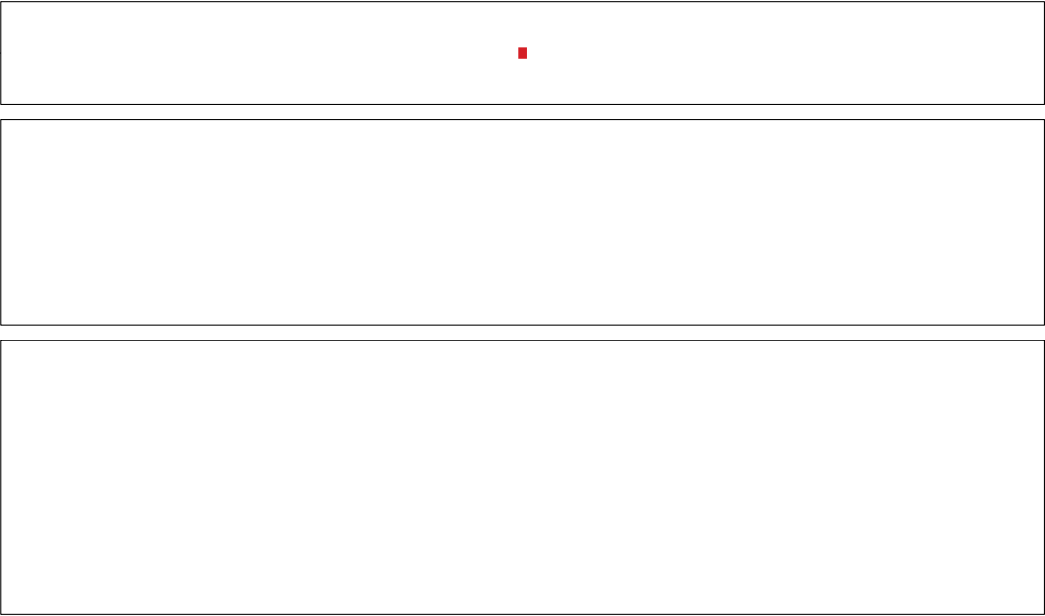


Genes

Coverage
(Norm. ATAC Signal Range (0-1.51) by Reads in TSS)



L5.6.NR4A2



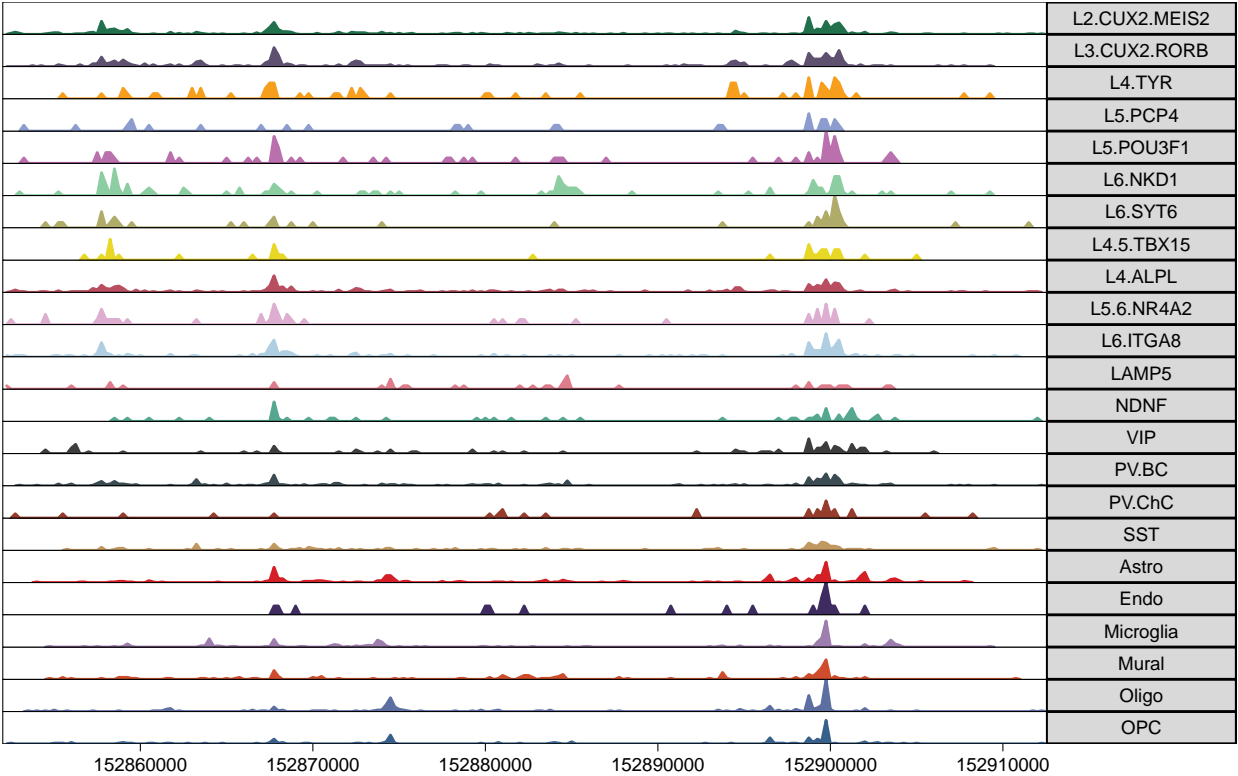
Peaks

LoopTrack

GeneTrack

chr7:152852025-152912526 Marker Peak Tracks for: L5.6.NR4A2

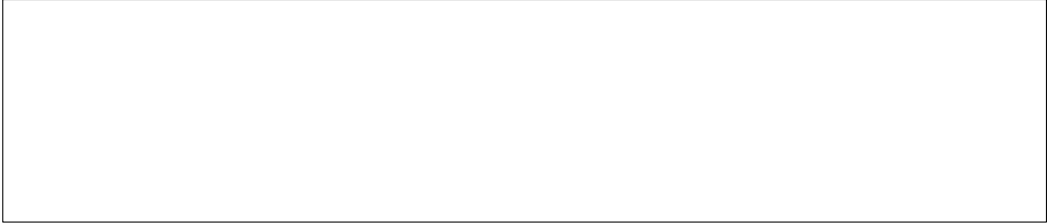
Coverage
(Norm. ATAC Signal Range 0-1.42) by ReadsInTSS)



L5.6.NR4A2



Peaks

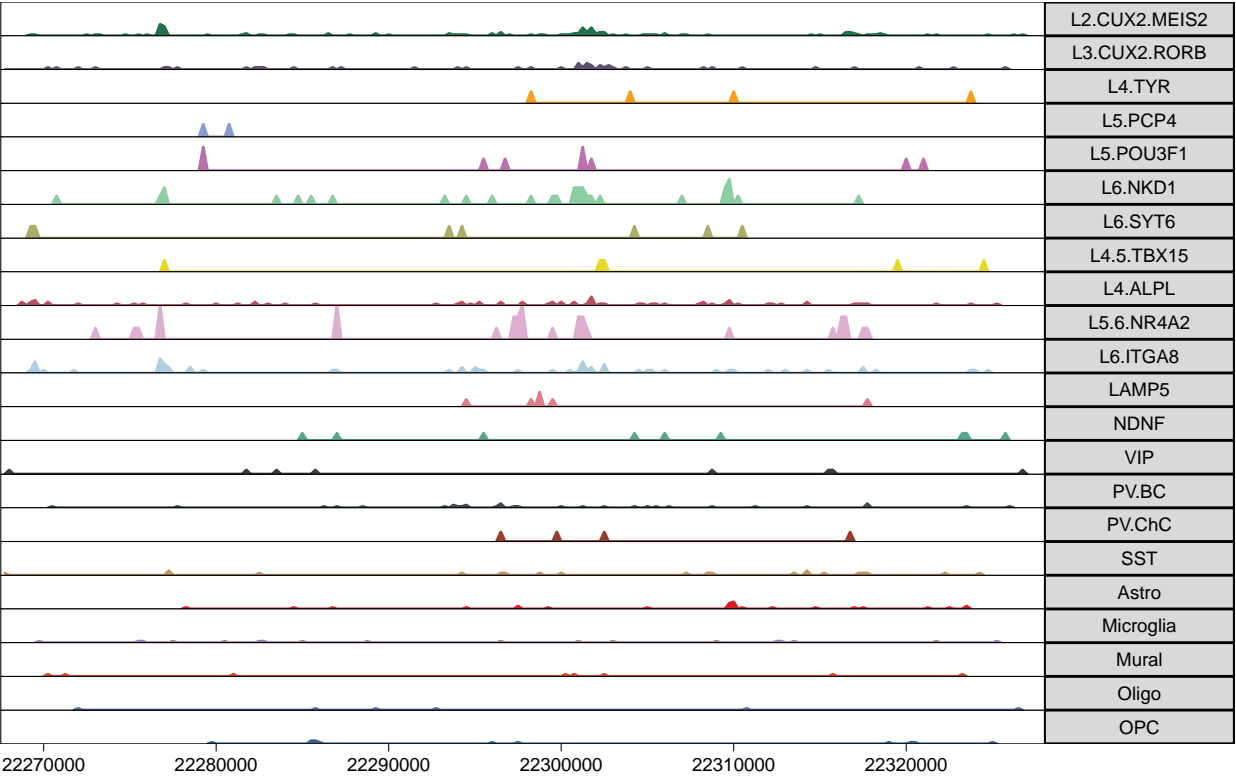


LoopTrack

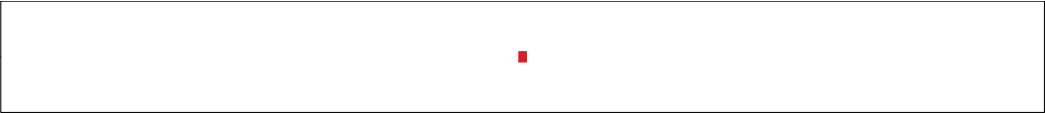


Genes

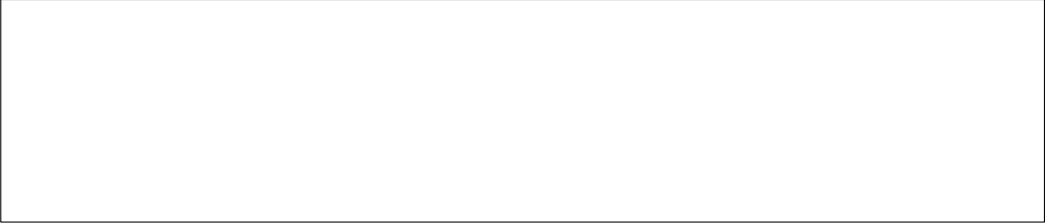
Coverage
(Norm. ATAC Signal Range 0–0.68) by ReadsInTSS)



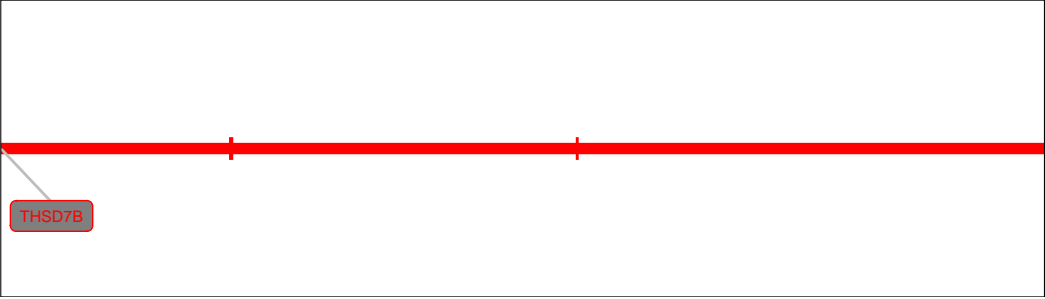
L5.6.NR4A2



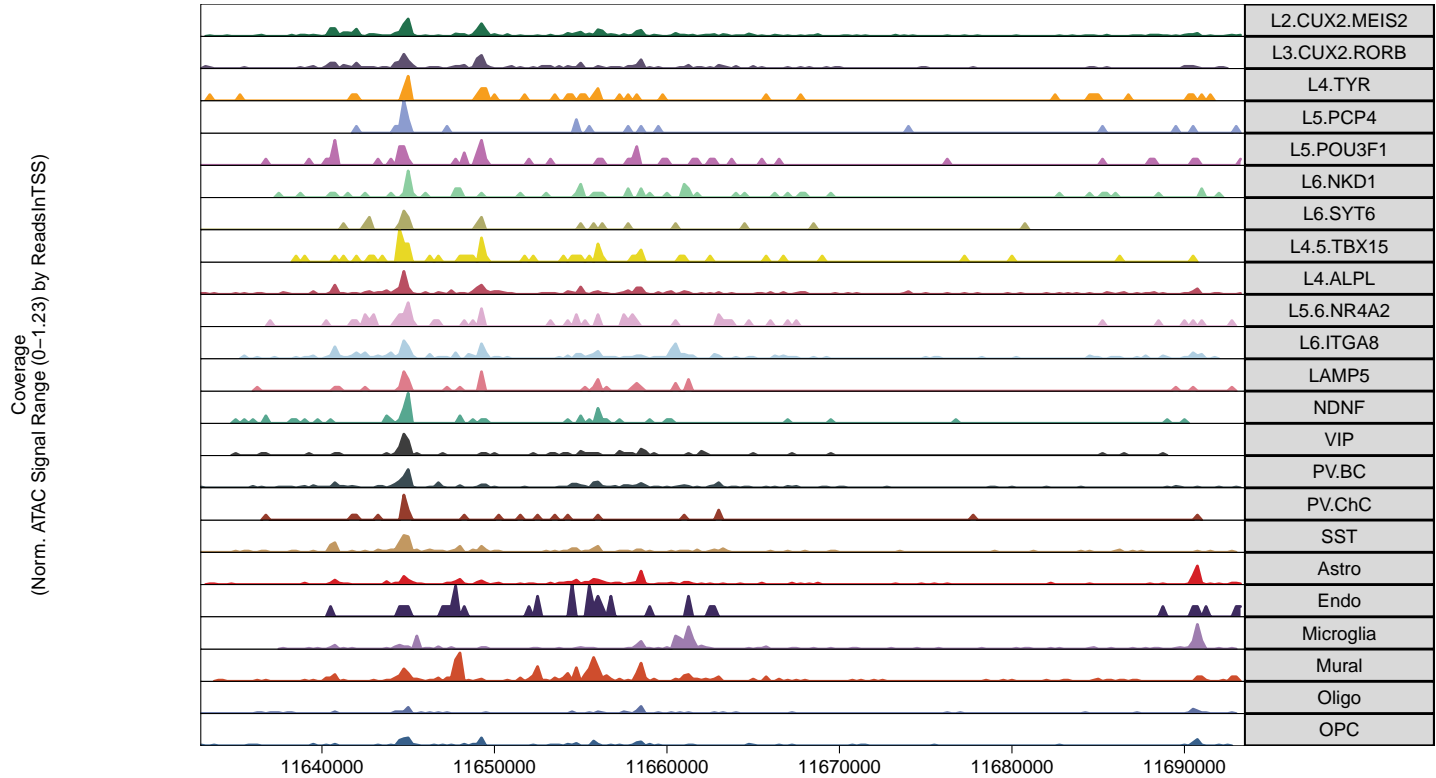
Peaks



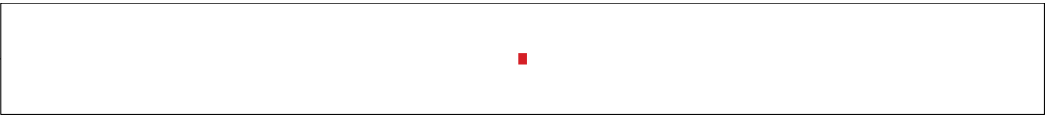
LoopTrack



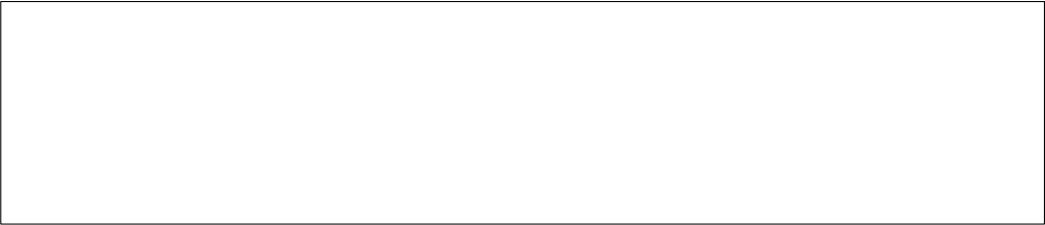
Genes



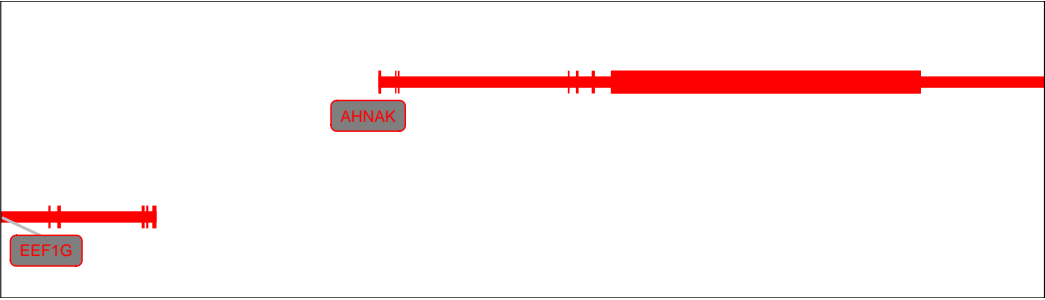
L5.6.NR4A2



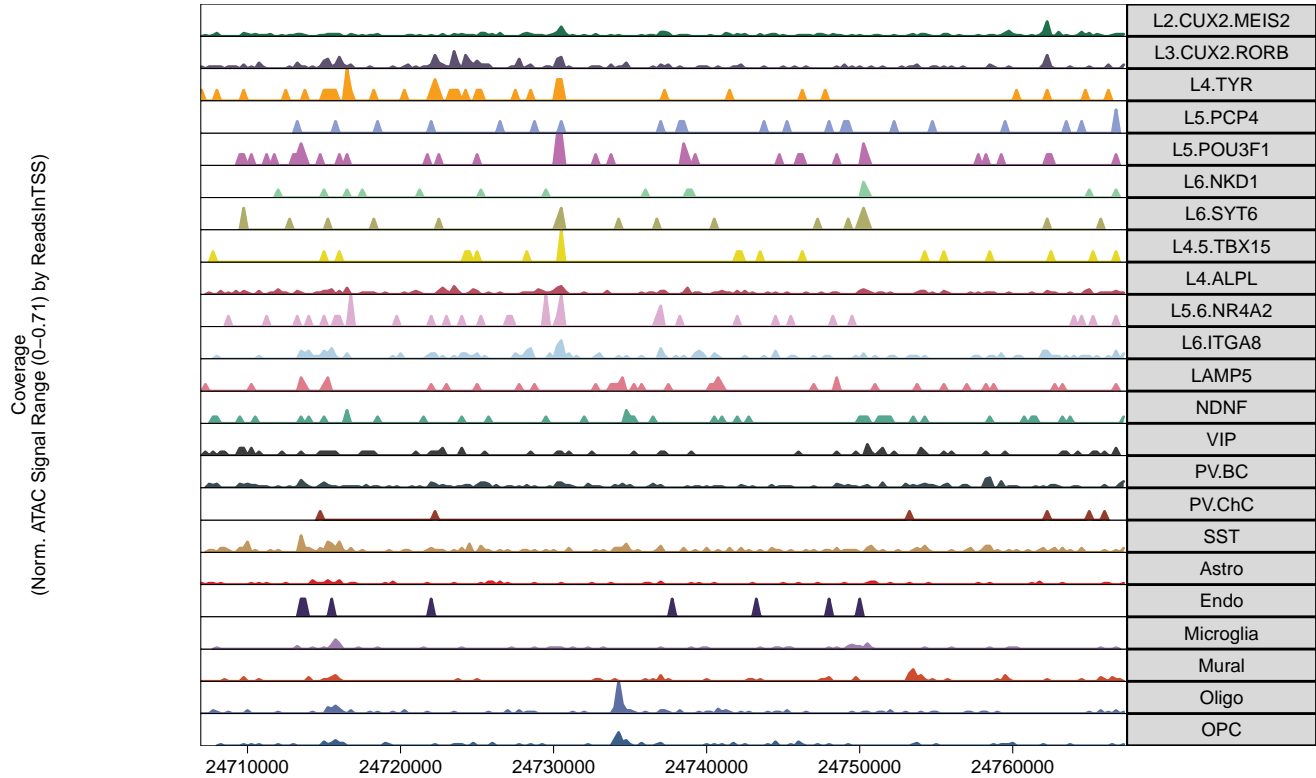
Peaks



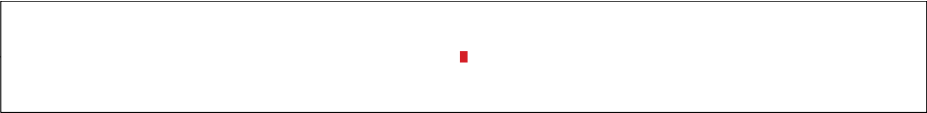
LoopTrack



Genes



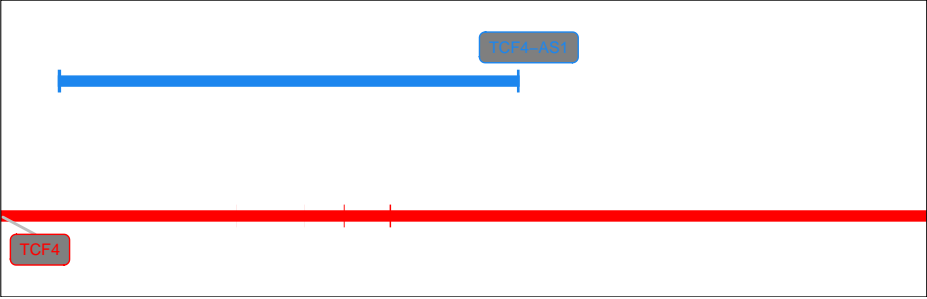
L5.6.NR4A2



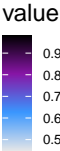
Peaks



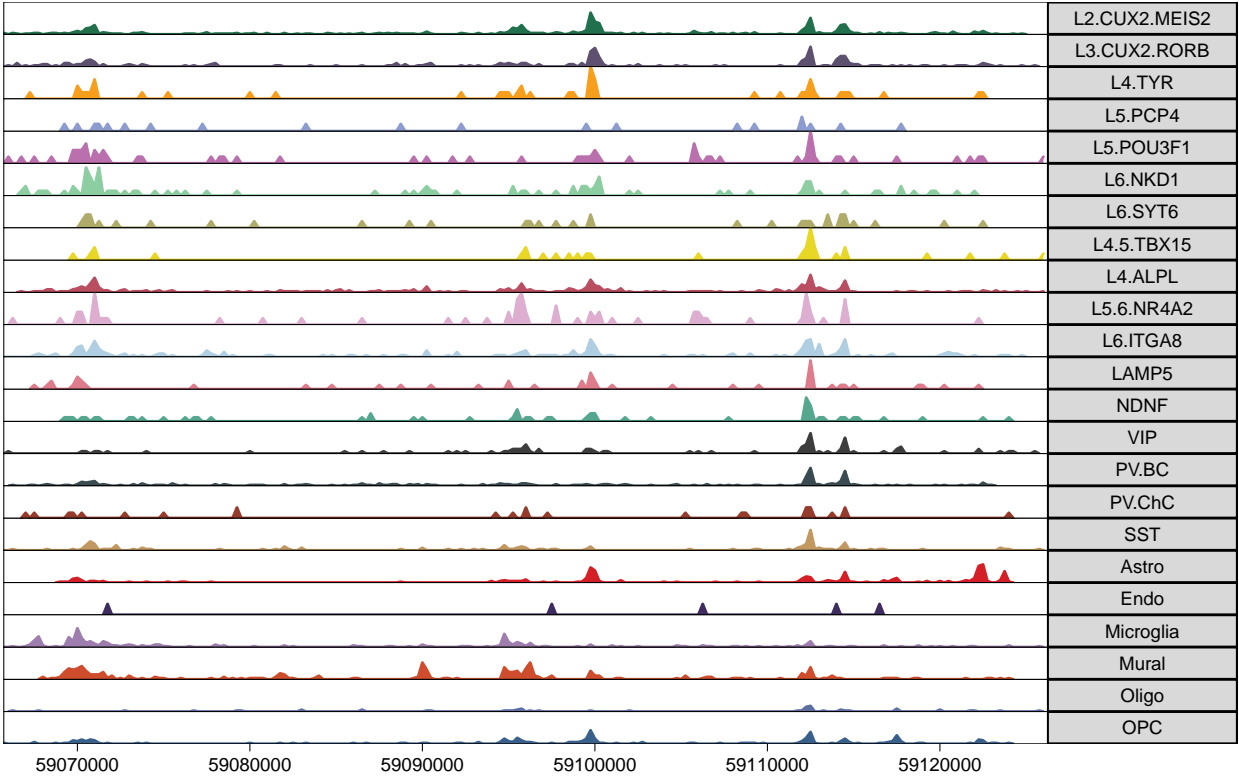
Peak2GeneLinks



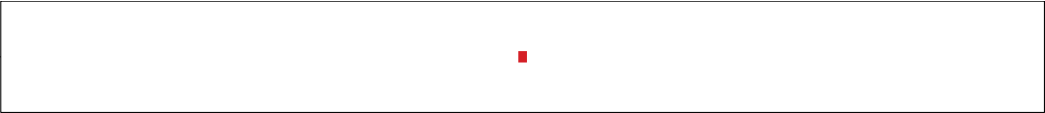
Genes



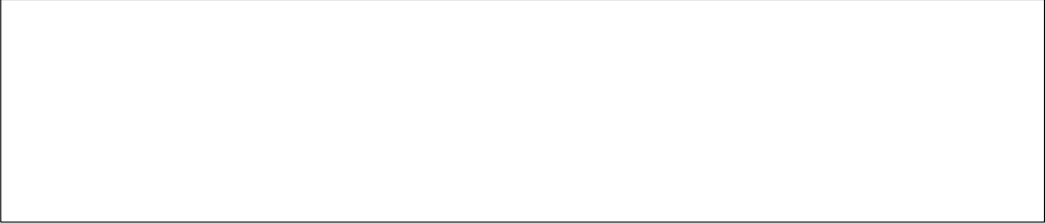
Coverage
(Norm. ATAC Signal Range 0–1.16) by ReadsInTSS)



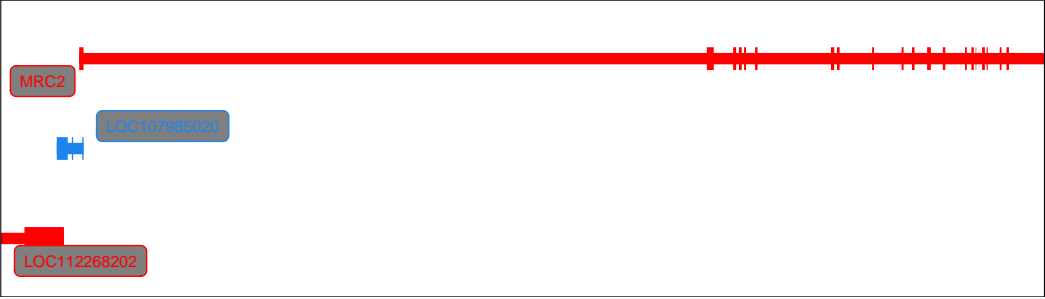
L5.6.NR4A2



Peaks



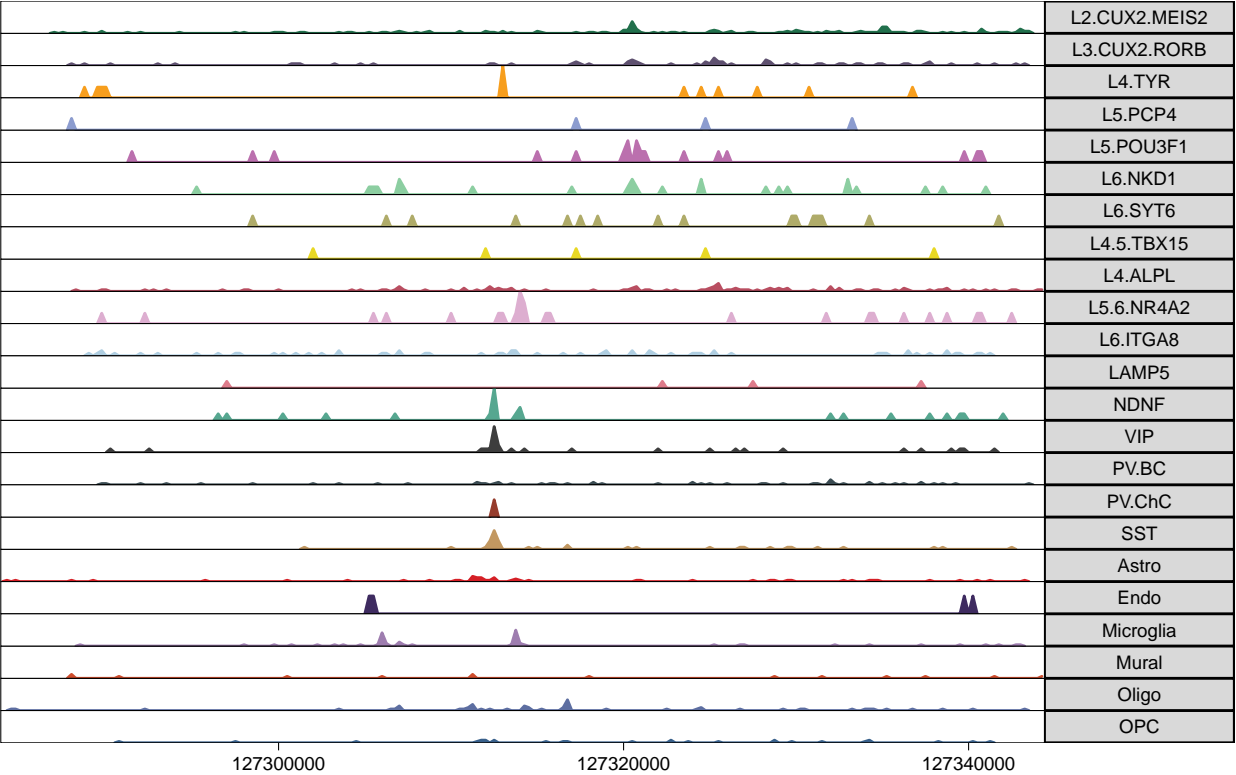
LoopTrack



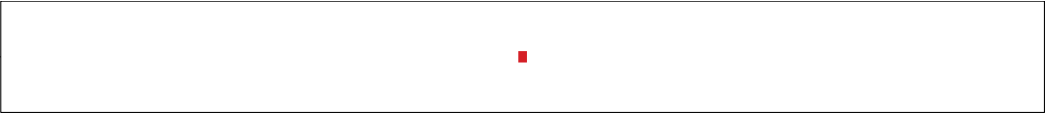
Genes

chr1:127283897-127344398 Marker Peak Tracks for: L5.6.NR4A2

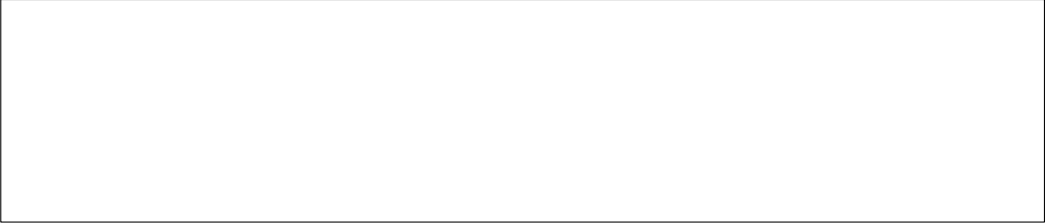
Coverage
(Norm. ATAC Signal Range (0-0.71) by ReadsInTSS)



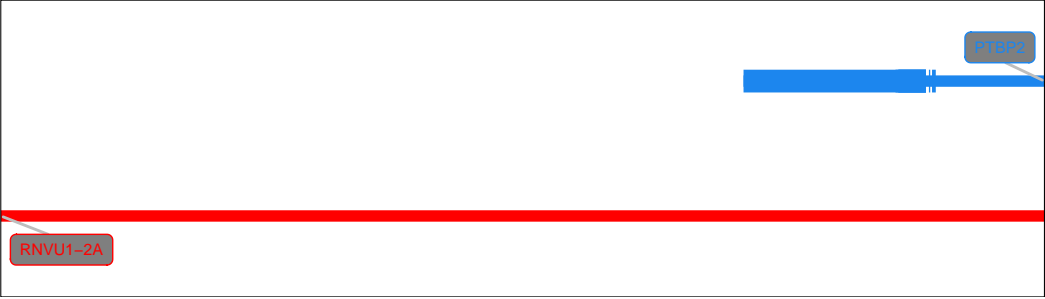
L5.6.NR4A2



Peaks

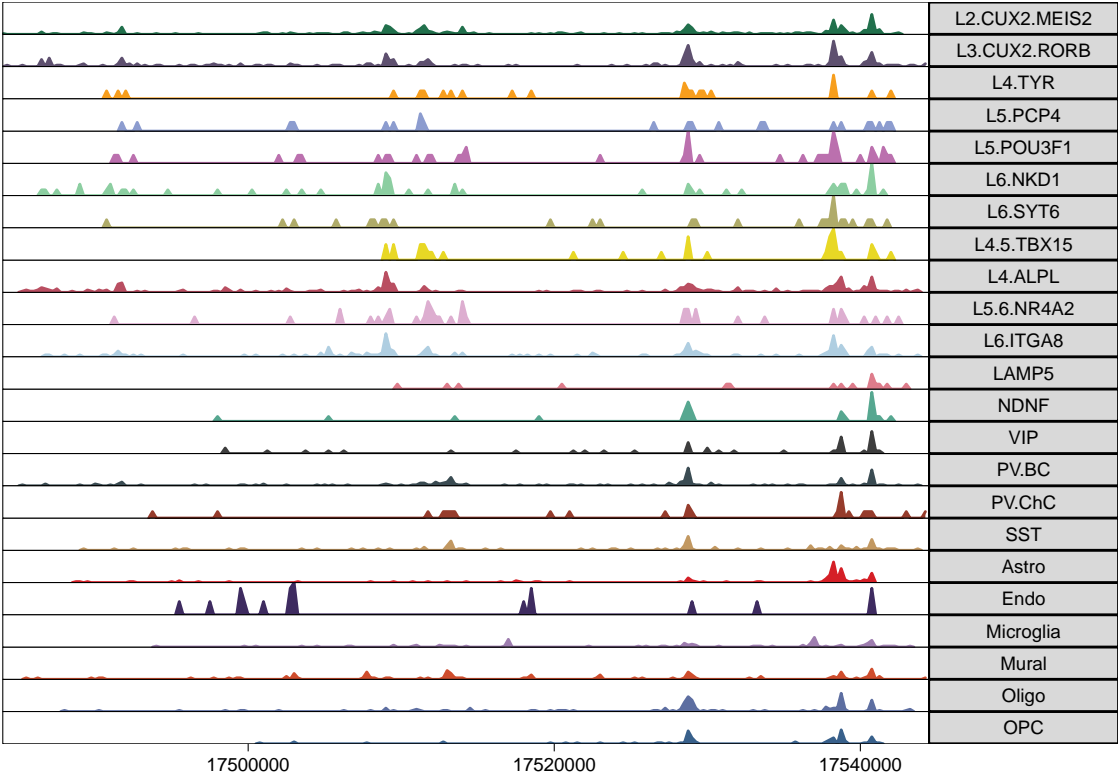


LoopTrack

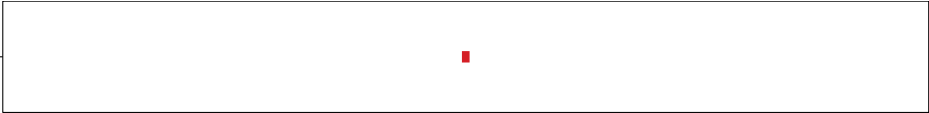


Genes

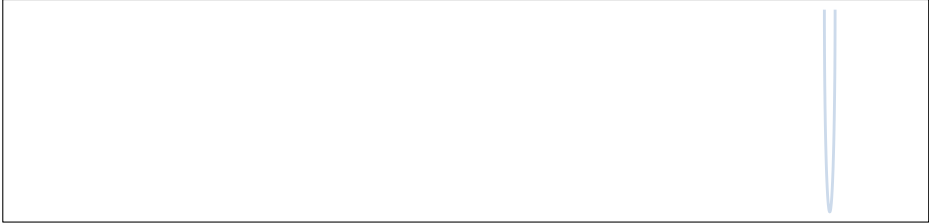
Coverage
(Norm. ATAC Signal Range (0–0.95) by ReadsInTSS)



L5.6.NR4A2



Peaks

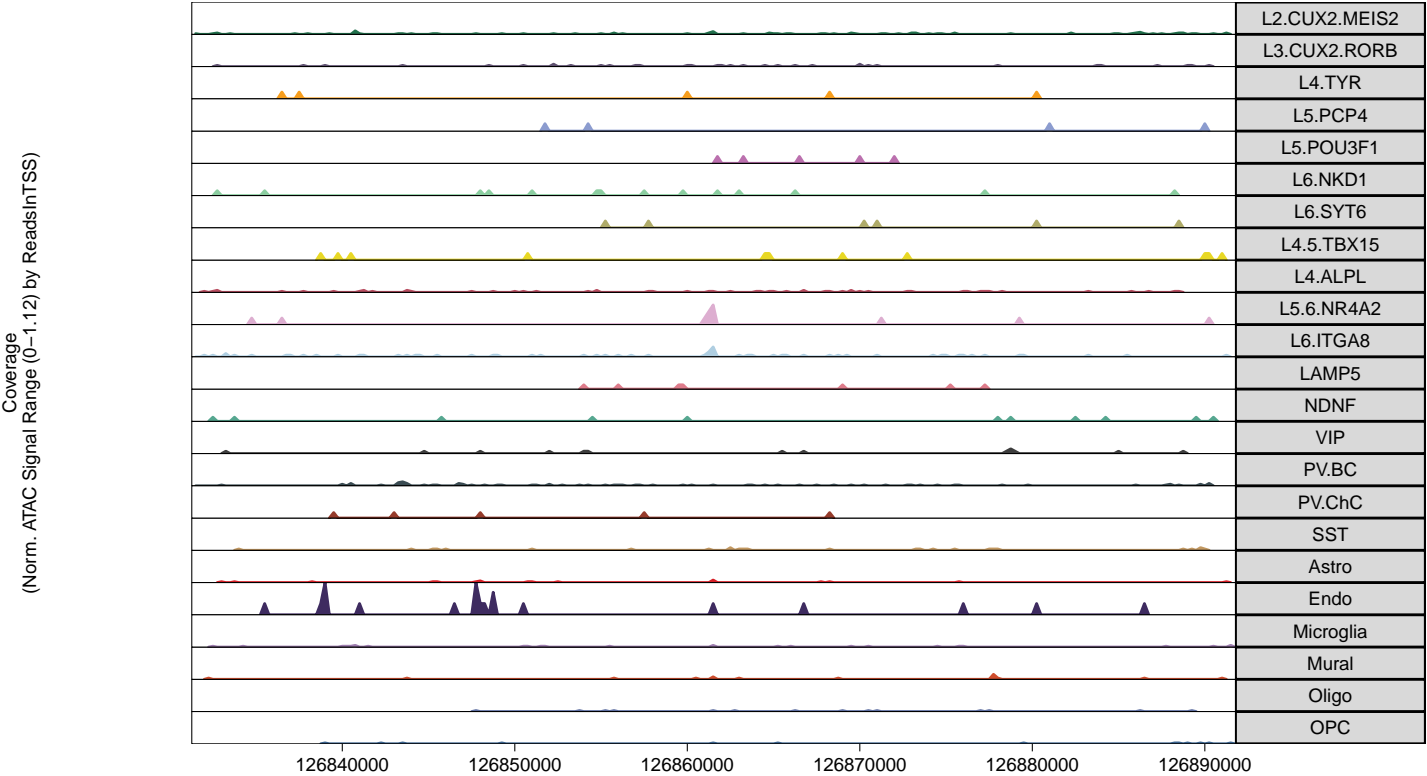


Peak2GeneLinks

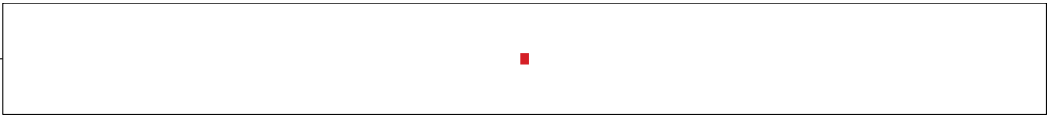


Genes

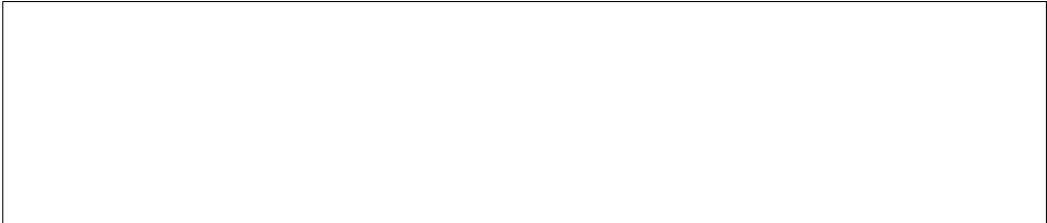
chr8:126831275–126891776 Marker Peak Tracks for: L5.6.NR4A2



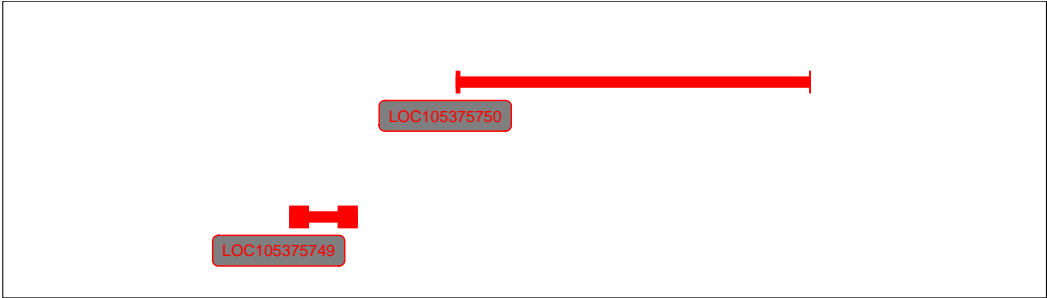
L5.6.NR4A2



Peaks

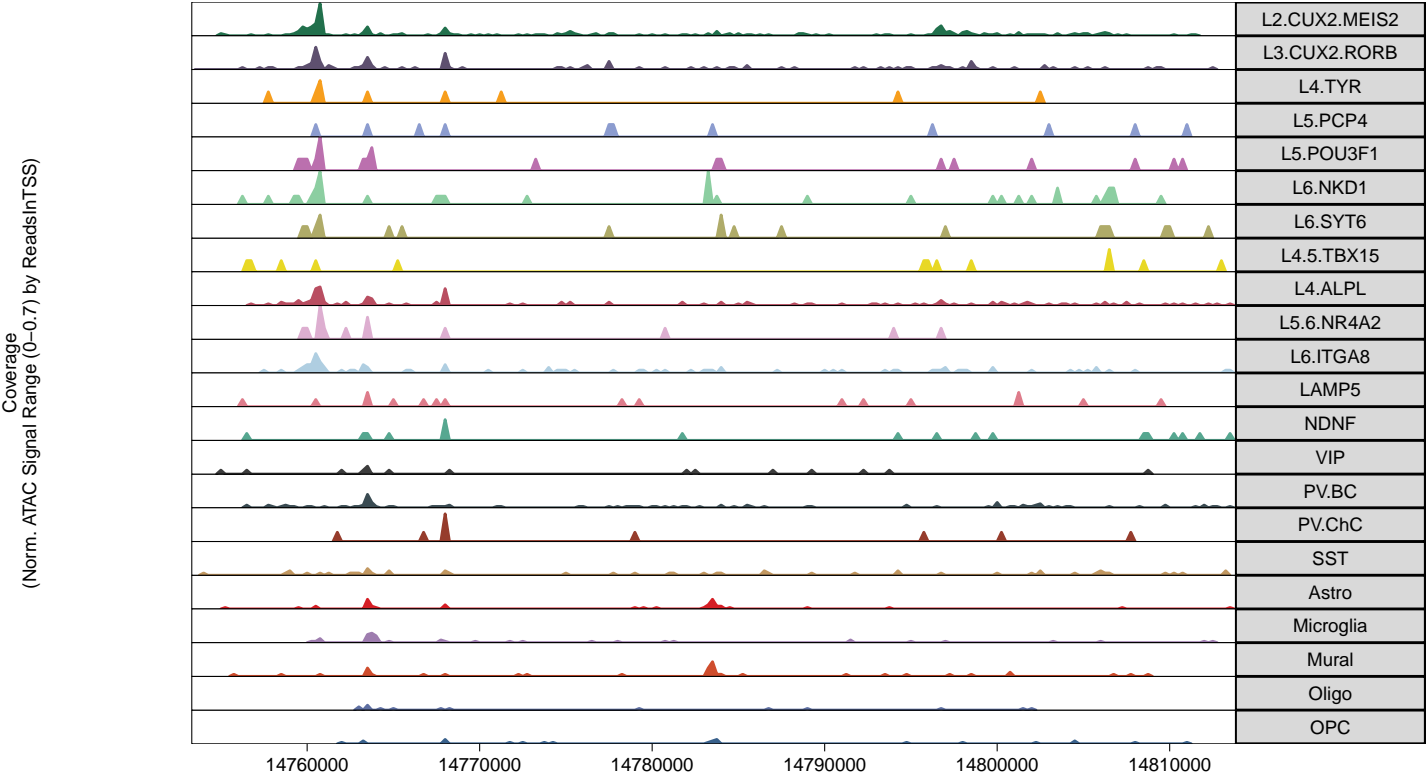


LoopTrack

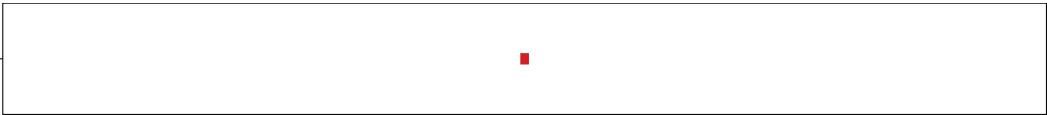


Genes

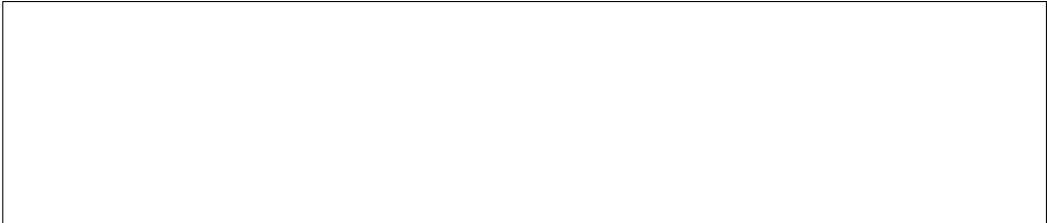
chr2:14753312–14813813 Marker Peak Tracks for: L5.6.NR4A2



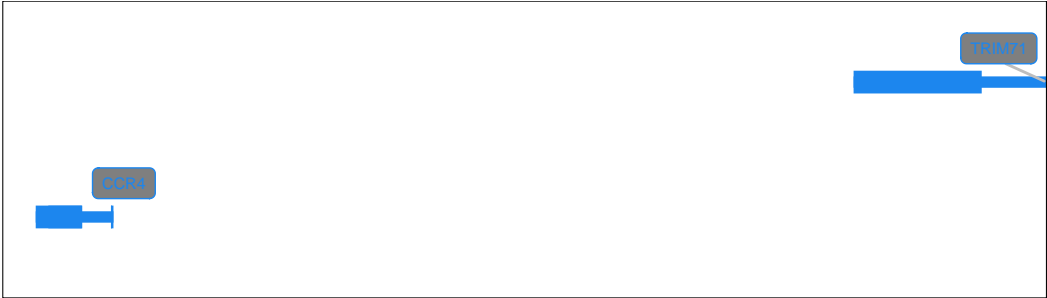
L5.6.NR4A2



Peaks

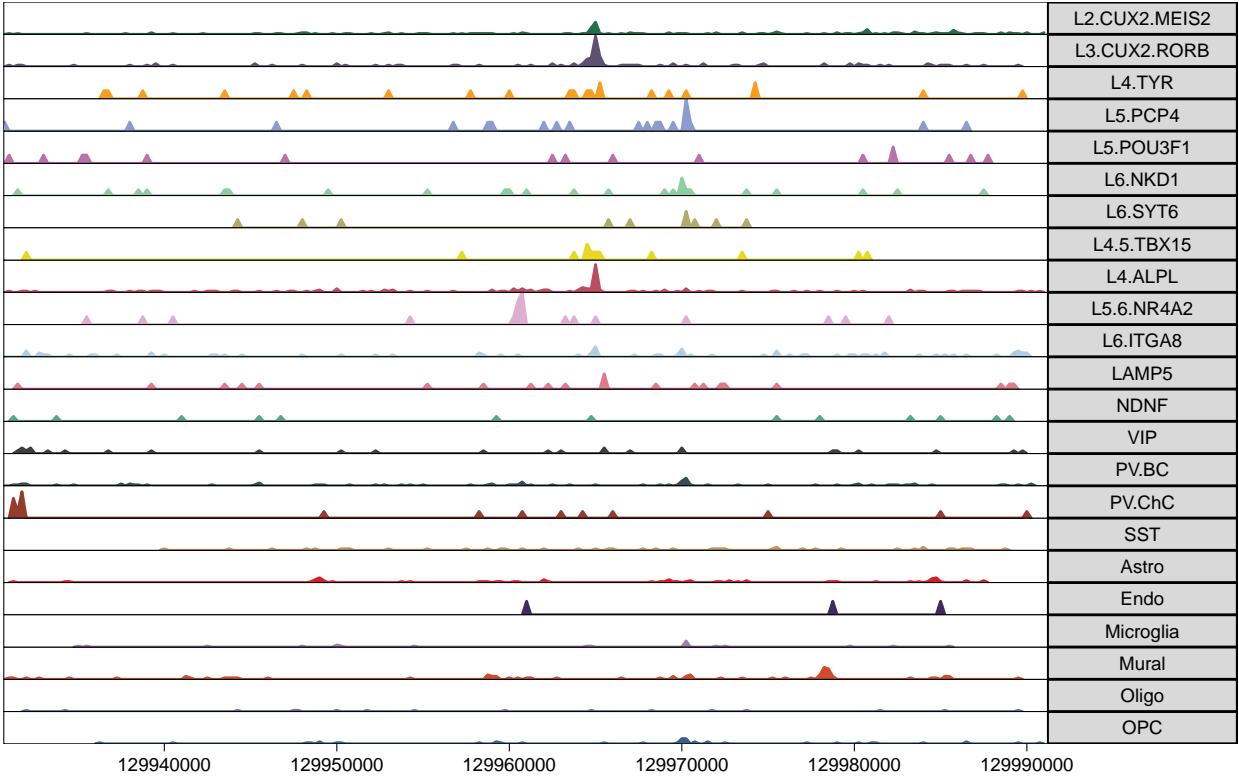


LoopTrack

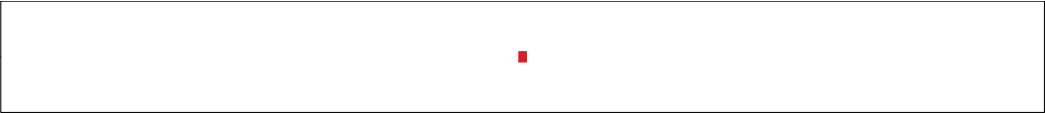


Genes

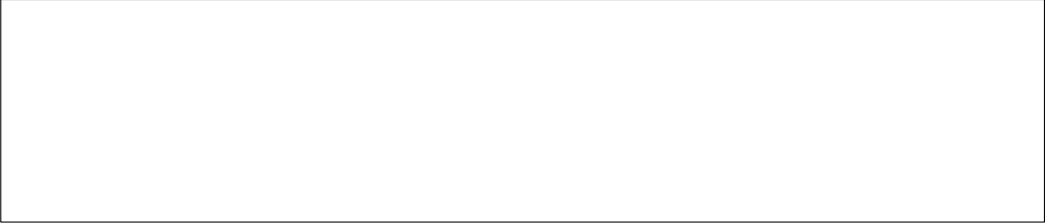
Coverage
(Norm. ATAC Signal Range 0–0.93) by ReadsInTSS)



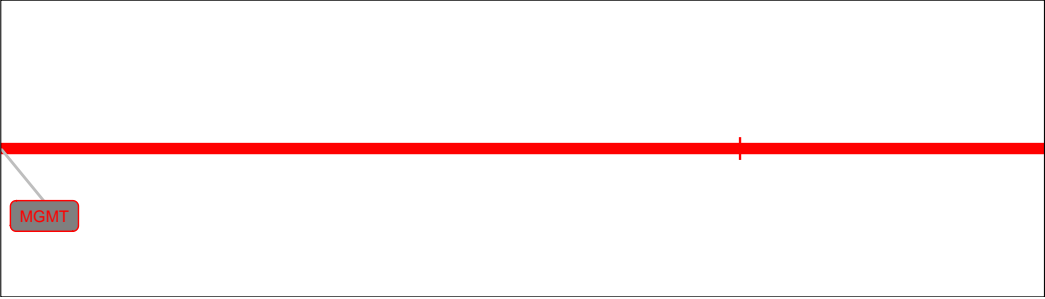
L5.6.NR4A2



Peaks

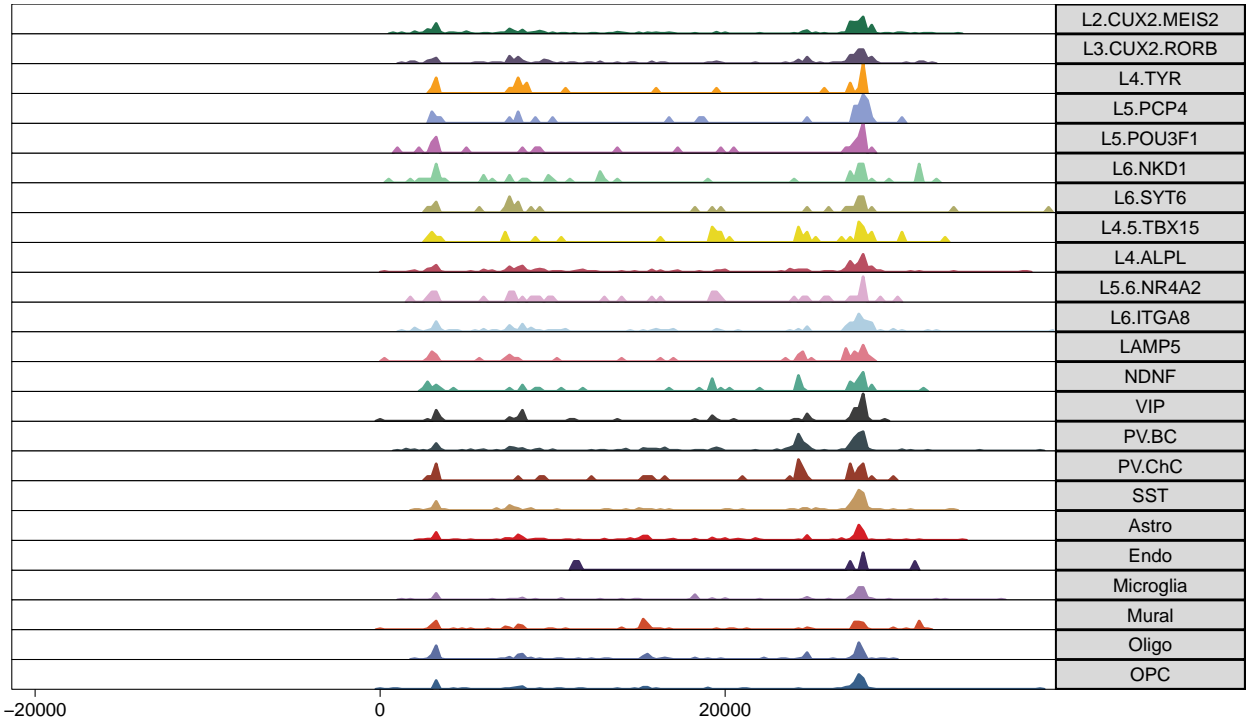


LoopTrack



Genes

Coverage
(Norm. ATAC Signal Range (0–1.34) by Reads in TSS)



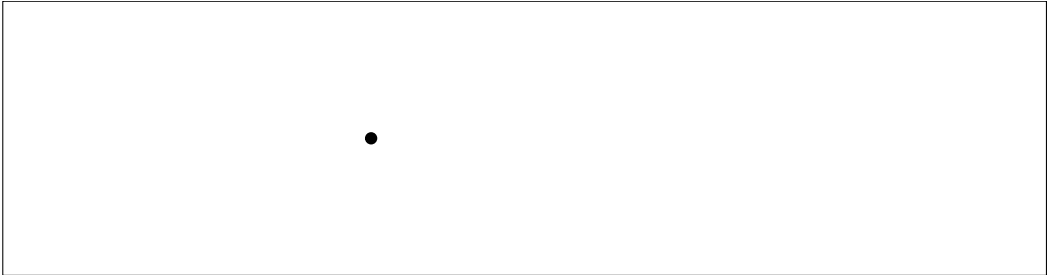
L5.6.NR4A2



Peaks

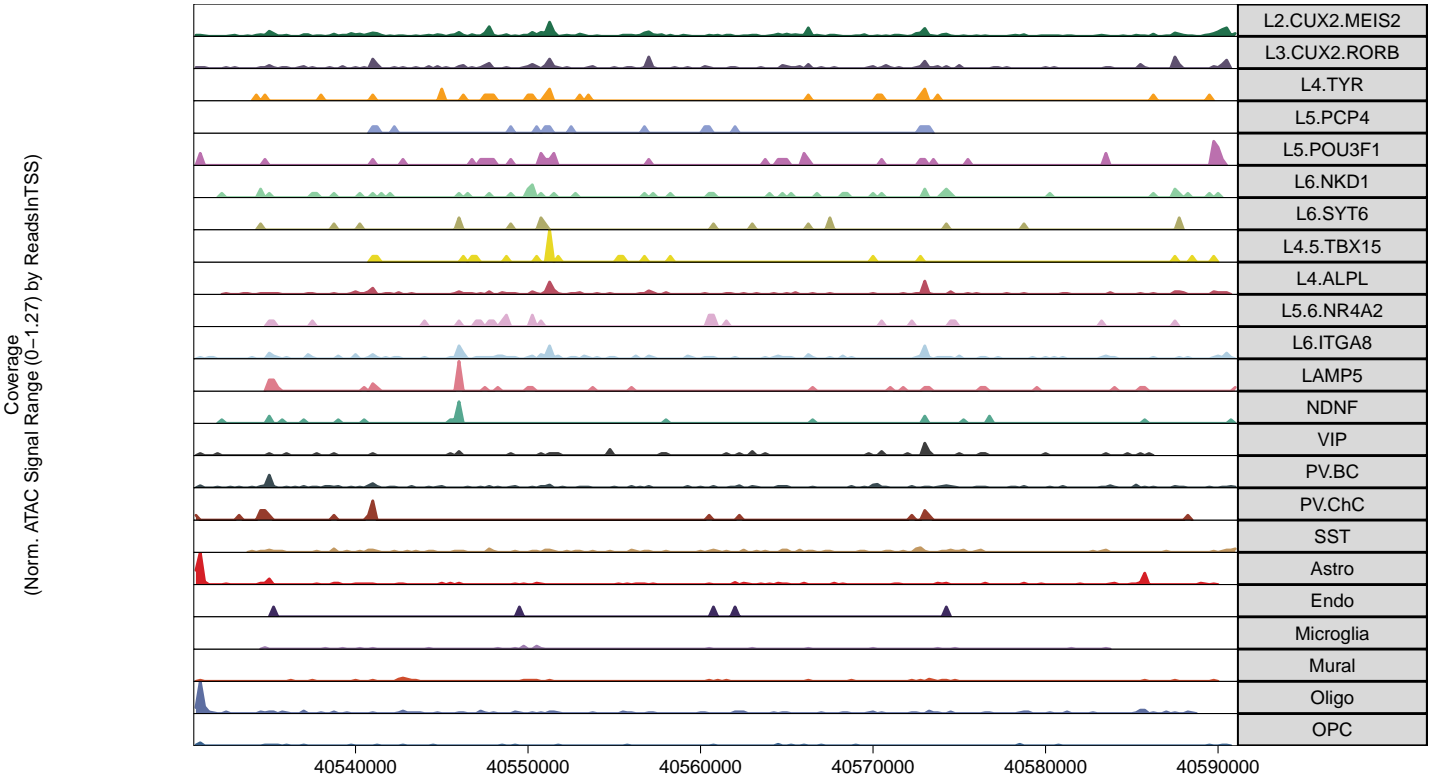


LoopTrack

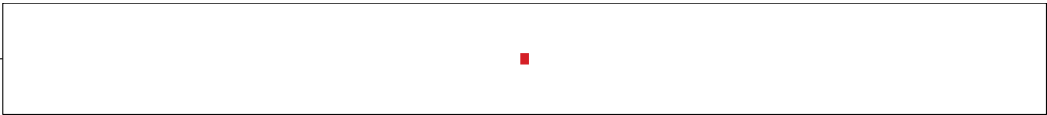


GeneTrack

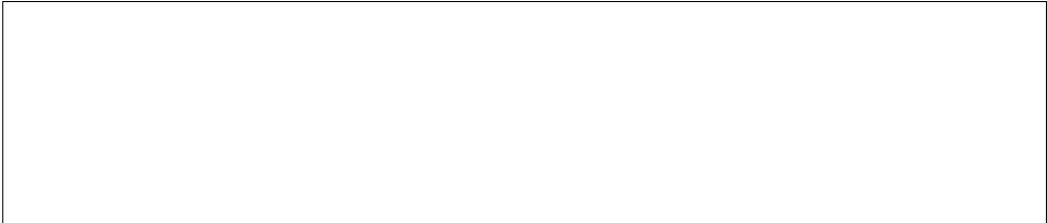
strand ● ·



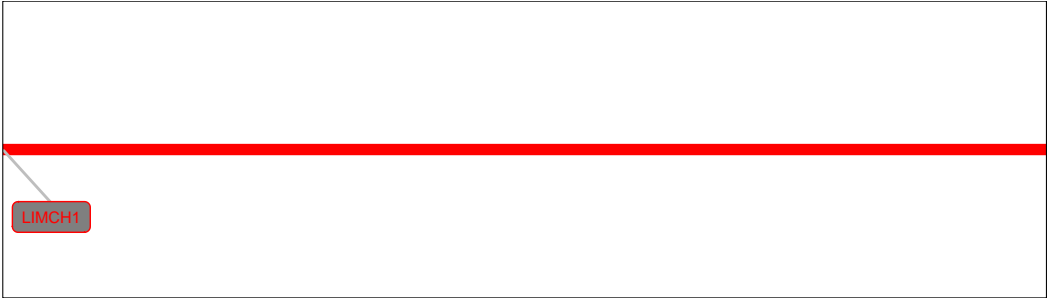
L5.6.NR4A2



Peaks



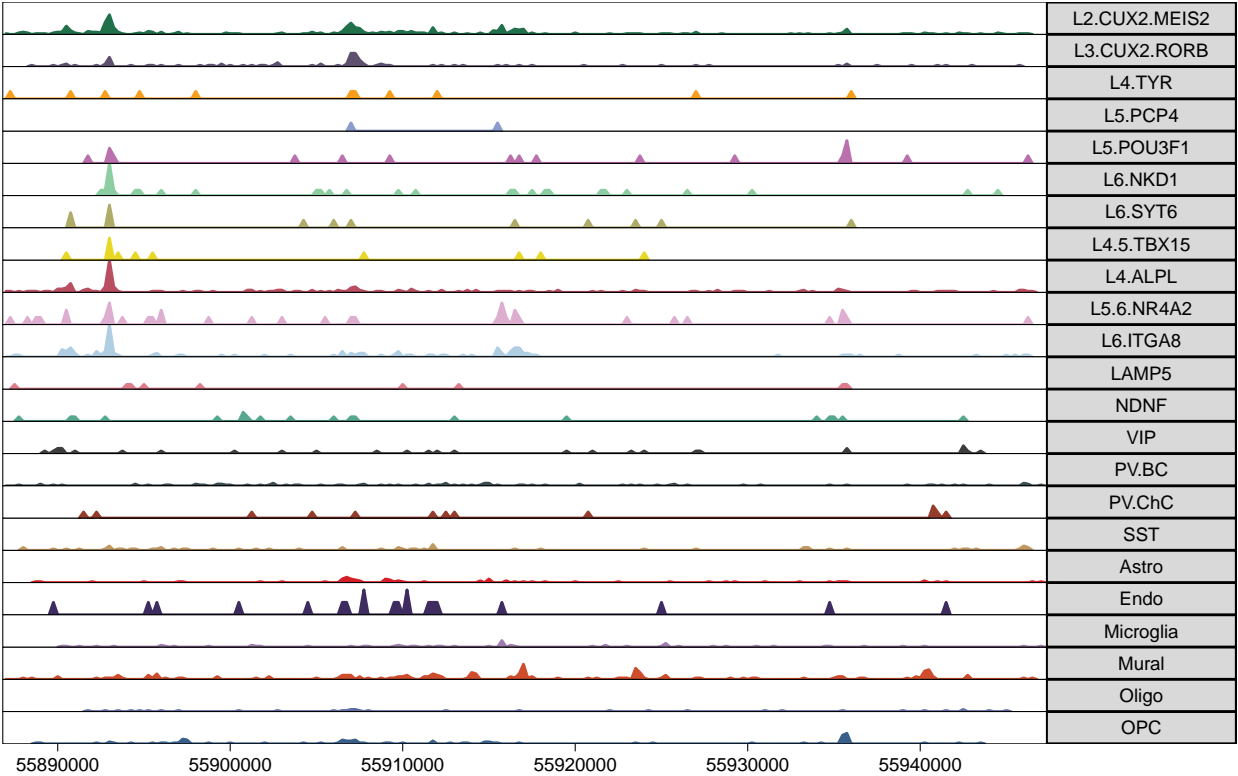
LoopTrack



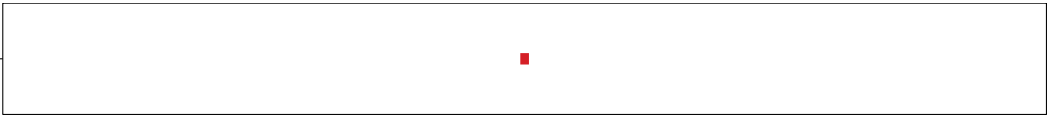
Genes

LIMCH1

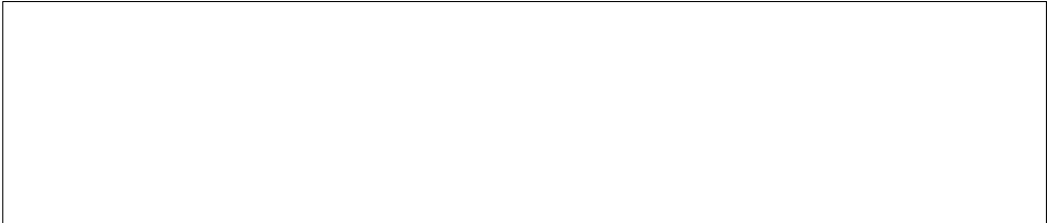
Coverage
(Norm. ATAC Signal Range 0-1.01) by ReadsInTSS)



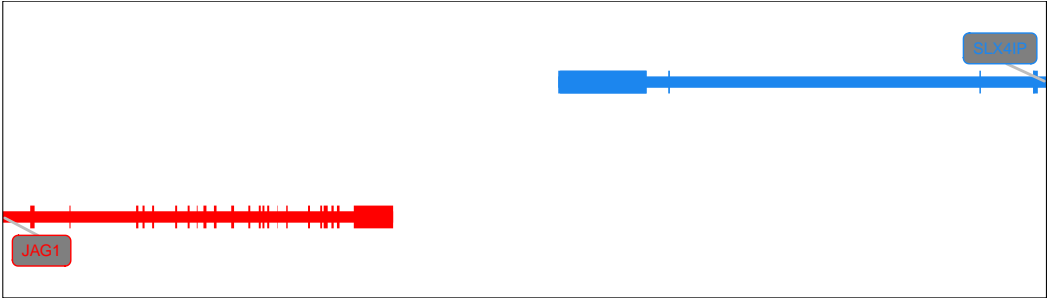
L5.6.NR4A2



Peaks

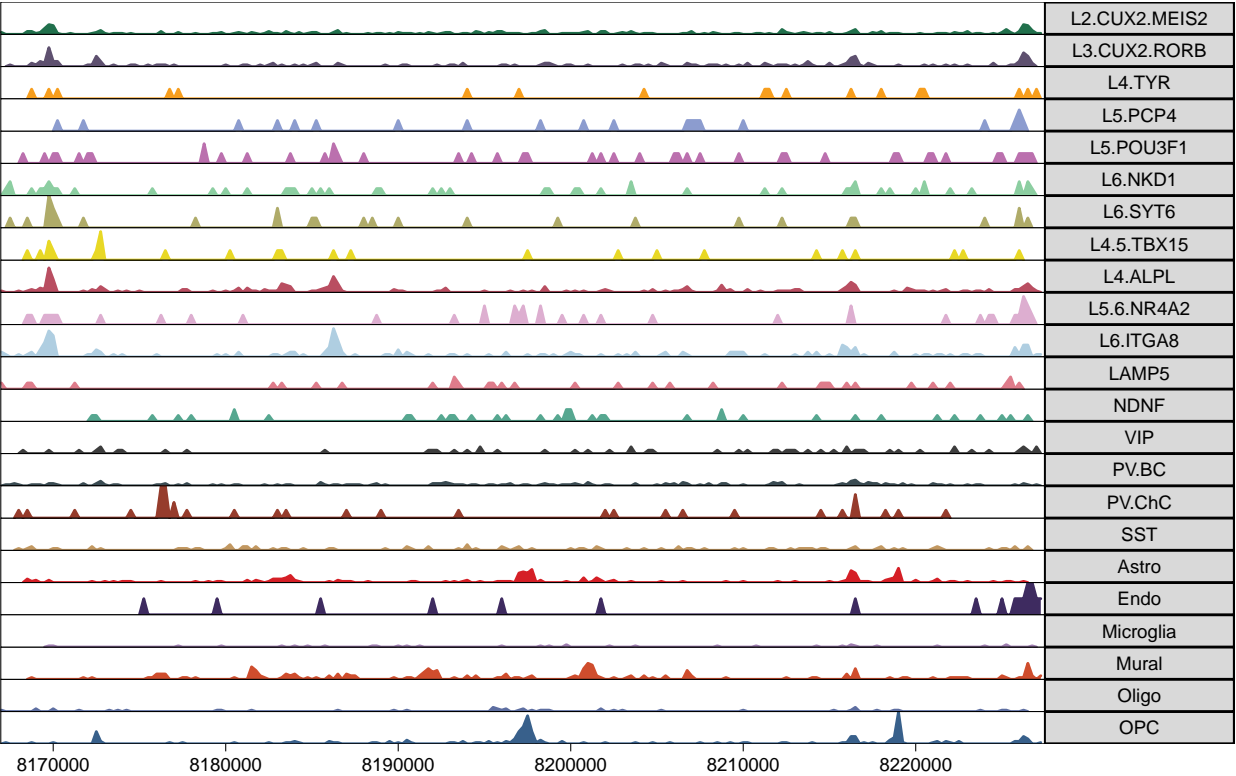


LoopTrack

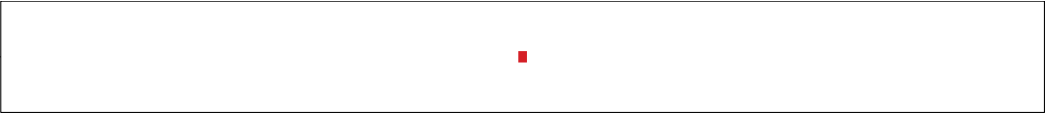


Genes

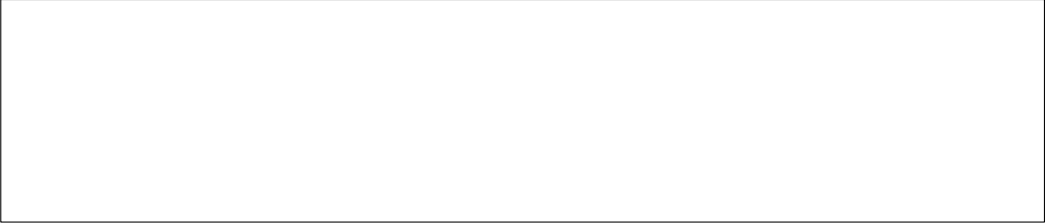
Coverage
(Norm. ATAC Signal Range (0-0.78) by ReadsInTSS)



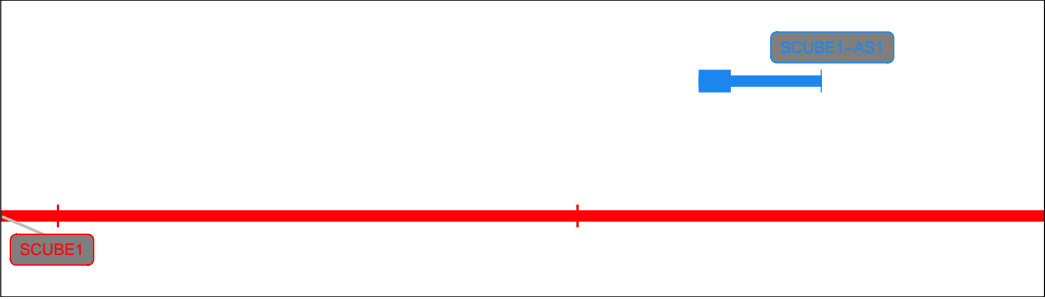
L5.6.NR4A2



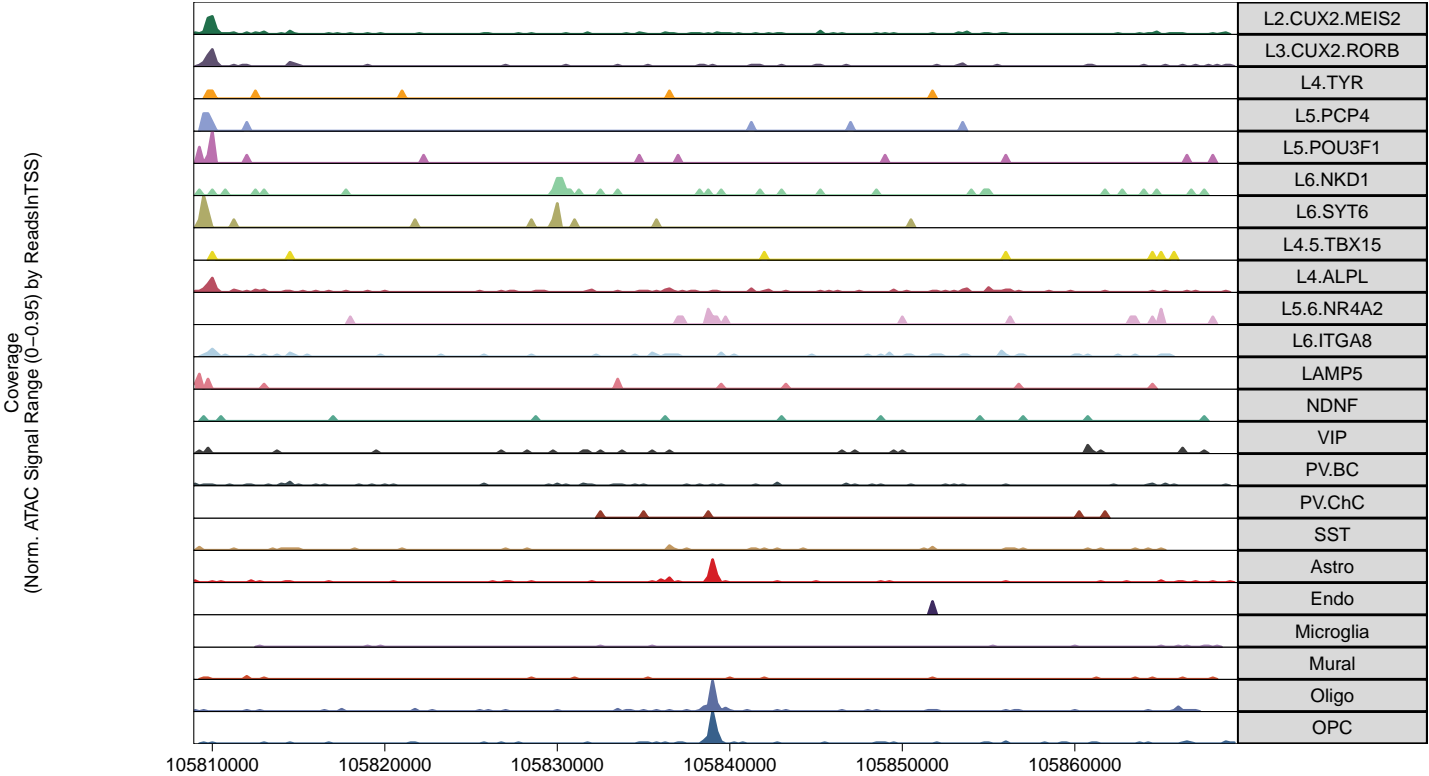
Peaks



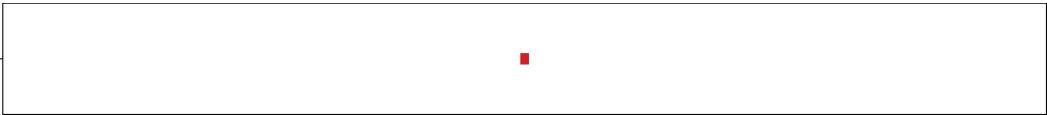
LoopTrack



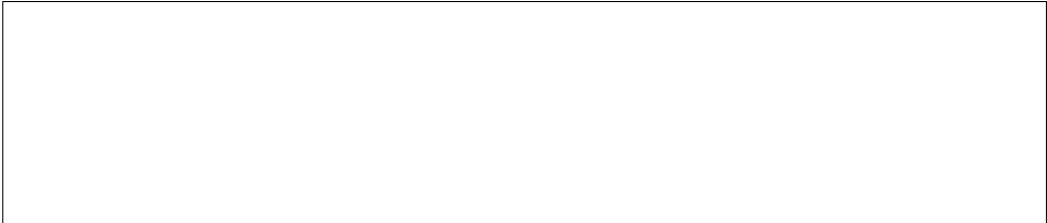
Genes



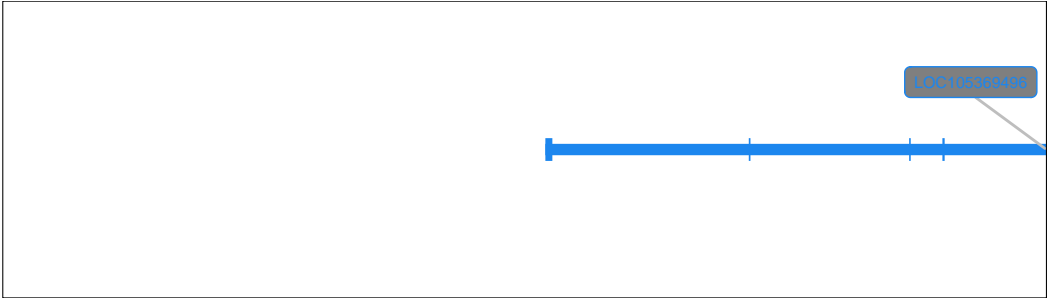
L5.6.NR4A2



Peaks

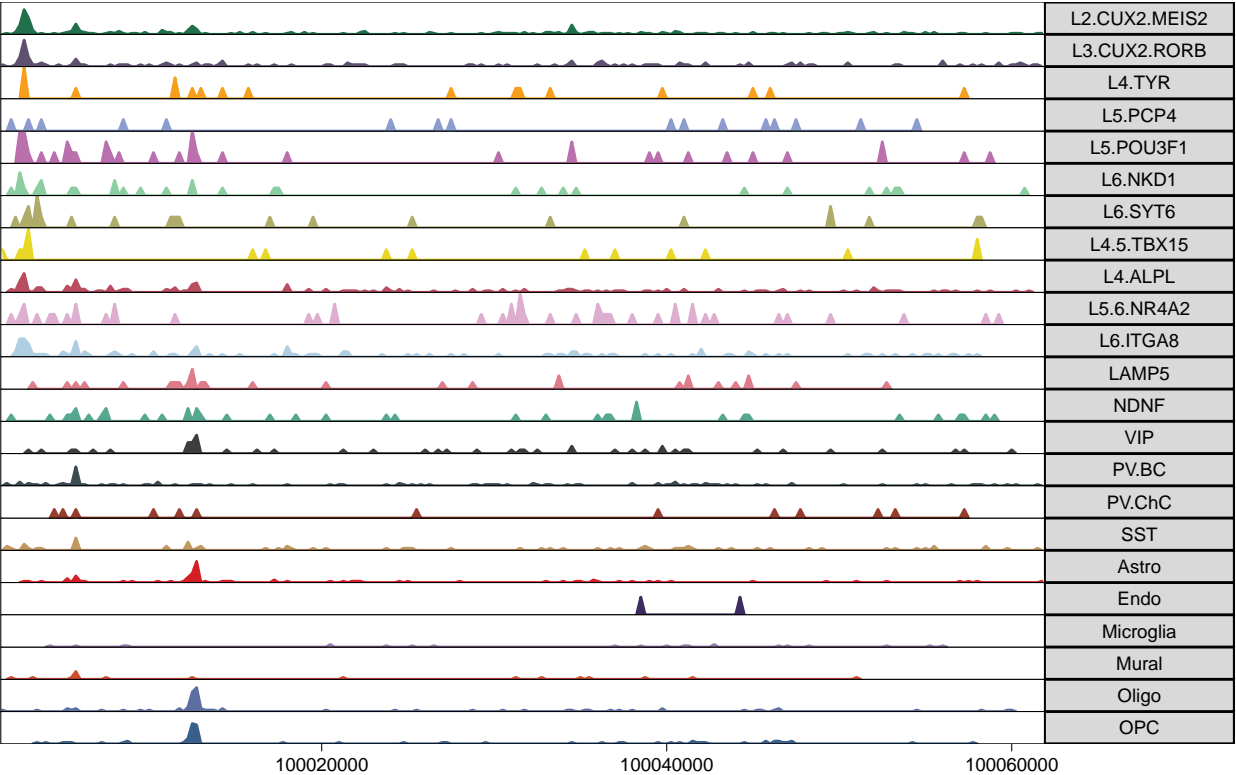


LoopTrack

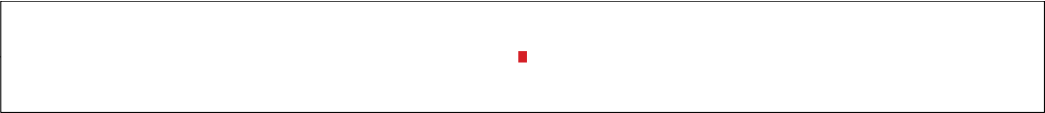


Genes

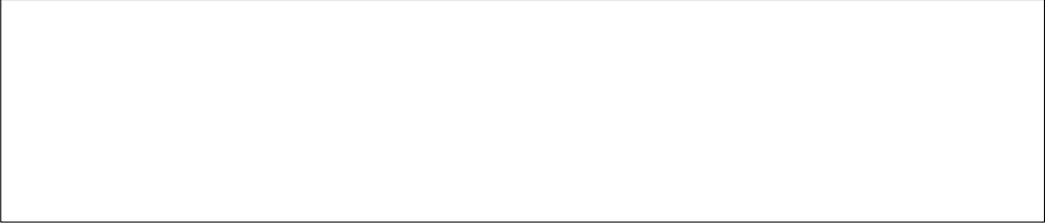
Coverage
(Norm. ATAC Signal Range (0-0.71) by ReadsInTSS)



L5.6.NR4A2



Peaks

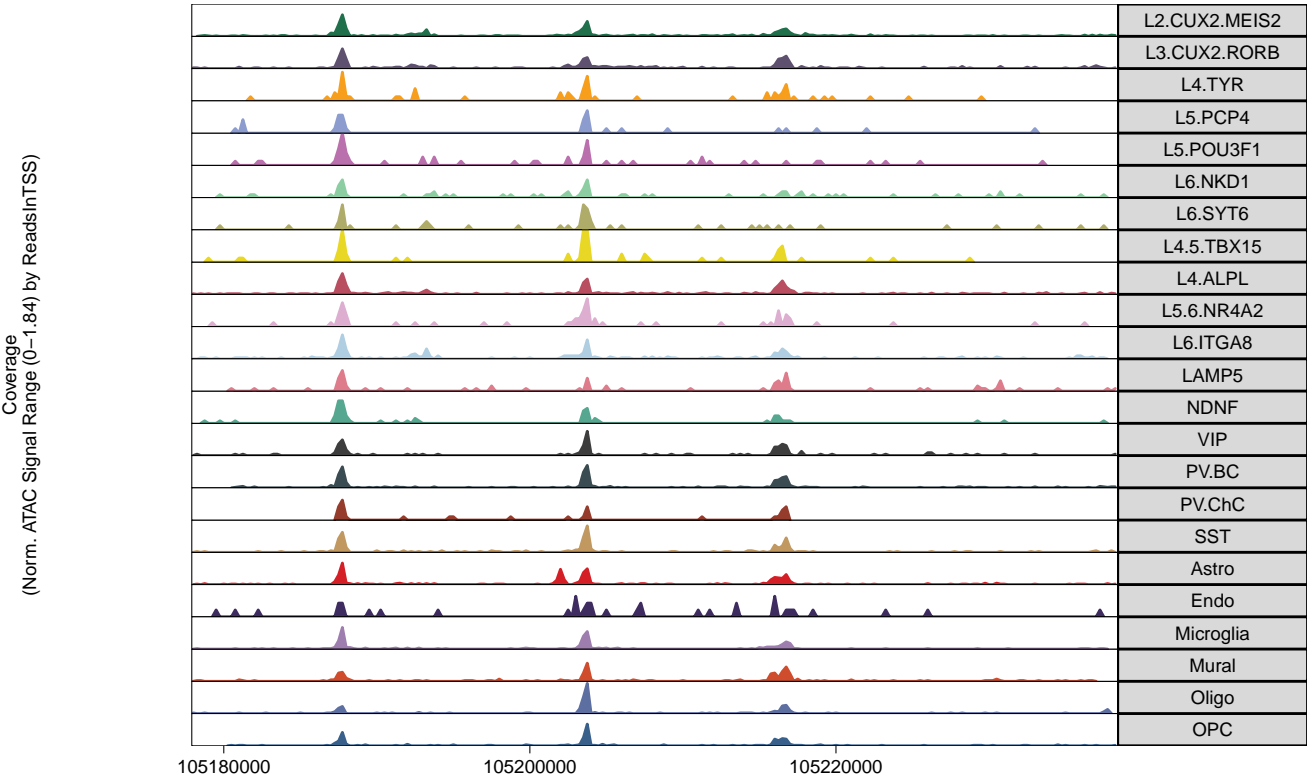


LoopTrack



Genes

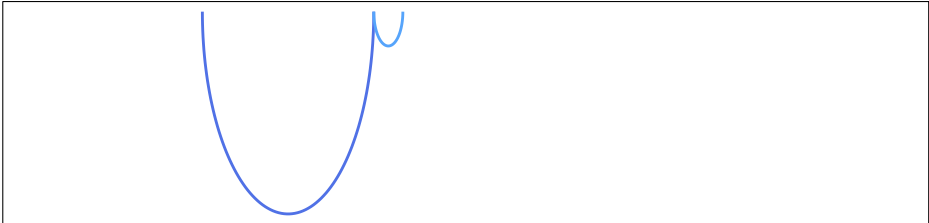
chr2:105177909-105238410 Marker Peak Tracks for: L5.6.NR4A2



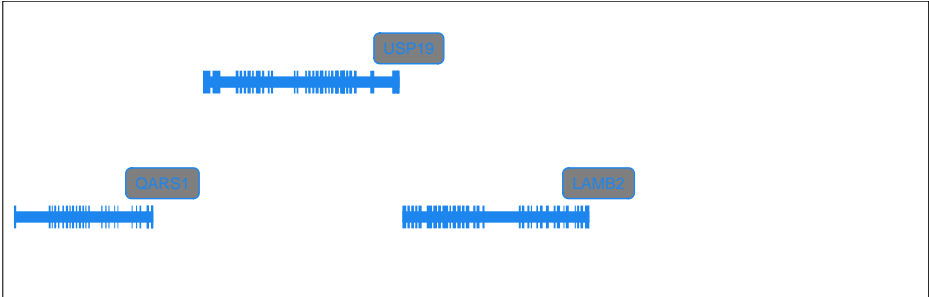
L5.6.NR4A2



Peaks



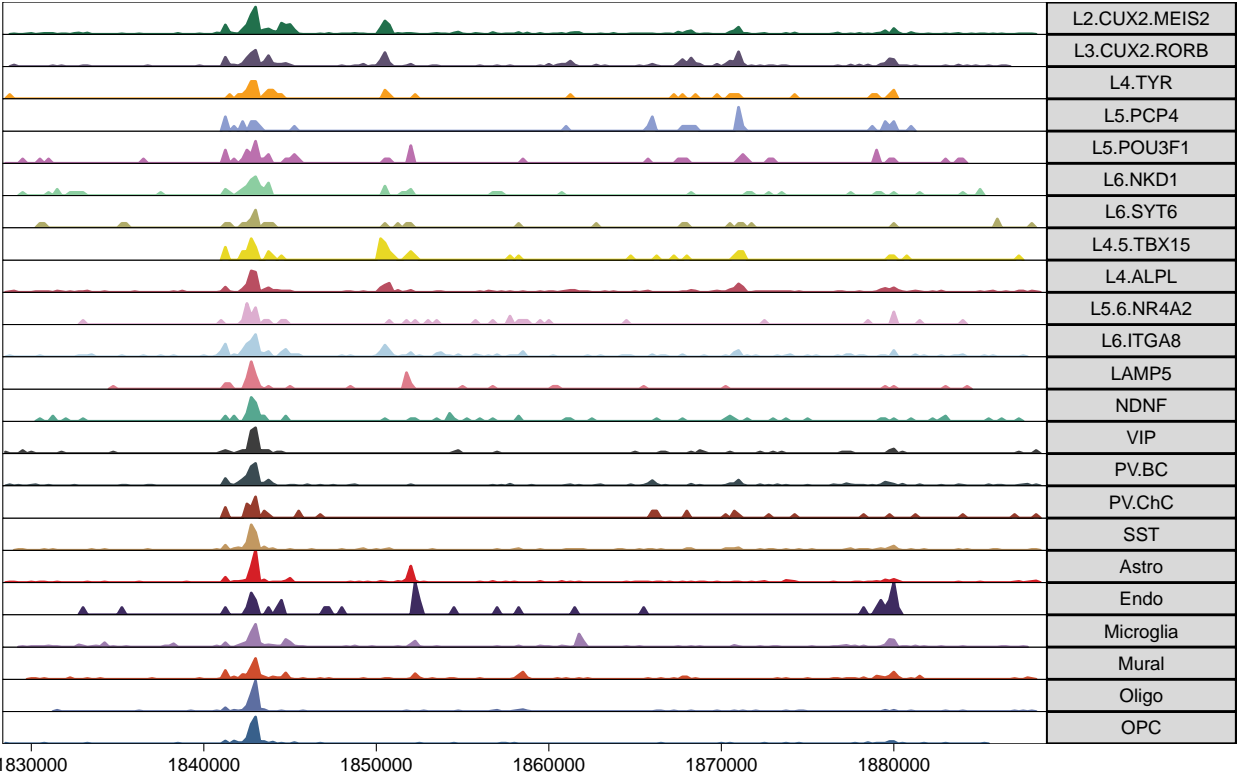
Peak2GeneLinks



Genes

chr3:1828346–1888847 Marker Peak Tracks for: L5.6.NR4A2

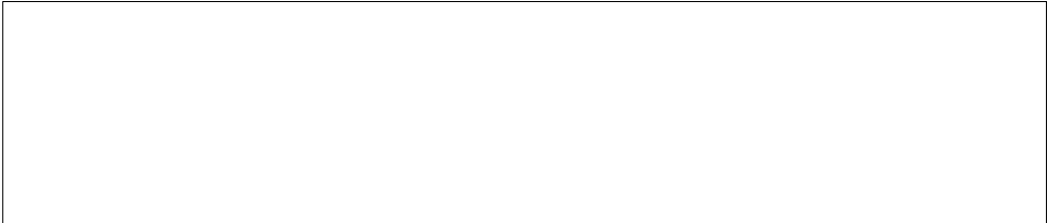
Coverage
(Norm. ATAC Signal Range 0–1.74) by ReadsInTSS)



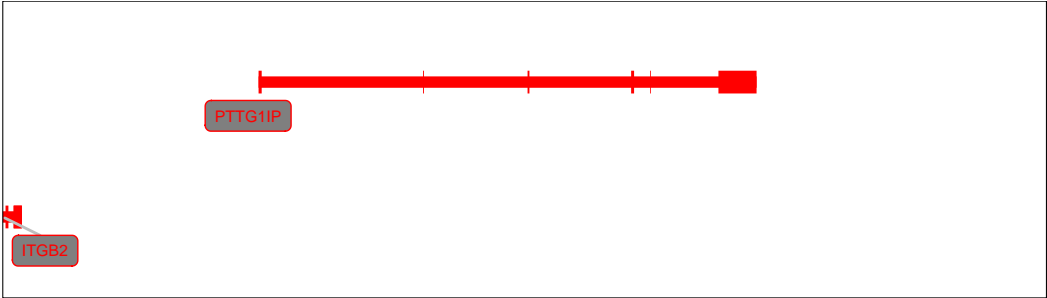
L5.6.NR4A2



Peaks

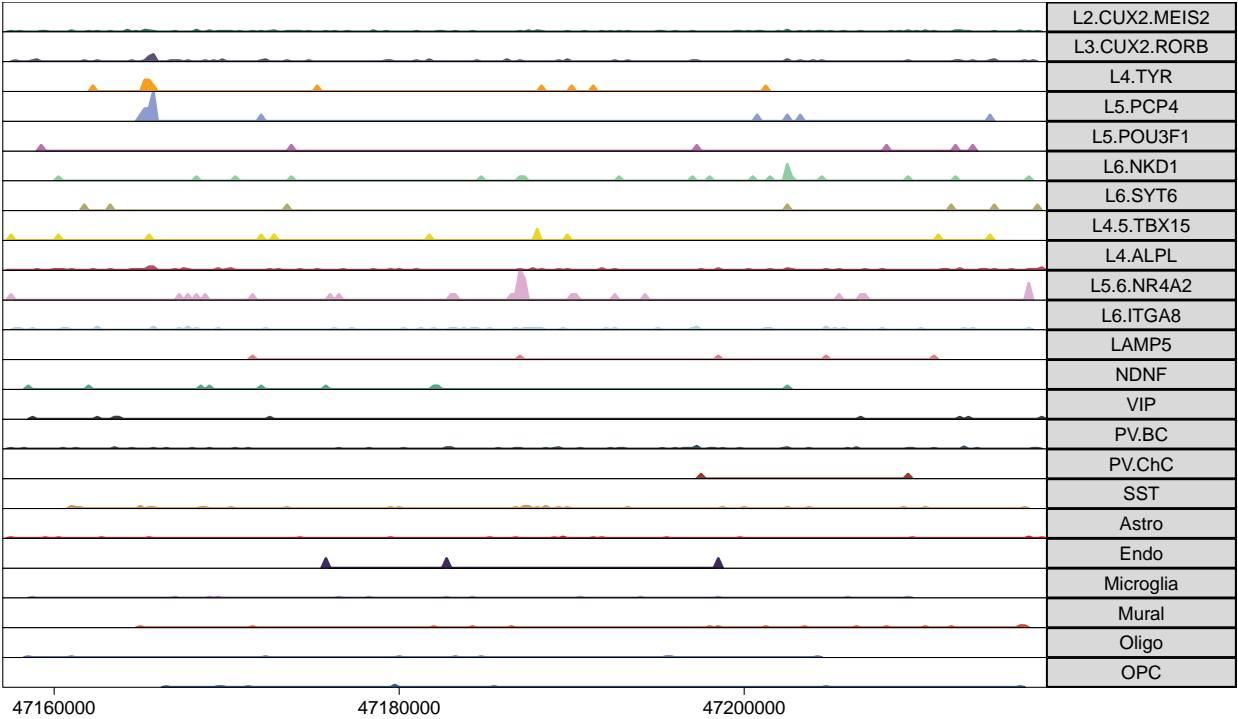


LoopTrack

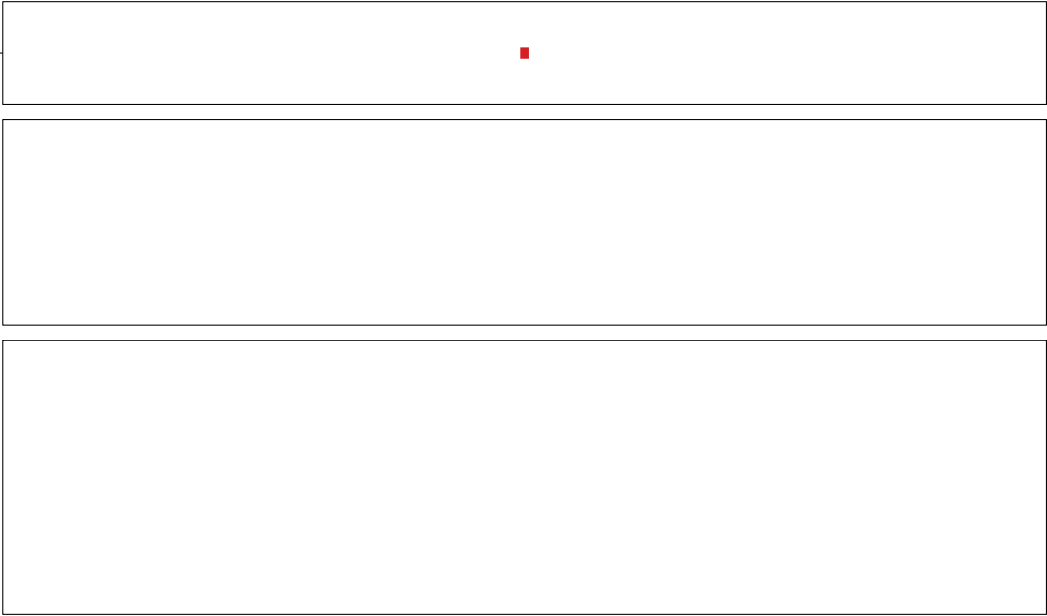


Genes

Coverage
(Norm. ATAC Signal Range (0–1.21) by Reads in TSS)

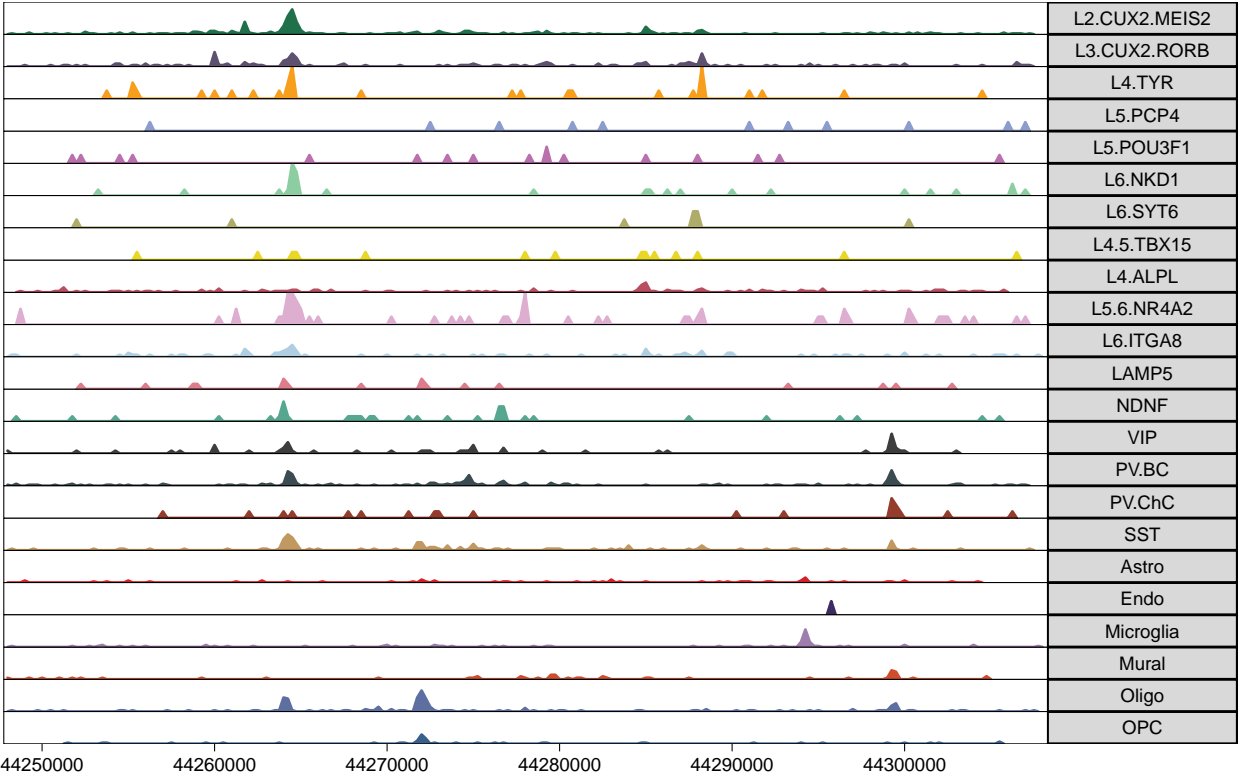


L5.6.NR4A2

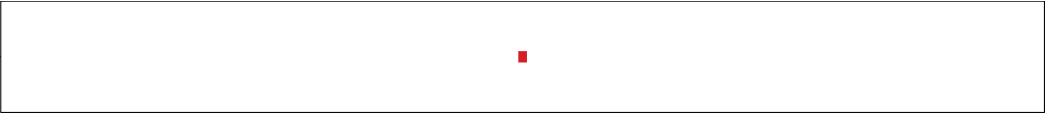


chr3:44247780–44308281 Marker Peak Tracks for: L5.6.NR4A2

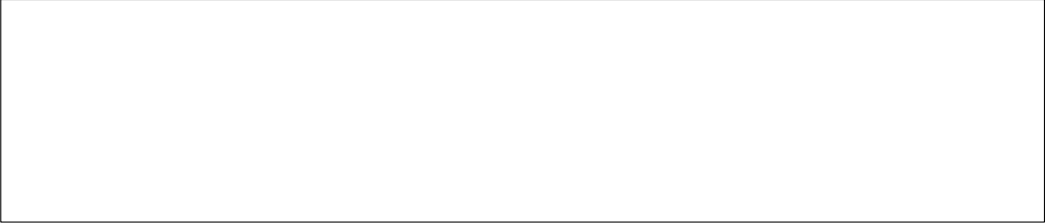
Coverage
(Norm. ATAC Signal Range 0–0.93) by ReadsInTSS)



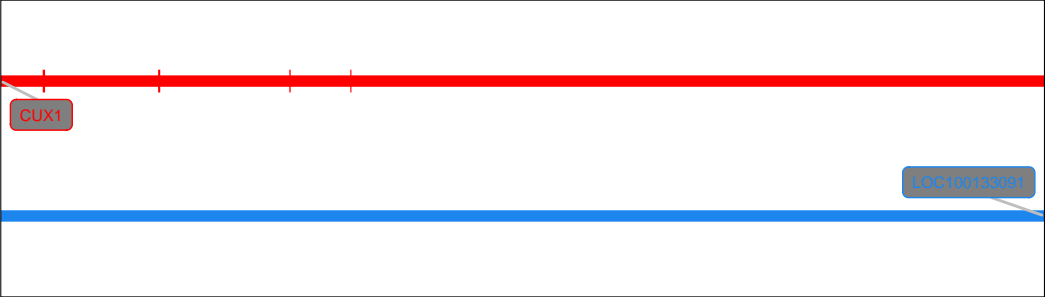
L5.6.NR4A2



Peaks

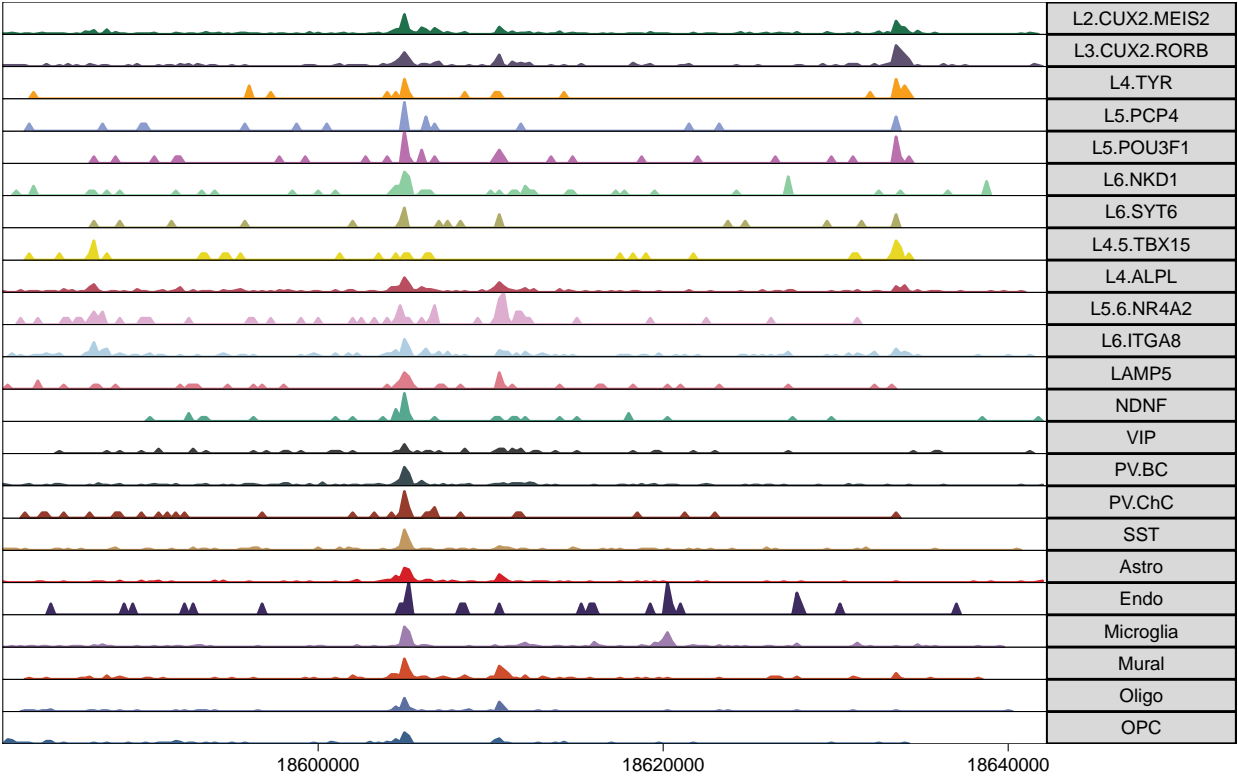


LoopTrack

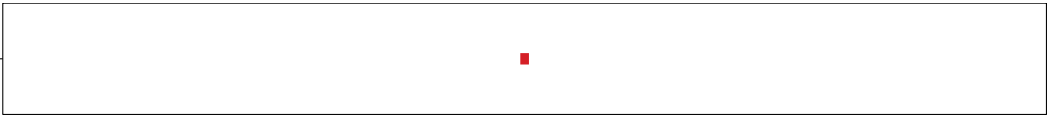


Genes

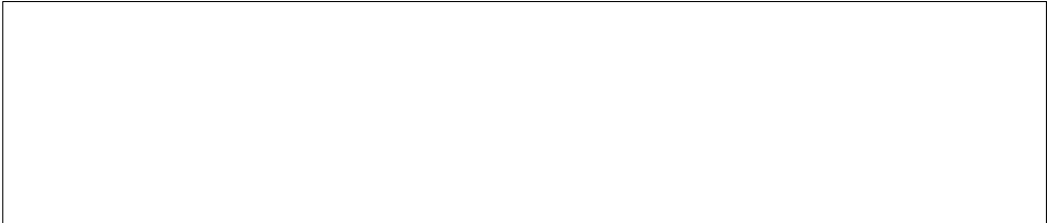
Coverage
(Norm. ATAC Signal Range 0-1.15) by ReadsInTSS)



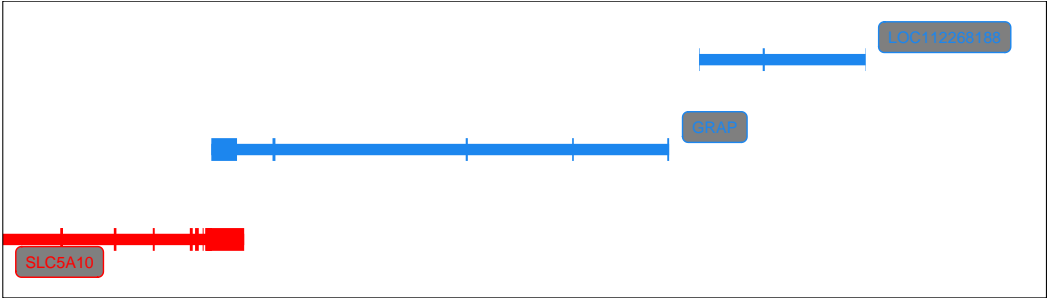
L5.6.NR4A2



Peaks

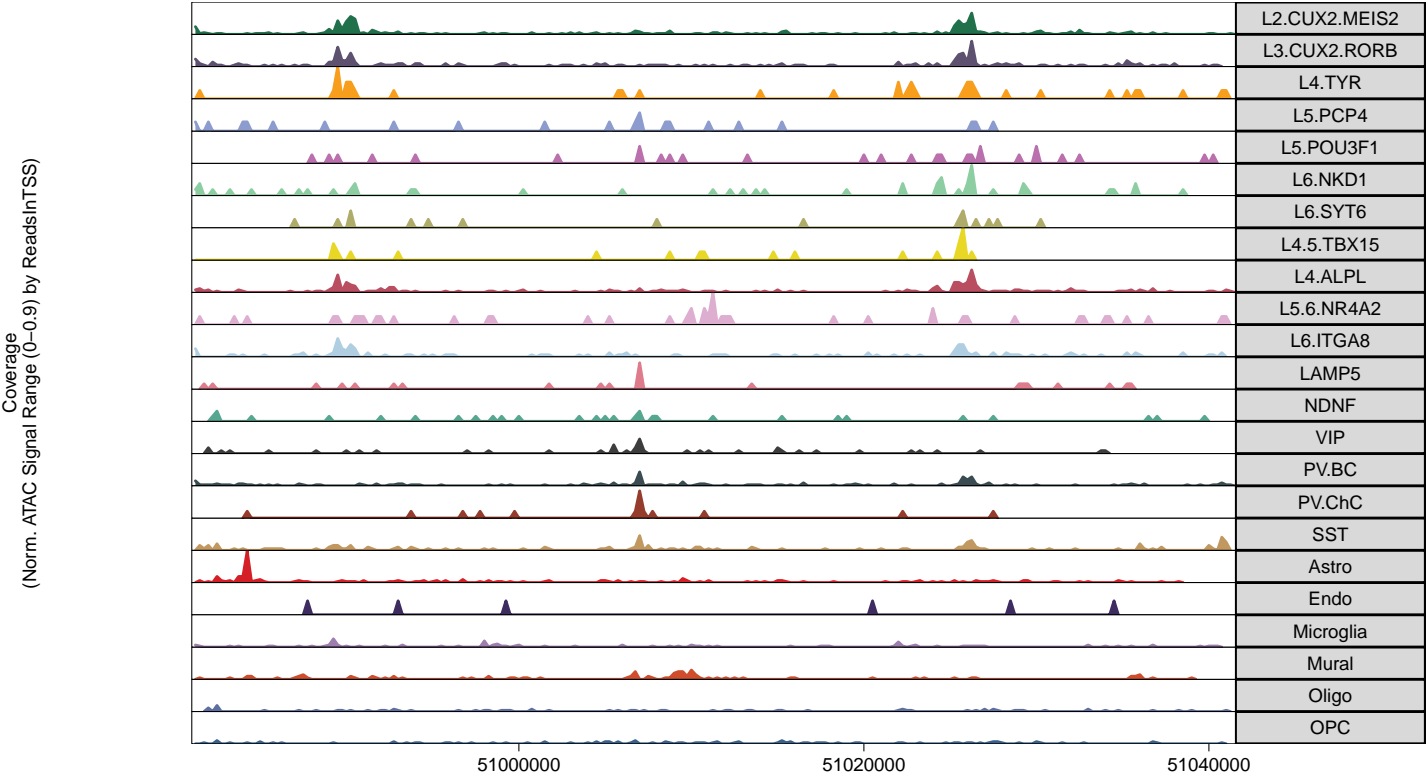


LoopTrack

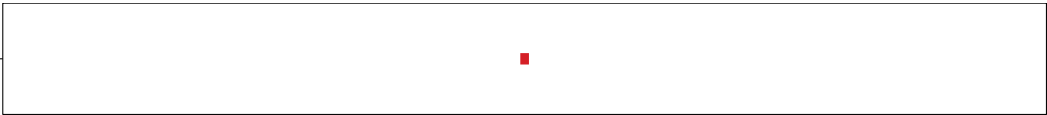


Genes

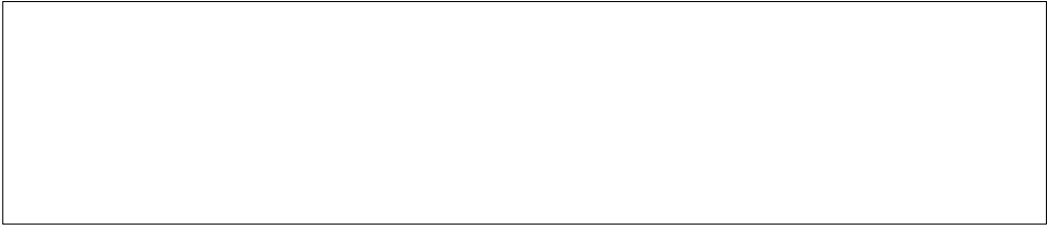
chr5:50981038–51041539 Marker Peak Tracks for: L5.6.NR4A2



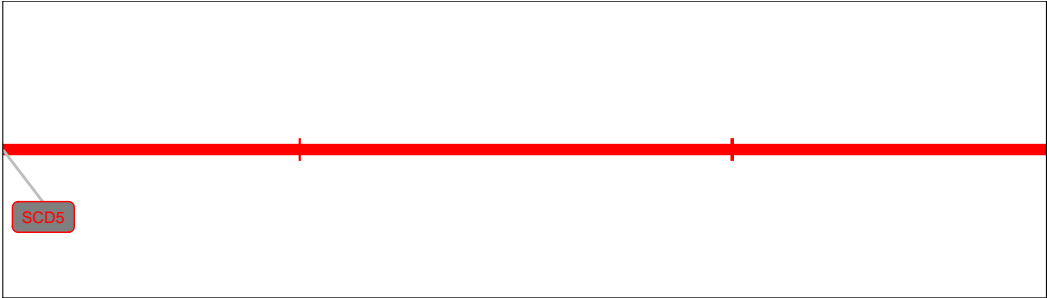
L5.6.NR4A2



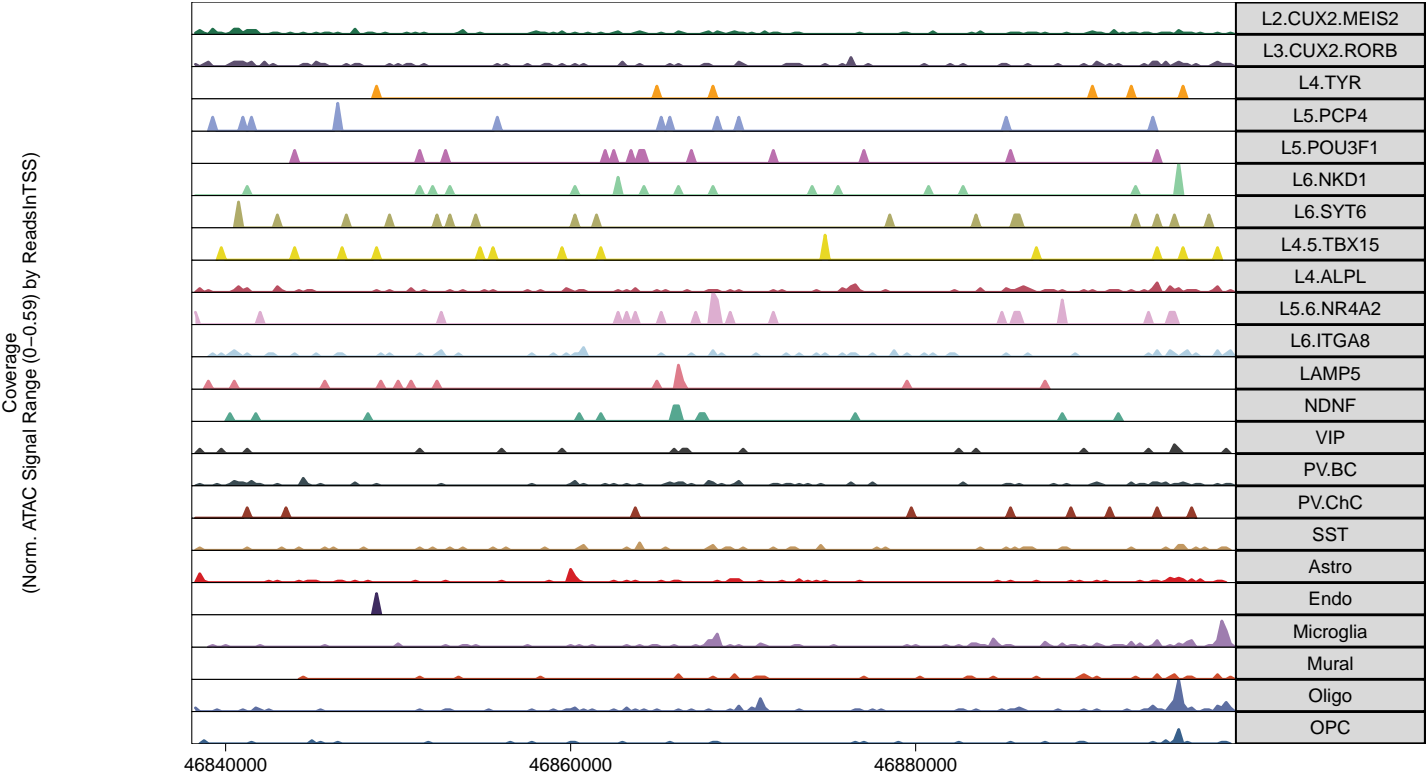
Peaks



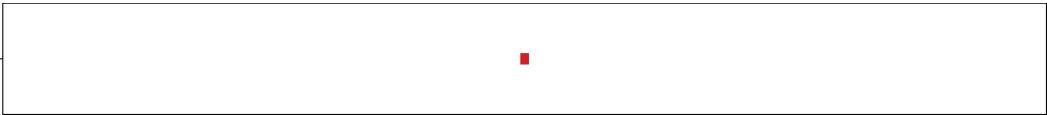
LoopTrack



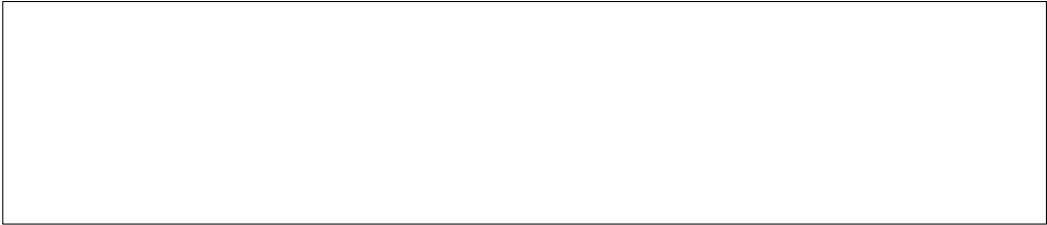
Genes



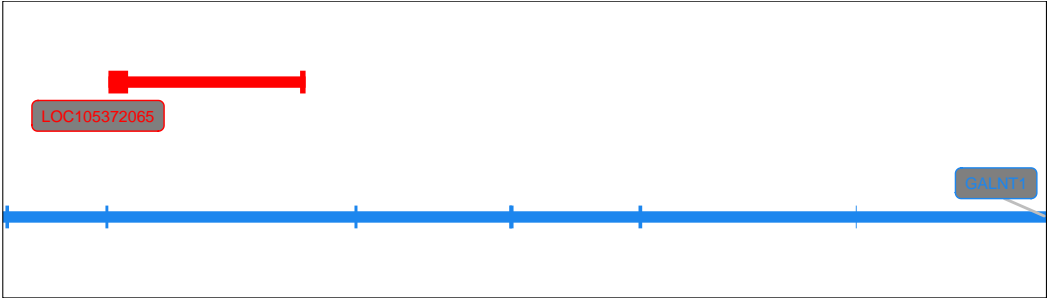
L5.6.NR4A2



Peaks

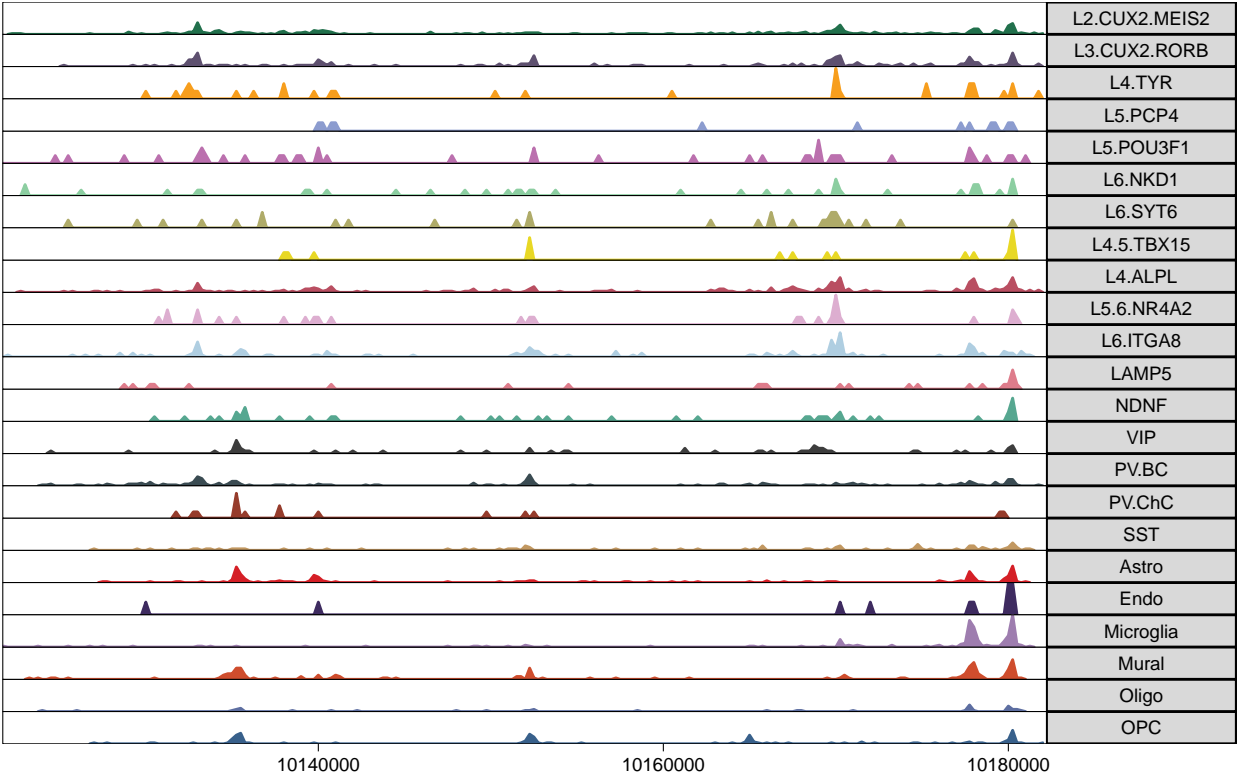


LoopTrack

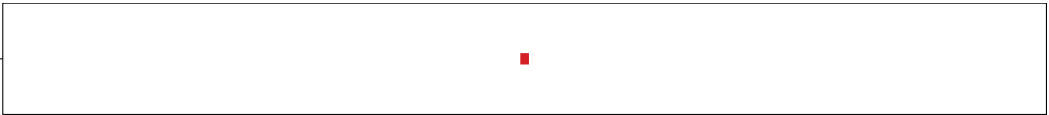


Genes

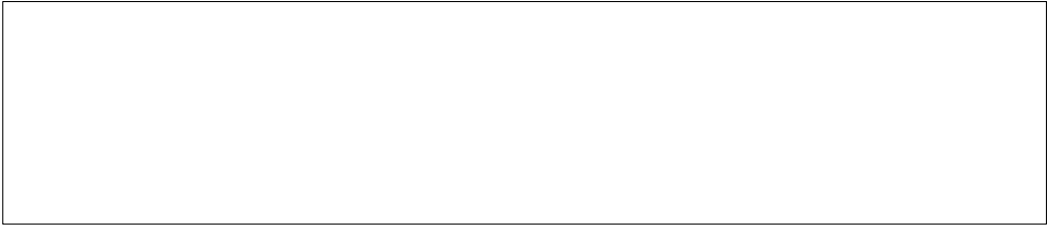
Coverage
(Norm. ATAC Signal Range 0–0.99) by ReadsInTSS)



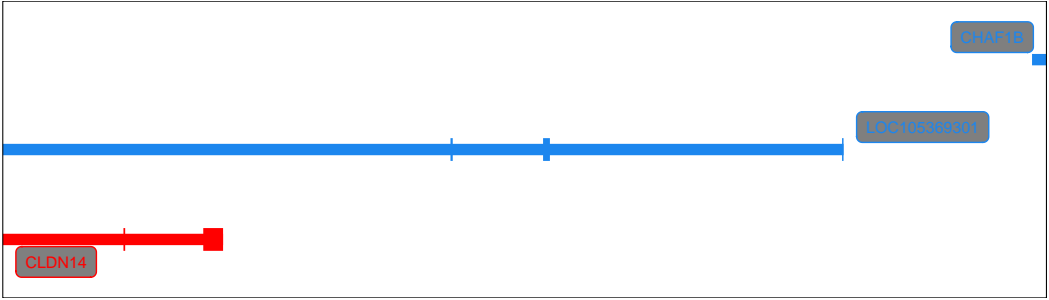
L5.6.NR4A2



Peaks



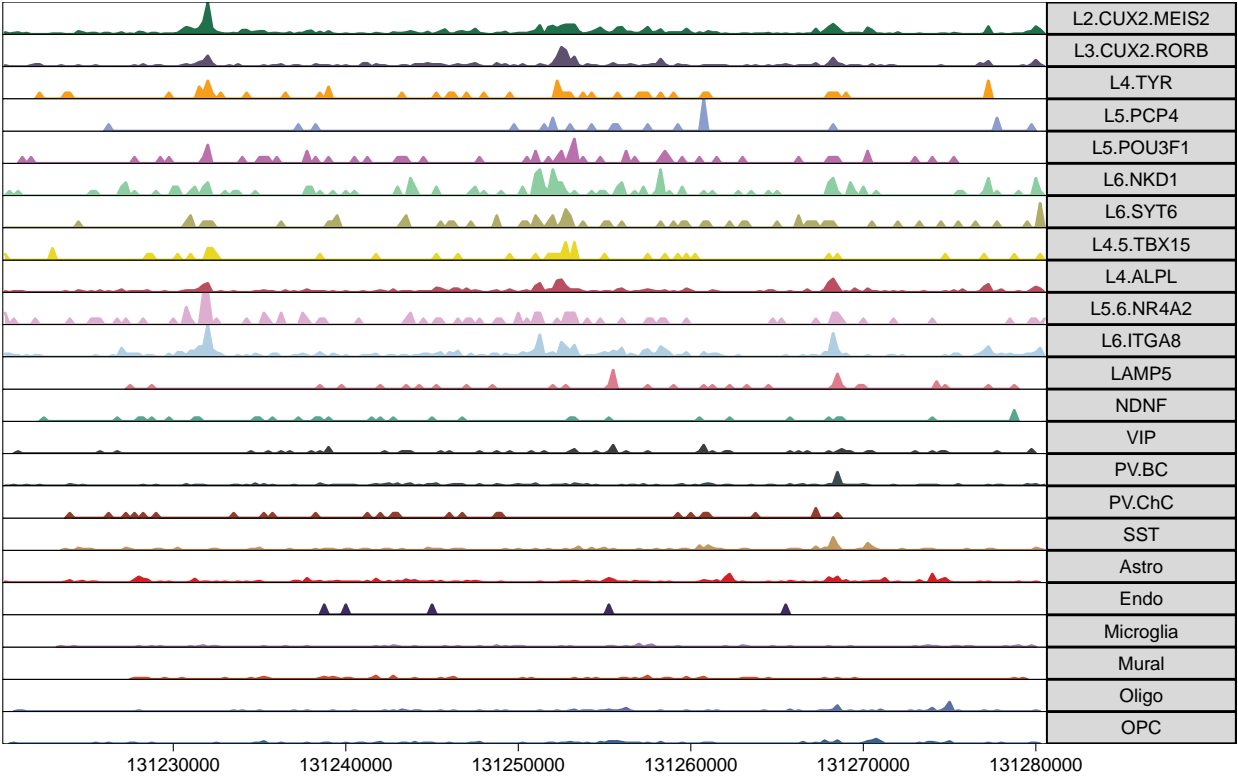
LoopTrack



Genes

chr4:131220115-131280616 Marker Peak Tracks for: L5.6.NR4A2

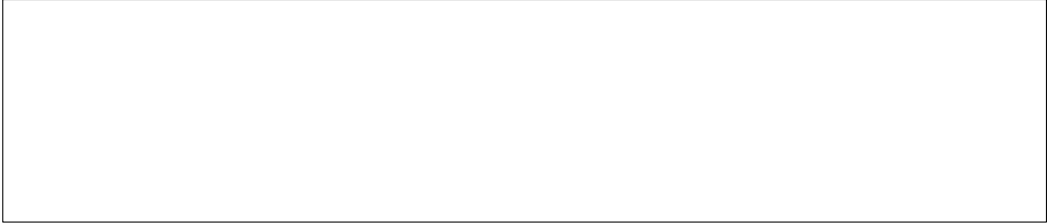
Coverage
(Norm. ATAC Signal Range 0-1.26) by ReadsInTSS)



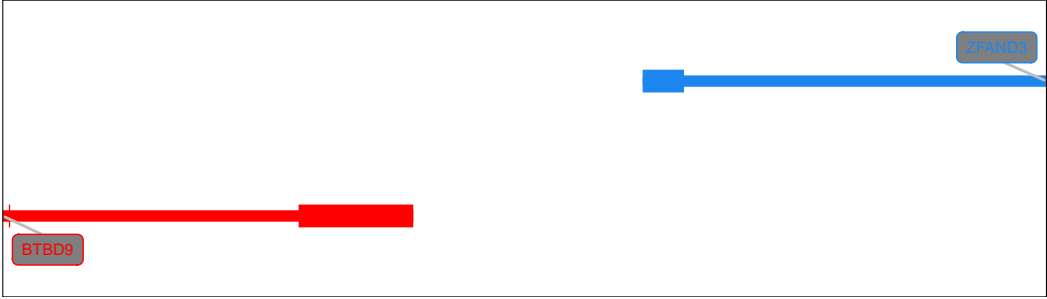
L5.6.NR4A2



Peaks

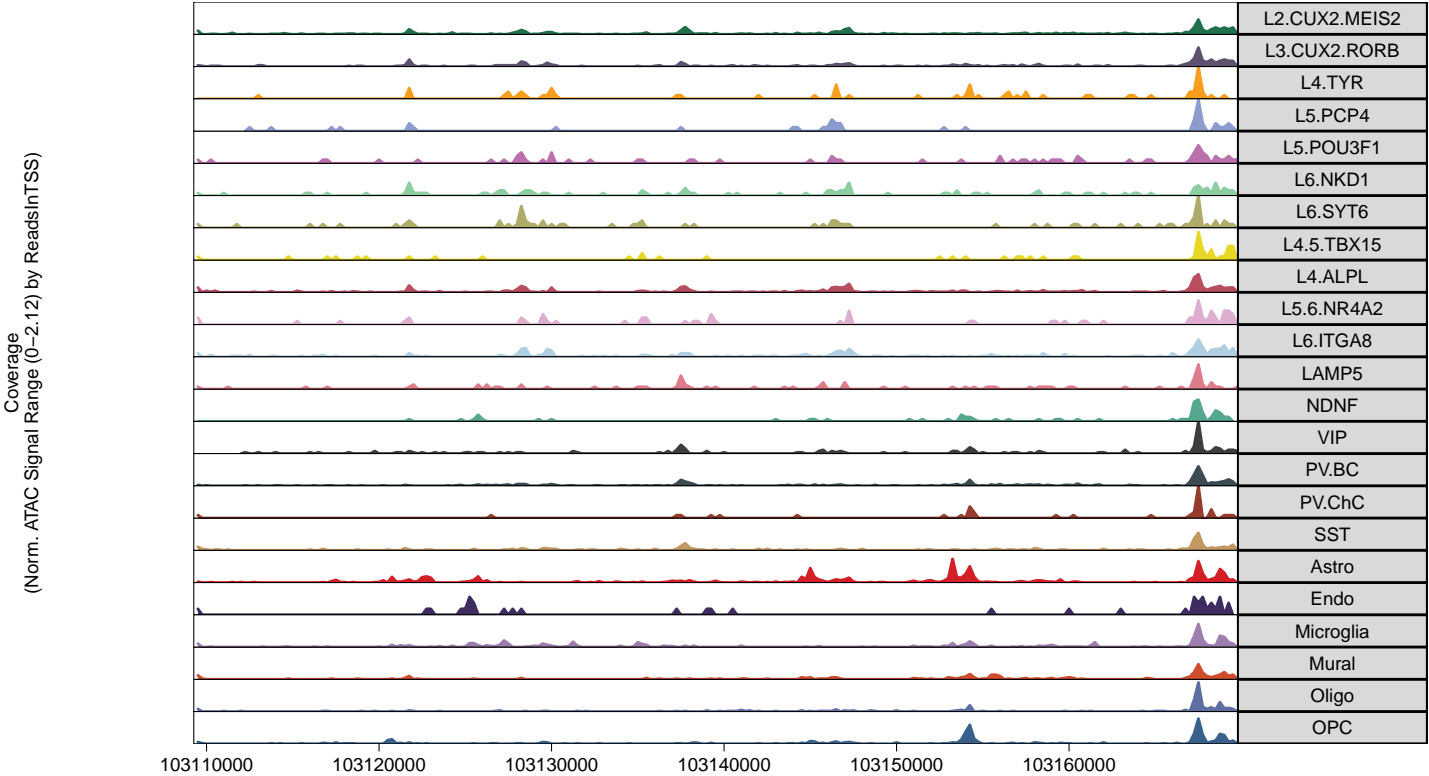


LoopTrack

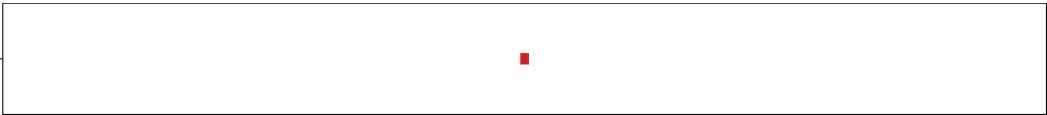


Genes

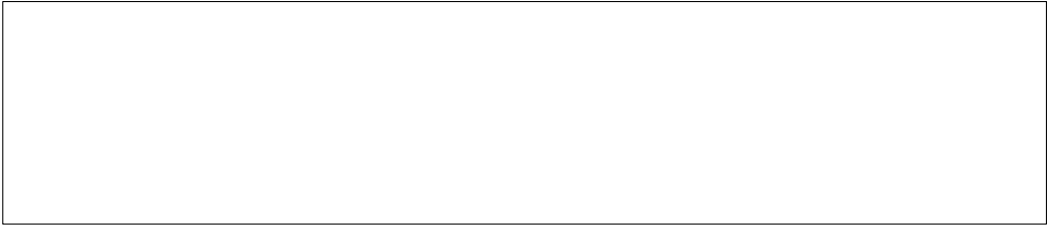
chr9:103109261–103169762 Marker Peak Tracks for: L5.6.NR4A2



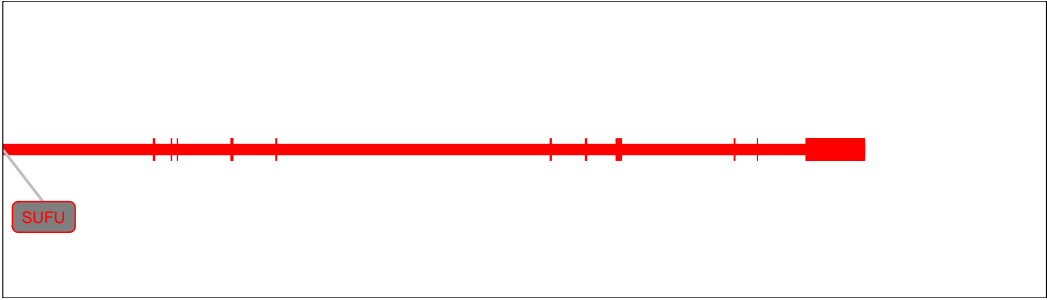
L5.6.NR4A2



Peaks

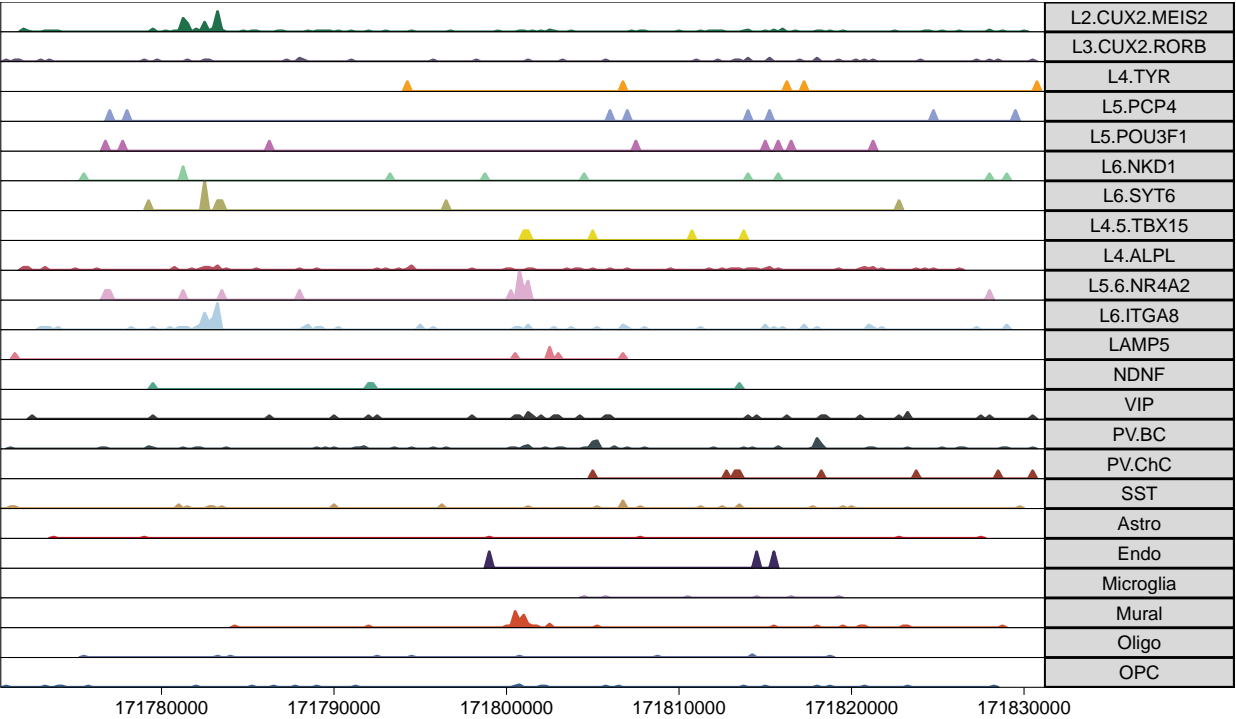


LoopTrack

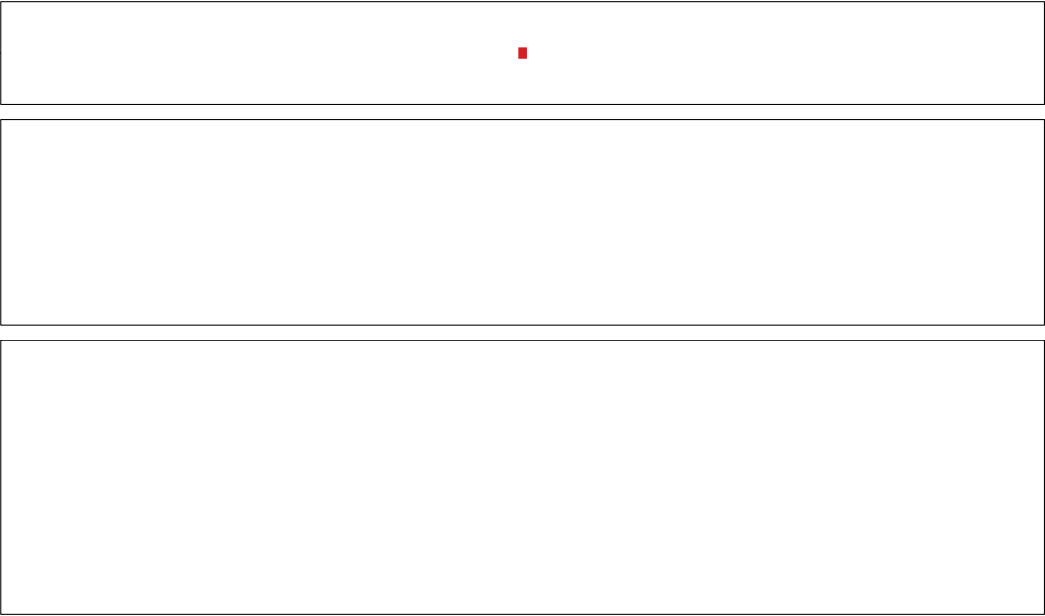


Genes

Coverage
(Norm. ATAC Signal Range (0–0.71) by Reads in TSS)



L5.6.NR4A2



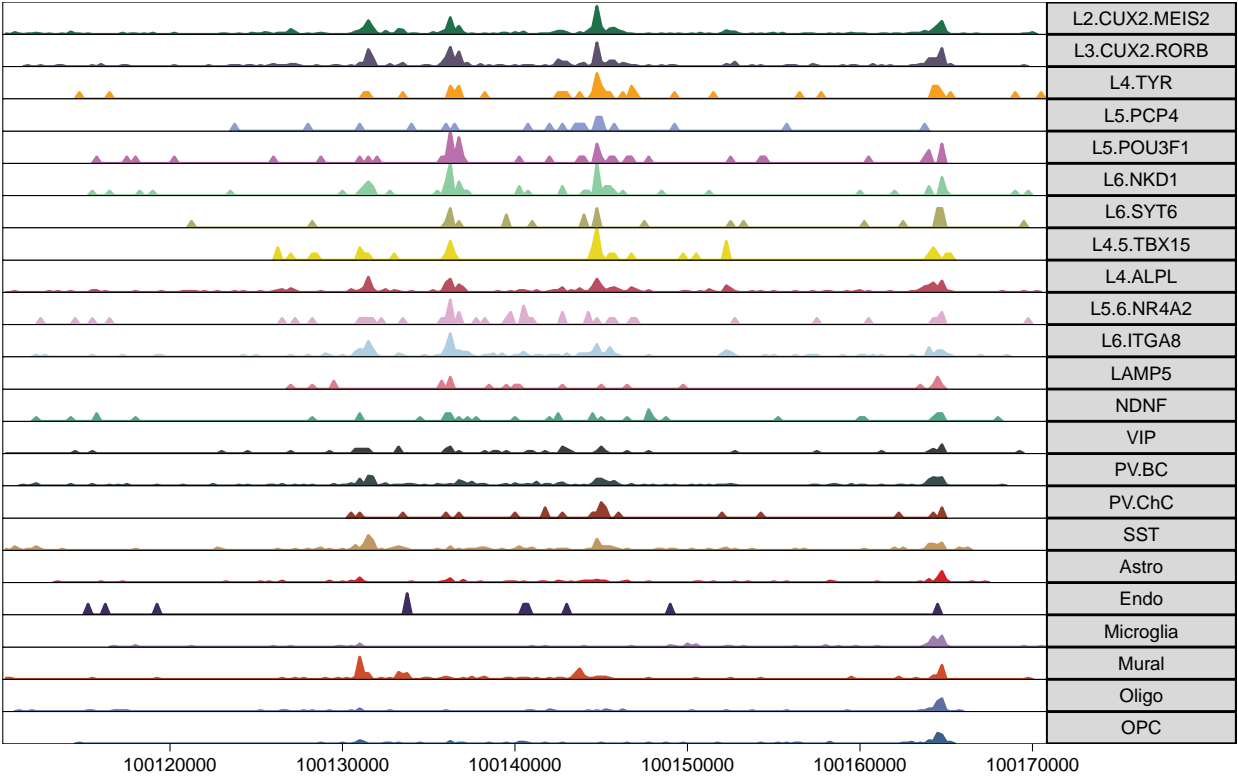
Peaks

LoopTrack

GeneTrack

chr8:100110308–100170809 Marker Peak Tracks for: L5.6.NR4A2

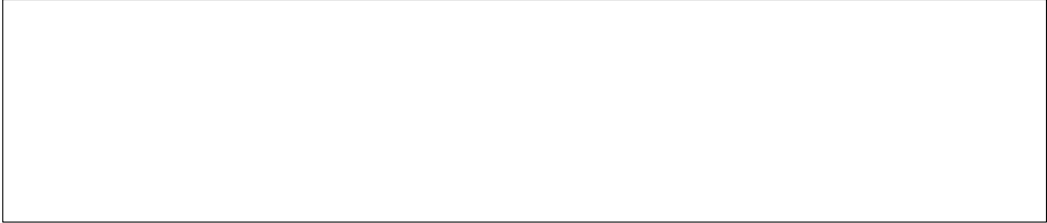
Coverage
(Norm. ATAC Signal Range (0–1.19) by ReadsInTSS)



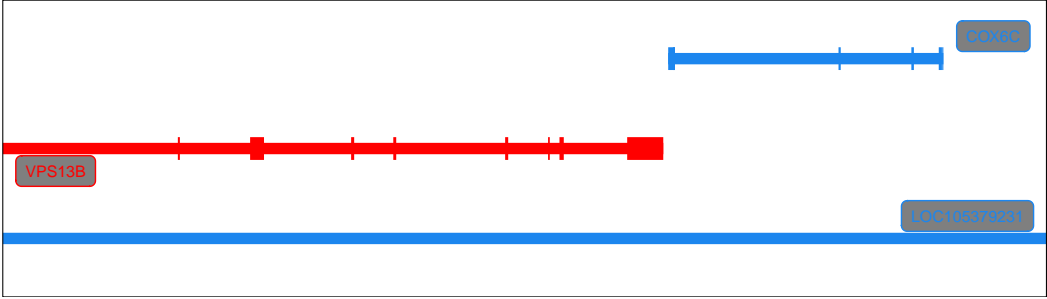
L5.6.NR4A2



Peaks

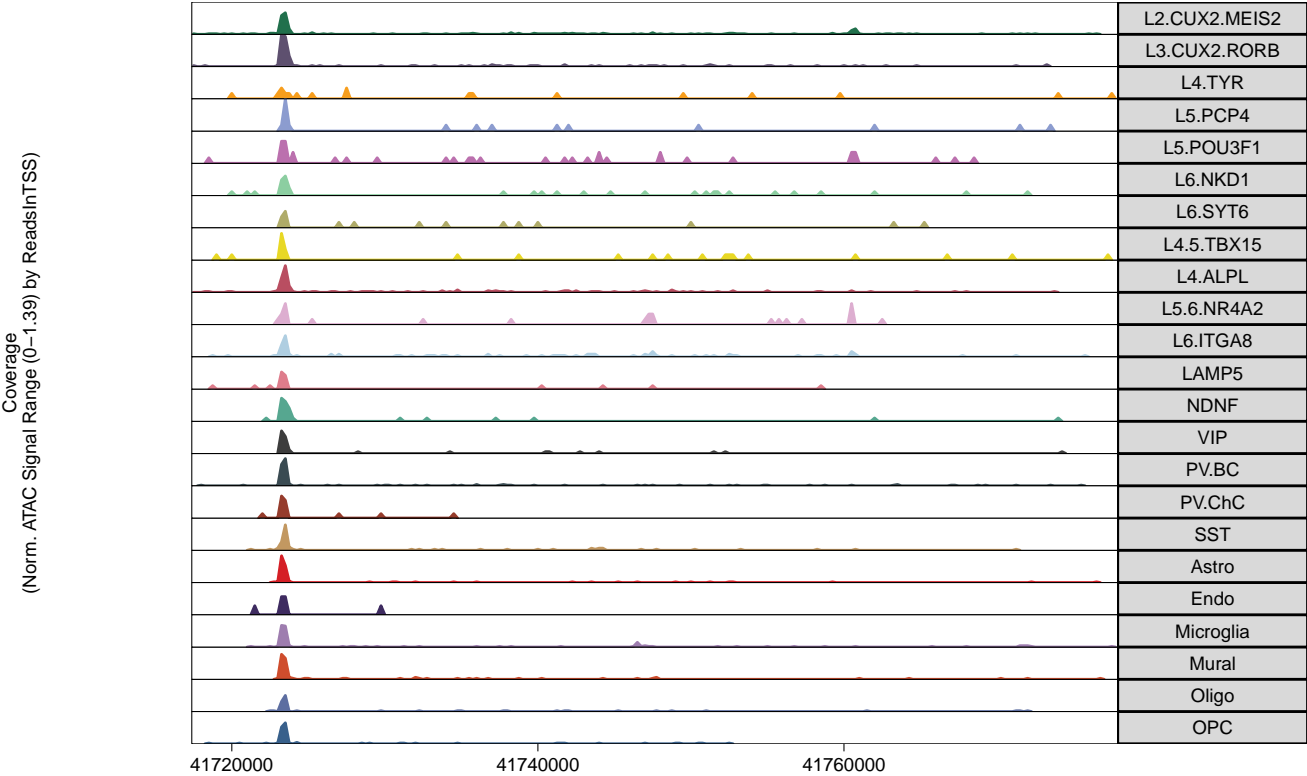


LoopTrack



Genes

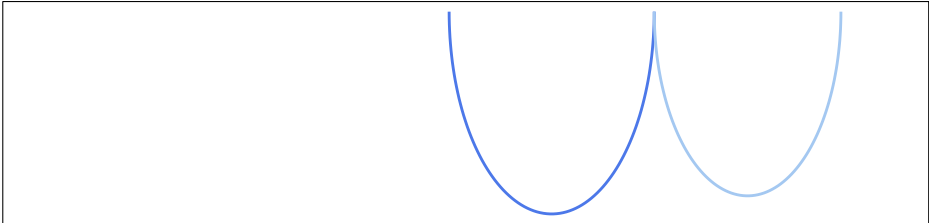
chr3:41717385–41777886 Marker Peak Tracks for: L5.6.NR4A2



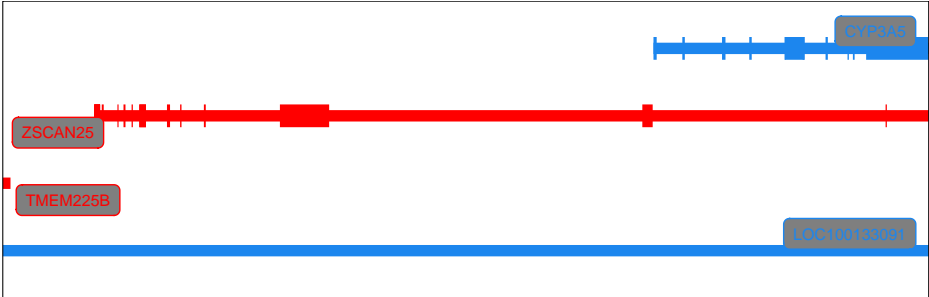
L5.6.NR4A2



Peaks

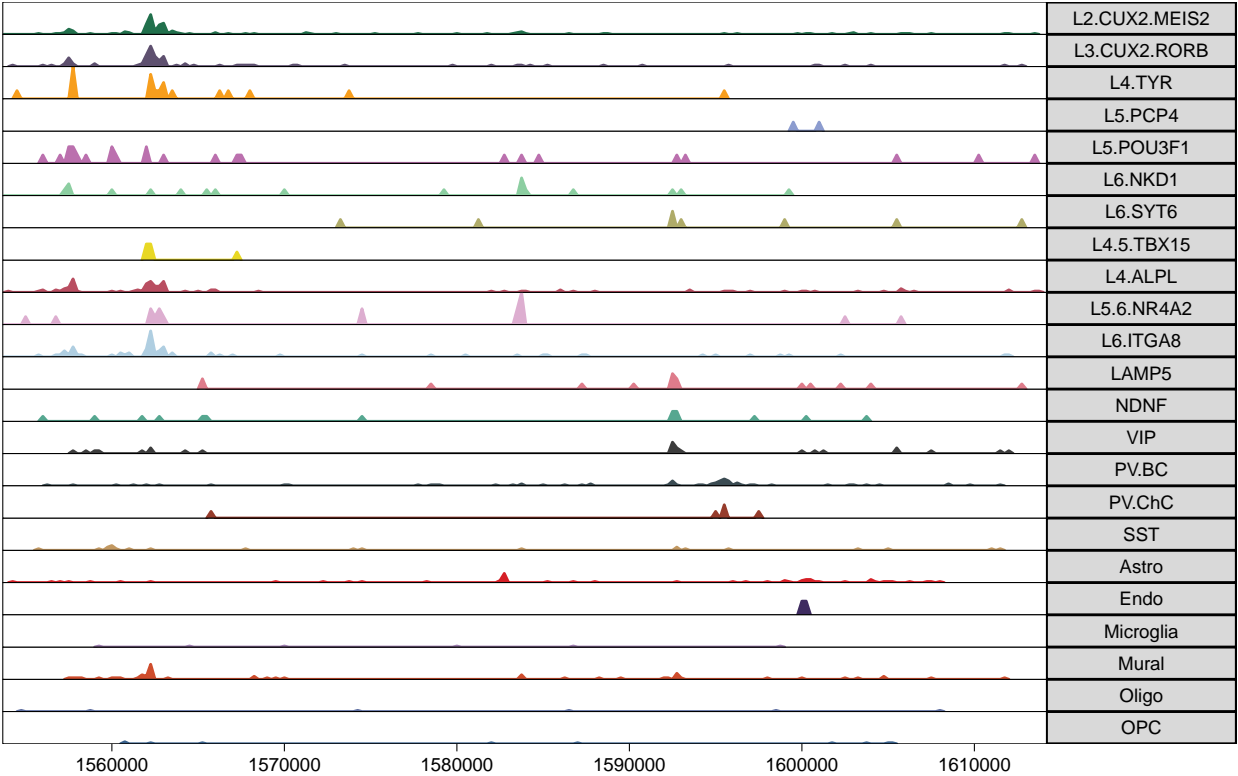


Peak2GeneLinks

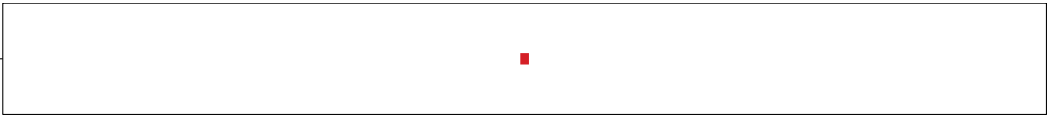


Genes

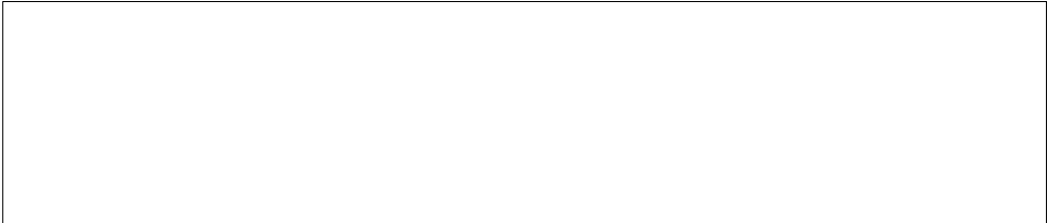
Coverage
(Norm. ATAC Signal Range (0–0.92) by ReadsInTSS)



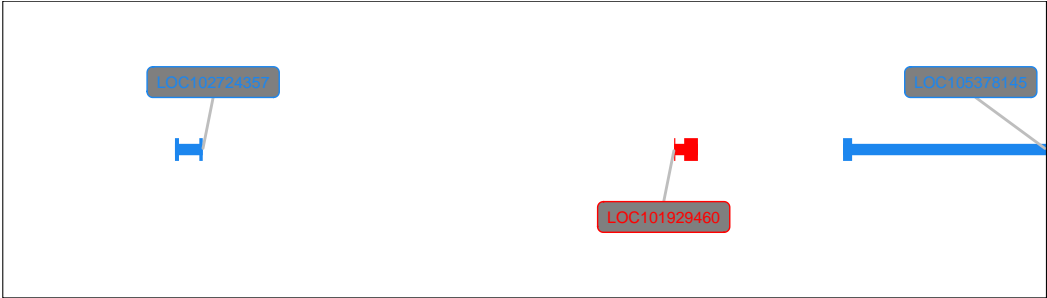
L5.6.NR4A2



Peaks

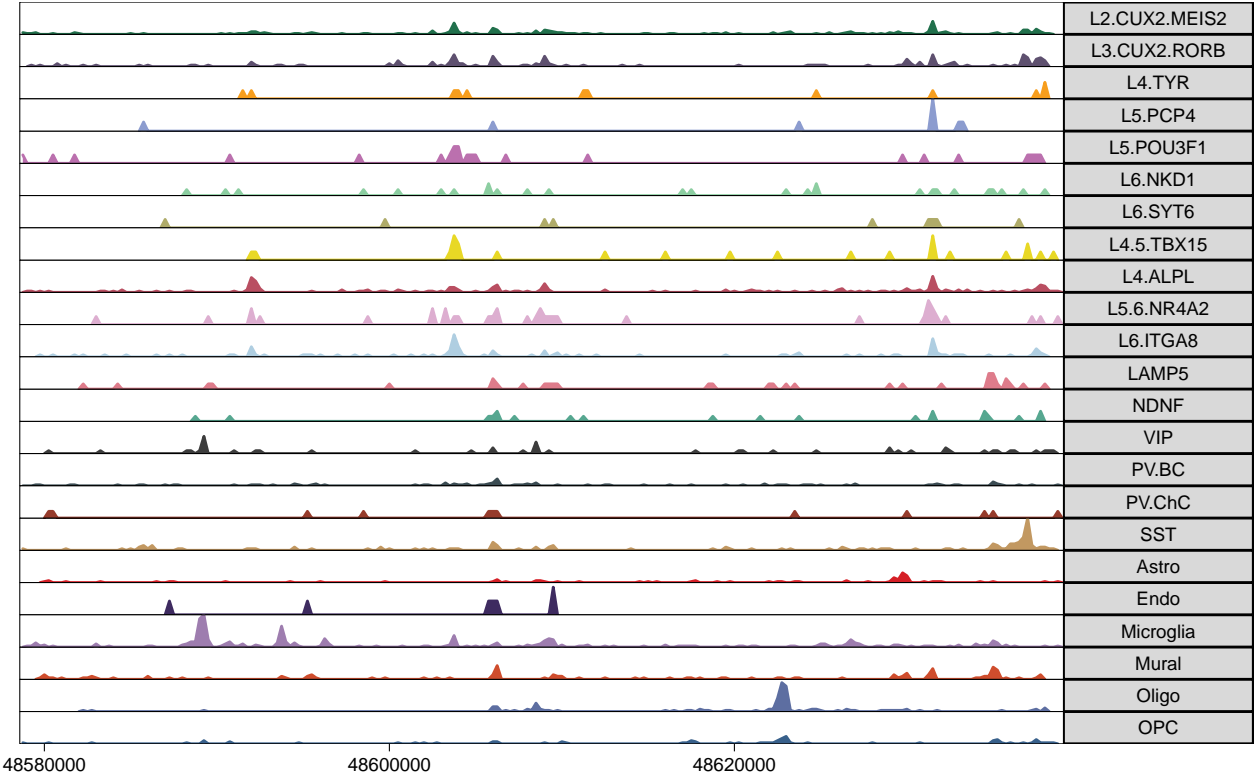


LoopTrack

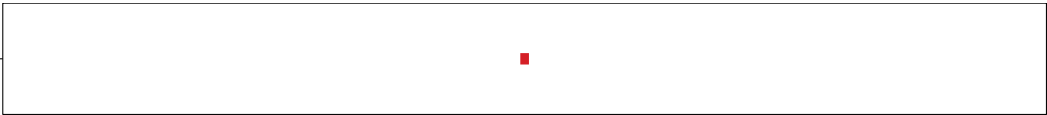


Genes

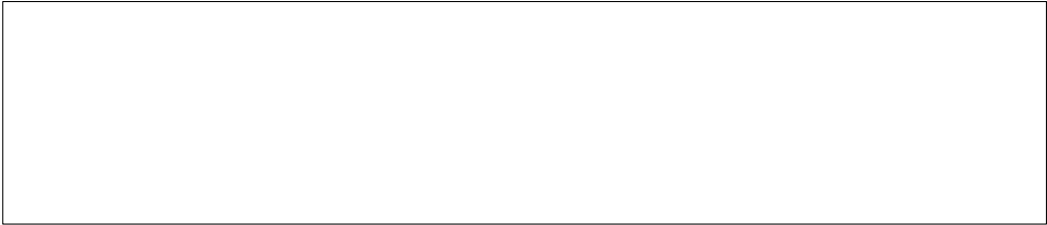
Coverage
(Norm. ATAC Signal Range (0-0.91) by ReadsInTSS)



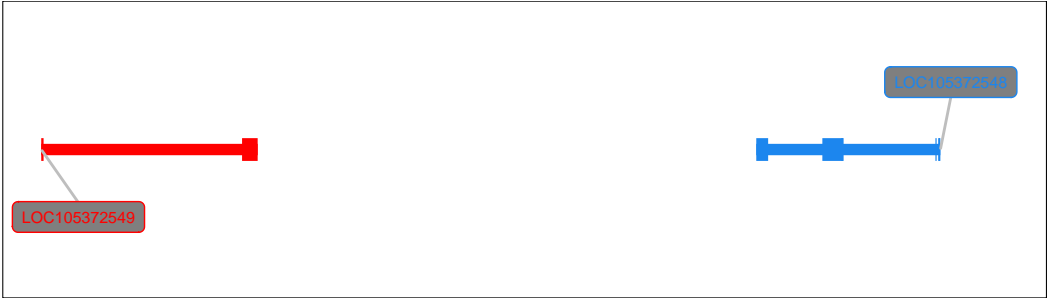
L5.6.NR4A2



Peaks

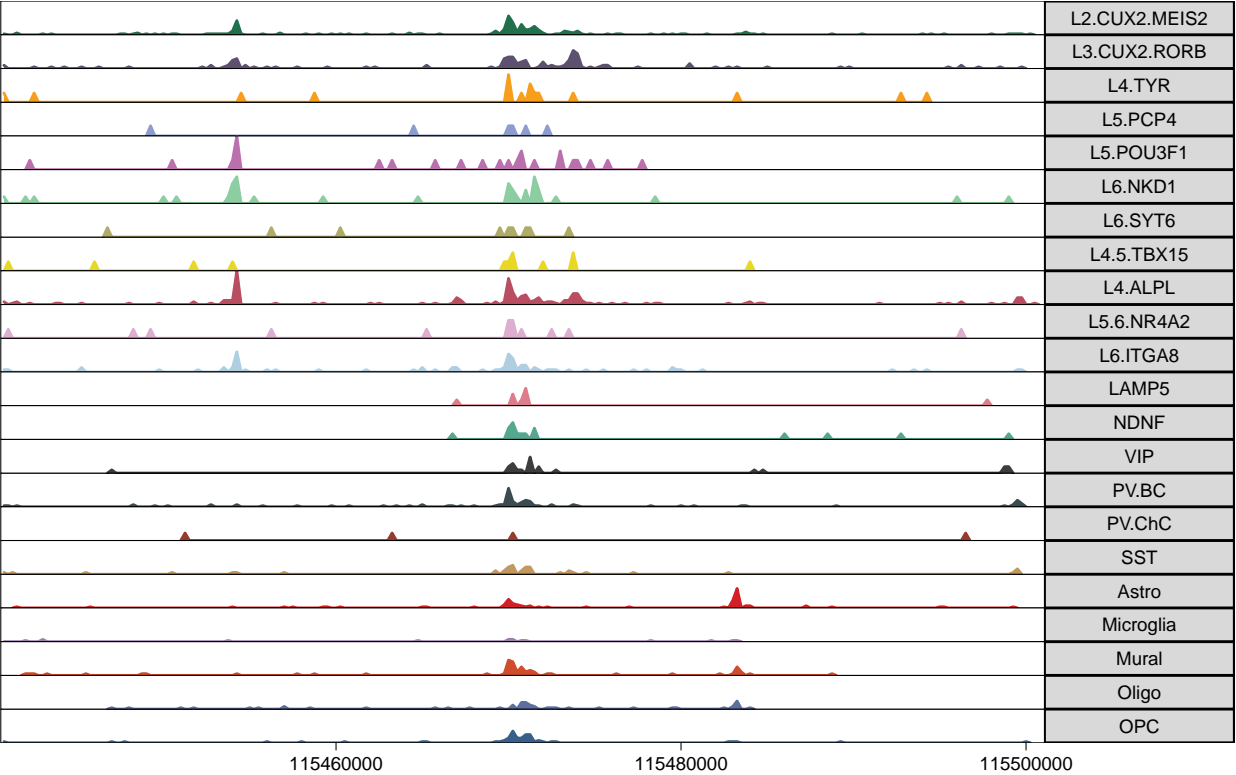


LoopTrack

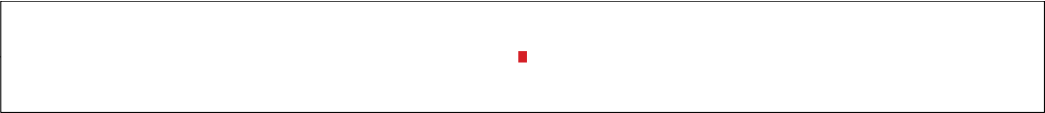


Genes

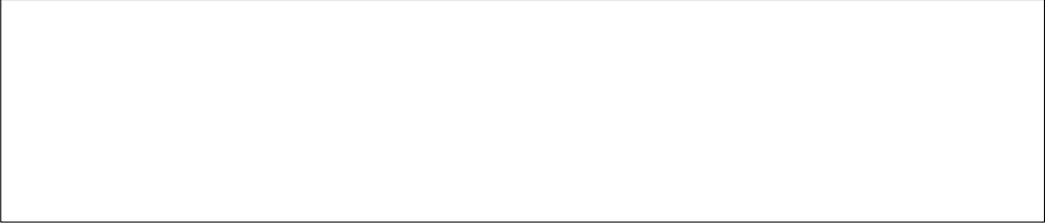
Coverage
(Norm. ATAC Signal Range 0–0.86) by ReadsInTSS)



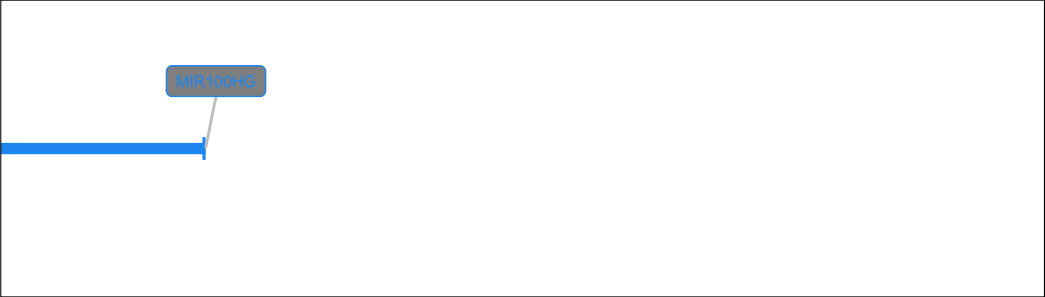
L5.6.NR4A2



Peaks

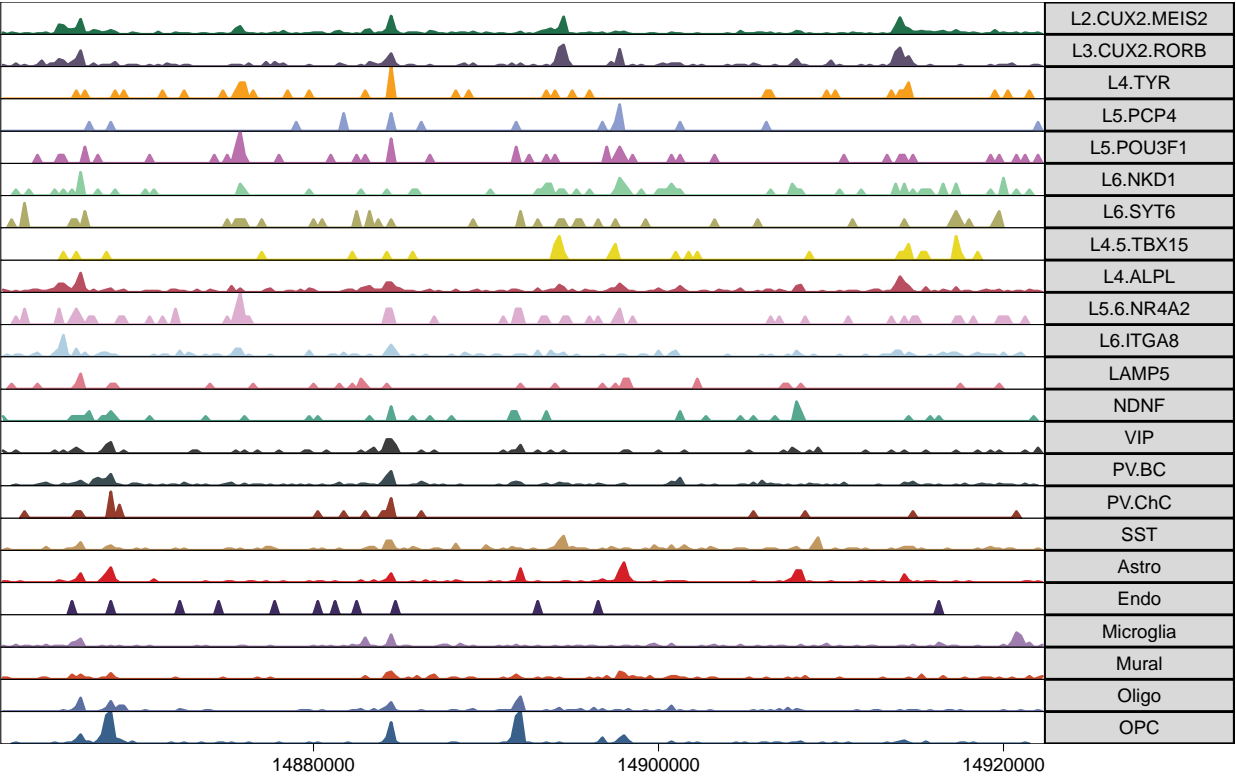


LoopTrack

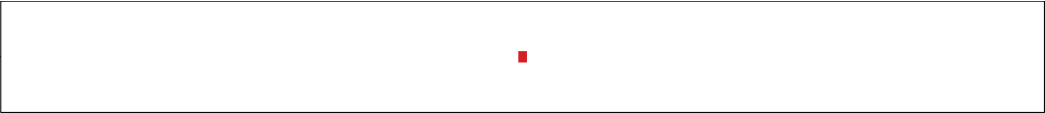


Genes

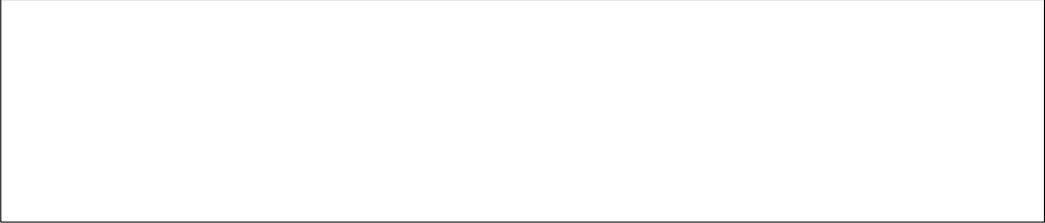
Coverage
(Norm. ATAC Signal Range (0–0.94) by ReadsInTSS)



L5.6.NR4A2



Peaks

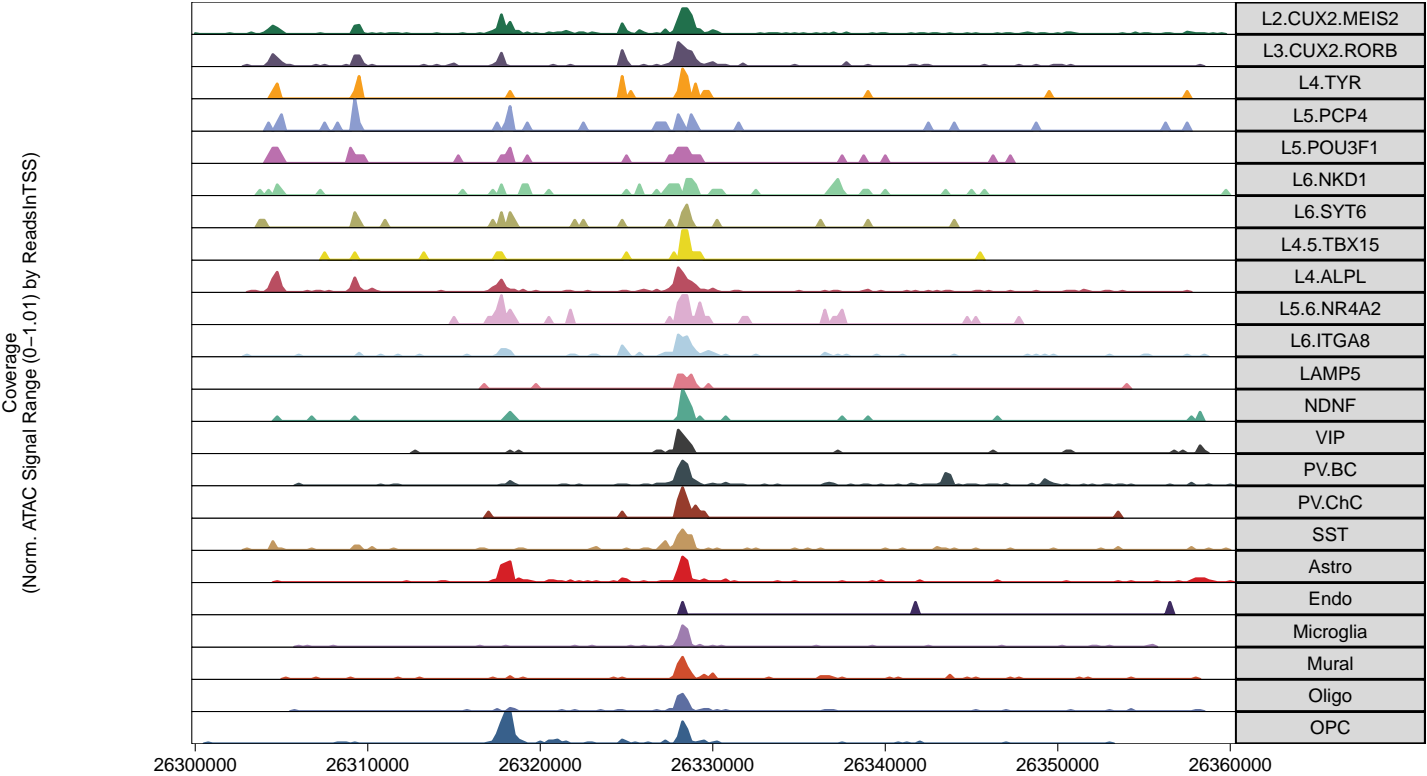


LoopTrack

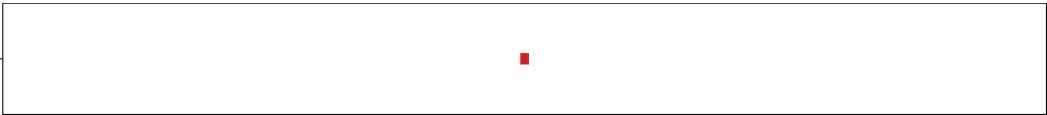


Genes

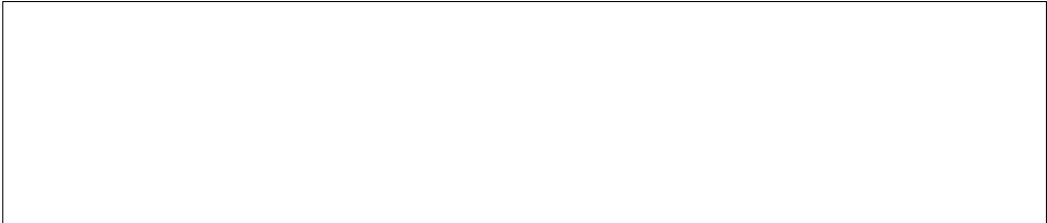
chr4:26299799–26360300 Marker Peak Tracks for: L5.6.NR4A2



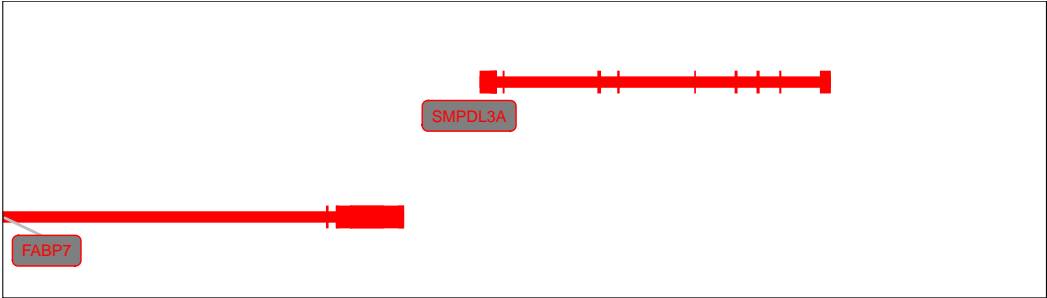
L5.6.NR4A2



Peaks

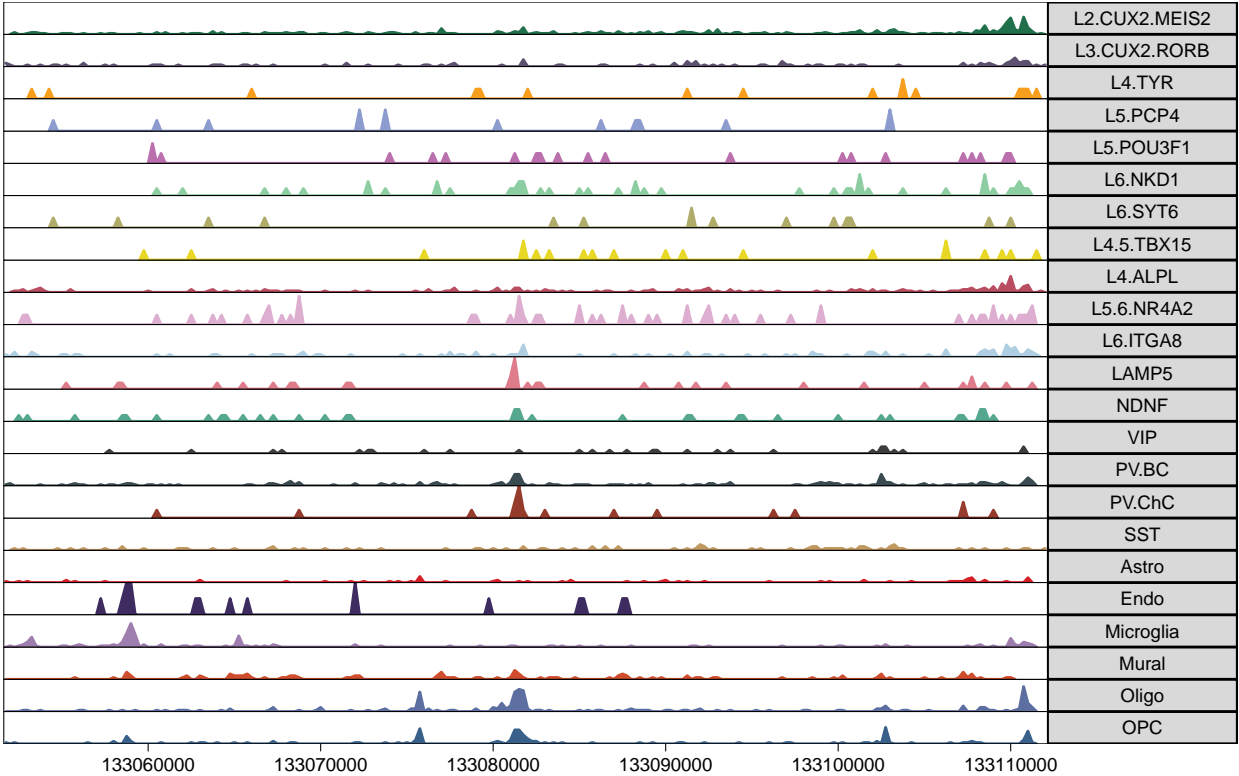


LoopTrack

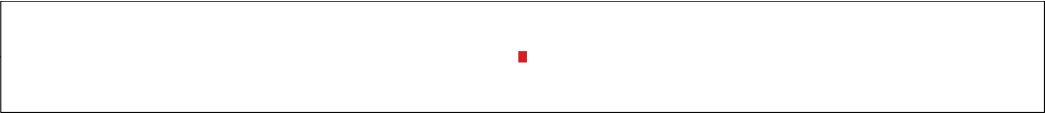


Genes

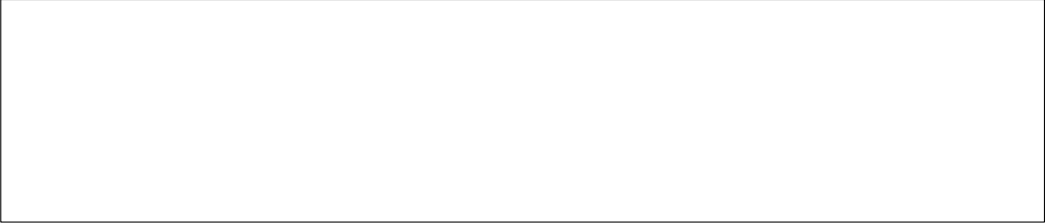
Coverage
(Norm. ATAC Signal Range (0–0.77) by ReadsInTSS)



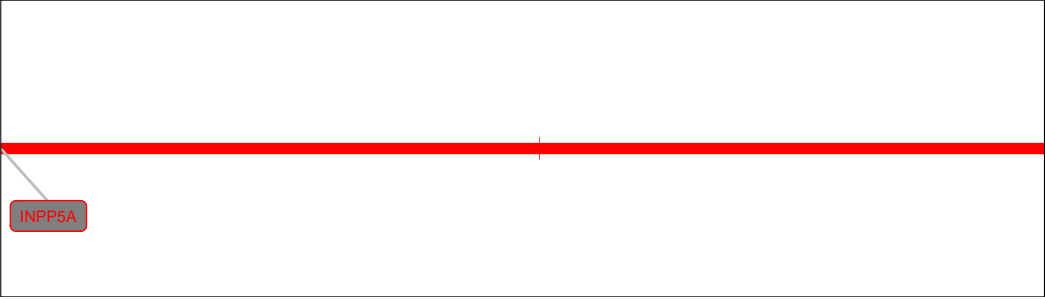
L5.6.NR4A2



Peaks



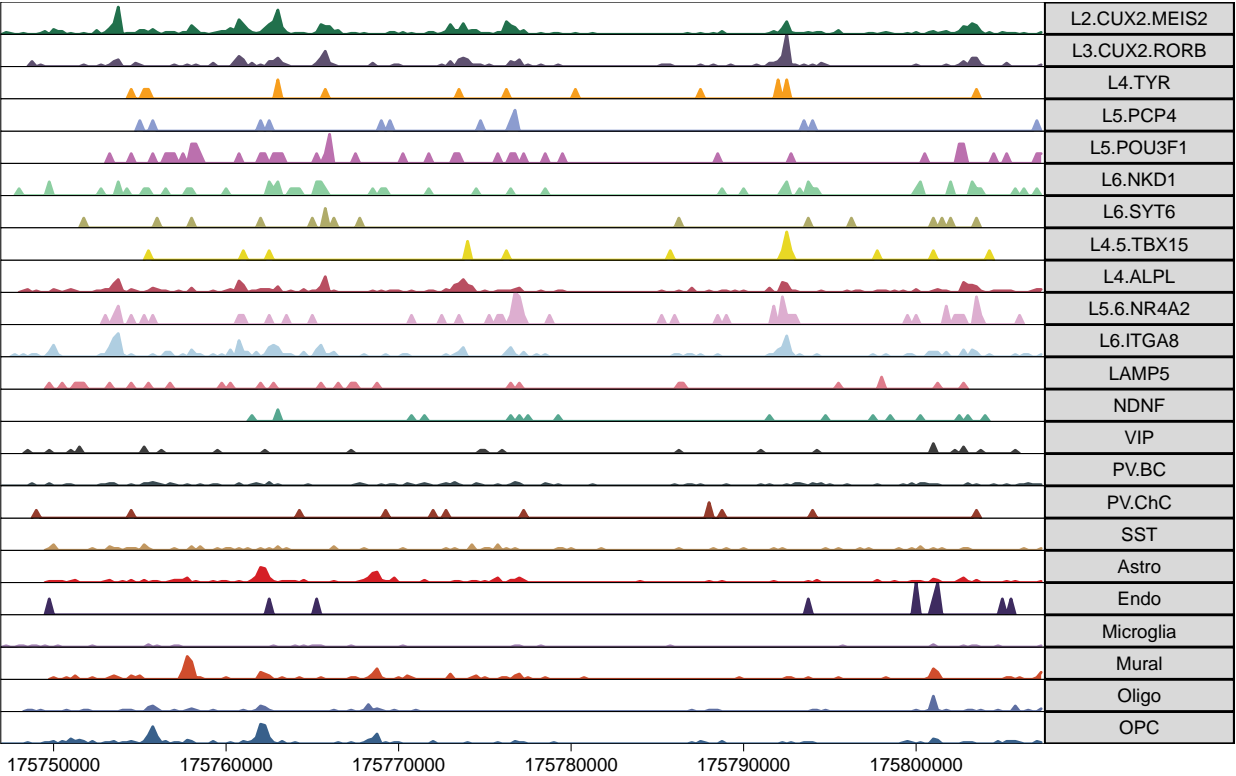
LoopTrack



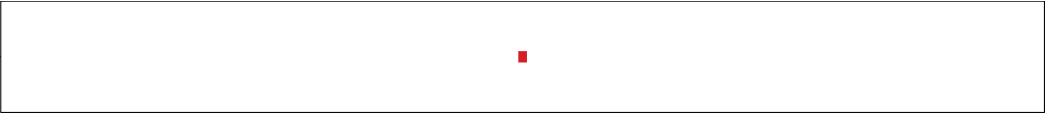
Genes

chr6:175746939–175807440 Marker Peak Tracks for: L5.6.NR4A2

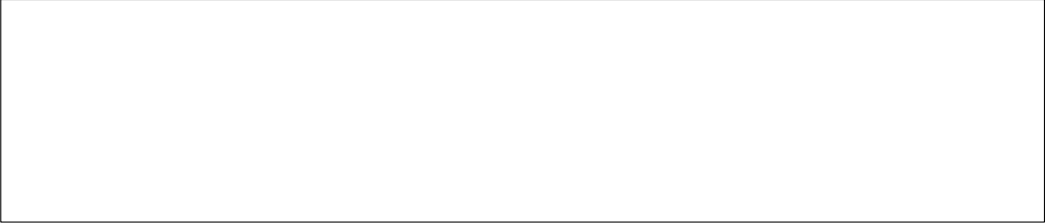
Coverage
(Norm. ATAC Signal Range (0–0.79) by ReadsInTSS)



L5.6.NR4A2



Peaks

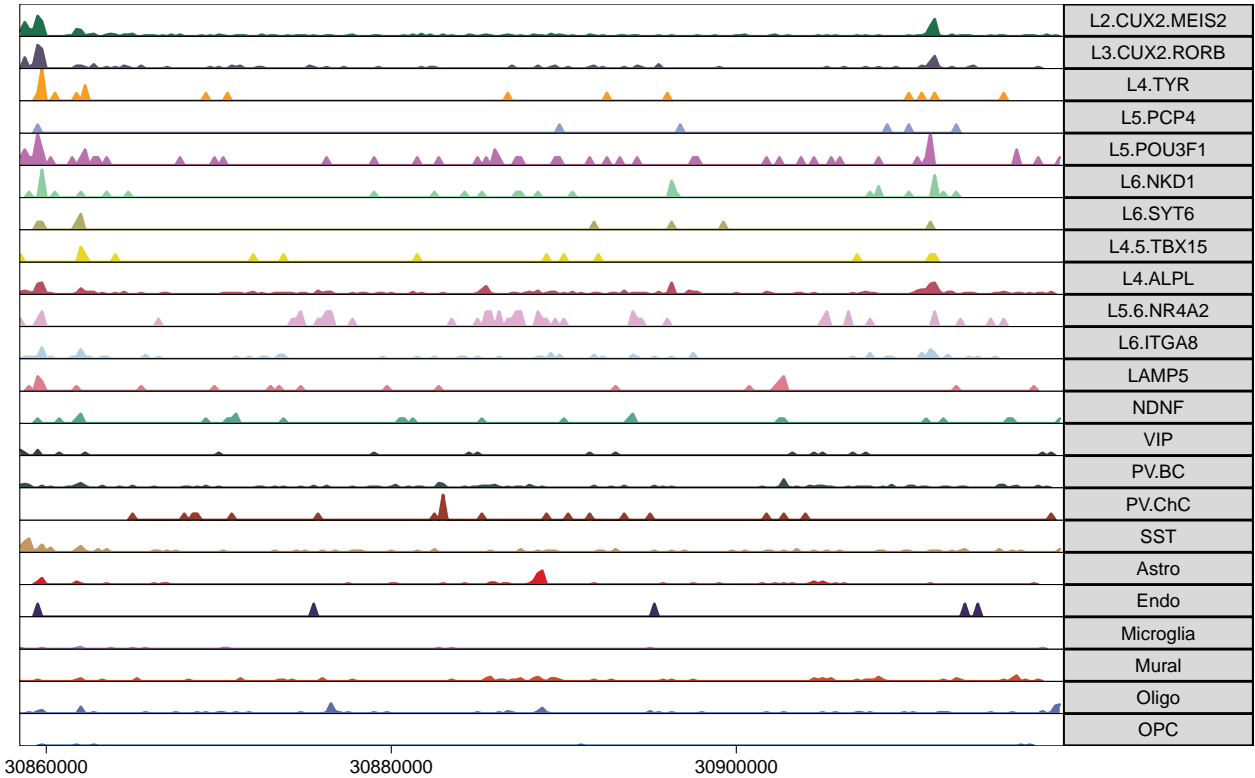


LoopTrack

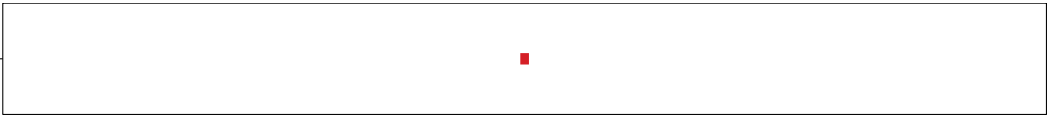


Genes

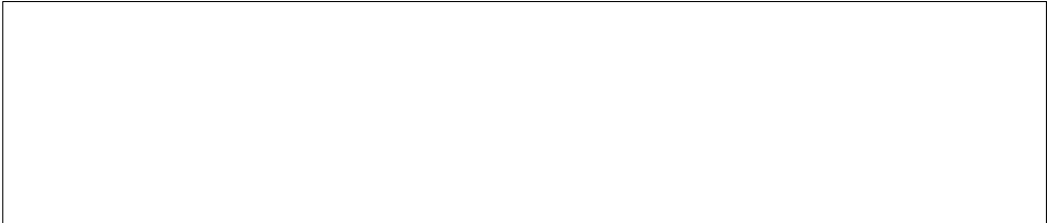
Coverage
(Norm. ATAC Signal Range (0–0.98) by ReadsInTSS)



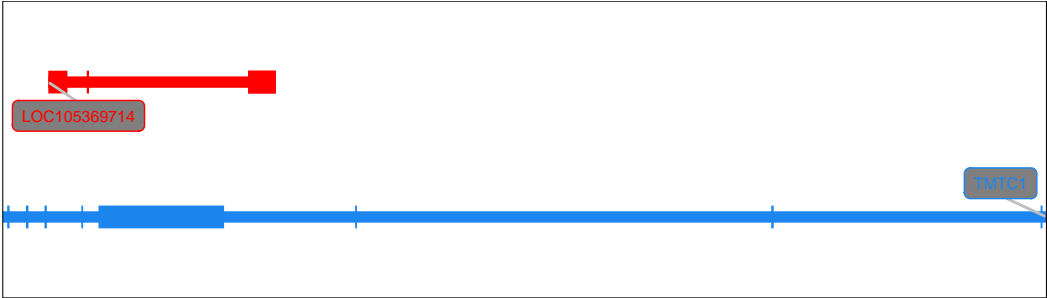
L5.6.NR4A2



Peaks

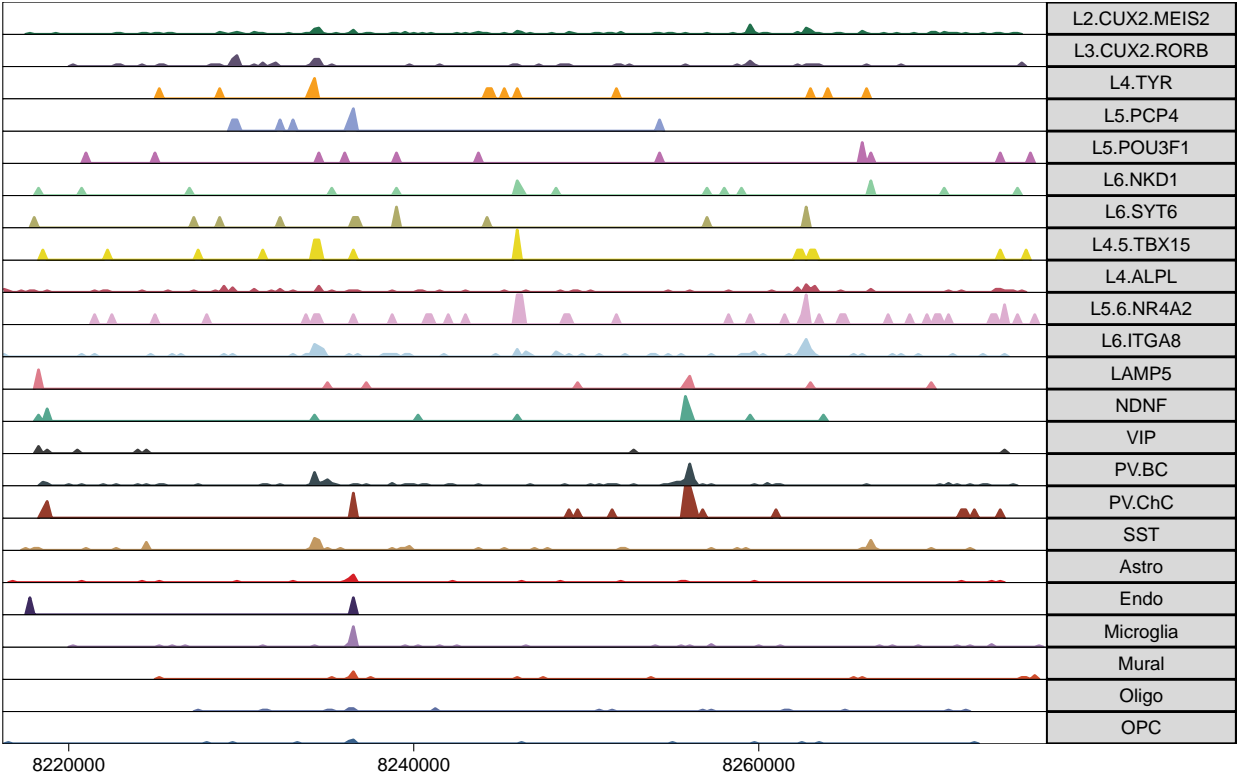


LoopTrack

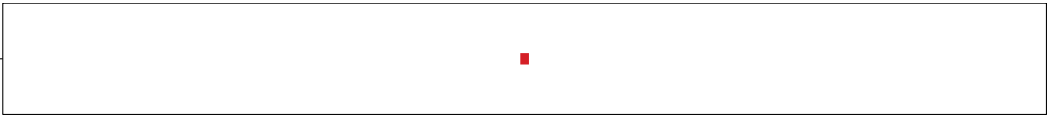


Genes

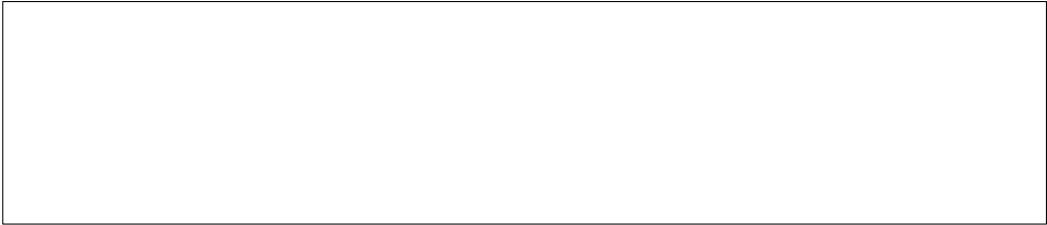
Coverage
(Norm. ATAC Signal Range (0-0.74) by ReadsInTSS)



L5.6.NR4A2



Peaks

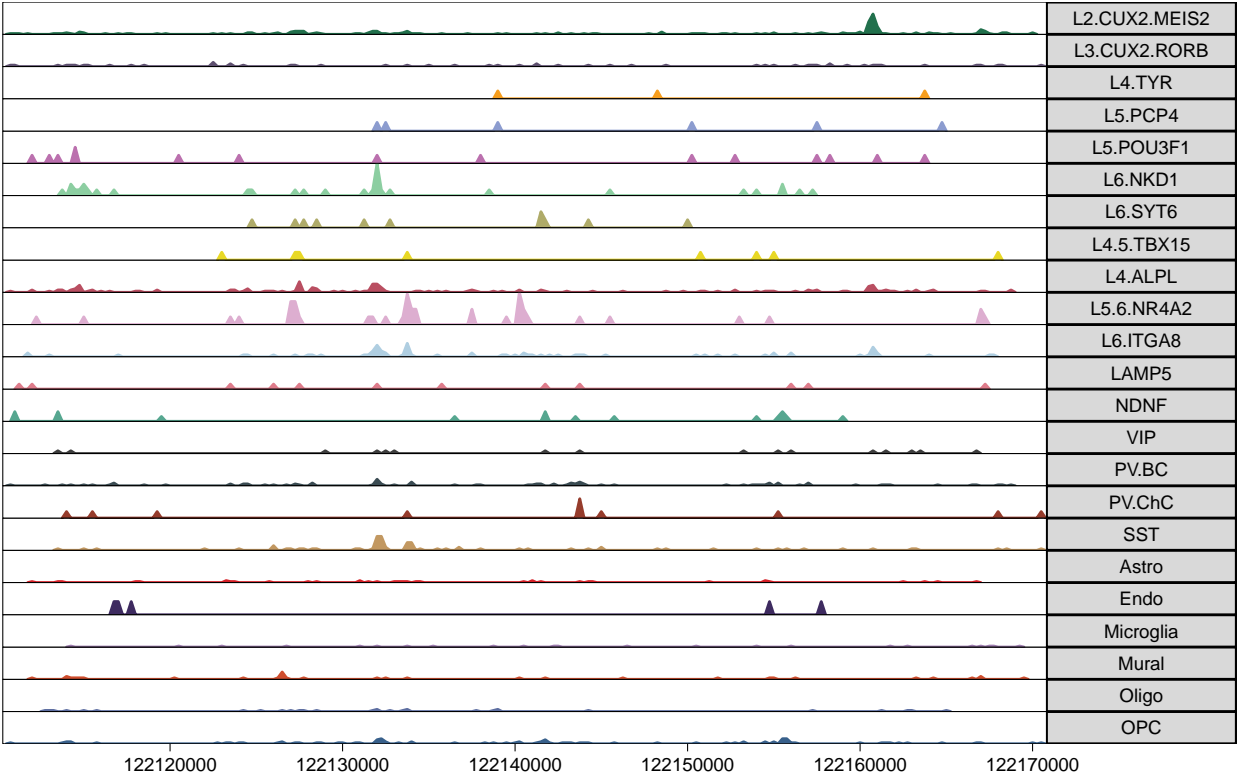


LoopTrack

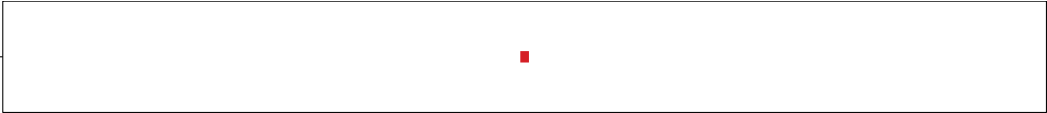


Genes

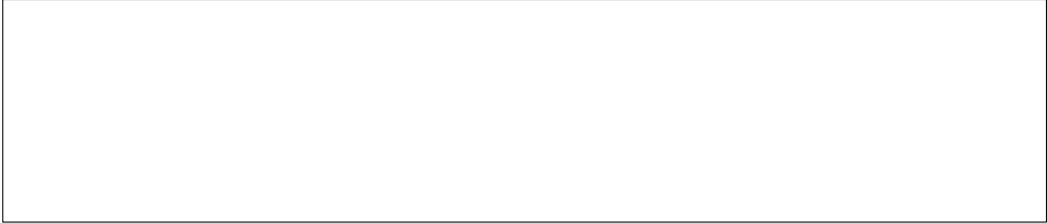
Coverage
(Norm. ATAC Signal Range (0–0.95) by ReadsInTSS)



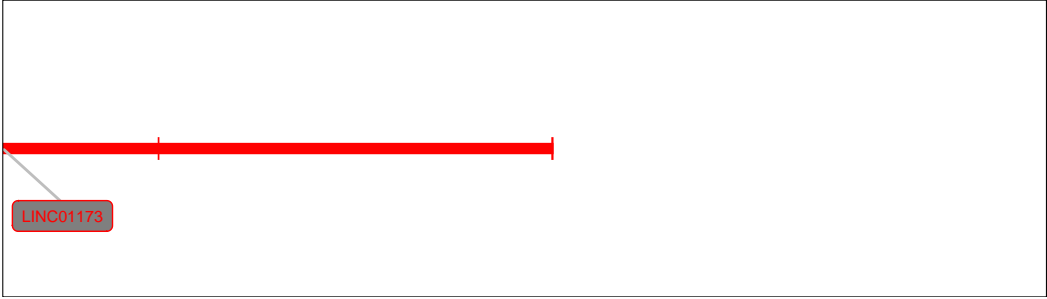
L5.6.NR4A2



Peaks



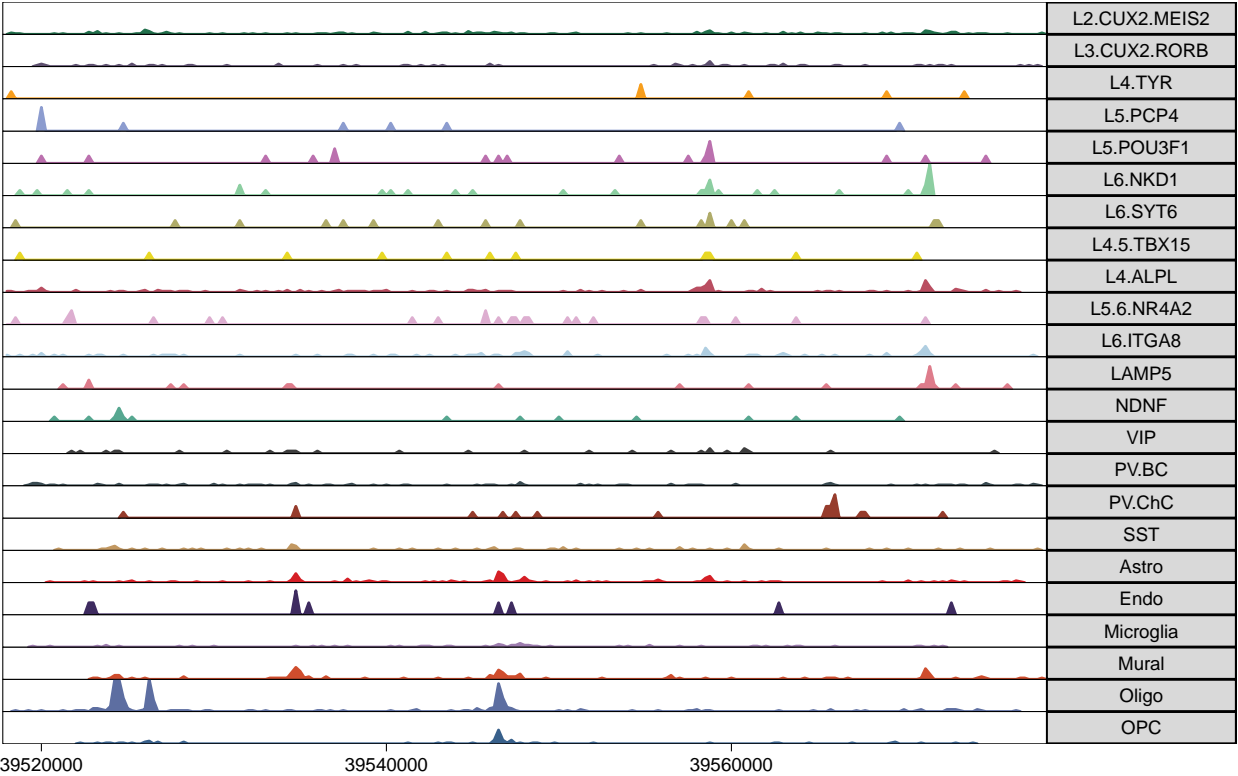
LoopTrack



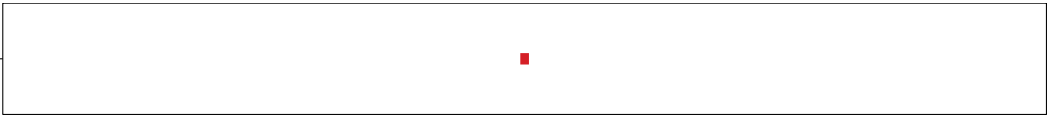
Genes

chr1:39517761-39578262 Marker Peak Tracks for: L5.6.NR4A2

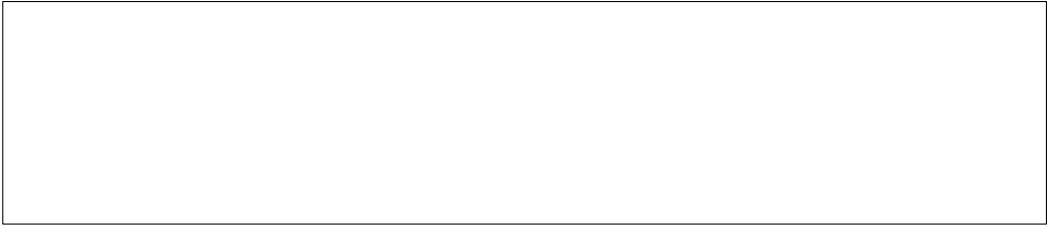
Coverage
(Norm. ATAC Signal Range 0-1.05) by ReadsInTSS)



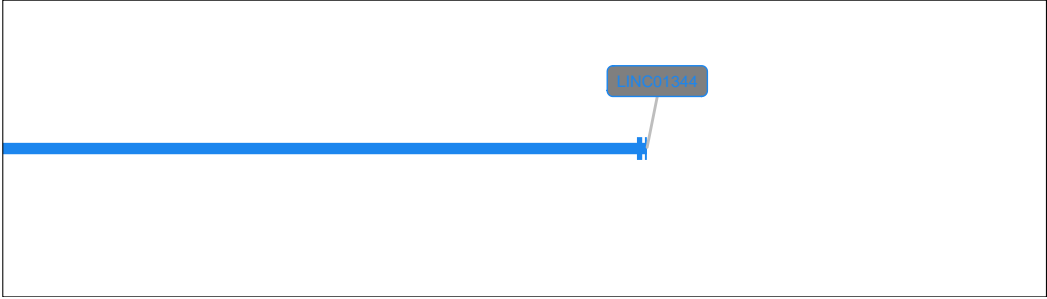
L5.6.NR4A2



Peaks



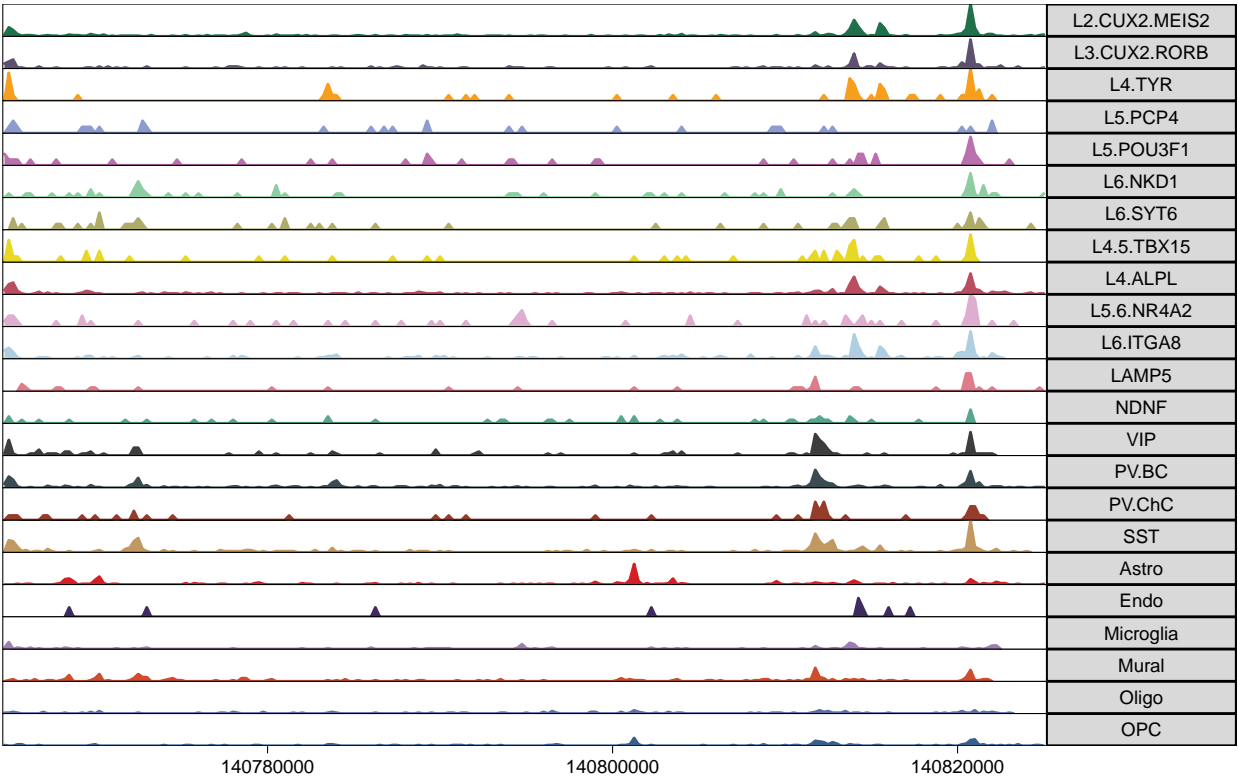
LoopTrack



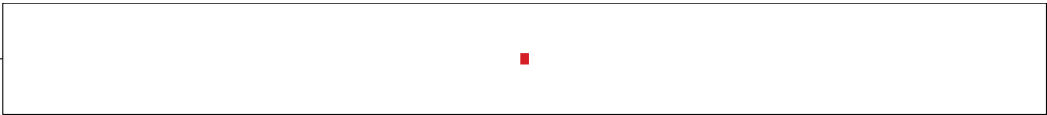
Genes

chr8:140764650–140825151 Marker Peak Tracks for: L5.6.NR4A2

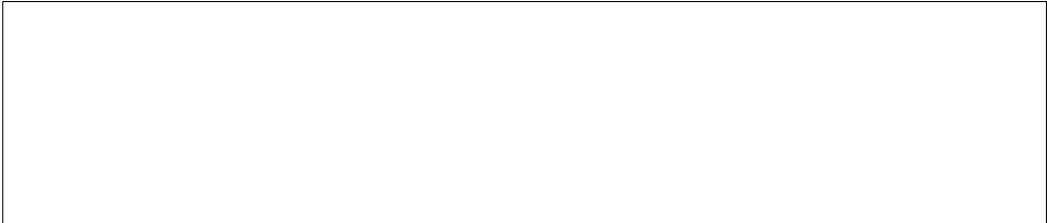
Coverage
(Norm. ATAC Signal Range 0–1.35) by ReadsInTSS)



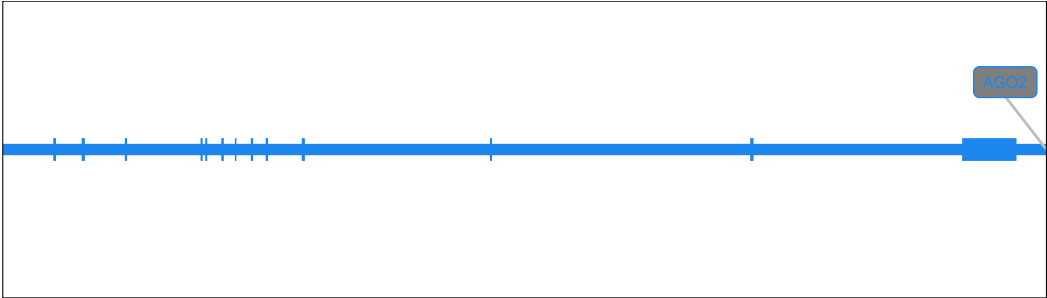
L5.6.NR4A2



Peaks

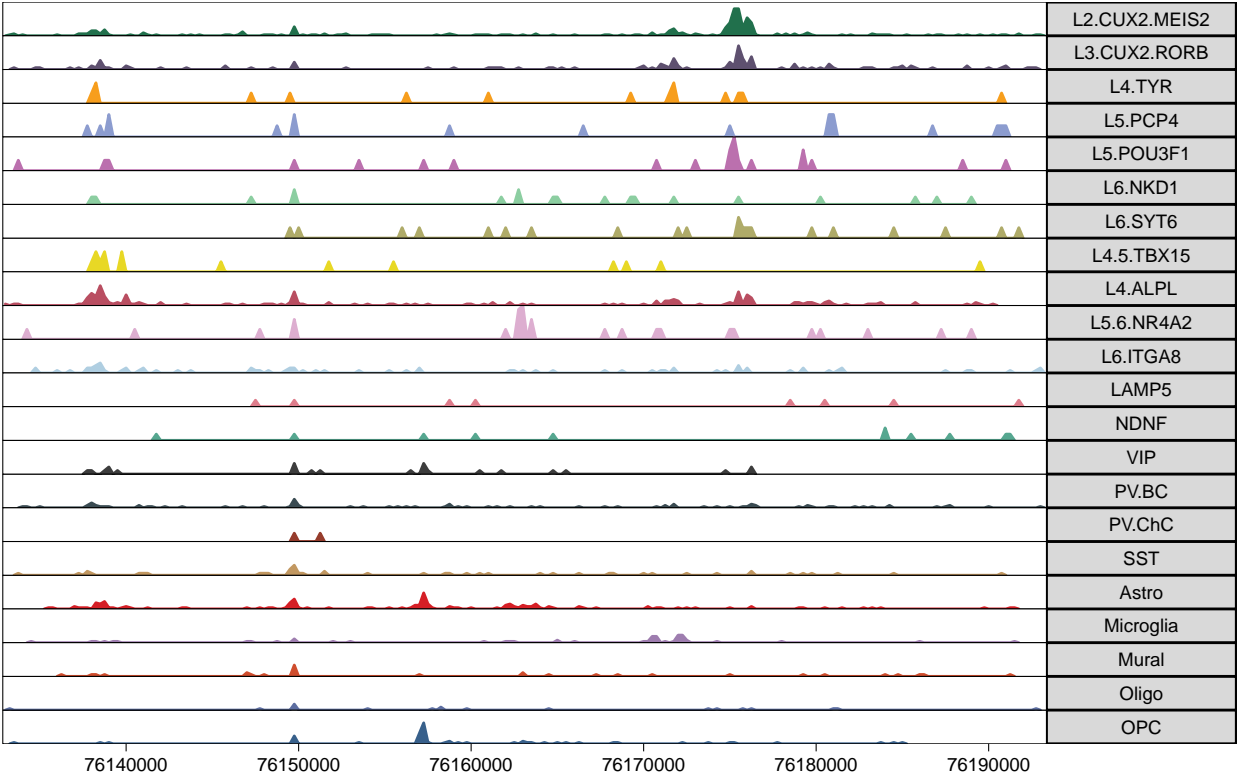


LoopTrack

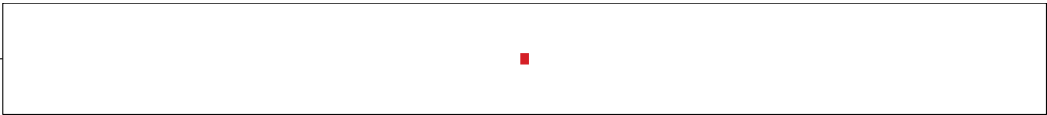


Genes

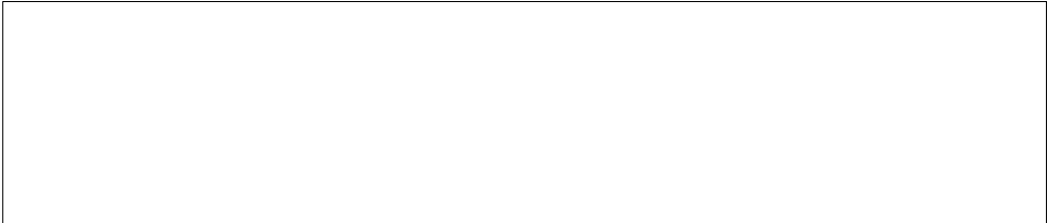
Coverage
(Norm. ATAC Signal Range (0-0.77) by ReadsInTSS)



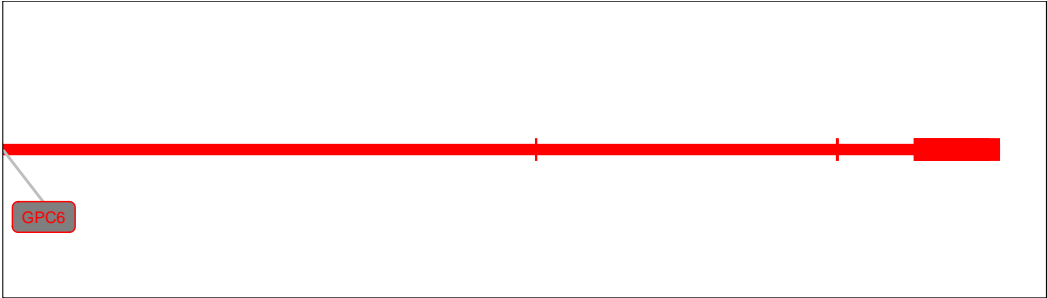
L5.6.NR4A2



Peaks

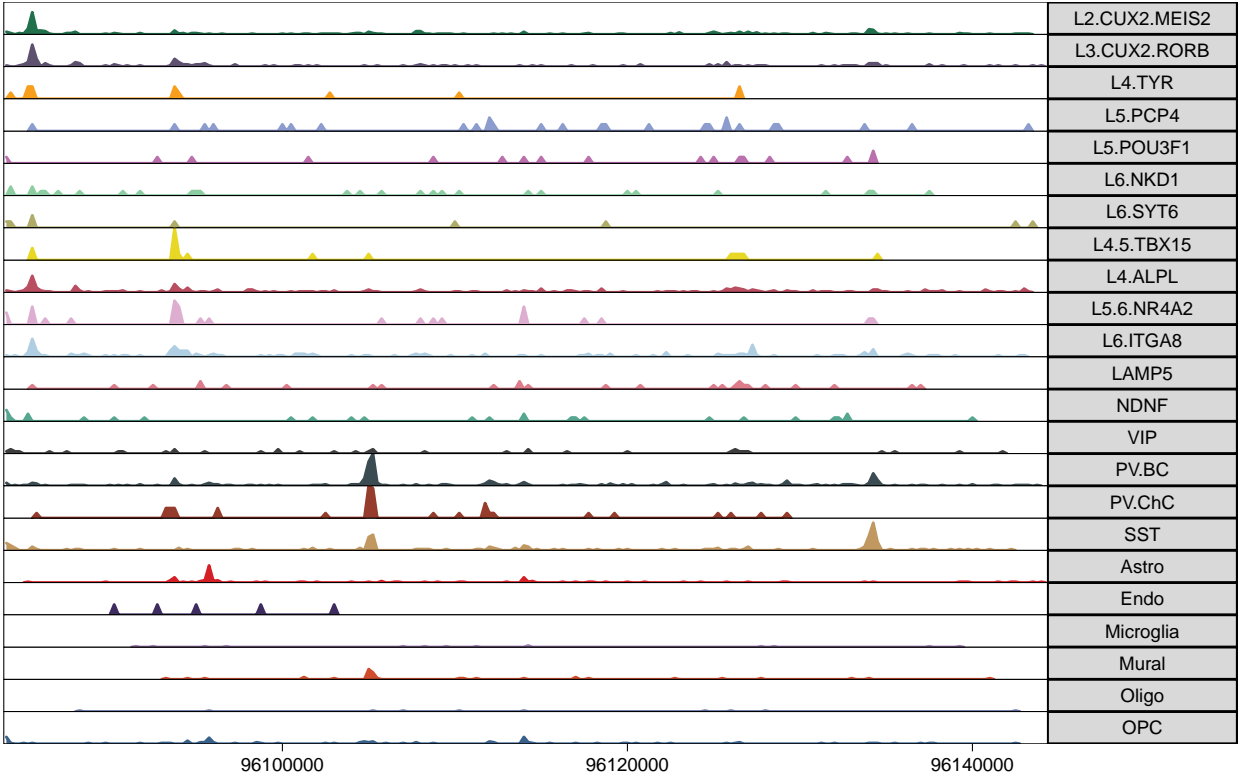


LoopTrack

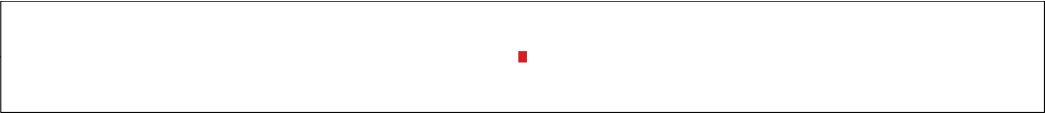


Genes

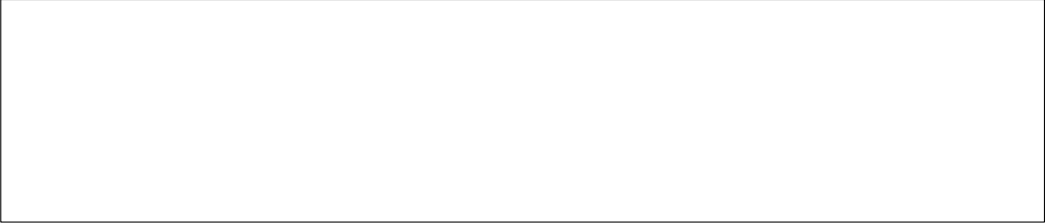
Coverage
(Norm. ATAC Signal Range 0–1.23) by ReadsInTSS)



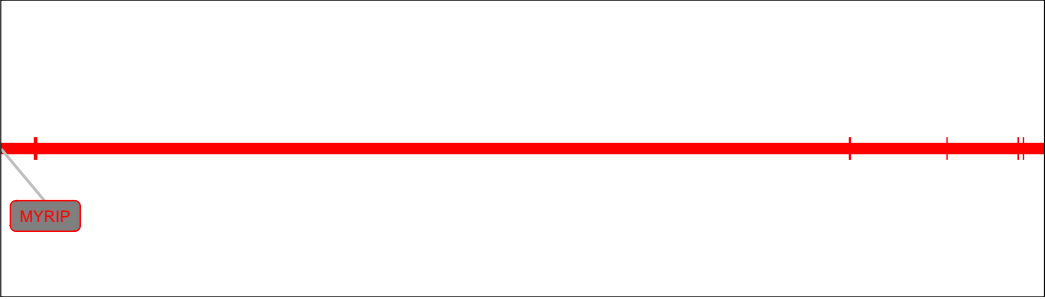
L5.6.NR4A2



Peaks



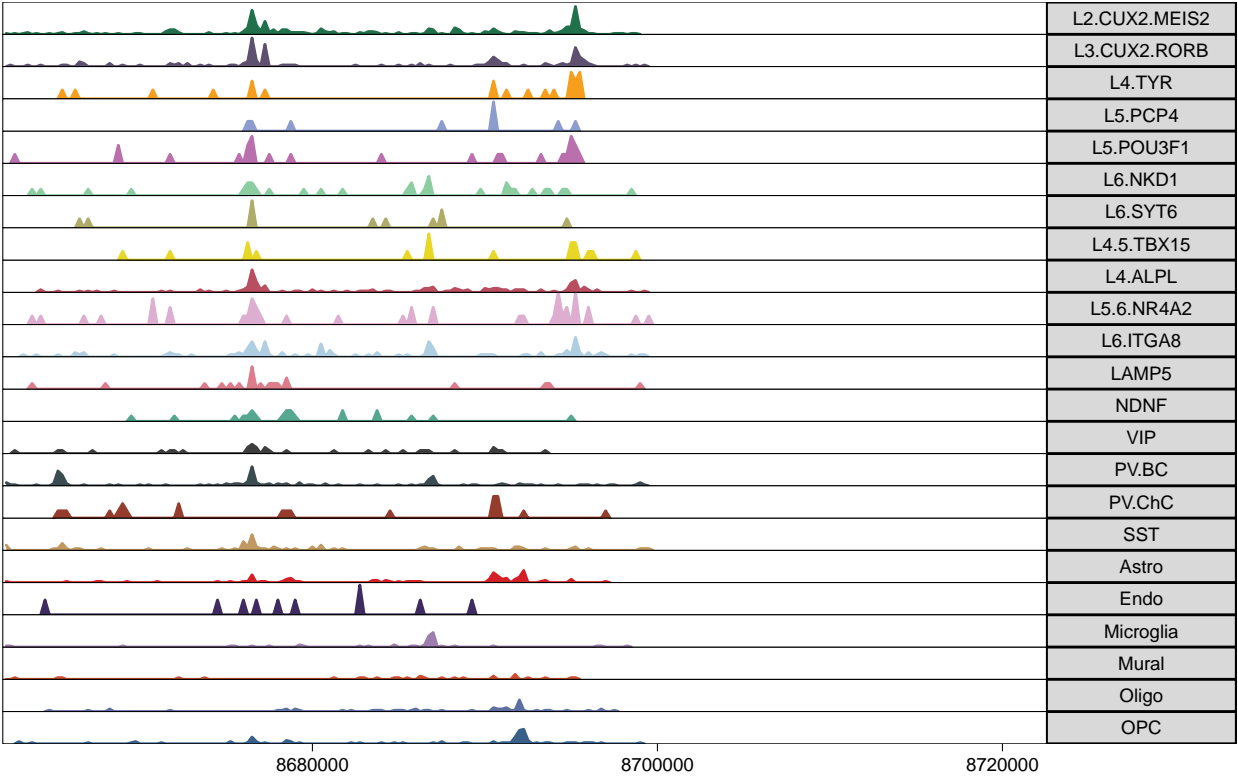
LoopTrack



Genes

chr17:8662051–8722552 Marker Peak Tracks for: L5.6.NR4A2

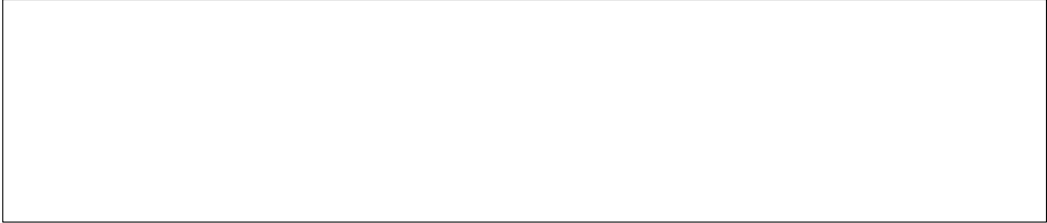
Coverage
(Norm. ATAC Signal Range (0–0.85) by ReadsInTSS)



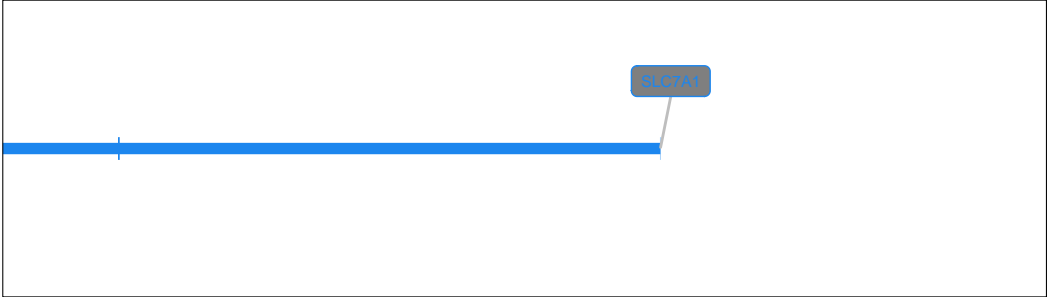
L5.6.NR4A2



Peaks



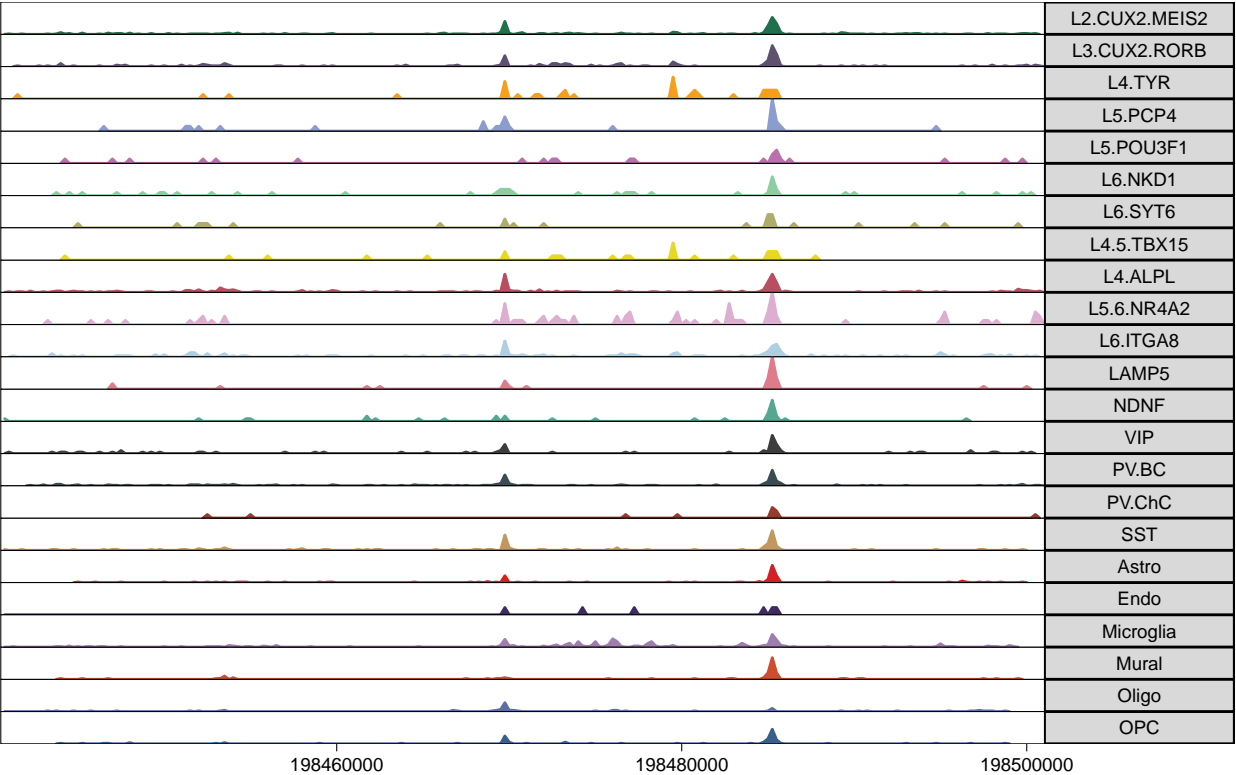
LoopTrack



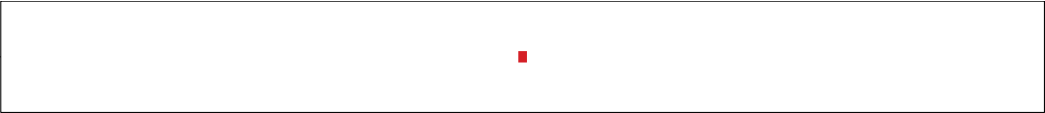
Genes

chr1:198440527-198501028 Marker Peak Tracks for: L5.6.NR4A2

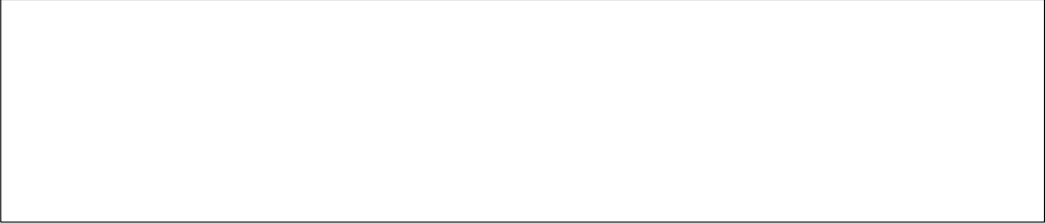
Coverage
(Norm. ATAC Signal Range 0-1.73) by ReadsInTSS)



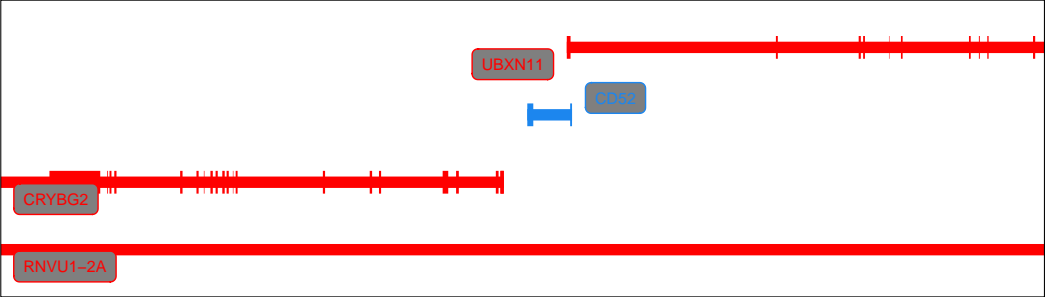
L5.6.NR4A2



Peaks

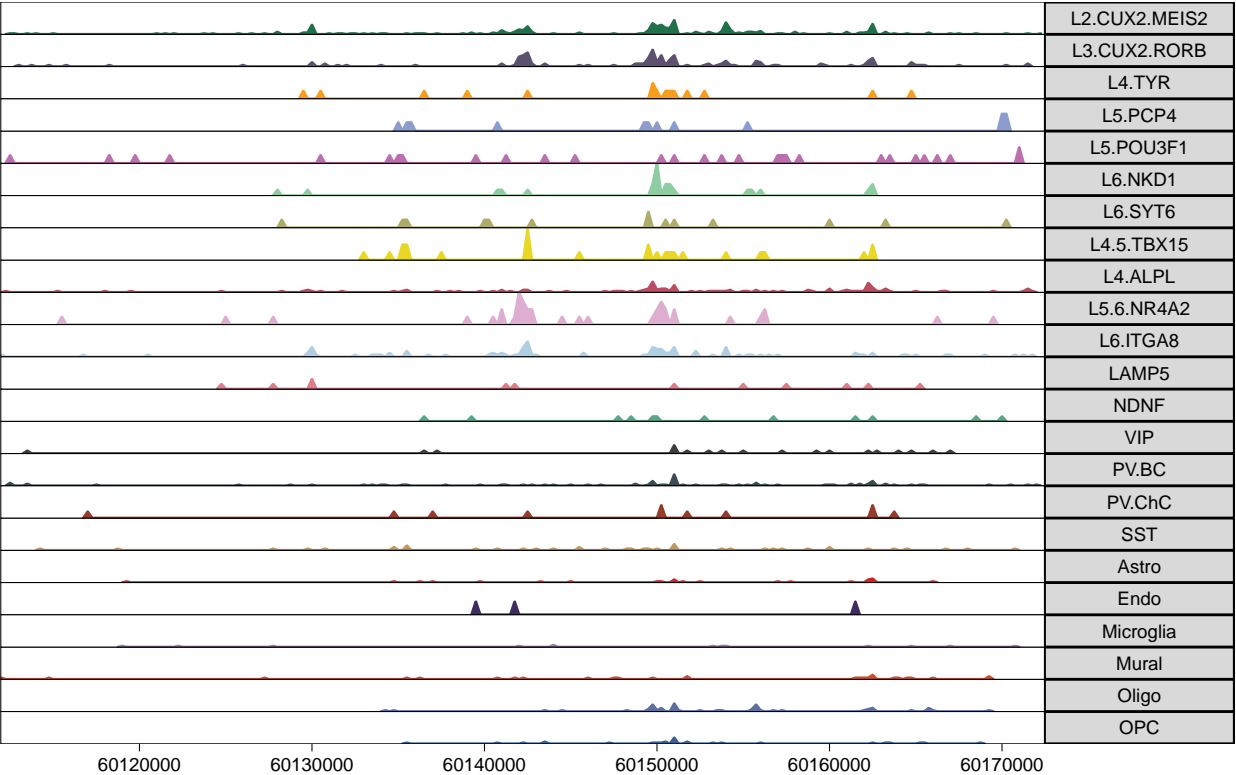


LoopTrack

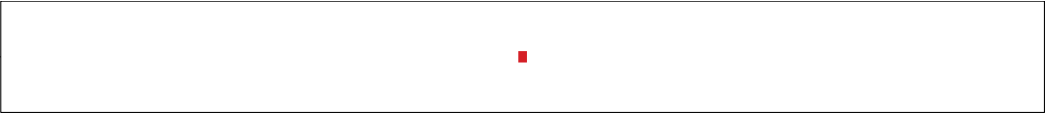


Genes

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



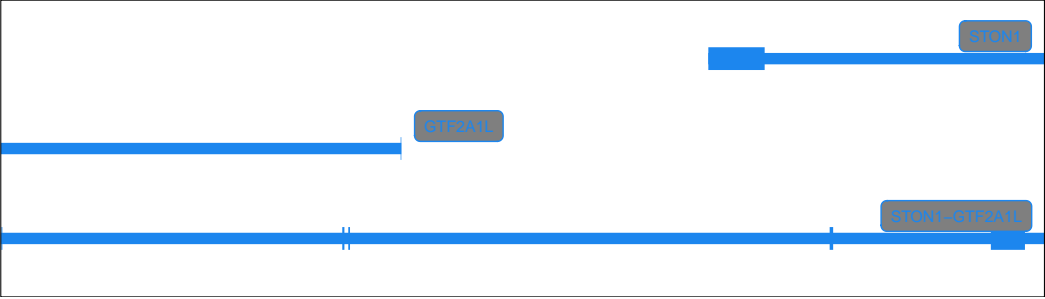
L5.6.NR4A2



Peaks



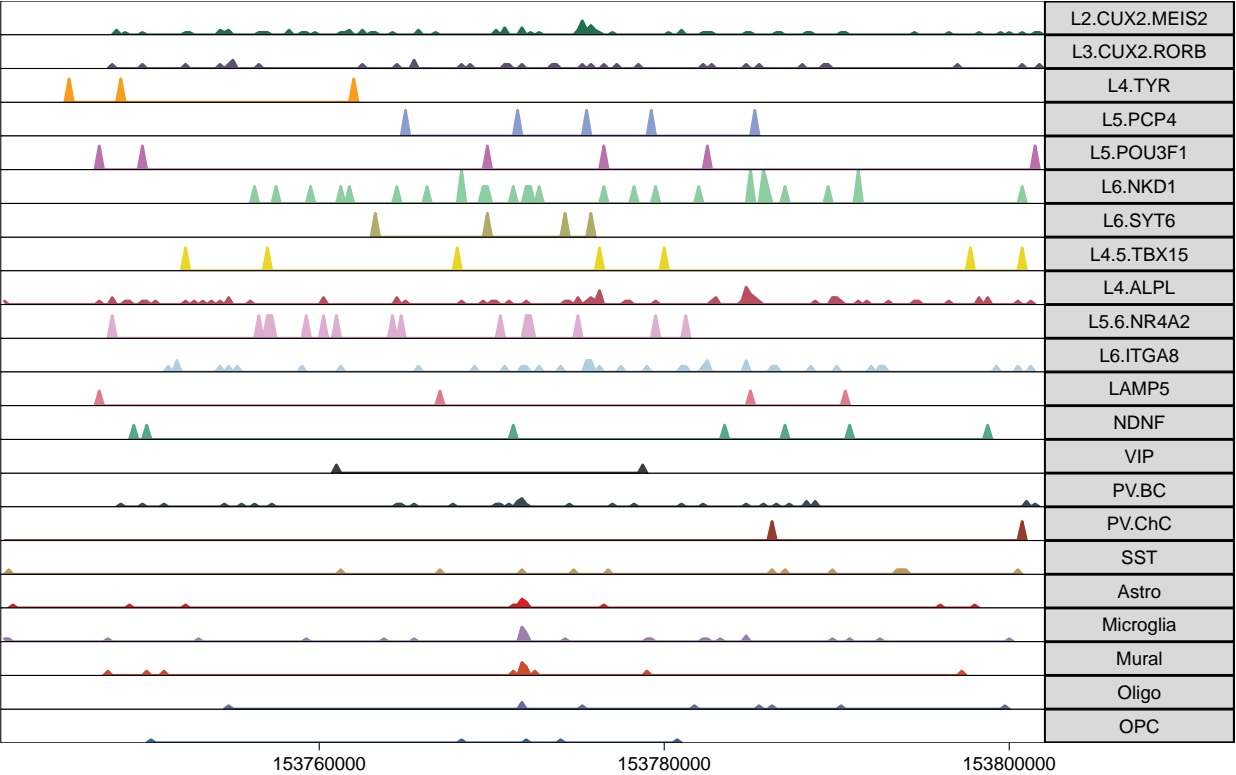
LoopTrack



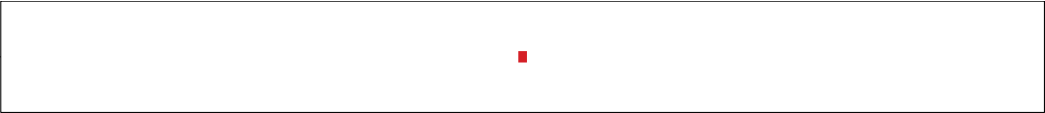
Genes

chr1:153741540-153802041 Marker Peak Tracks for: L5.6.NR4A2

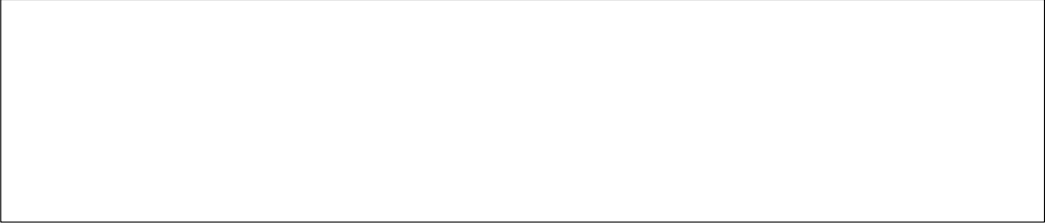
Coverage
(Norm. ATAC Signal Range (0-0.34) by ReadsInTSS)



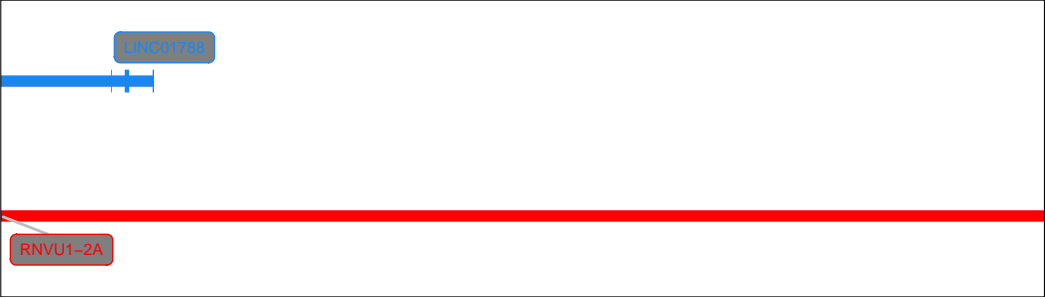
L5.6.NR4A2



Peaks

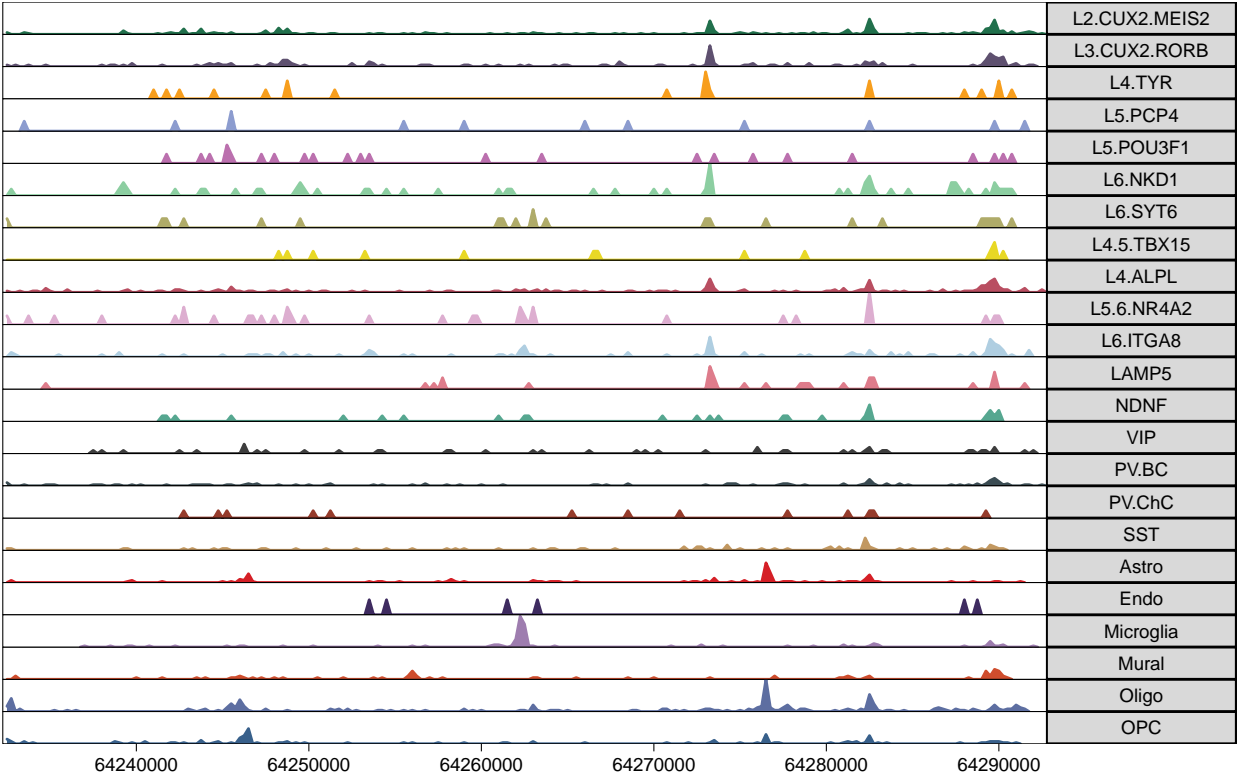


LoopTrack

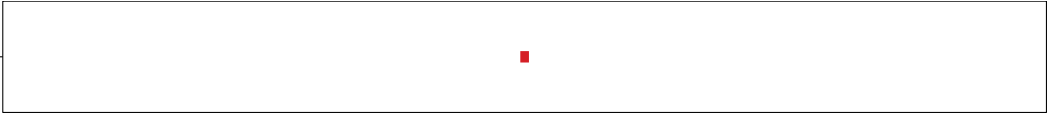


Genes

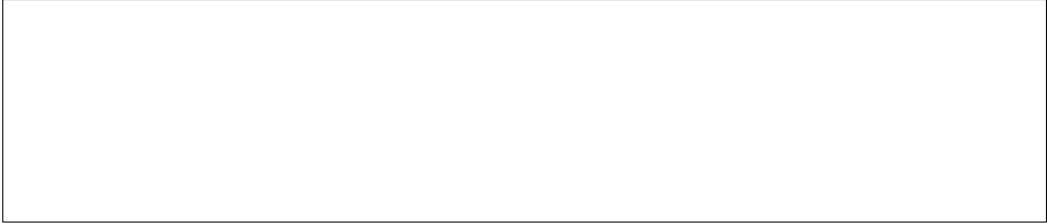
Coverage
(Norm. ATAC Signal Range 0–0.84) by ReadsInTSS)



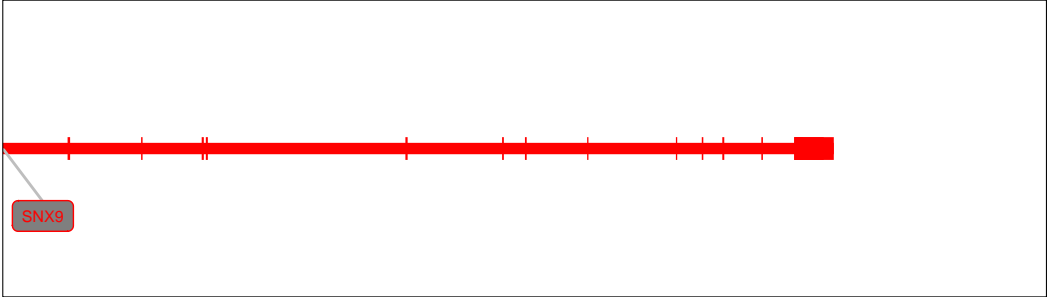
L5.6.NR4A2



Peaks

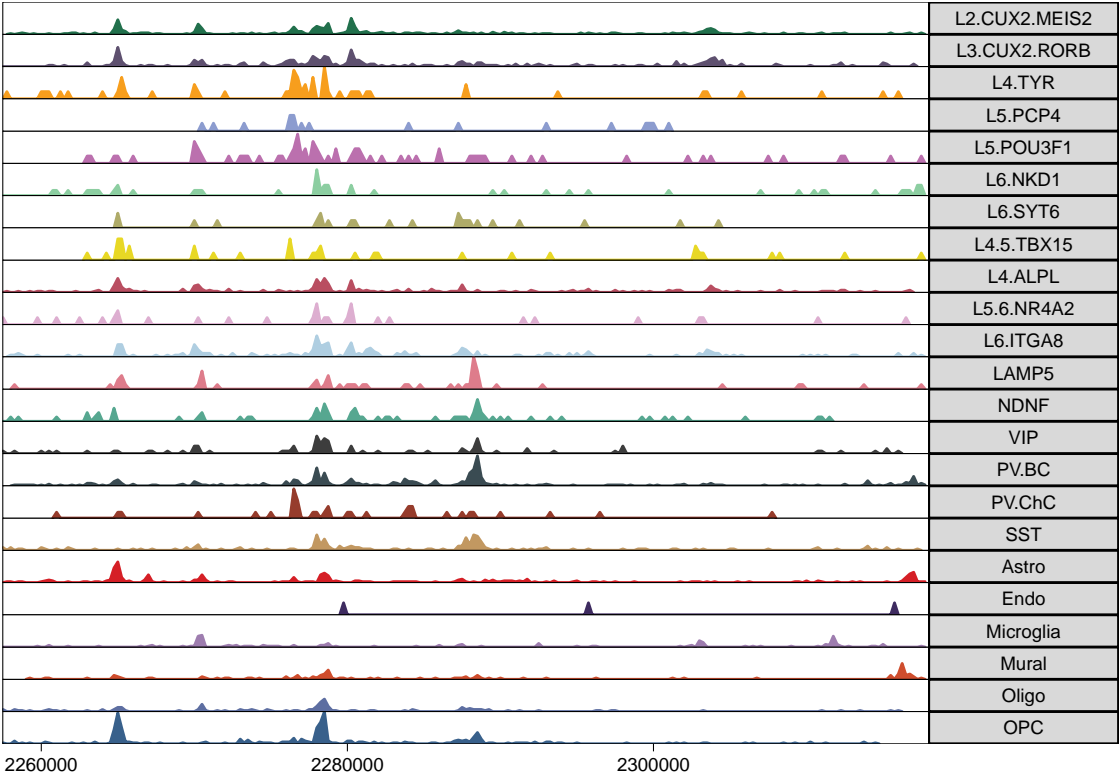


LoopTrack



Genes

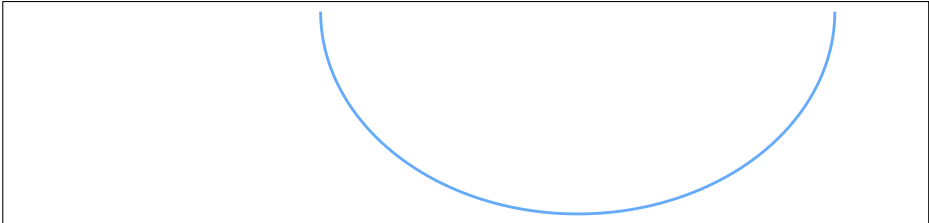
Coverage
(Norm. ATAC Signal Range 0–1.05) by ReadsInTSS)



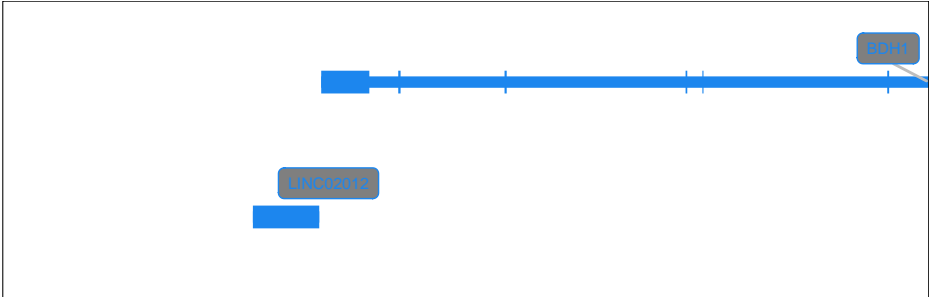
L5.6.NR4A2



Peaks

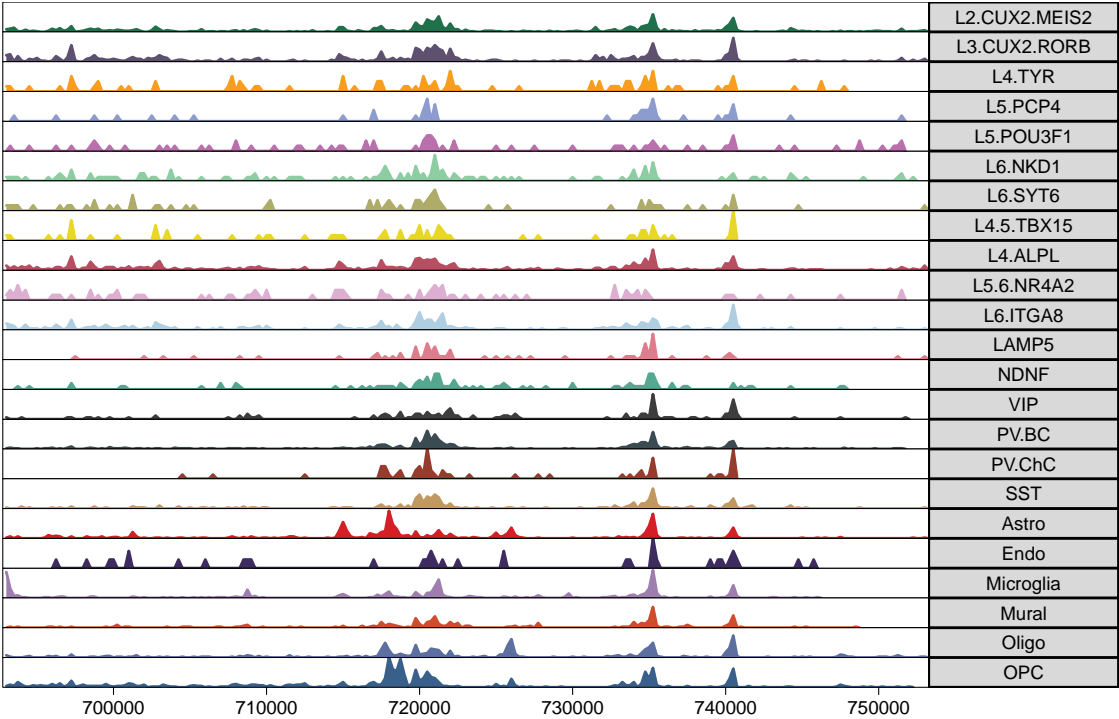


Peak2GeneLinks



Genes

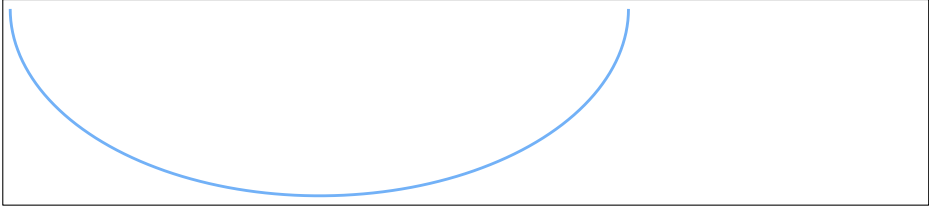
Coverage
(Norm. ATAC Signal Range (0–1.37) by ReadslnTSS)



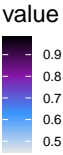
L5.6.NR4A2



Peaks



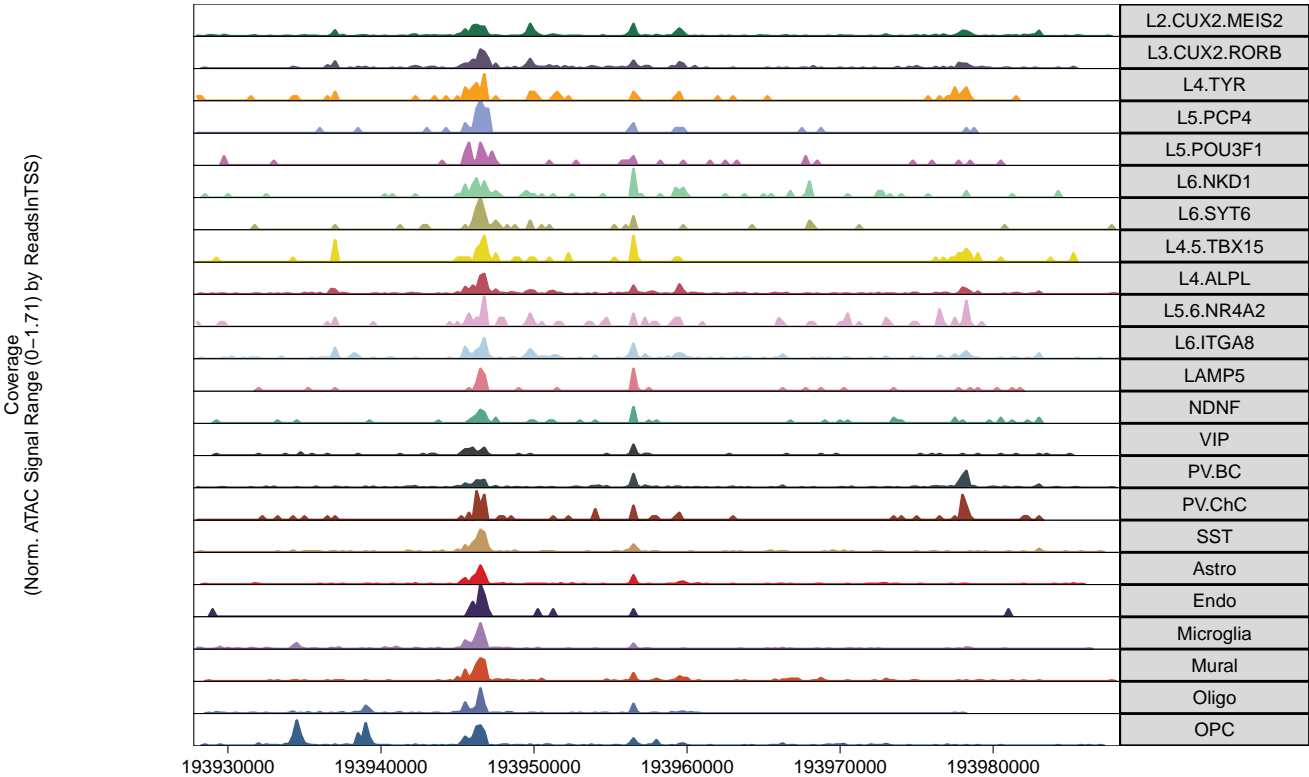
Peak2GeneLinks



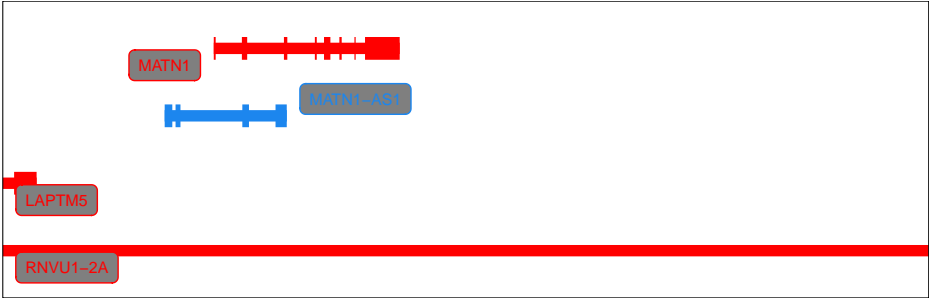
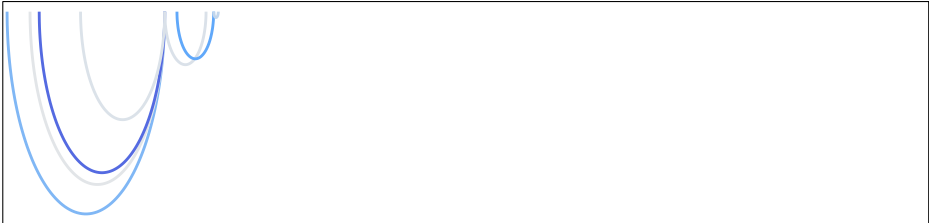
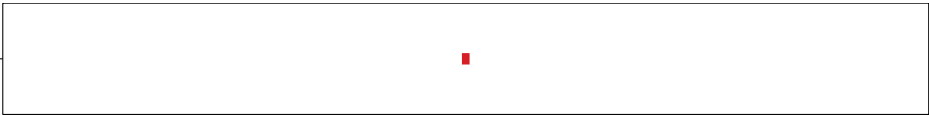
GeneTrack

strand ● ·

chr1:193927769–193988270 Marker Peak Tracks for: L5.6.NR4A2

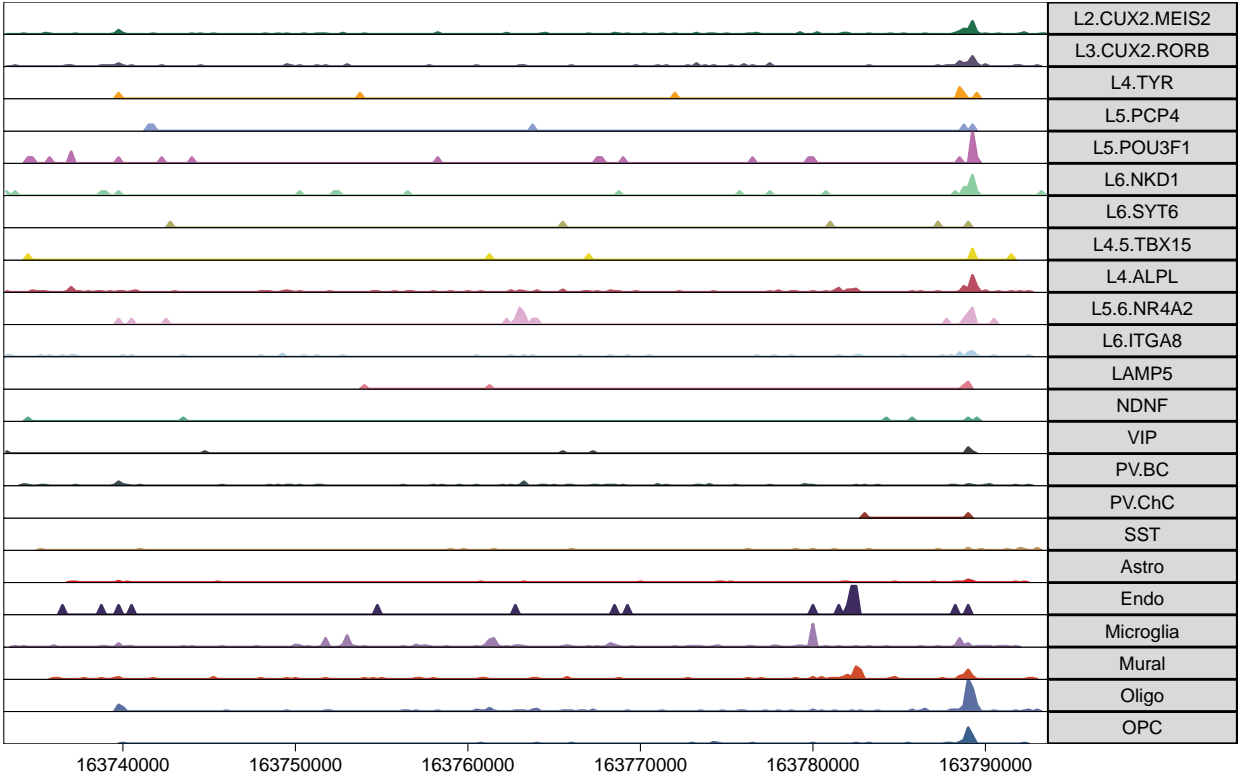


L5.6.NR4A2

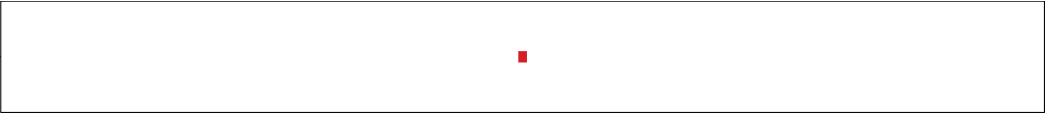


chr2:163733094–163793595 Marker Peak Tracks for: L5.6.NR4A2

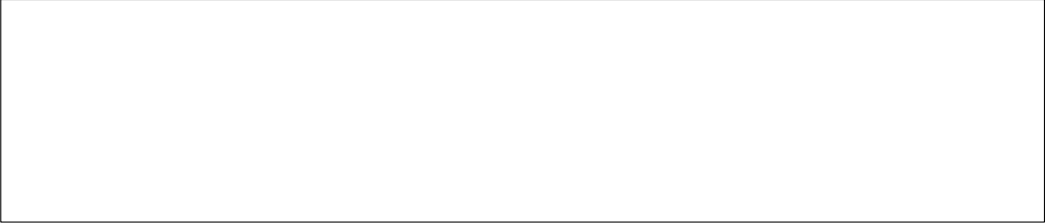
Coverage
(Norm. ATAC Signal Range 0–1.33) by ReadsInTSS)



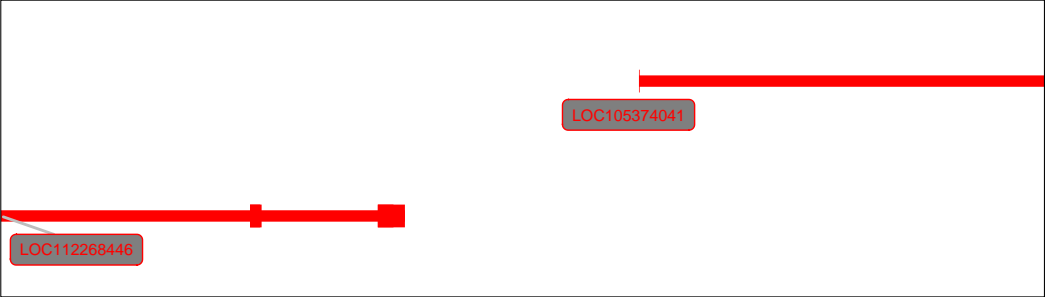
L5.6.NR4A2



Peaks

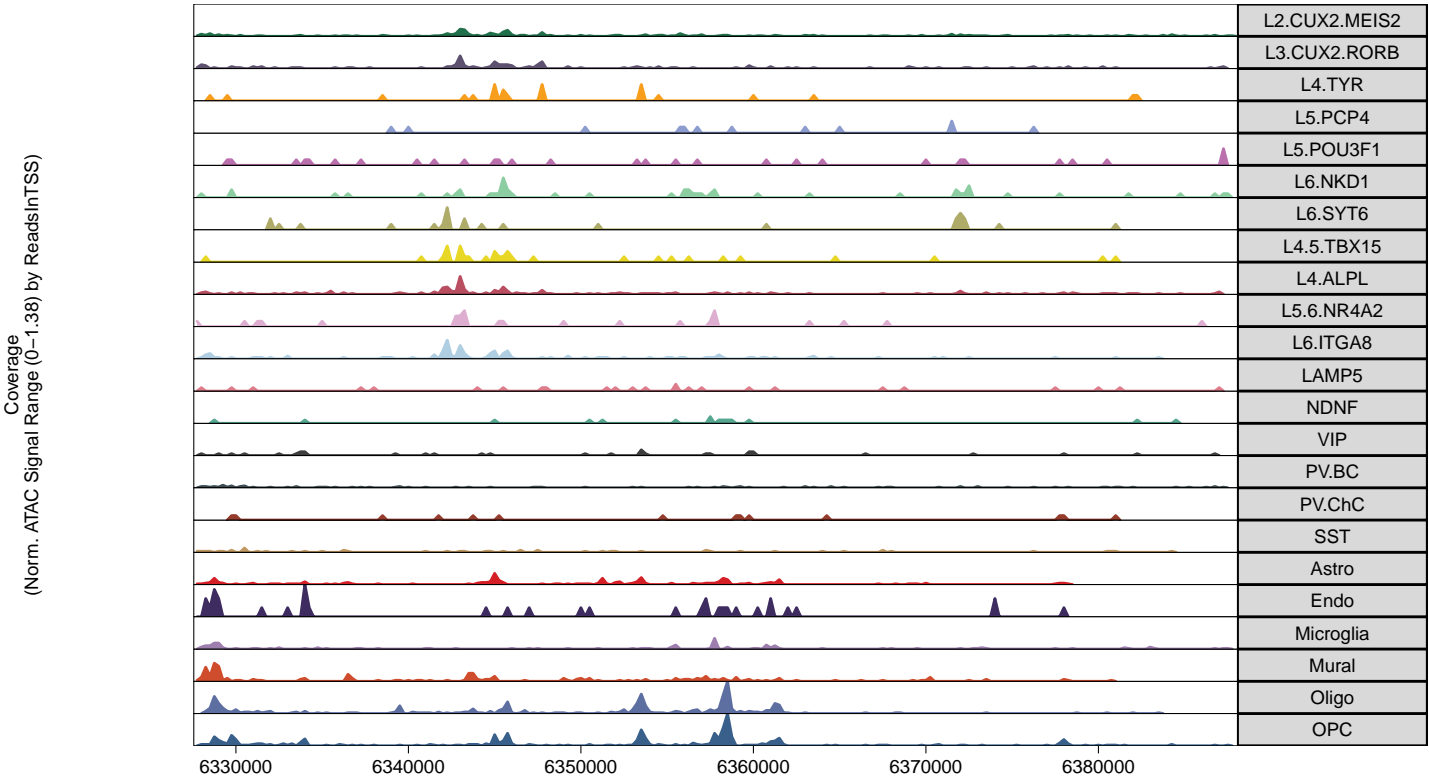


LoopTrack



Genes

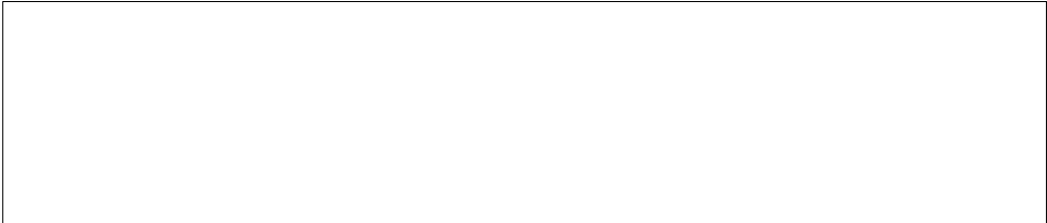
chr11:6327558–6388059 Marker Peak Tracks for: L5.6.NR4A2



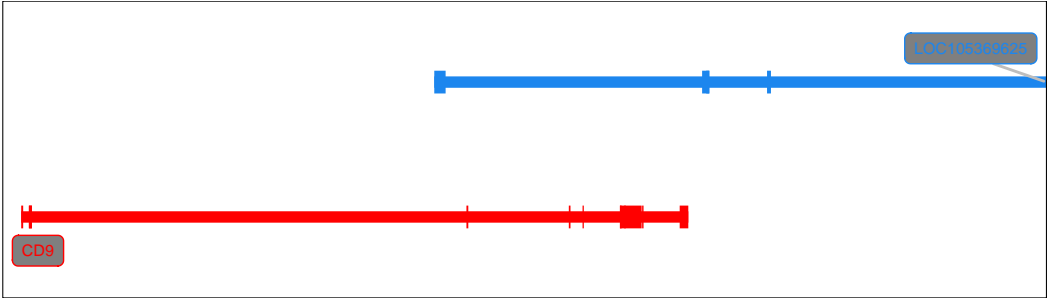
L5.6.NR4A2



Peaks

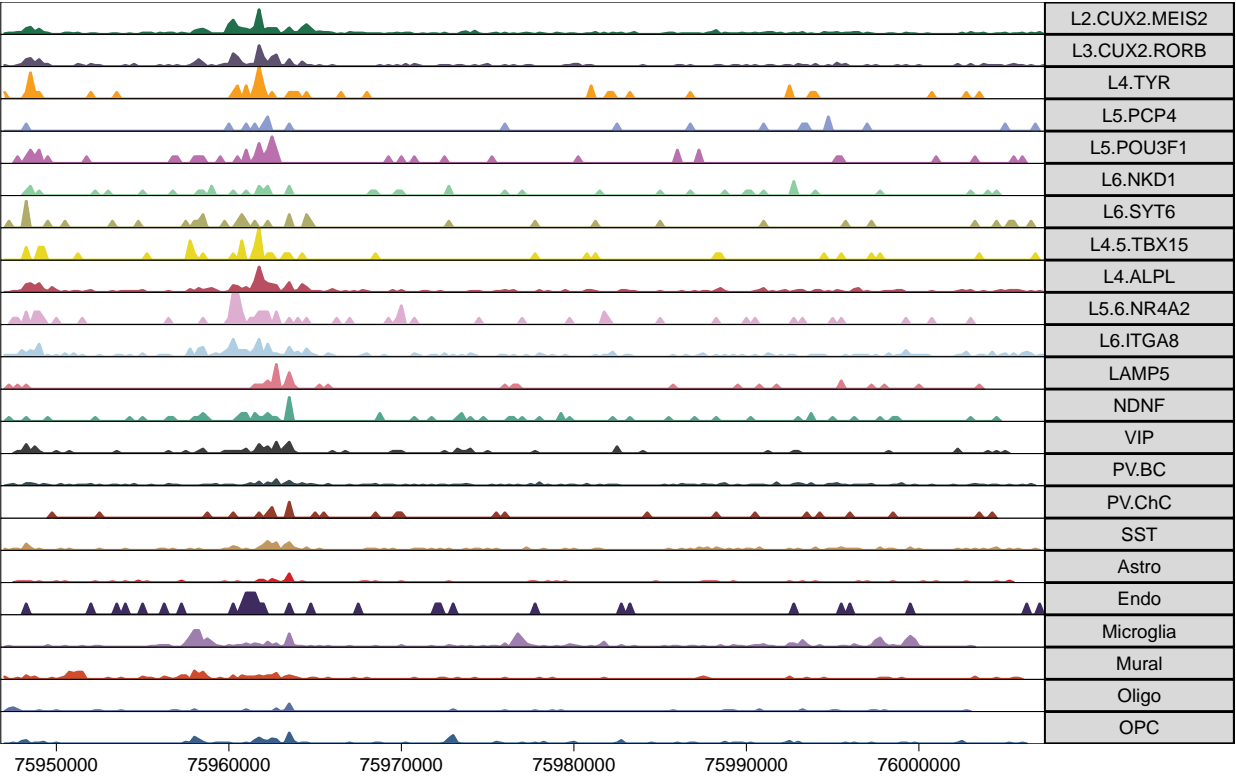


LoopTrack



Genes

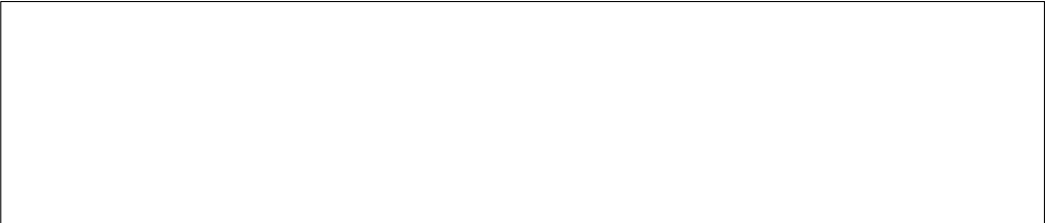
Coverage
(Norm. ATAC Signal Range (0-1.15) by ReadsInTSS)



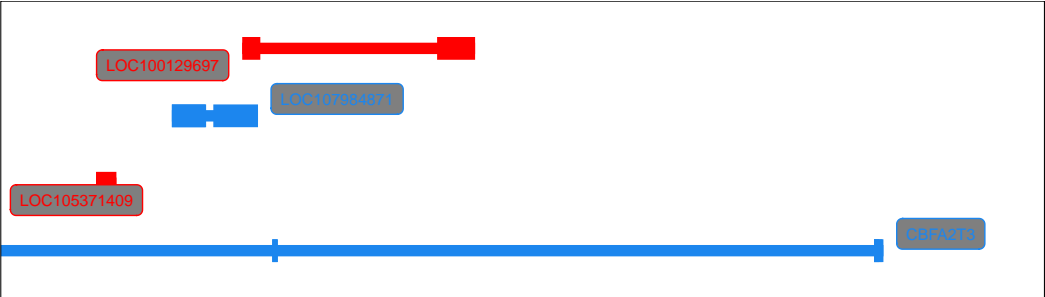
L5.6.NR4A2



Peaks

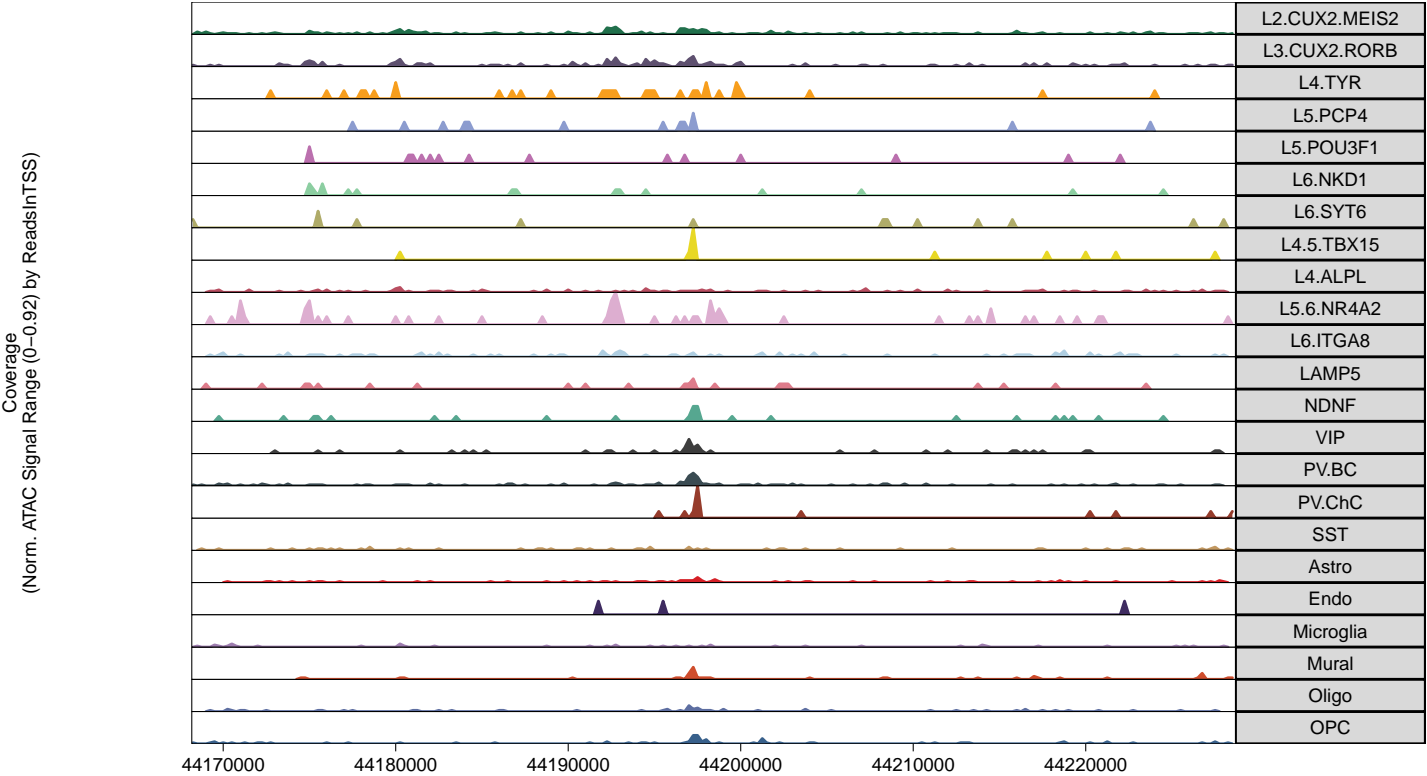


LoopTrack

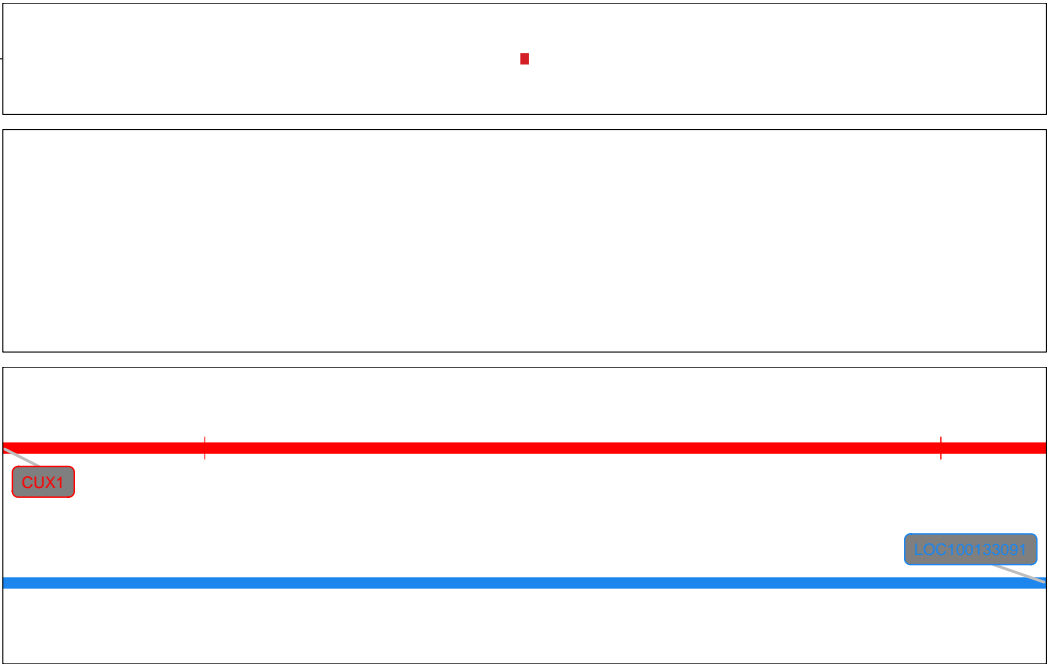


Genes

chr3:44168178–44228679 Marker Peak Tracks for: L5.6.NR4A2

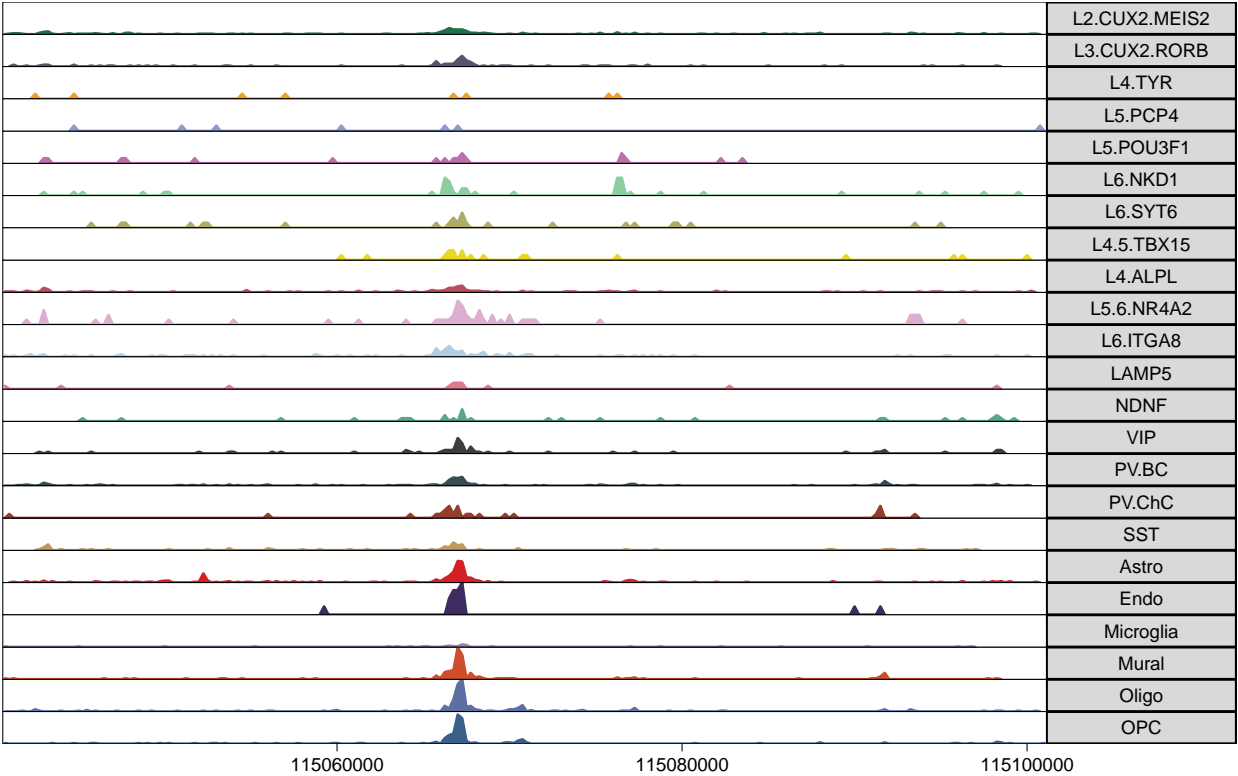


L5.6.NR4A2

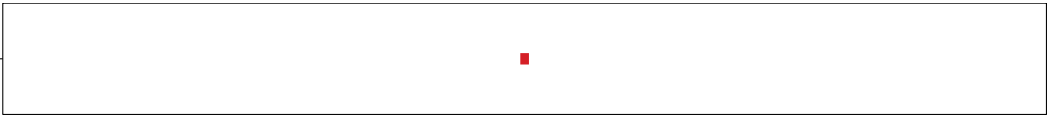


chr7:115040623–115101124 Marker Peak Tracks for: L5.6.NR4A2

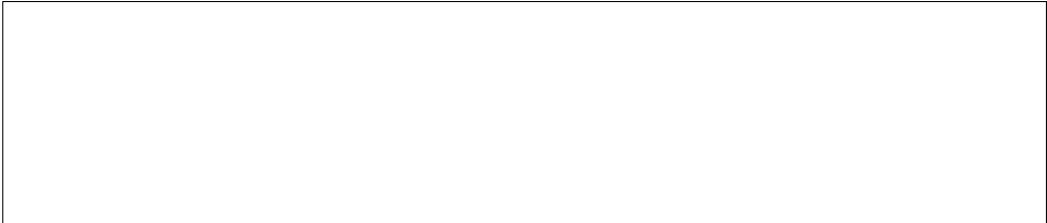
Coverage
(Norm. ATAC Signal Range 0–1.54) by ReadsInTSS)



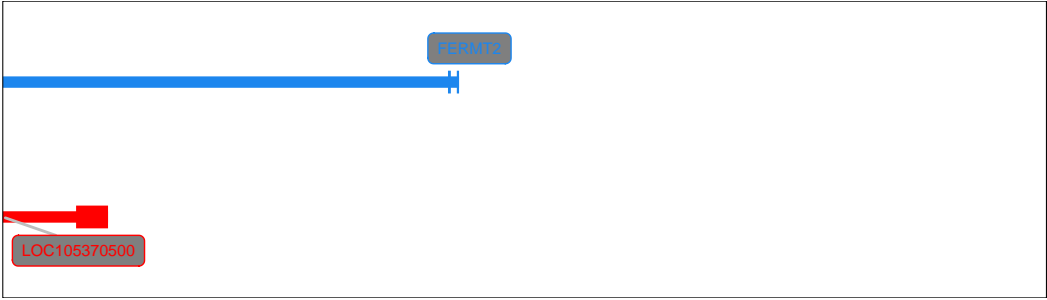
L5.6.NR4A2



Peaks



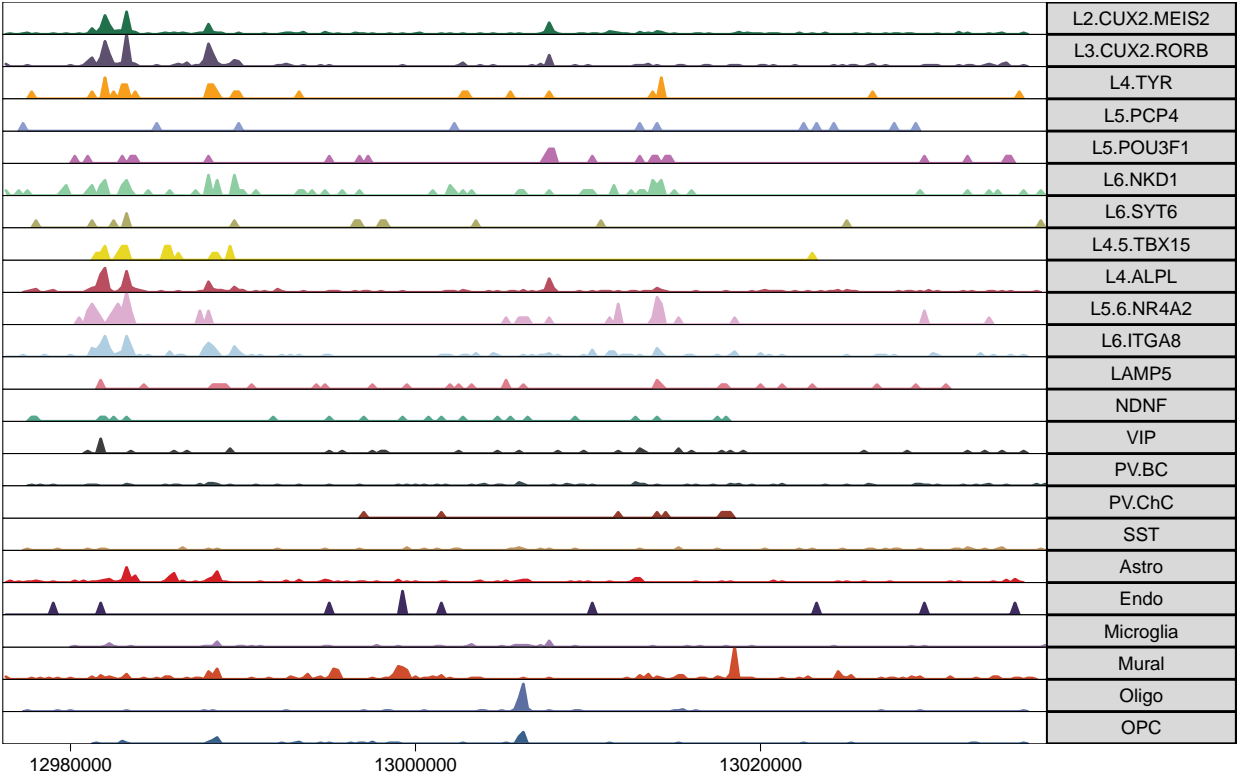
LoopTrack



Genes

chr1:12976074–13036575 Marker Peak Tracks for: L5.6.NR4A2

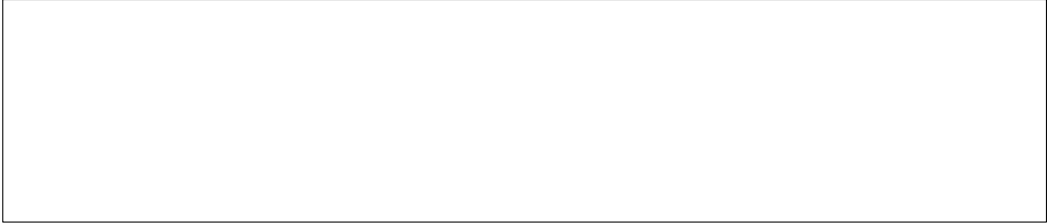
Coverage
(Norm. ATAC Signal Range 0–1.07) by ReadsInTSS)



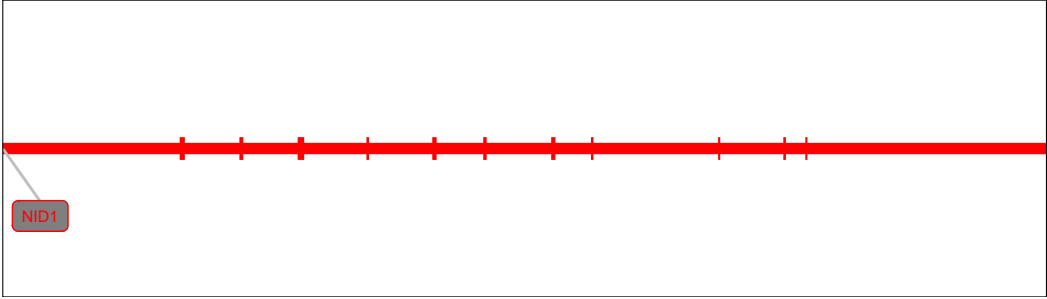
L5.6.NR4A2



Peaks

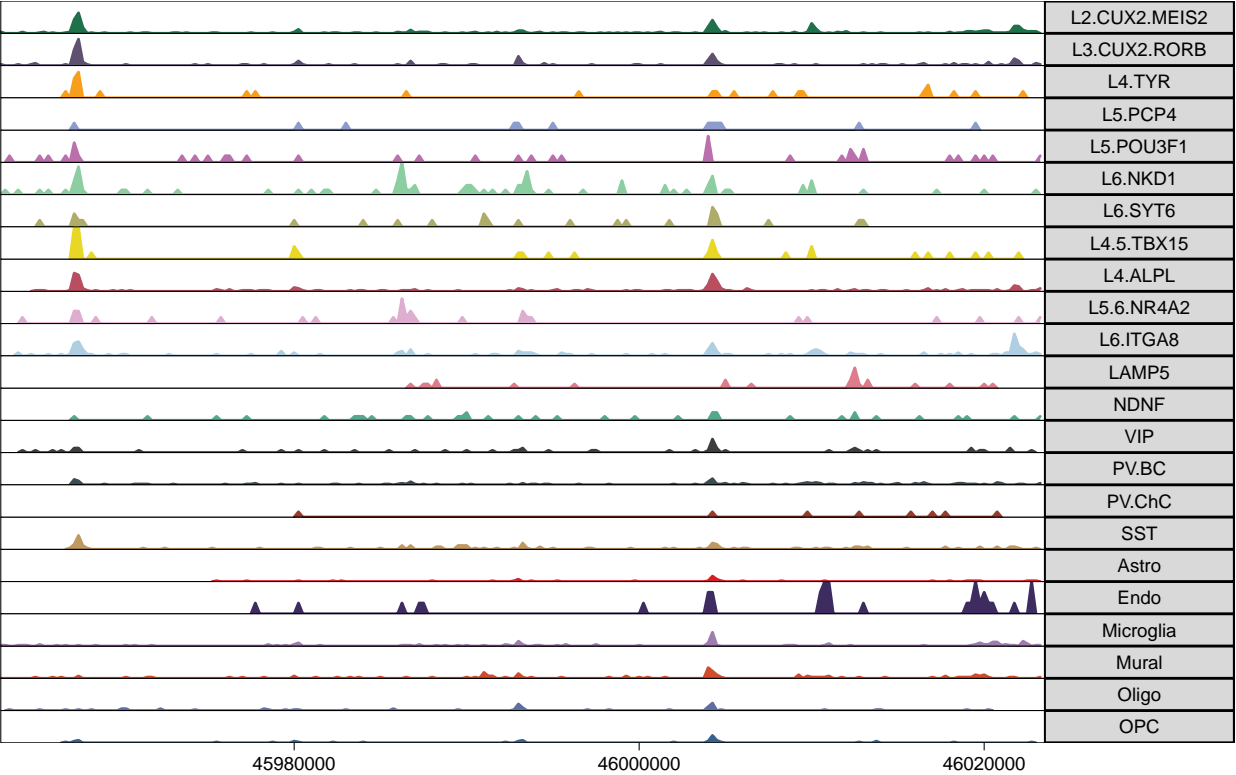


LoopTrack

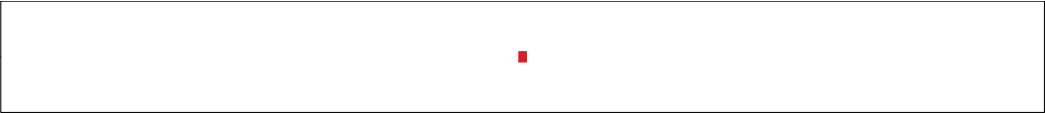


Genes

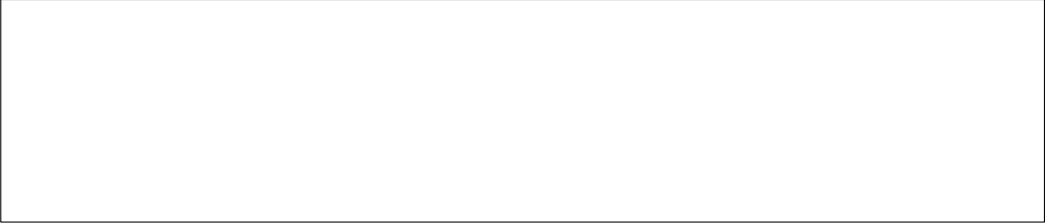
Coverage
(Norm. ATAC Signal Range 0–1.17) by ReadsInTSS)



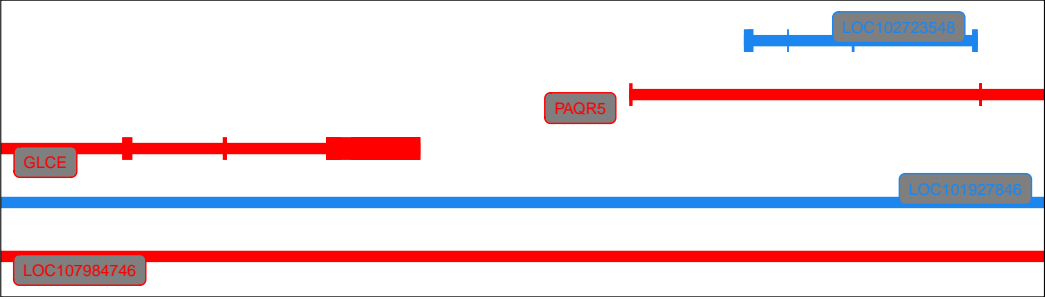
L5.6.NR4A2



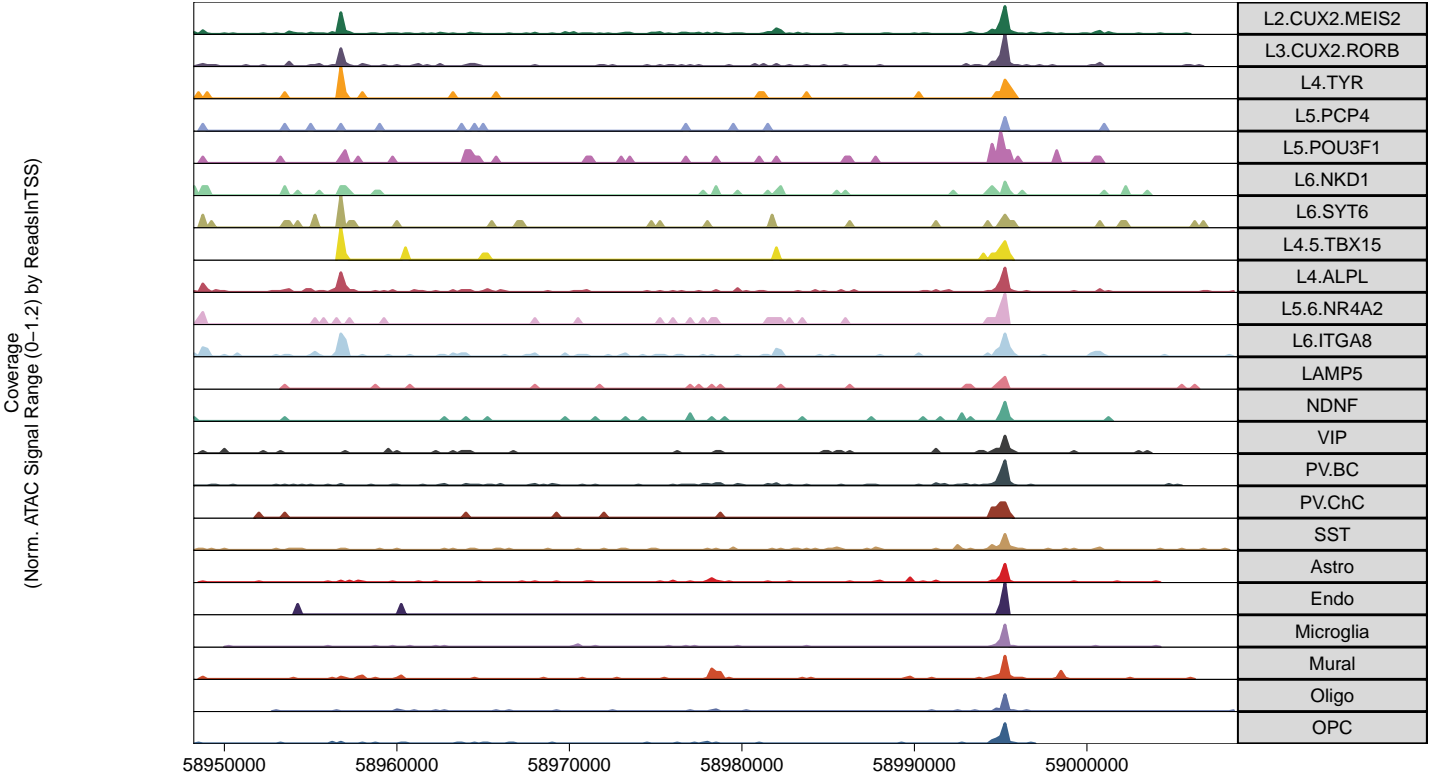
Peaks



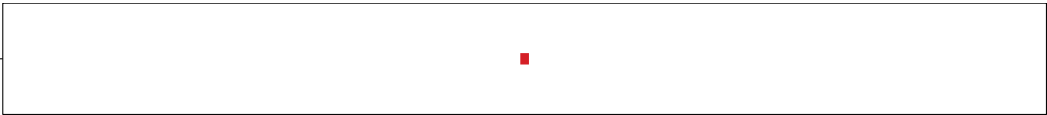
LoopTrack



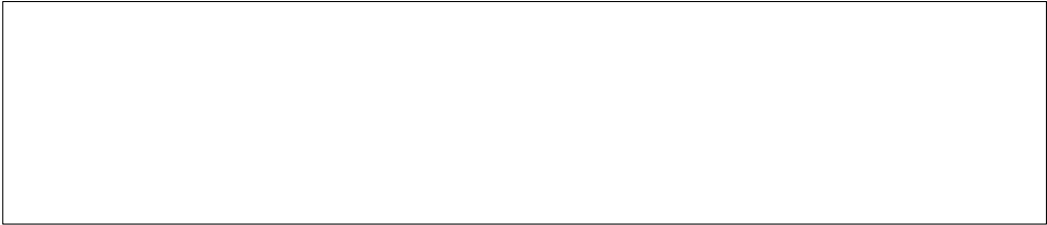
Genes



L5.6.NR4A2



Peaks

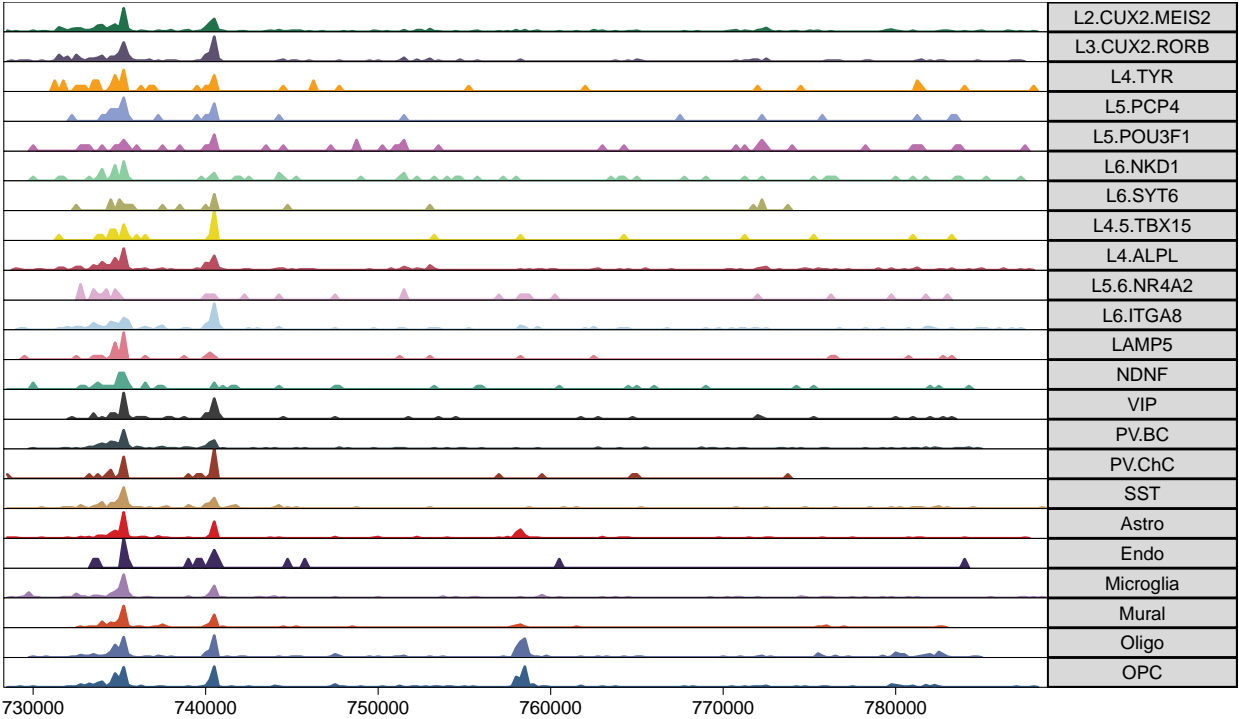


LoopTrack



Genes

Coverage
(Norm. ATAC Signal Range (0–1.3) by ReadslnTSS)



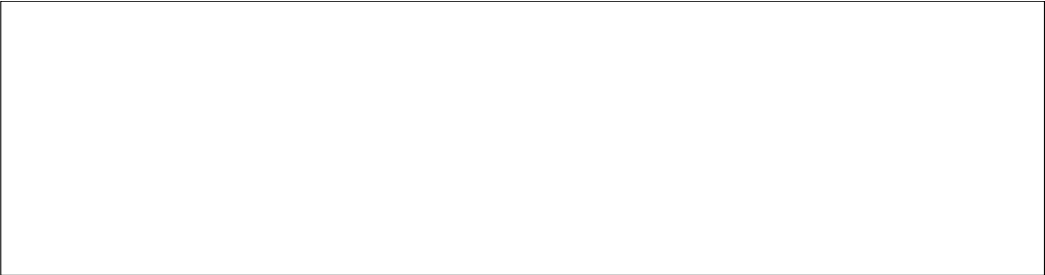
L5.6.NR4A2



Peaks

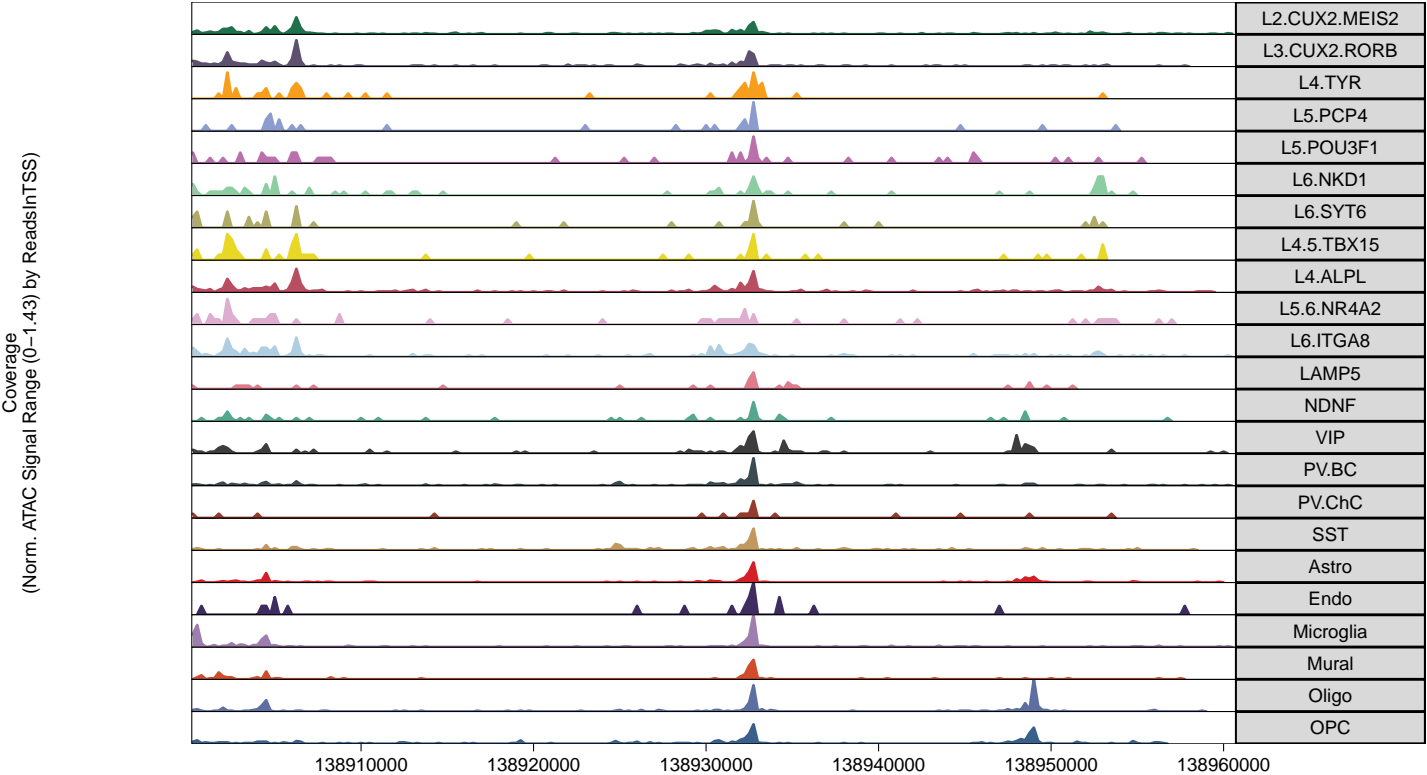


LoopTrack

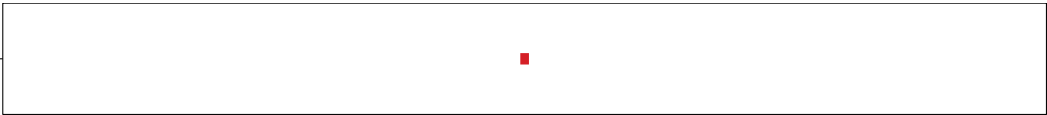


GeneTrack

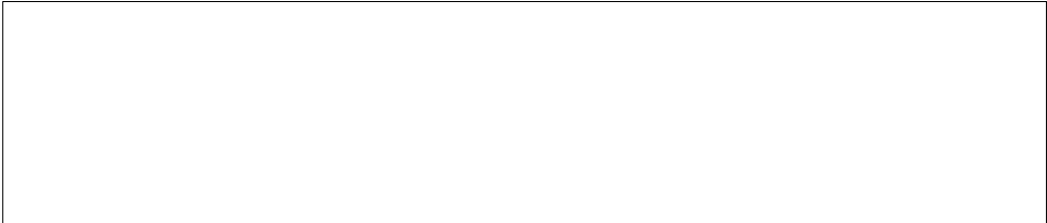
chr7:138900187-138960688 Marker Peak Tracks for: L5.6.NR4A2



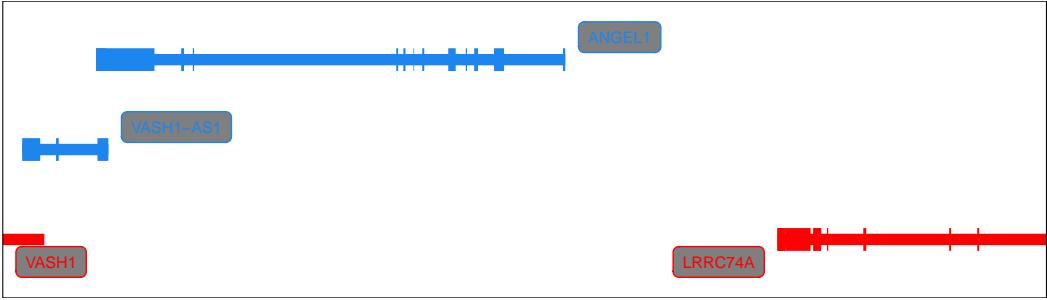
L5.6.NR4A2



Peaks

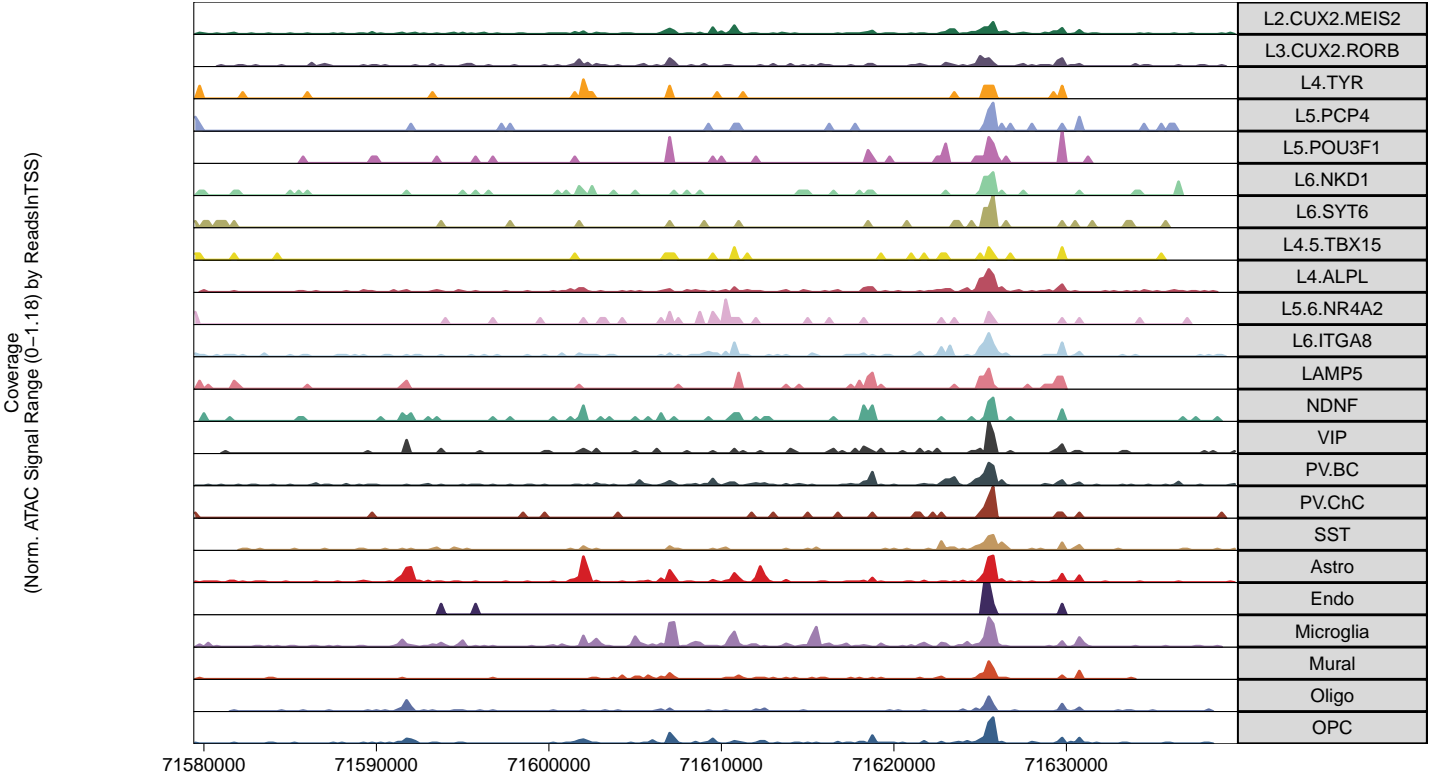


LoopTrack

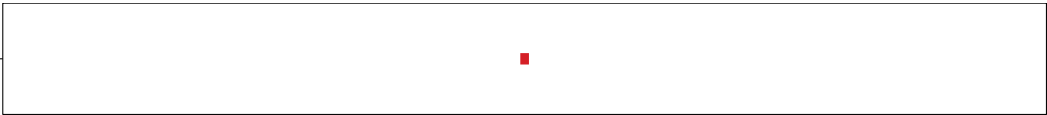


Genes

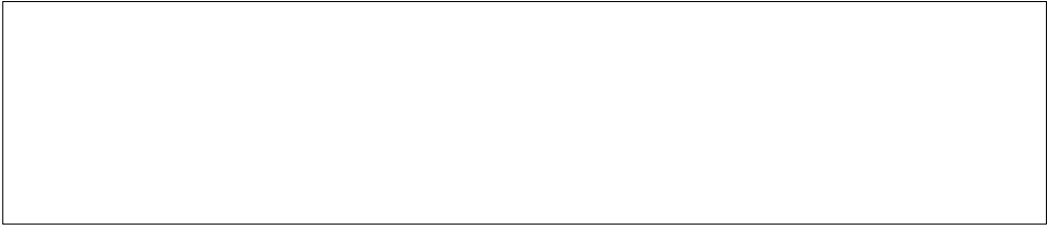
chr20:71579413–71639914 Marker Peak Tracks for: L5.6.NR4A2



L5.6.NR4A2



Peaks



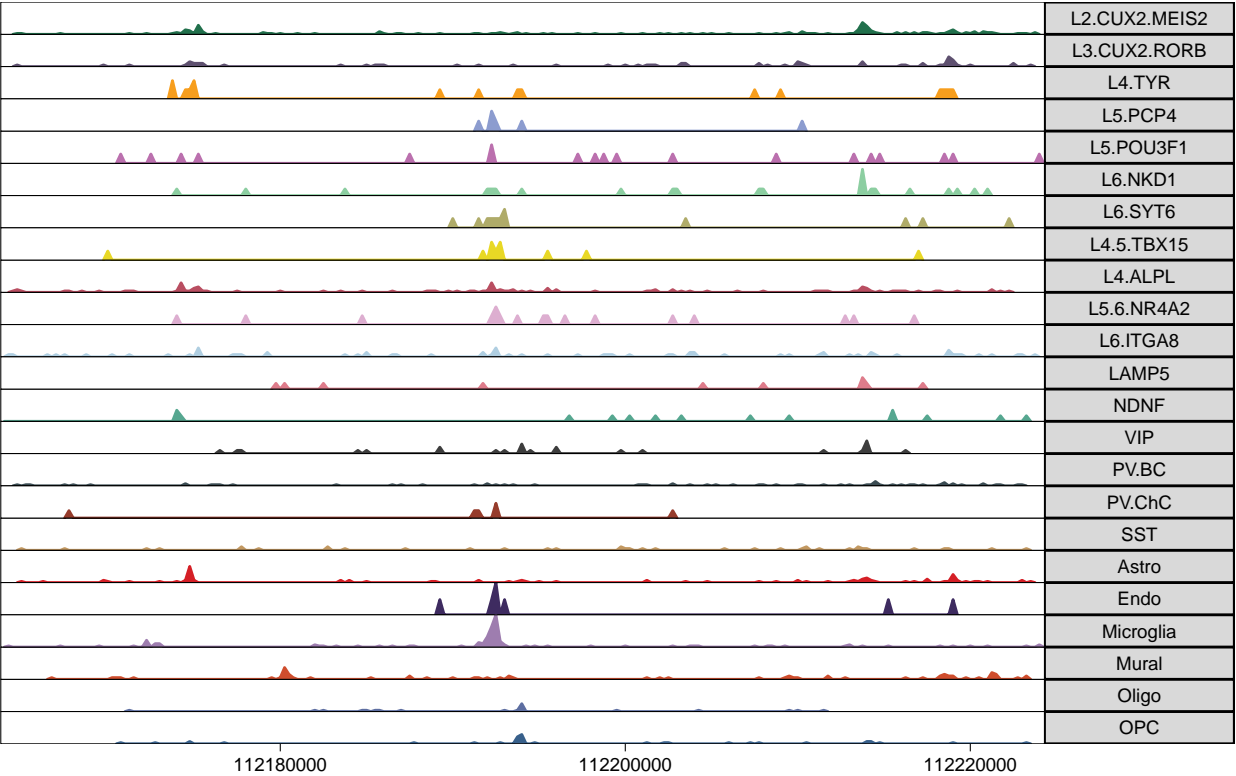
LoopTrack



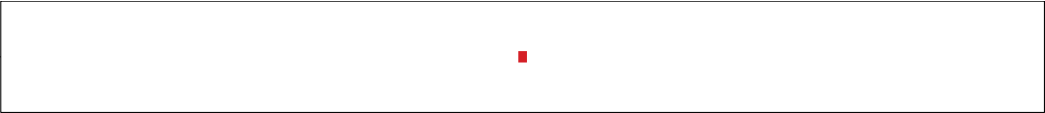
Genes

CUX1

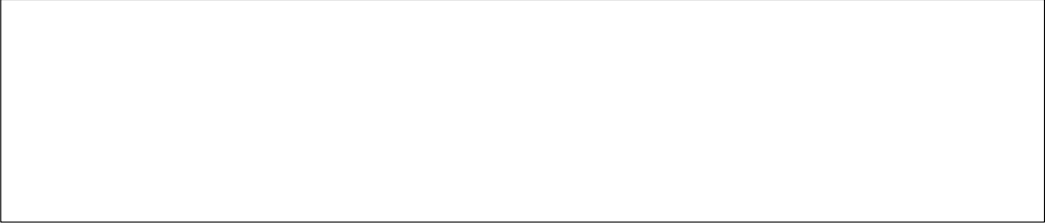
Coverage
(Norm. ATAC Signal Range 0–0.83) by ReadsInTSS)



L5.6.NR4A2



Peaks

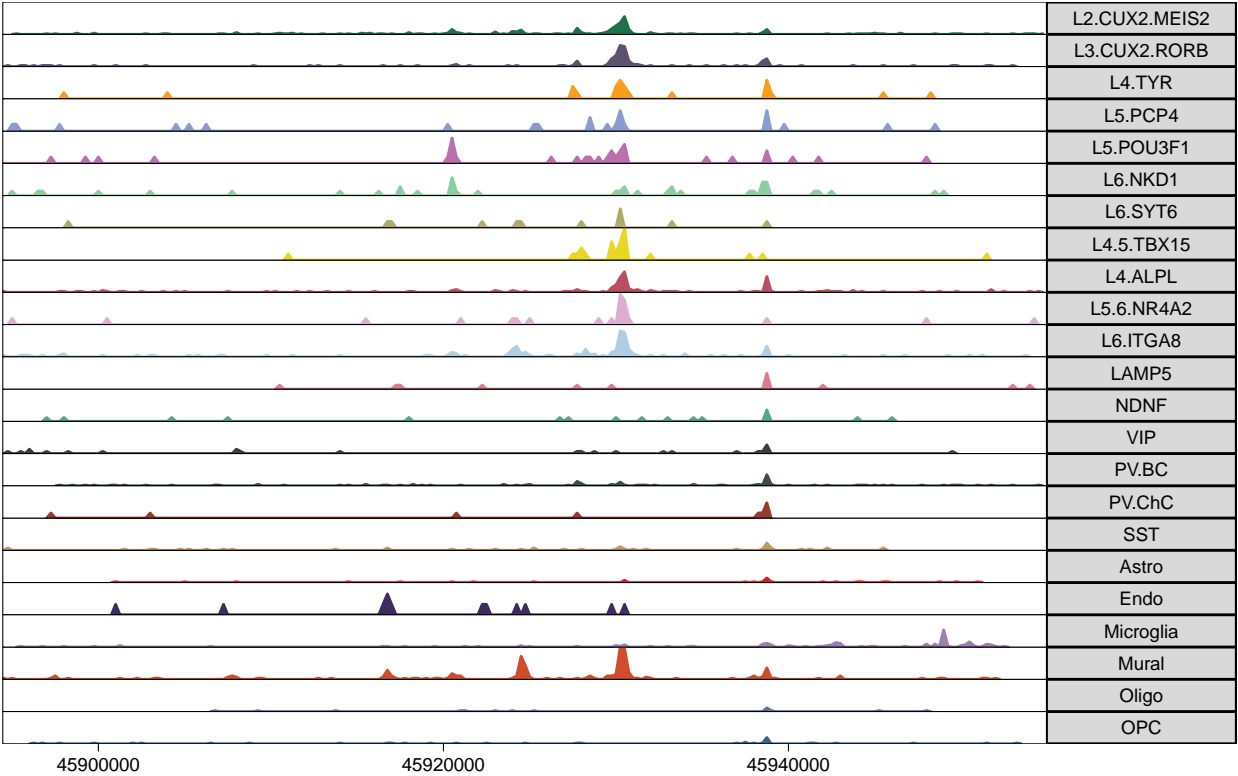


LoopTrack

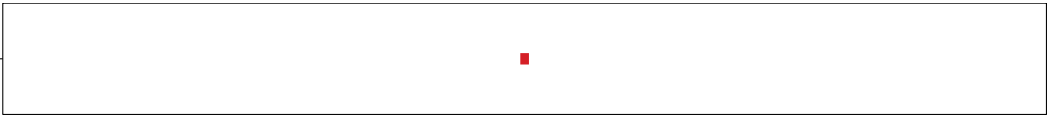


Genes

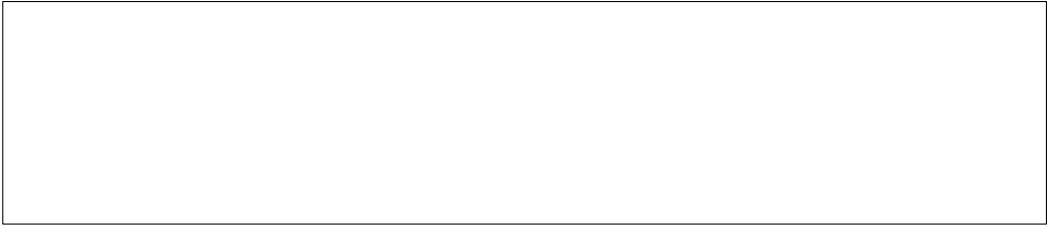
Coverage
(Norm. ATAC Signal Range 0-1.21) by ReadsInTSS)



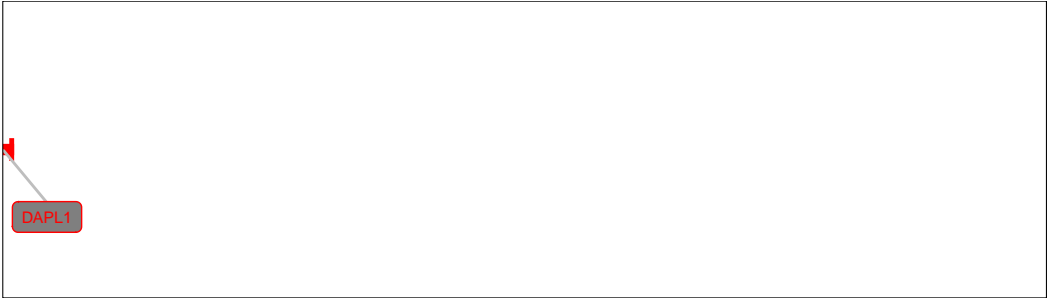
L5.6.NR4A2



Peaks



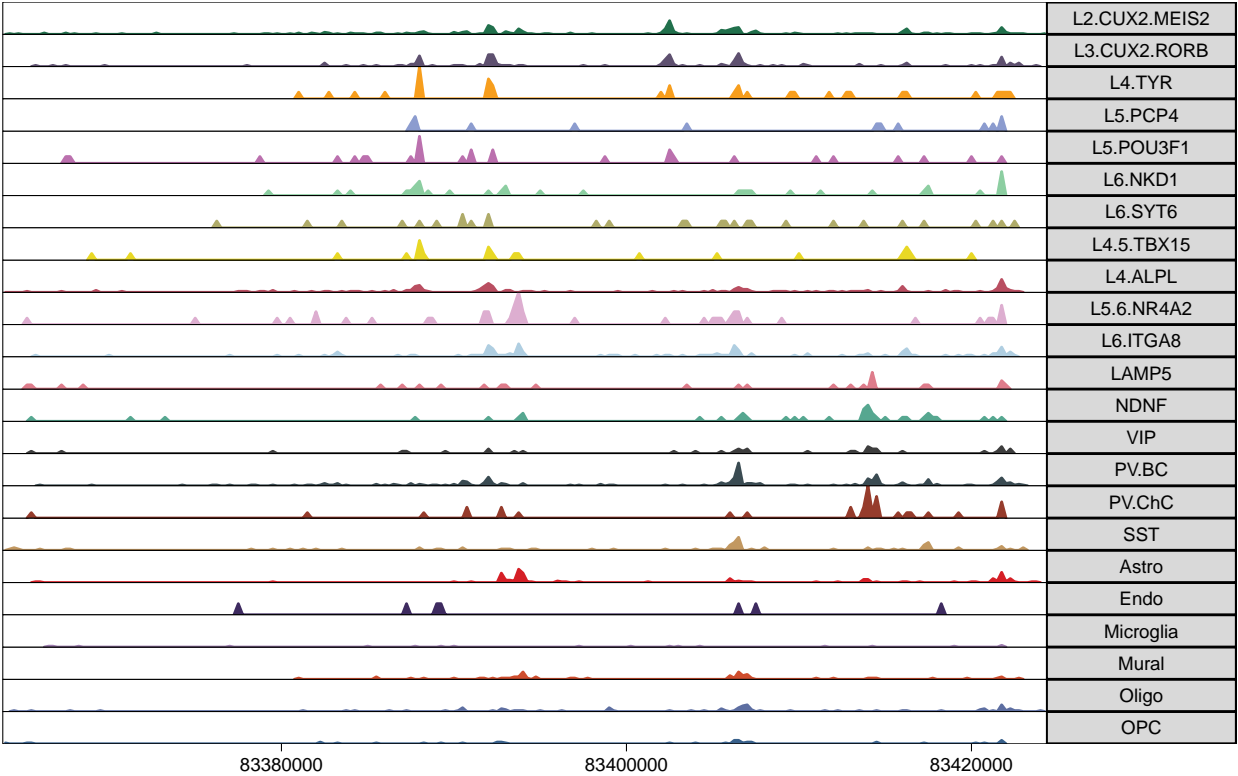
LoopTrack



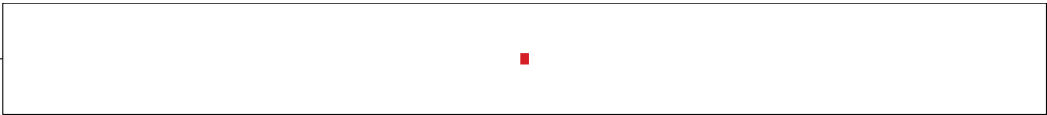
Genes

chr4:83363845–83424346 Marker Peak Tracks for: L5.6.NR4A2

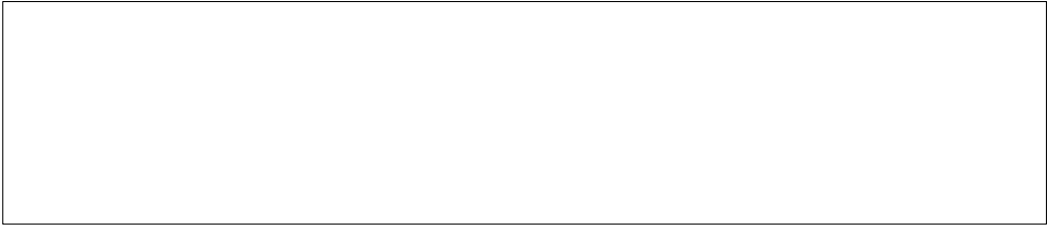
Coverage
(Norm. ATAC Signal Range 0–1.14) by ReadsInTSS)



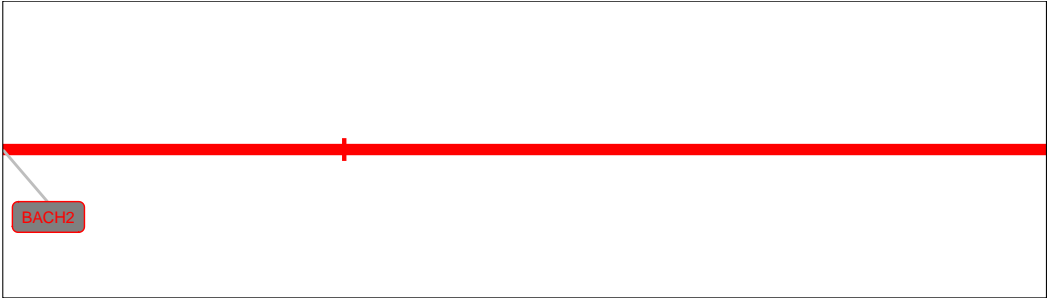
L5.6.NR4A2



Peaks

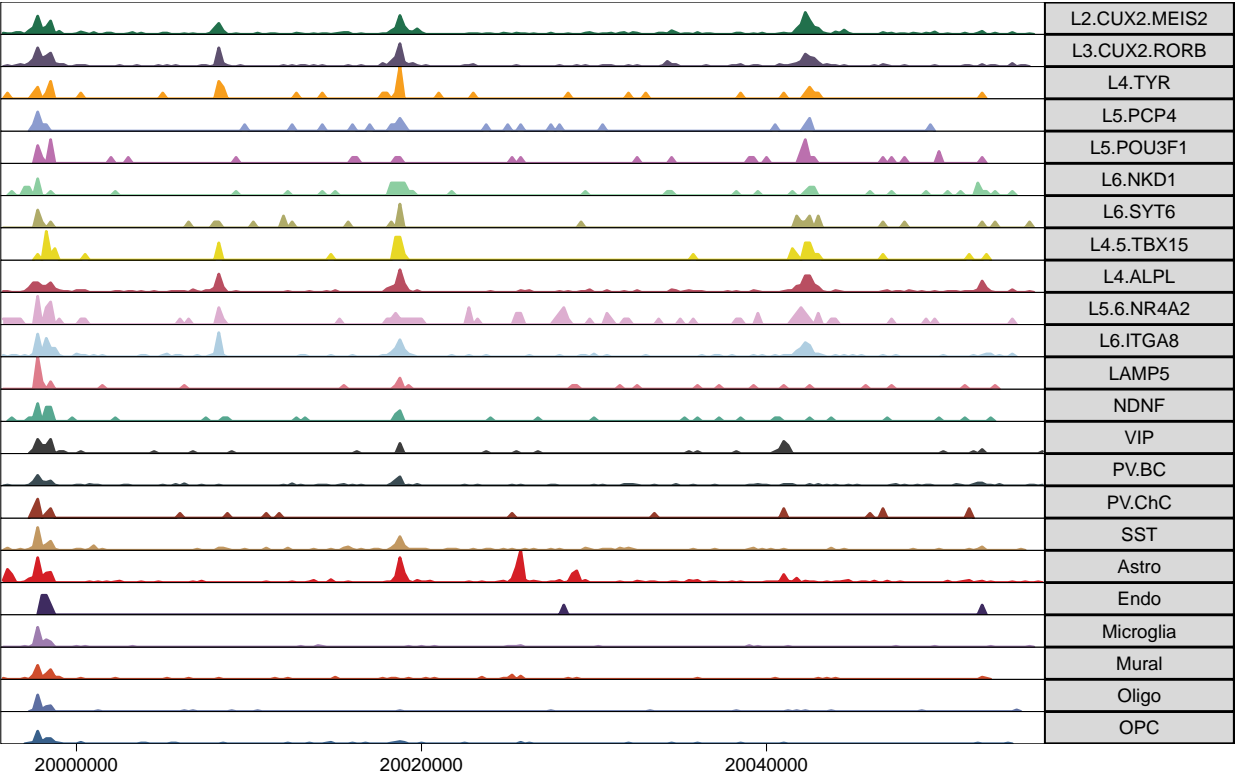


LoopTrack

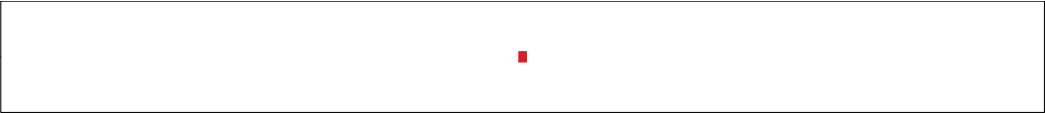


Genes

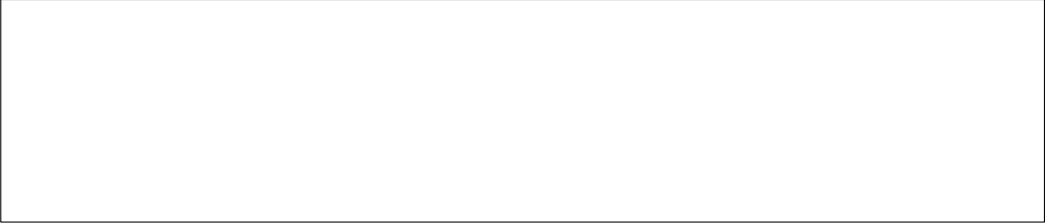
Coverage
(Norm. ATAC Signal Range 0–1.29) by ReadsInTSS)



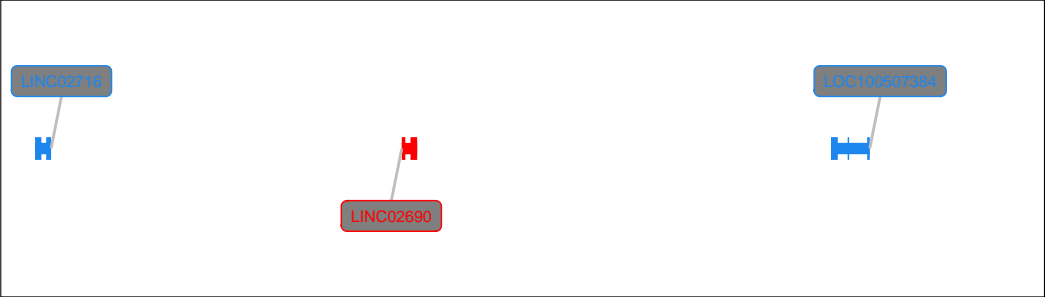
L5.6.NR4A2



Peaks

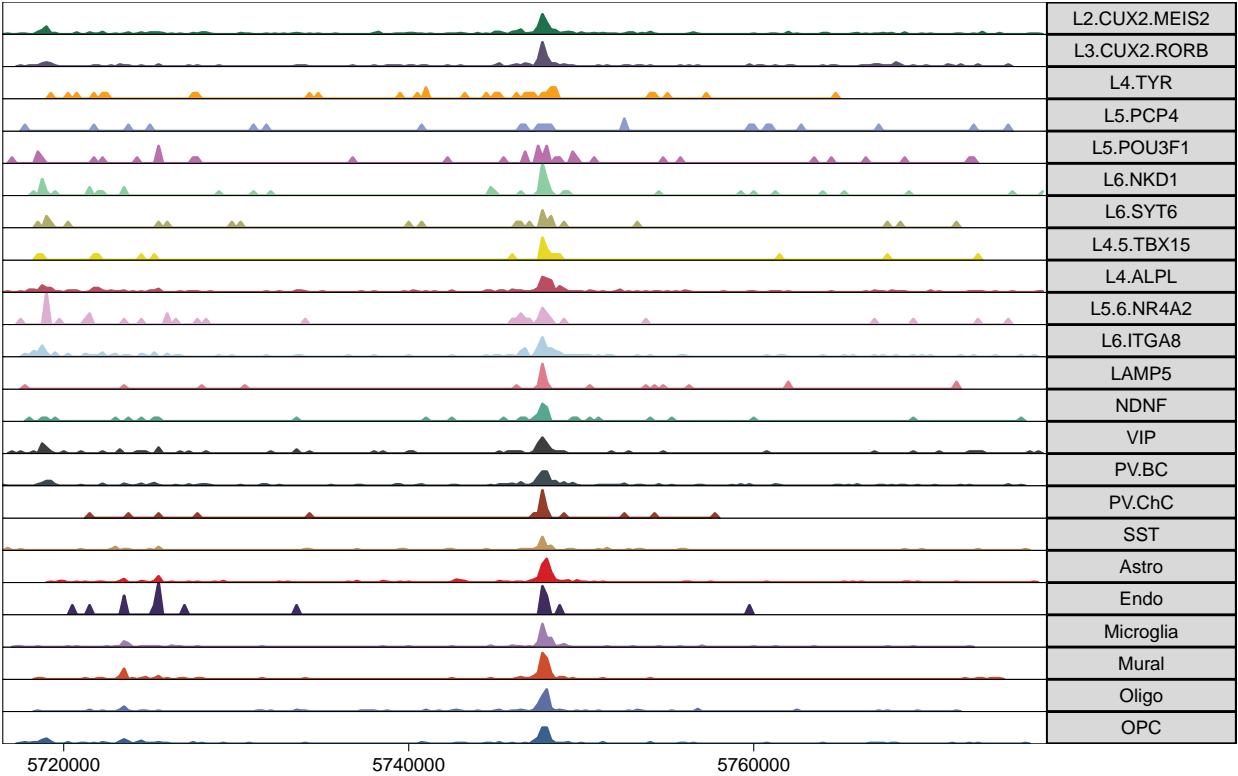


LoopTrack

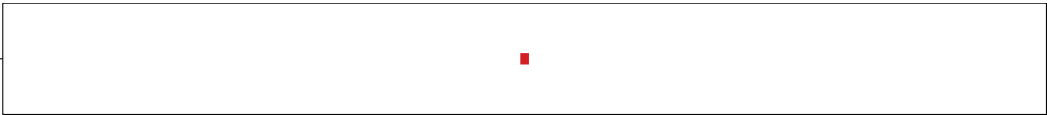


Genes

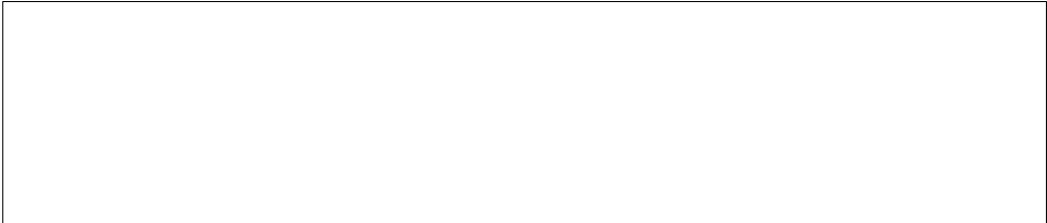
Coverage
(Norm. ATAC Signal Range 0–1.33) by ReadsInTSS)



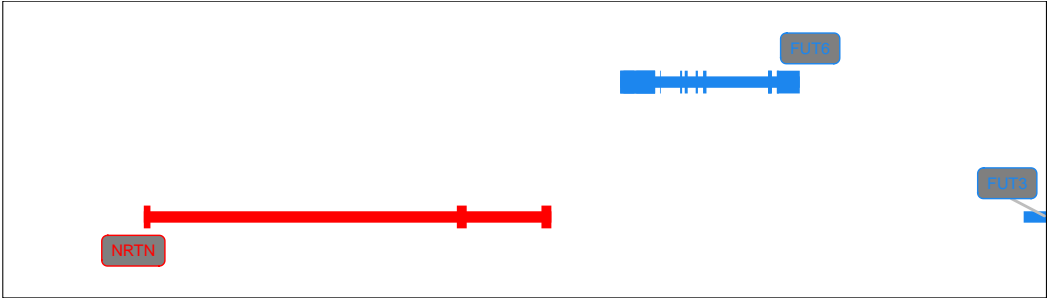
L5.6.NR4A2



Peaks

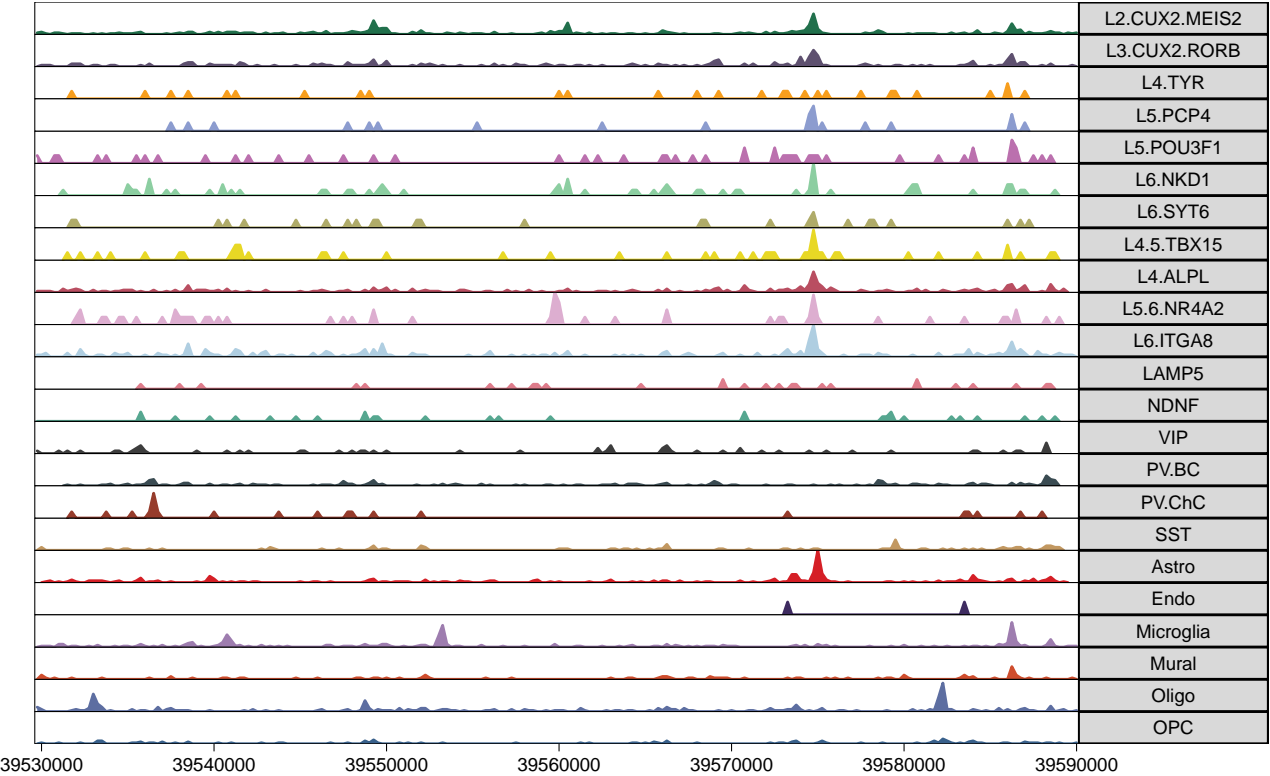


LoopTrack



Genes

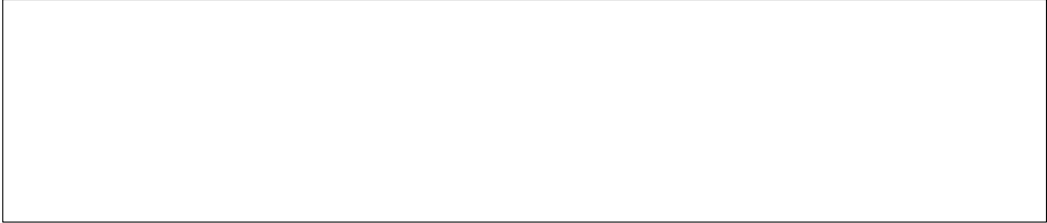
Coverage
(Norm. ATAC Signal Range (0–0.99) by ReadsInTSS)



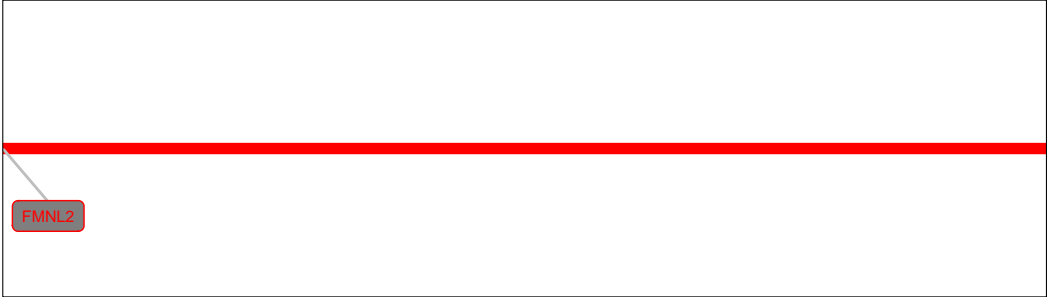
L5.6.NR4A2



Peaks

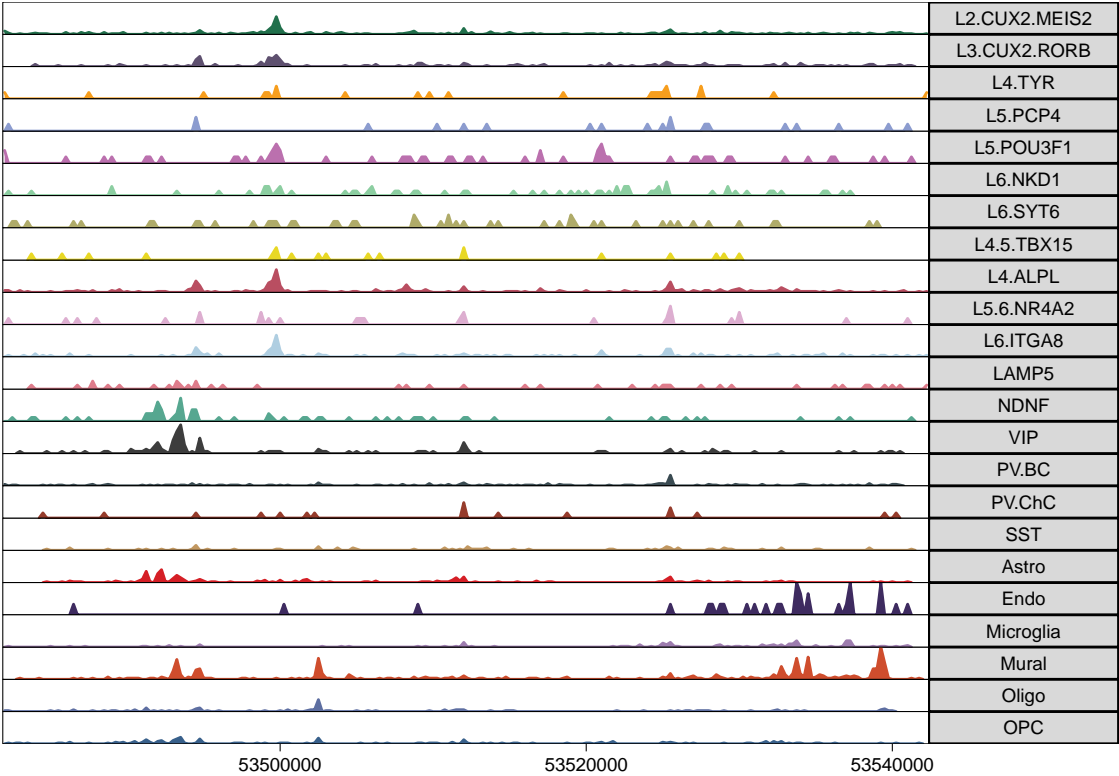


LoopTrack

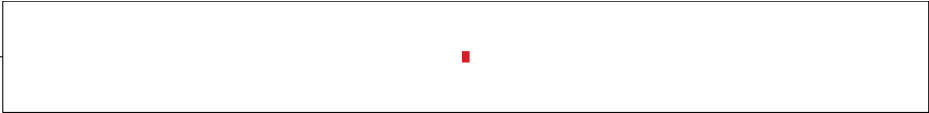


Genes

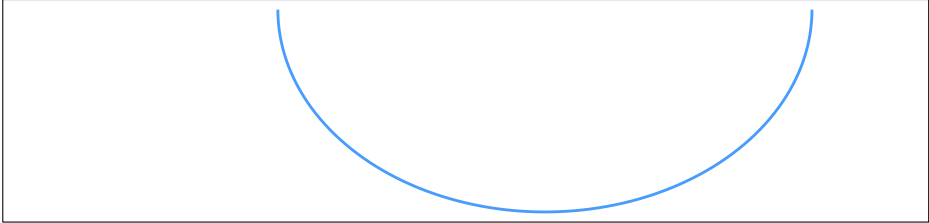
Coverage
(Norm. ATAC Signal Range 0-1.19) by ReadsInTSS)



L5.6.NR4A2



Peaks



Peak2GeneLinks

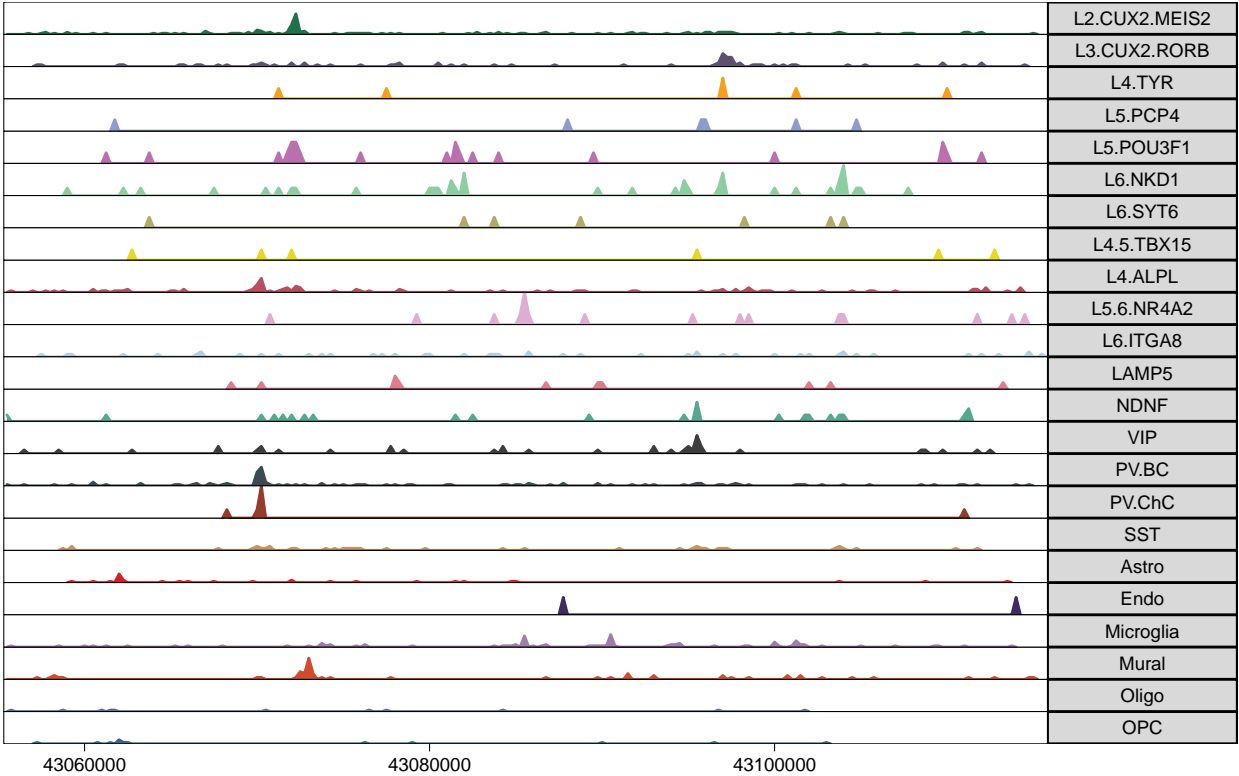


Genes

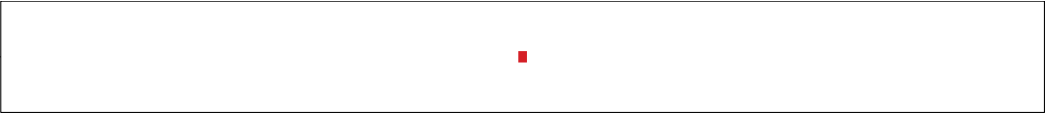


chr4:43055317–43115818 Marker Peak Tracks for: L5.6.NR4A2

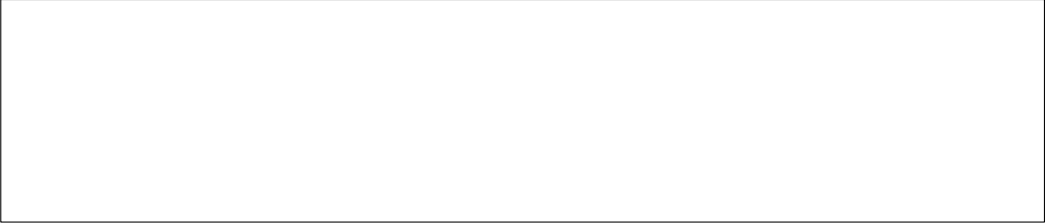
Coverage
(Norm. ATAC Signal Range (0–0.72) by ReadsInTSS)



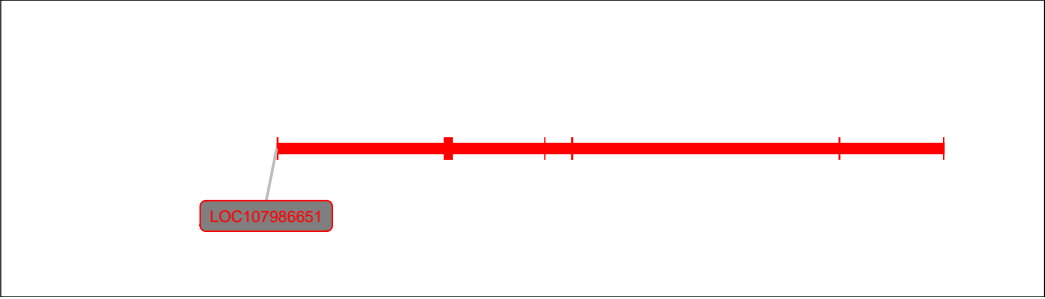
L5.6.NR4A2



Peaks

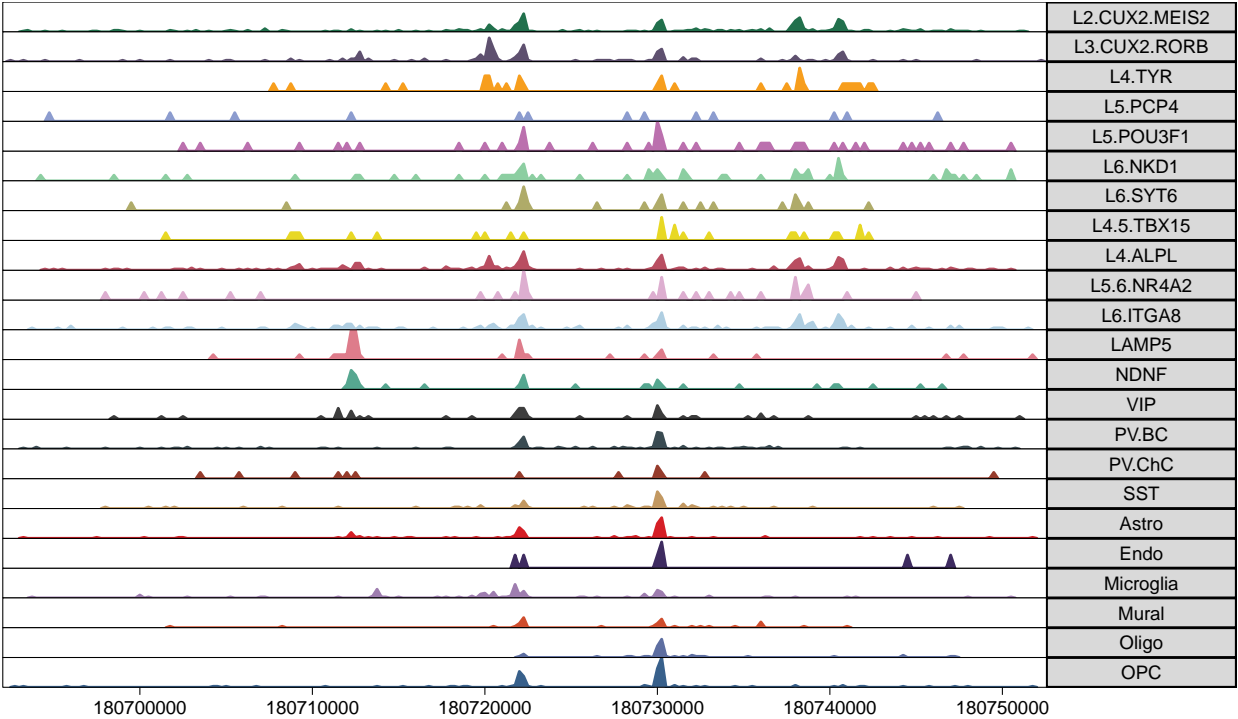


LoopTrack

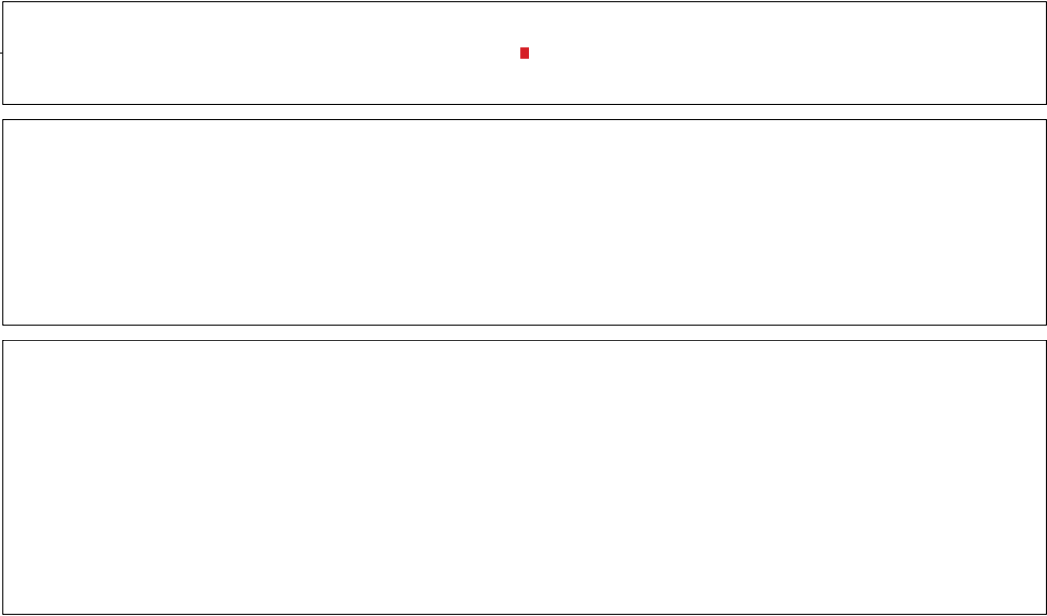


Genes

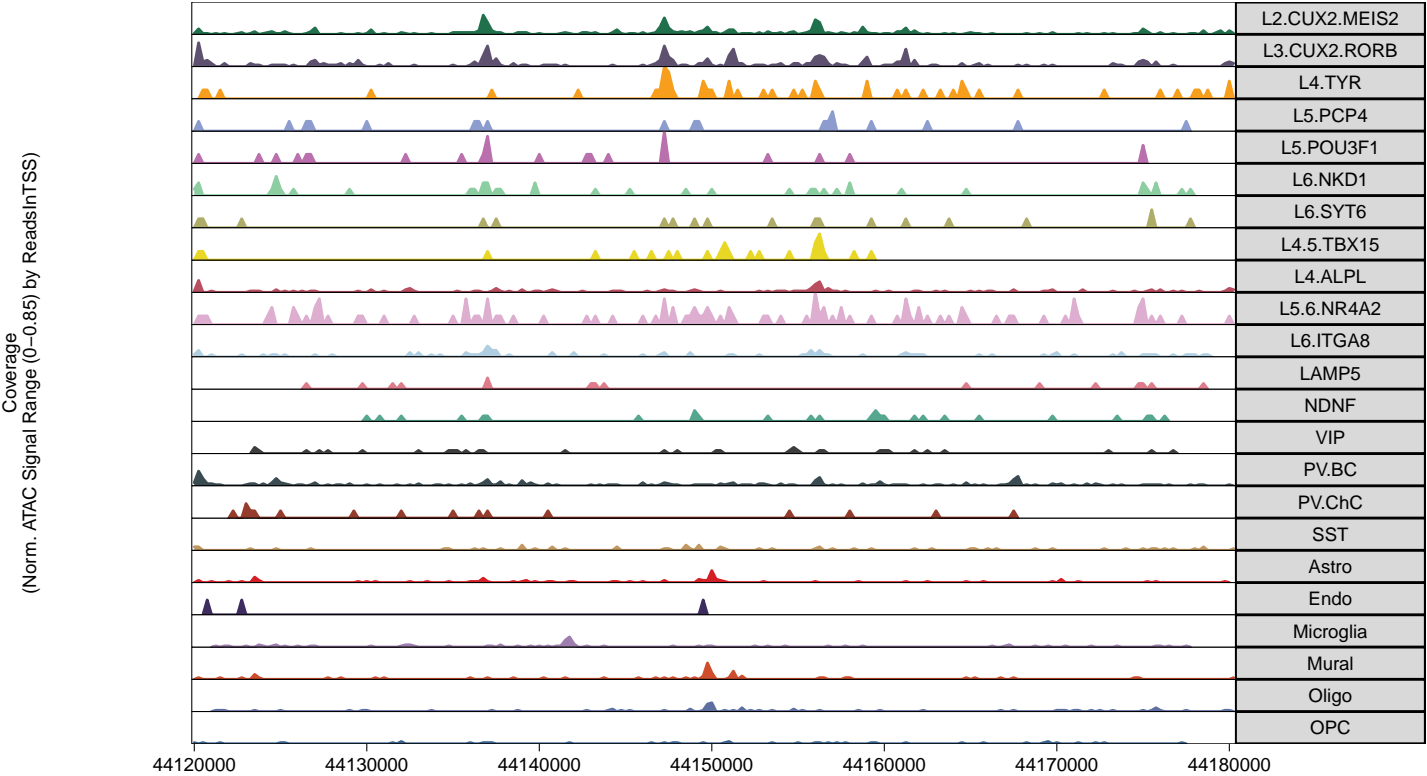
Coverage
(Norm. ATAC Signal Range (0–0.89) by Reads in TSS)



L5.6.NR4A2



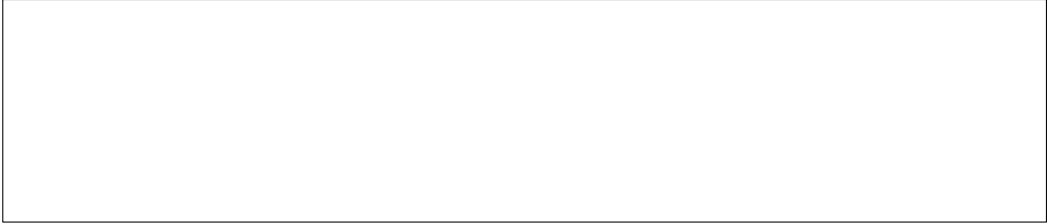
chr3:44119854–44180355 Marker Peak Tracks for: L5.6.NR4A2



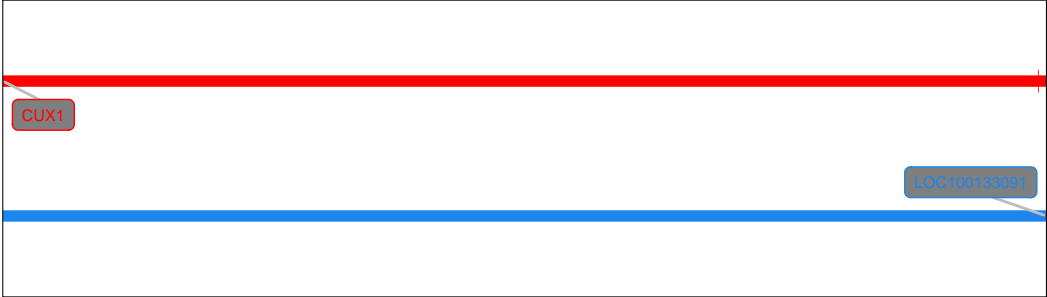
L5.6.NR4A2



Peaks

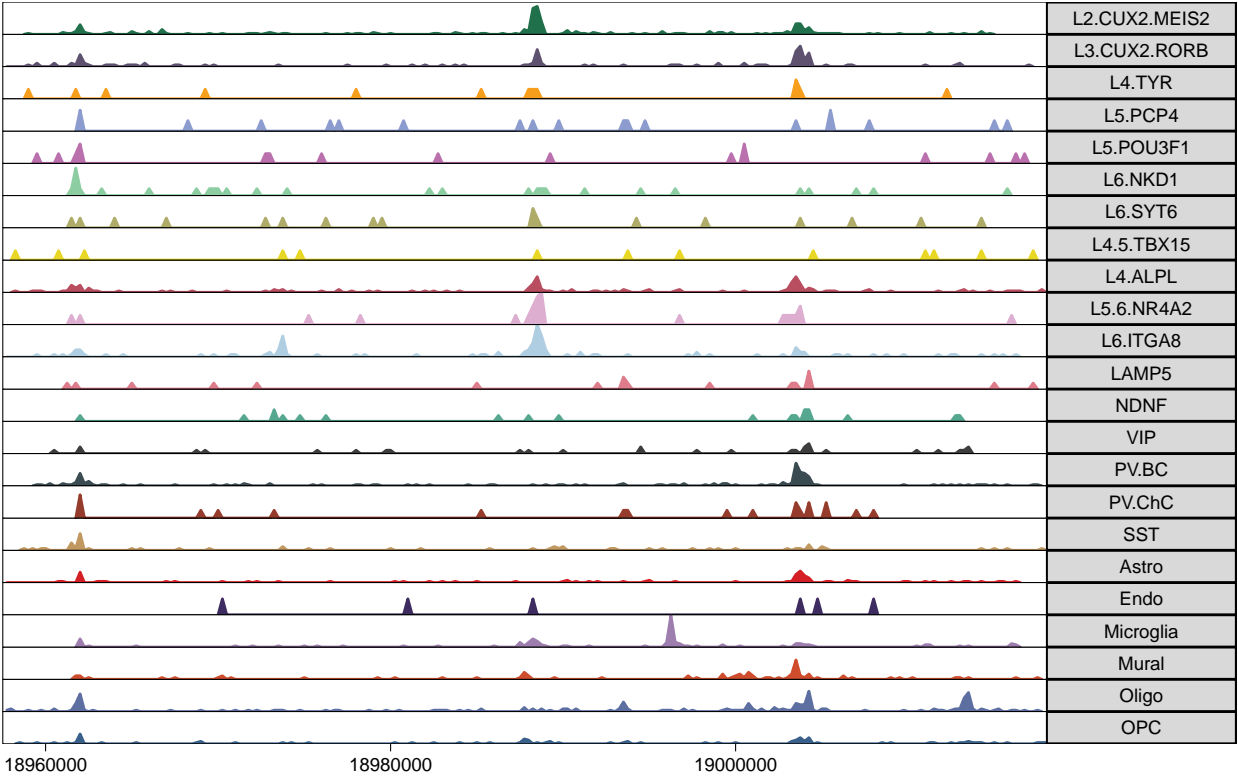


LoopTrack

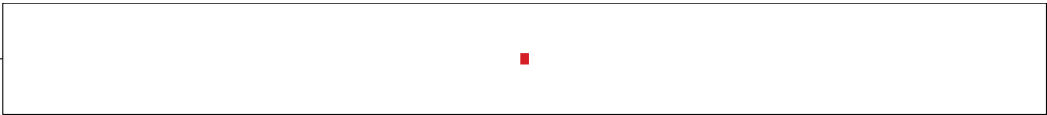


Genes

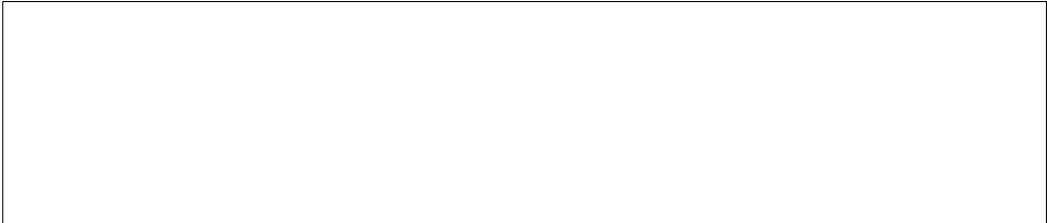
Coverage
(Norm. ATAC Signal Range (0–0.79) by ReadsInTSS)



L5.6.NR4A2



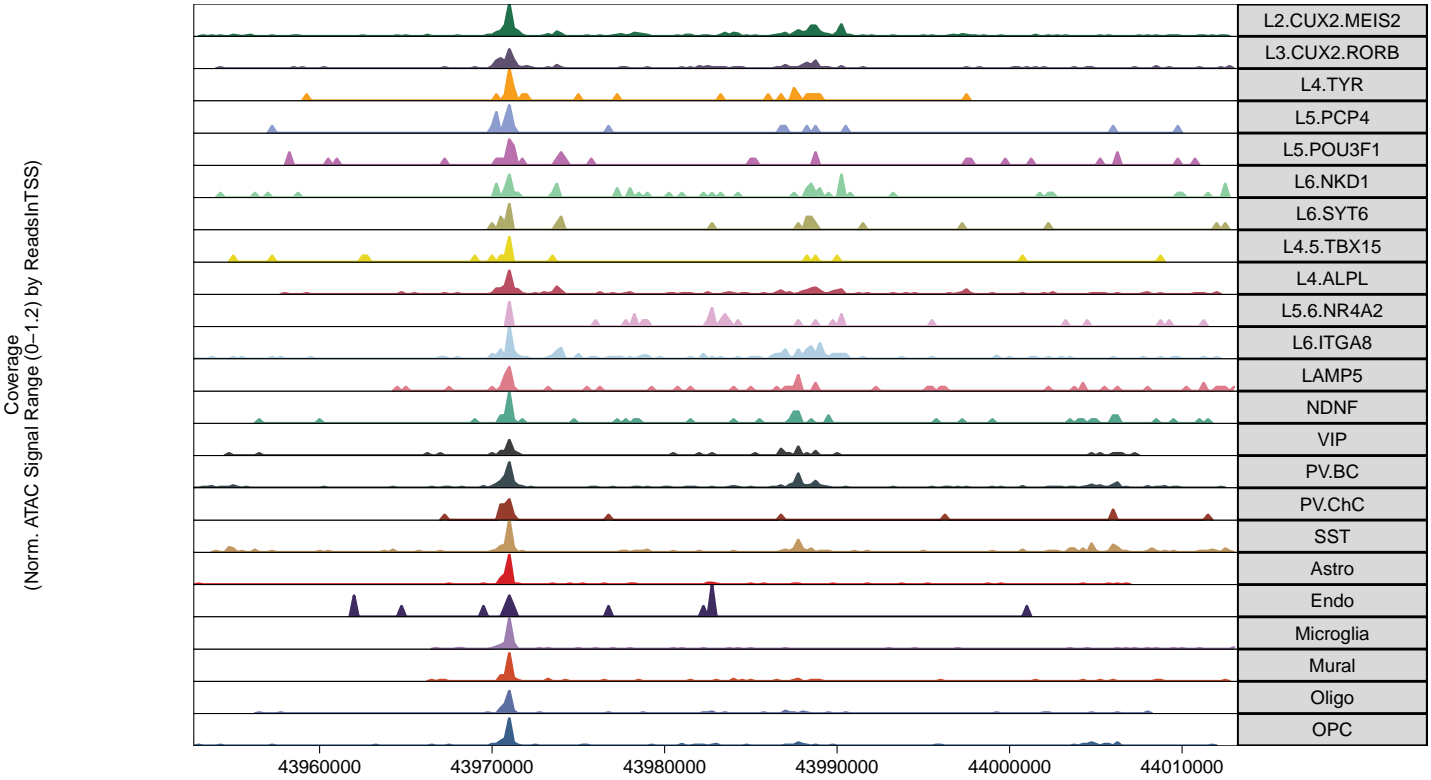
Peaks



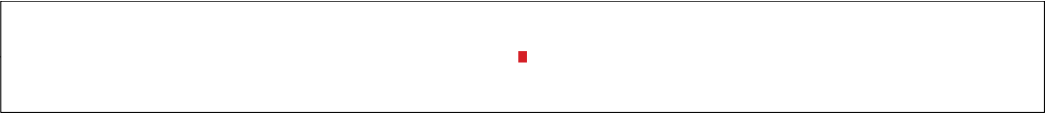
LoopTrack



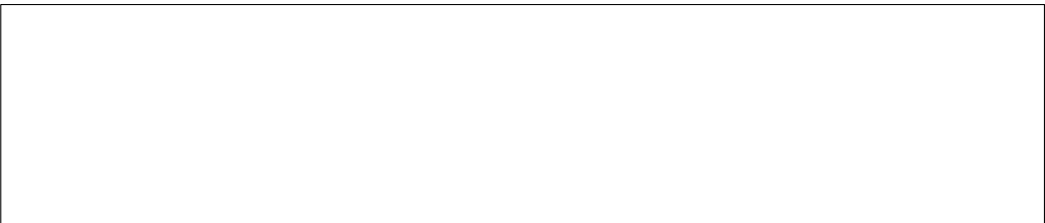
Genes



L5.6.NR4A2



Peaks



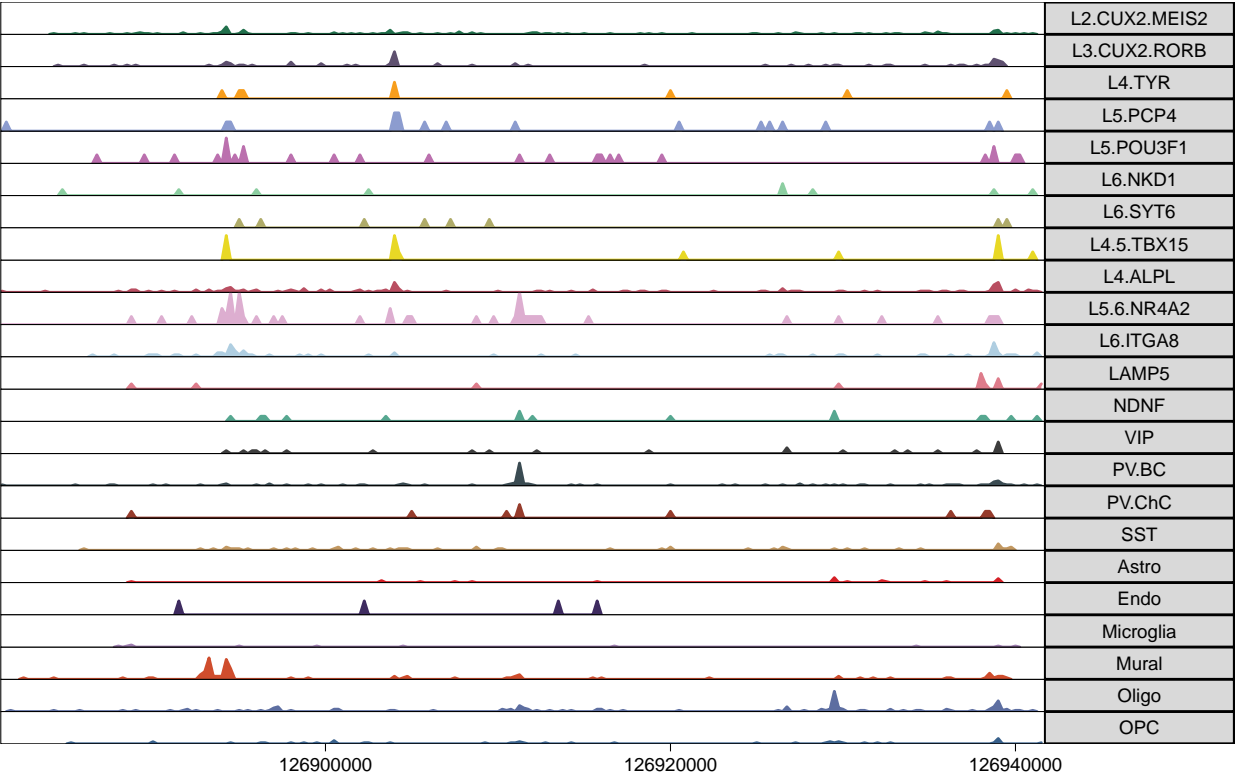
LoopTrack



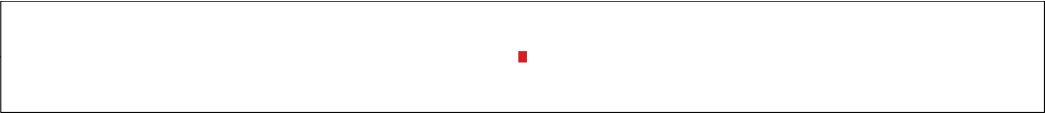
Genes

chr9:126881178–126941679 Marker Peak Tracks for: L5.6.NR4A2

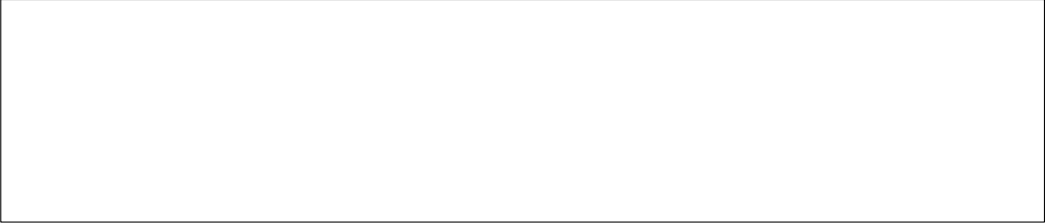
Coverage
(Norm. ATAC Signal Range (0–0.91) by ReadsInTSS)



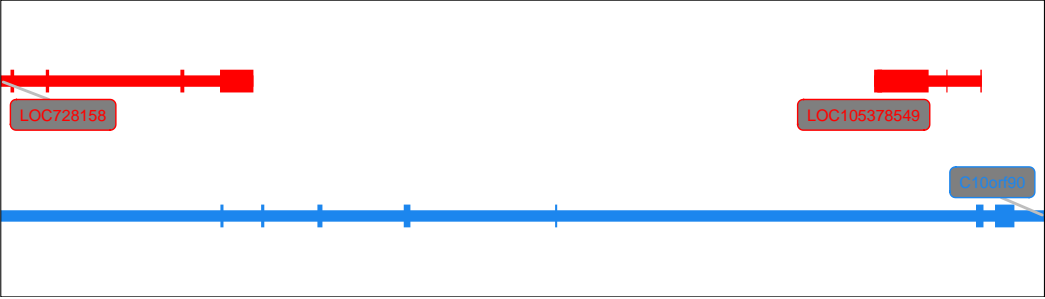
L5.6.NR4A2



Peaks



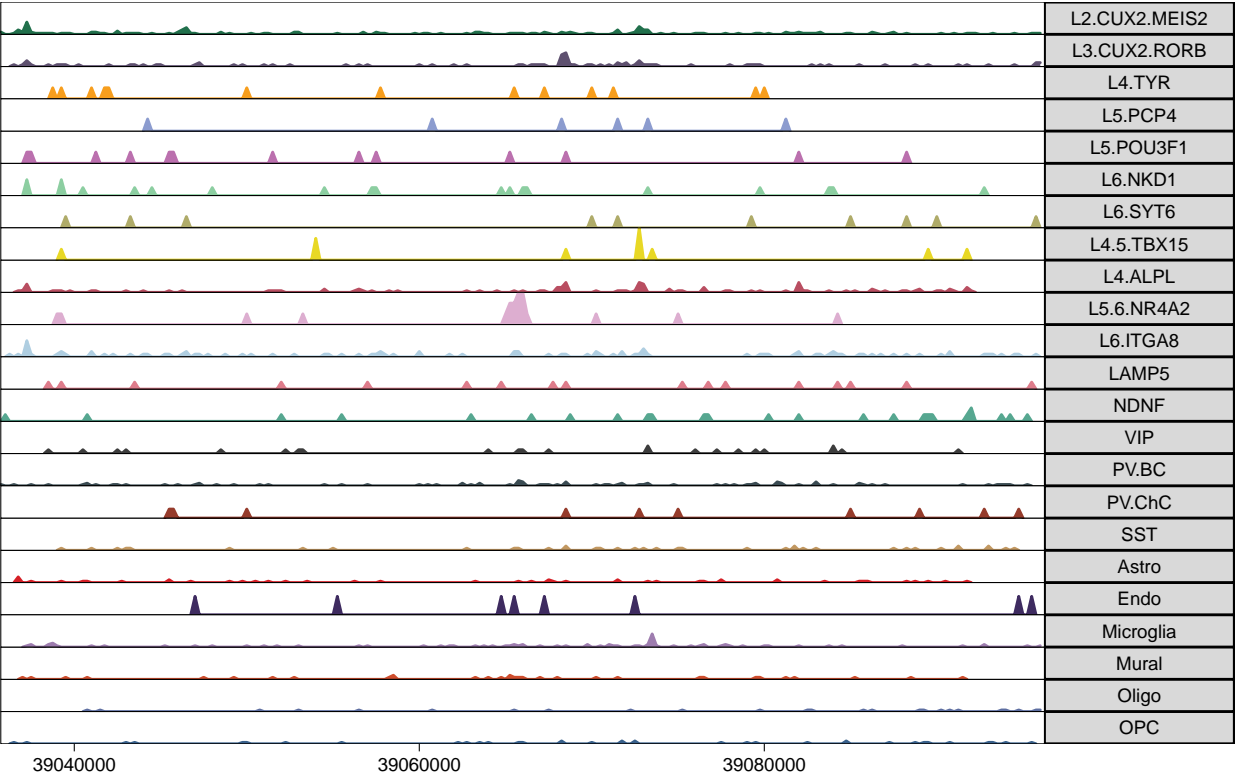
LoopTrack



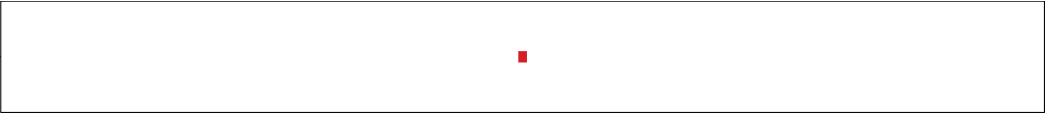
Genes

chr5:39035739–39096240 Marker Peak Tracks for: L5.6.NR4A2

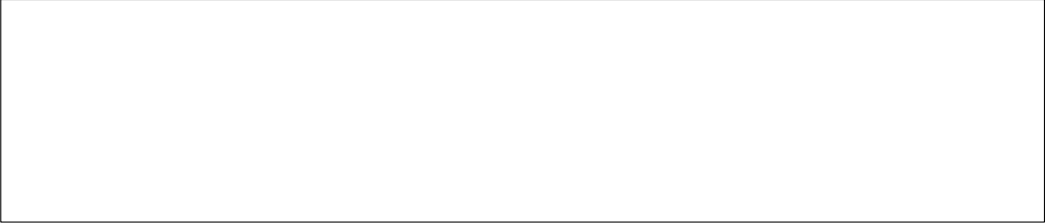
Coverage
(Norm. ATAC Signal Range 0–0.68) by ReadsInTSS)



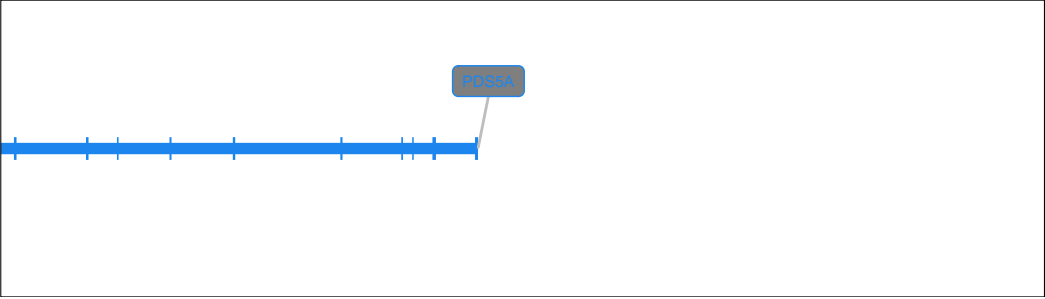
L5.6.NR4A2



Peaks

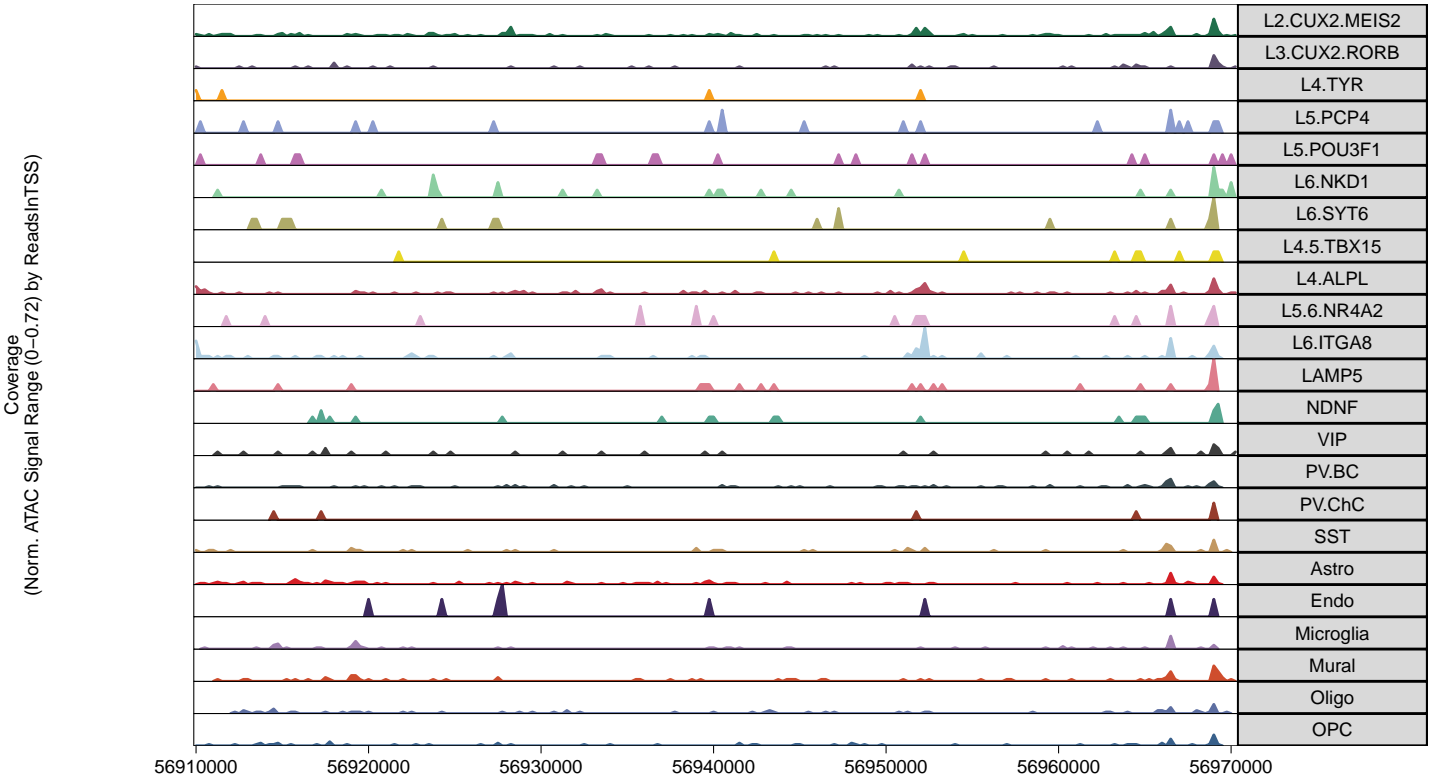


LoopTrack

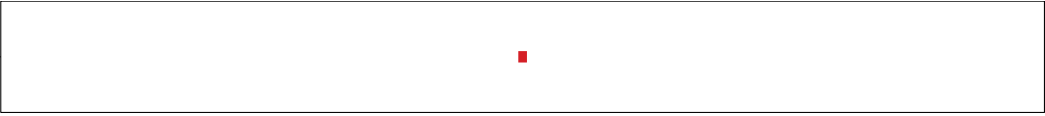


Genes

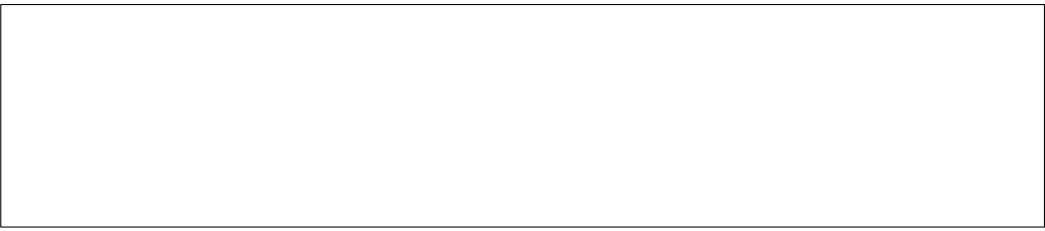
chr4:56909868–56970369 Marker Peak Tracks for: L5.6.NR4A2



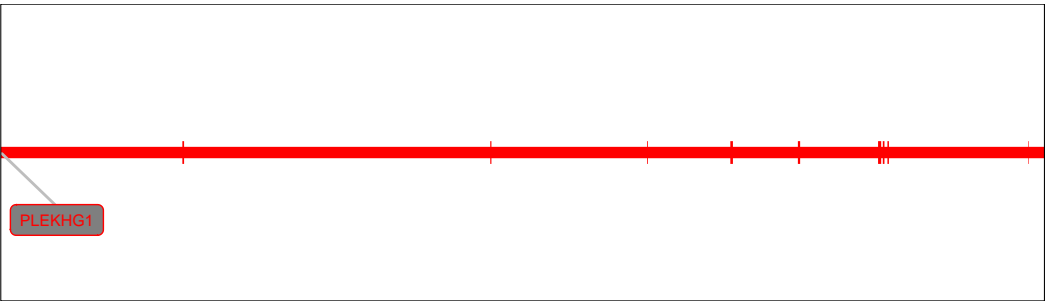
L5.6.NR4A2



Peaks



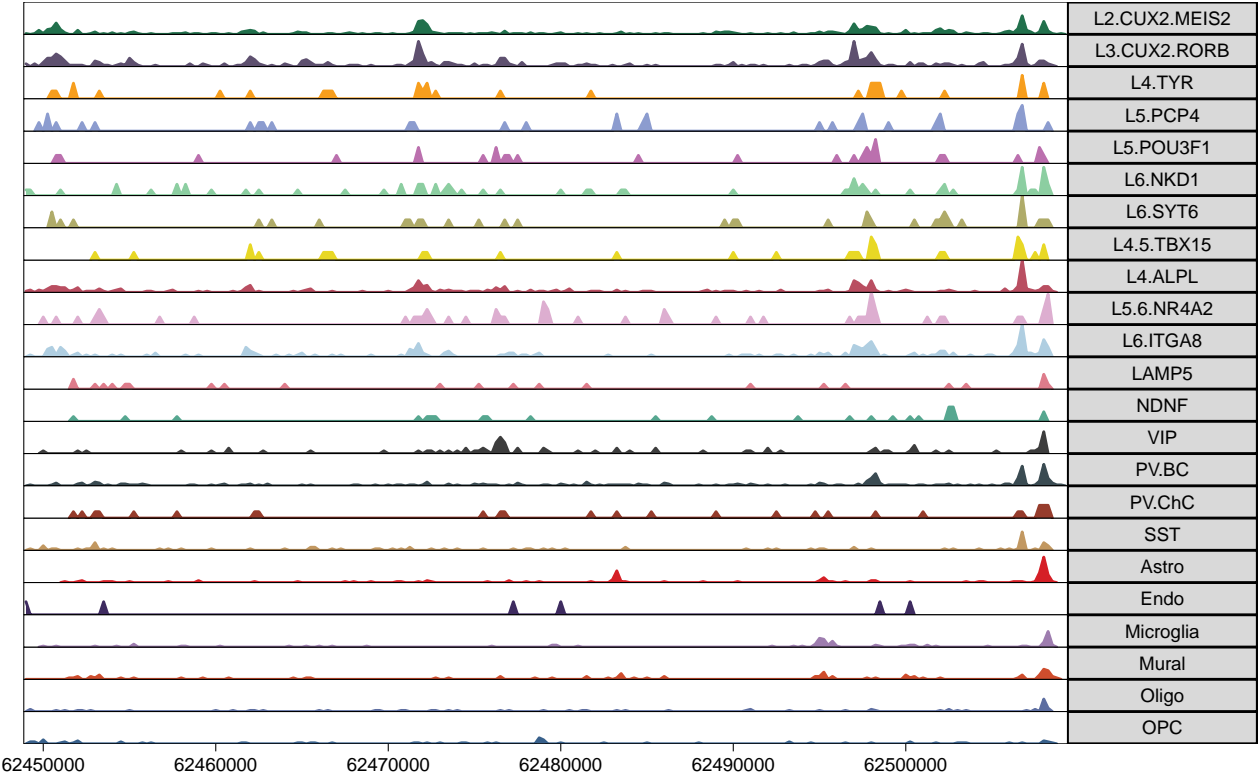
LoopTrack



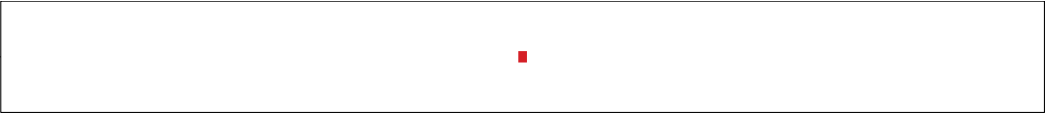
Genes

PLEKHG1

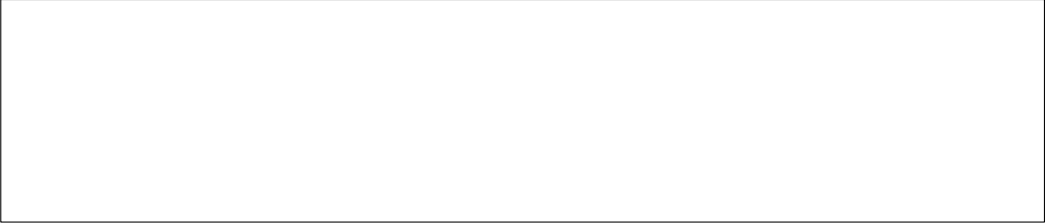
Coverage
(Norm. ATAC Signal Range 0–0.96) by ReadsInTSS)



L5.6.NR4A2



Peaks

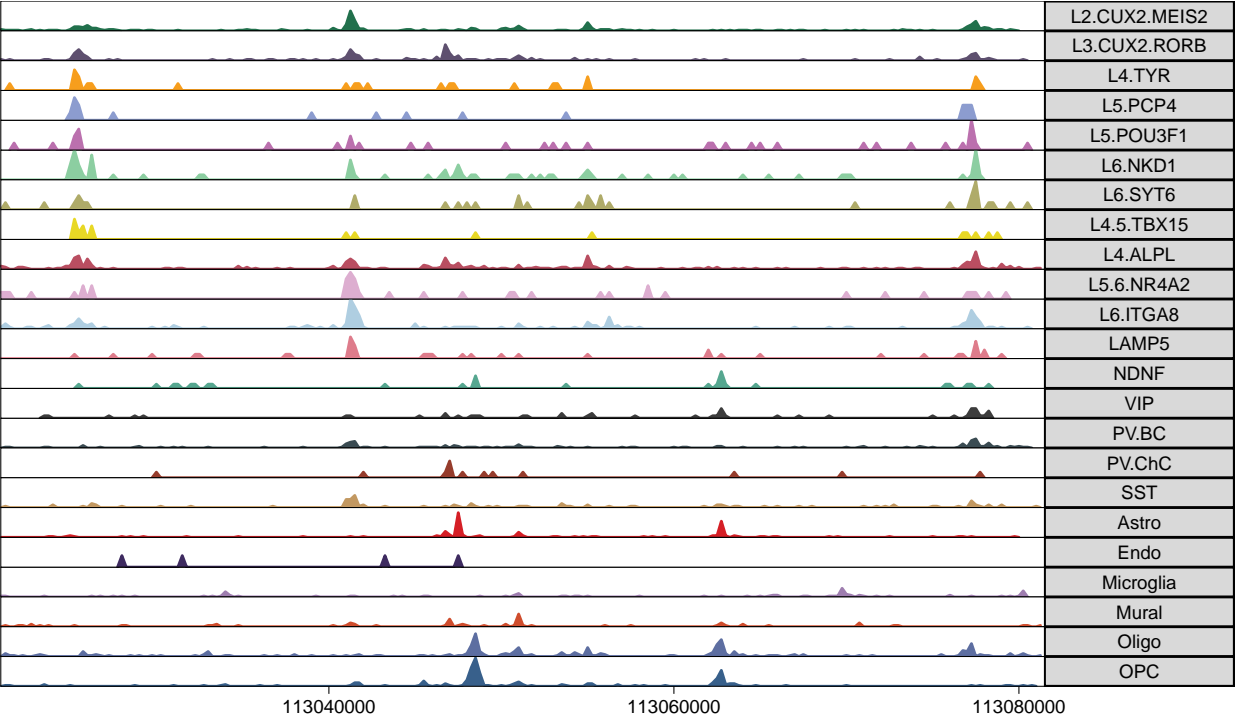


LoopTrack

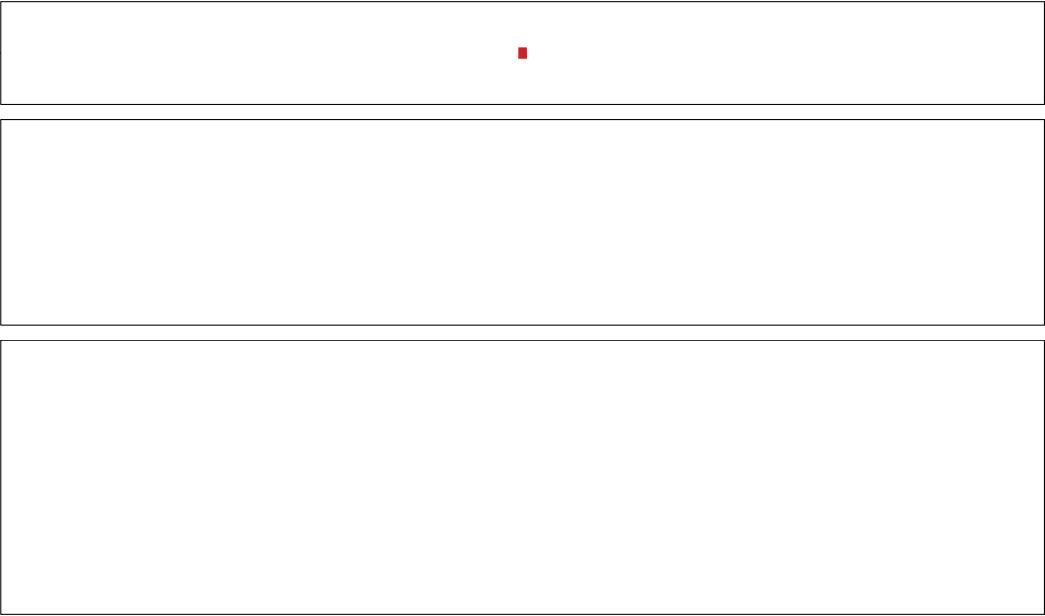


Genes

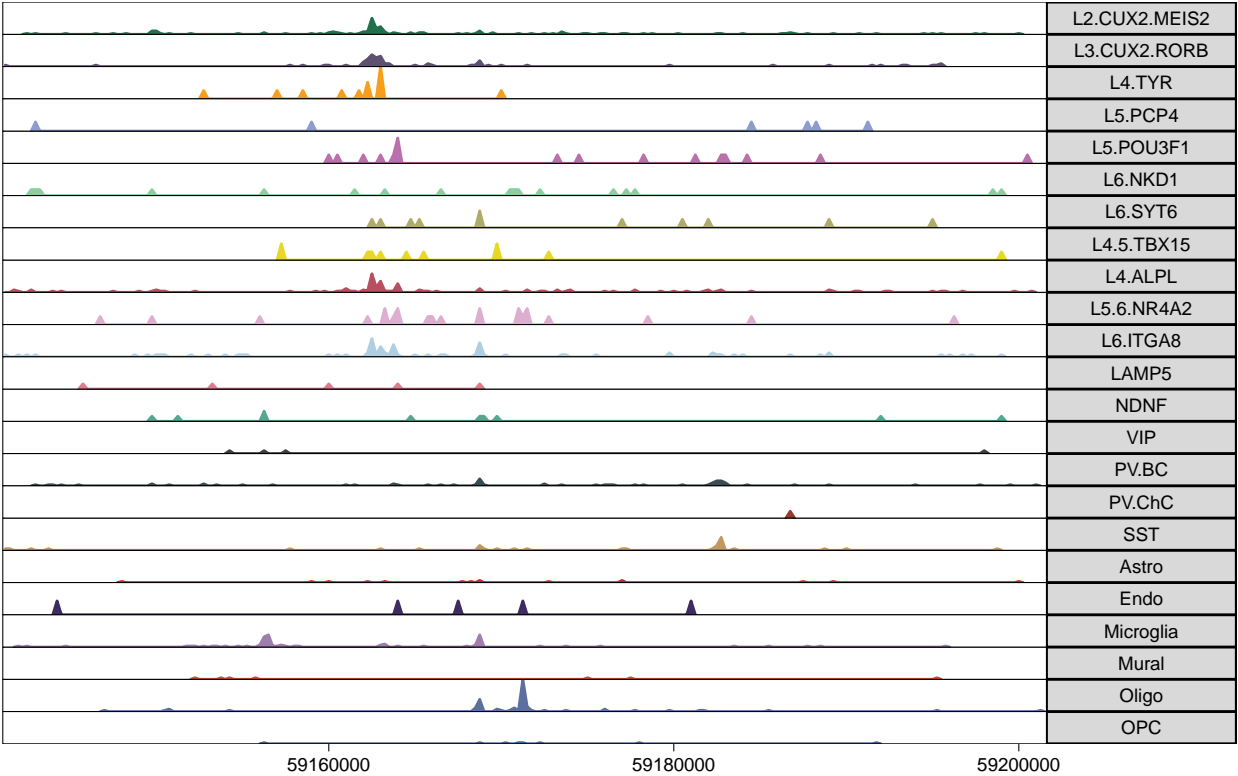
Coverage
(Norm. ATAC Signal Range (0–1.01) by Reads in TSS)



L5.6.NR4A2



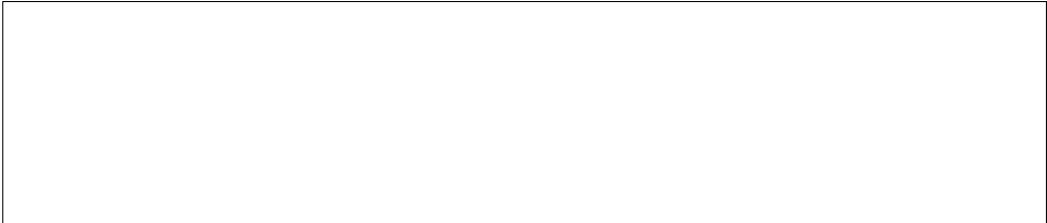
Coverage
(Norm. ATAC Signal Range (0–0.91) by ReadsInTSS)



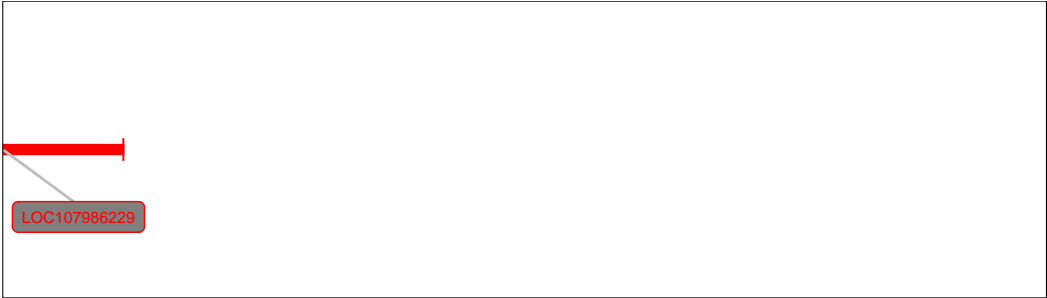
L5.6.NR4A2



Peaks

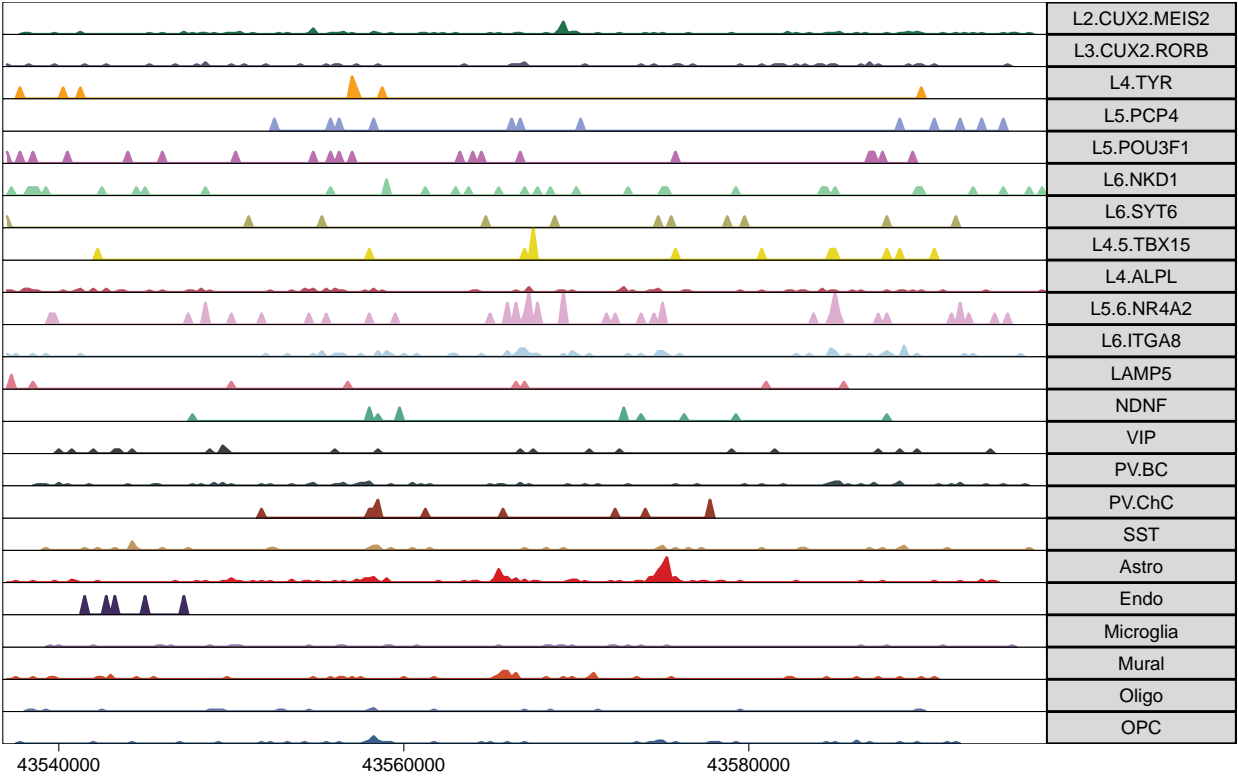


LoopTrack



Genes

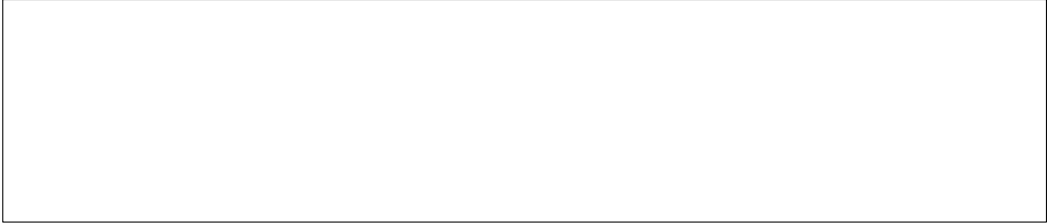
Coverage
(Norm. ATAC Signal Range 0–0.68) by ReadsInTSS)



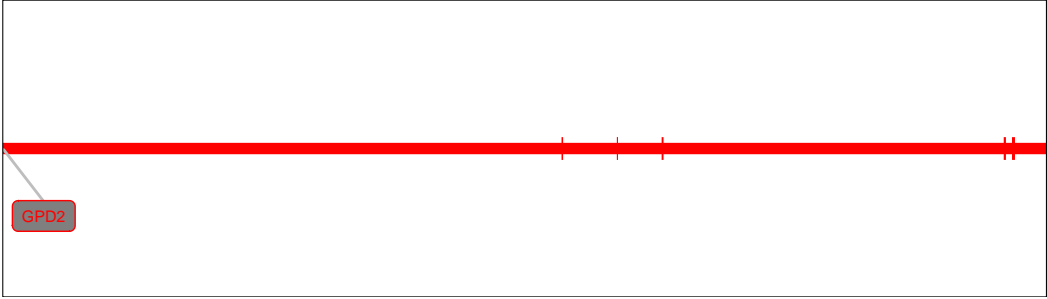
L5.6.NR4A2



Peaks

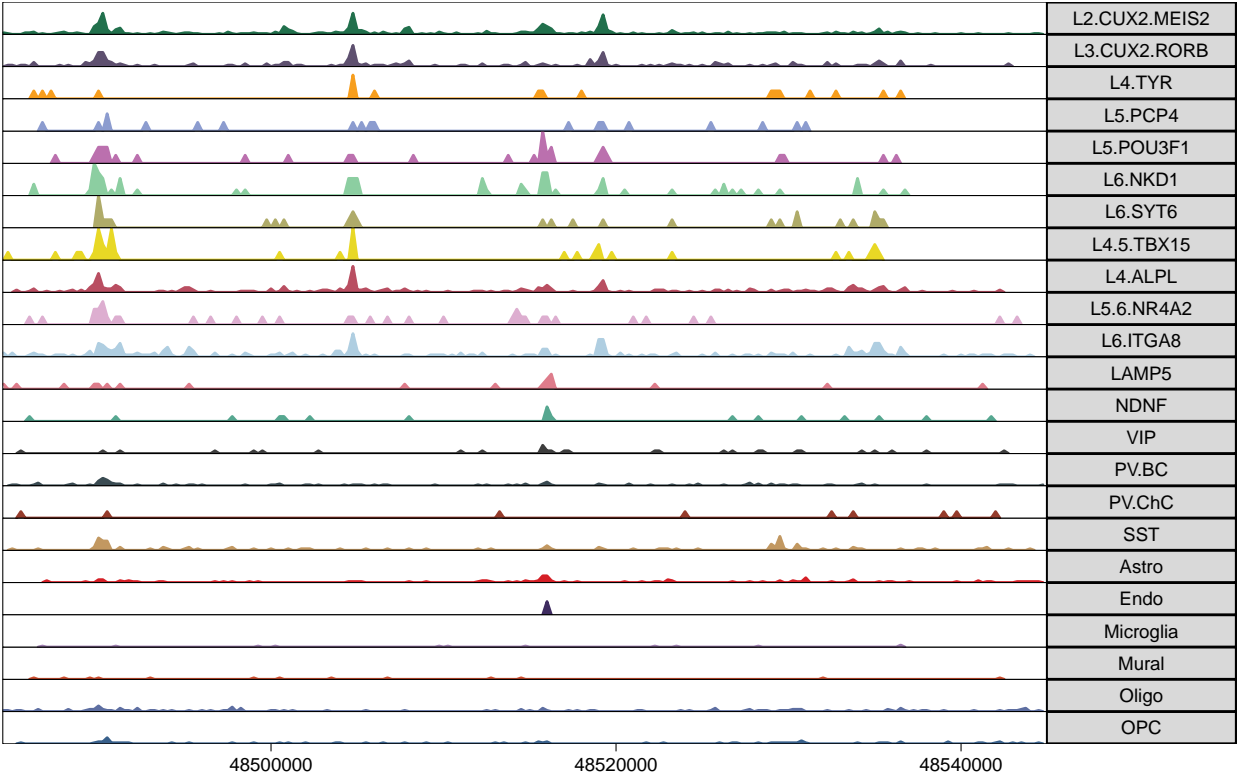


LoopTrack

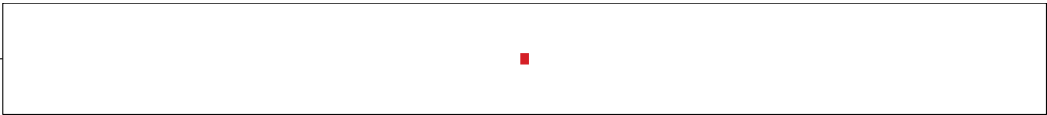


Genes

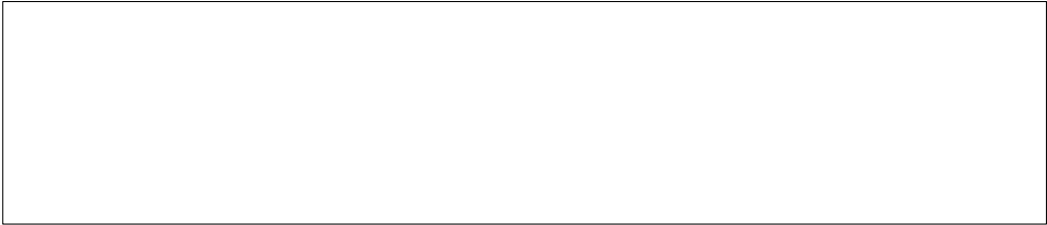
Coverage
(Norm. ATAC Signal Range (0–0.95) by ReadsInTSS)



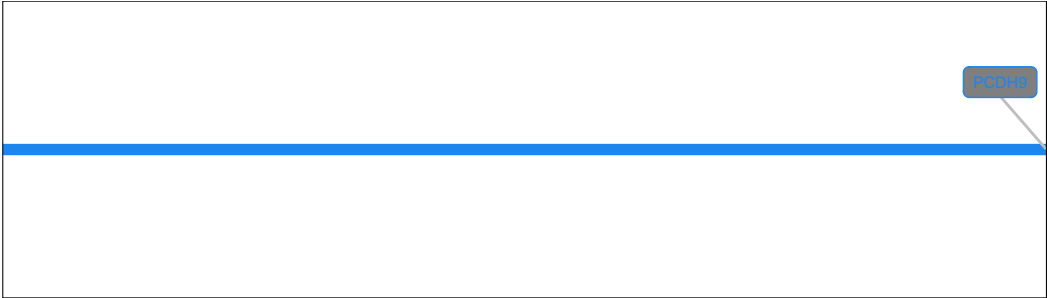
L5.6.NR4A2



Peaks



LoopTrack



Genes