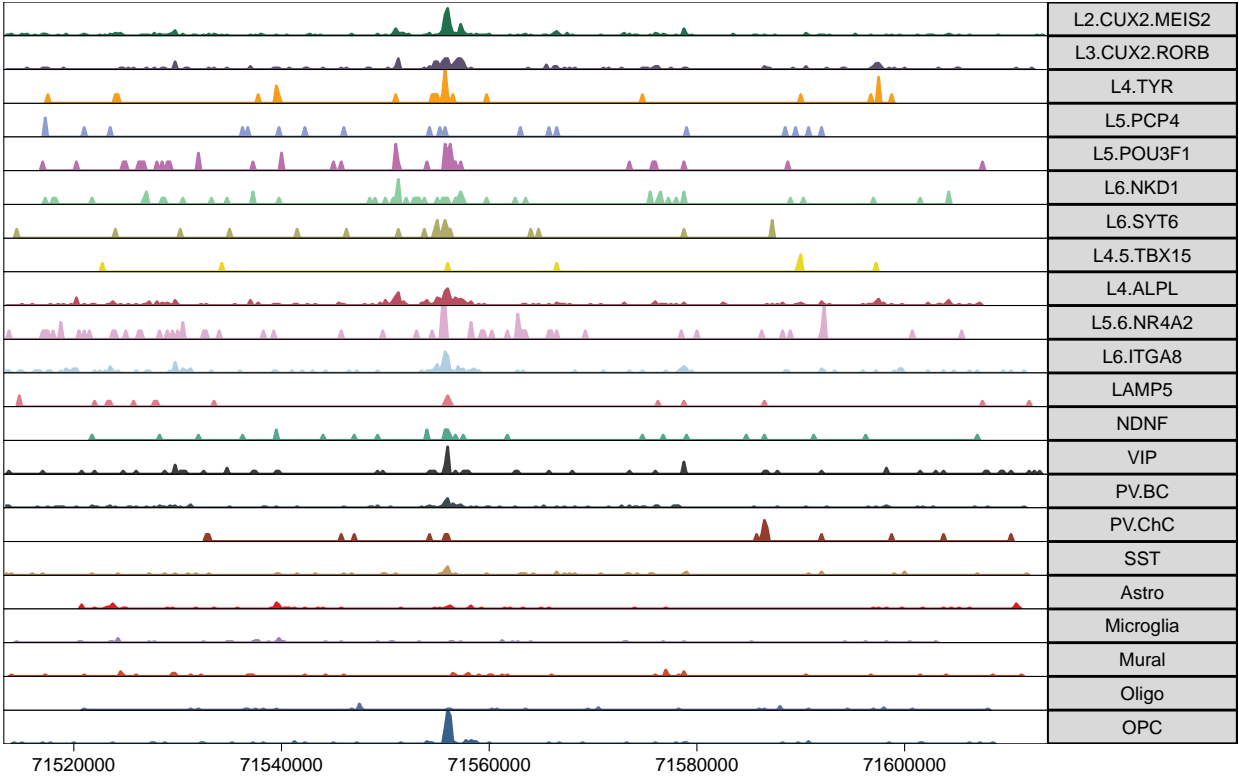
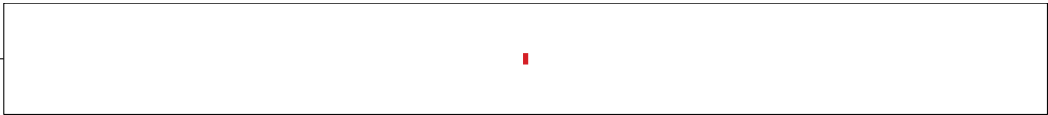


chr7:71513255–71613756 Marker Peak Tracks for: L5.6.NR4A2

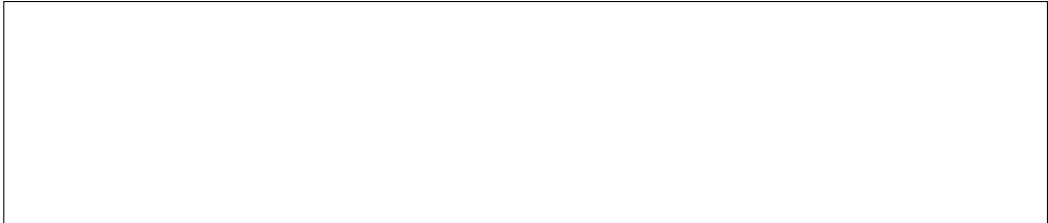
Coverage
(Norm. ATAC Signal Range (0–0.91) by ReadsInTSS)



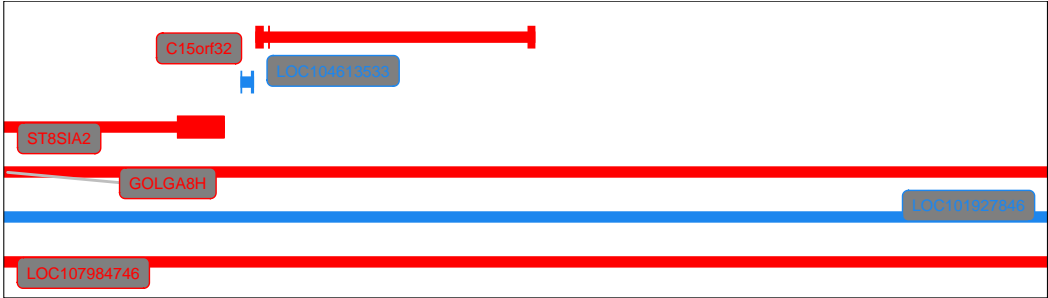
L5.6.NR4A2



Peaks

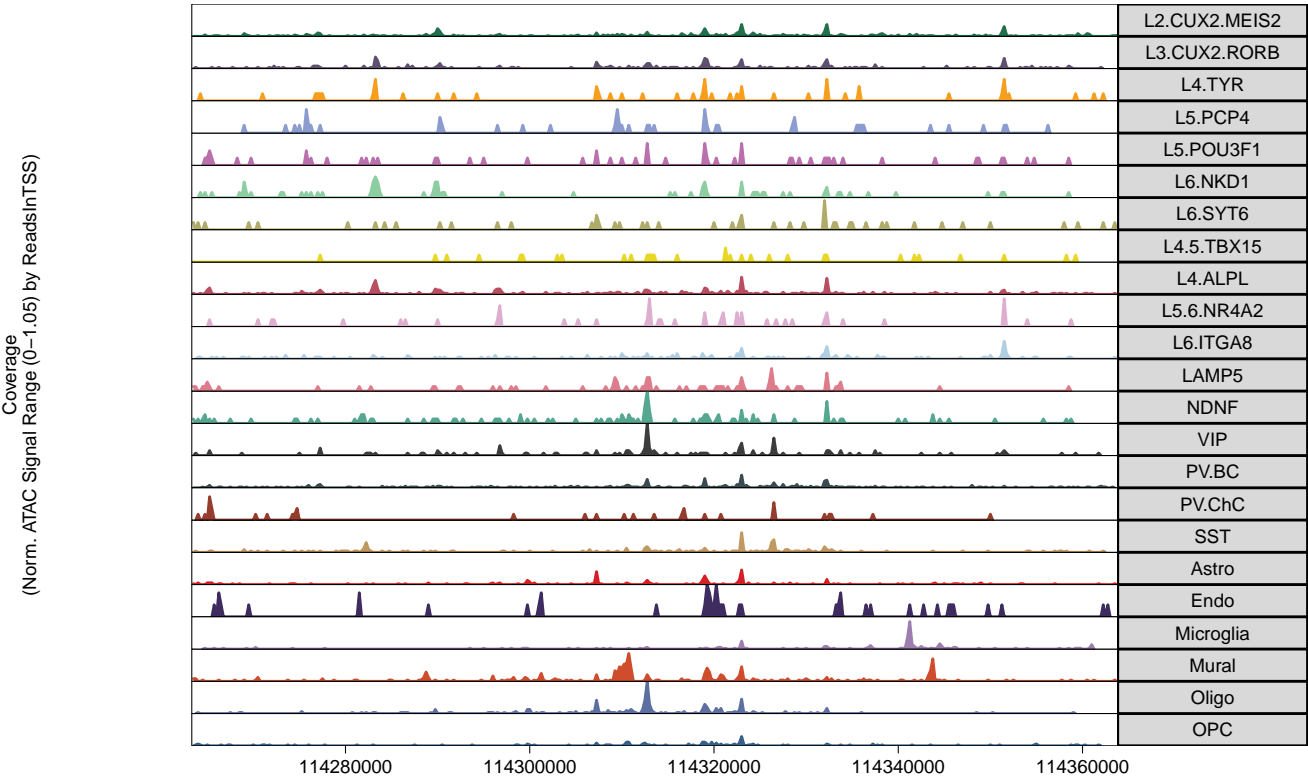


LoopTrack

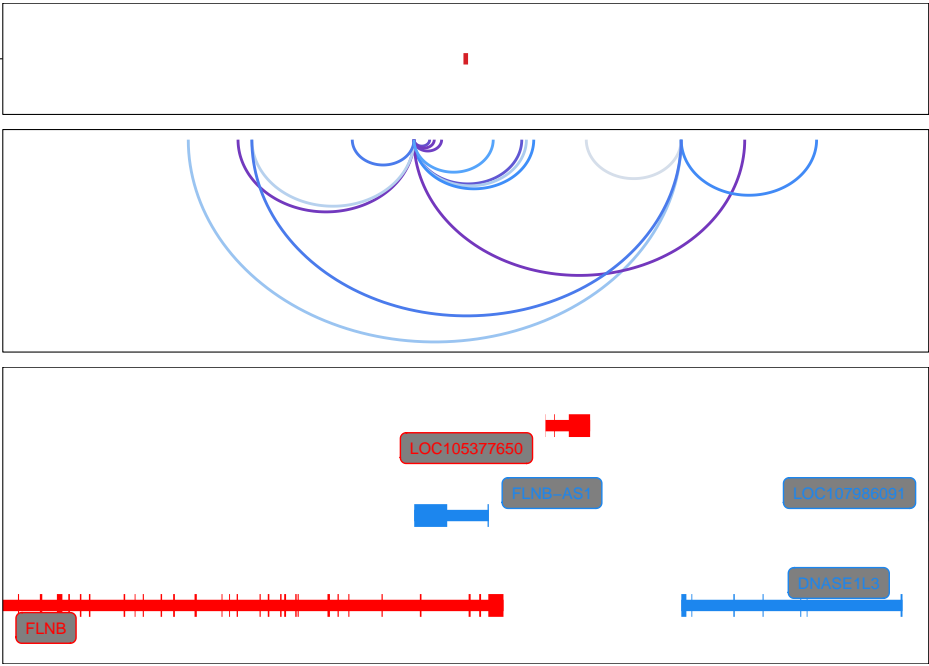


Genes

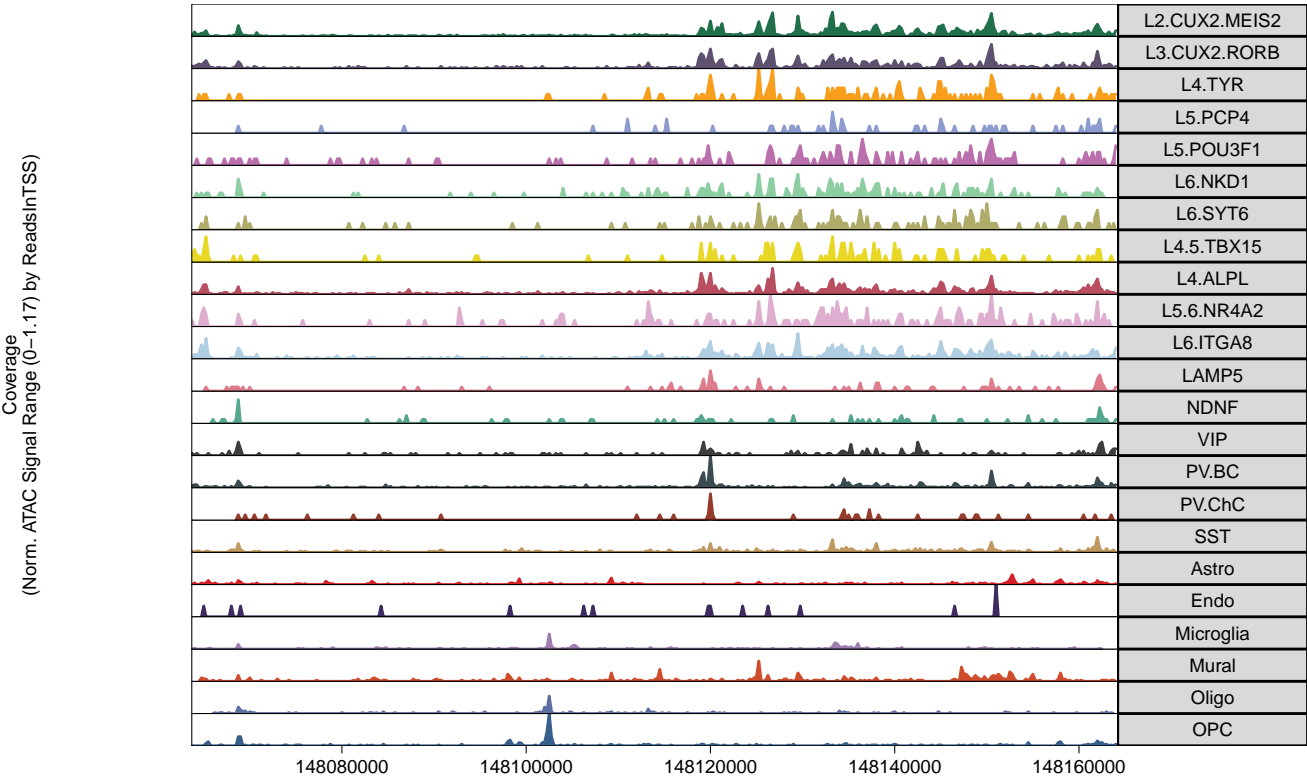
chr2:114263318–114363819 Marker Peak Tracks for: L5.6.NR4A2



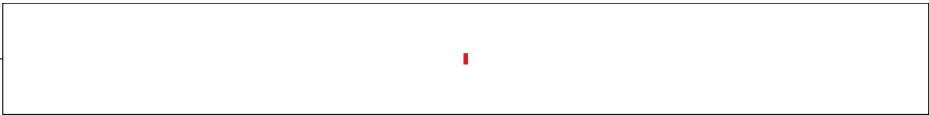
L5.6.NR4A2



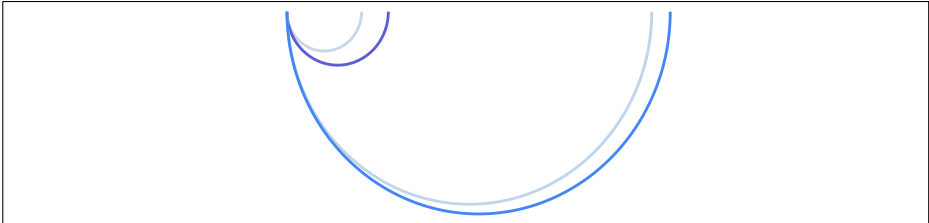
chr6:148063704–148164205 Marker Peak Tracks for: L5.6.NR4A2



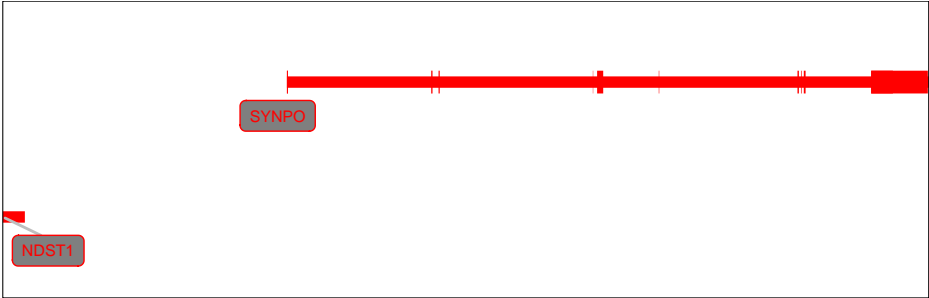
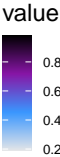
L5.6.NR4A2



Peaks

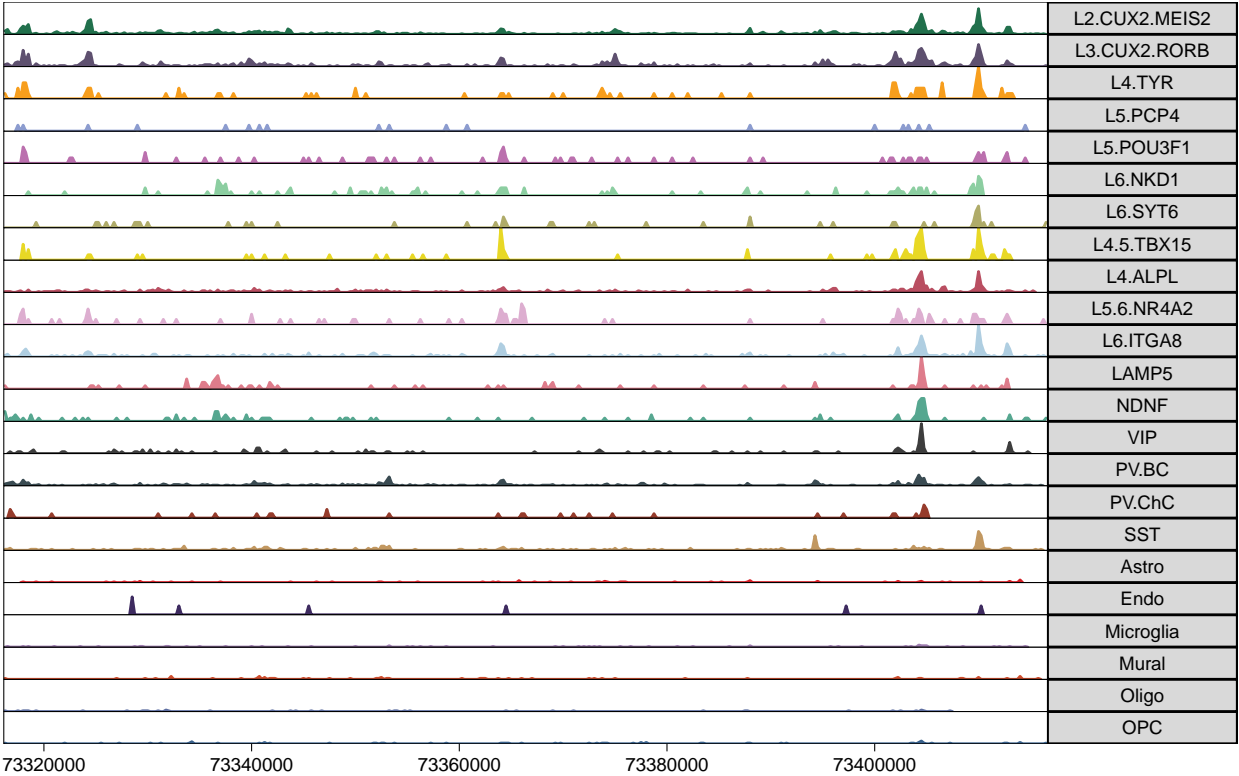


Peak2GeneLinks



Genes

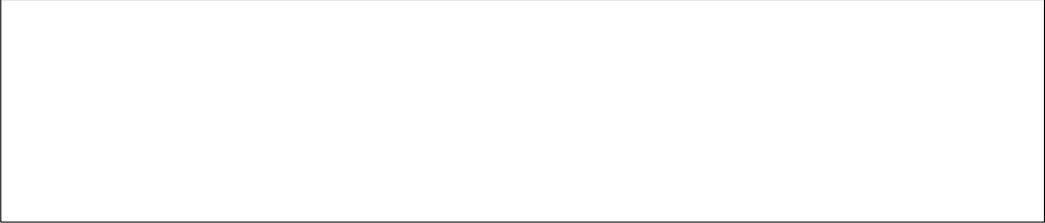
Coverage
(Norm. ATAC Signal Range 0-1.41) by ReadsInTSS)



L5.6.NR4A2



Peaks

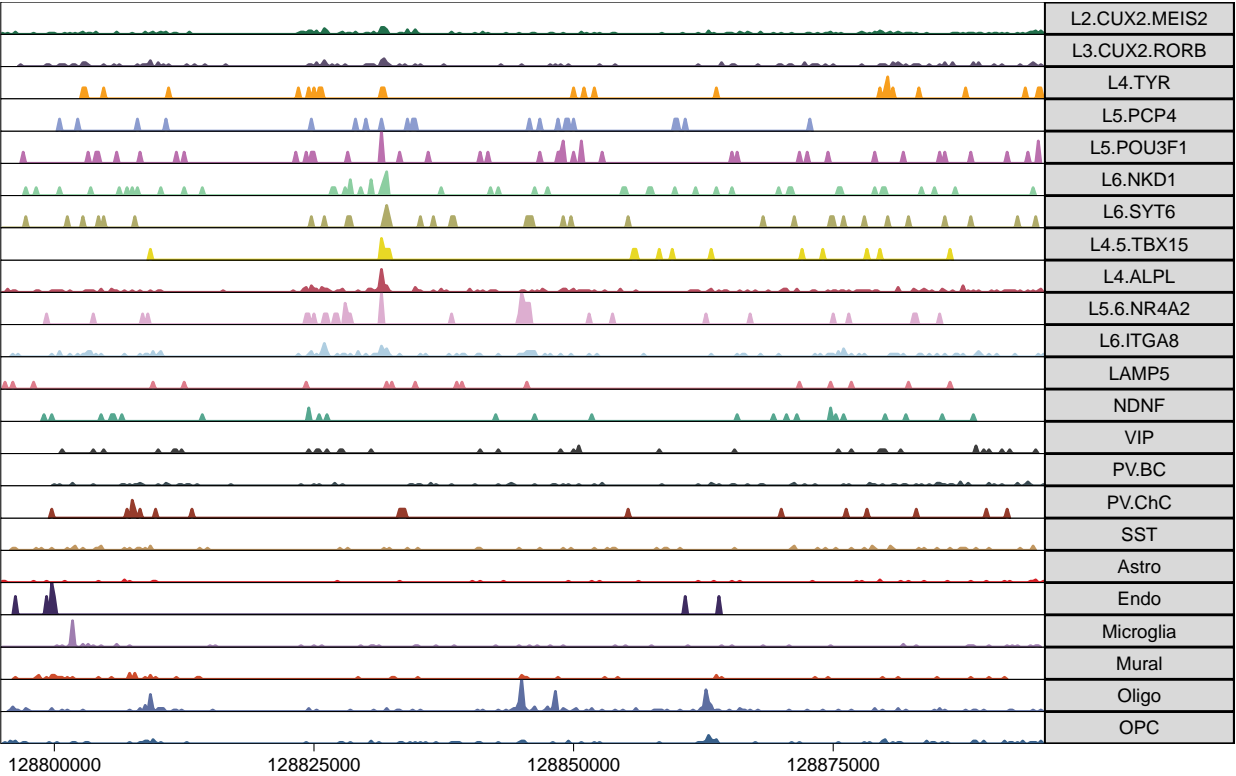


LoopTrack

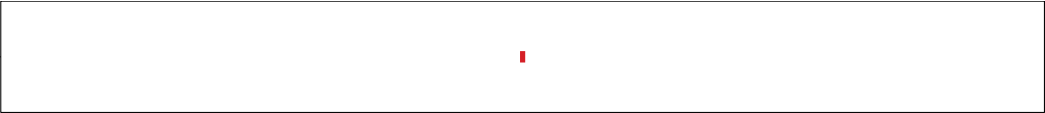


Genes

Coverage
(Norm. ATAC Signal Range 0–0.69) by ReadsInTSS)



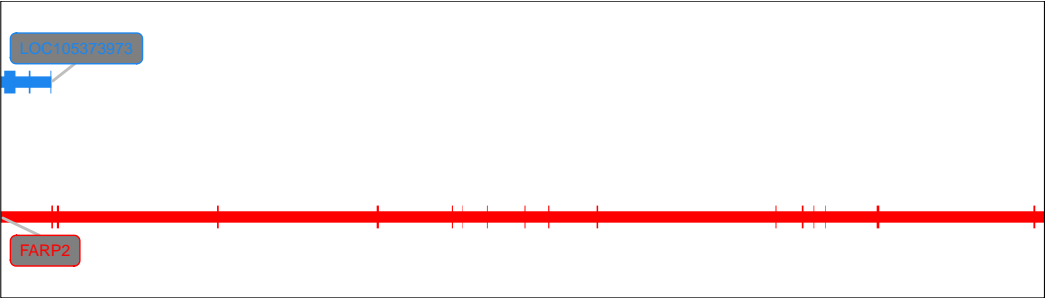
L5.6.NR4A2



Peaks

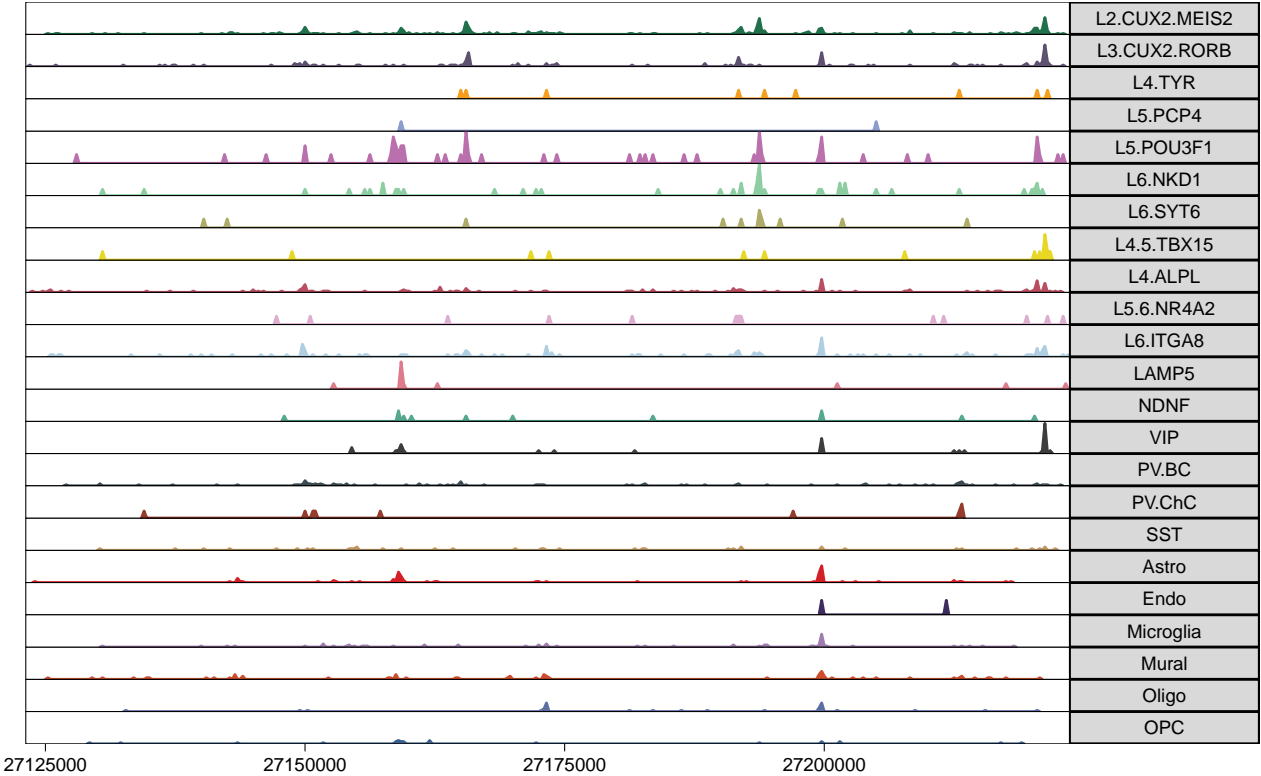


LoopTrack

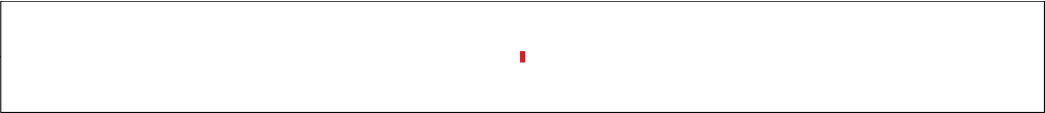


Genes

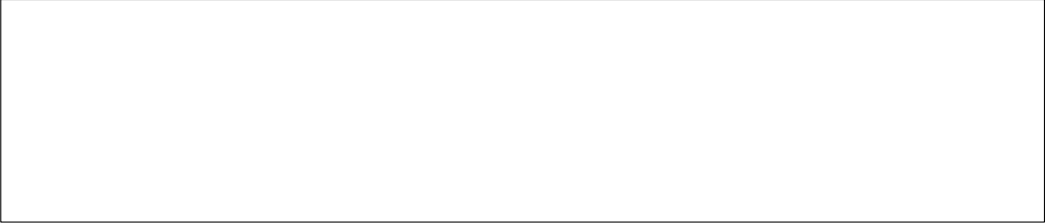
Coverage
(Norm. ATAC Signal Range 0–0.87) by ReadsInTSS)



L5.6.NR4A2



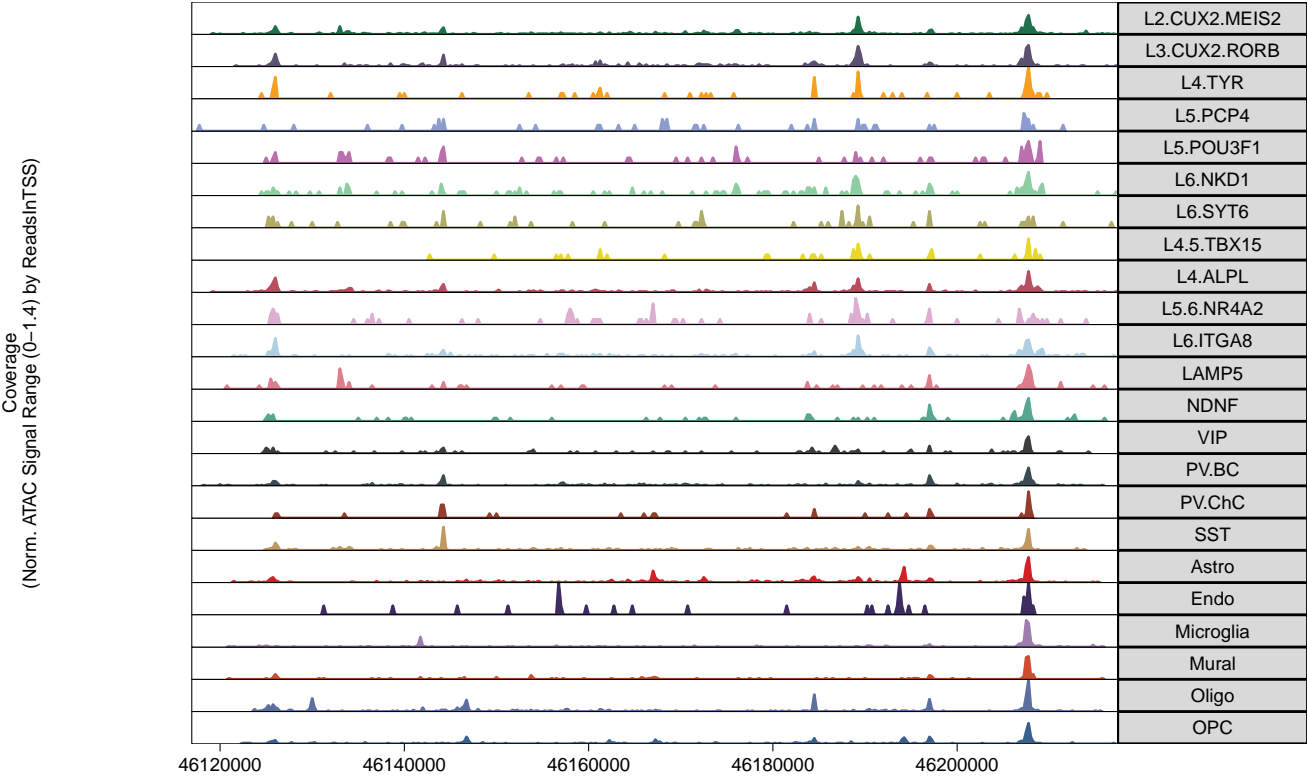
Peaks



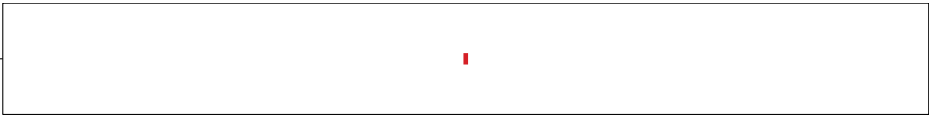
LoopTrack



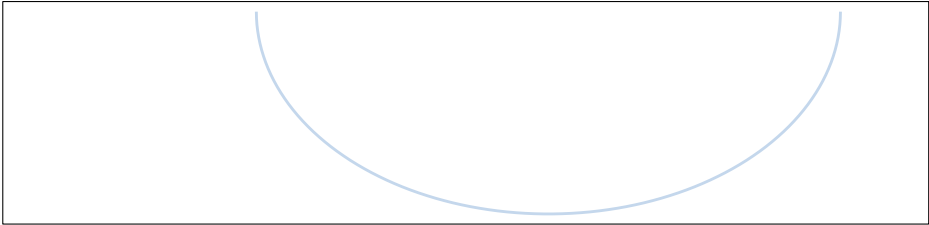
Genes



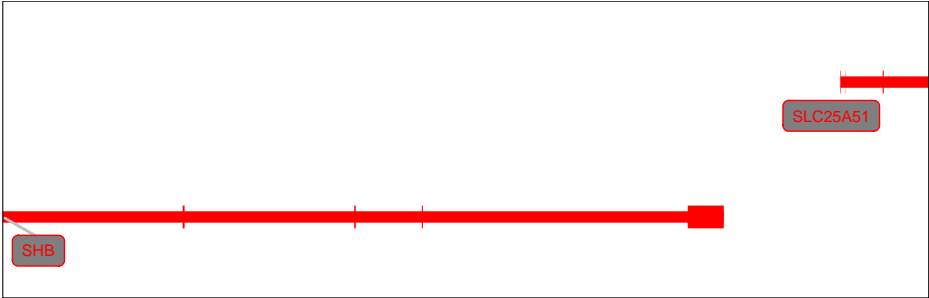
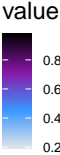
L5.6.NR4A2



Peaks

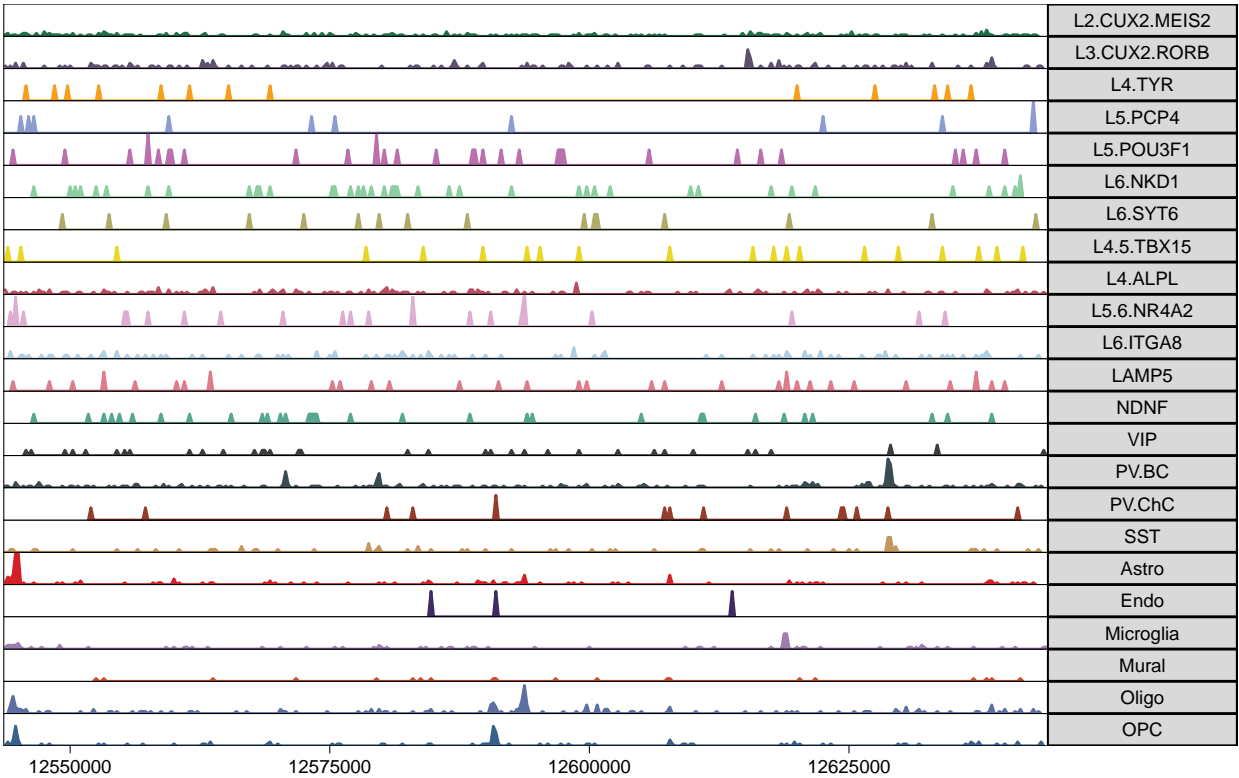


Peak2GeneLinks



Genes

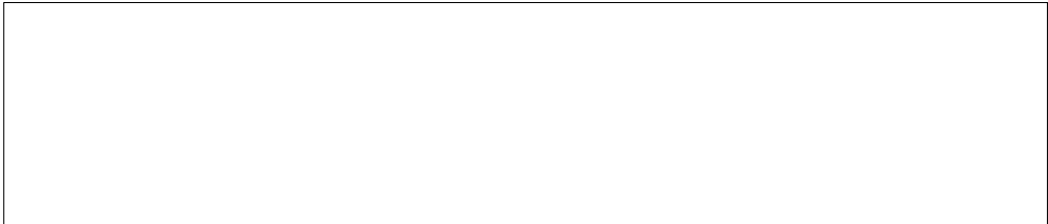
Coverage
(Norm. ATAC Signal Range (0–0.5) by Reads in TSS)



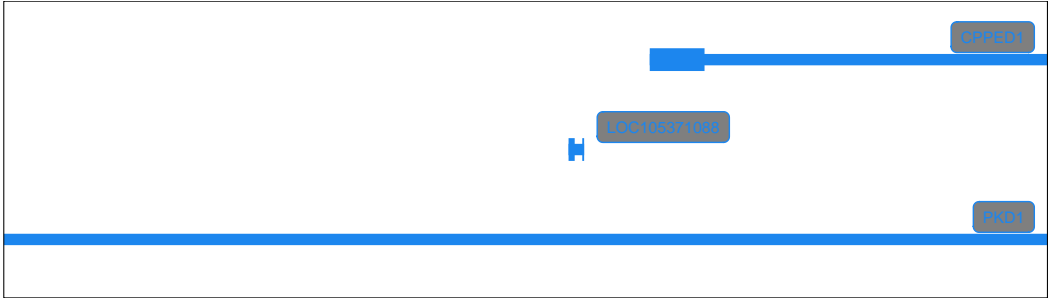
L5.6.NR4A2



Peaks

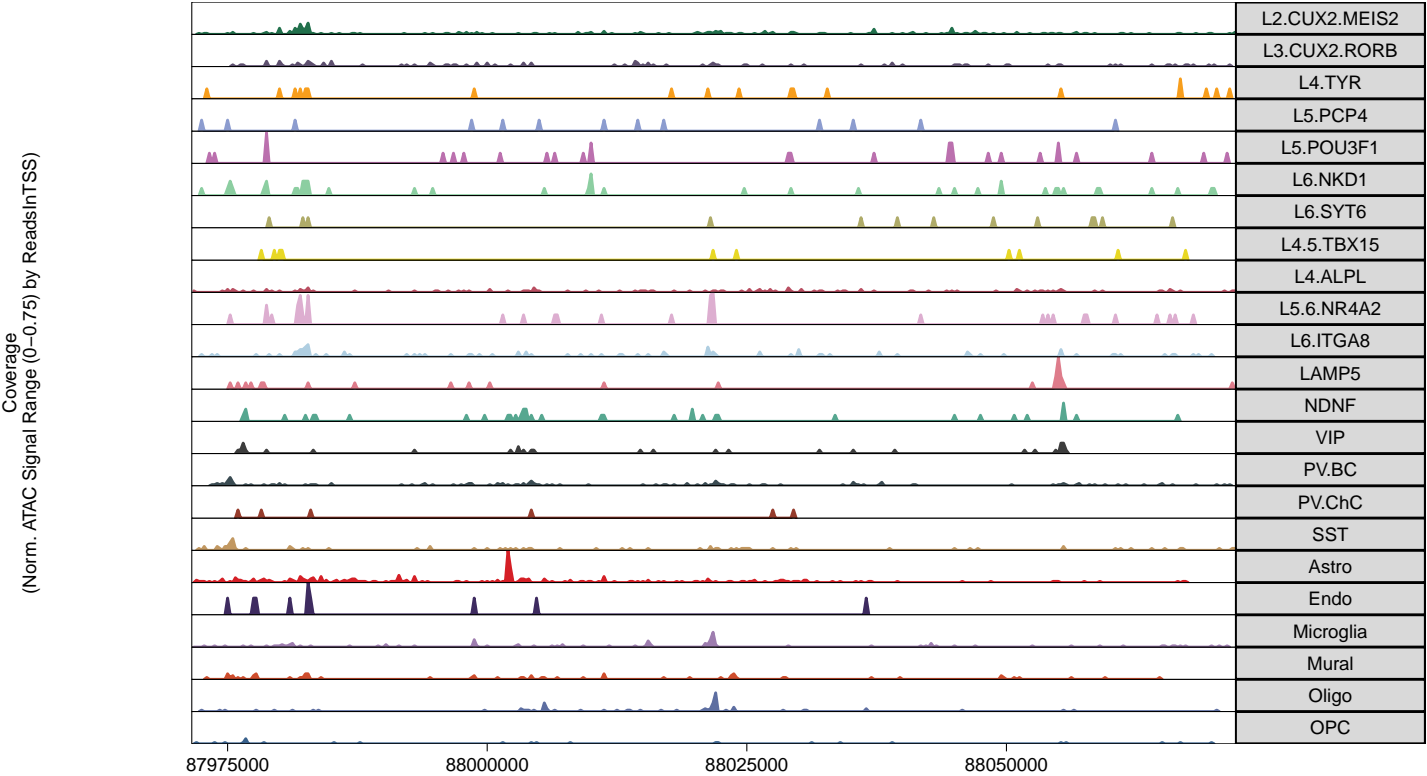


LoopTrack

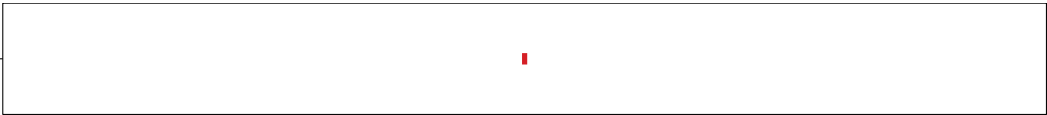


Genes

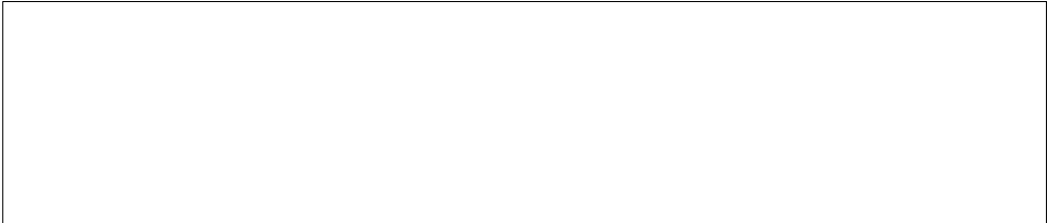
chr6:87971552–88072053 Marker Peak Tracks for: L5.6.NR4A2



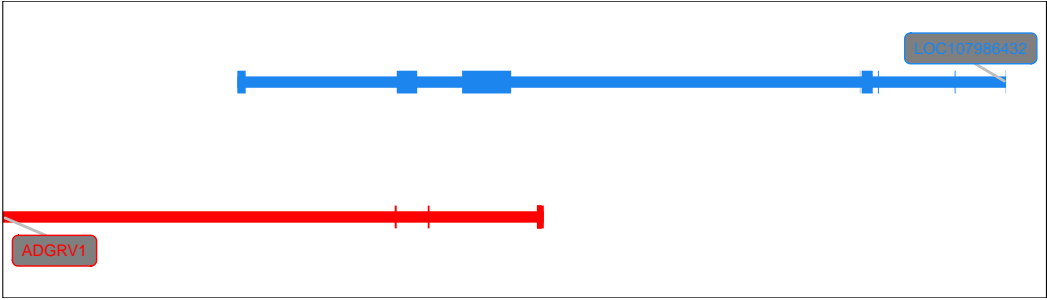
L5.6.NR4A2



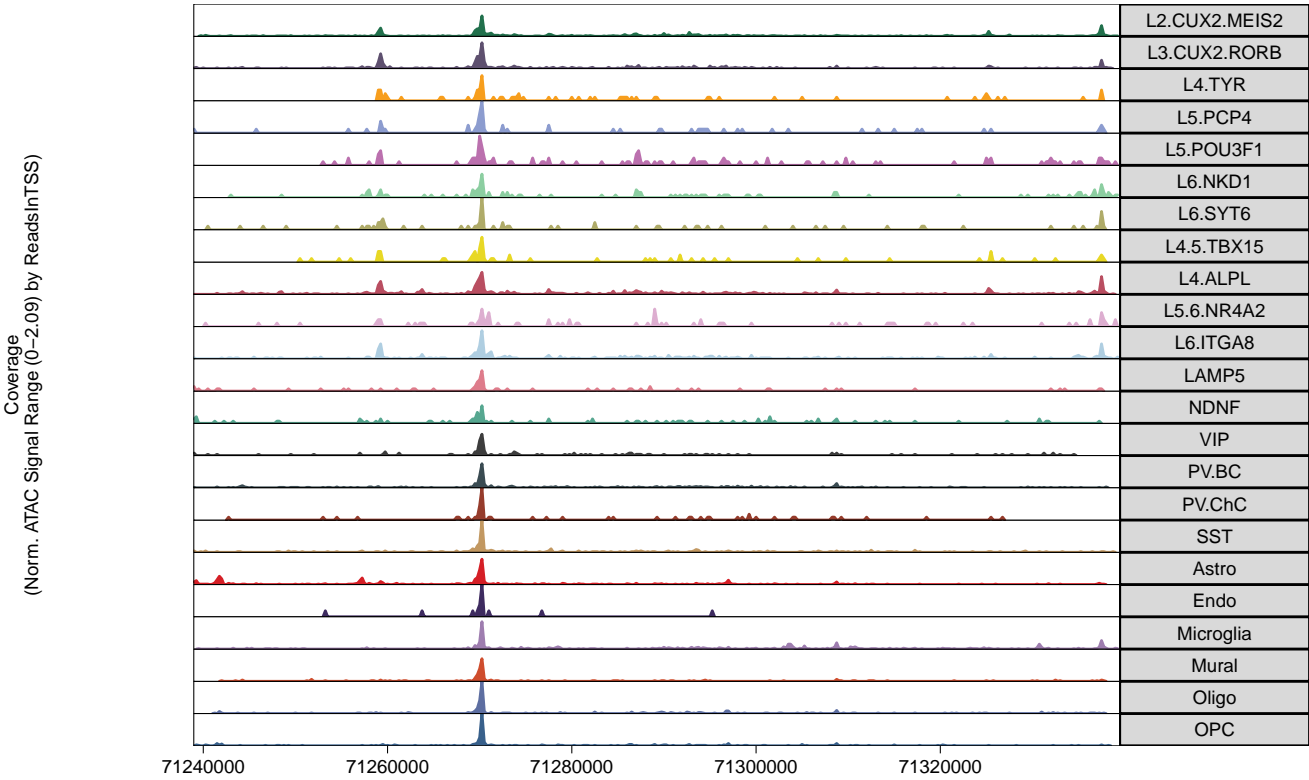
Peaks



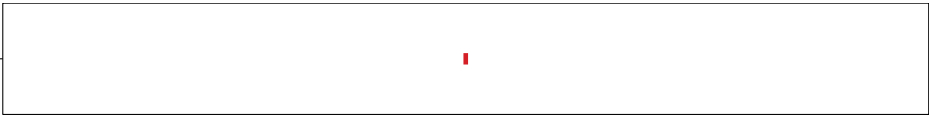
LoopTrack



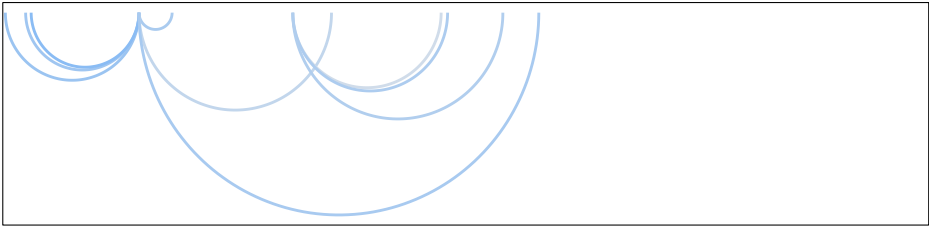
Genes



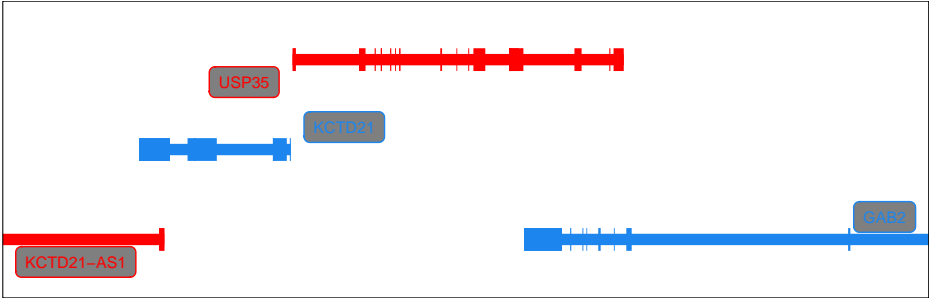
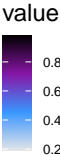
L5.6.NR4A2



Peaks

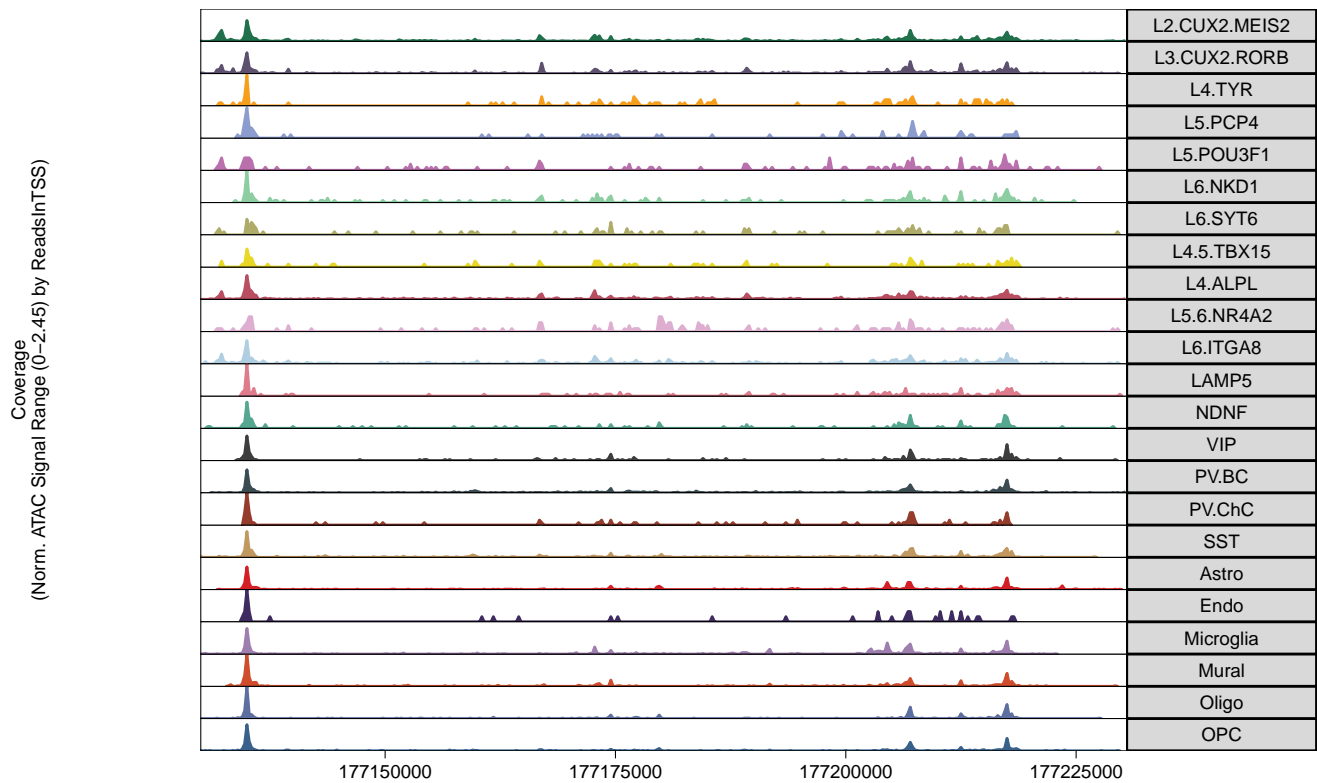


Peak2GeneLinks

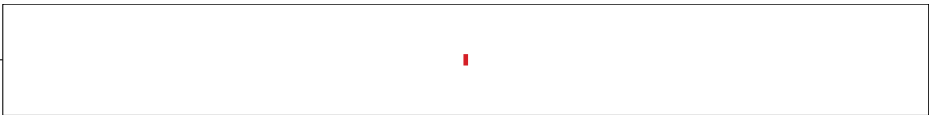


Genes

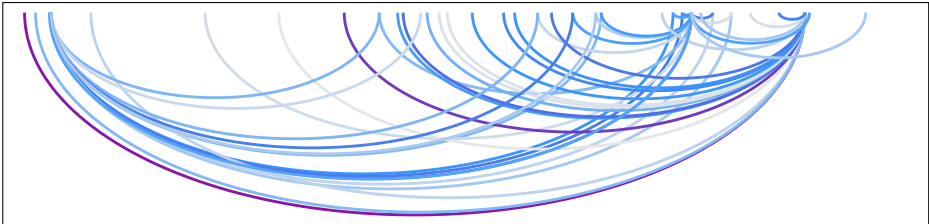
chr6:177129991-177230492 Marker Peak Tracks for: L5.6.NR4A2



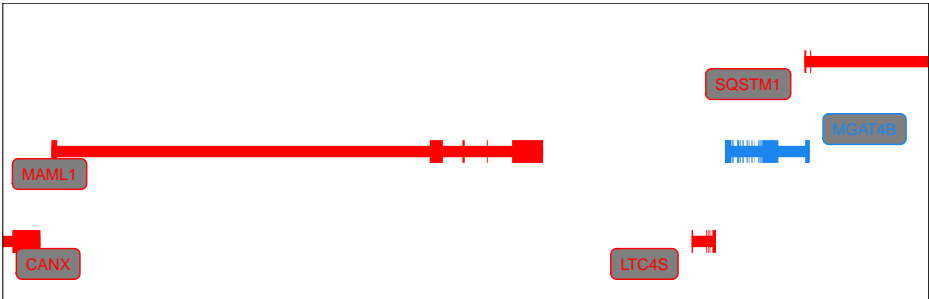
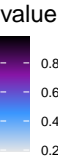
L5.6.NR4A2



Peaks



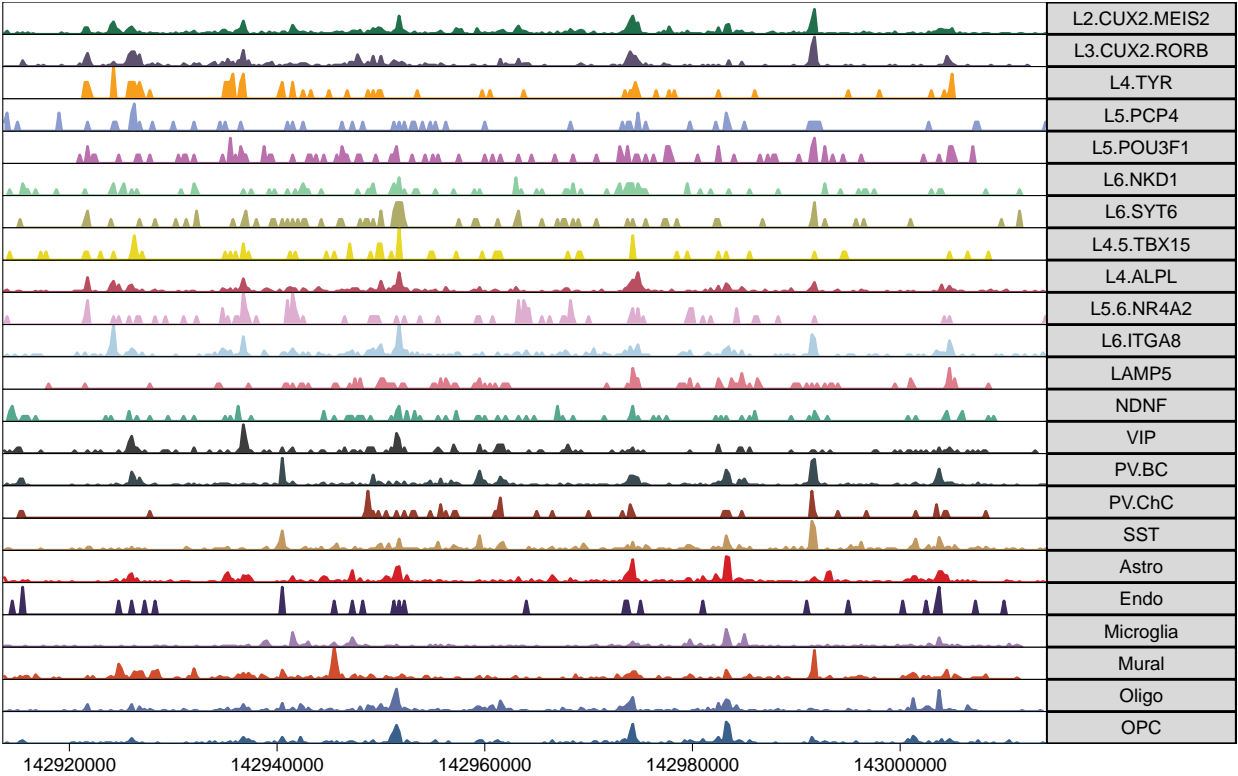
Peak2GeneLinks



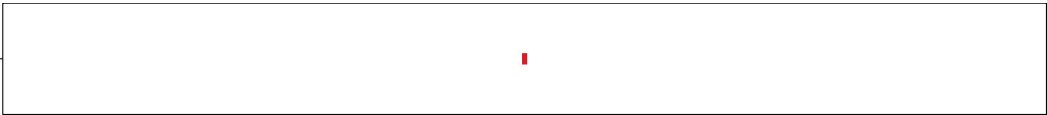
Genes

chr2:142913586–143014087 Marker Peak Tracks for: L5.6.NR4A2

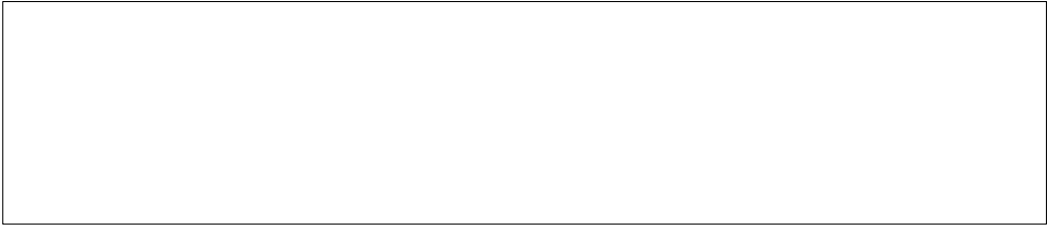
Coverage
(Norm. ATAC Signal Range (0–0.91) by ReadsInTSS)



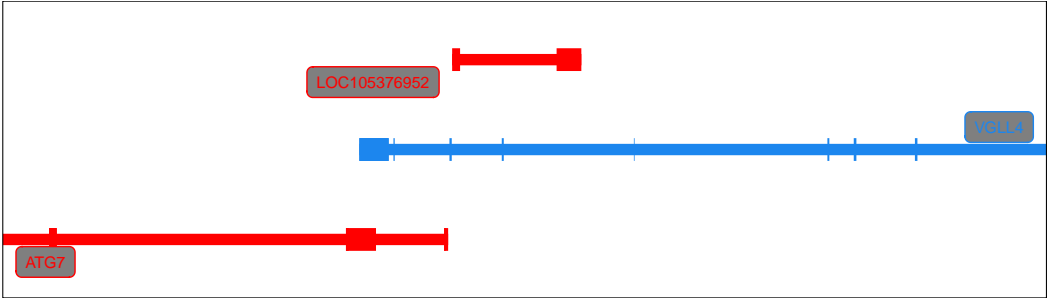
L5.6.NR4A2



Peaks



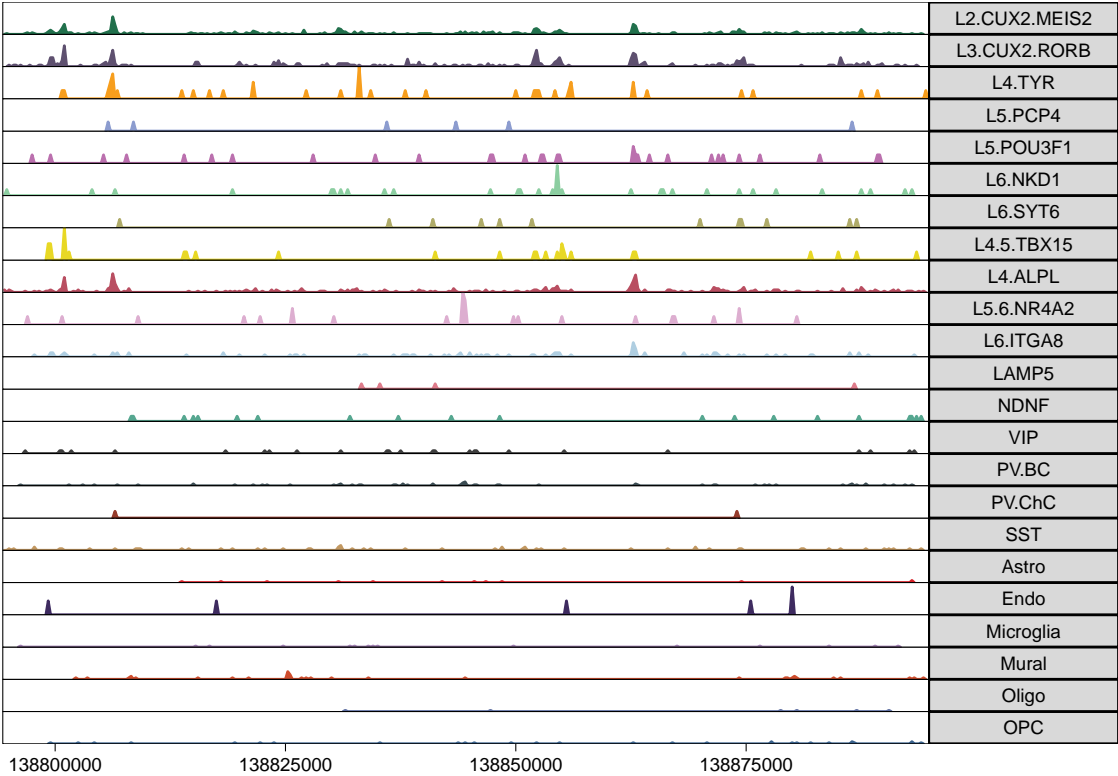
LoopTrack



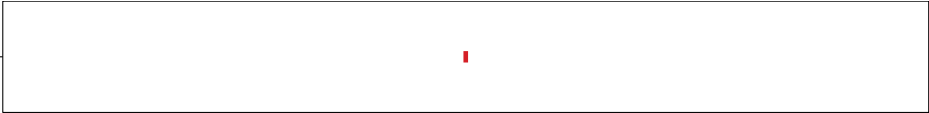
Genes

chr2:138794317-138894818 Marker Peak Tracks for: L5.6.NR4A2

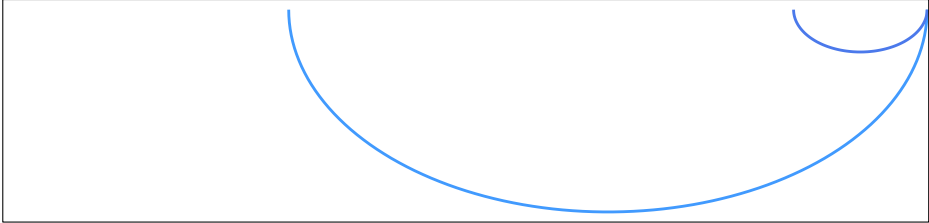
Coverage
(Norm. ATAC Signal Range (0-0.91) by ReadsInTSS)



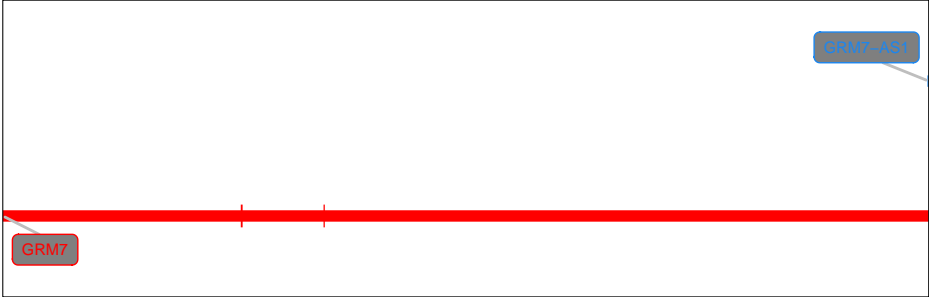
L5.6.NR4A2



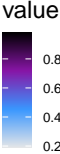
Peaks



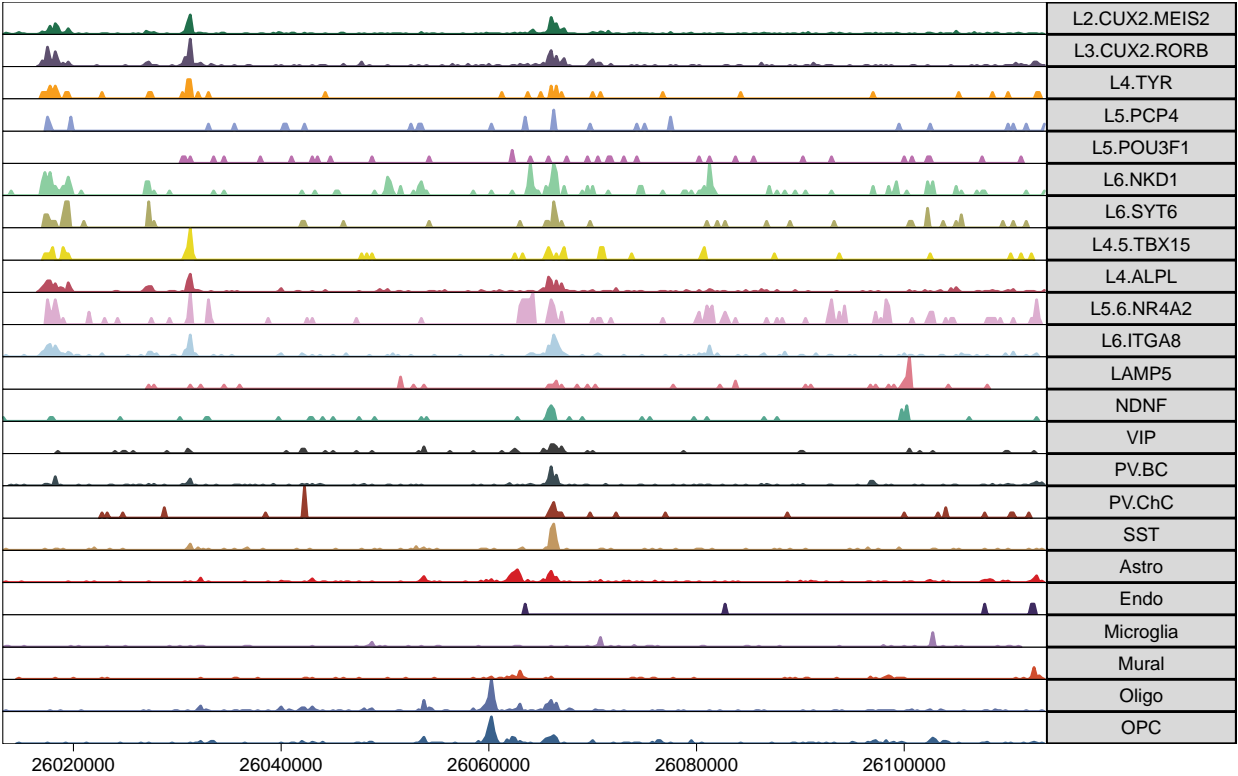
Peak2GeneLinks



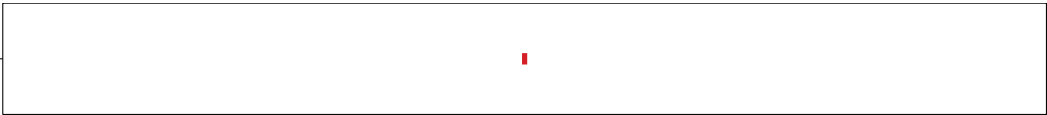
Genes



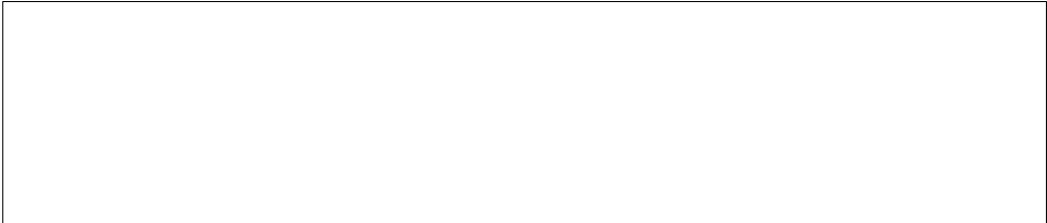
Coverage
(Norm. ATAC Signal Range 0–1.18) by ReadsInTSS)



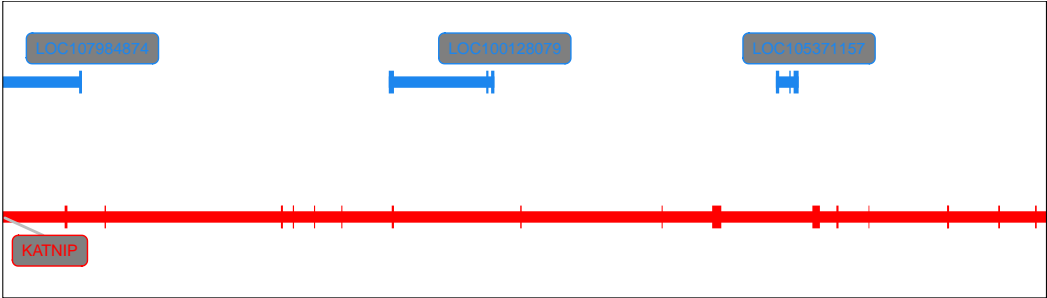
L5.6.NR4A2



Peaks

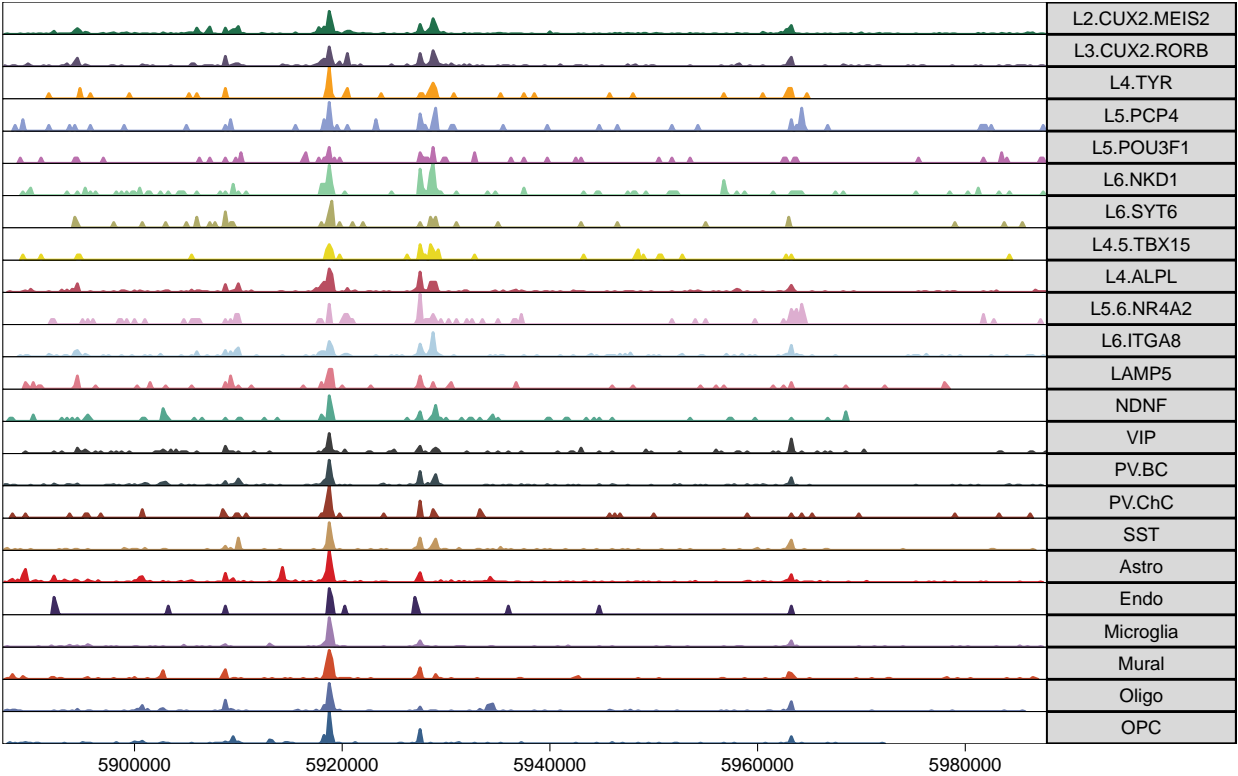


LoopTrack



Genes

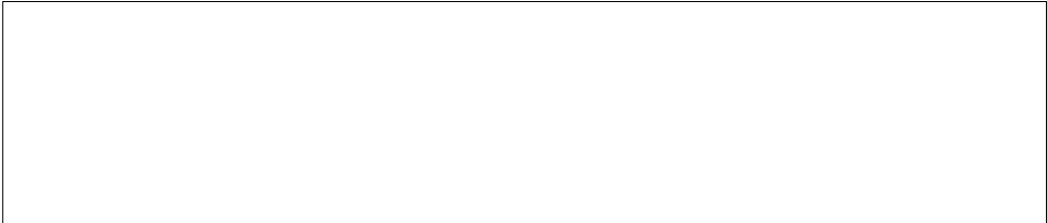
Coverage
(Norm. ATAC Signal Range 0–1.44) by ReadsInTSS)



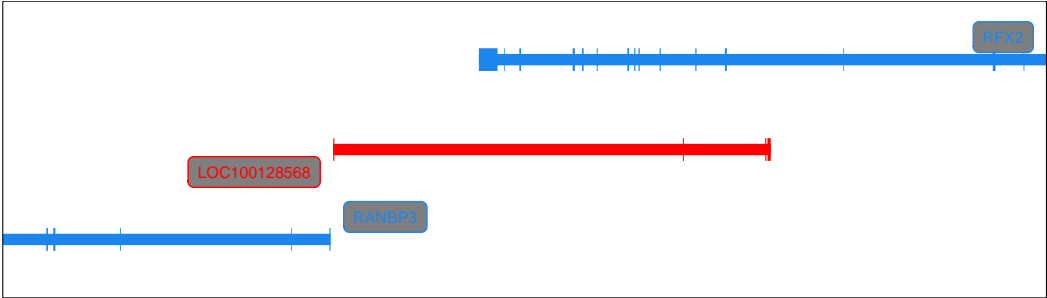
L5.6.NR4A2



Peaks



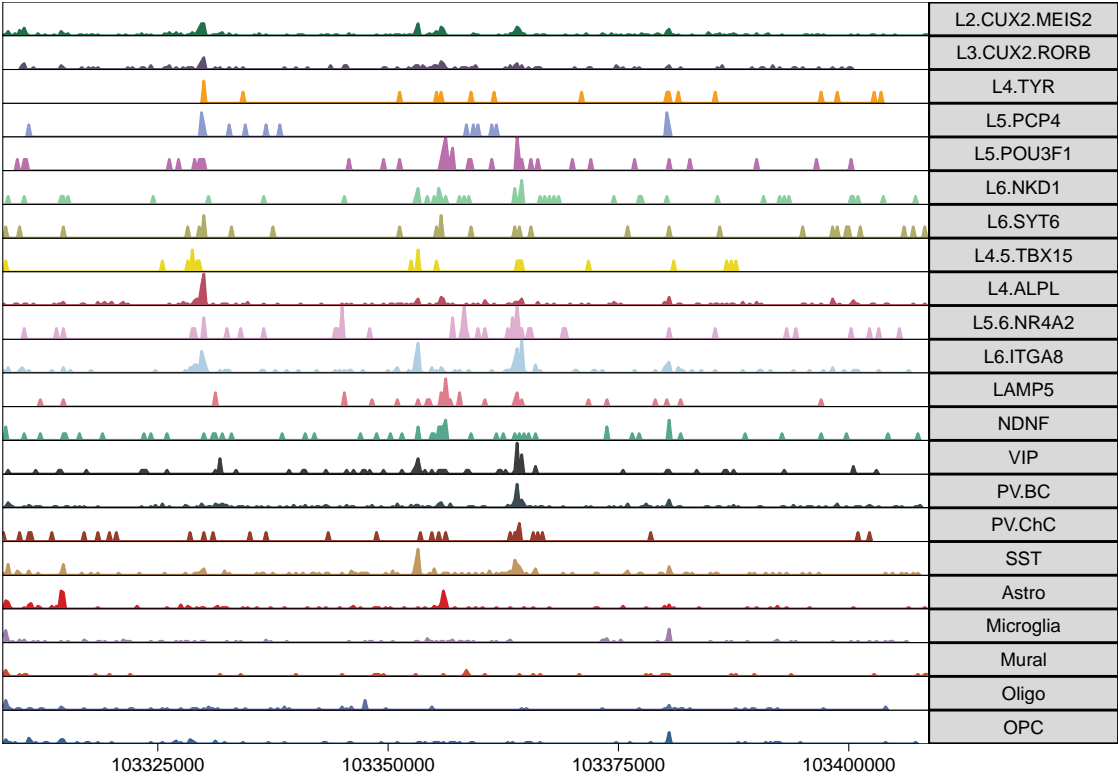
LoopTrack



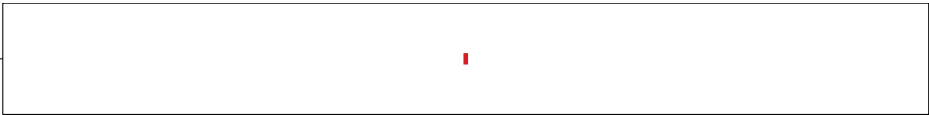
Genes

chr14:103308176–103408677 Marker Peak Tracks for: L5.6.NR4A2

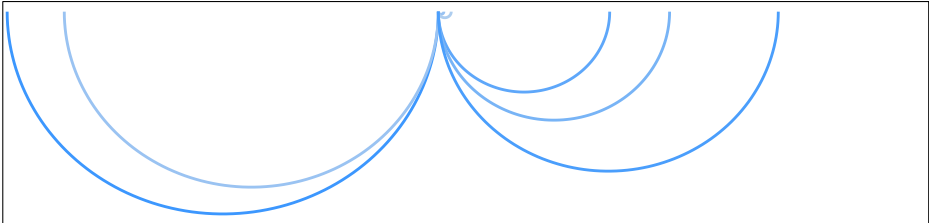
Coverage
(Norm. ATAC Signal Range (0–0.71) by ReadsInTSS)



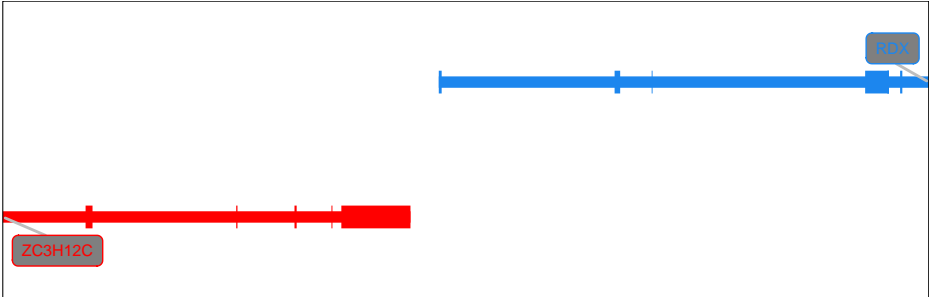
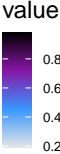
L5.6.NR4A2



Peaks



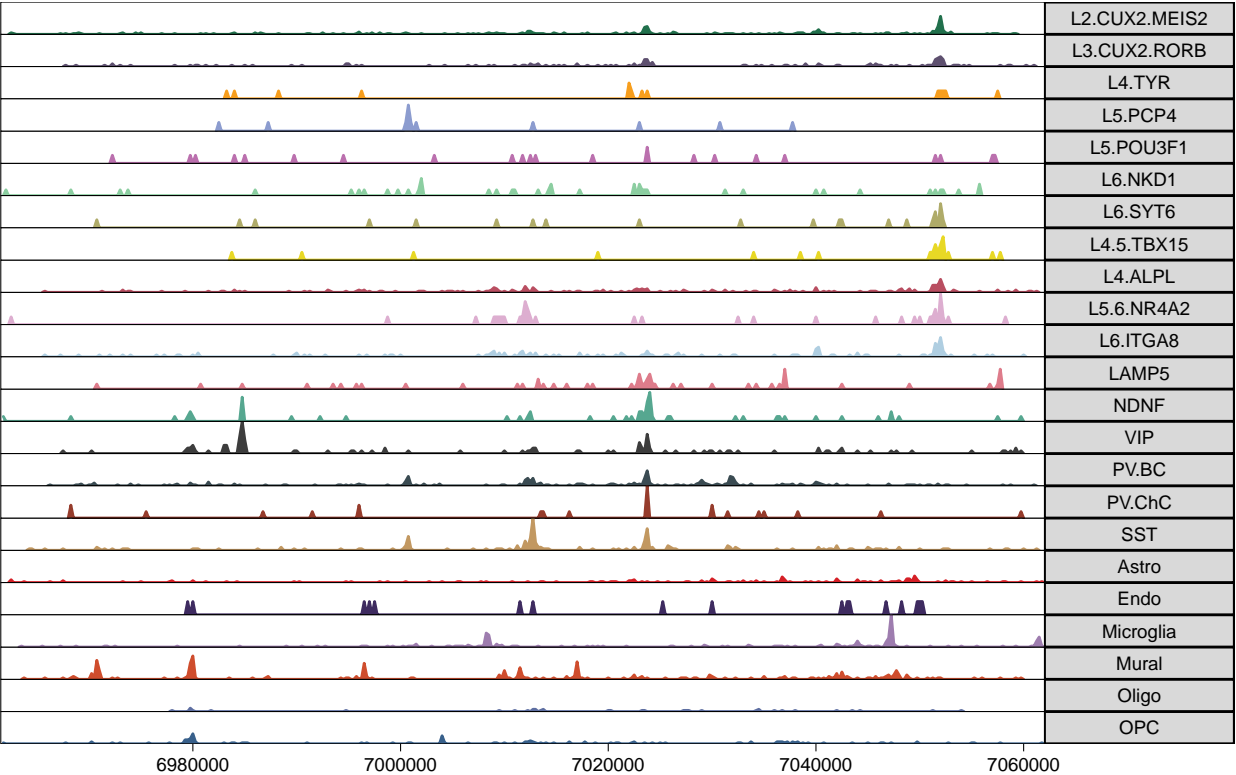
Peak2GeneLinks



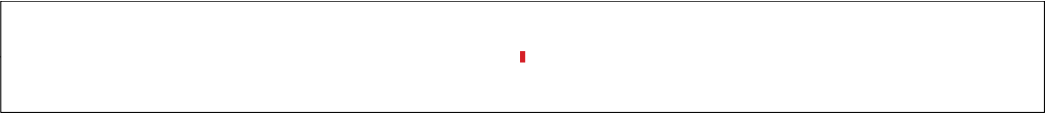
Genes

chr4:6961503–7062004 Marker Peak Tracks for: L5.6.NR4A2

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



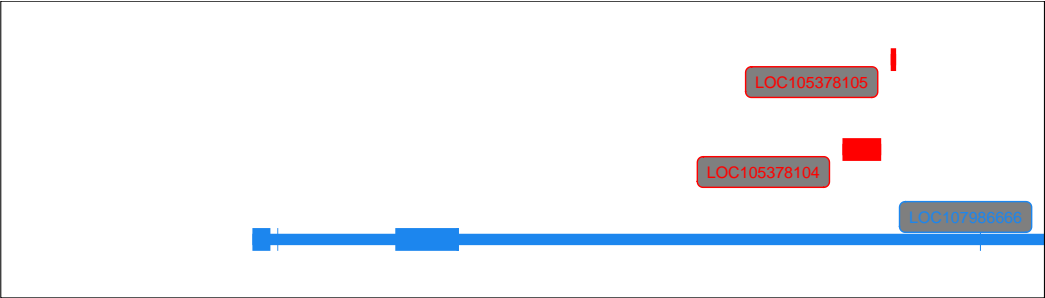
L5.6.NR4A2



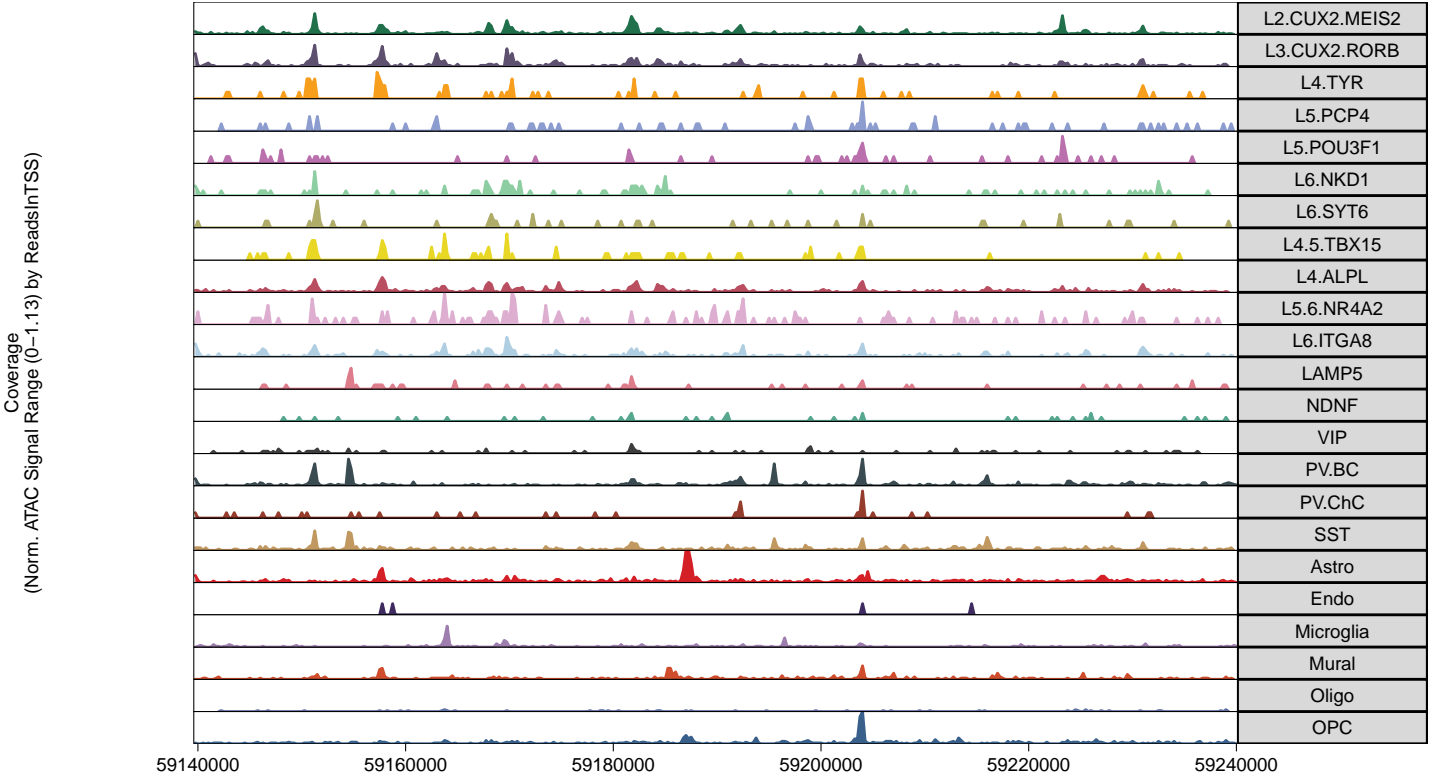
Peaks



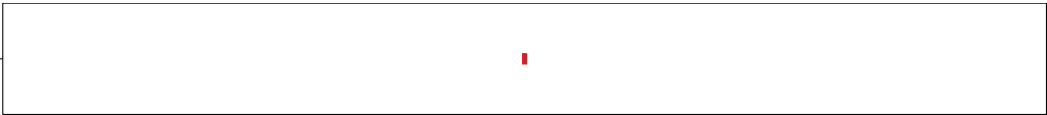
LoopTrack



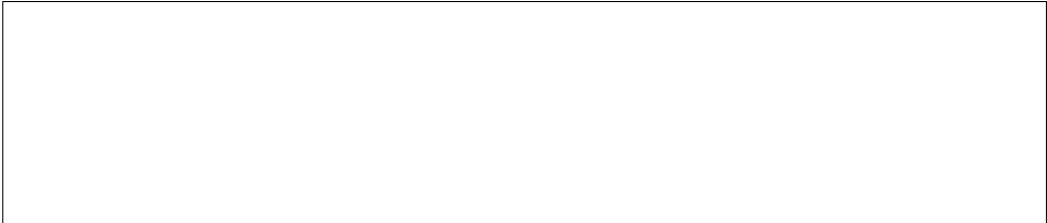
Genes



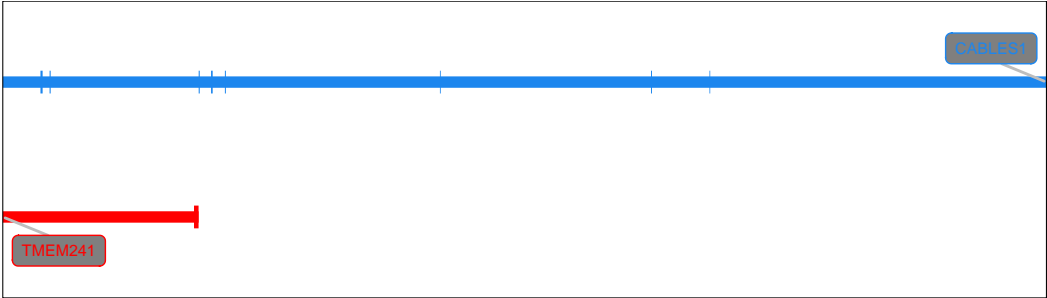
L5.6.NR4A2



Peaks

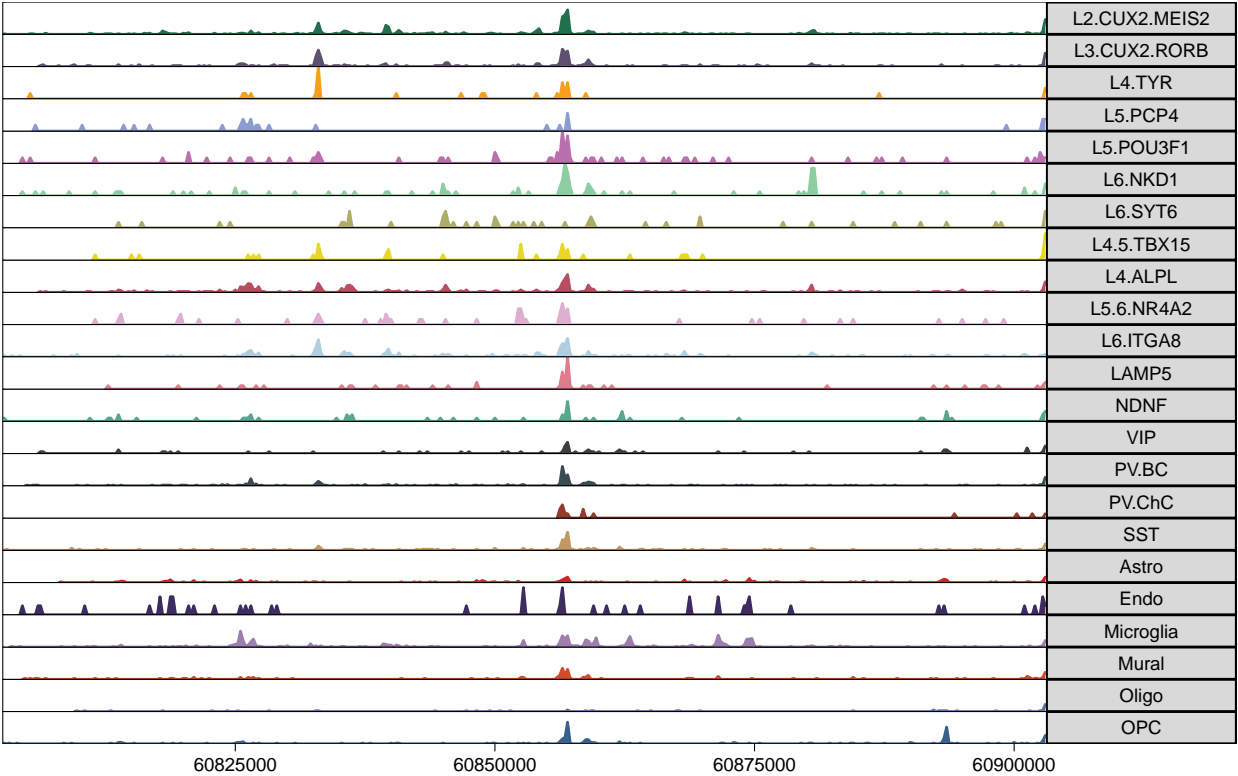


LoopTrack

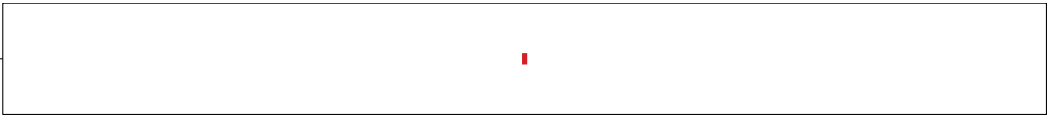


Genes

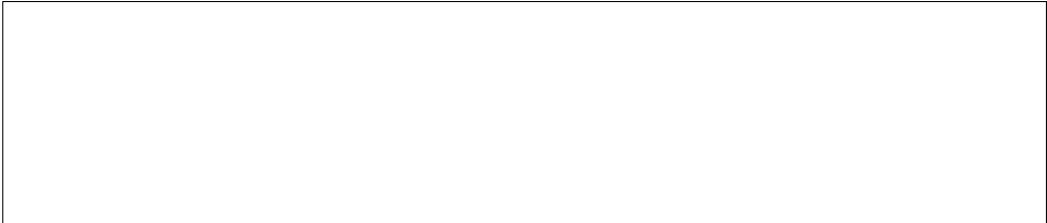
Coverage
(Norm. ATAC Signal Range (0-1.39) by ReadsInTSS)



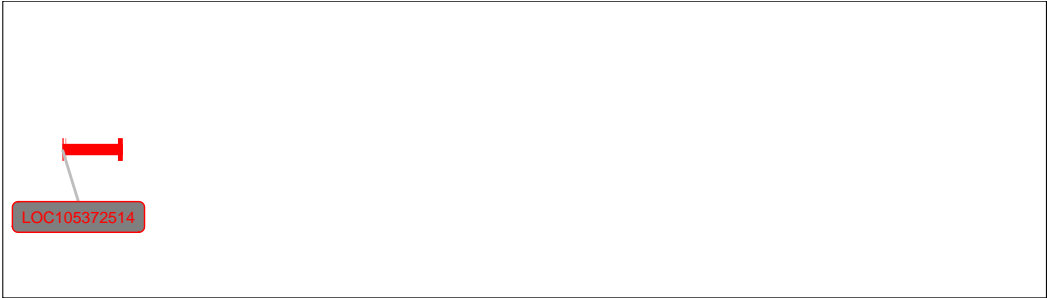
L5.6.NR4A2



Peaks

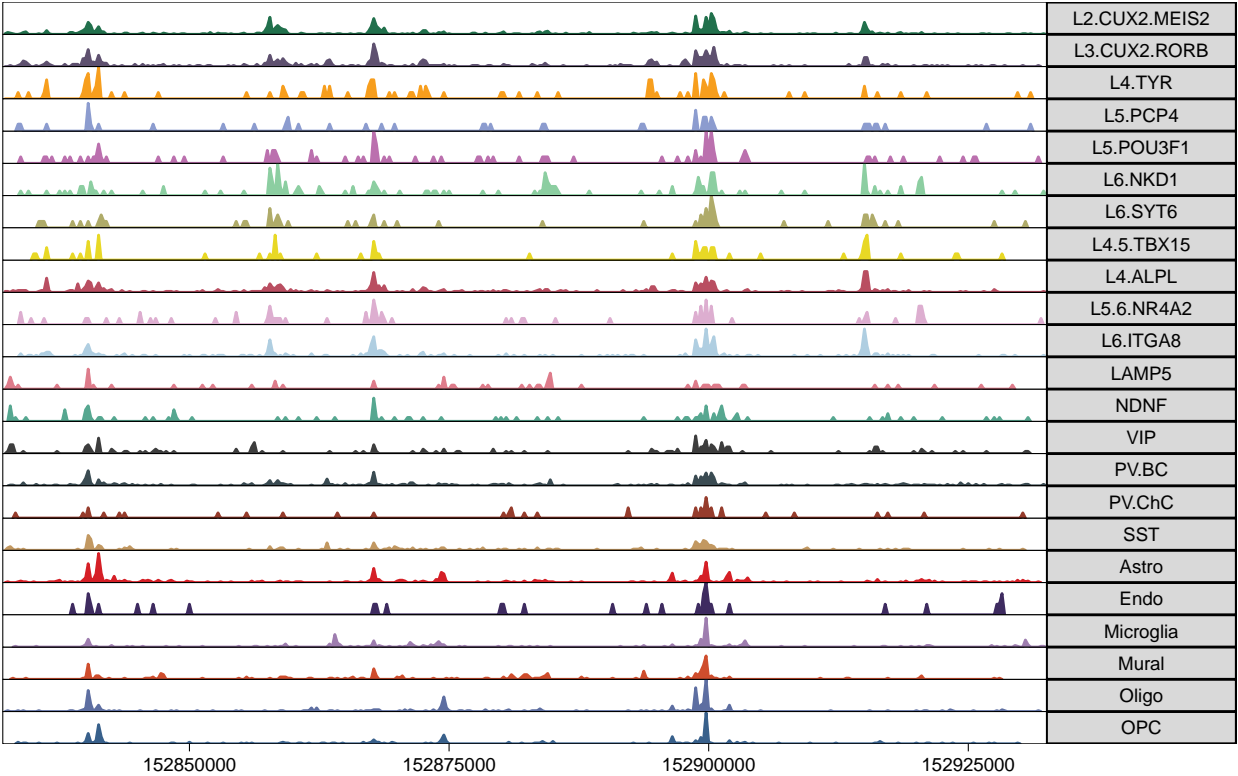


LoopTrack

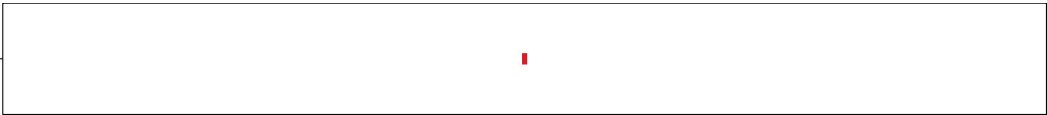


Genes

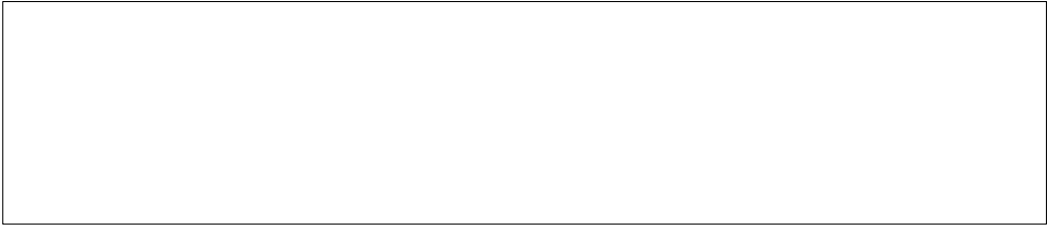
Coverage
(Norm. ATAC Signal Range (0-1.19) by ReadsInTSS)



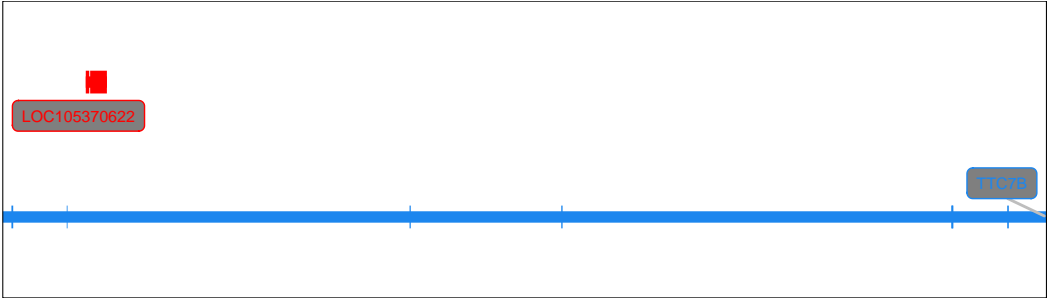
L5.6.NR4A2



Peaks

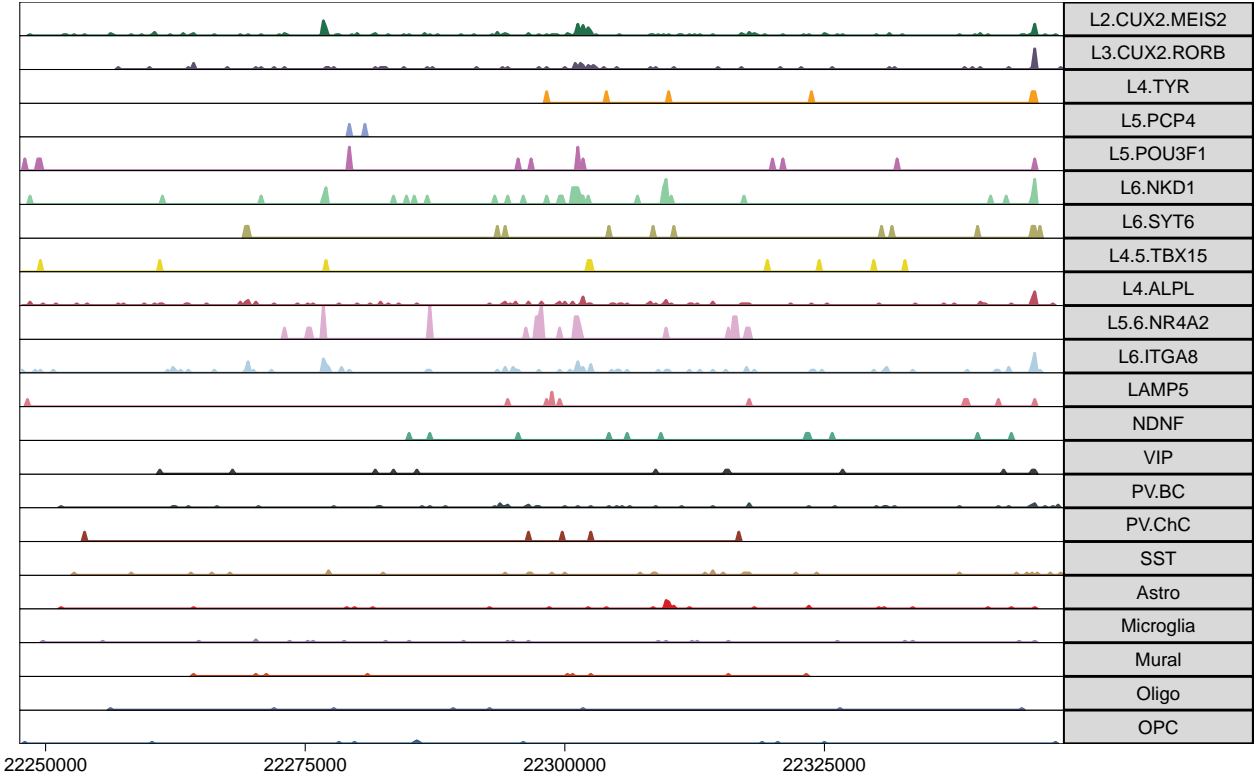


LoopTrack

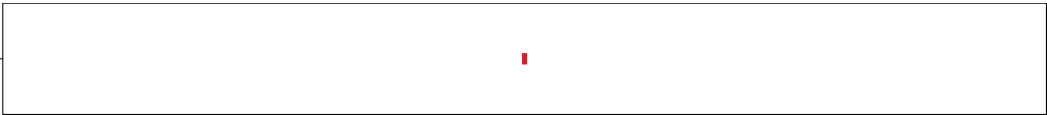


Genes

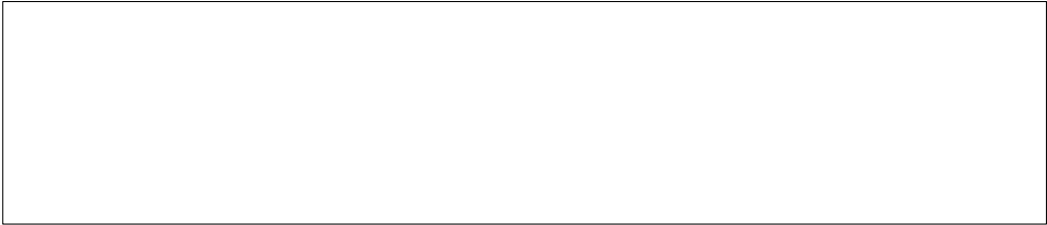
Coverage
(Norm. ATAC Signal Range 0–0.68) by ReadsInTSS)



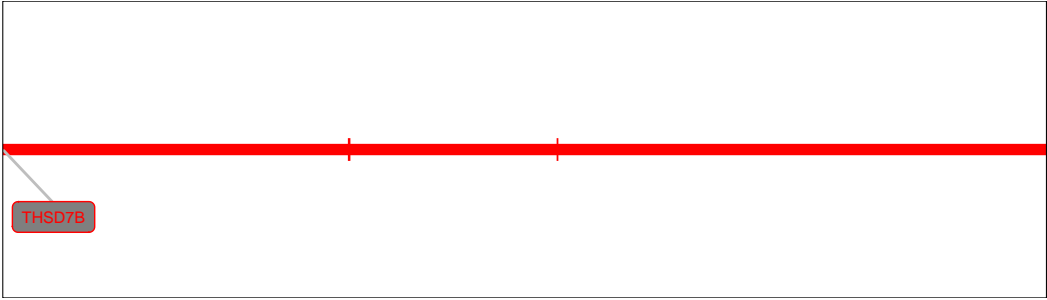
L5.6.NR4A2



Peaks

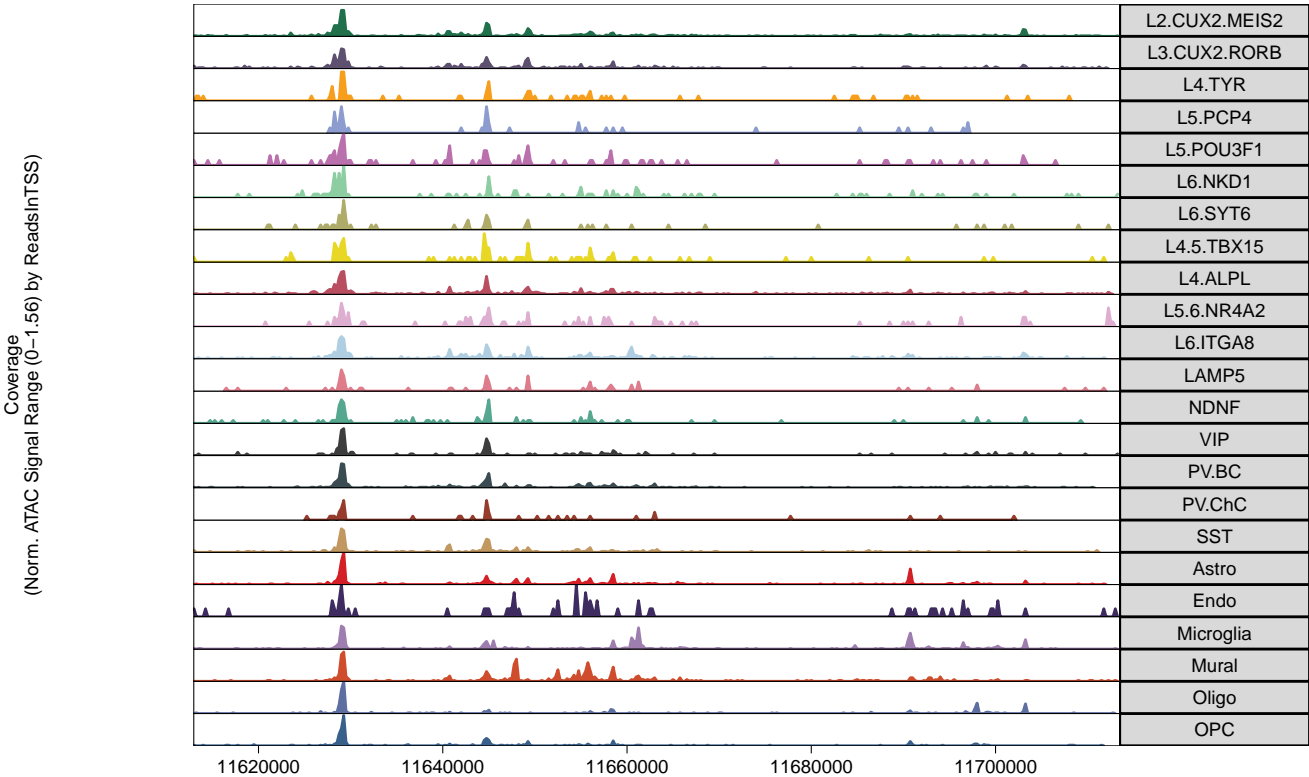


LoopTrack

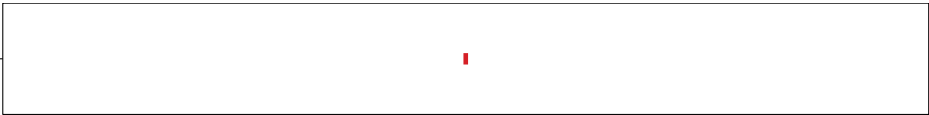


Genes

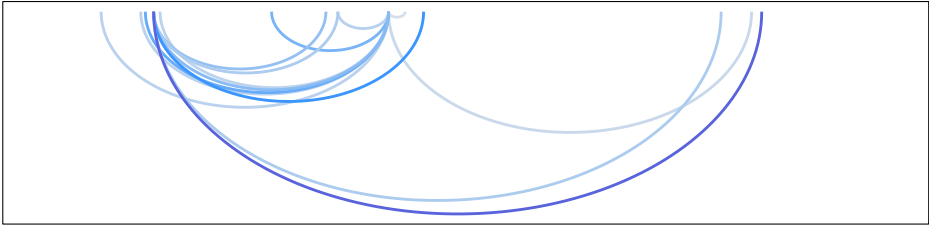
chr14:11612976–11713477 Marker Peak Tracks for: L5.6.NR4A2



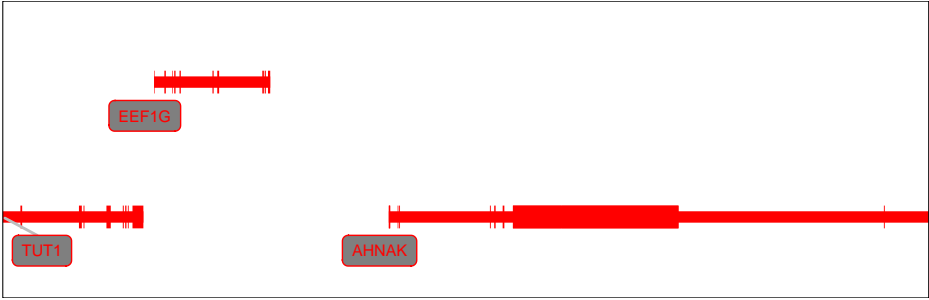
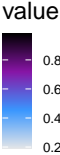
L5.6.NR4A2



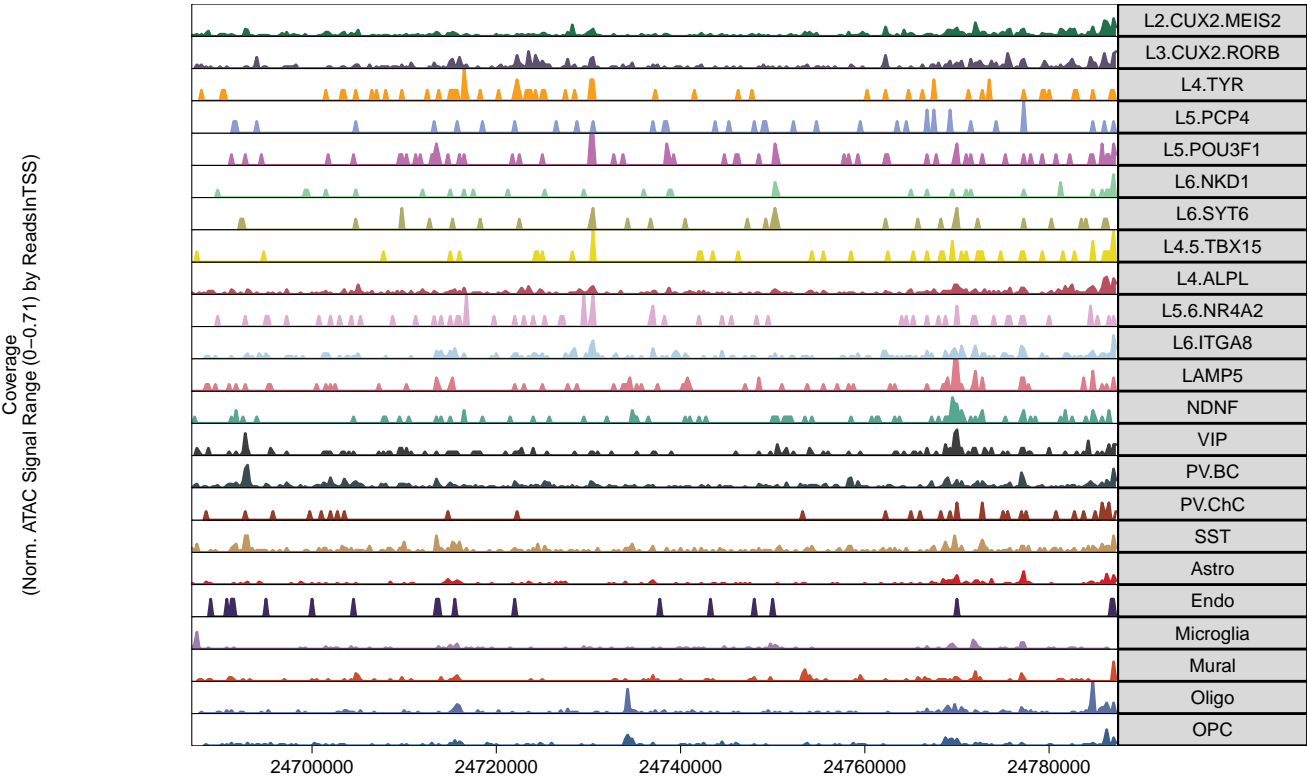
Peaks



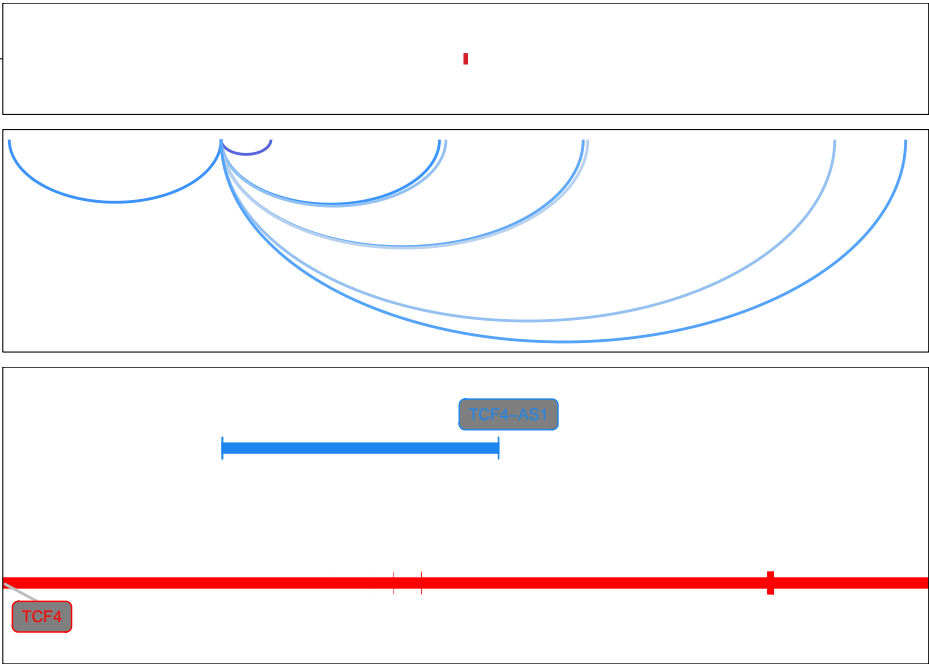
Peak2GeneLinks



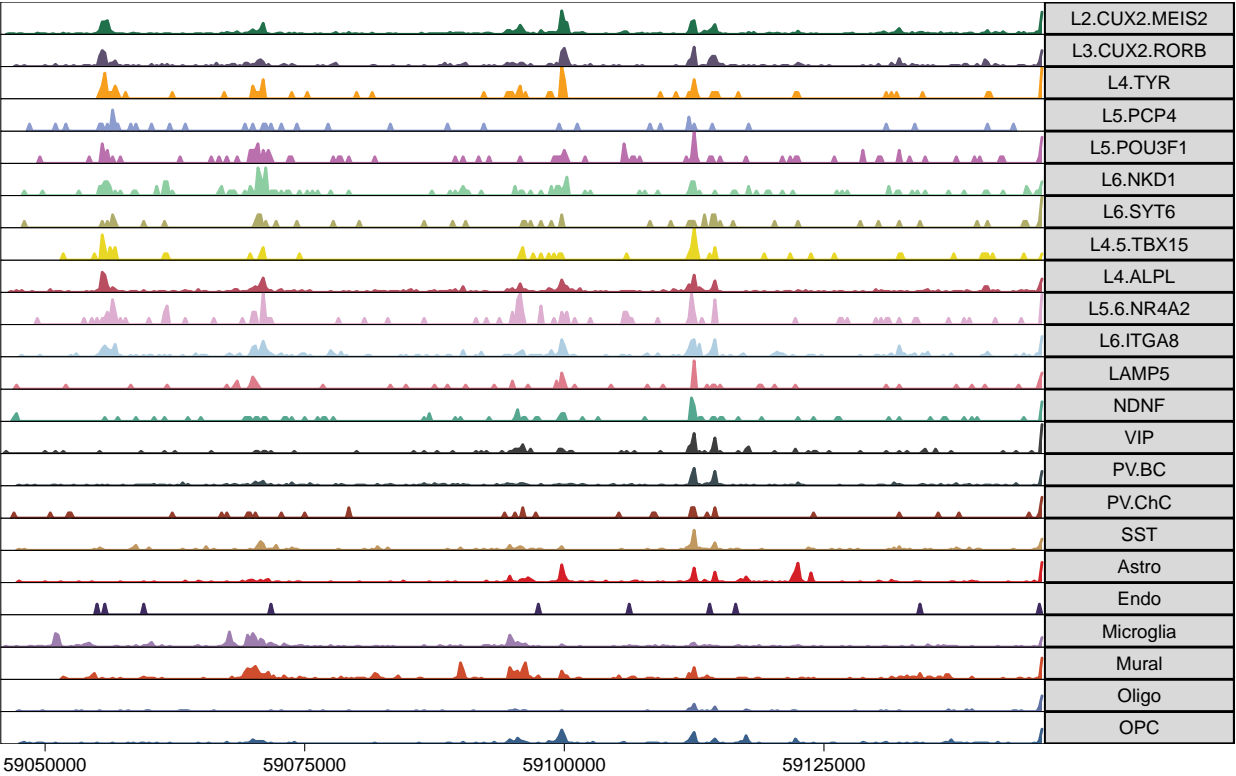
Genes



L5.6.NR4A2



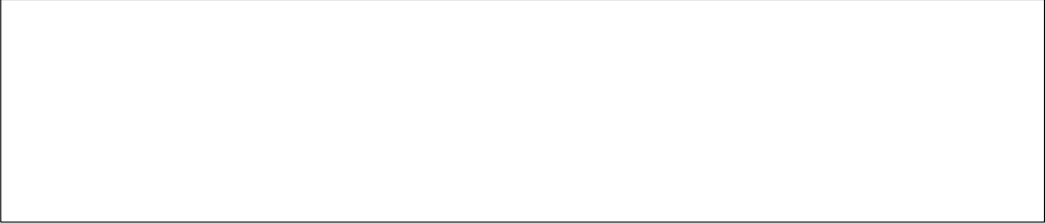
Coverage
(Norm. ATAC Signal Range 0–1.17) by ReadsInTSS)



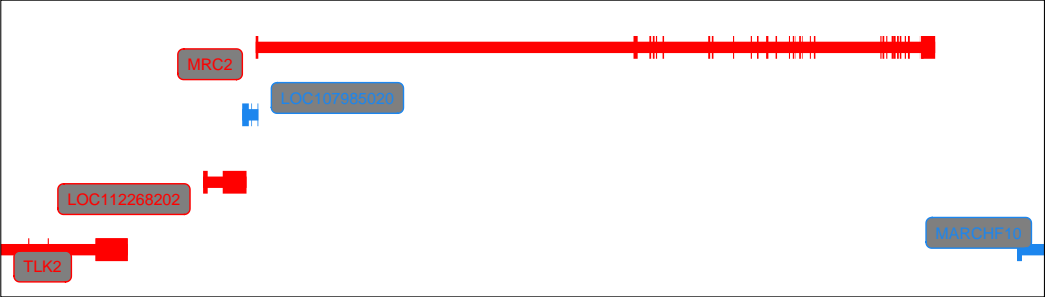
L5.6.NR4A2



Peaks



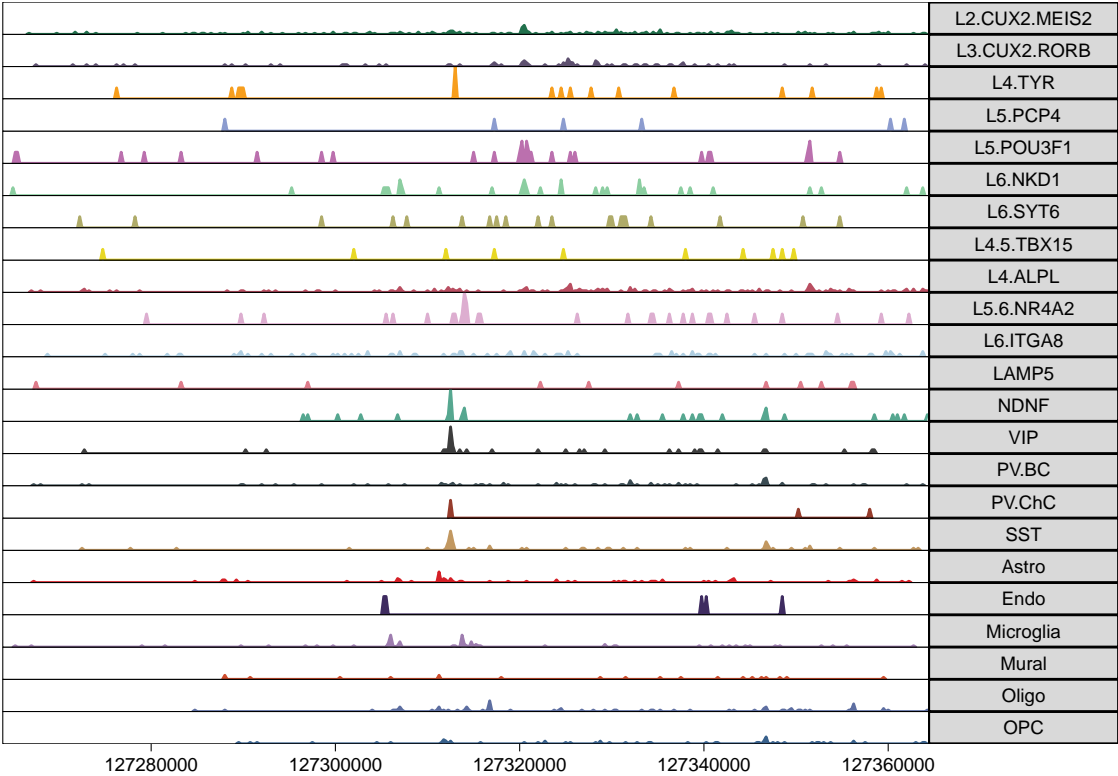
LoopTrack



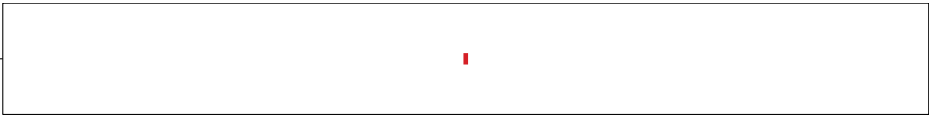
Genes

chr1:127263897-127364398 Marker Peak Tracks for: L5.6.NR4A2

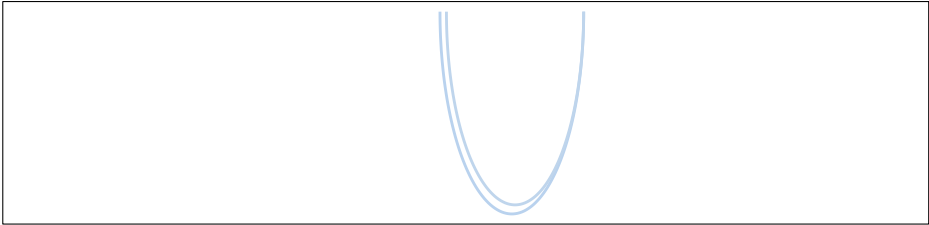
Coverage
(Norm. ATAC Signal Range (0-0.69) by ReadsInTSS)



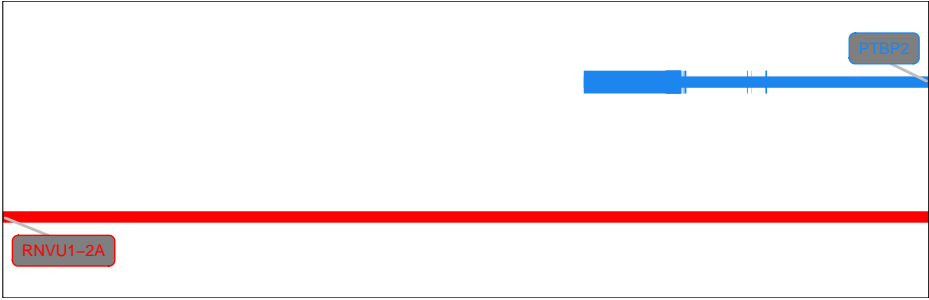
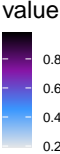
L5.6.NR4A2



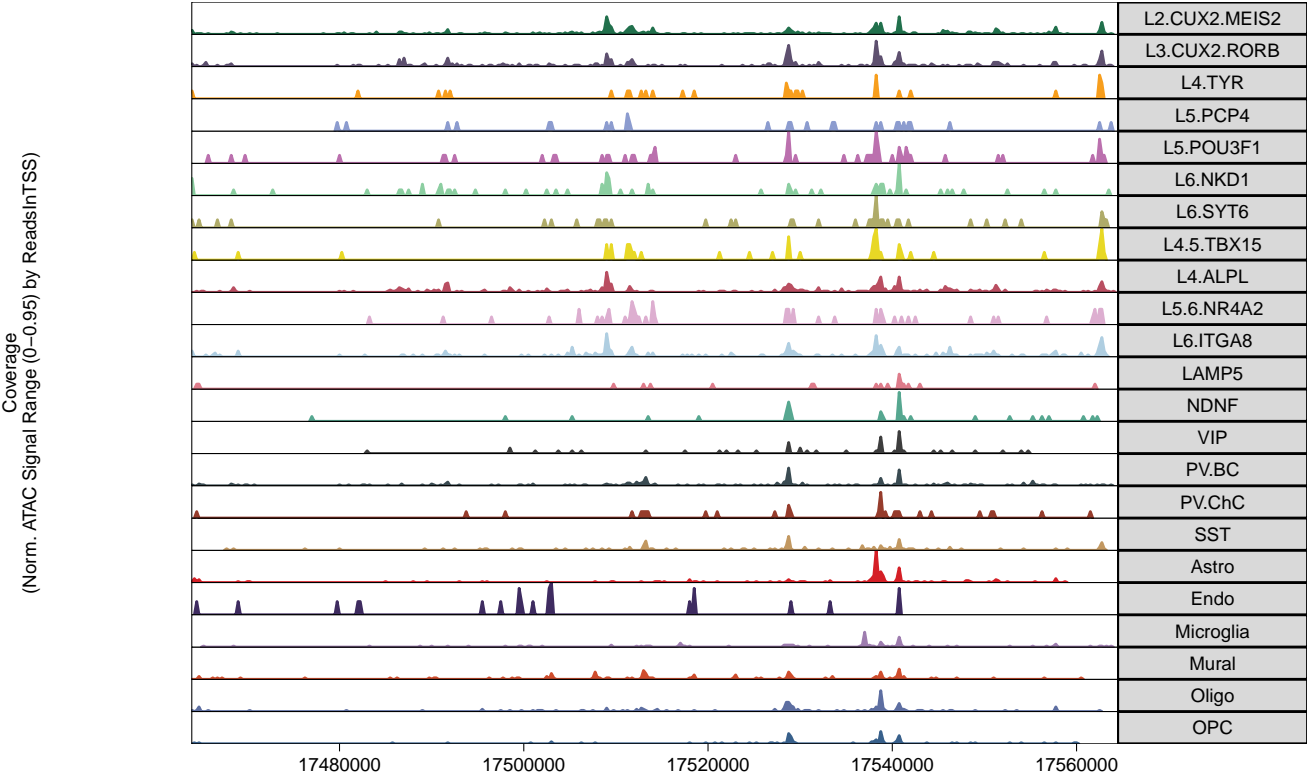
Peaks



Peak2GeneLinks



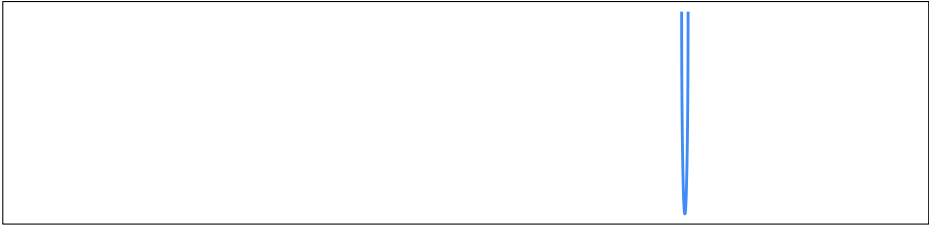
Genes



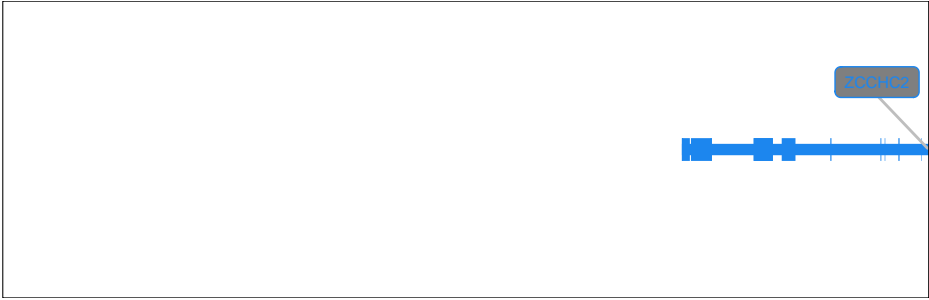
L5.6.NR4A2



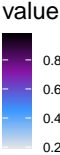
Peaks



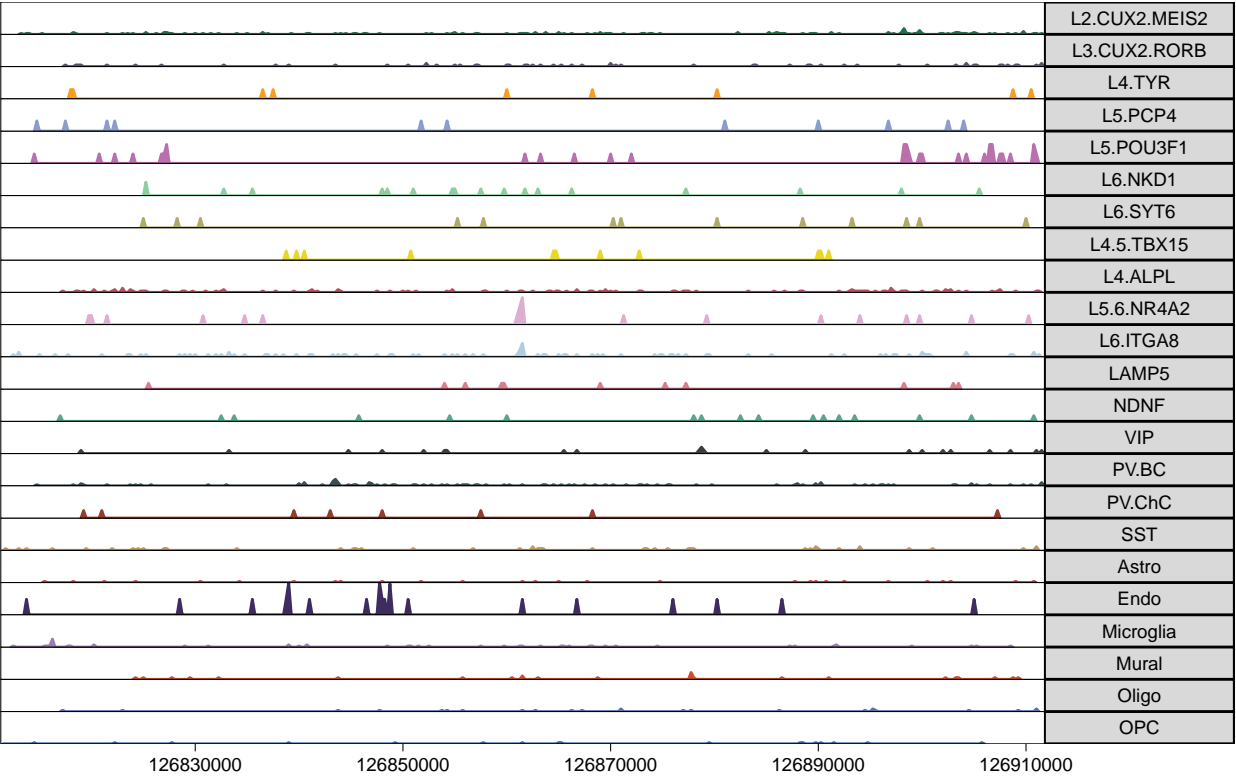
Peak2GeneLinks



Genes



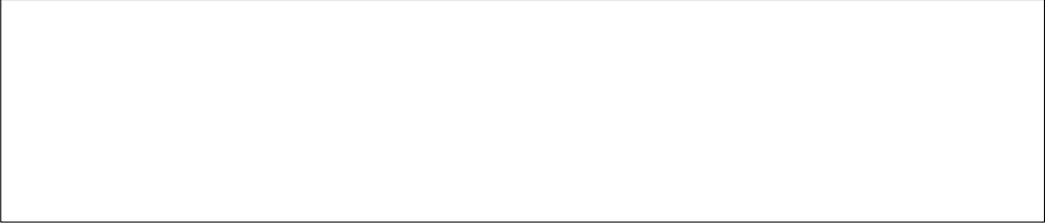
Coverage
(Norm. ATAC Signal Range (0–0.81) by ReadsInTSS)



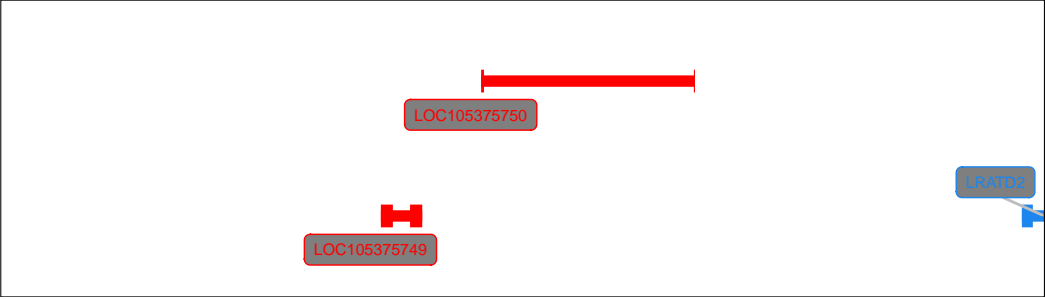
L5.6.NR4A2



Peaks

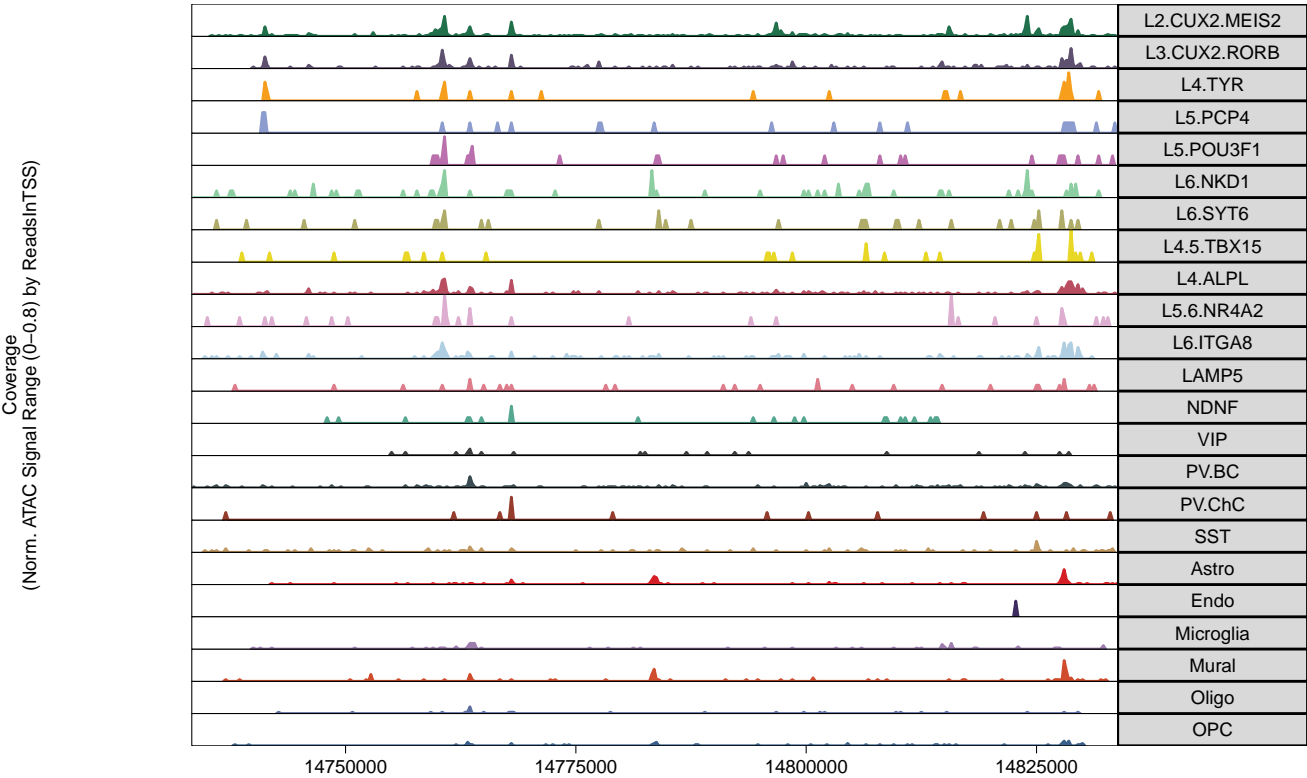


LoopTrack

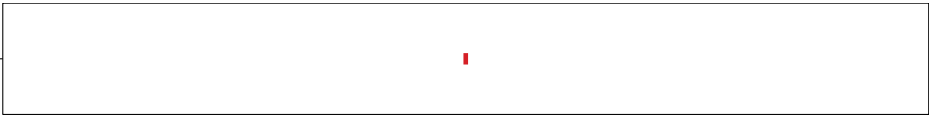


Genes

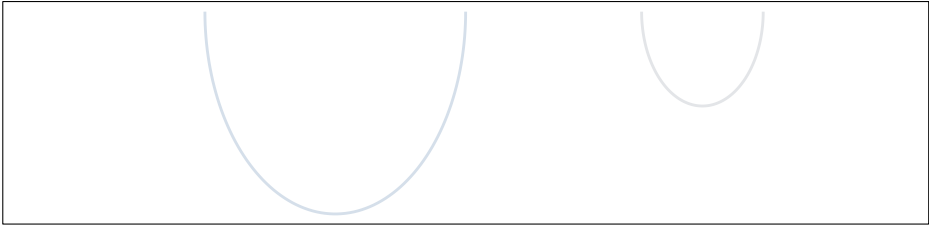
chr2:14733312–14833813 Marker Peak Tracks for: L5.6.NR4A2



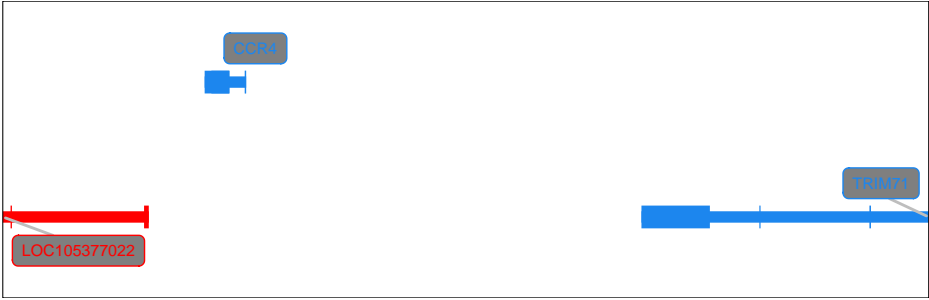
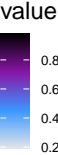
L5.6.NR4A2



Peaks

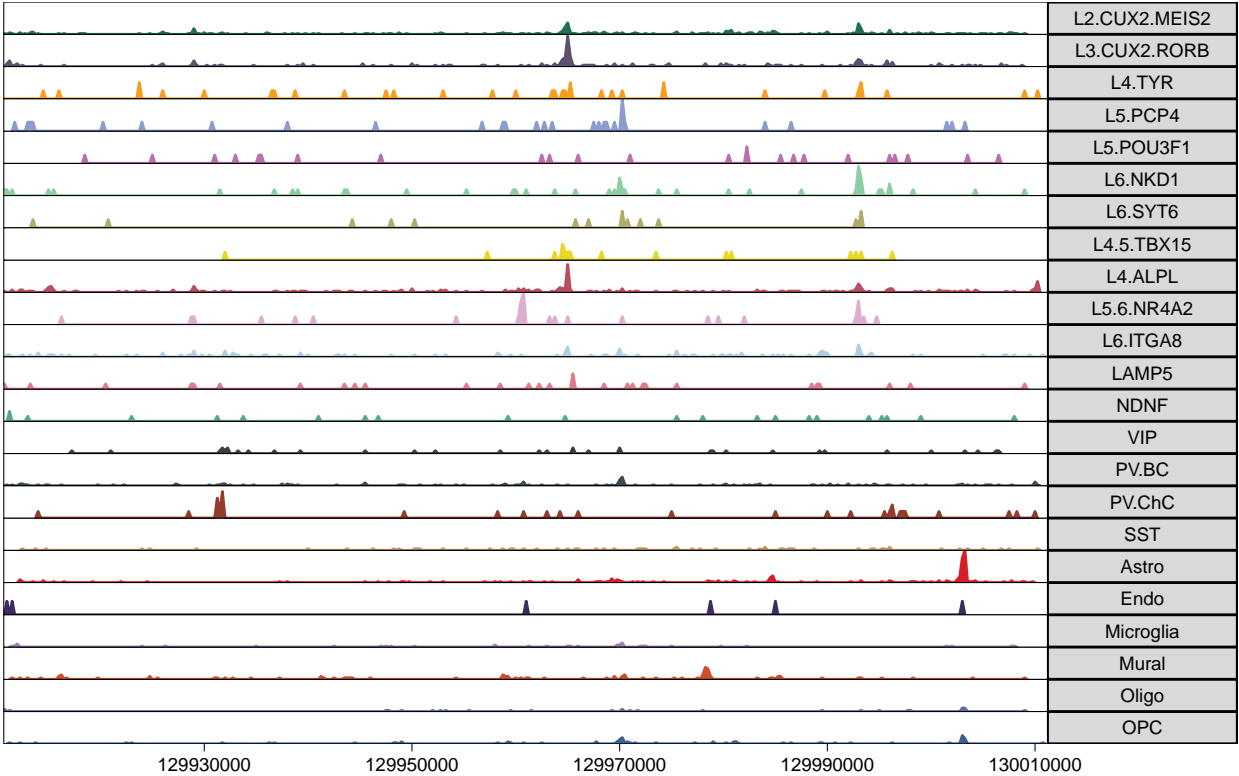


Peak2GeneLinks

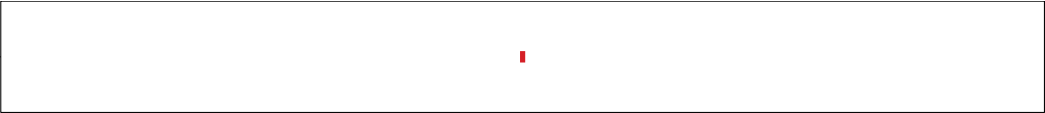


Genes

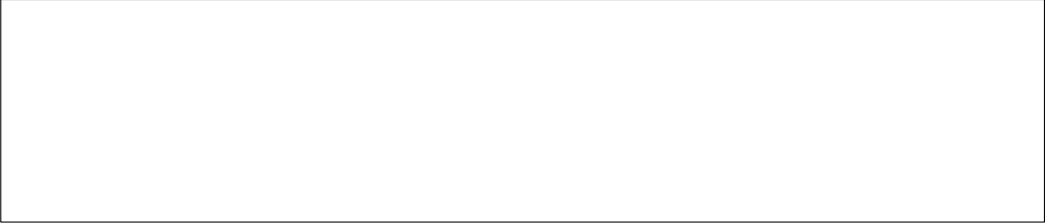
Coverage
(Norm. ATAC Signal Range 0–0.93) by ReadsInTSS)



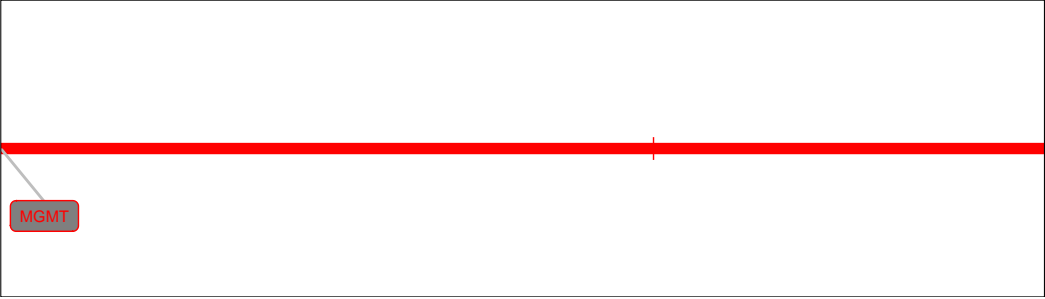
L5.6.NR4A2



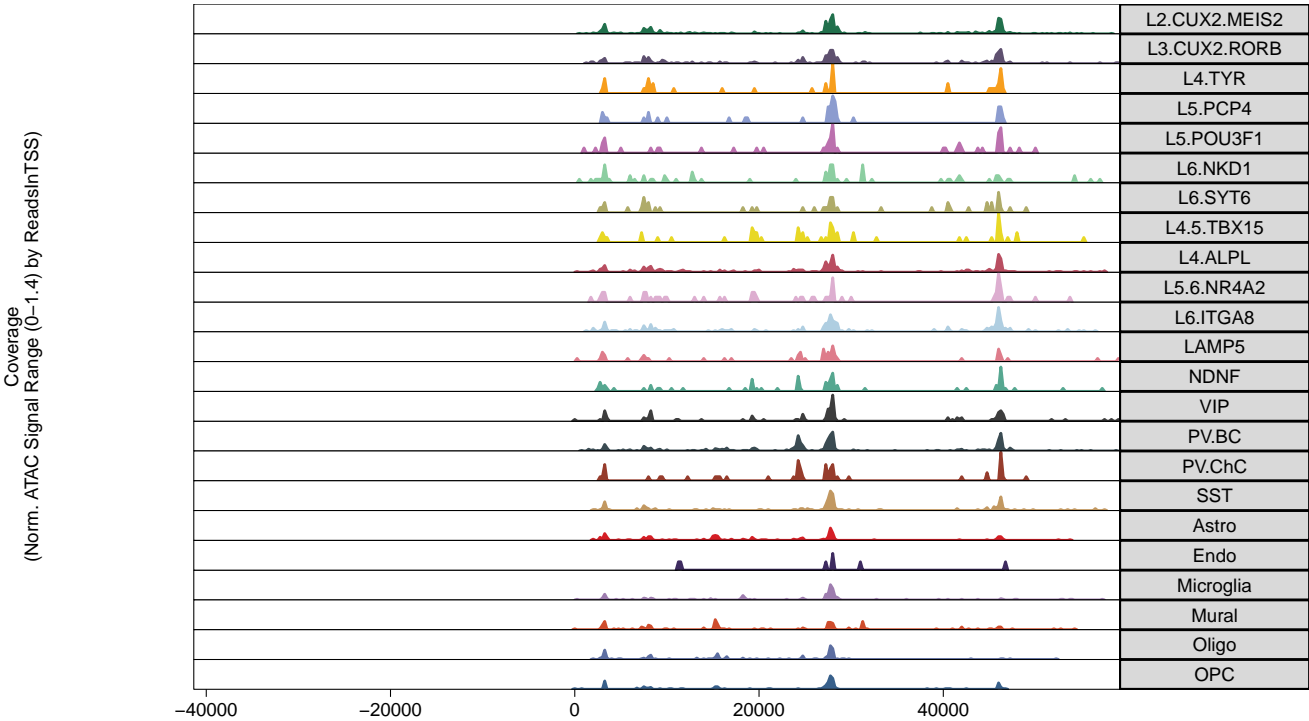
Peaks



LoopTrack



Genes



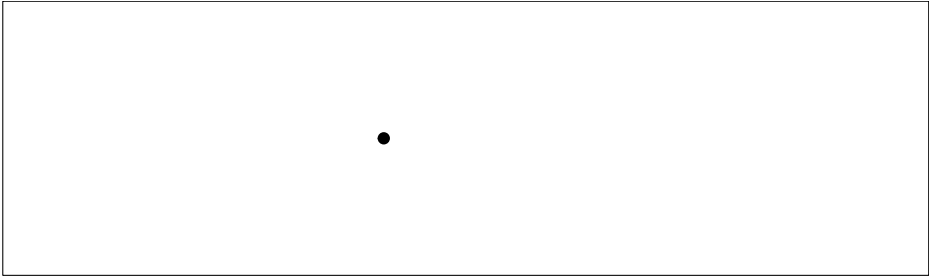
L5.6.NR4A2



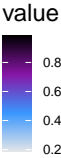
Peaks



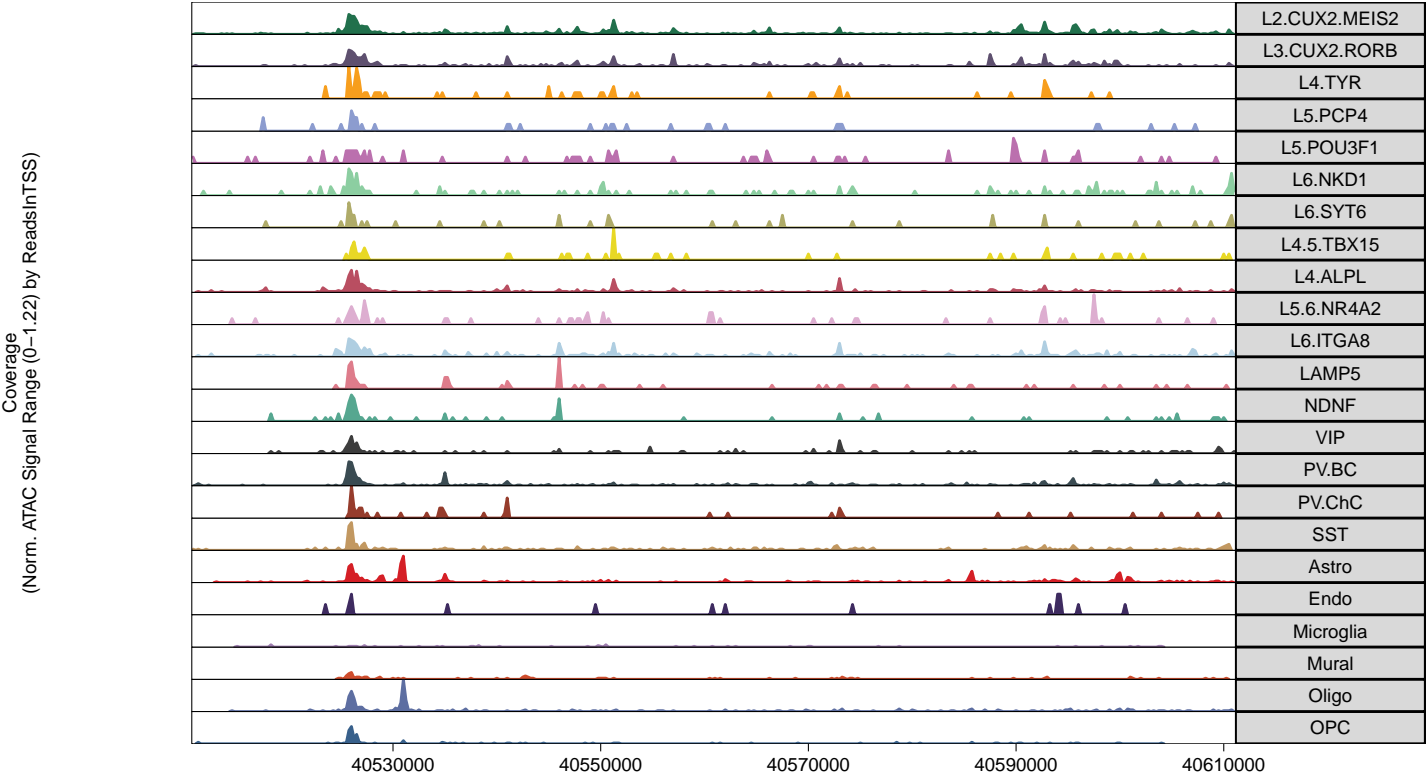
Peak2GeneLinks



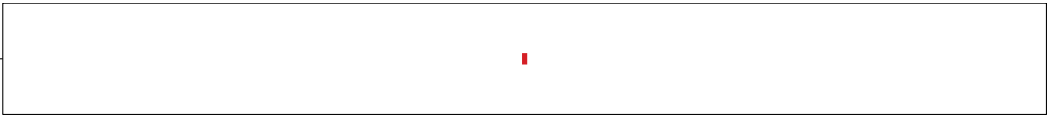
GeneTrack



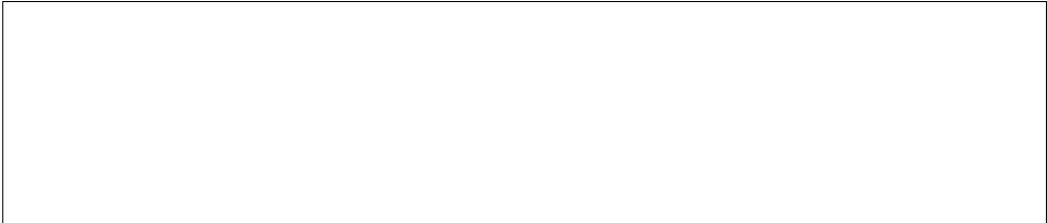
strand ● ·



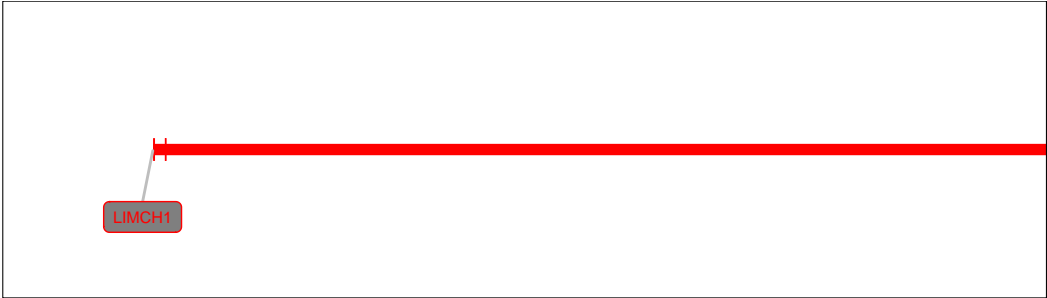
L5.6.NR4A2



Peaks



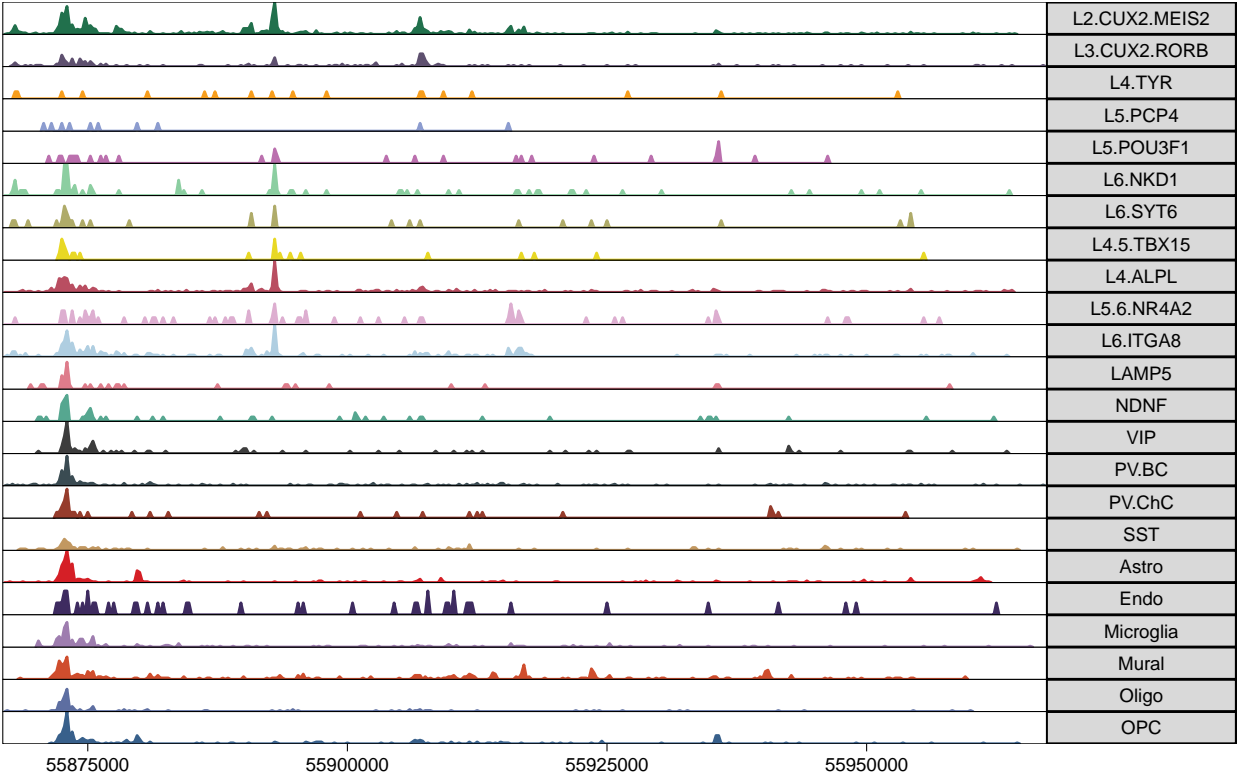
LoopTrack



Genes

chr10:55866816-55967317 Marker Peak Tracks for: L5.6.NR4A2

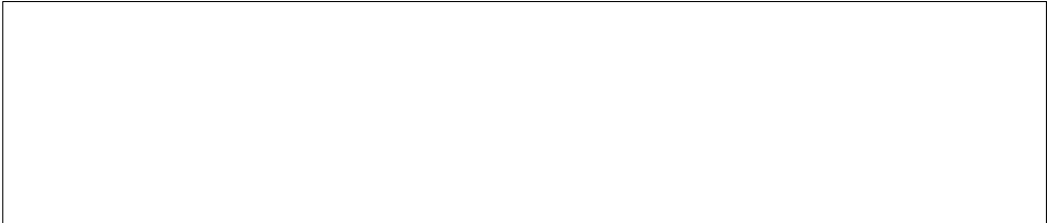
Coverage
(Norm. ATAC Signal Range 0-1.06) by ReadsInTSS)



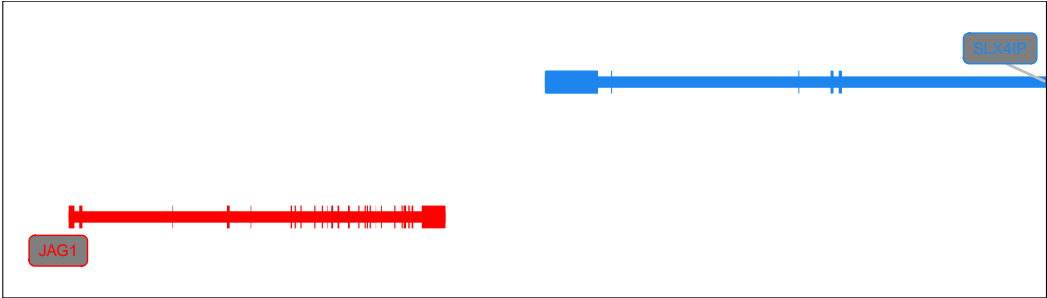
L5.6.NR4A2



Peaks



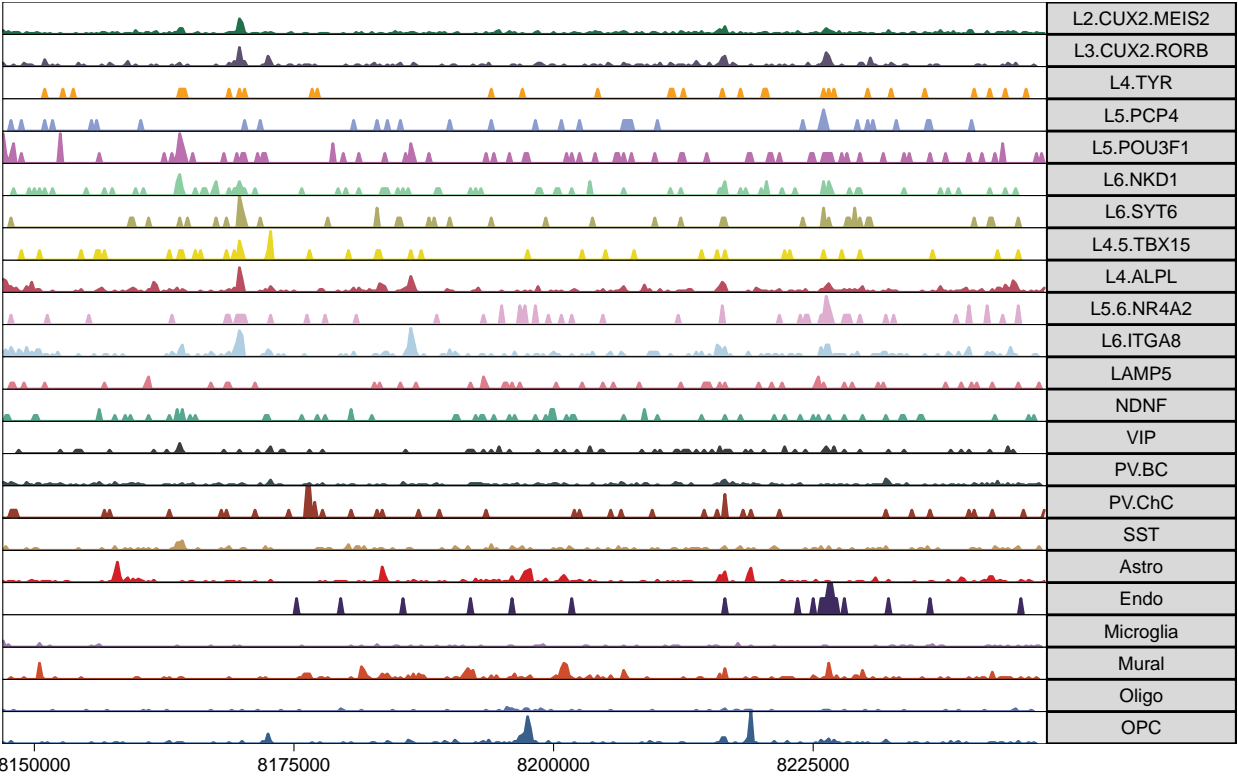
LoopTrack



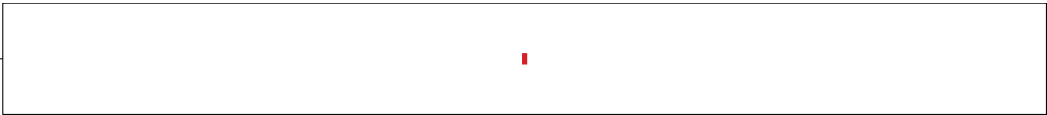
Genes

chr10:8146961–8247462 Marker Peak Tracks for: L5.6.NR4A2

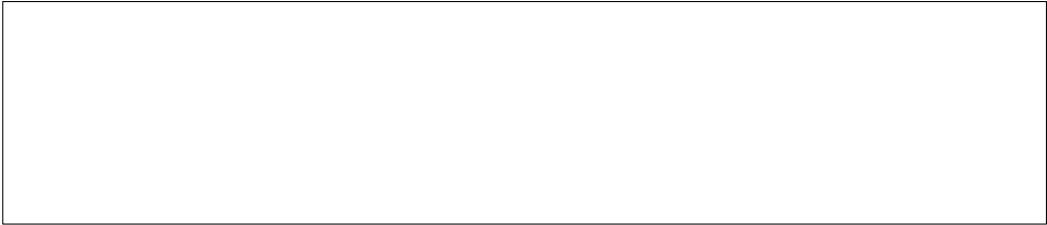
Coverage
(Norm. ATAC Signal Range 0–0.78) by ReadsInTSS)



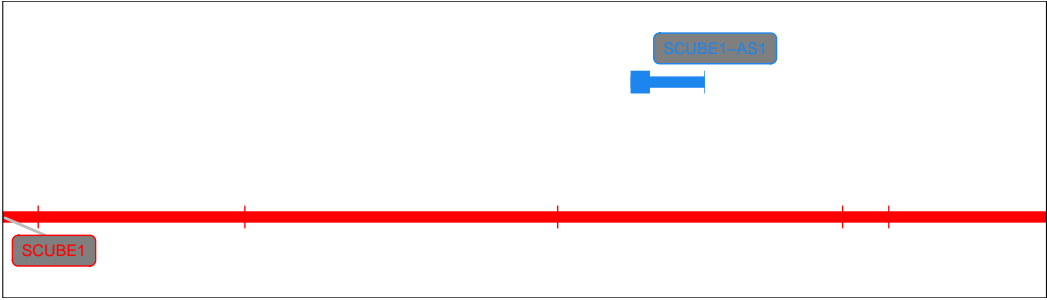
L5.6.NR4A2



Peaks

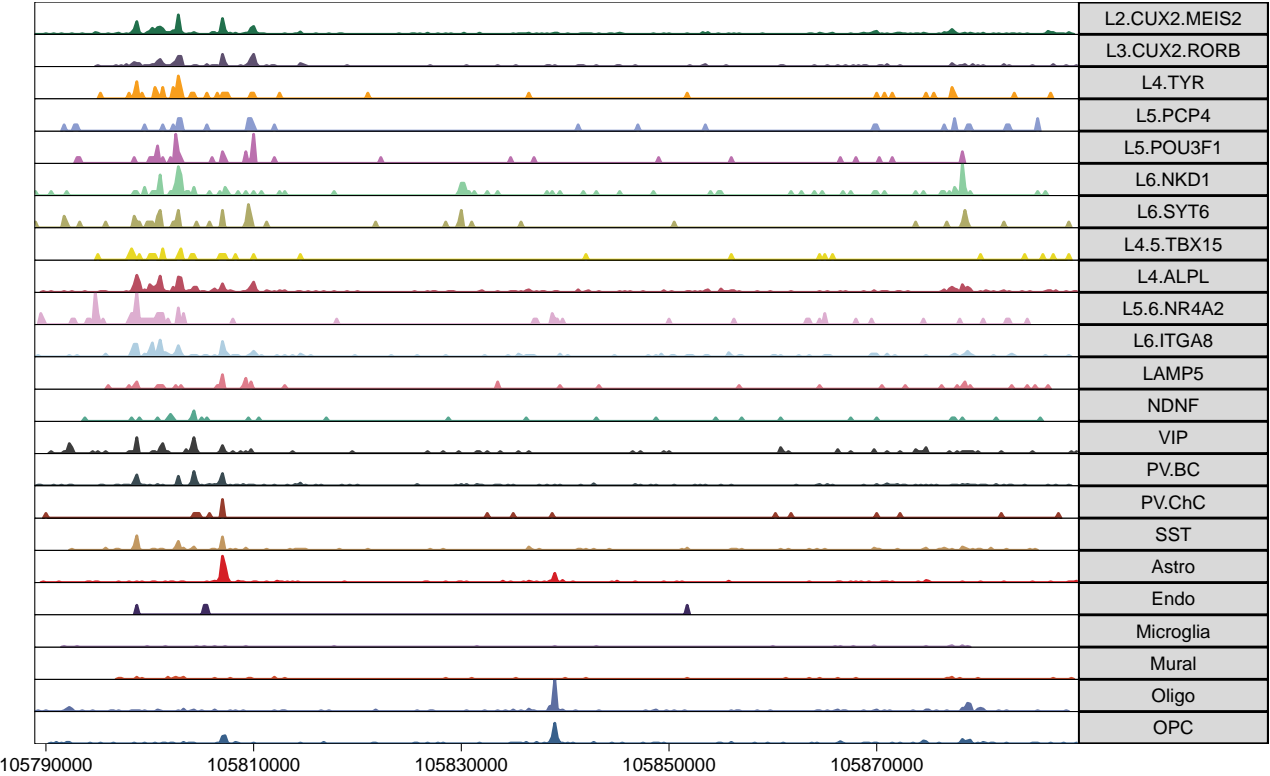


LoopTrack

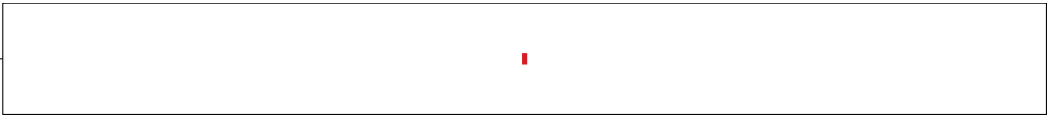


Genes

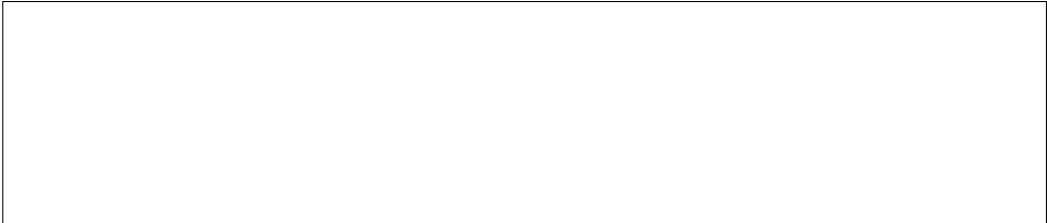
Coverage
(Norm. ATAC Signal Range 0–1.32) by ReadsInTSS)



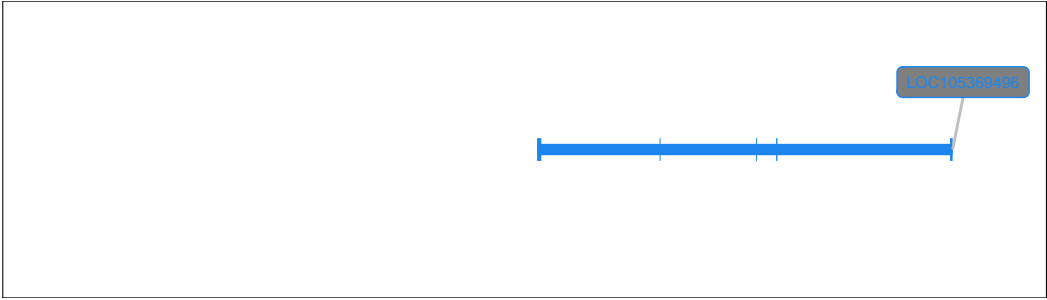
L5.6.NR4A2



Peaks

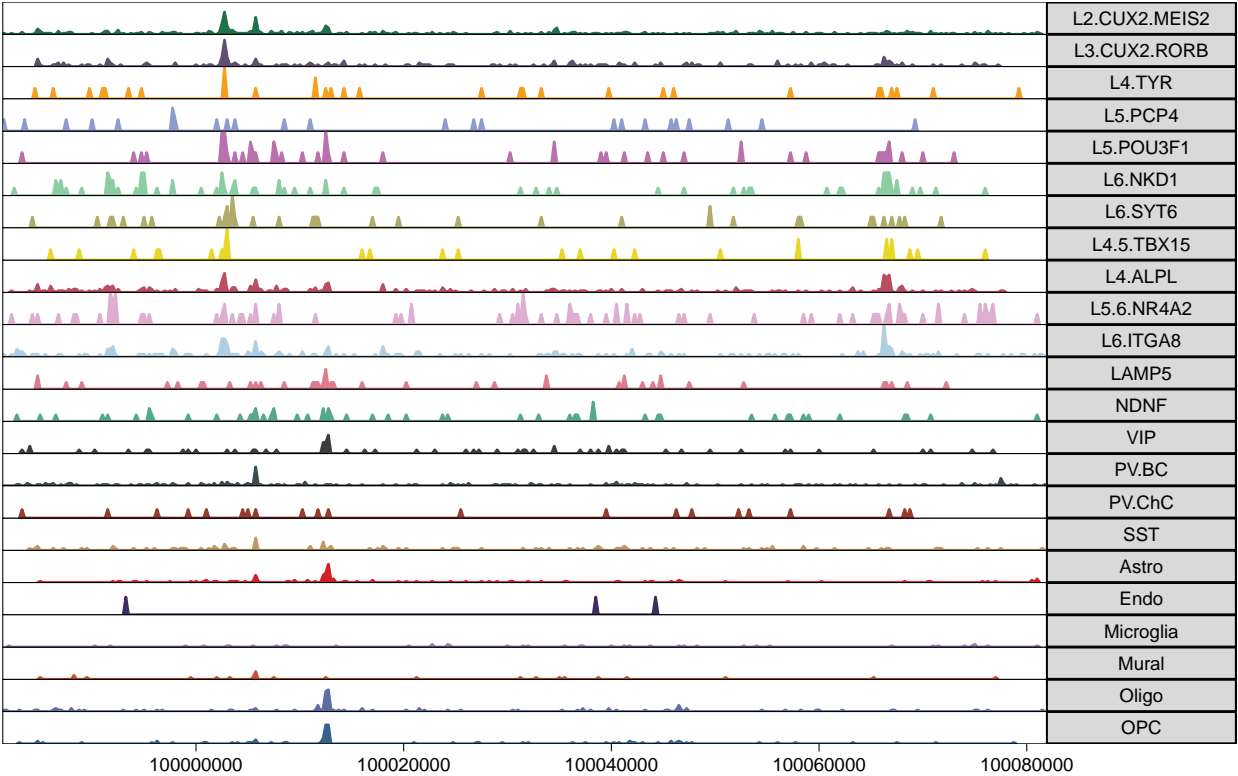


LoopTrack

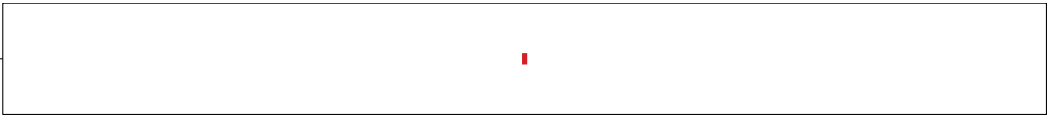


Genes

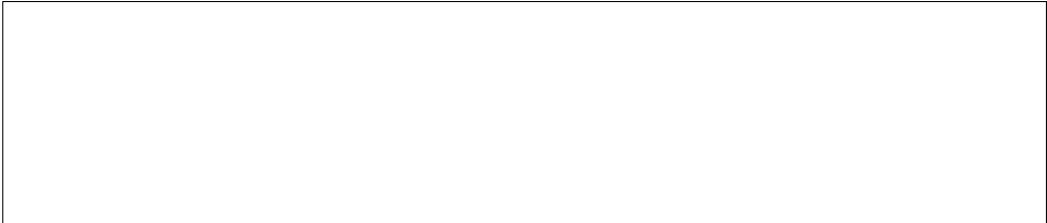
Coverage
(Norm. ATAC Signal Range (0-0.71) by ReadsInTSS)



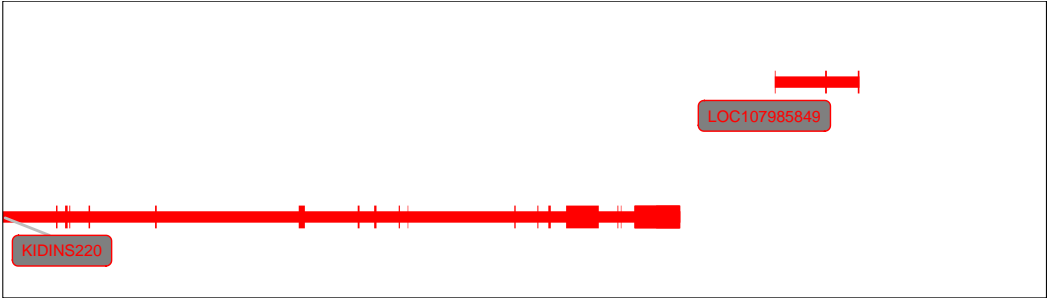
L5.6.NR4A2



Peaks

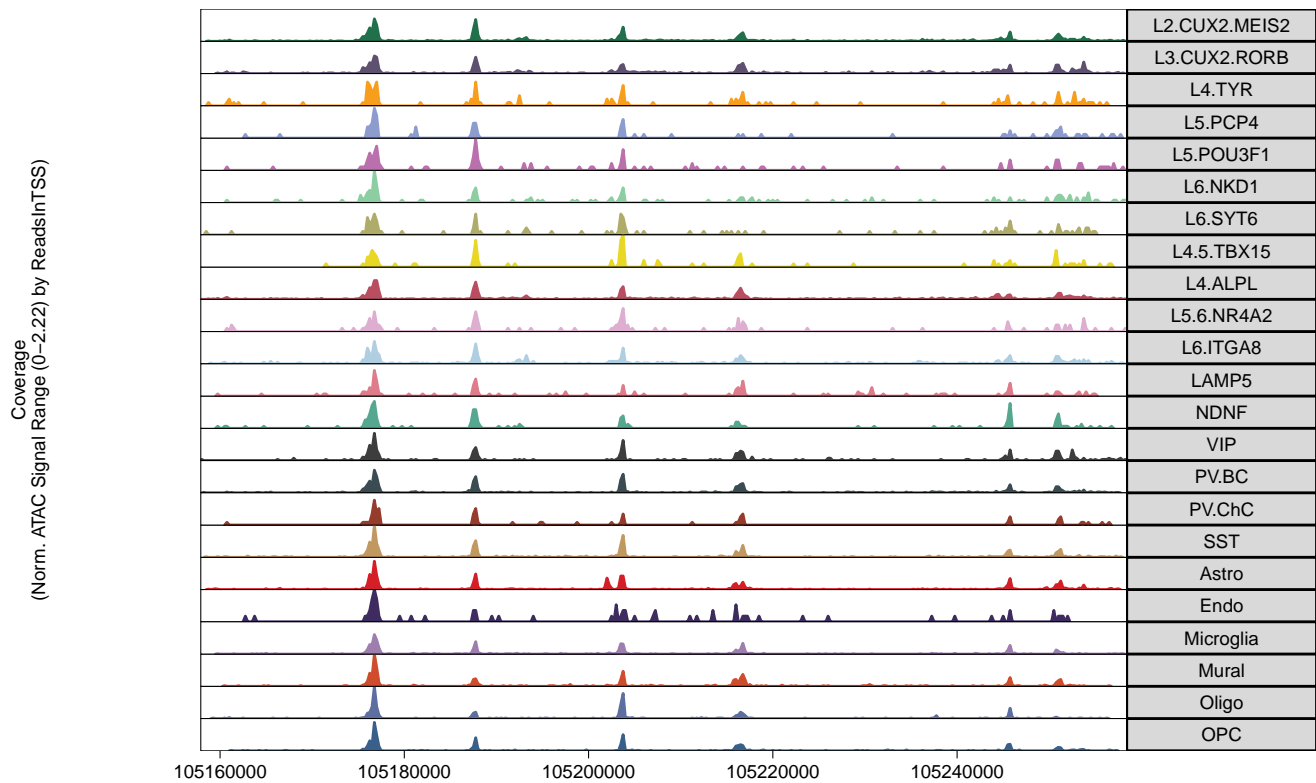


LoopTrack



Genes

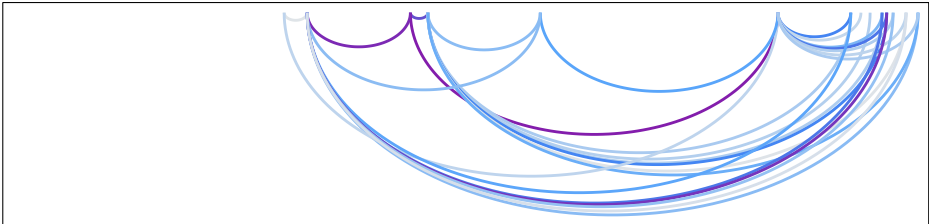
chr2:105157909–105258410 Marker Peak Tracks for: L5.6.NR4A2



L5.6.NR4A2

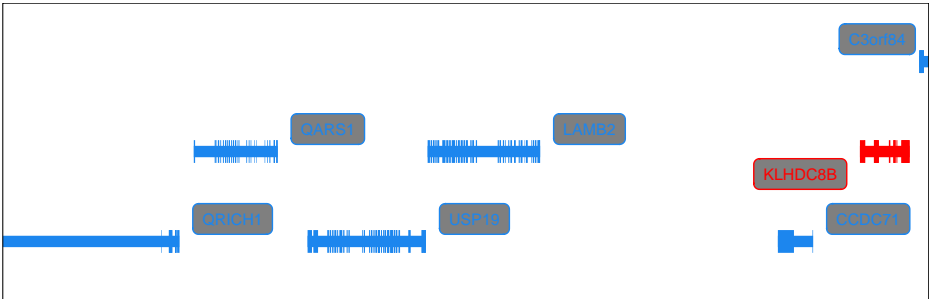
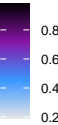


Peaks



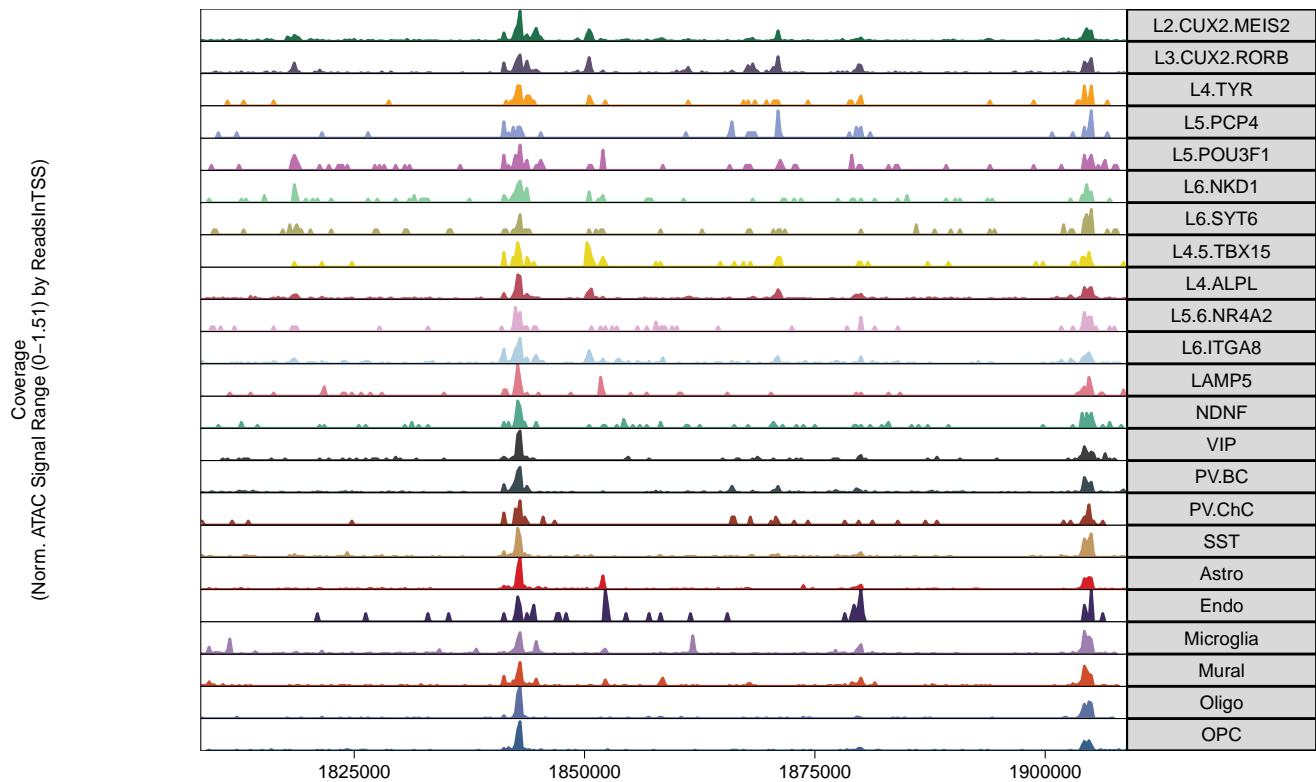
Peak2GeneLinks

value



Genes

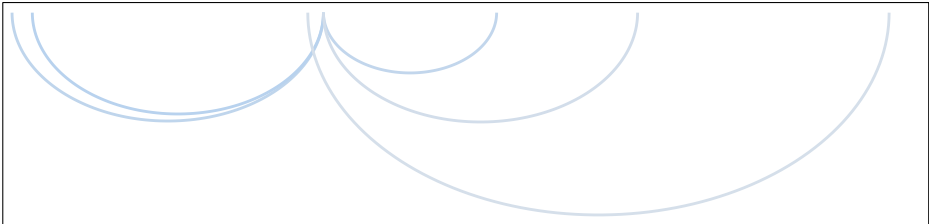
chr3:1808346–1908847 Marker Peak Tracks for: L5.6.NR4A2



L5.6.NR4A2

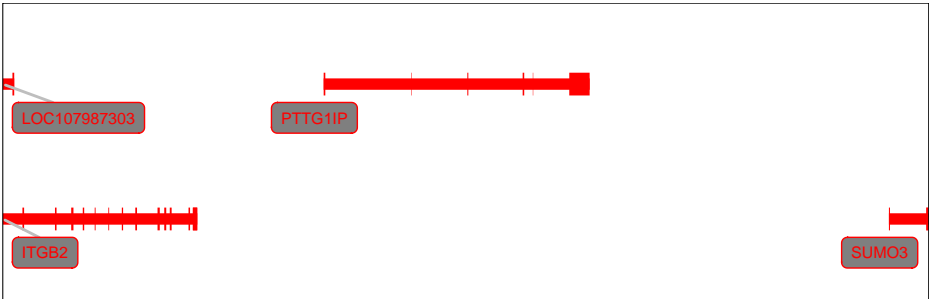


Peaks



Peak2GeneLinks

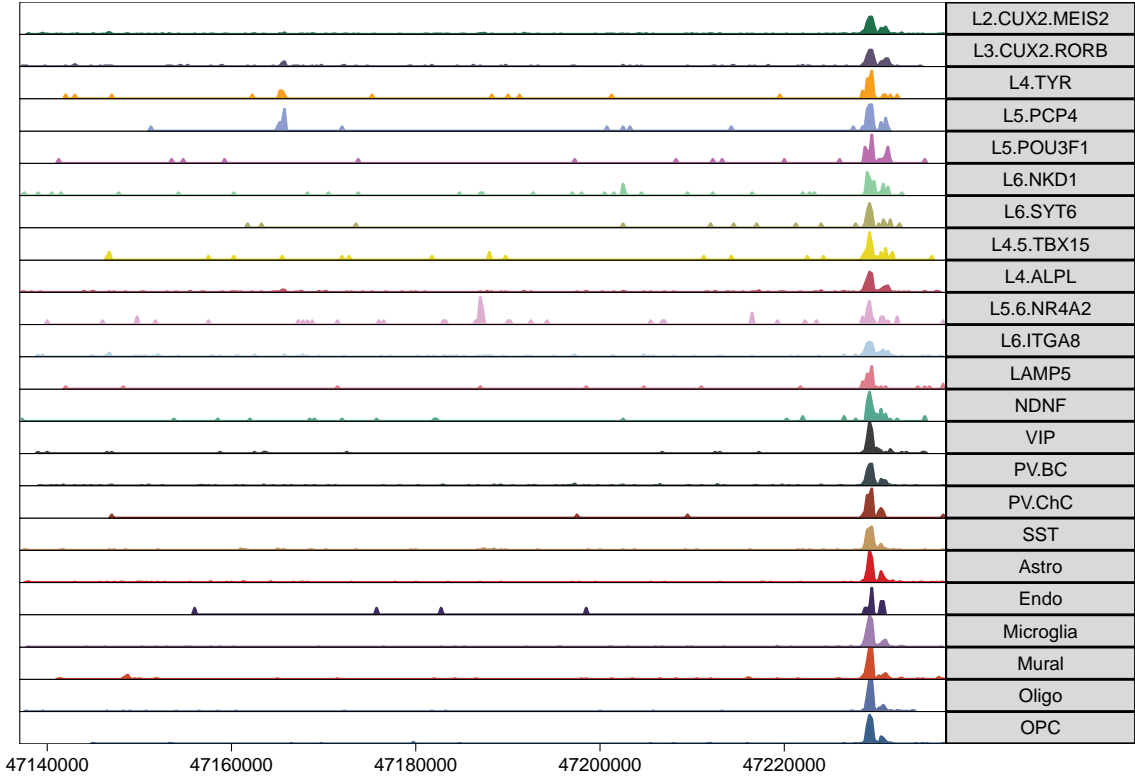
value



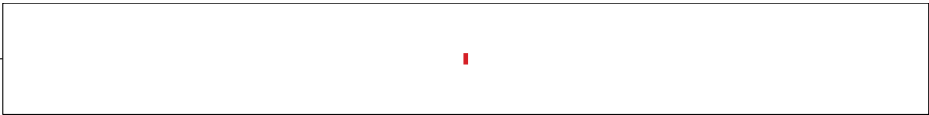
Genes

chr13:47137021-47237522 Marker Peak Tracks for: L5.6.NR4A2

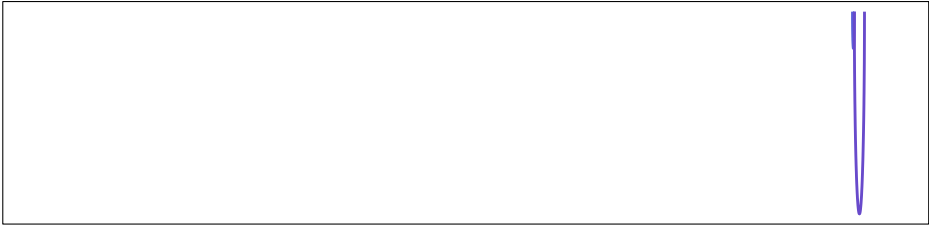
Coverage
(Norm. ATAC Signal Range 0-1.88) by ReadsInTSS)



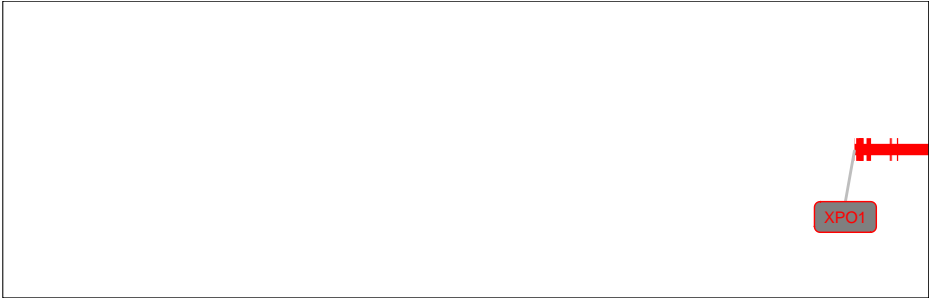
L5.6.NR4A2



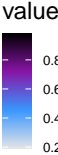
Peaks



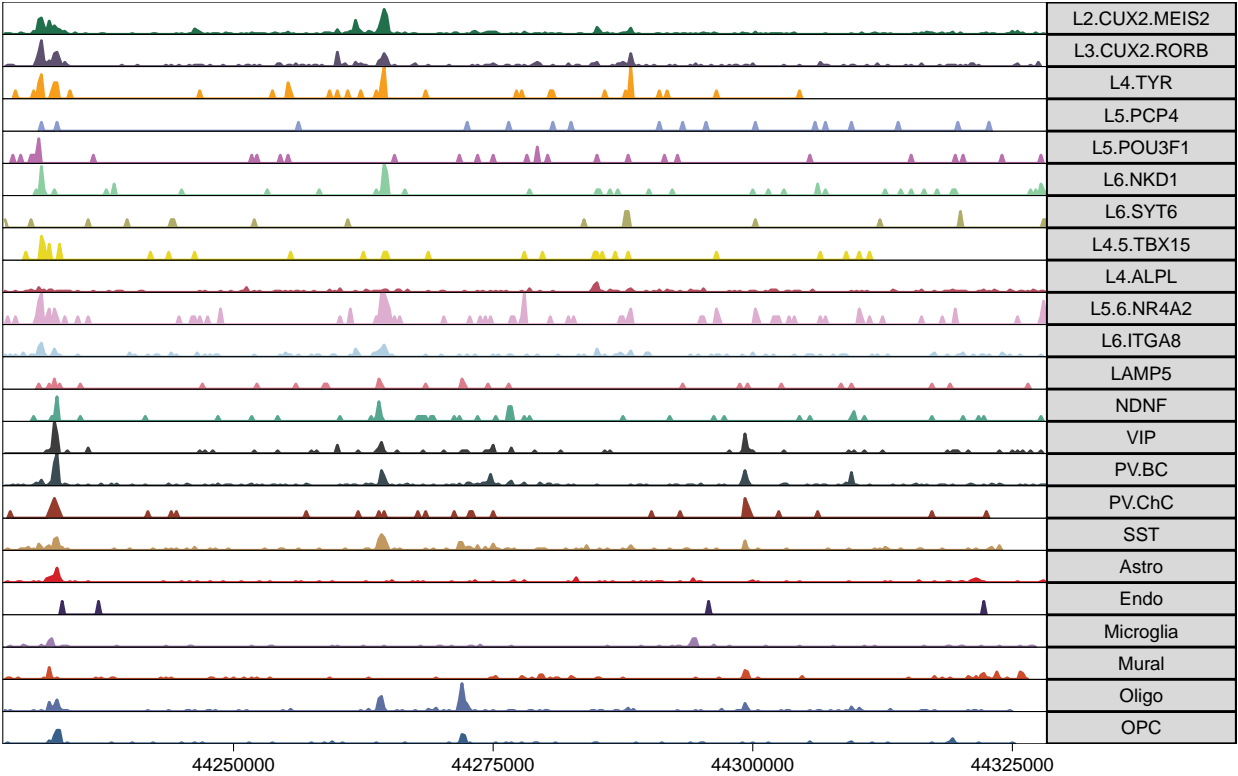
Peak2GeneLinks



Genes



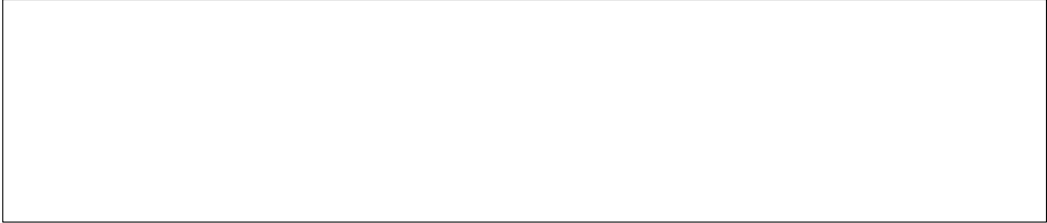
Coverage
(Norm. ATAC Signal Range (0–0.93) by ReadsInTSS)



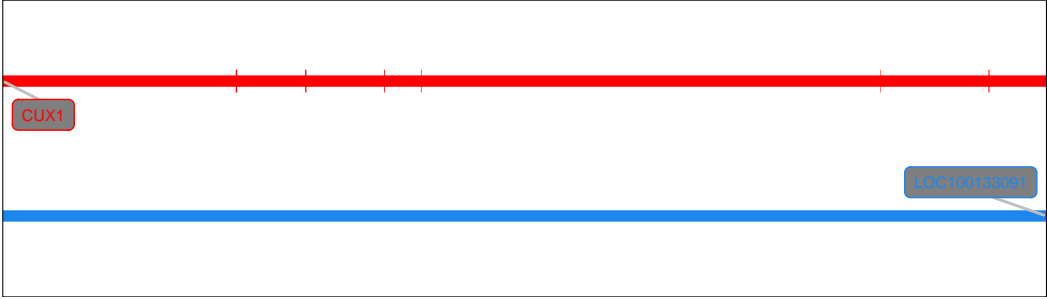
L5.6.NR4A2

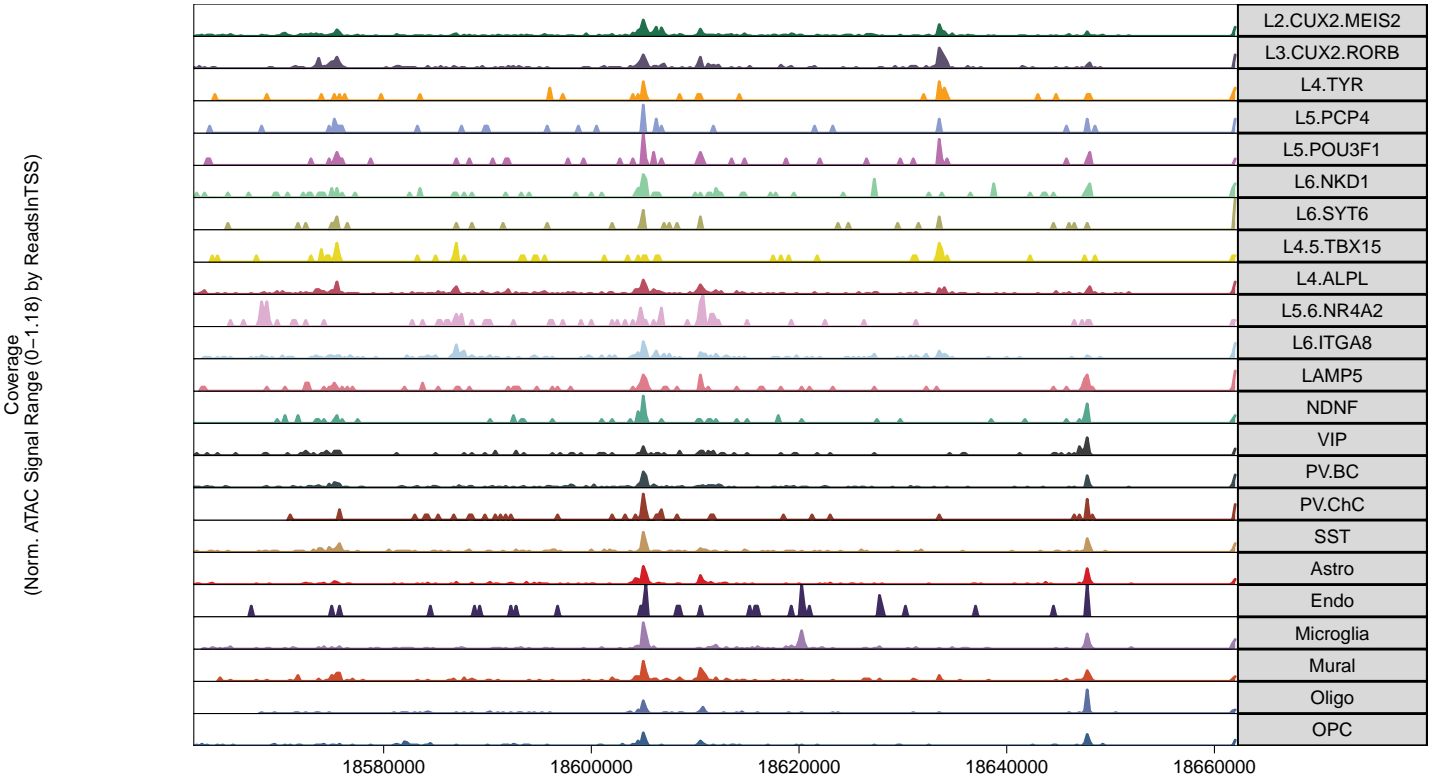


Peaks



LoopTrack

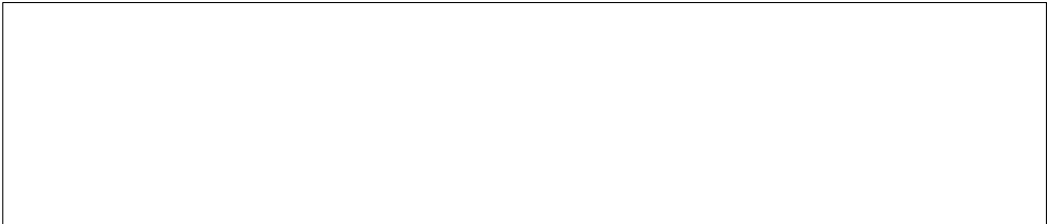




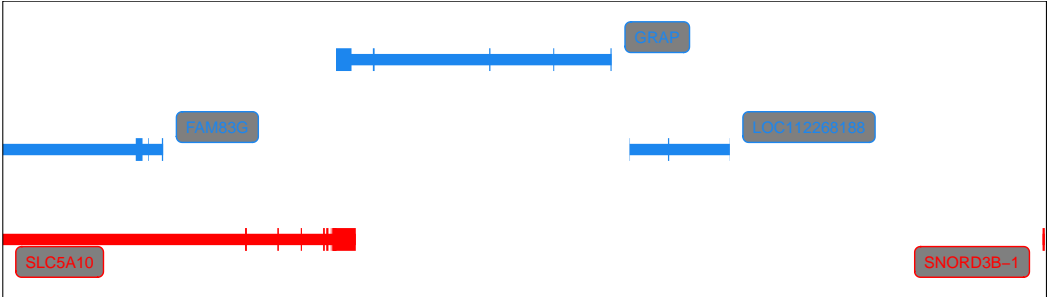
L5.6.NR4A2



Peaks



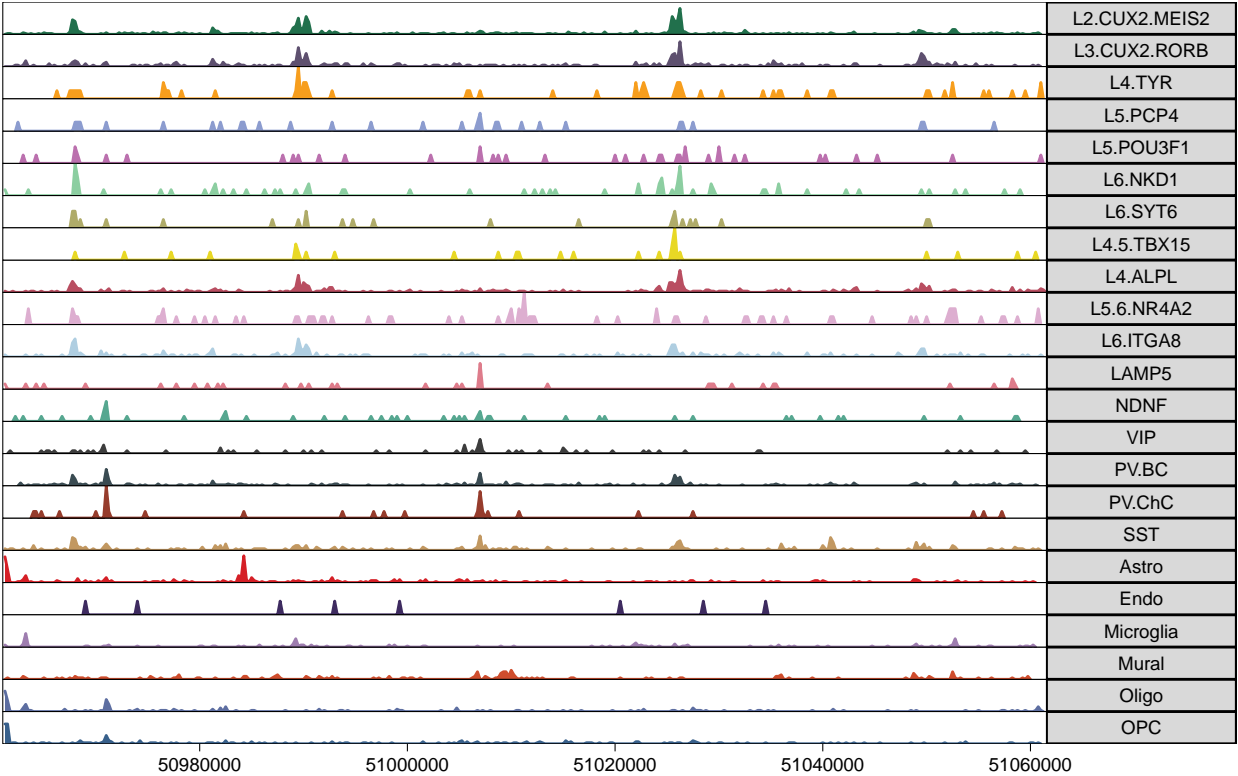
LoopTrack



Genes

chr5:50961038–51061539 Marker Peak Tracks for: L5.6.NR4A2

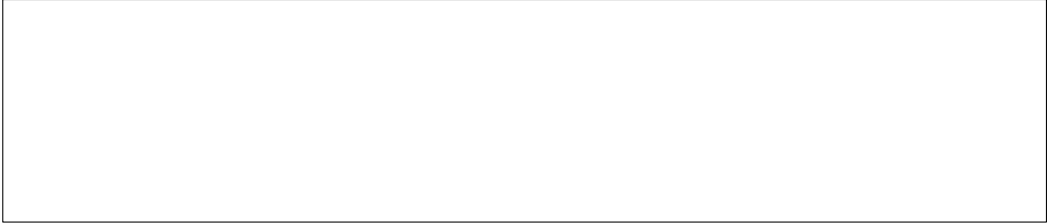
Coverage
(Norm. ATAC Signal Range (0–0.92) by ReadsInTSS)



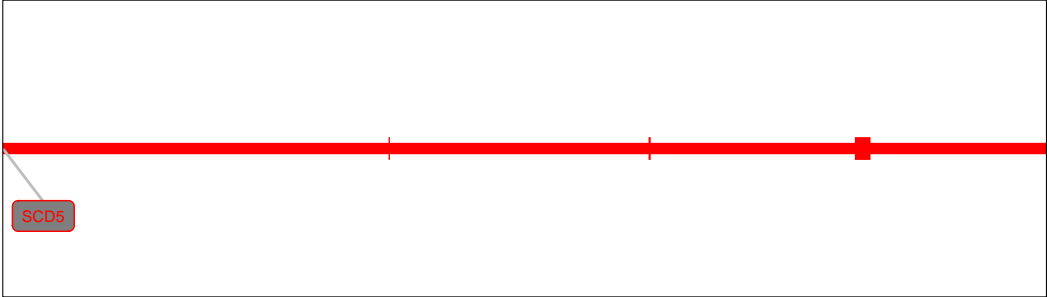
L5.6.NR4A2



Peaks

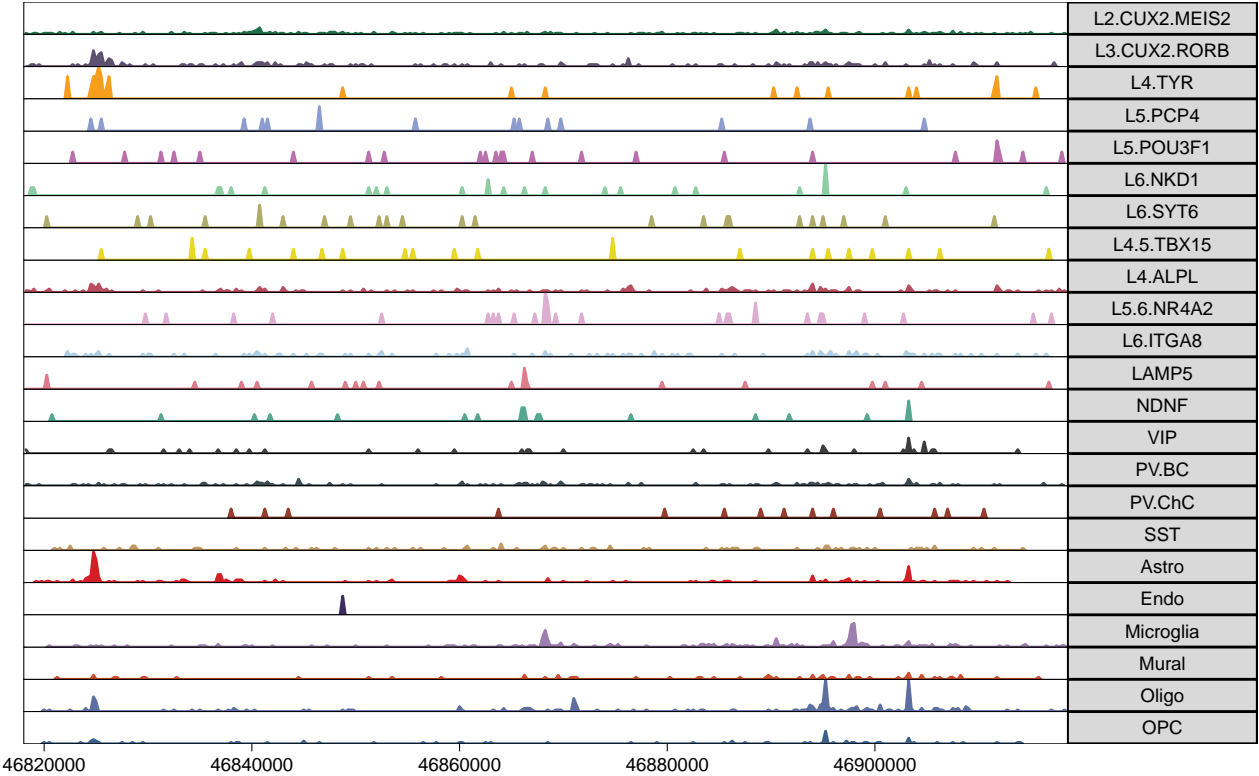


LoopTrack

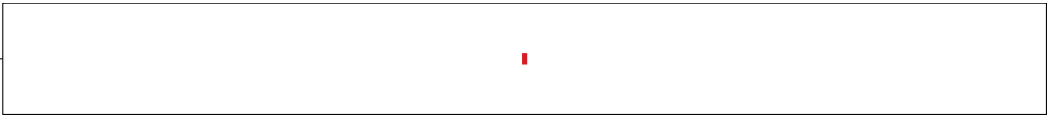


Genes

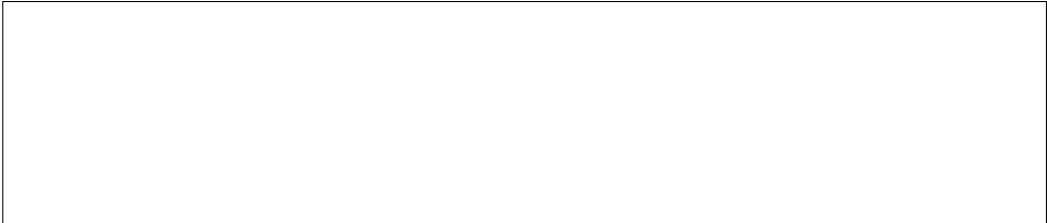
Coverage
(Norm. ATAC Signal Range 0–0.68) by ReadsInTSS)



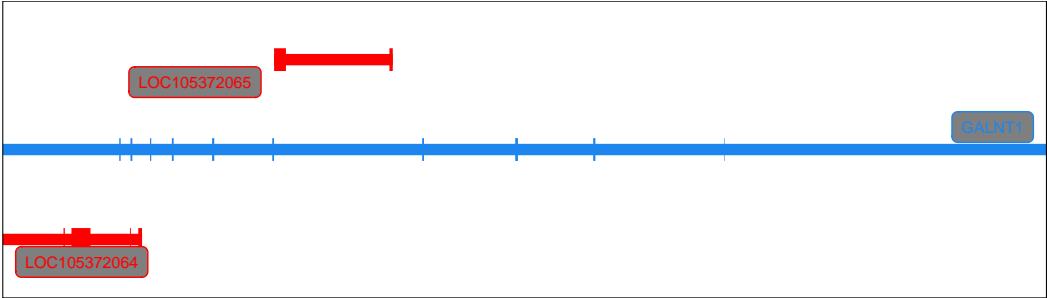
L5.6.NR4A2



Peaks

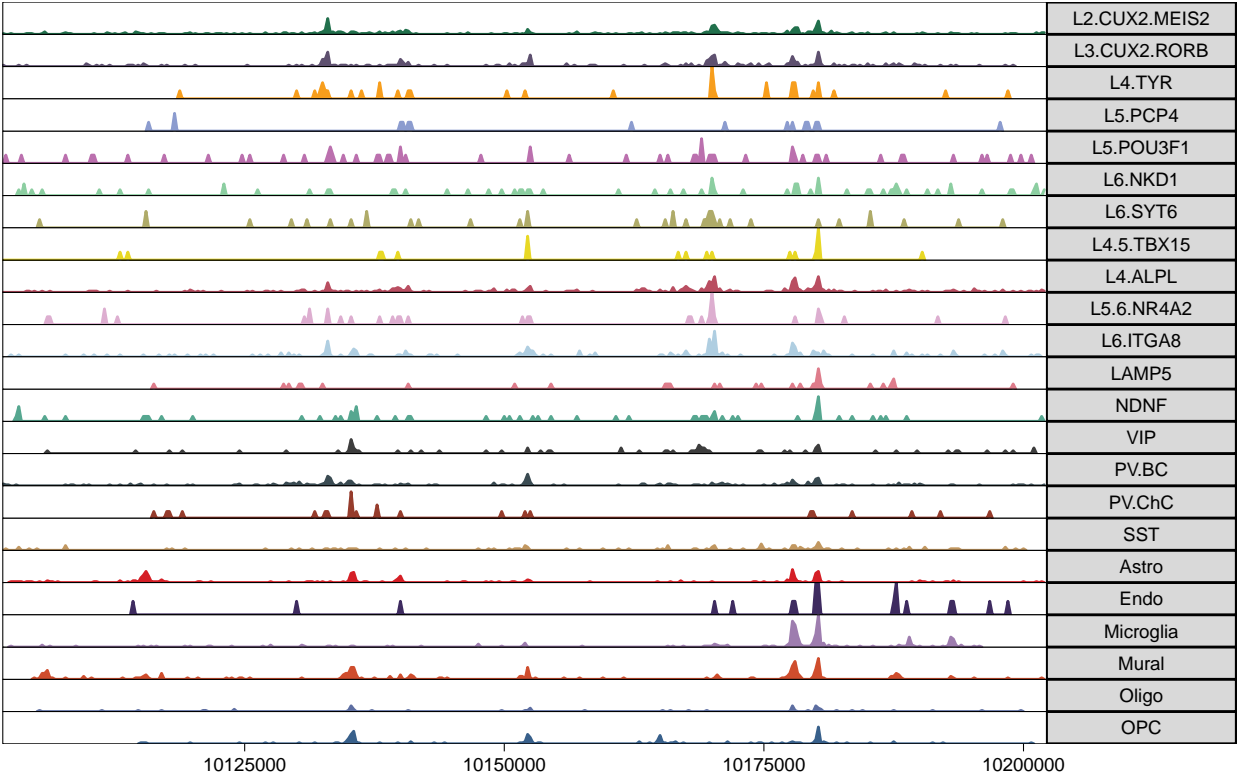


LoopTrack



Genes

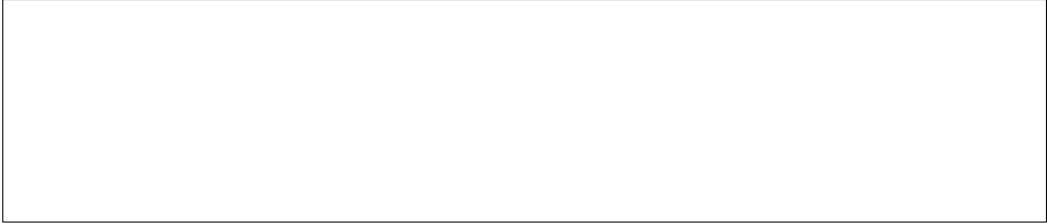
Coverage
(Norm. ATAC Signal Range (0–0.93) by ReadsInTSS)



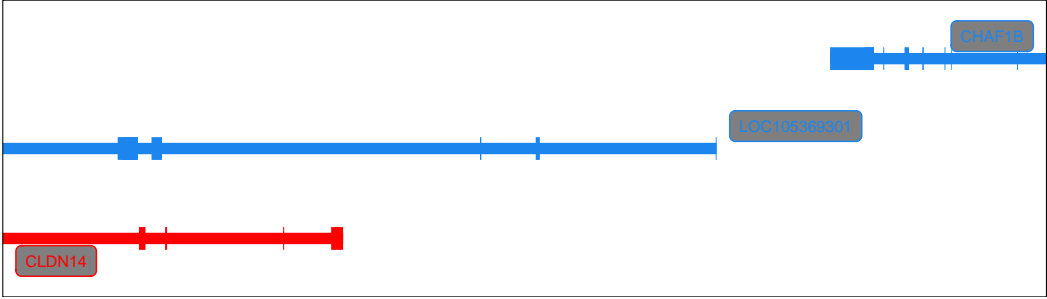
L5.6.NR4A2



Peaks



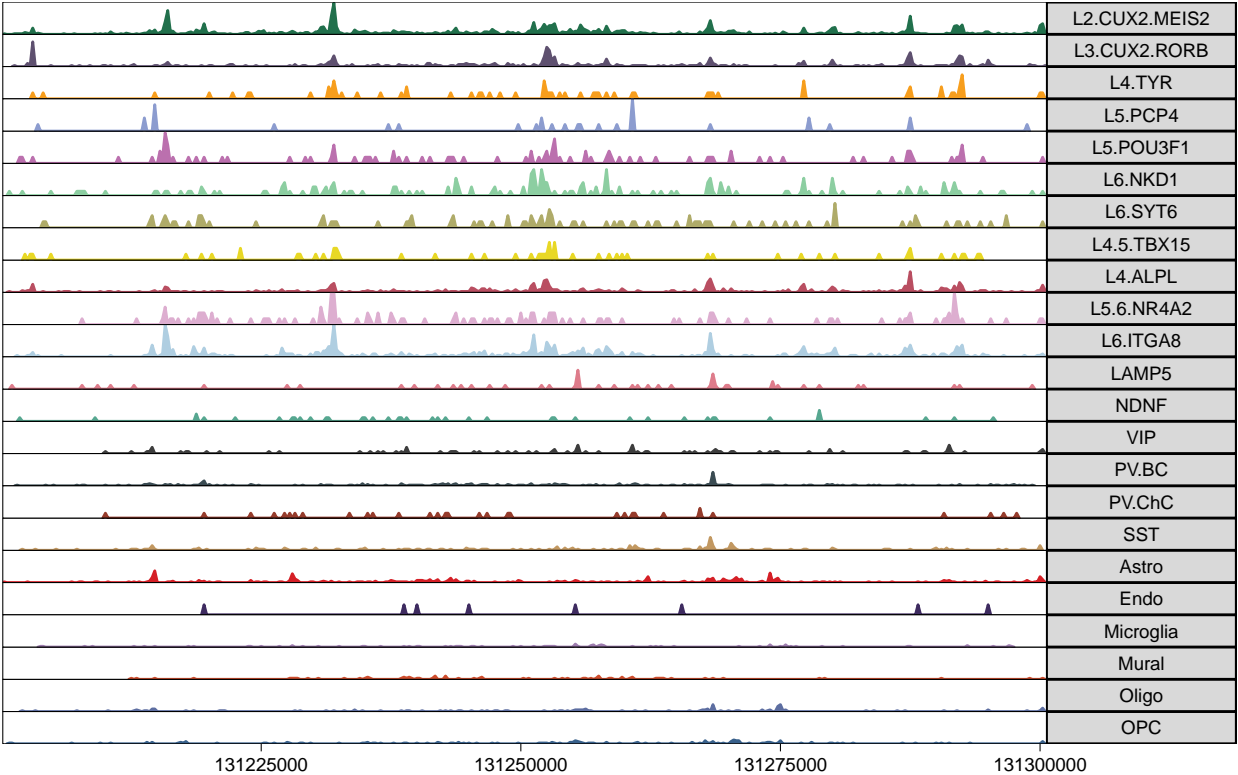
LoopTrack



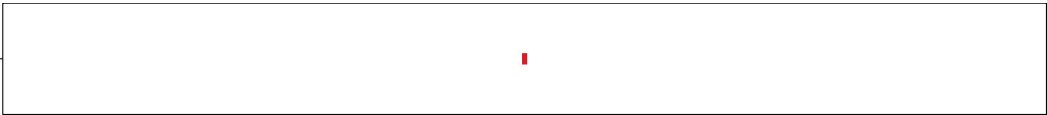
Genes

chr4:131200115-131300616 Marker Peak Tracks for: L5.6.NR4A2

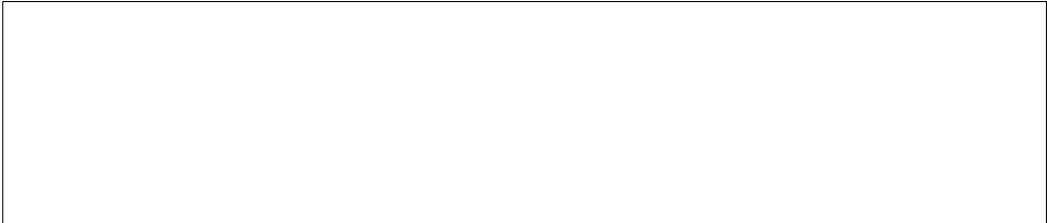
Coverage
(Norm. ATAC Signal Range 0-1.25) by ReadsInTSS)



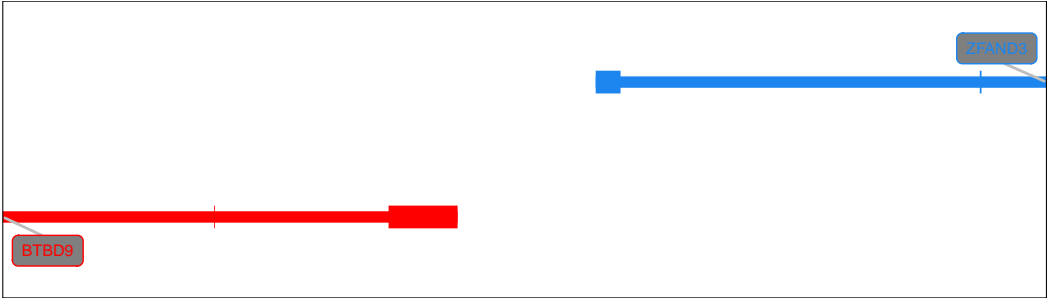
L5.6.NR4A2



Peaks

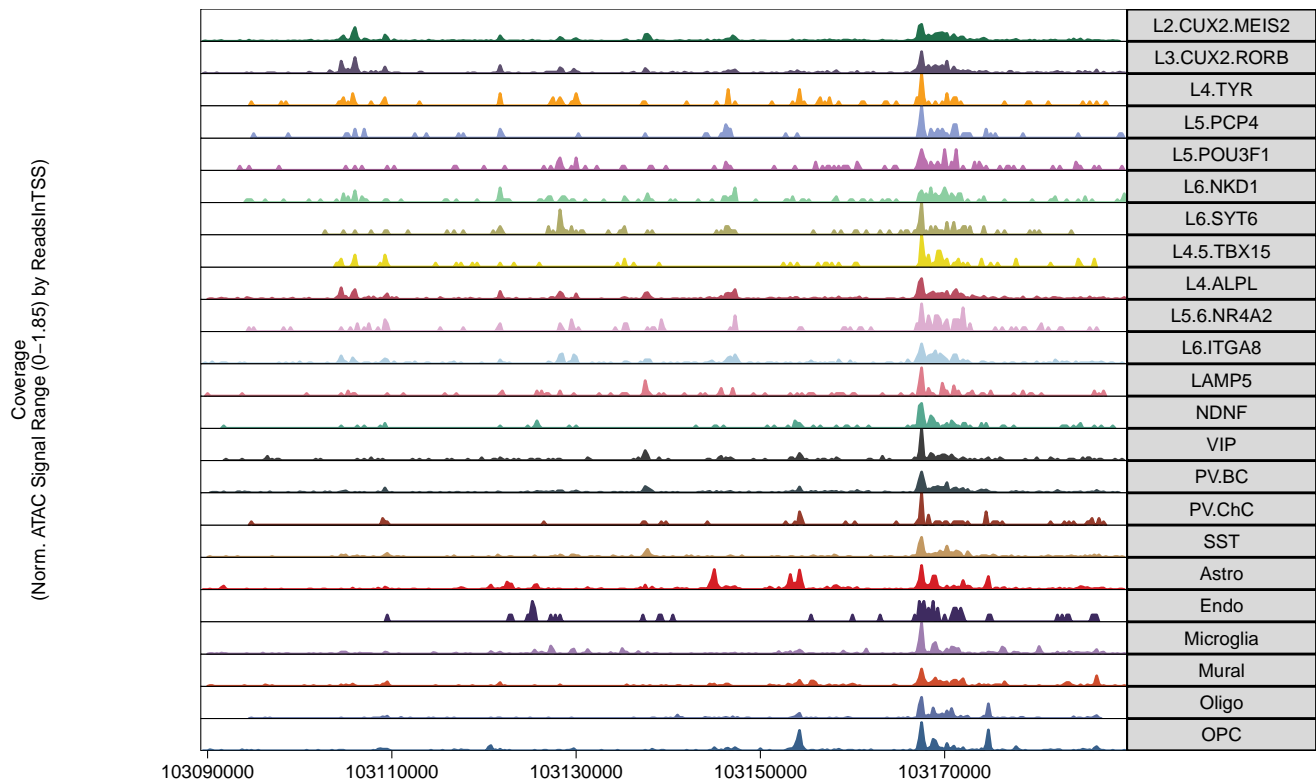


LoopTrack



Genes

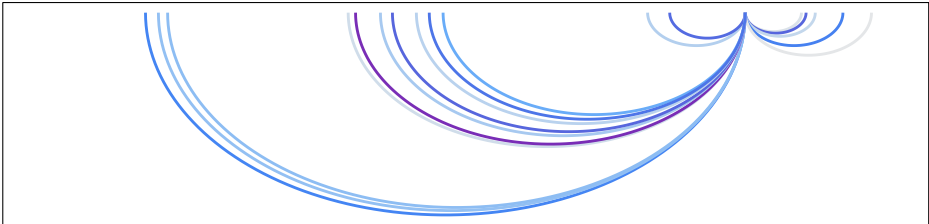
chr9:103089261–103189762 Marker Peak Tracks for: L5.6.NR4A2



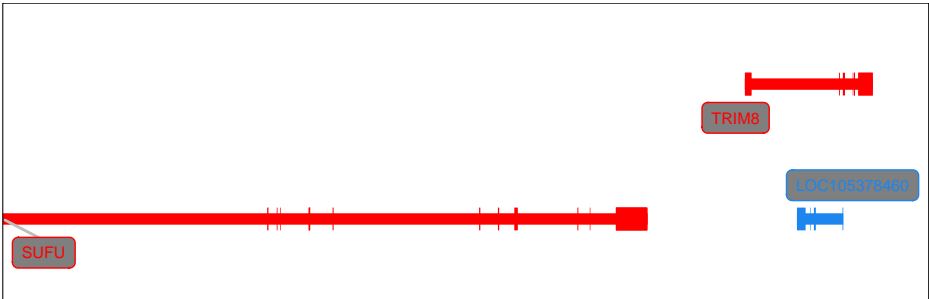
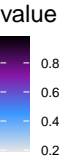
L5.6.NR4A2



Peaks

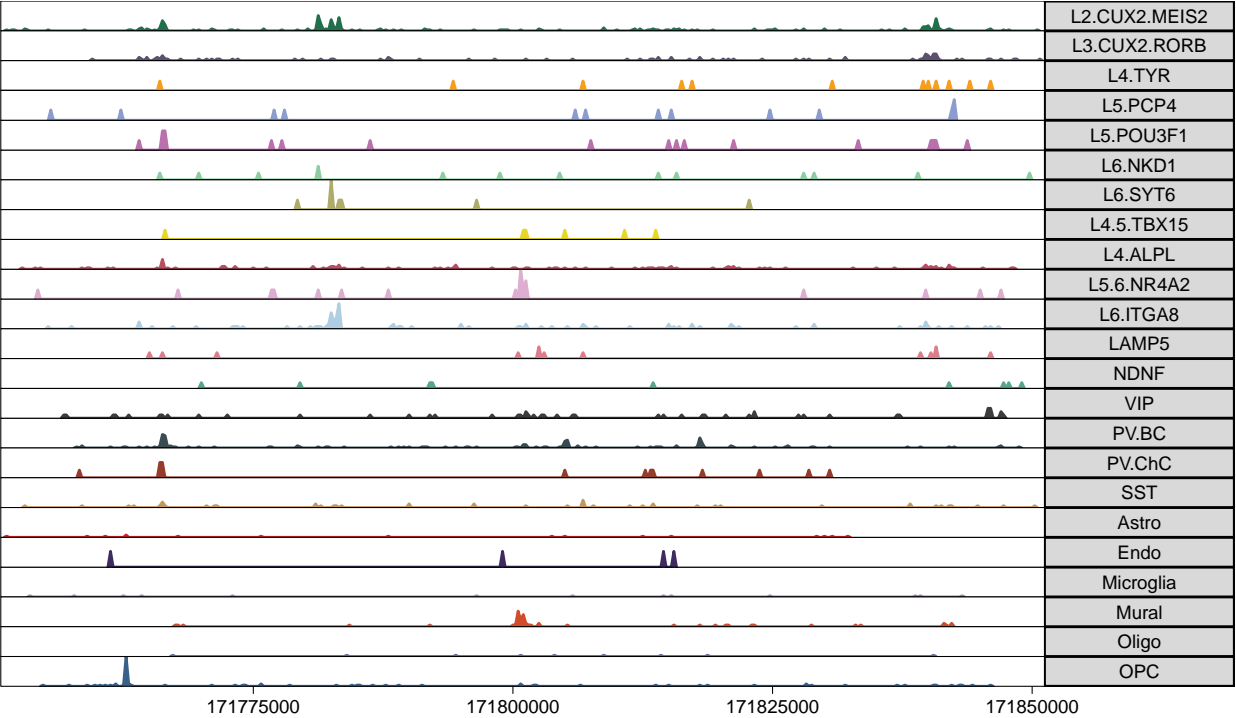


Peak2GeneLinks

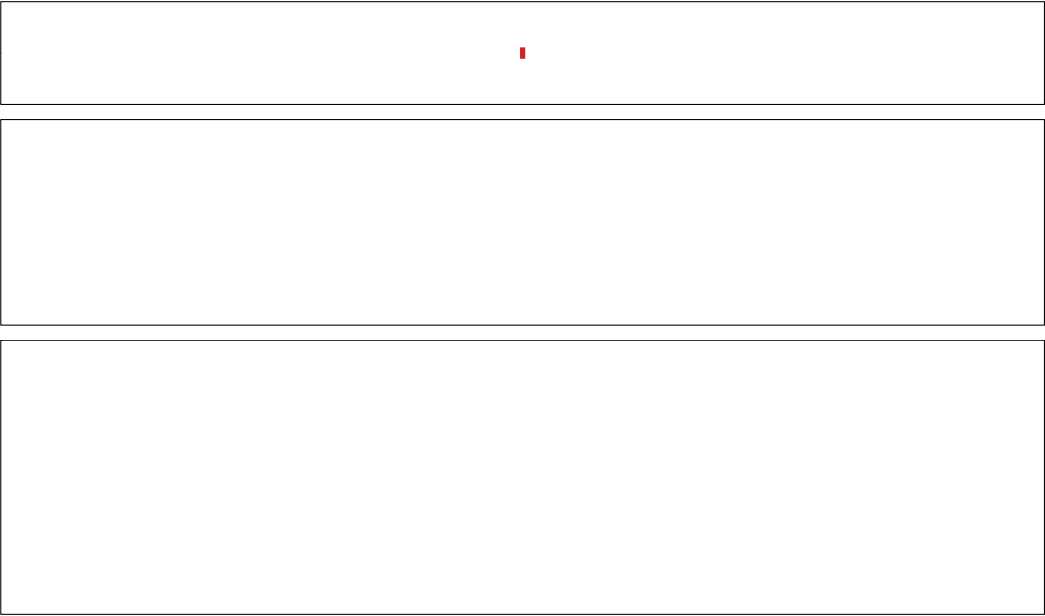


Genes

Coverage
(Norm. ATAC Signal Range (0–0.74) by Reads in TSS)



L5.6.NR4A2



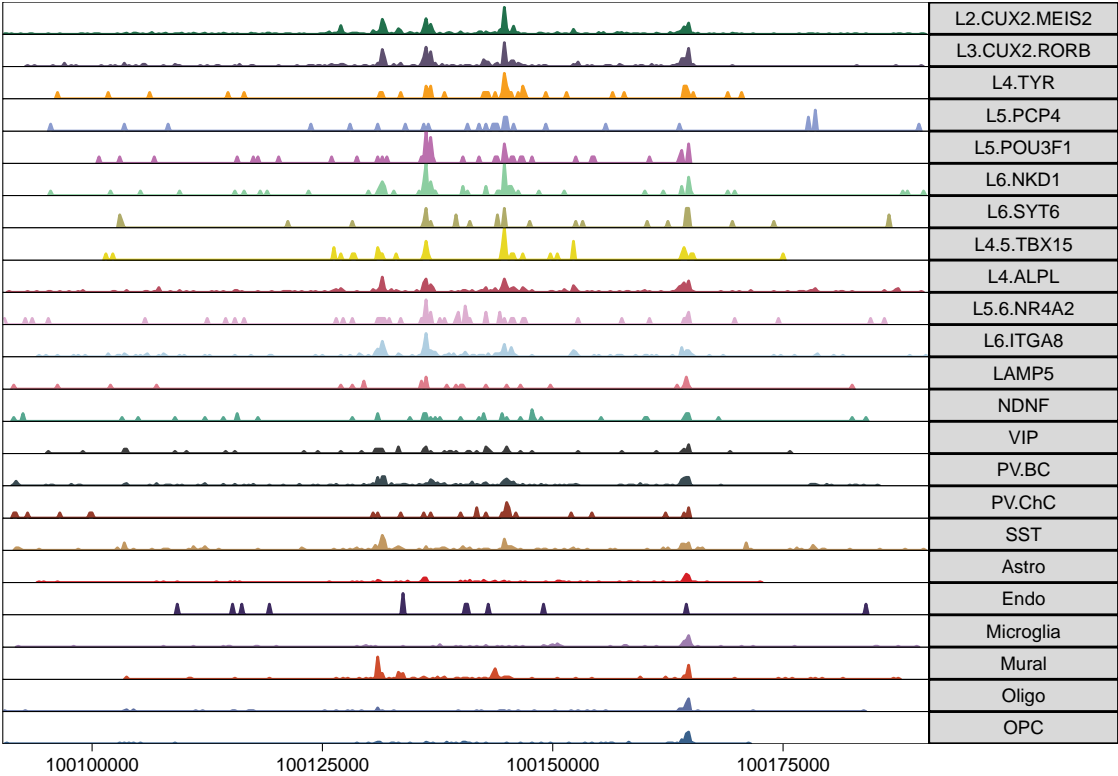
Peaks

LoopTrack

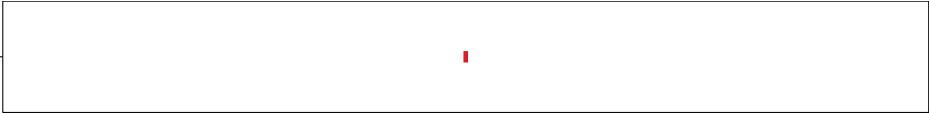
GeneTrack

chr8:100090308–100190809 Marker Peak Tracks for: L5.6.NR4A2

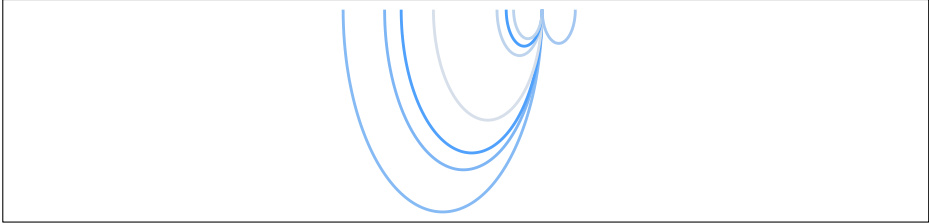
Coverage
(Norm. ATAC Signal Range (0–1.18) by ReadsInTSS)



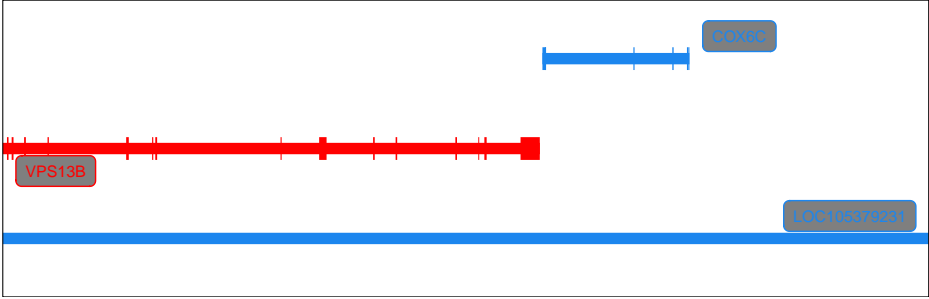
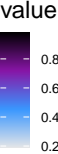
L5.6.NR4A2



Peaks

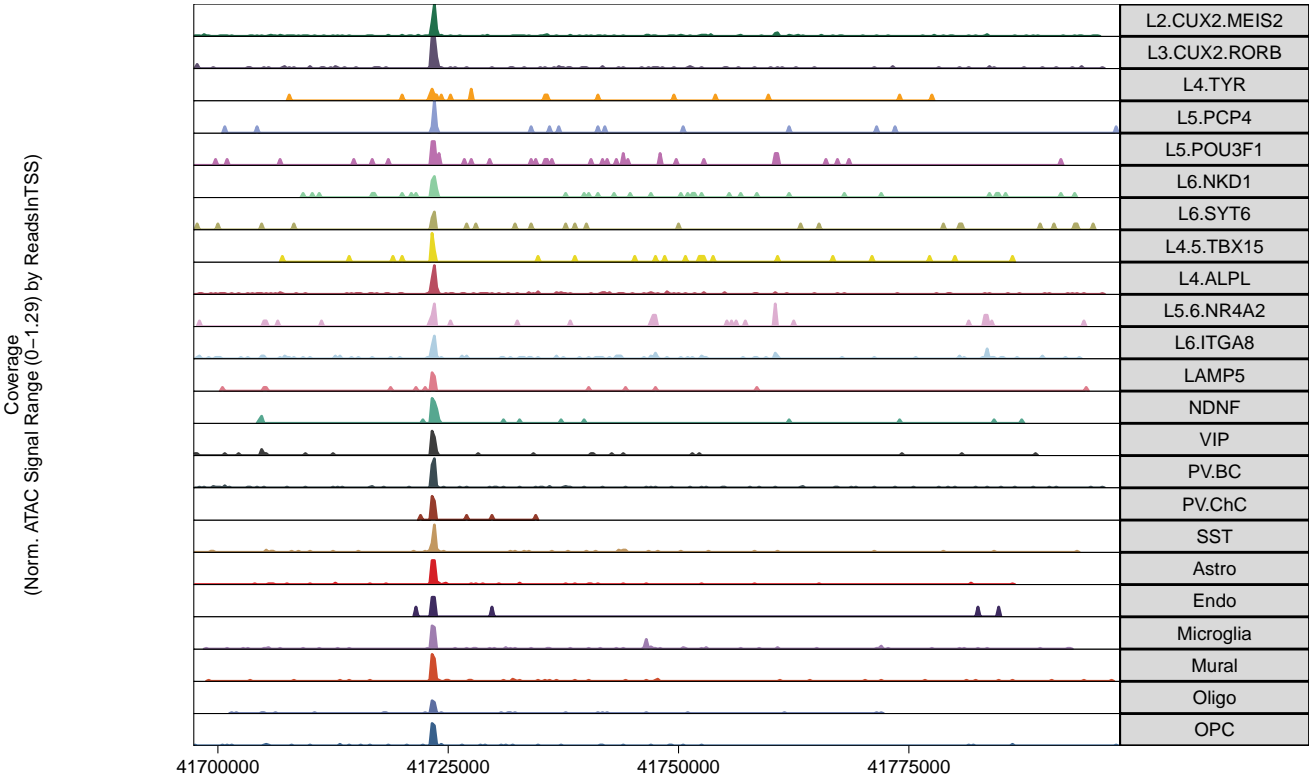


Peak2GeneLinks



Genes

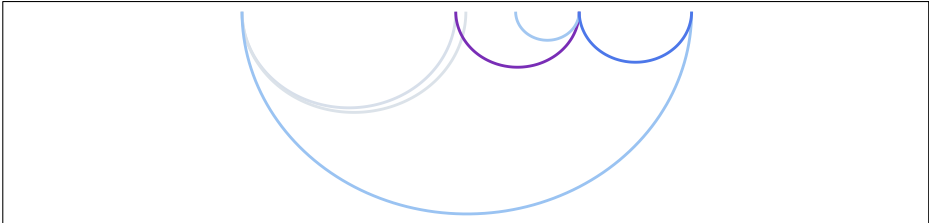
chr3:41697385–41797886 Marker Peak Tracks for: L5.6.NR4A2



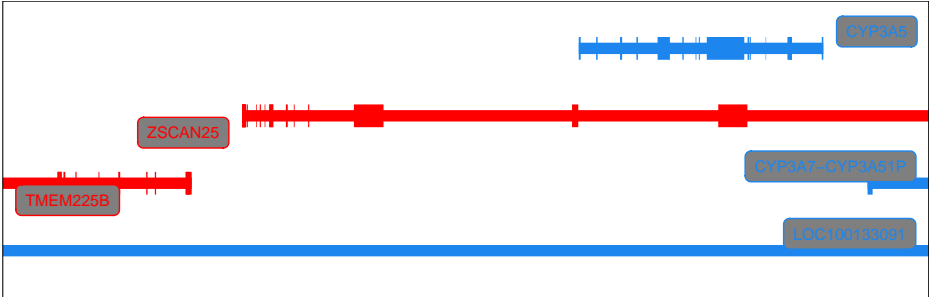
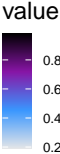
L5.6.NR4A2



Peaks

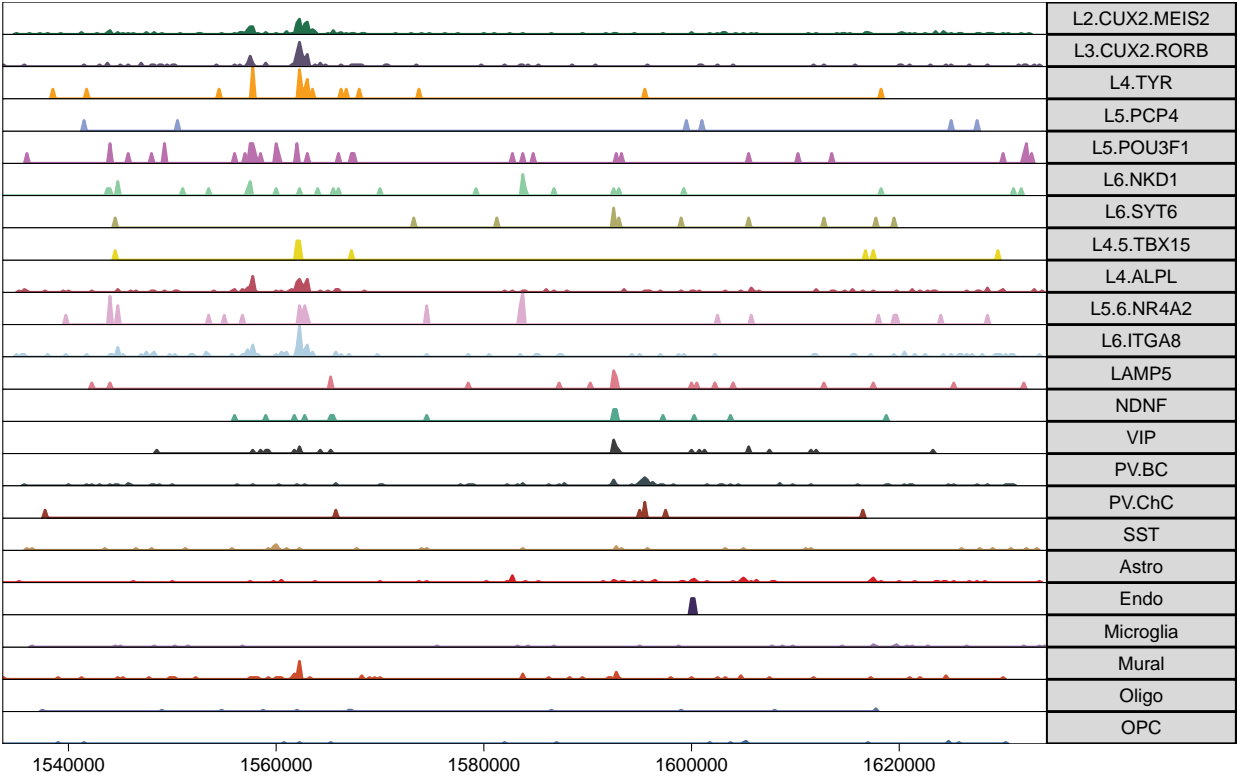


Peak2GeneLinks

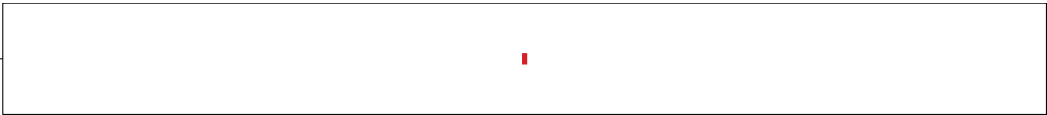


Genes

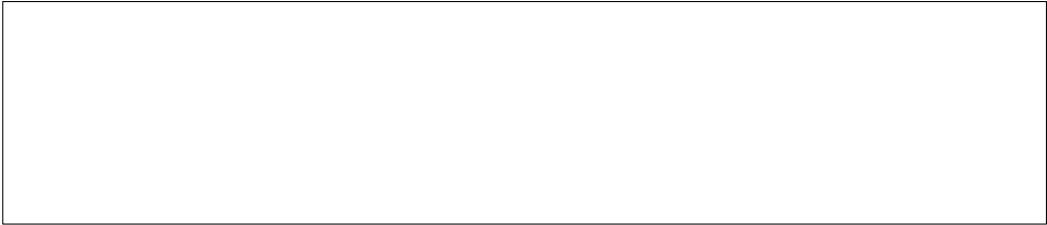
Coverage
(Norm. ATAC Signal Range (0–0.77) by ReadsInTSS)



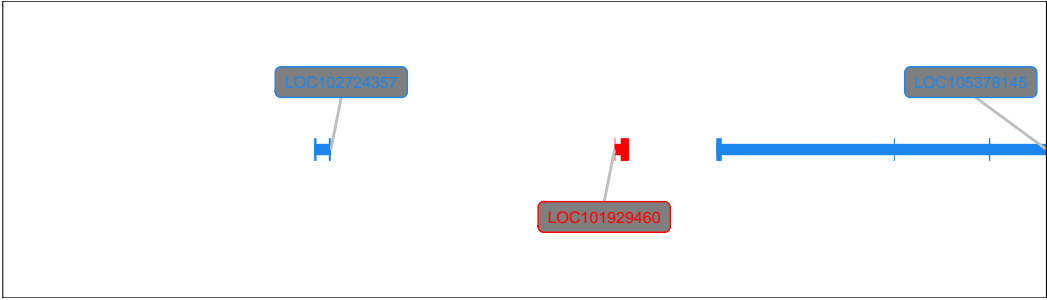
L5.6.NR4A2



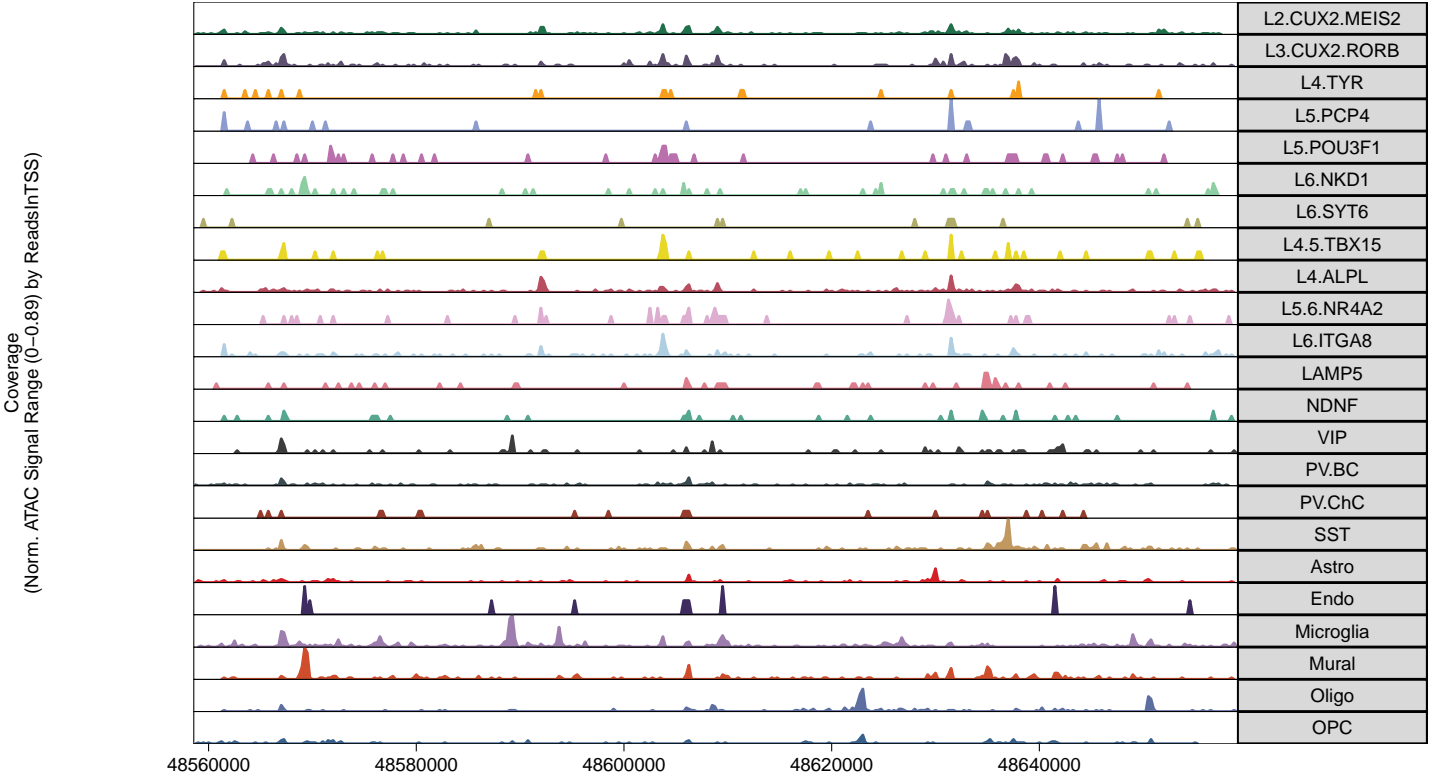
Peaks



LoopTrack



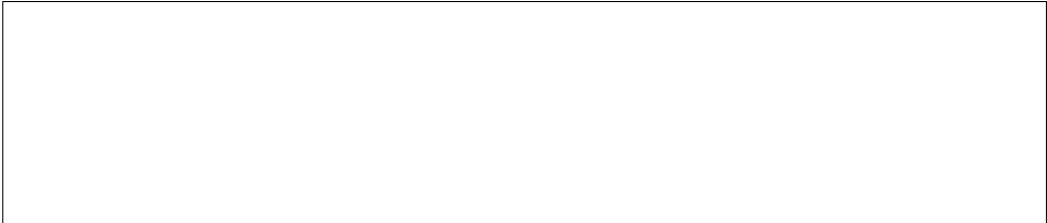
Genes



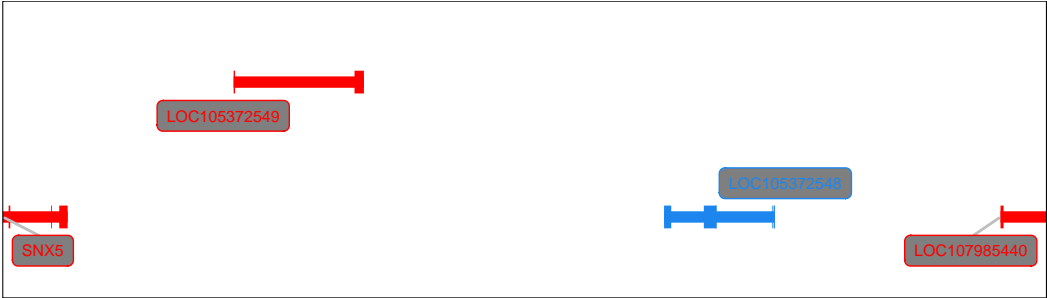
L5.6.NR4A2



Peaks

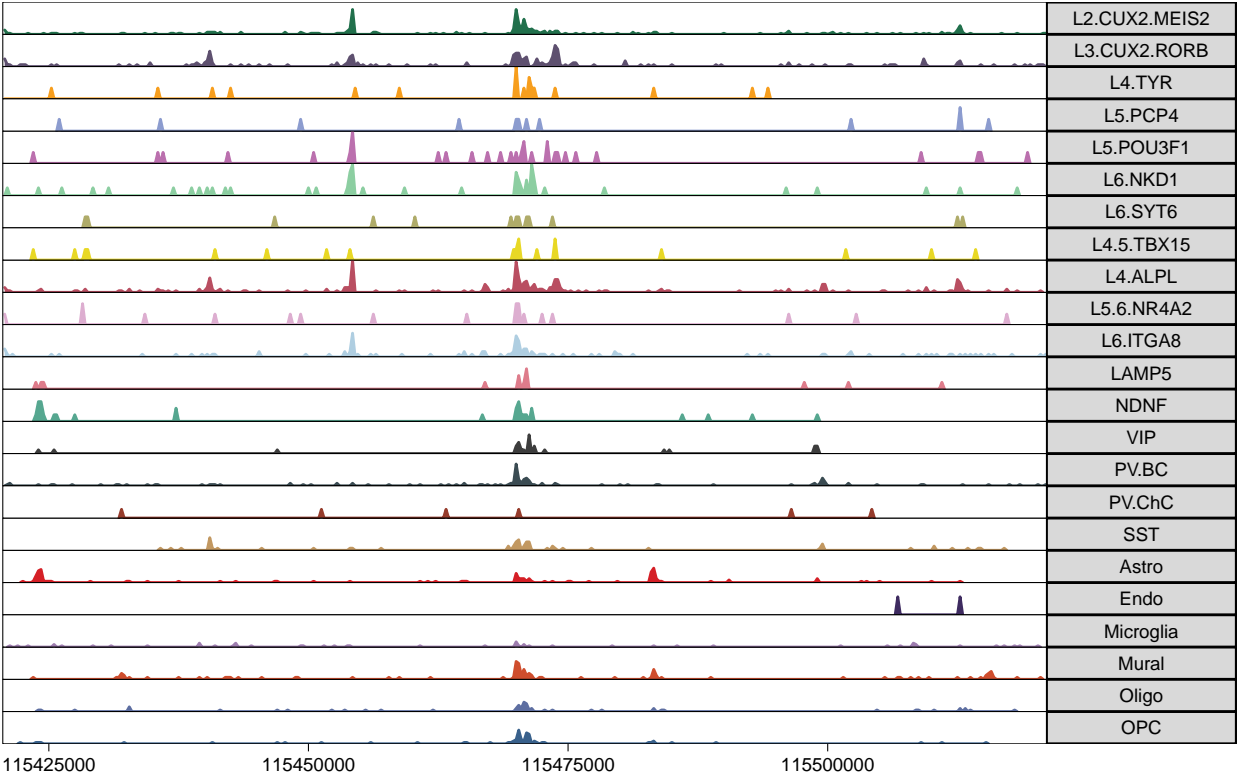


LoopTrack

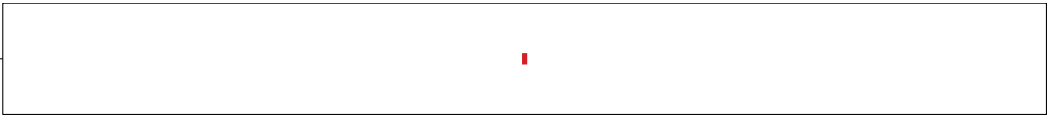


Genes

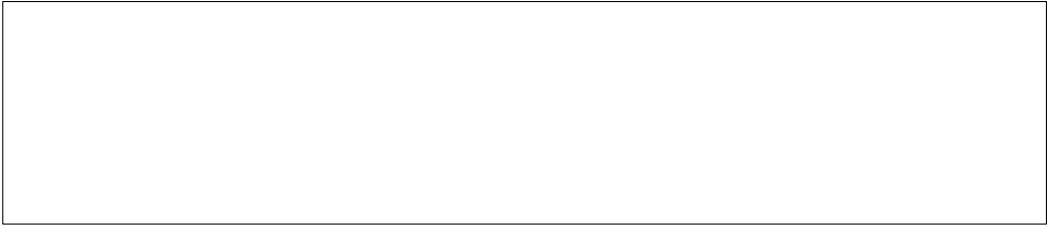
Coverage
(Norm. ATAC Signal Range (0–0.7) by Reads in TSS)



L5.6.NR4A2



Peaks



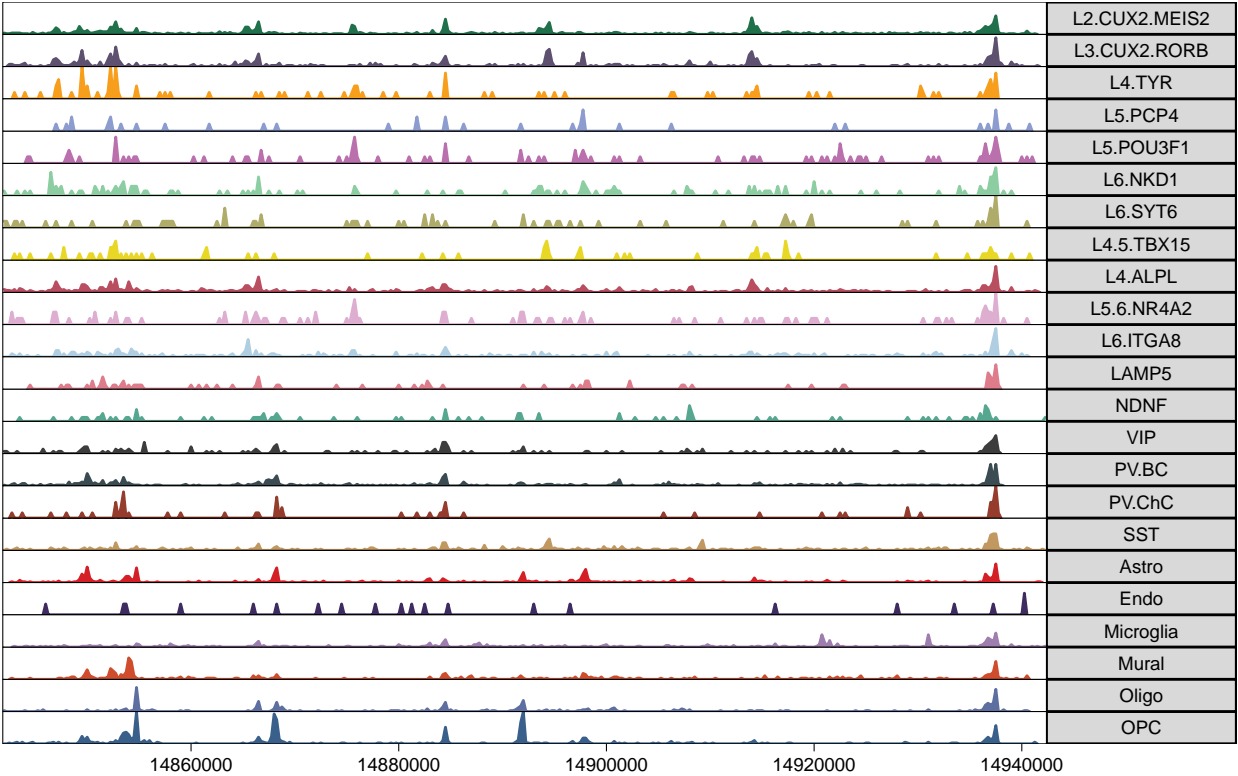
LoopTrack



Genes

chr19:14841872–14942373 Marker Peak Tracks for: L5.6.NR4A2

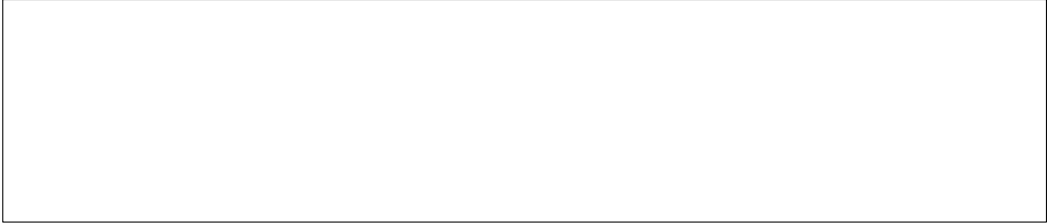
Coverage
(Norm. ATAC Signal Range 0–1.17) by ReadsInTSS)



L5.6.NR4A2



Peaks



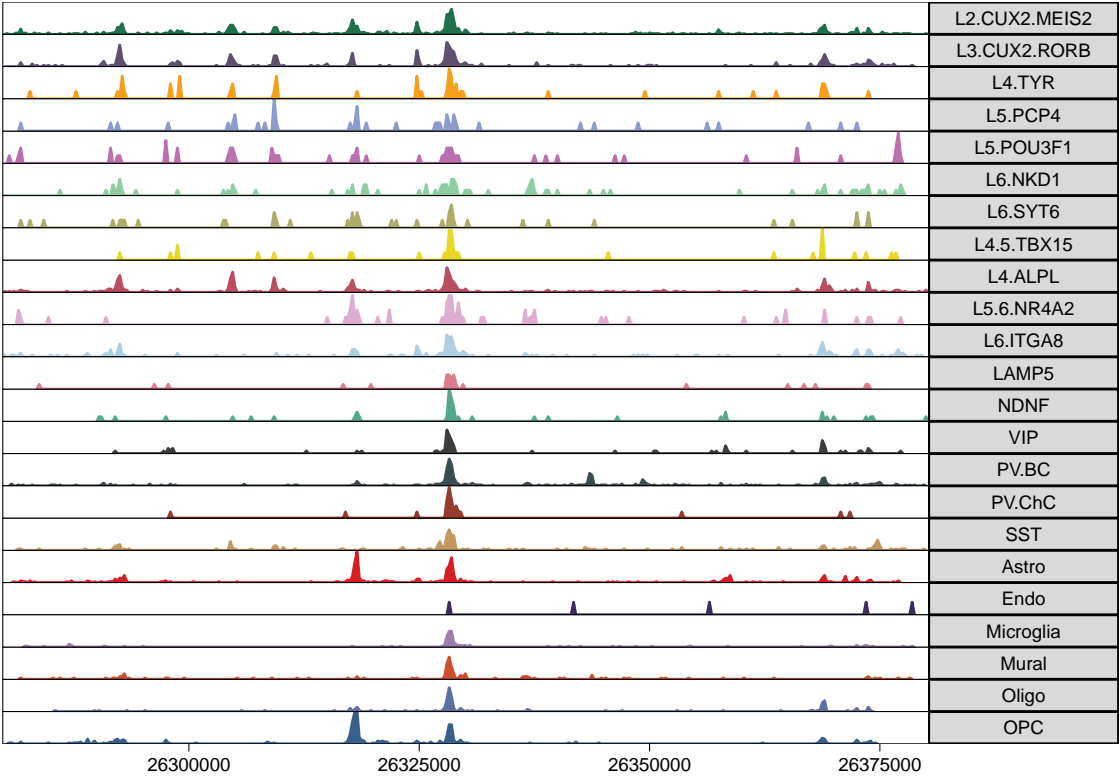
LoopTrack



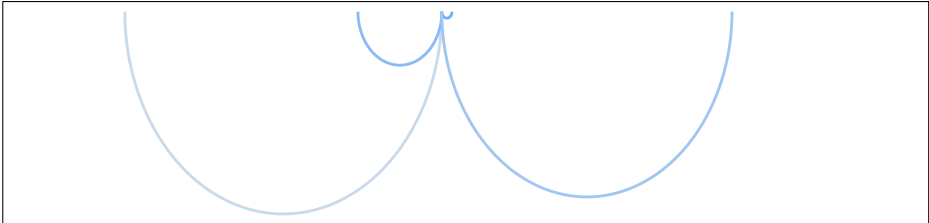
Genes

chr4:26279799–26380300 Marker Peak Tracks for: L5.6.NR4A2

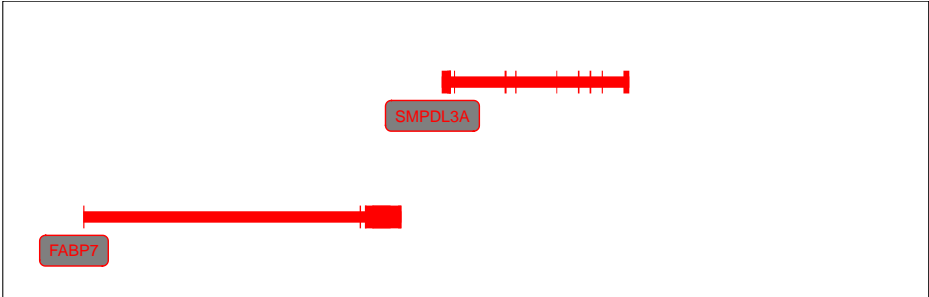
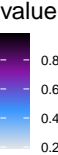
Coverage
(Norm. ATAC Signal Range (0–1) by ReadsInTSS)



L5.6.NR4A2

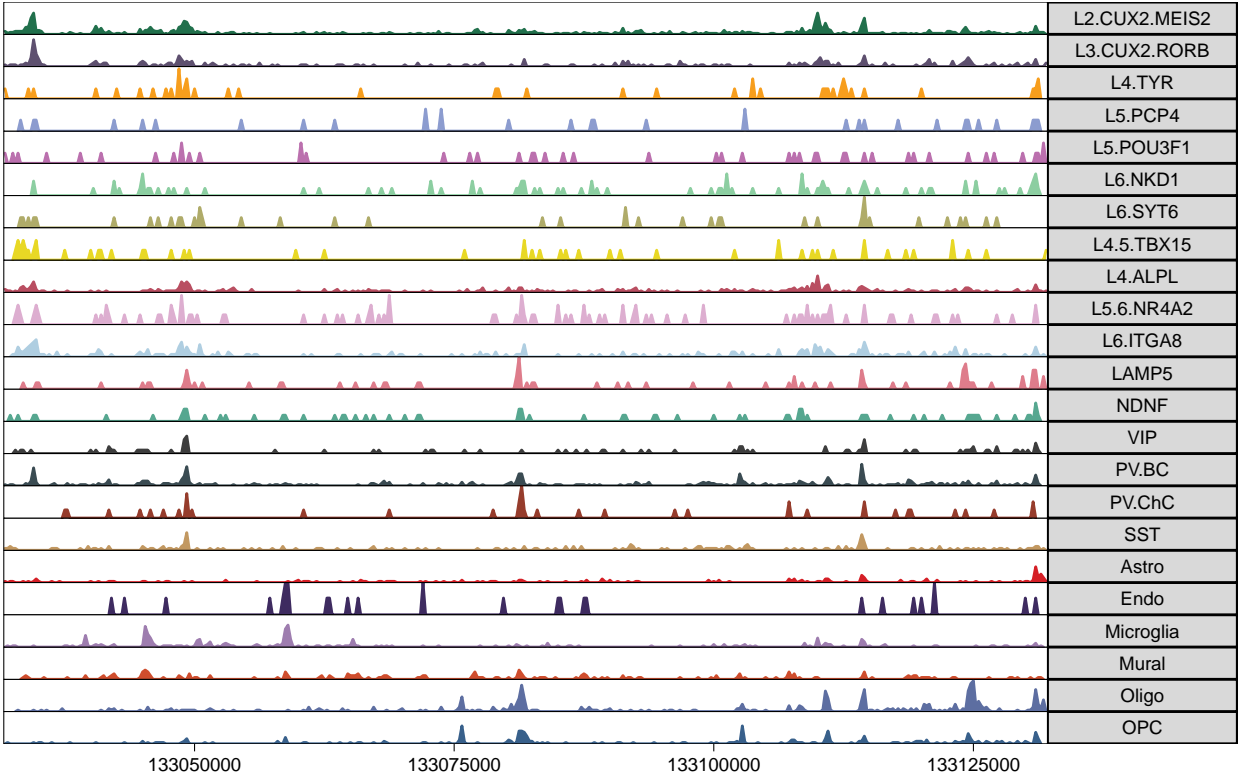


Peak2GeneLinks

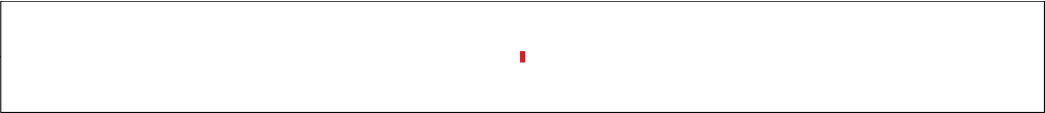


Genes

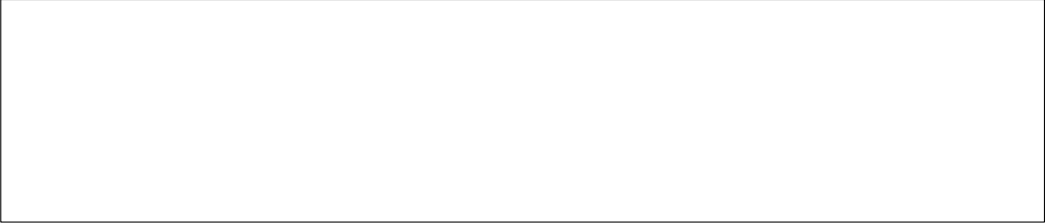
Coverage
(Norm. ATAC Signal Range (0-0.75) by ReadsInTSS)



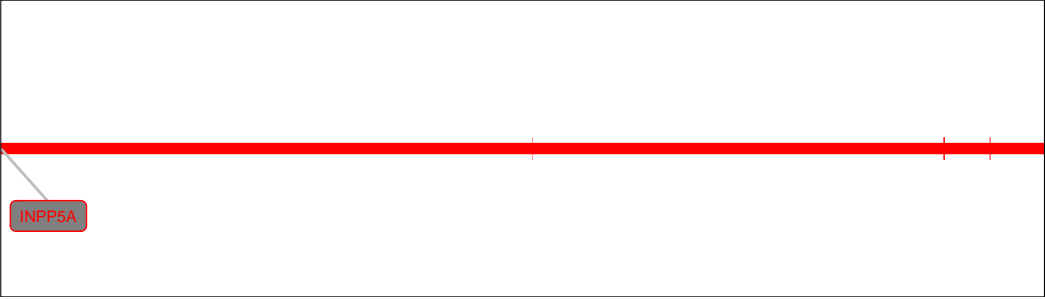
L5.6.NR4A2



Peaks



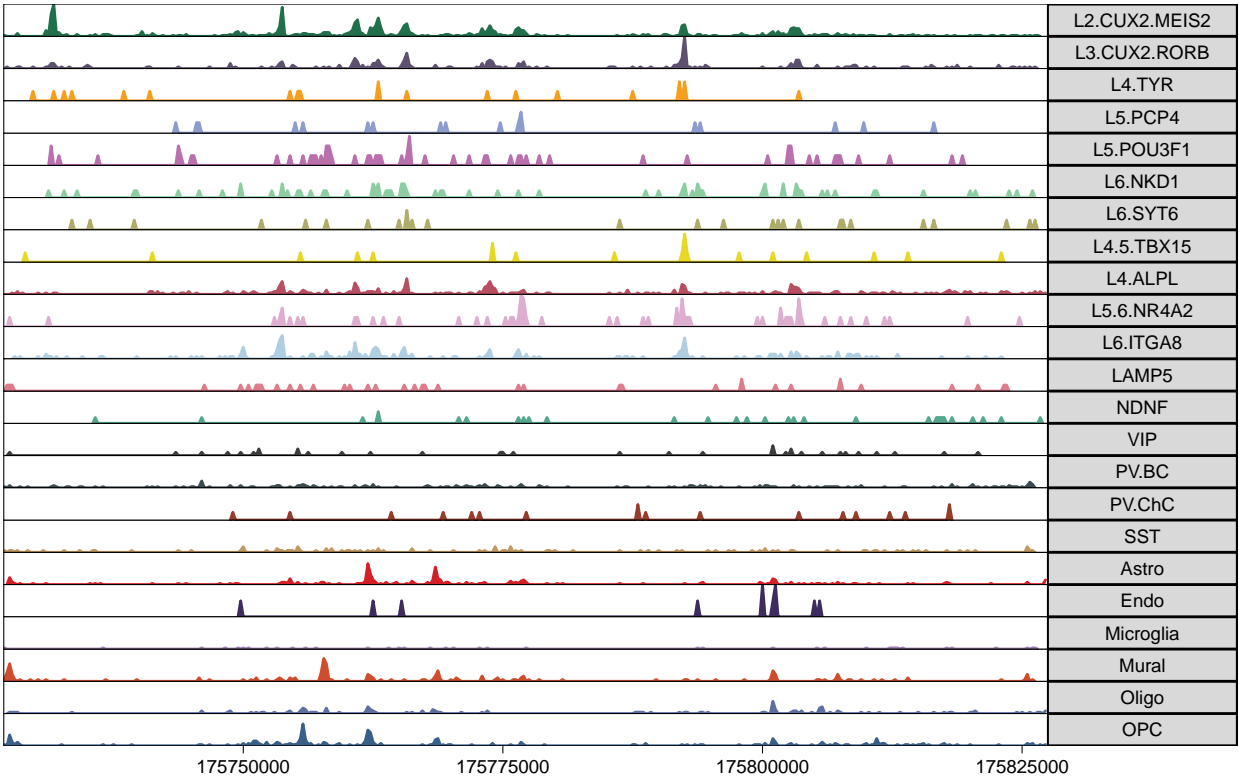
LoopTrack



Genes

chr6:175726939–175827440 Marker Peak Tracks for: L5.6.NR4A2

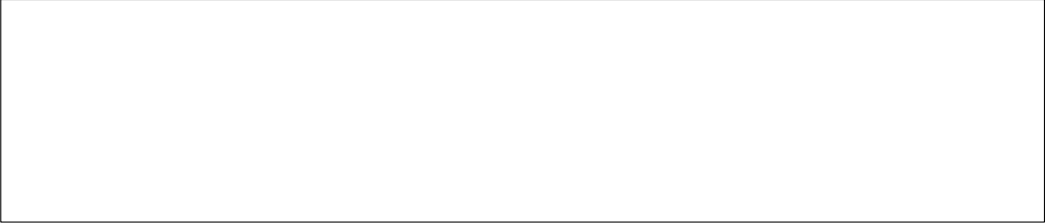
Coverage
(Norm. ATAC Signal Range (0–0.79) by ReadsInTSS)



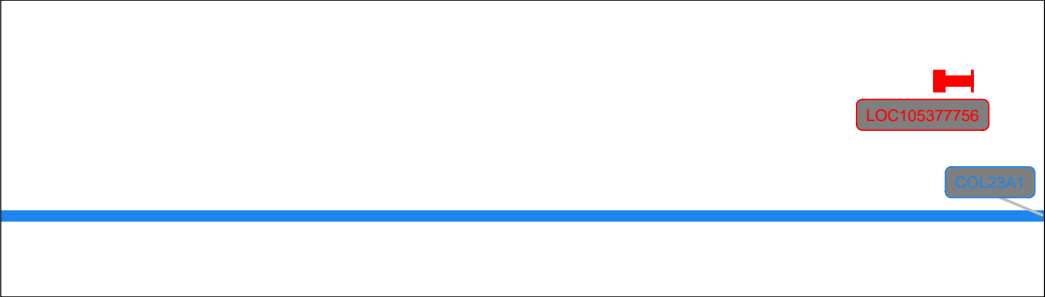
L5.6.NR4A2



Peaks

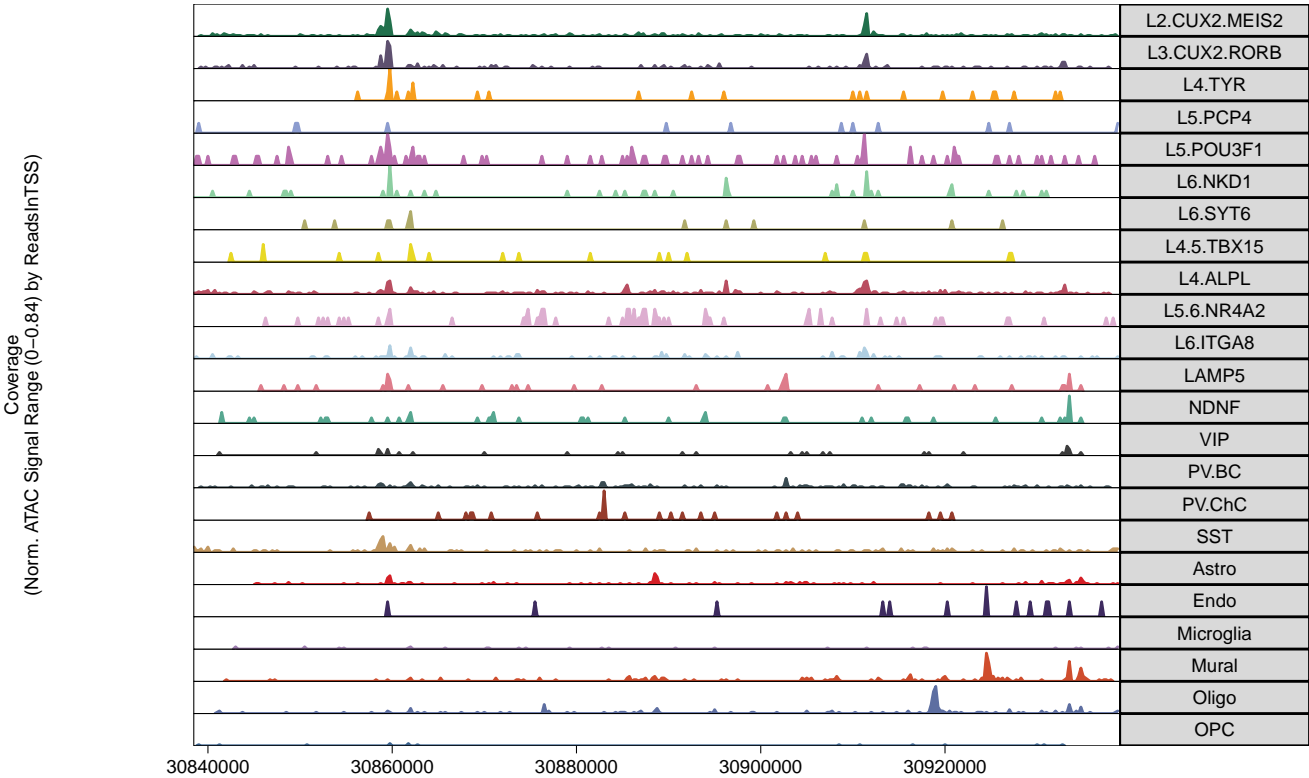


LoopTrack

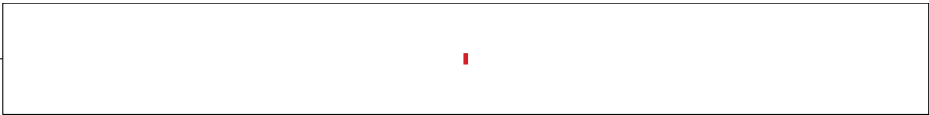


Genes

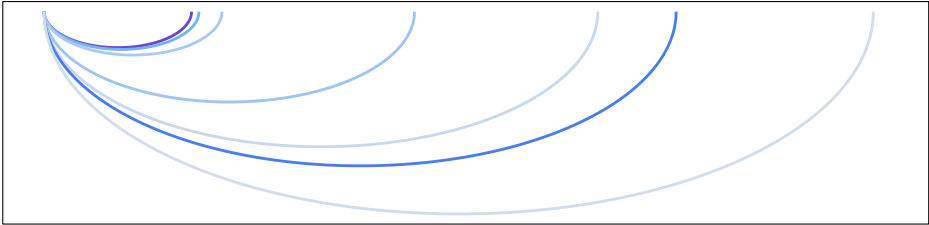
chr11:30838463–30938964 Marker Peak Tracks for: L5.6.NR4A2



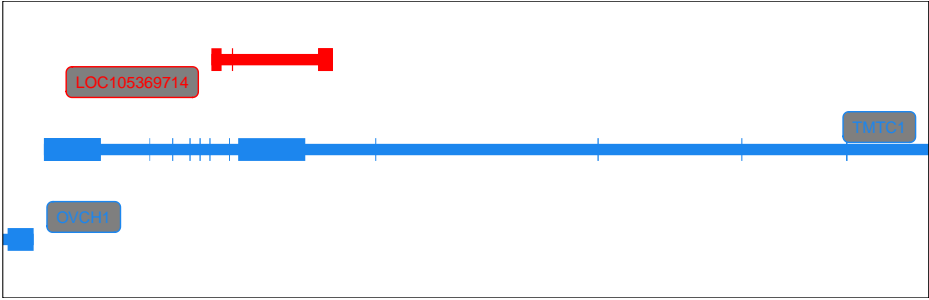
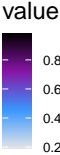
L5.6.NR4A2



Peaks

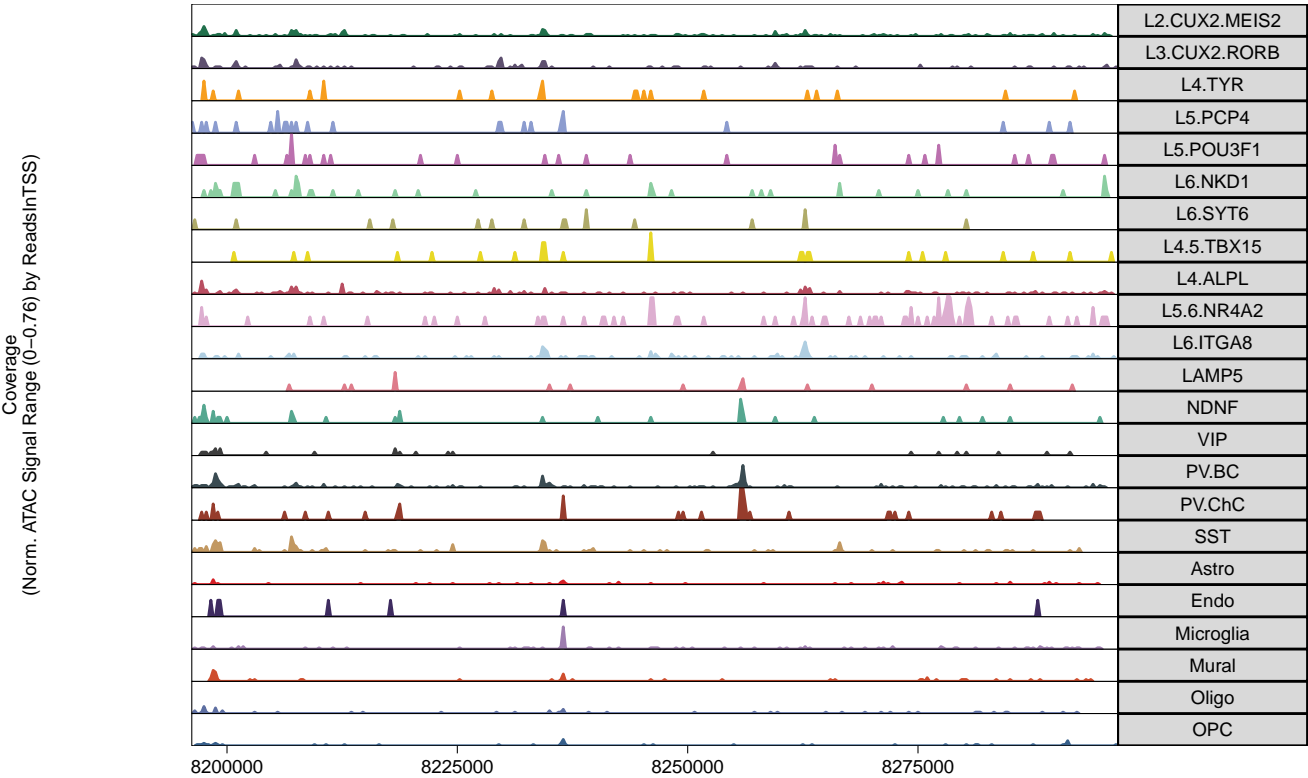


Peak2GeneLinks

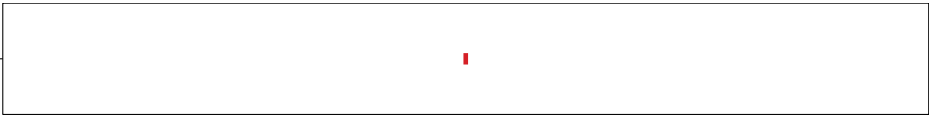


Genes

chr9:8196177–8296678 Marker Peak Tracks for: L5.6.NR4A2



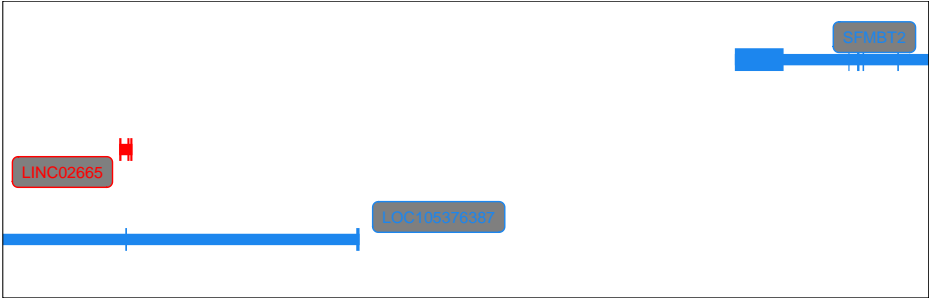
L5.6.NR4A2



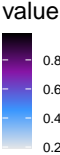
Peaks



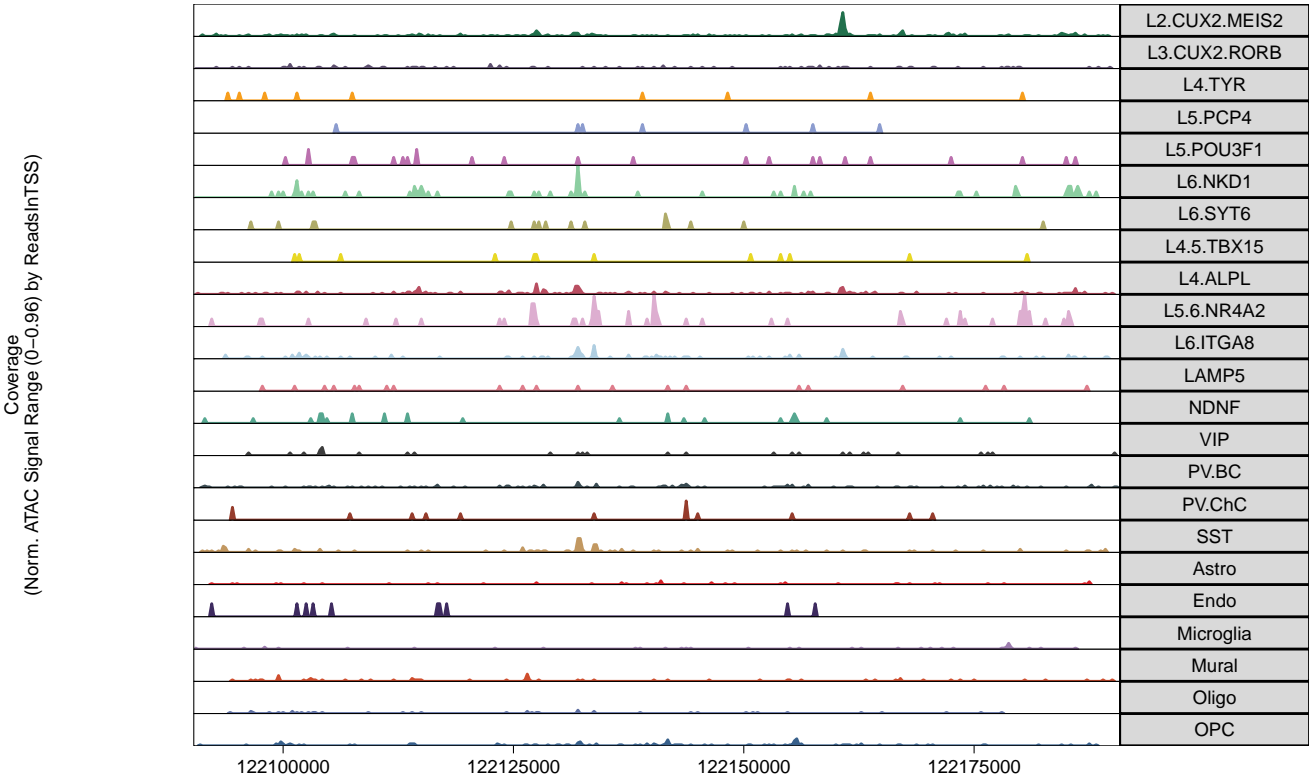
Peak2GeneLinks



Genes



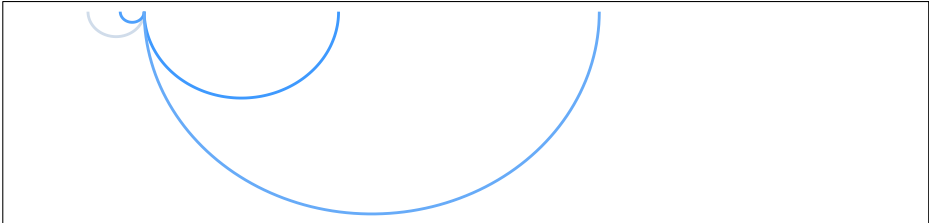
chr12:122090303–122190804 Marker Peak Tracks for: L5.6.NR4A2



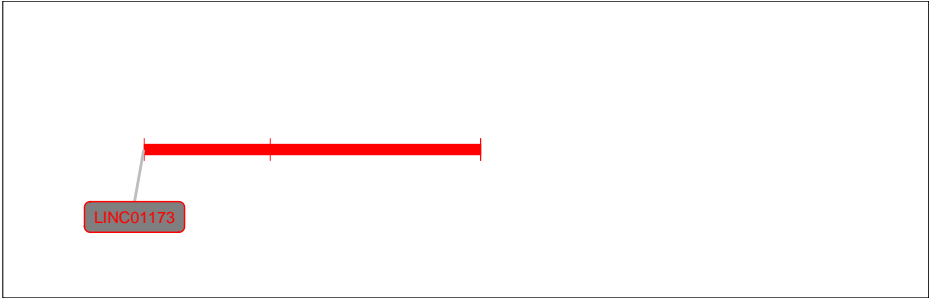
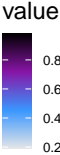
L5.6.NR4A2



Peaks



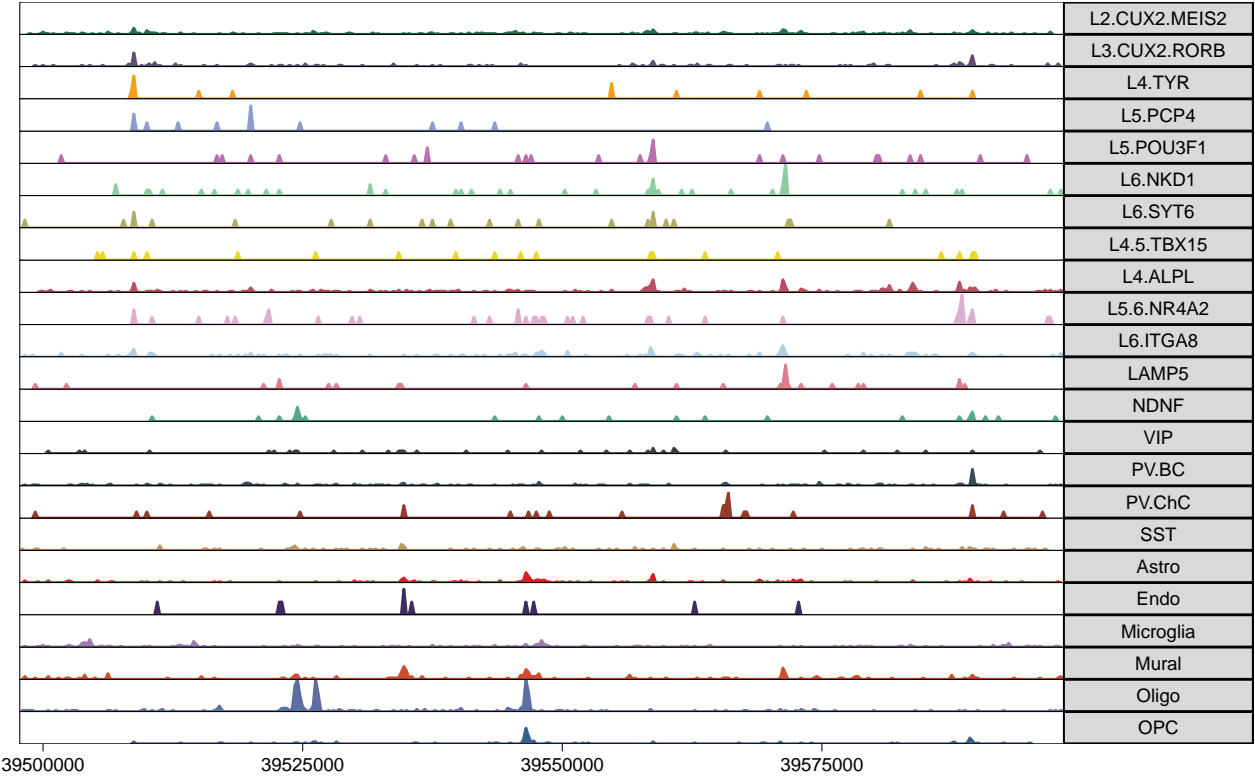
Peak2GeneLinks



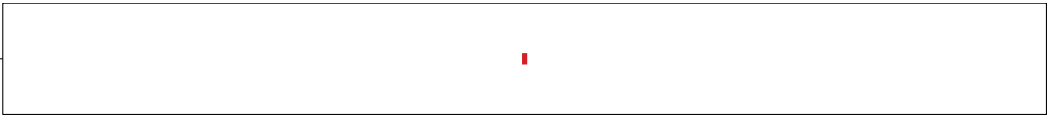
Genes

chr1:39497761–39598262 Marker Peak Tracks for: L5.6.NR4A2

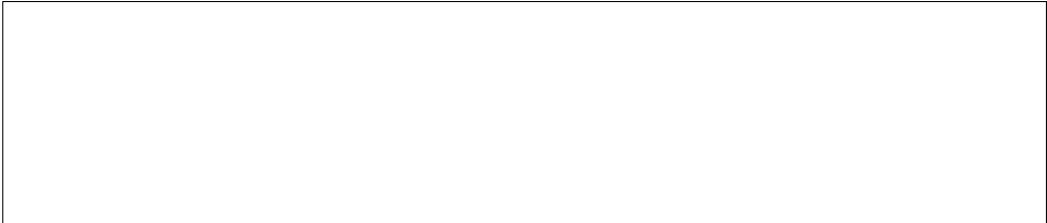
Coverage
(Norm. ATAC Signal Range 0–0.98) by ReadsInTSS)



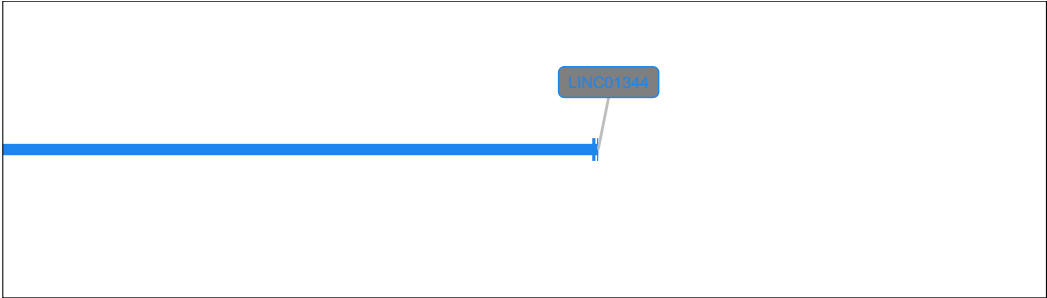
L5.6.NR4A2



Peaks



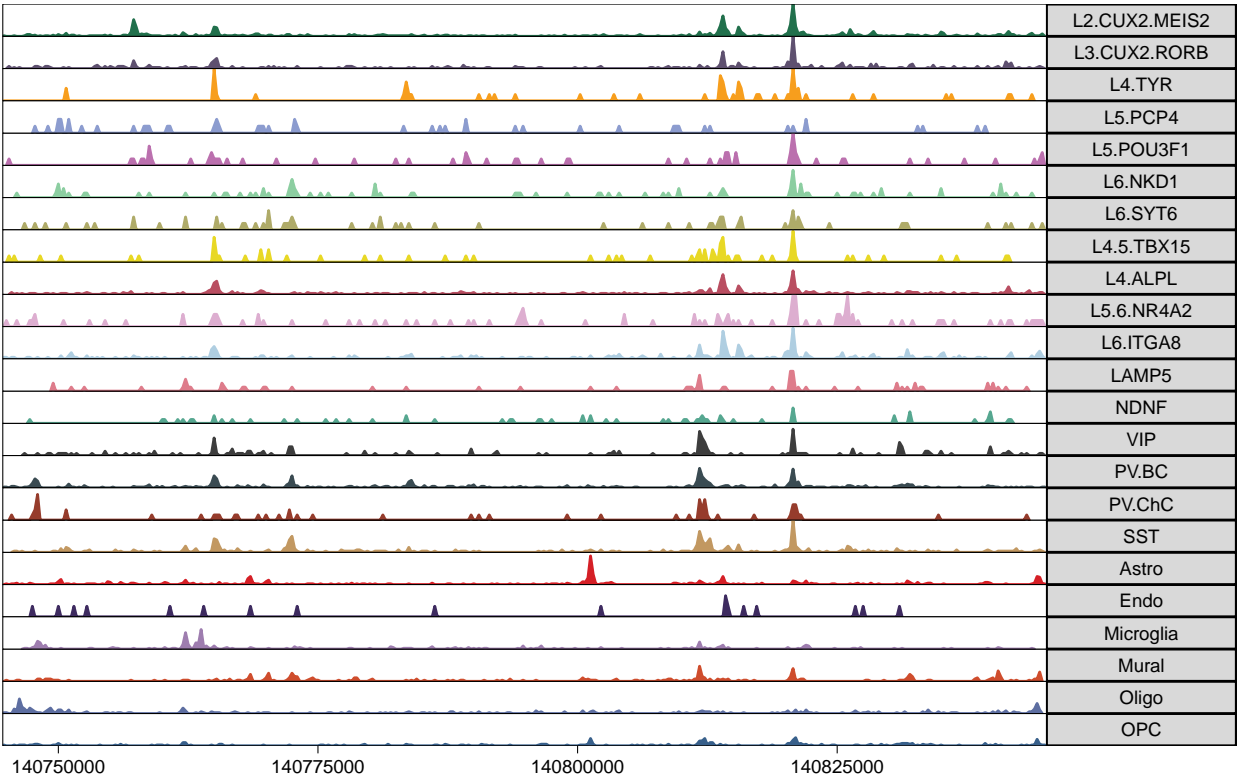
LoopTrack



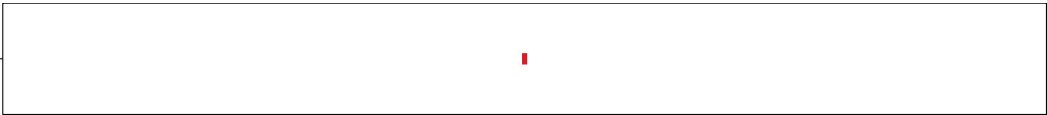
Genes

chr8:140744650–140845151 Marker Peak Tracks for: L5.6.NR4A2

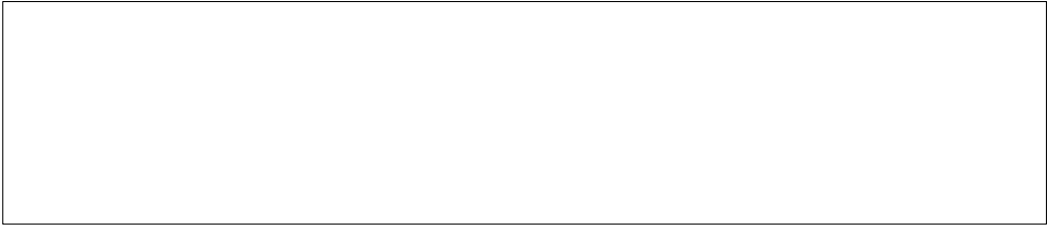
Coverage
(Norm. ATAC Signal Range (0–1.2) by ReadsInTSS)



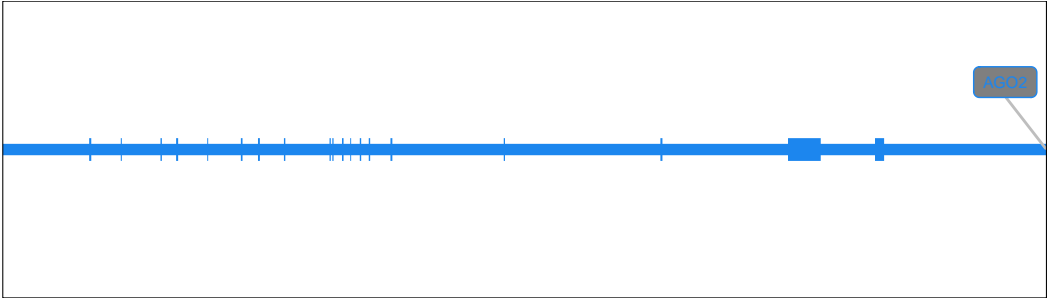
L5.6.NR4A2



Peaks

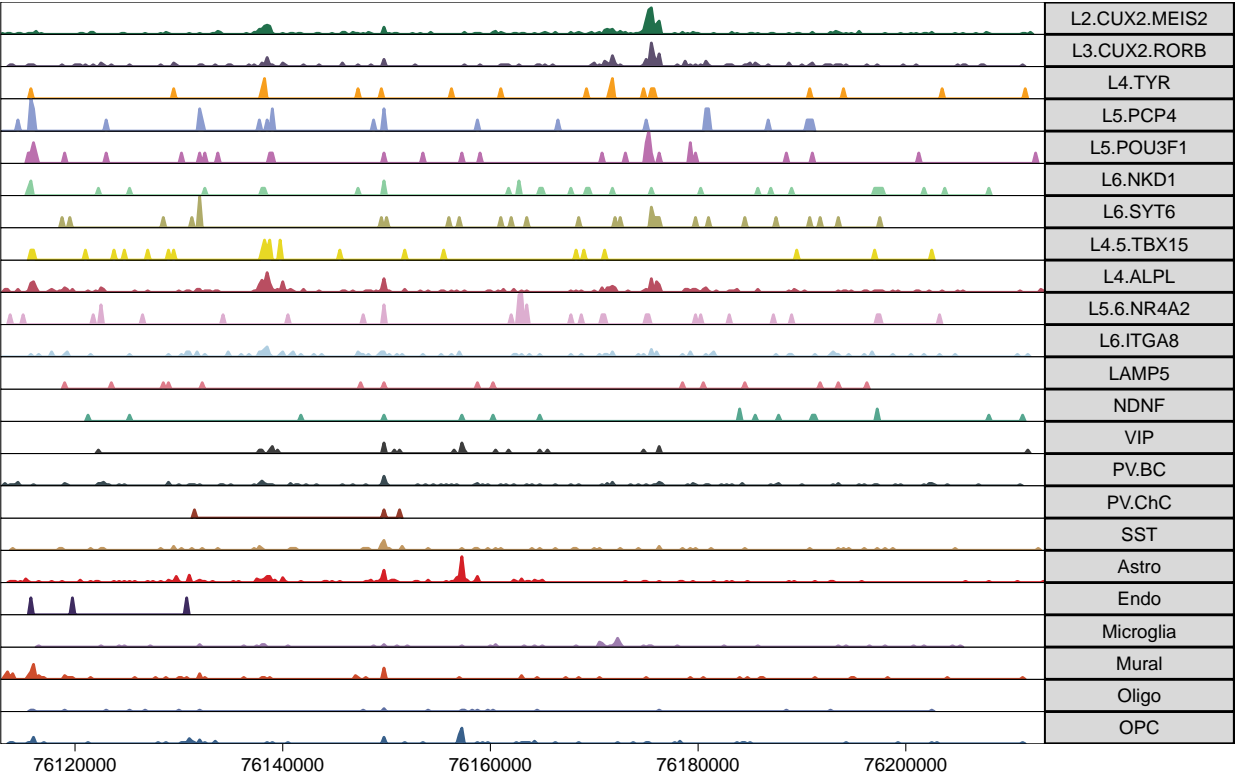


LoopTrack

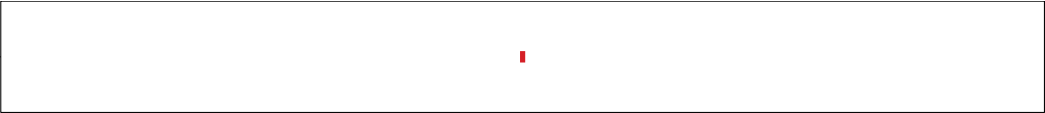


Genes

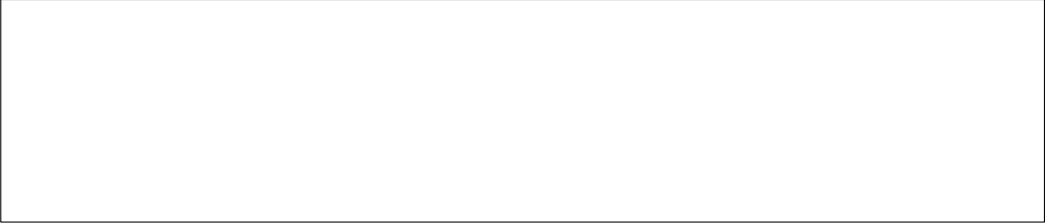
Coverage
(Norm. ATAC Signal Range 0–0.74) by ReadsInTSS)



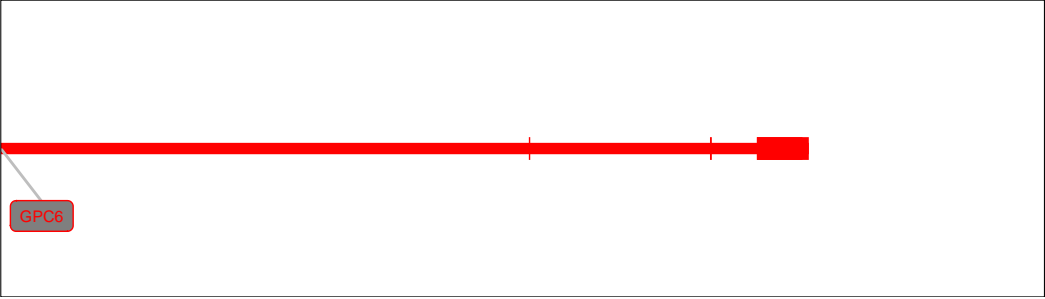
L5.6.NR4A2



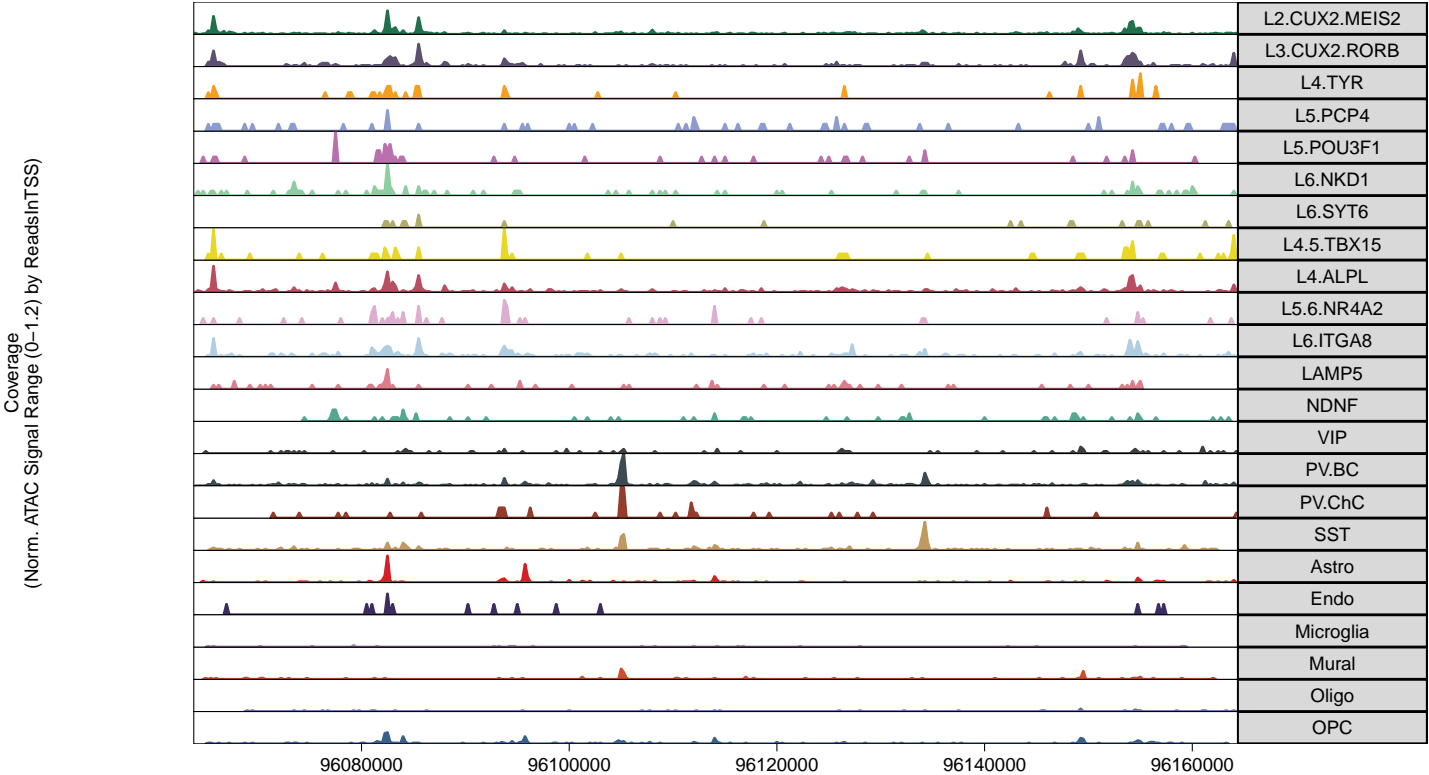
Peaks



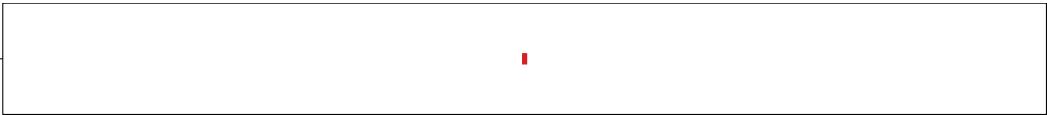
LoopTrack



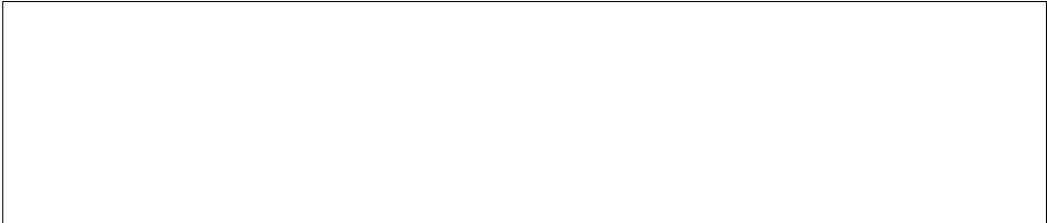
Genes



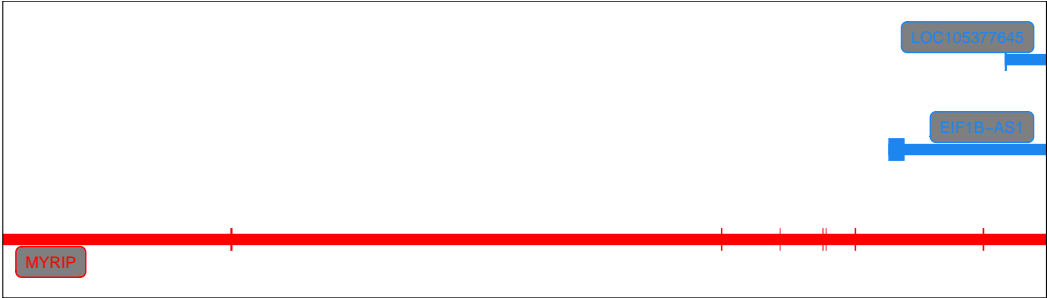
L5.6.NR4A2



Peaks



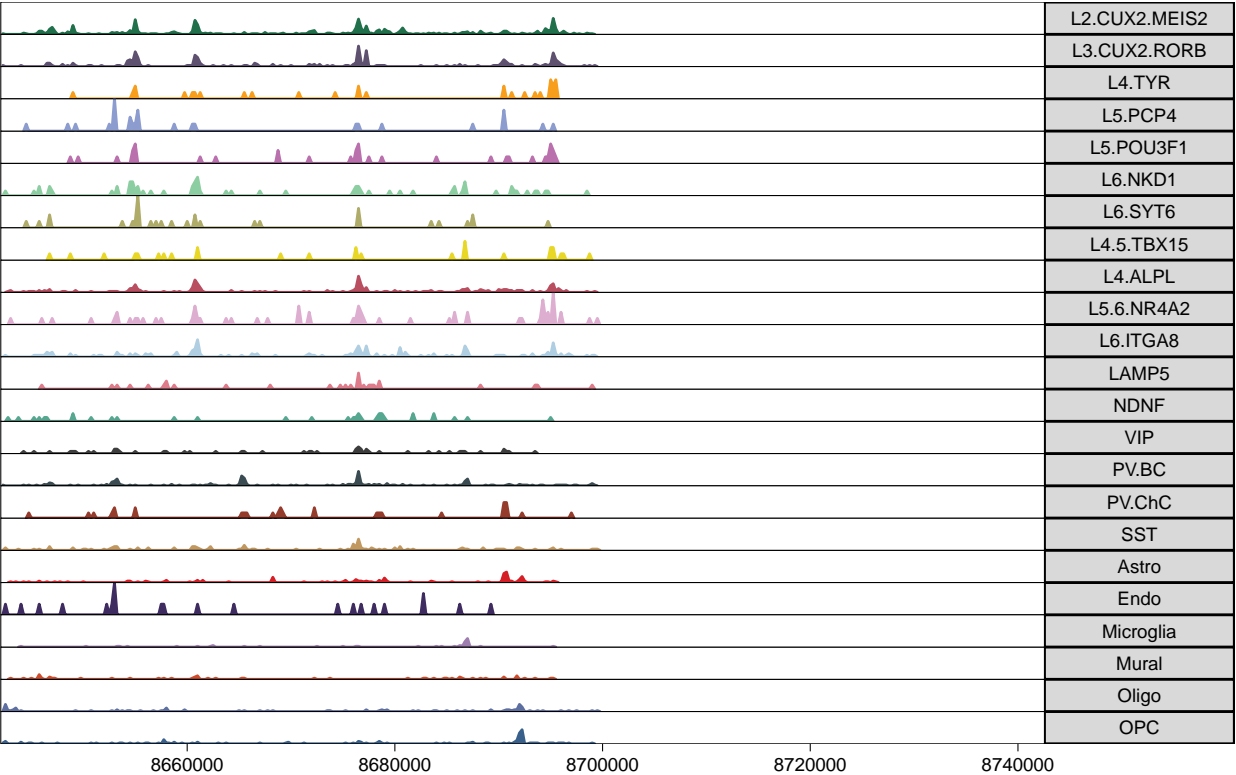
LoopTrack



Genes

chr17:8642051–8742552 Marker Peak Tracks for: L5.6.NR4A2

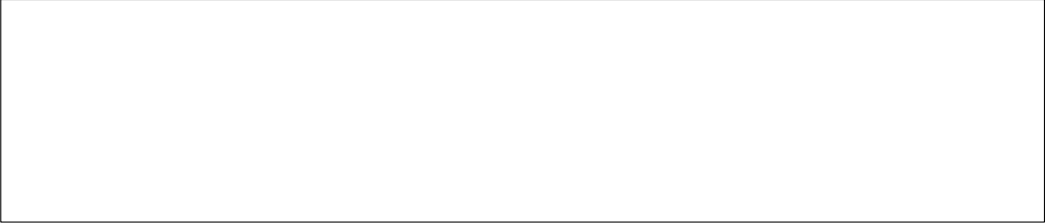
Coverage
(Norm. ATAC Signal Range (0–1.18) by ReadsInTSS)



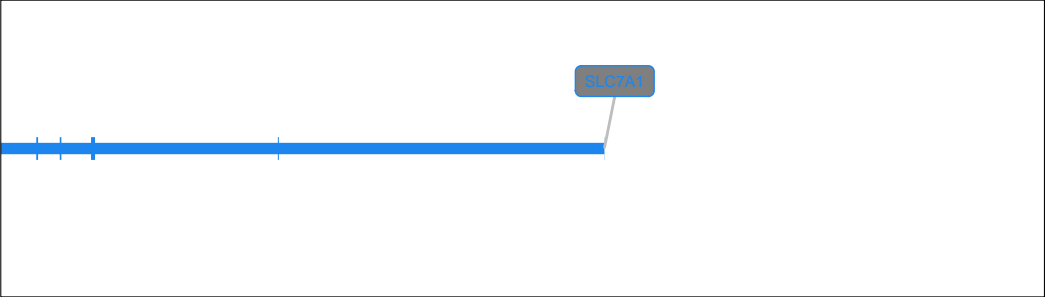
L5.6.NR4A2



Peaks

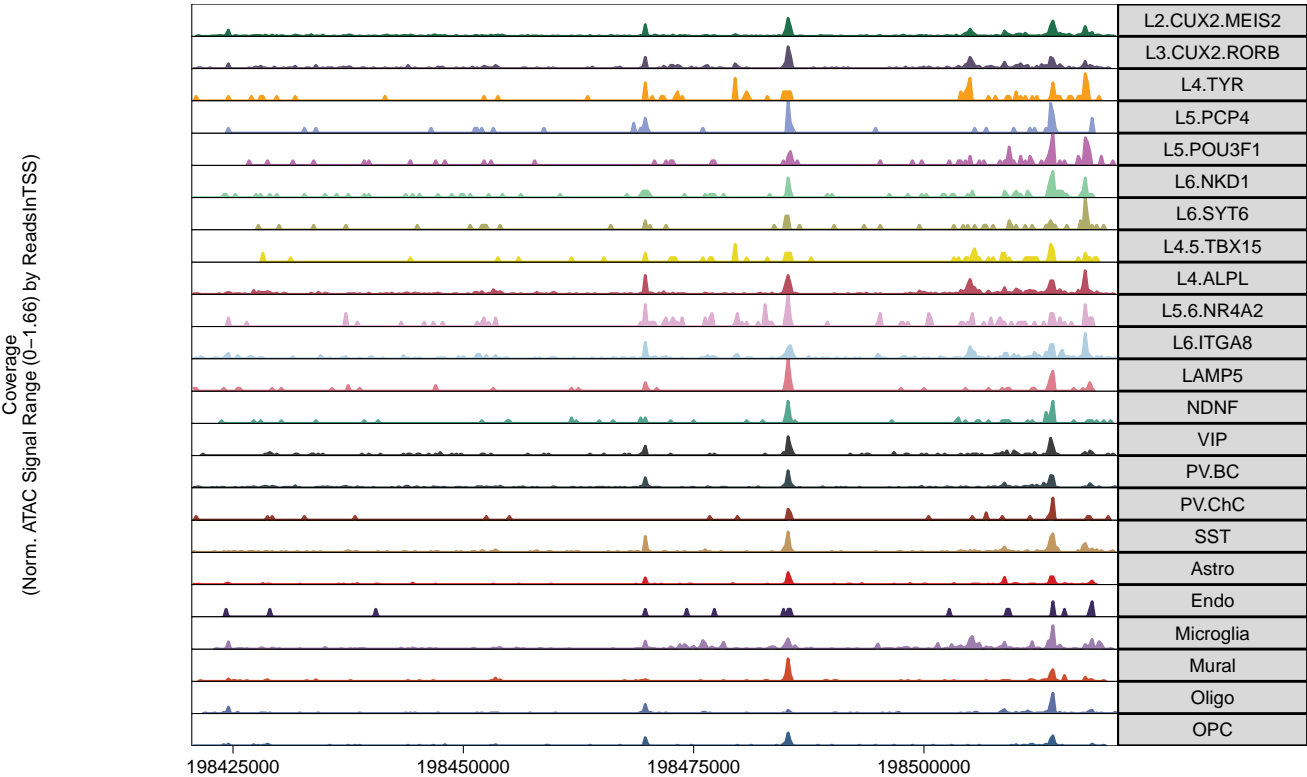


LoopTrack



Genes

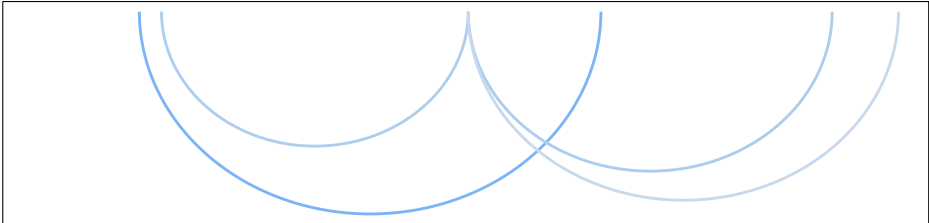
chr1:198420527-198521028 Marker Peak Tracks for: L5.6.NR4A2



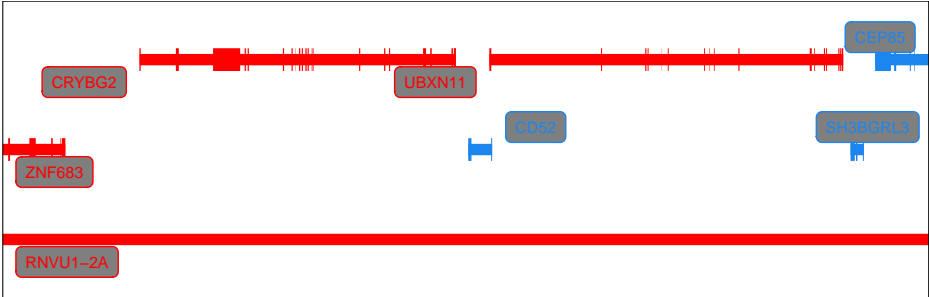
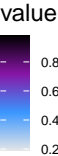
L5.6.NR4A2



Peaks

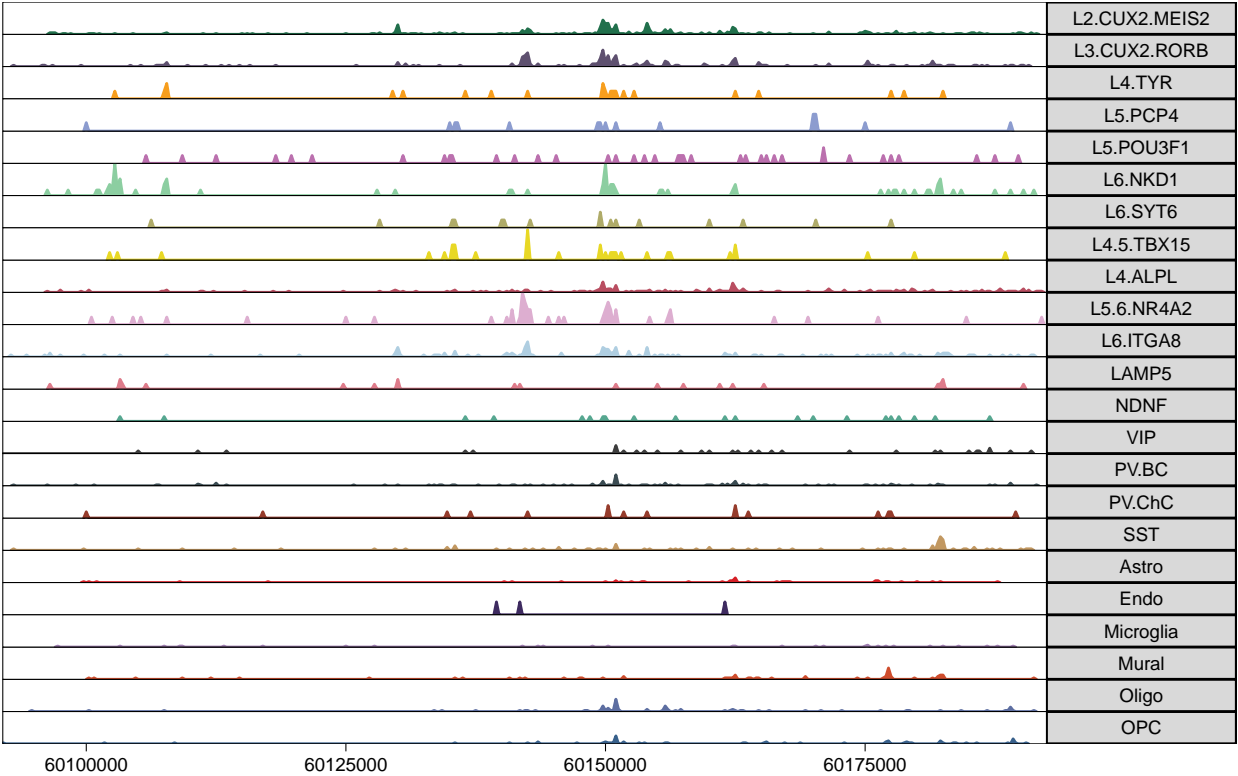


Peak2GeneLinks

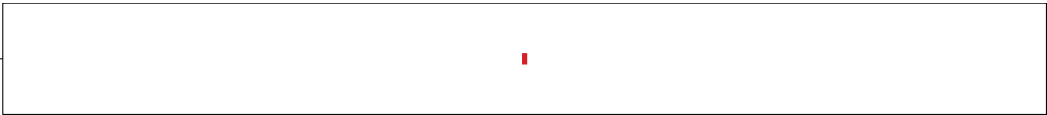


Genes

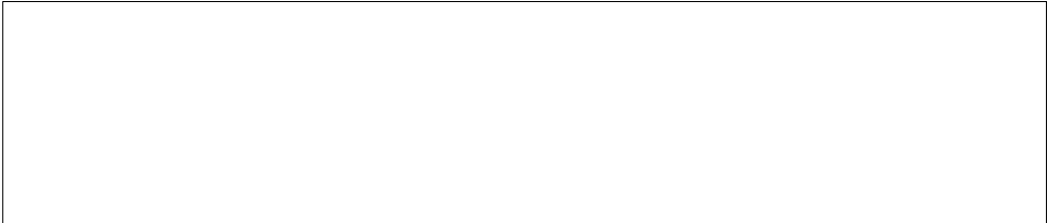
Coverage
(Norm. ATAC Signal Range (0–0.98) by ReadsInTSS)



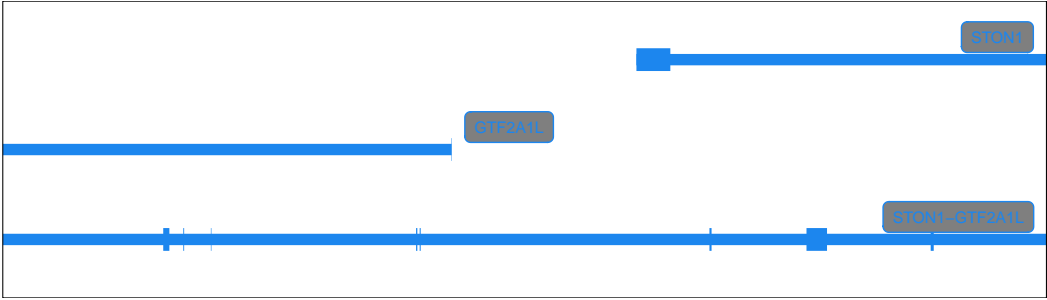
L5.6.NR4A2



Peaks



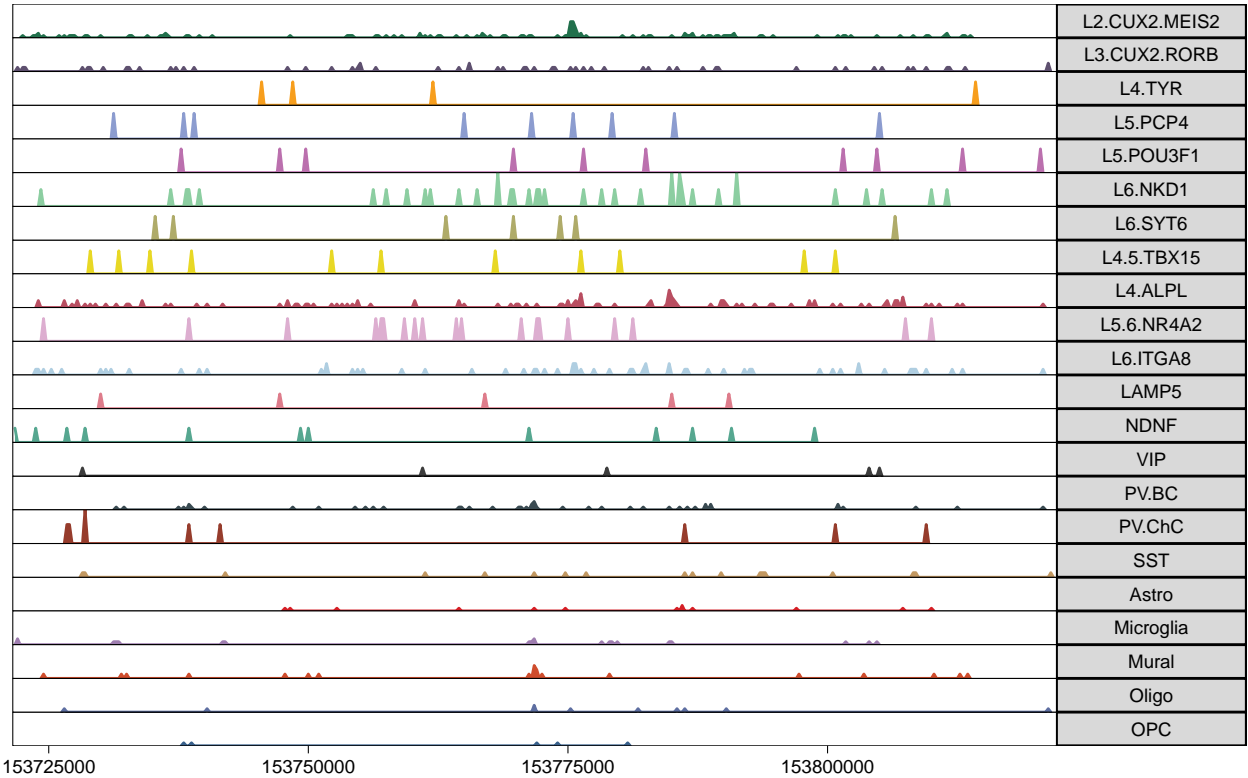
LoopTrack



Genes

chr1:153721540–153822041 Marker Peak Tracks for: L5.6.NR4A2

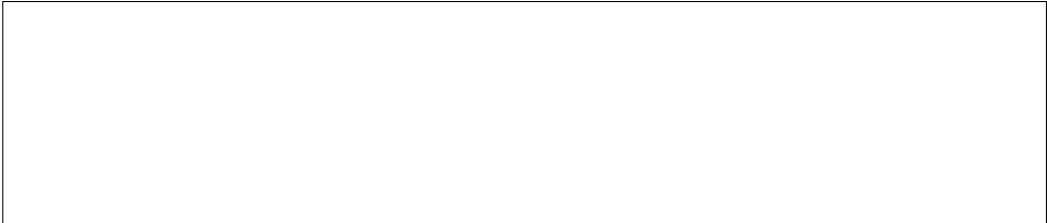
Coverage
(Norm. ATAC Signal Range (0–0.34) by ReadsInTSS)



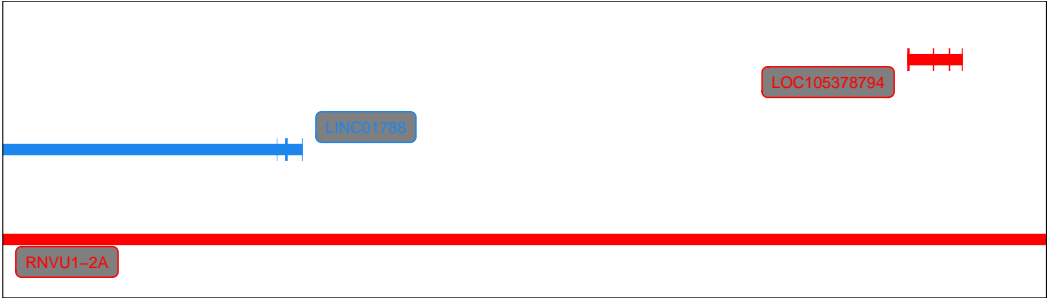
L5.6.NR4A2



Peaks



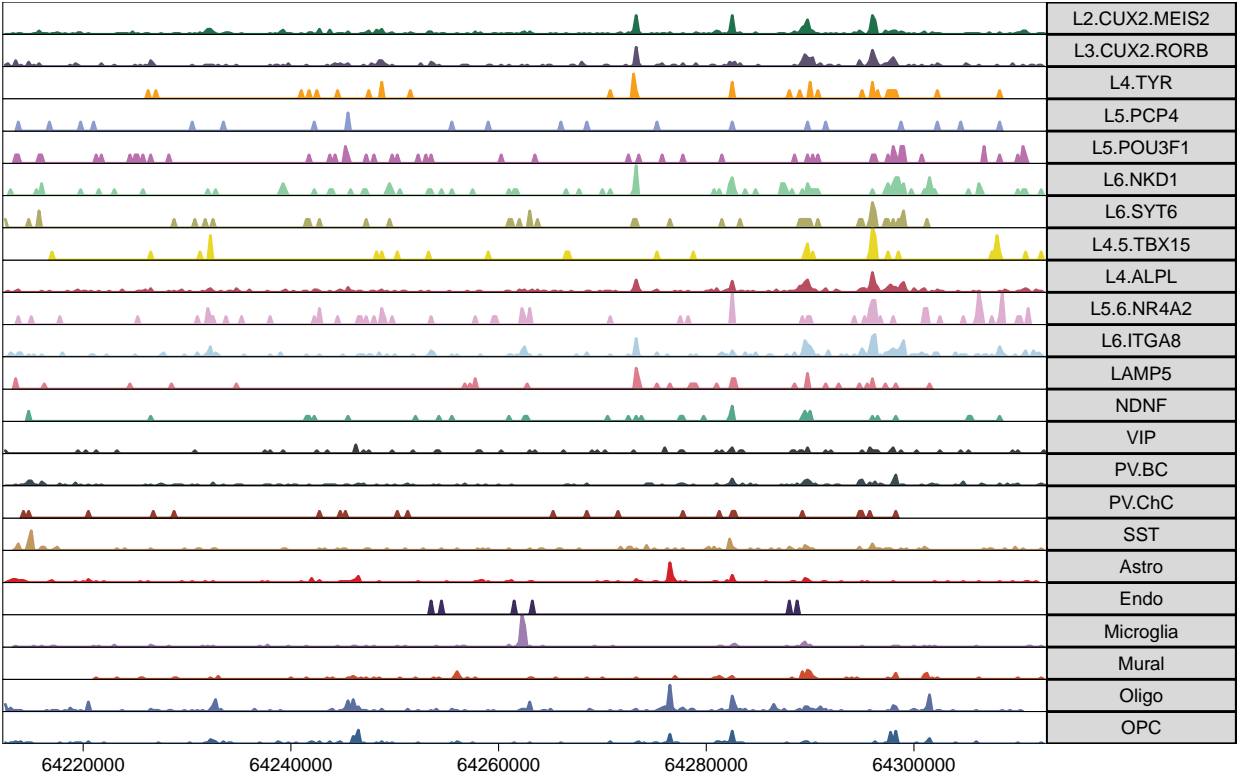
LoopTrack



Genes

chr4:64212257-64312758 Marker Peak Tracks for: L5.6.NR4A2

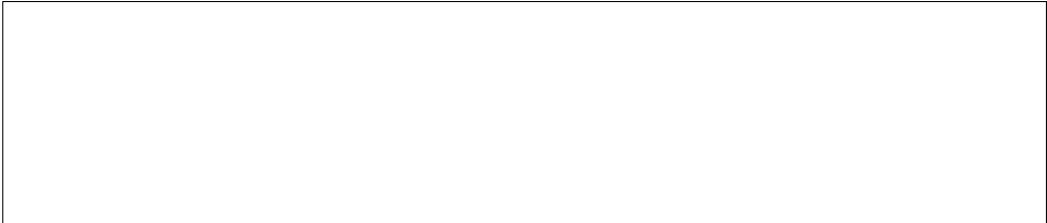
Coverage
(Norm. ATAC Signal Range (0-0.91) by ReadsInTSS)



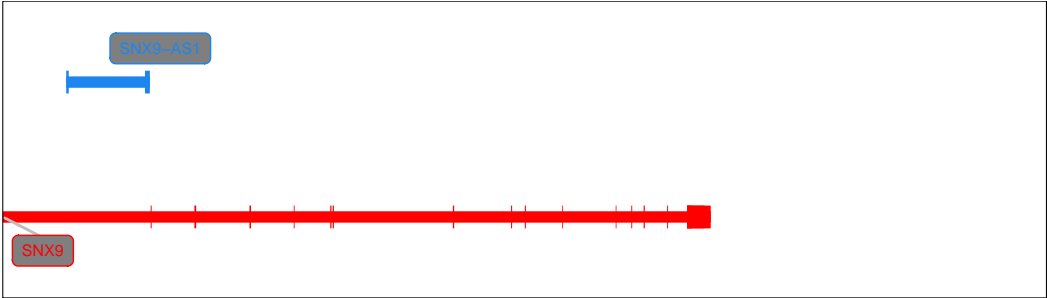
L5.6.NR4A2



Peaks

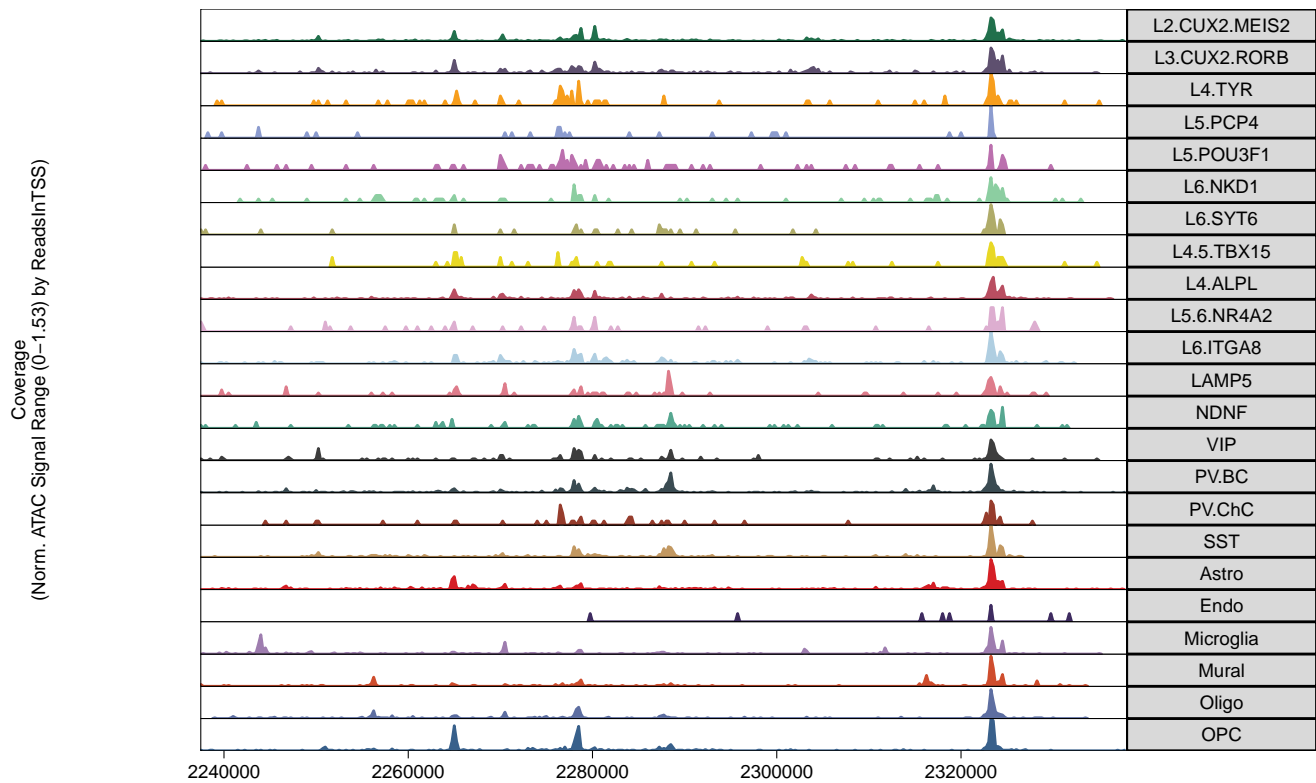


LoopTrack



Genes

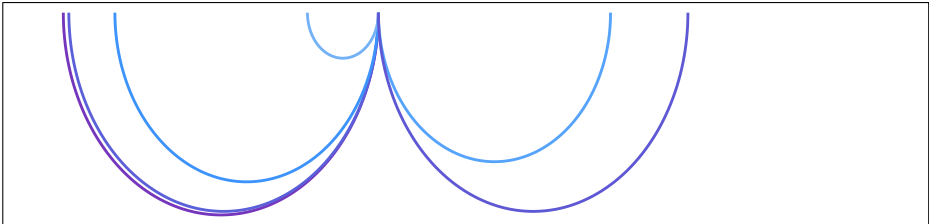
chr2:2237483–2337984 Marker Peak Tracks for: L5.6.NR4A2



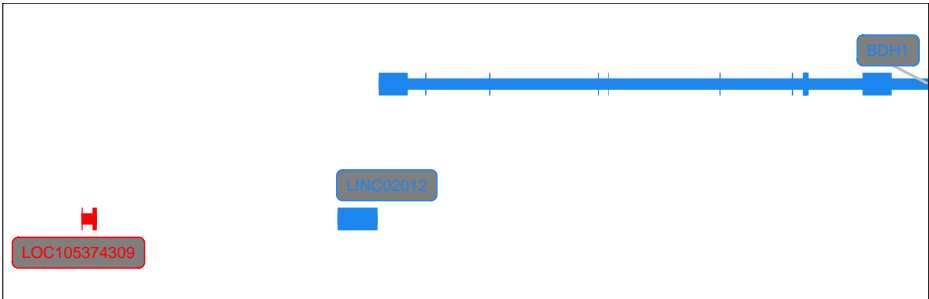
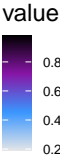
L5.6.NR4A2



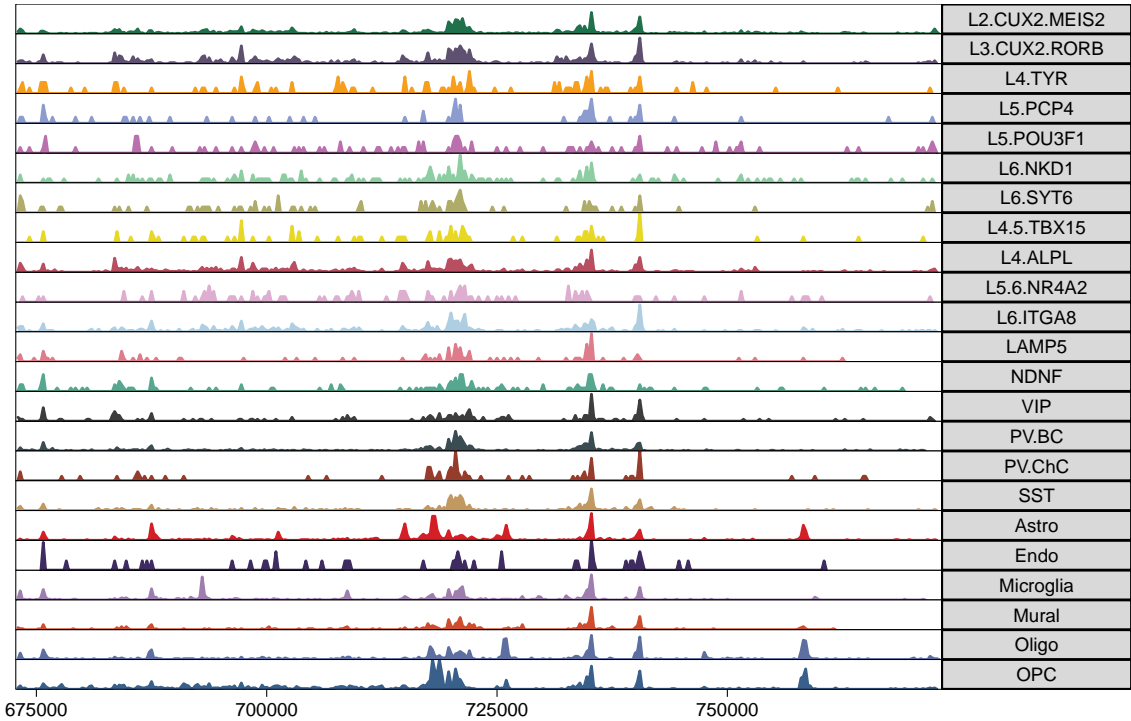
Peaks



Peak2GeneLinks



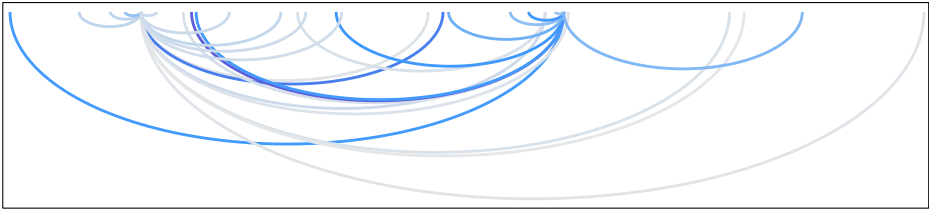
Coverage
(Norm. ATAC Signal Range (0–1.26) by Reads in TSS)



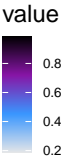
L5.6.NR4A2



Peaks



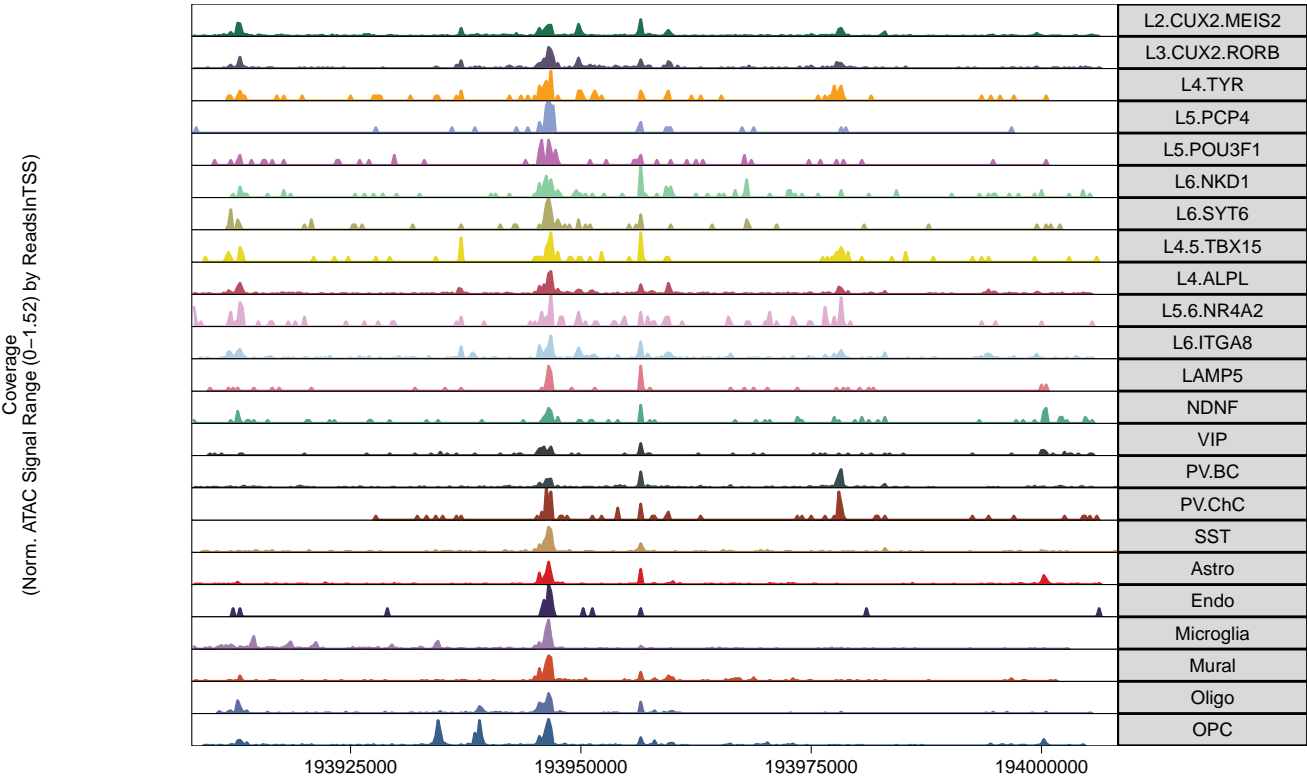
Peak2GeneLinks



GeneTrack

strand ● ·

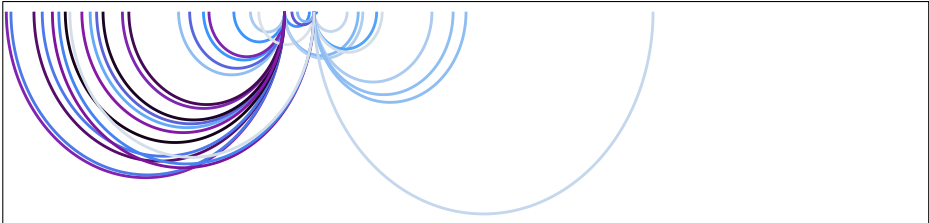
chr1:193907769-194008270 Marker Peak Tracks for: L5.6.NR4A2



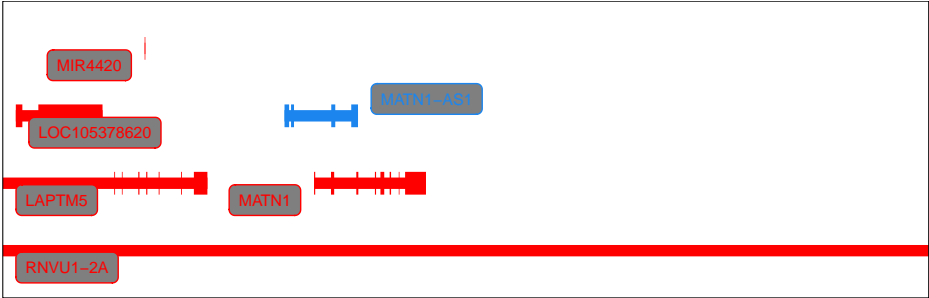
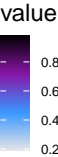
L5.6.NR4A2



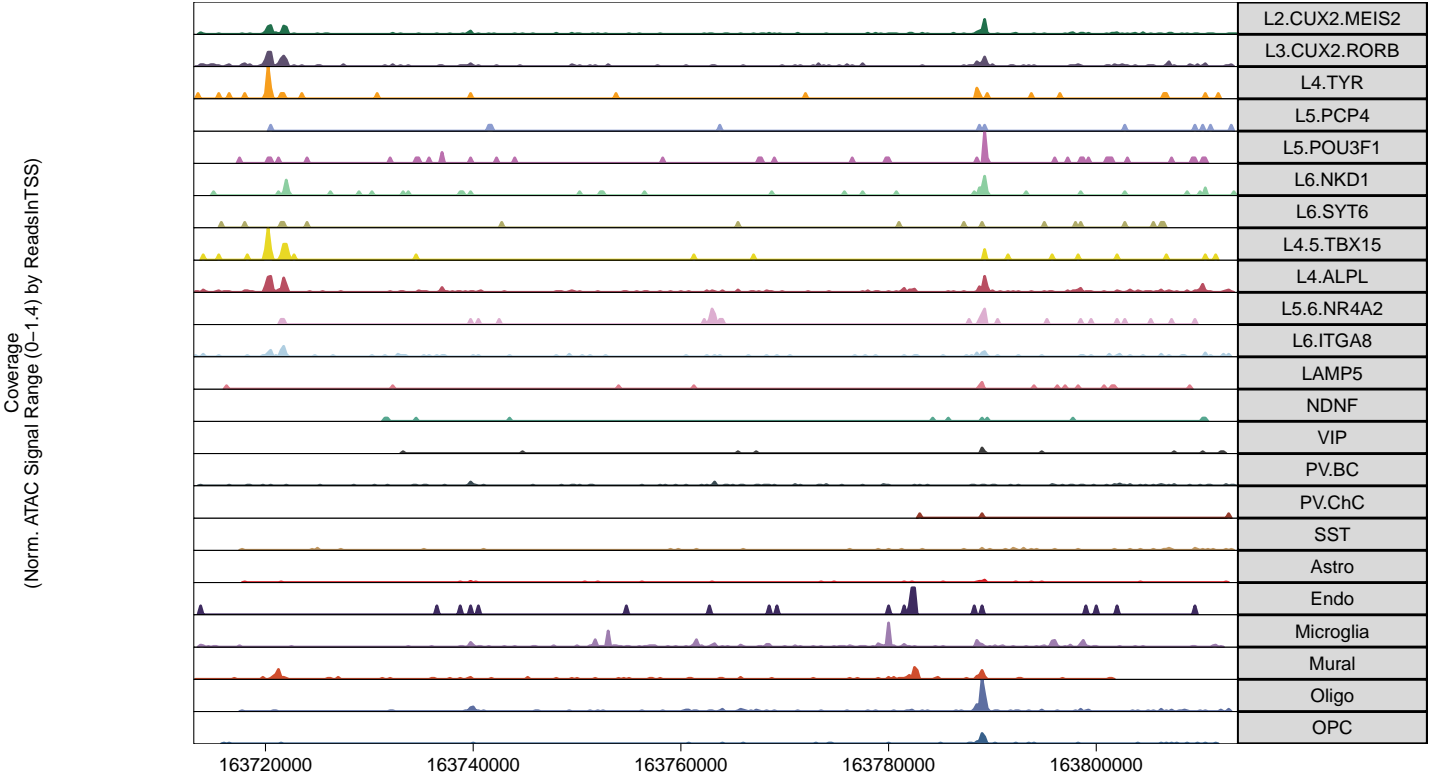
Peaks



Peak2GeneLinks



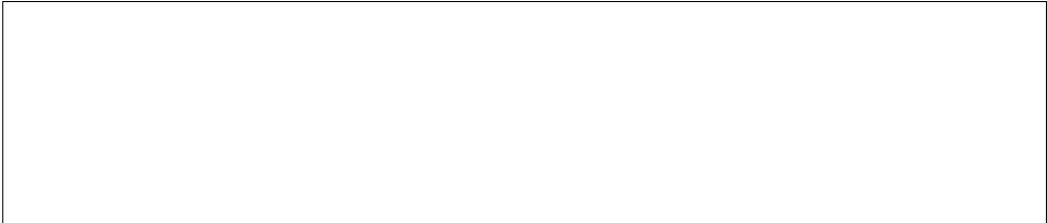
Genes



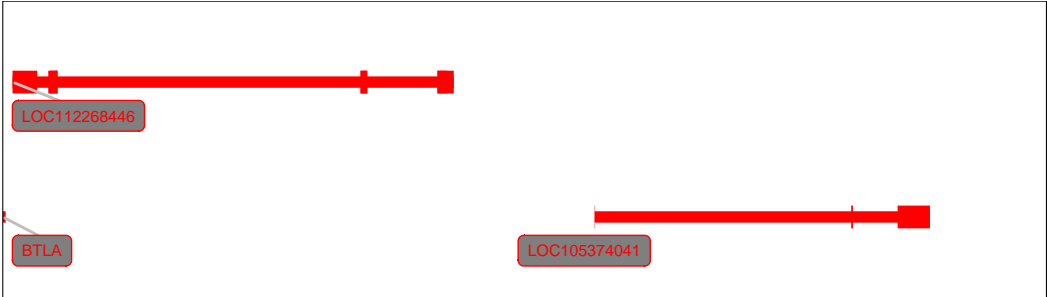
L5.6.NR4A2



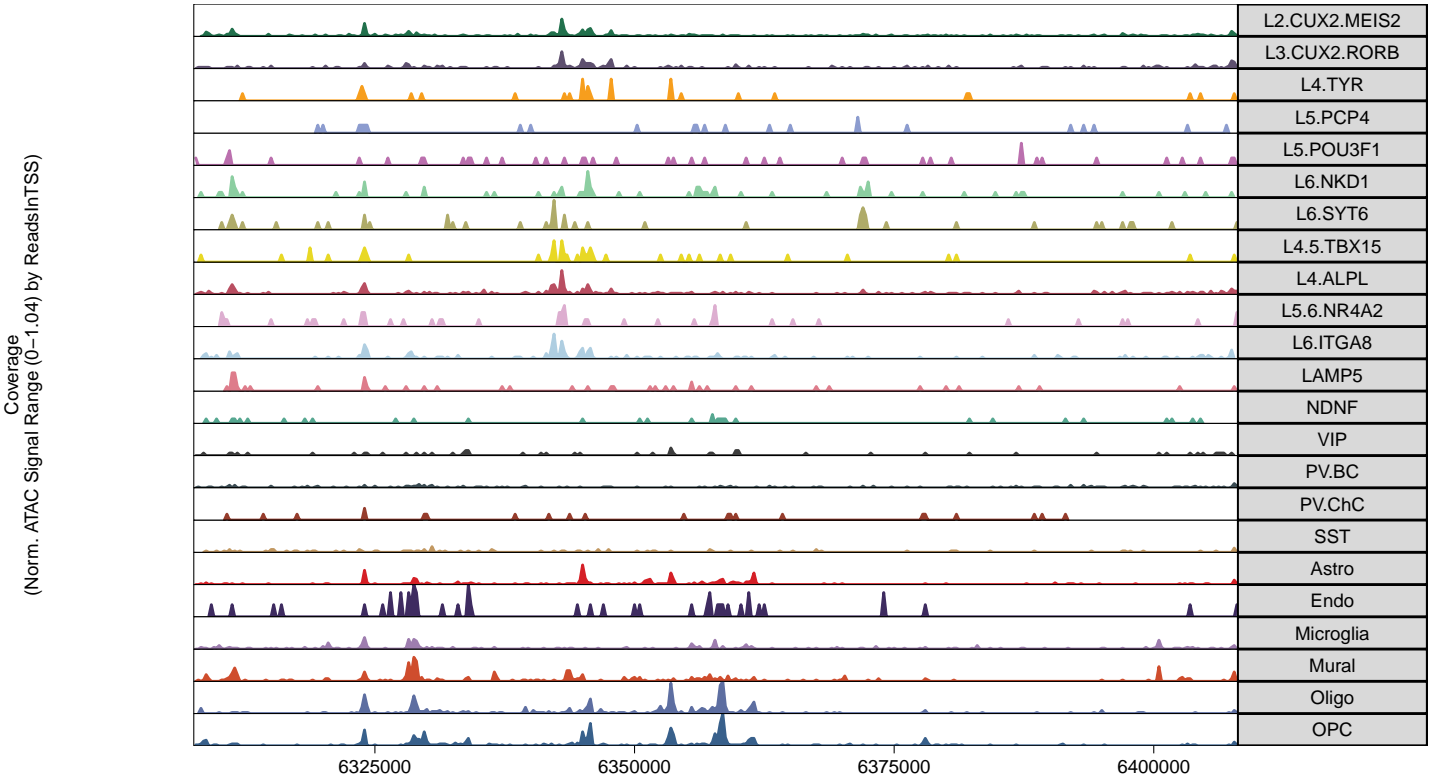
Peaks



LoopTrack



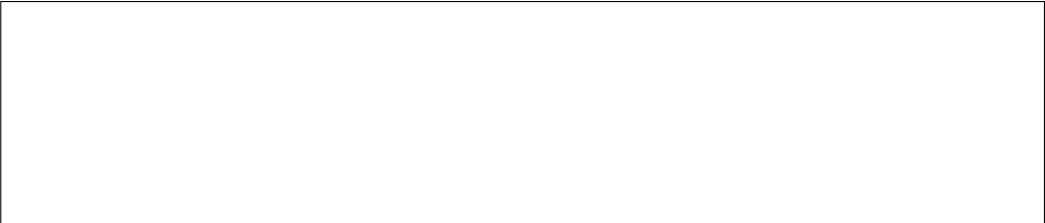
Genes



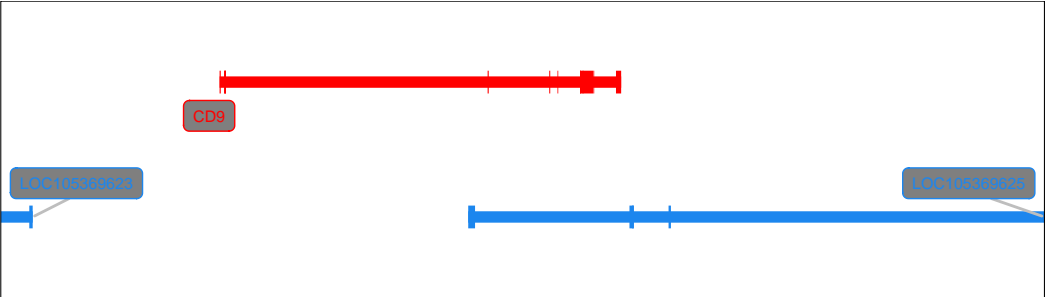
L5.6.NR4A2



Peaks

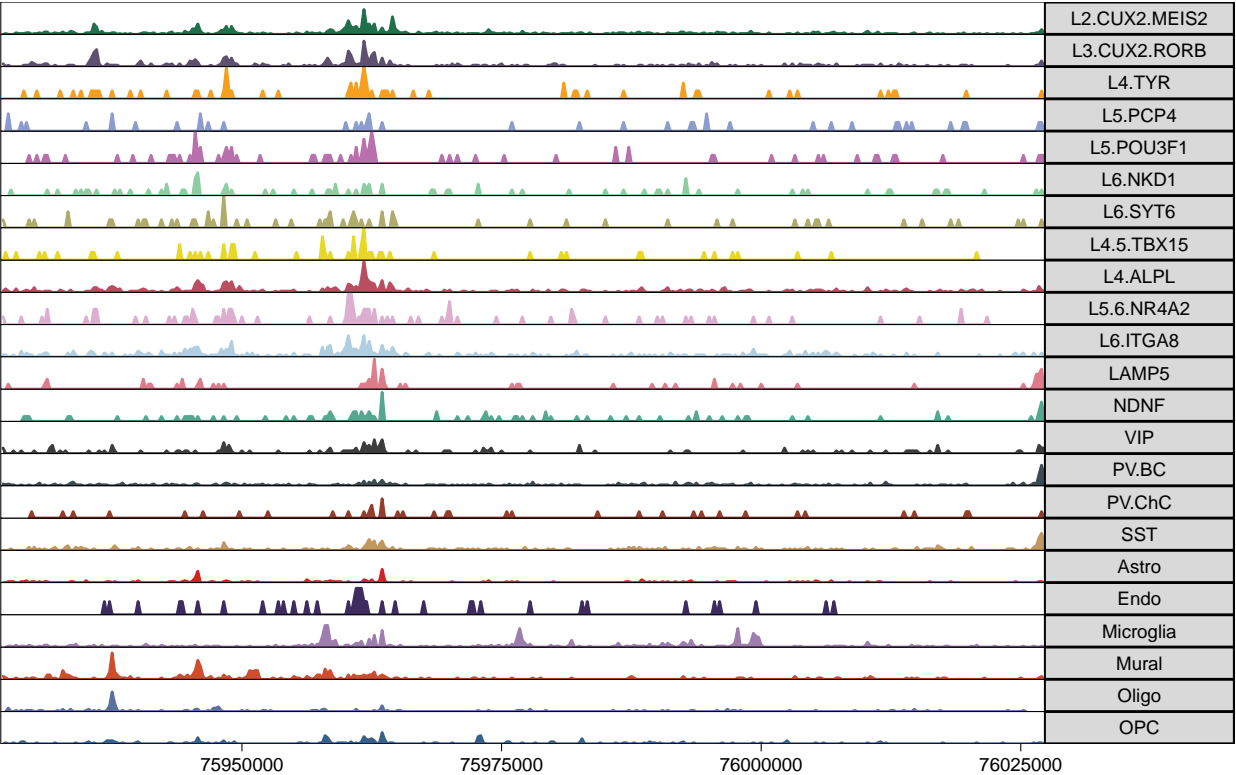


LoopTrack

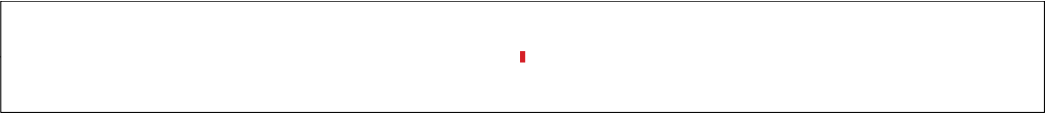


Genes

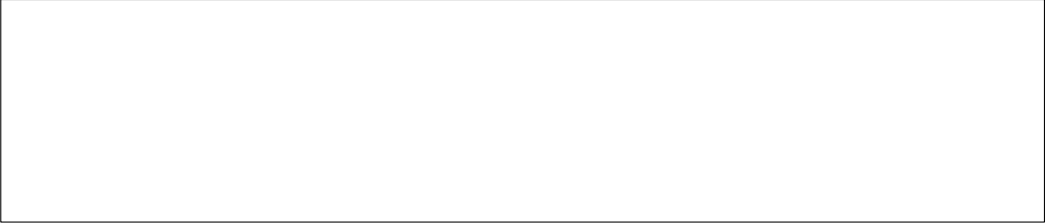
Coverage
(Norm. ATAC Signal Range (0-0.95) by ReadsInTSS)



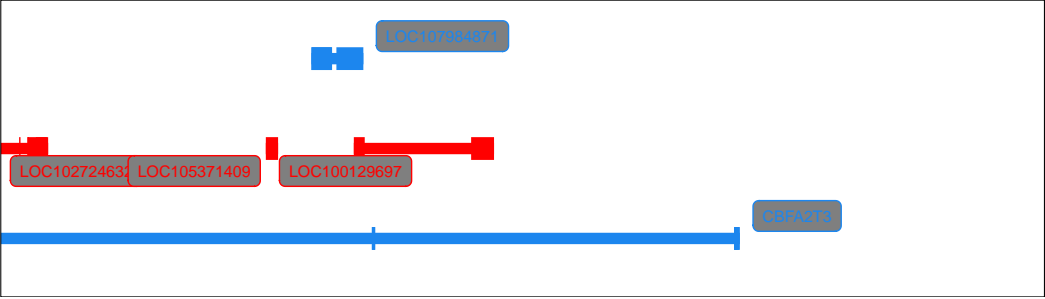
L5.6.NR4A2



Peaks

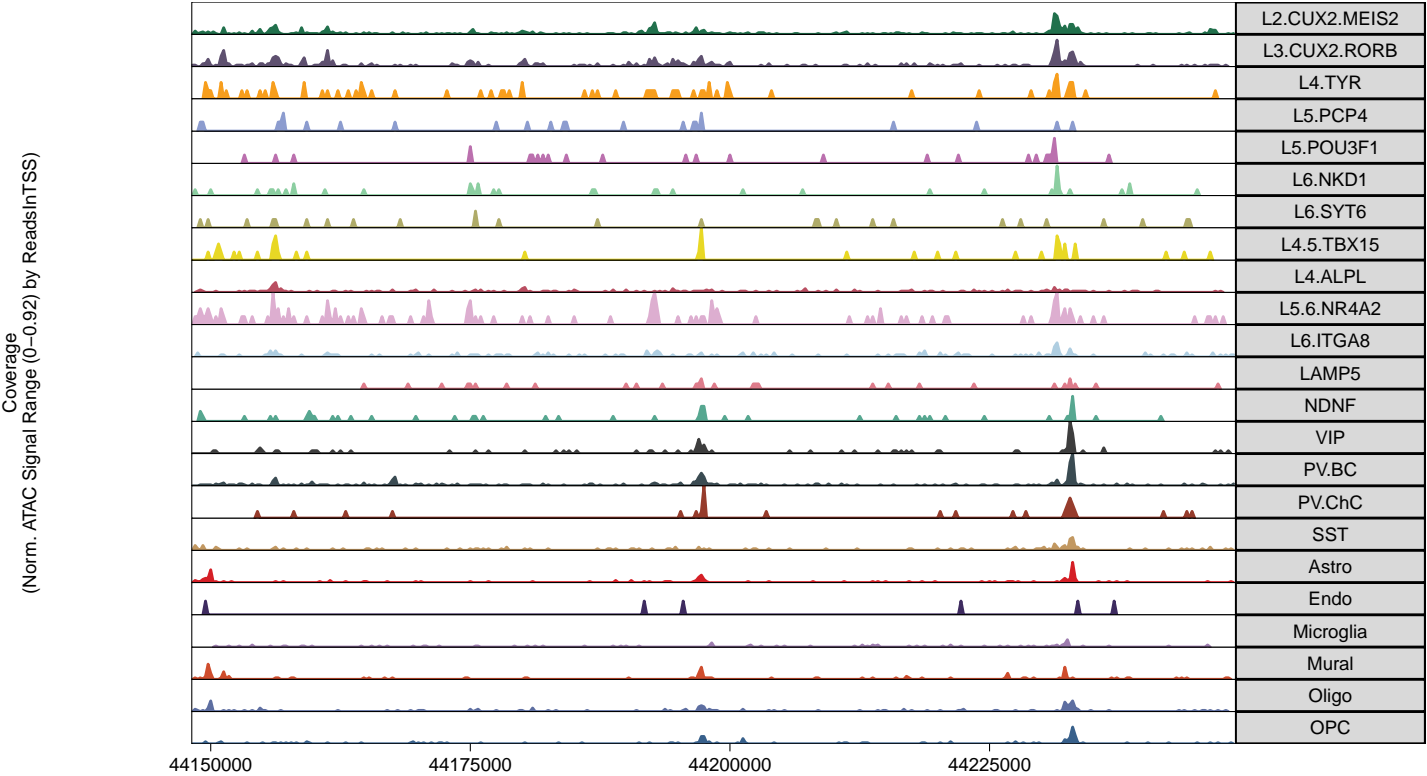


LoopTrack

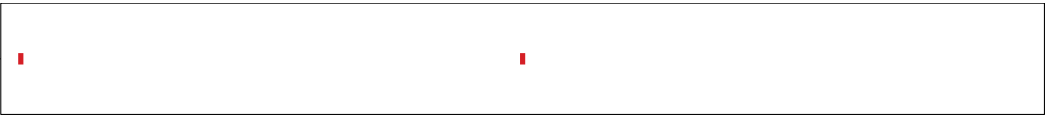


Genes

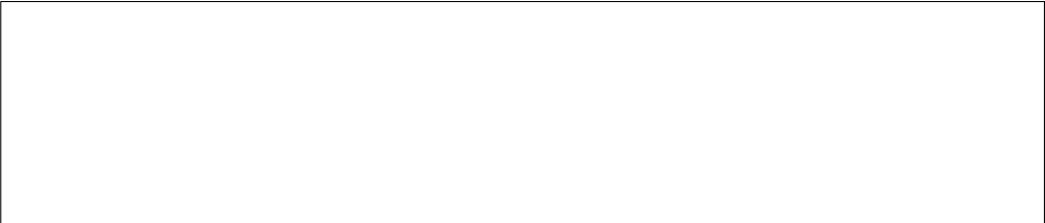
chr3:44148178–44248679 Marker Peak Tracks for: L5.6.NR4A2



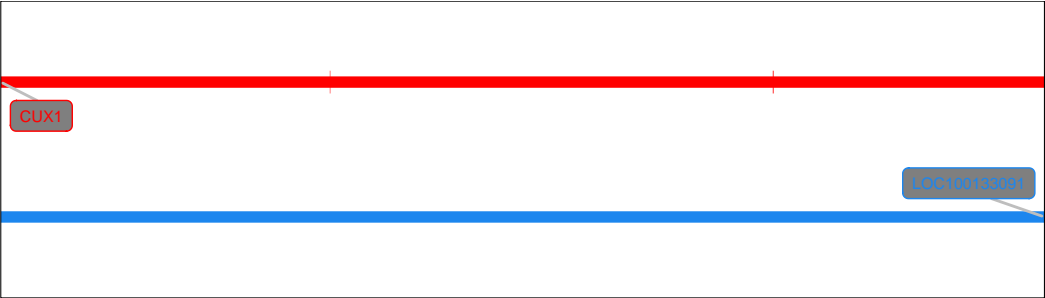
L5.6.NR4A2



Peaks



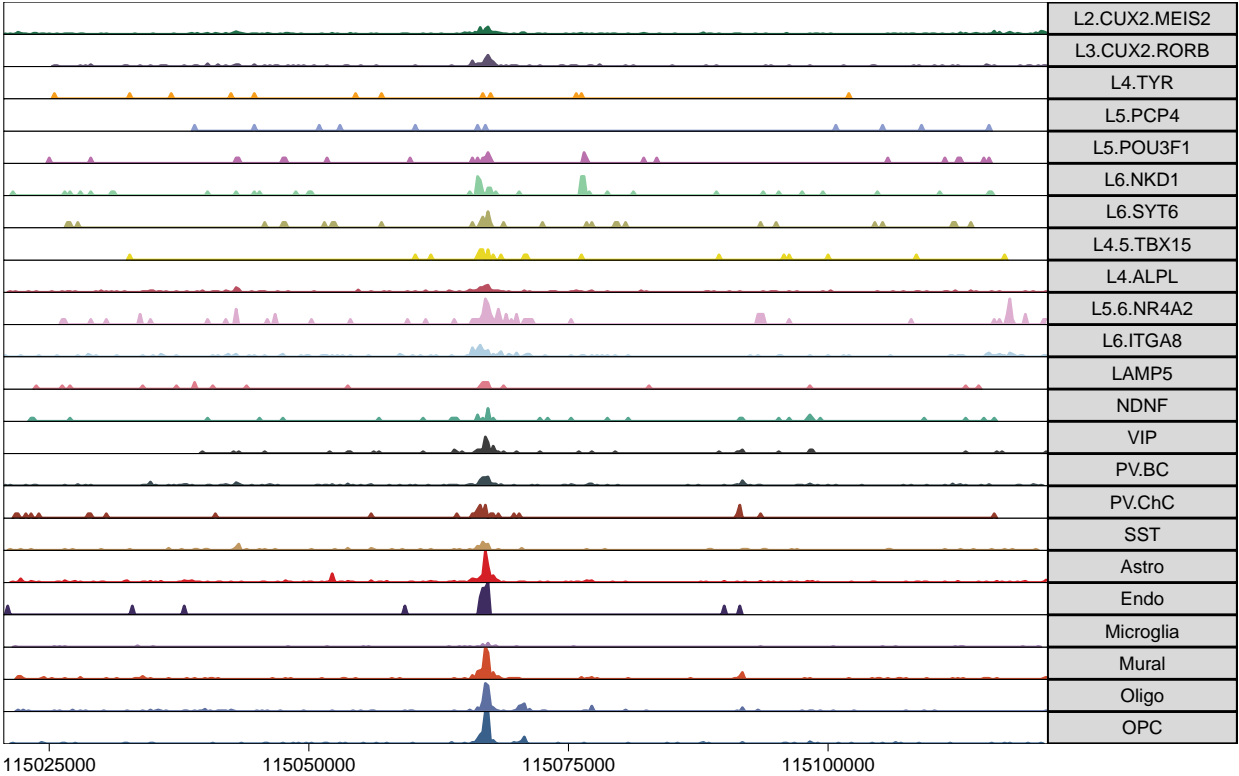
LoopTrack



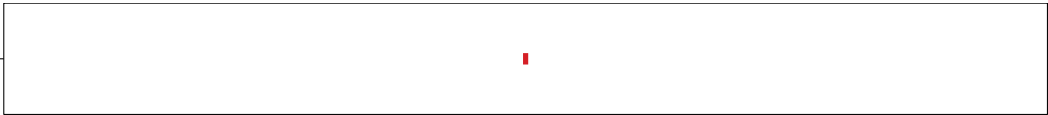
Genes

chr7:115020623–115121124 Marker Peak Tracks for: L5.6.NR4A2

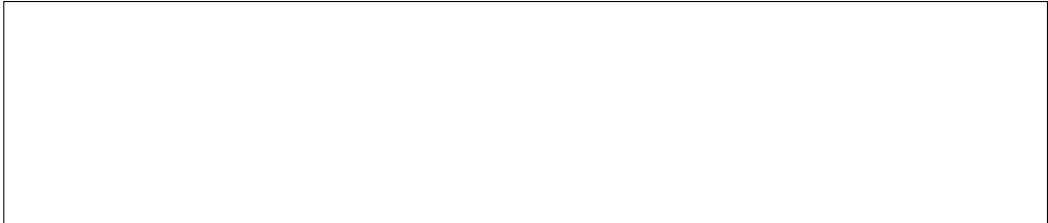
Coverage
(Norm. ATAC Signal Range 0–1.42) by ReadsInTSS)



L5.6.NR4A2



Peaks

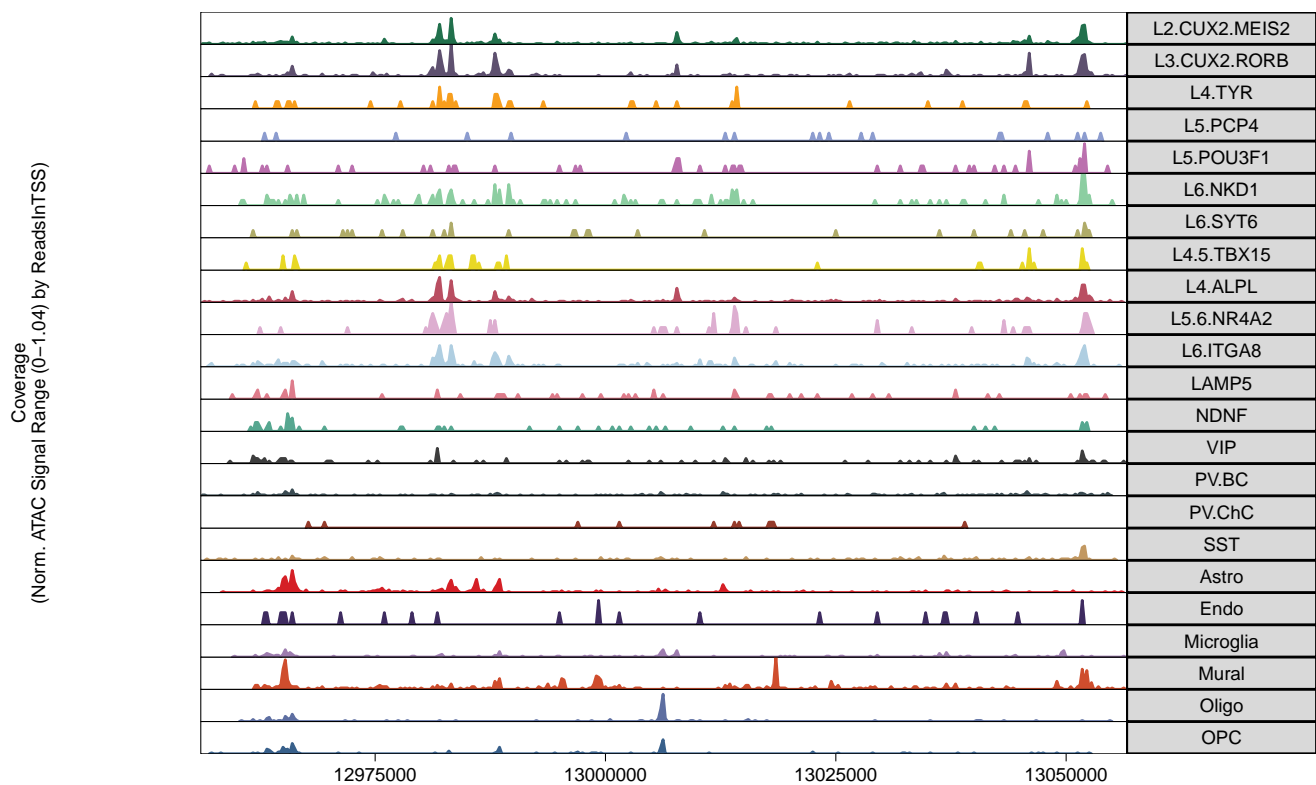


LoopTrack



Genes

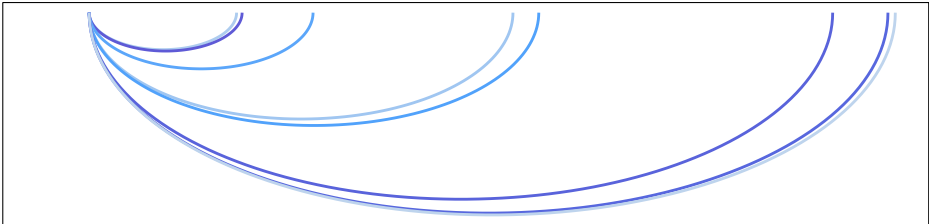
chr1:12956074–13056575 Marker Peak Tracks for: L5.6.NR4A2



L5.6.NR4A2

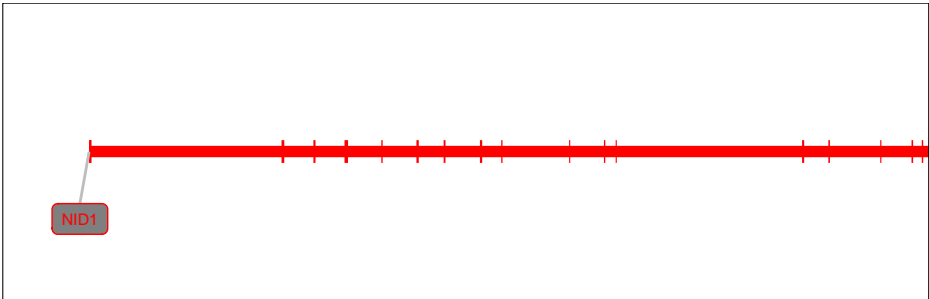
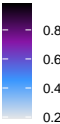


Peaks



Peak2GeneLinks

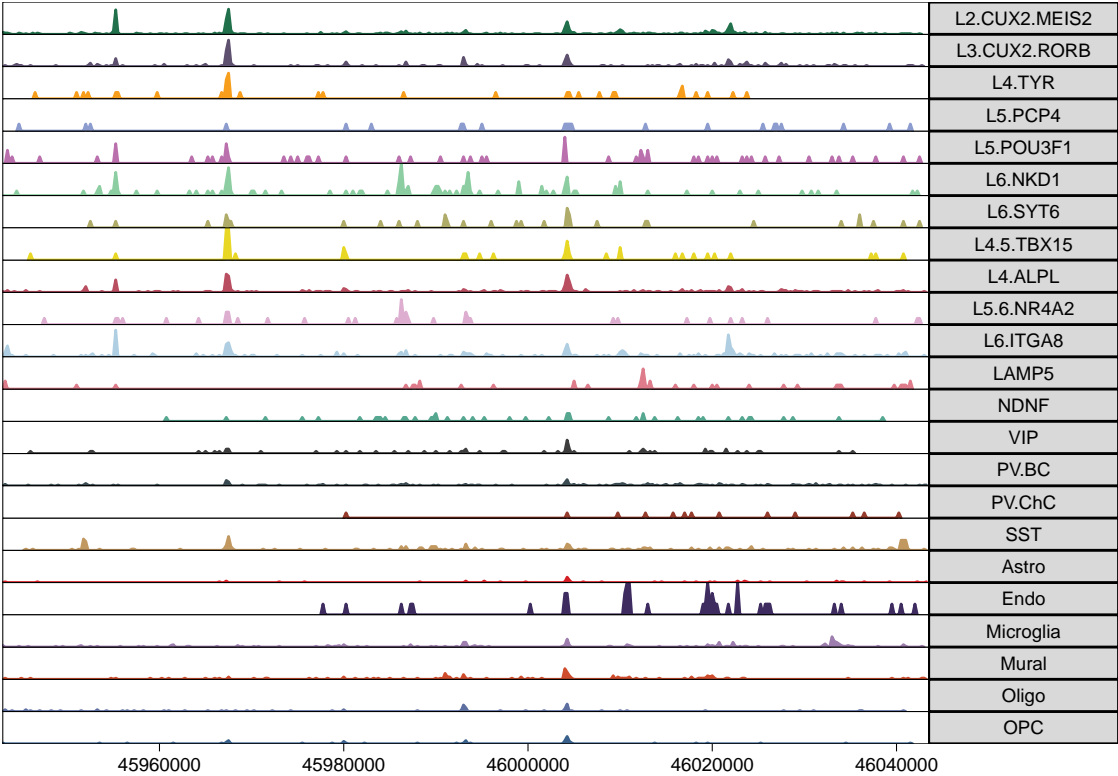
value



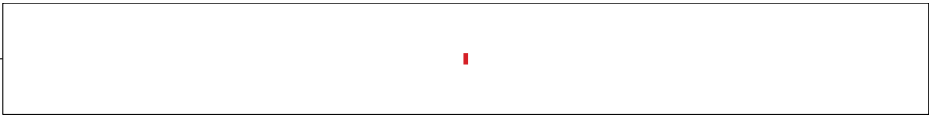
Genes

chr7:45942993–46043494 Marker Peak Tracks for: L5.6.NR4A2

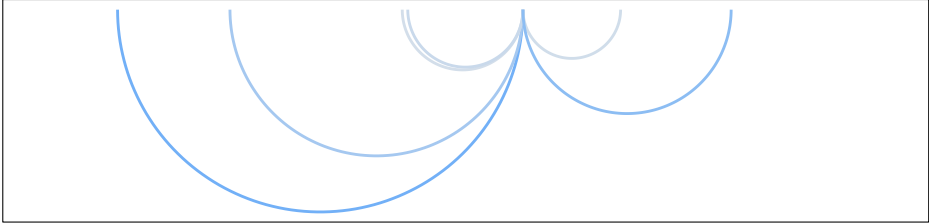
Coverage
(Norm. ATAC Signal Range (0–1.17) by ReadsInTSS)



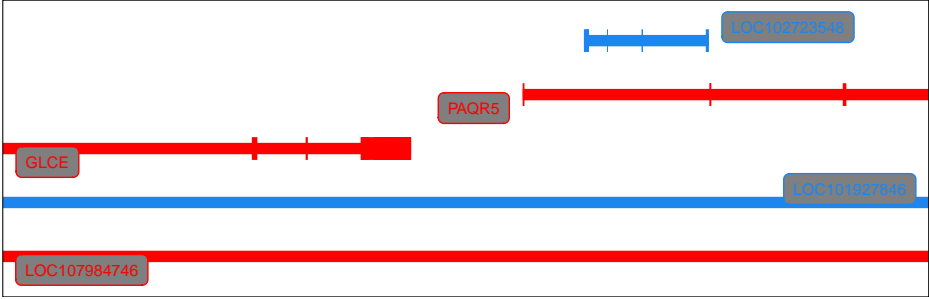
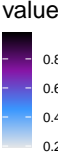
L5.6.NR4A2



Peaks

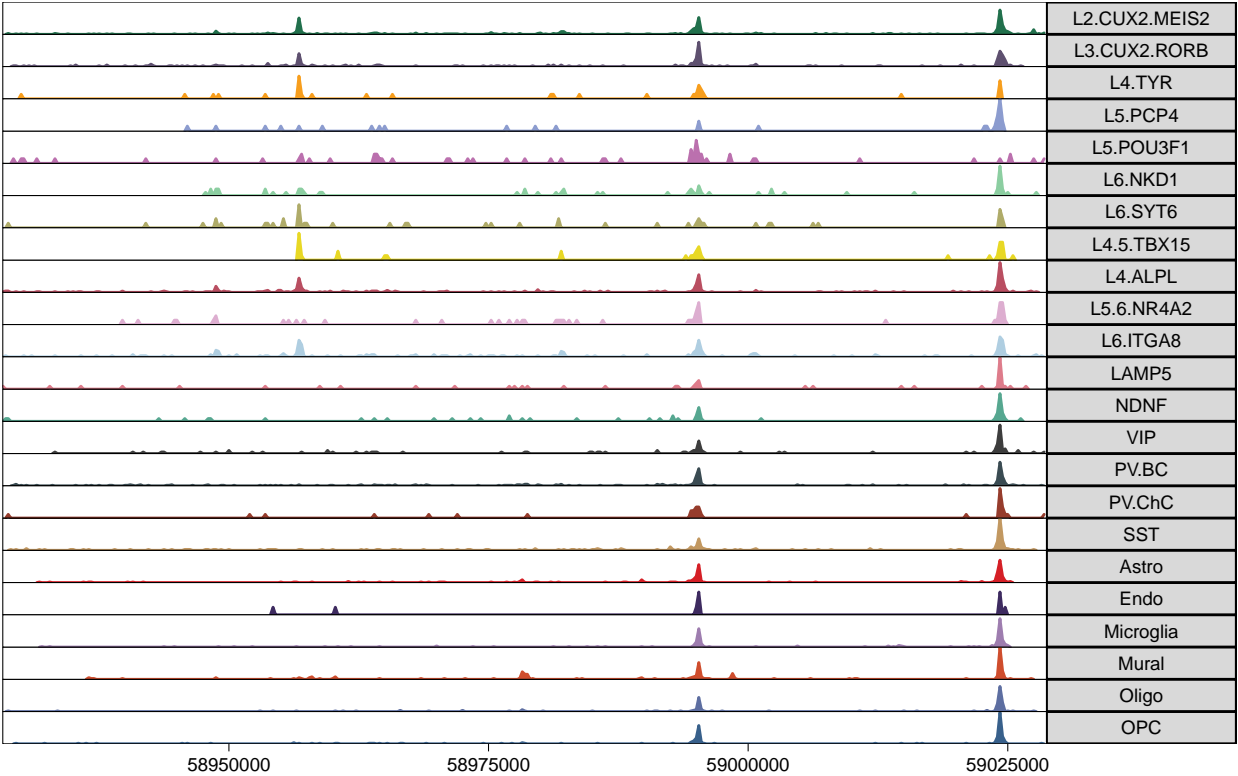


Peak2GeneLinks

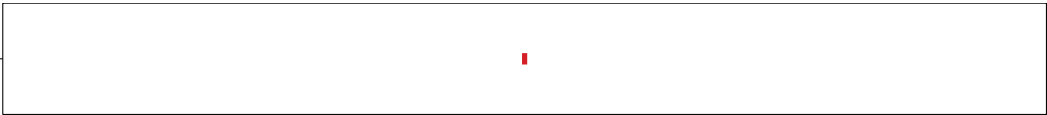


Genes

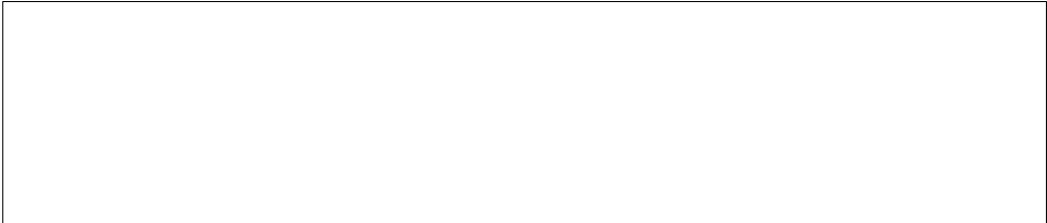
Coverage
(Norm. ATAC Signal Range 0-1.66) by ReadsInTSS)



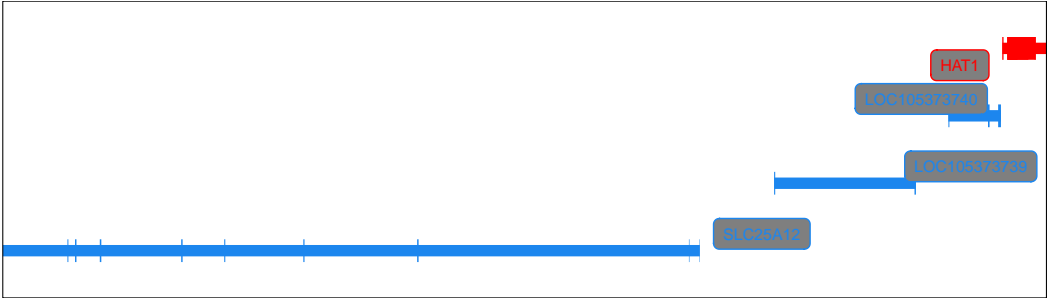
L5.6.NR4A2



Peaks

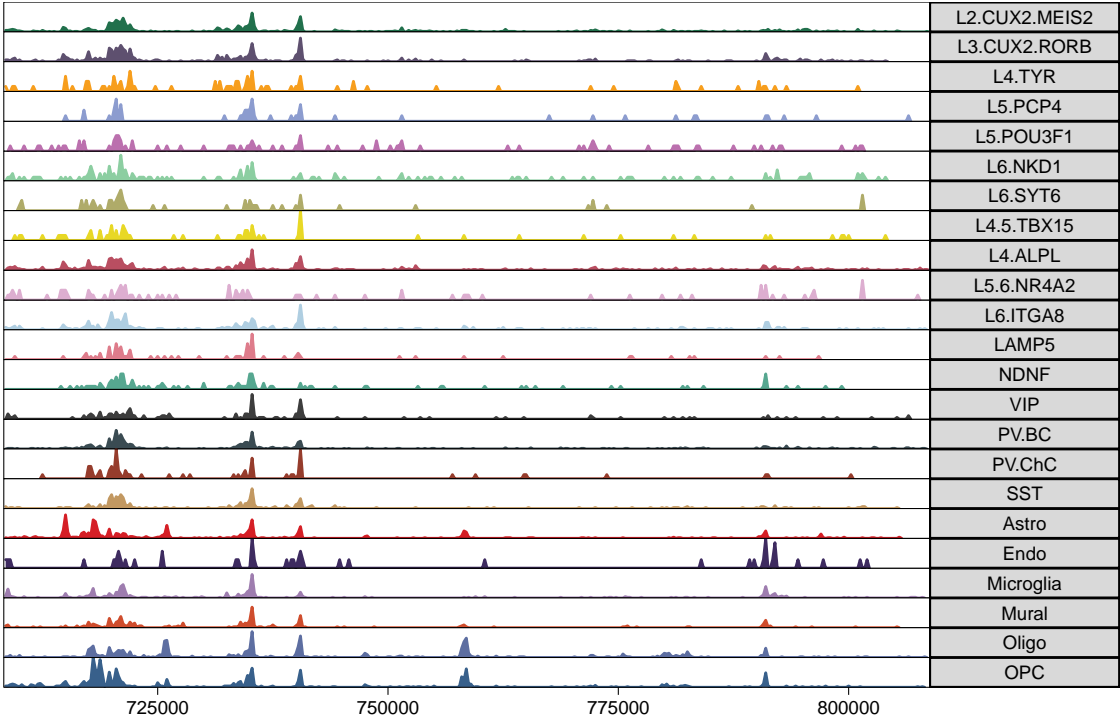


LoopTrack



Genes

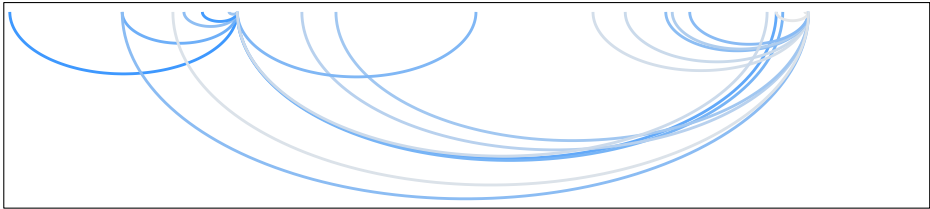
Coverage
(Norm. ATAC Signal Range (0–1.37) by ReadsInTSS)



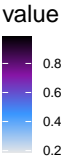
L5.6.NR4A2



Peaks



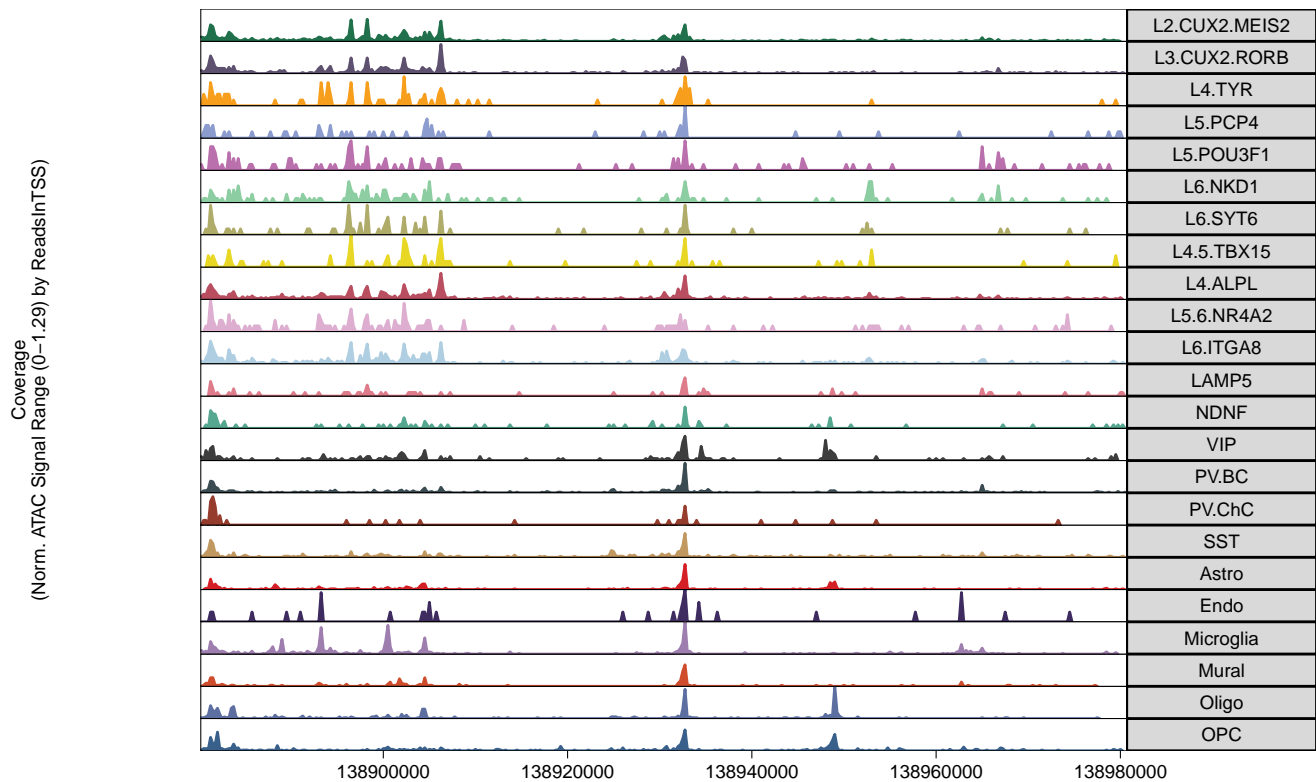
Peak2GeneLinks



GeneTrack

strand ● ·

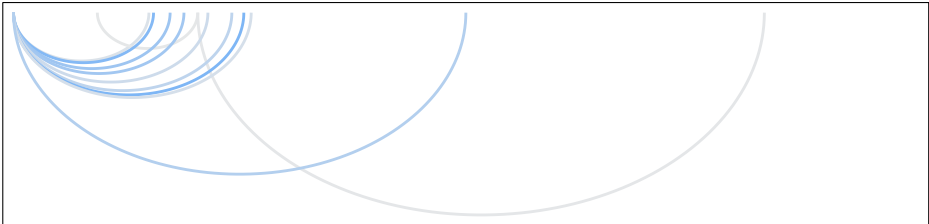
chr7:138880187-138980688 Marker Peak Tracks for: L5.6.NR4A2



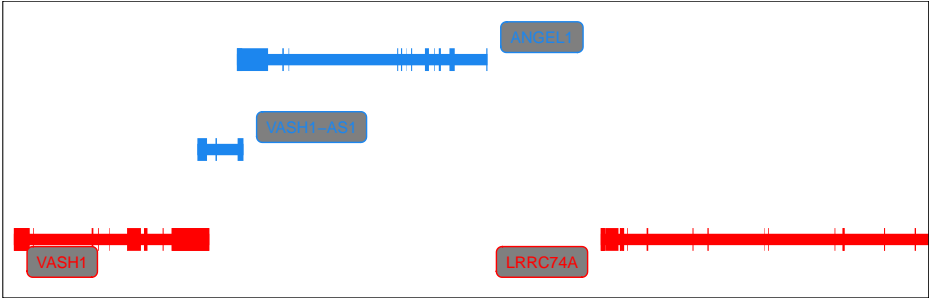
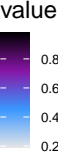
L5.6.NR4A2



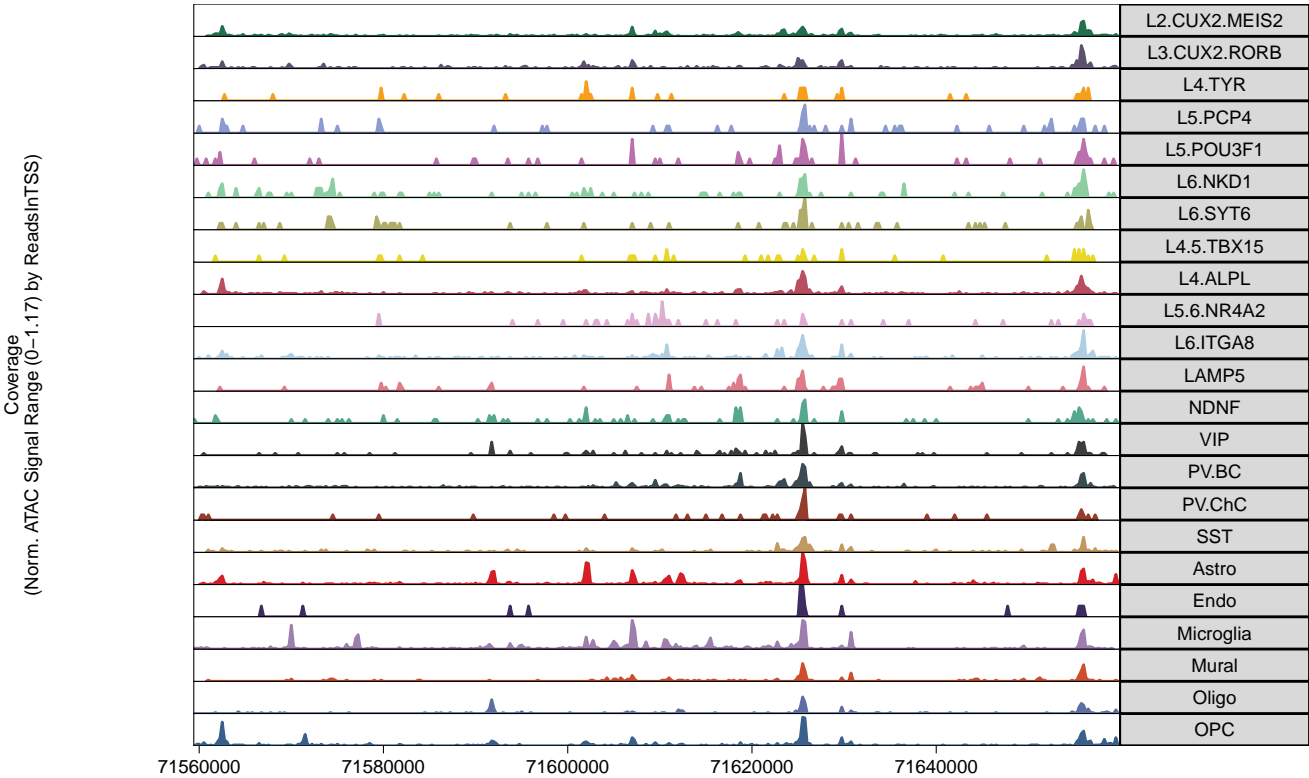
Peaks



Peak2GeneLinks



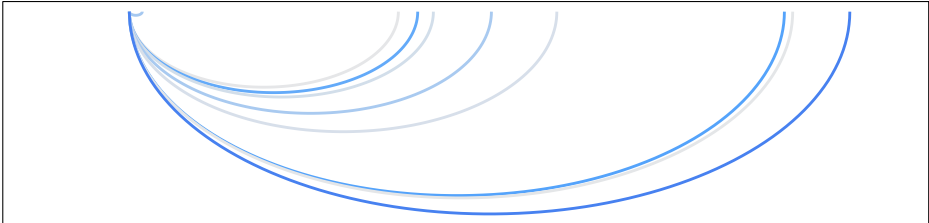
Genes



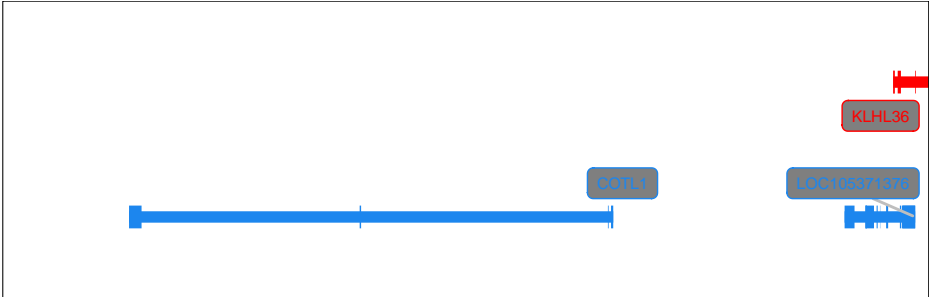
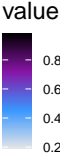
L5.6.NR4A2



Peaks

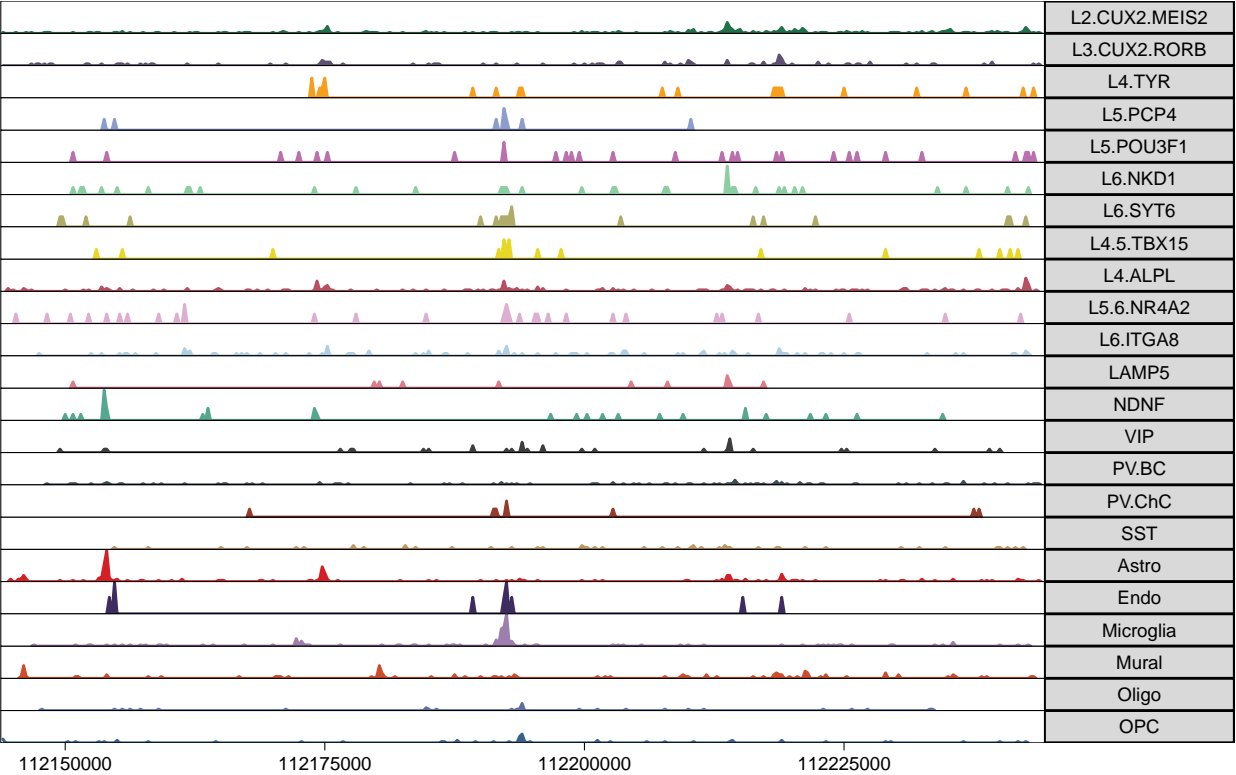


Peak2GeneLinks



Genes

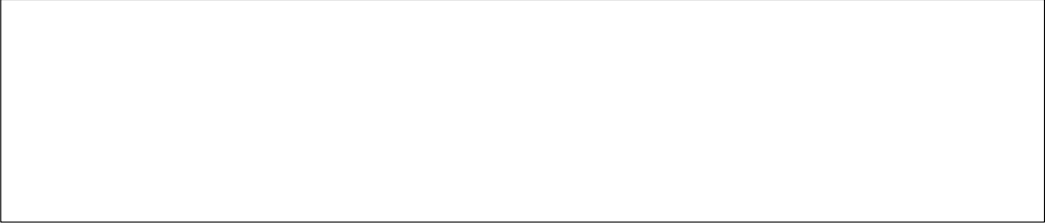
Coverage
(Norm. ATAC Signal Range (0–0.76) by ReadsInTSS)



L5.6.NR4A2



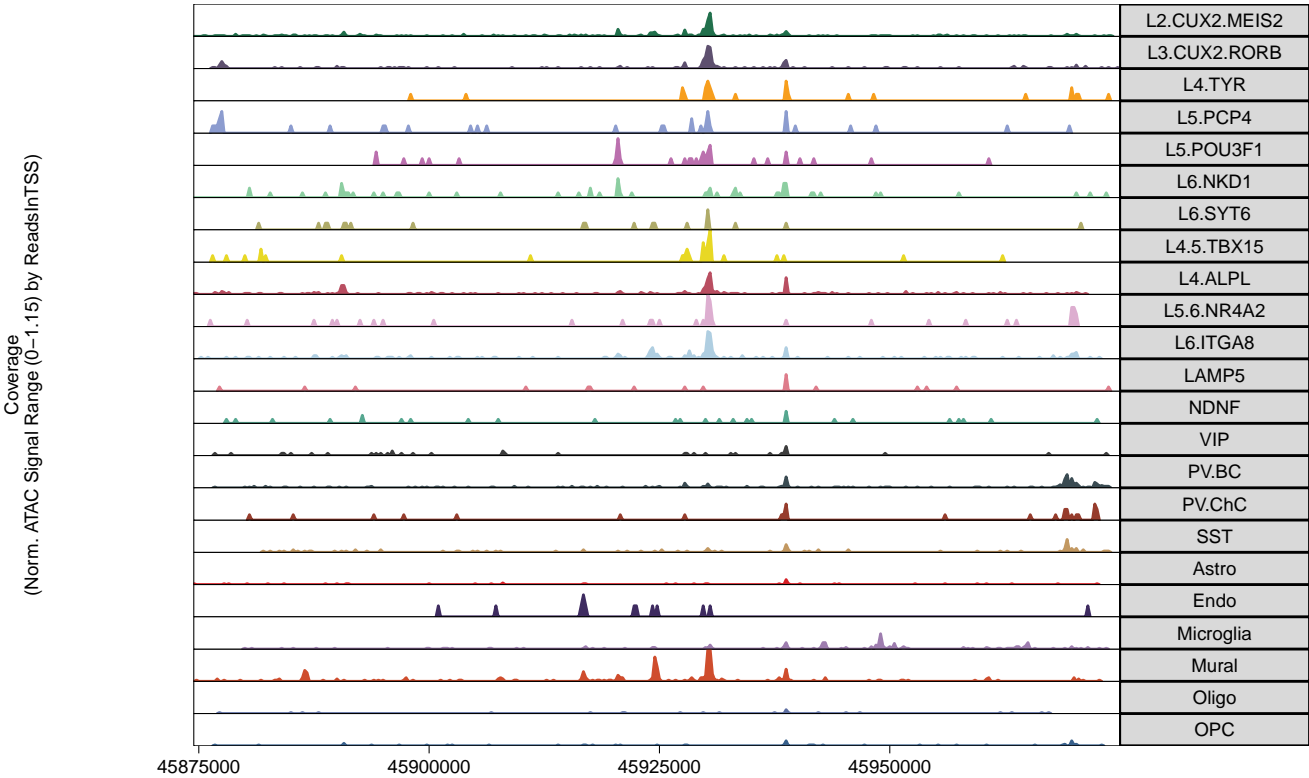
Peaks



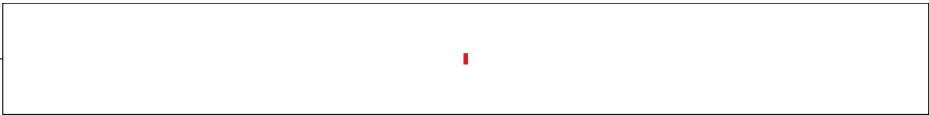
LoopTrack



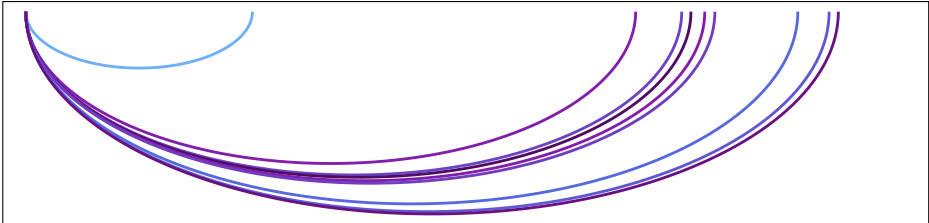
Genes



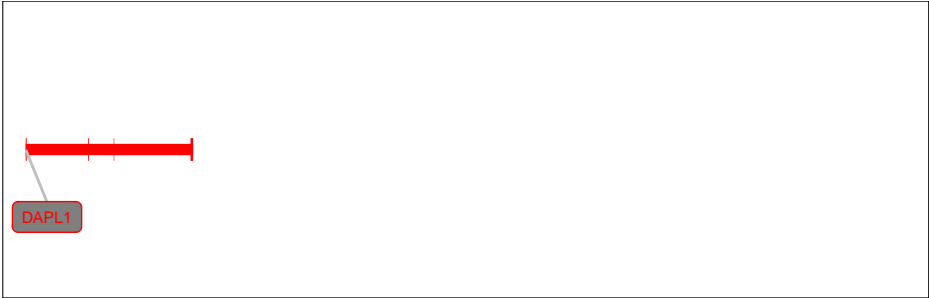
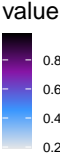
L5.6.NR4A2



Peaks



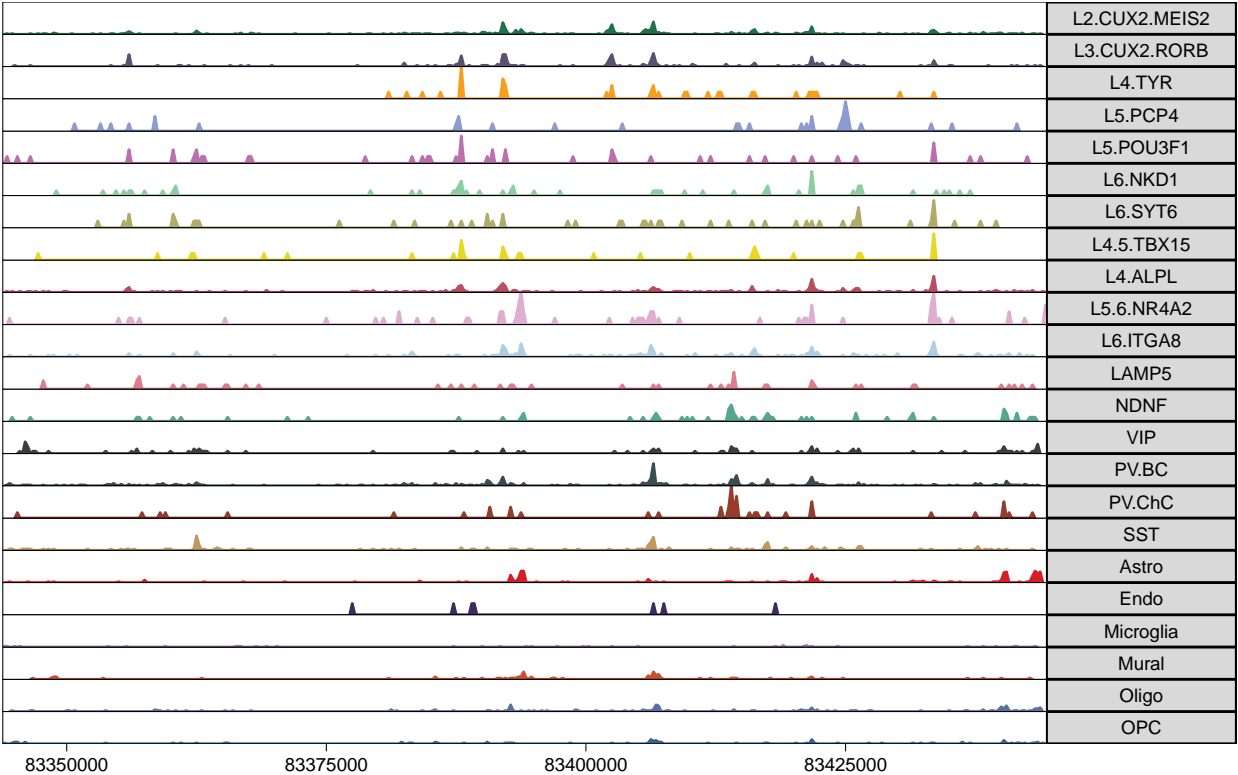
Peak2GeneLinks



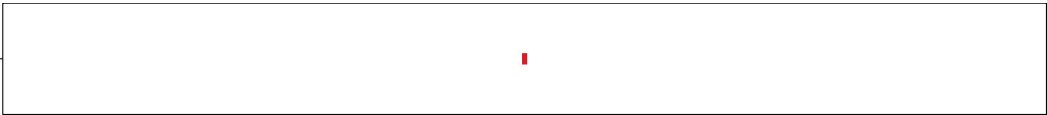
Genes

chr4:83343845–83444346 Marker Peak Tracks for: L5.6.NR4A2

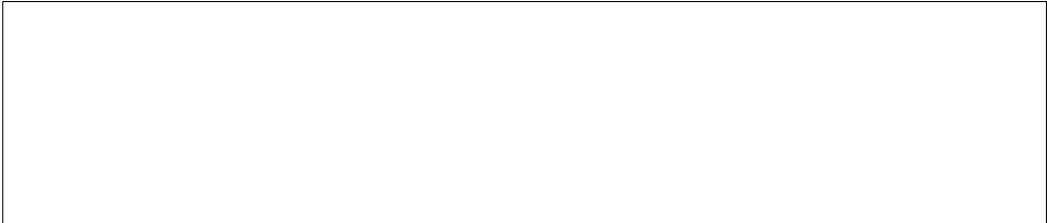
Coverage
(Norm. ATAC Signal Range 0–1.12) by ReadsInTSS)



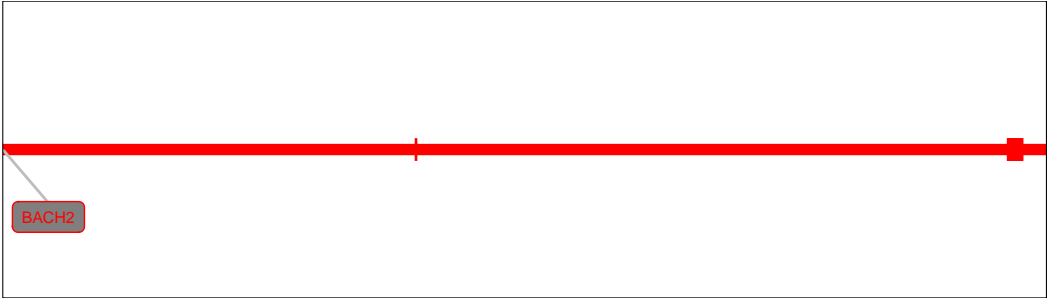
L5.6.NR4A2



Peaks

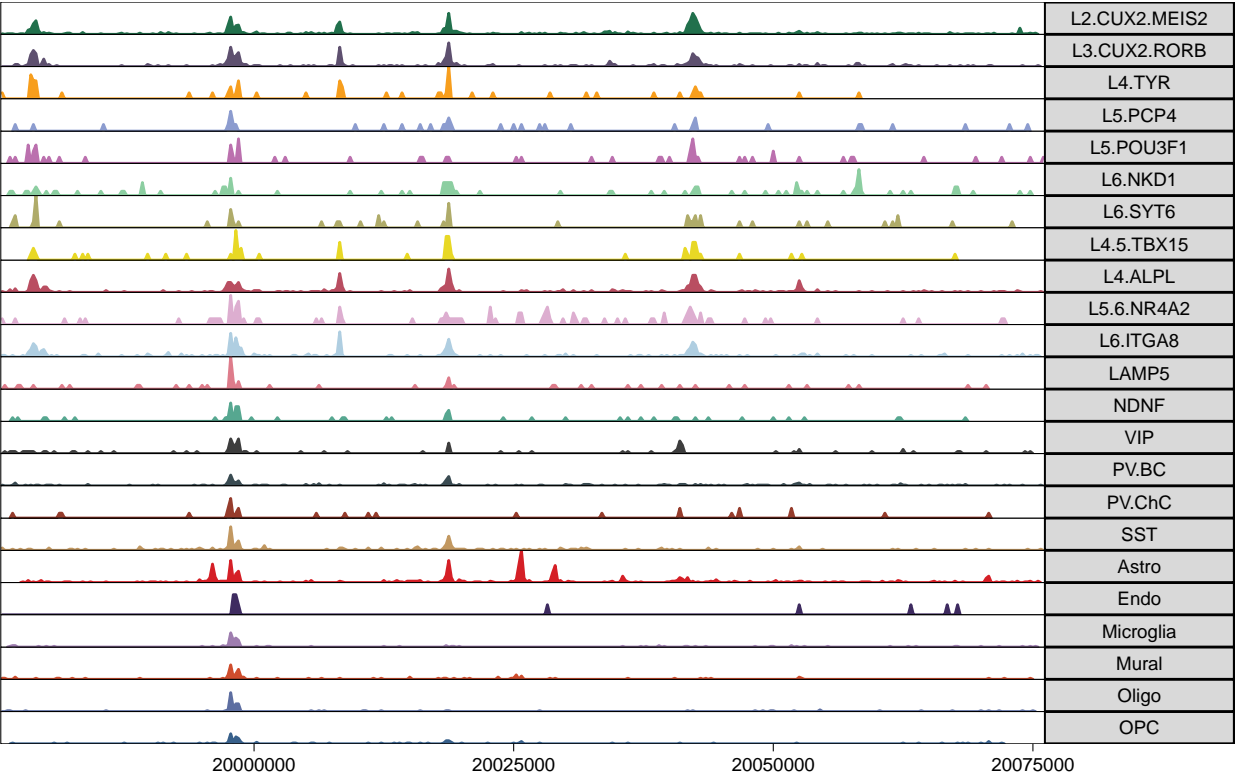


LoopTrack

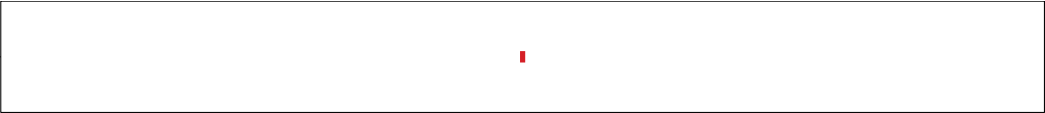


Genes

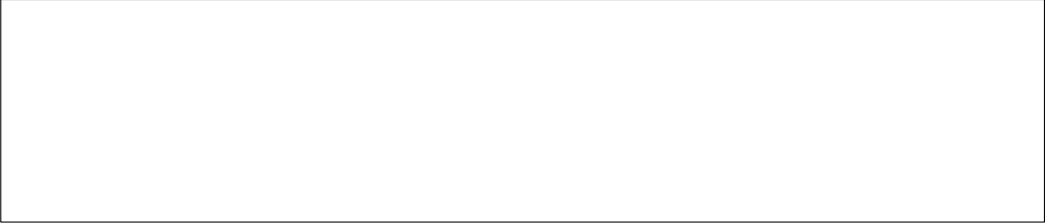
Coverage
(Norm. ATAC Signal Range 0–1.25) by ReadsInTSS)



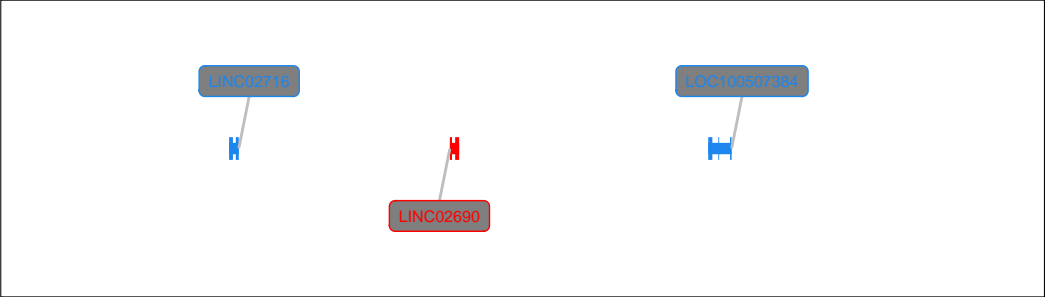
L5.6.NR4A2



Peaks

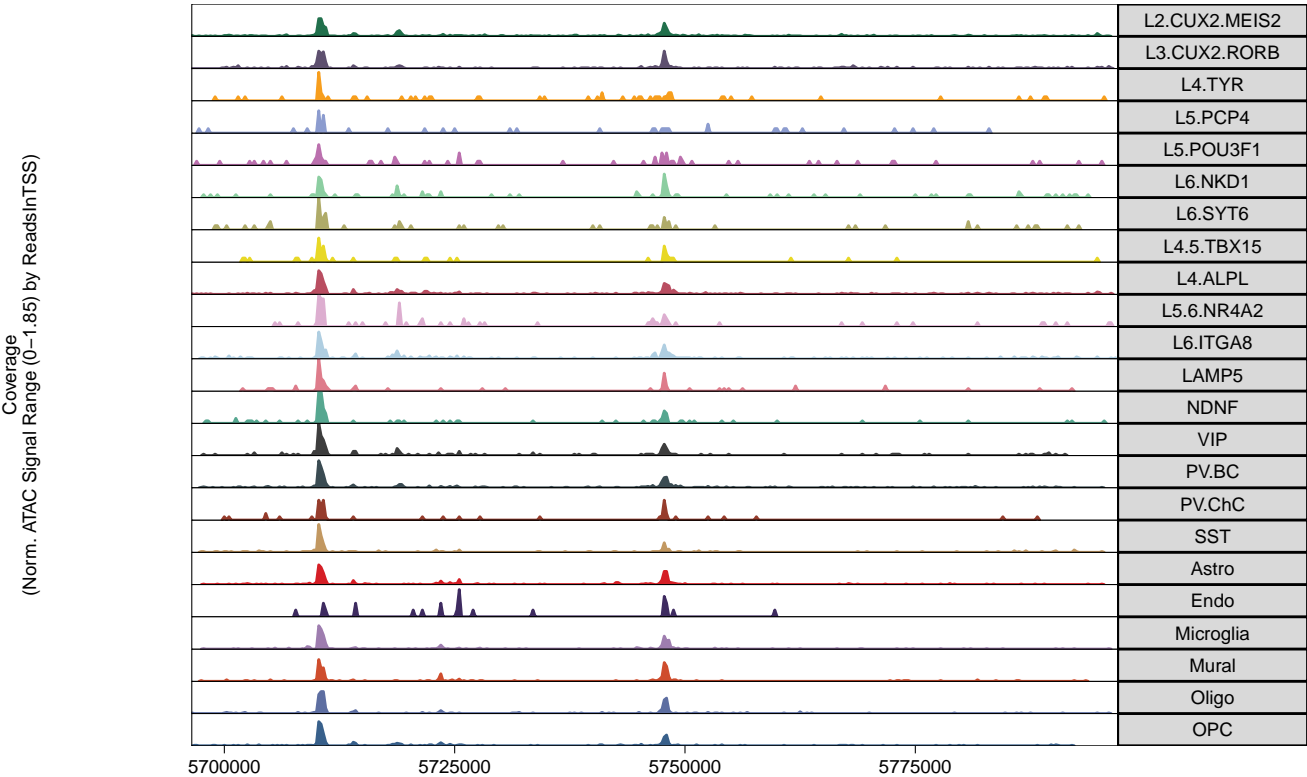


LoopTrack

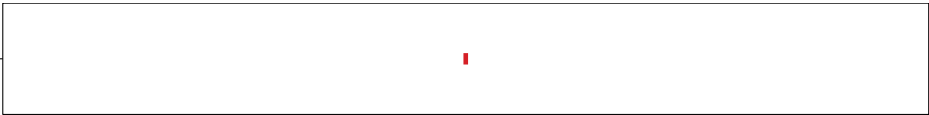


Genes

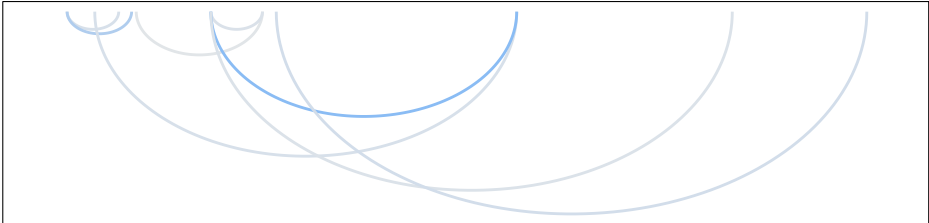
chr19:5696467–5796968 Marker Peak Tracks for: L5.6.NR4A2



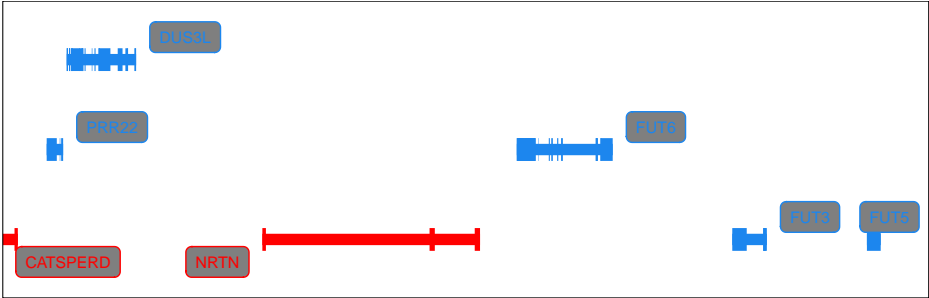
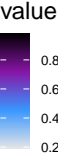
L5.6.NR4A2



Peaks

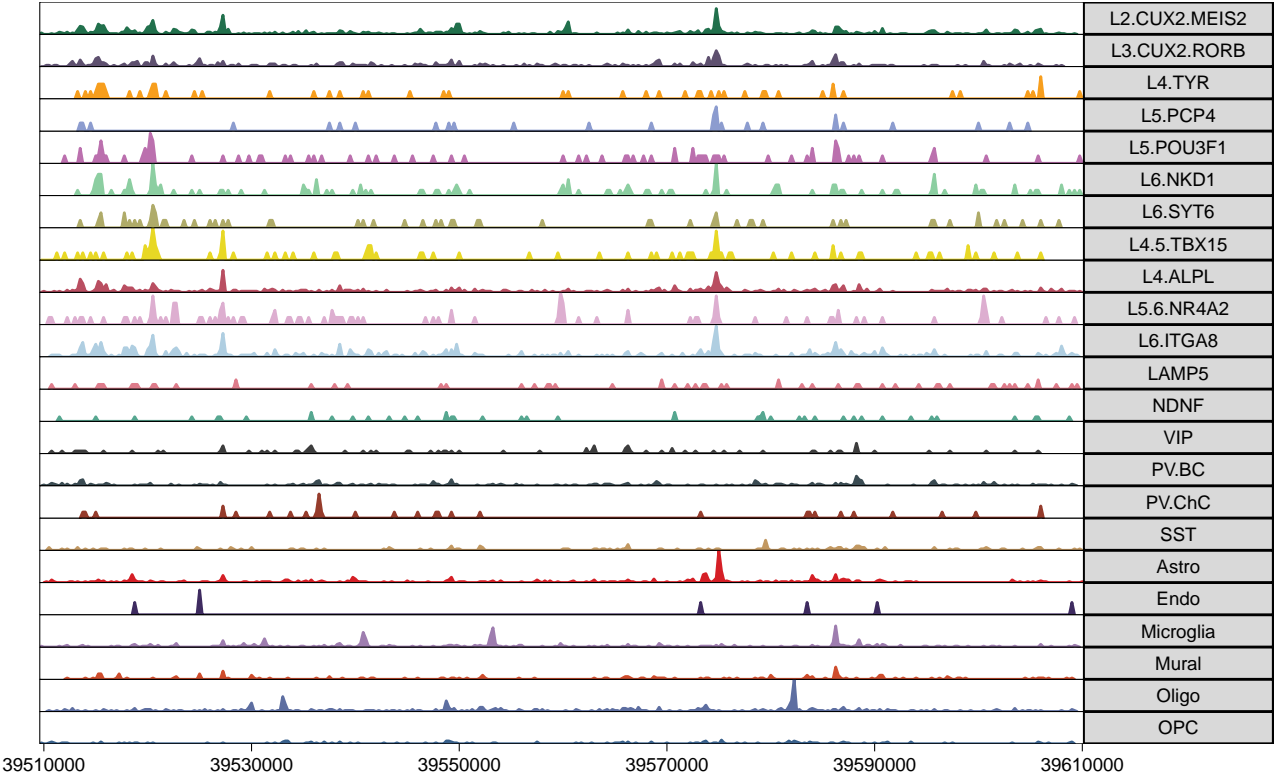


Peak2GeneLinks



Genes

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



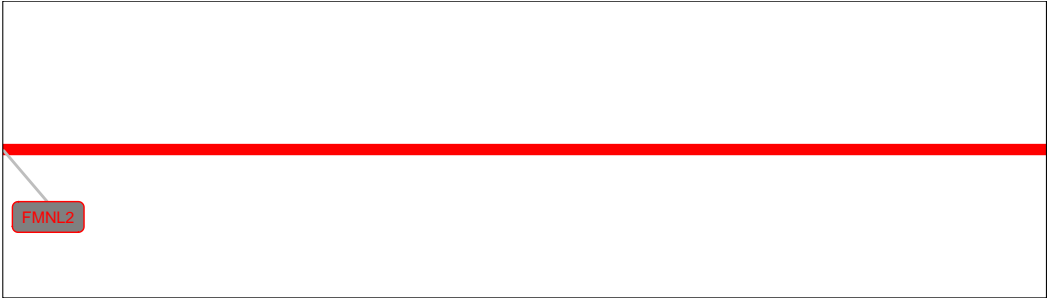
L5.6.NR4A2



Peaks

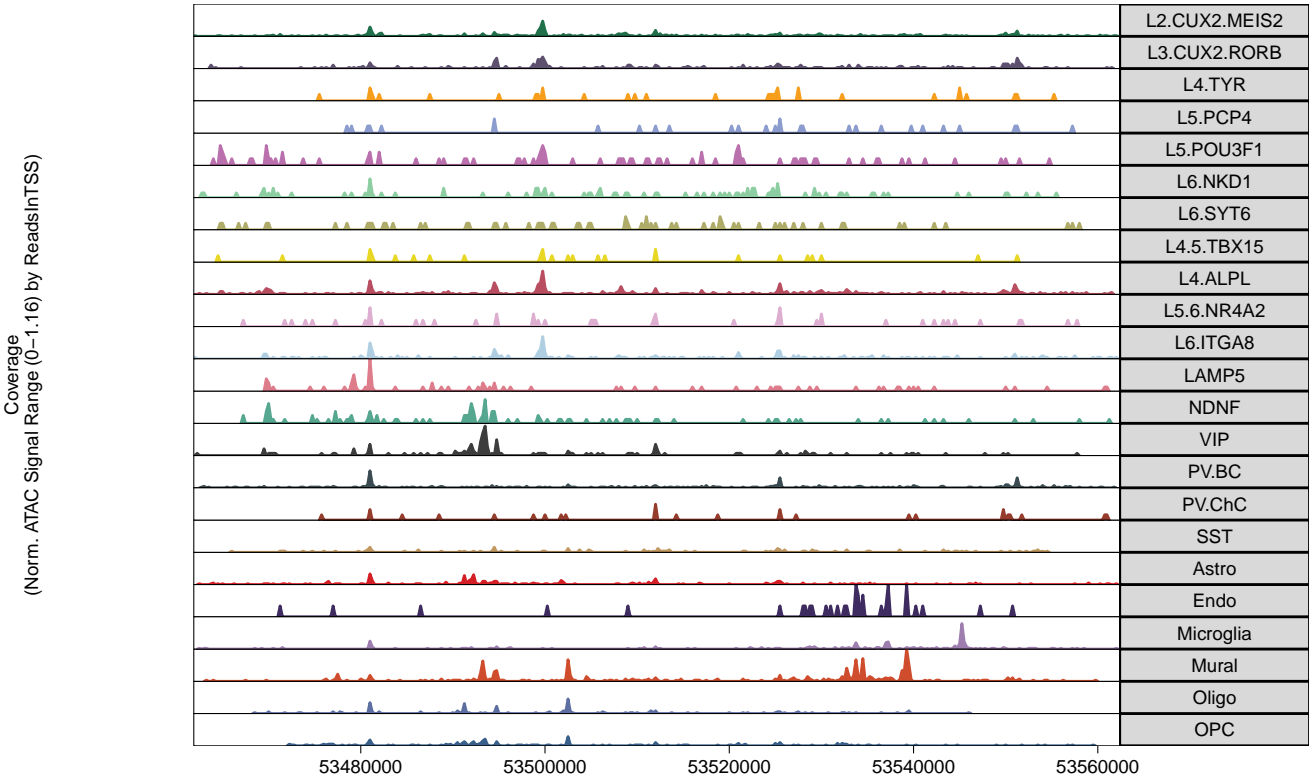


LoopTrack



Genes

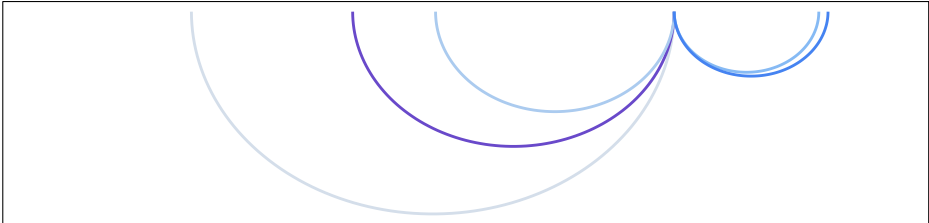
chr20:53461877-53562378 Marker Peak Tracks for: L5.6.NR4A2



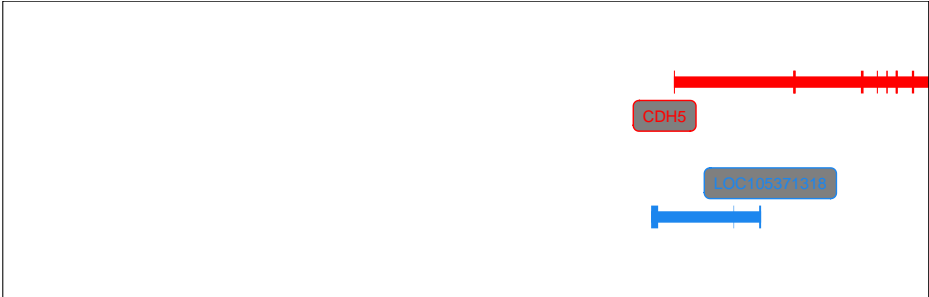
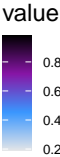
L5.6.NR4A2



Peaks



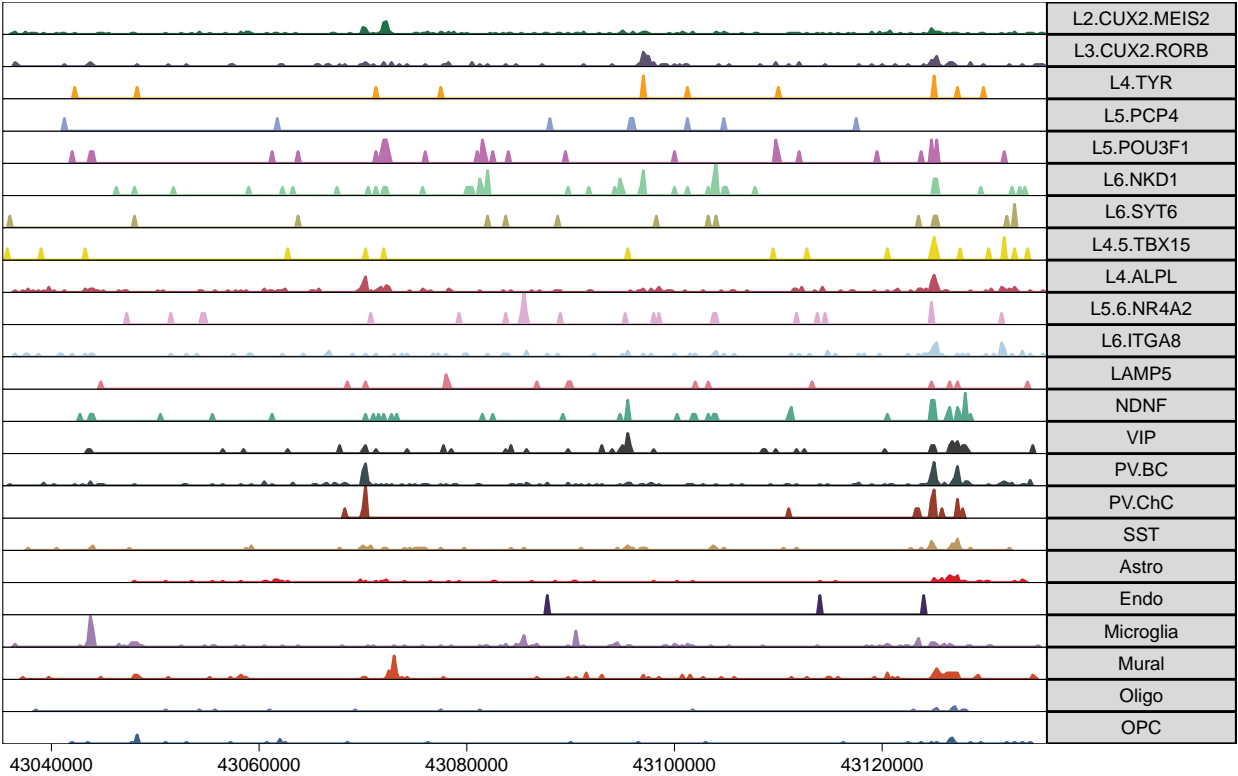
Peak2GeneLinks



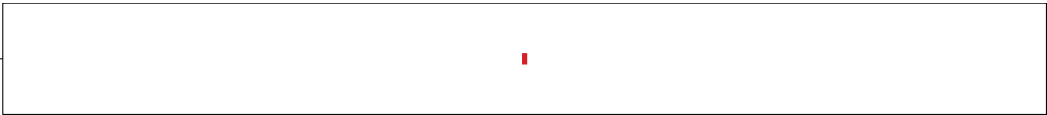
Genes

chr4:43035317–43135818 Marker Peak Tracks for: L5.6.NR4A2

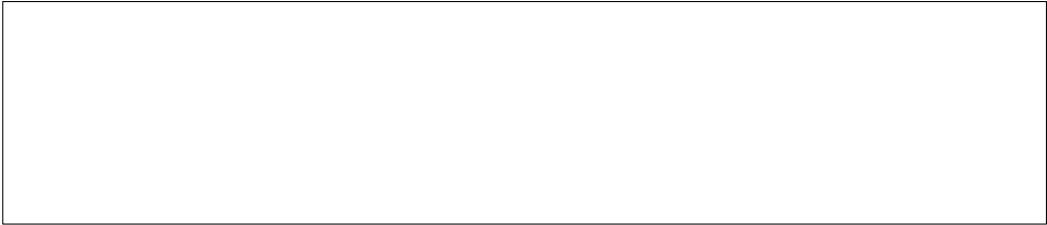
Coverage
(Norm. ATAC Signal Range 0–0.65) by ReadsInTSS)



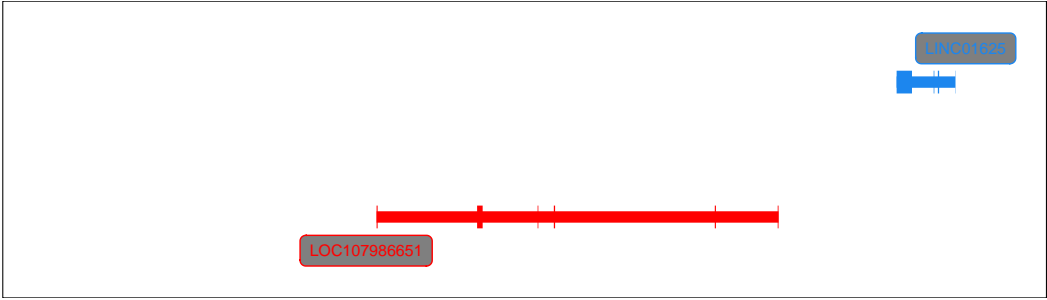
L5.6.NR4A2



Peaks

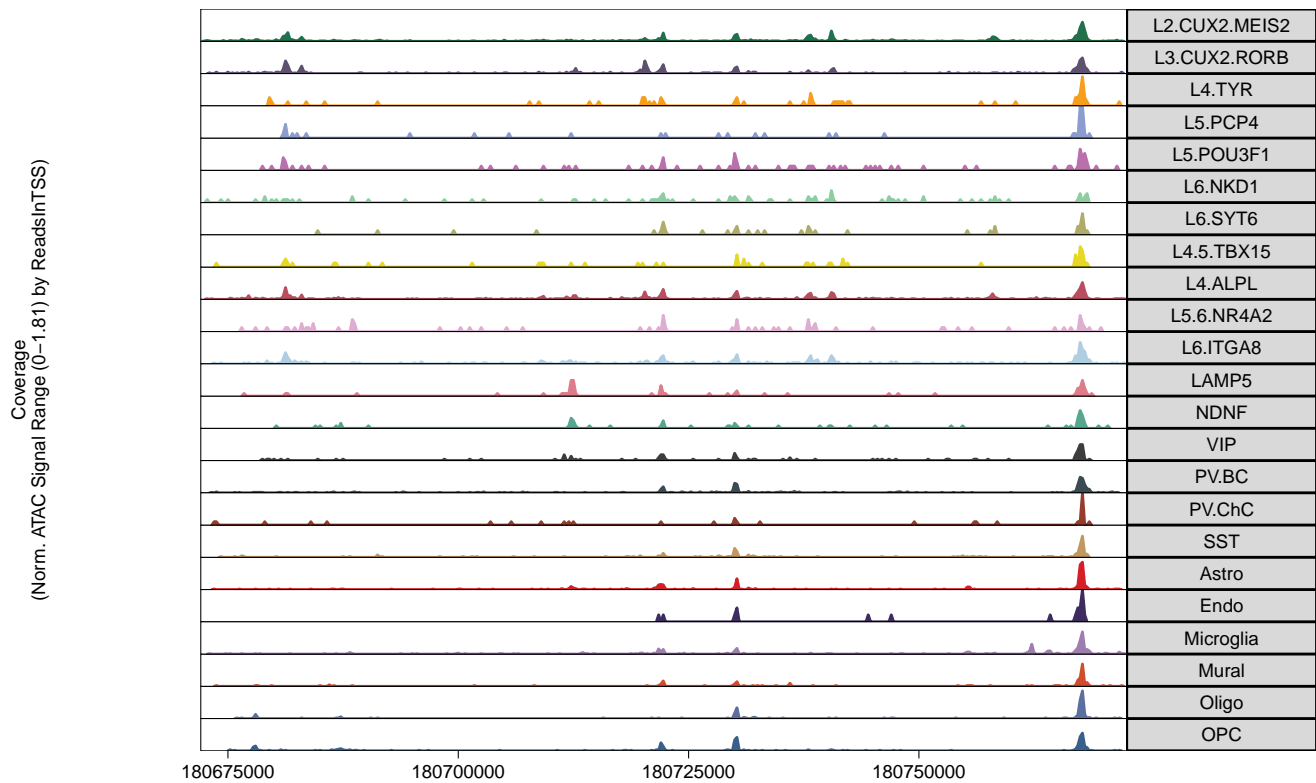


LoopTrack

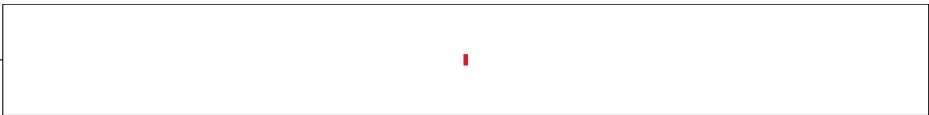


Genes

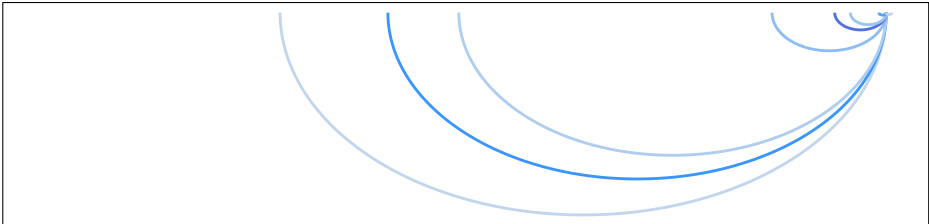
chr5:180672053–180772554 Marker Peak Tracks for: L5.6.NR4A2



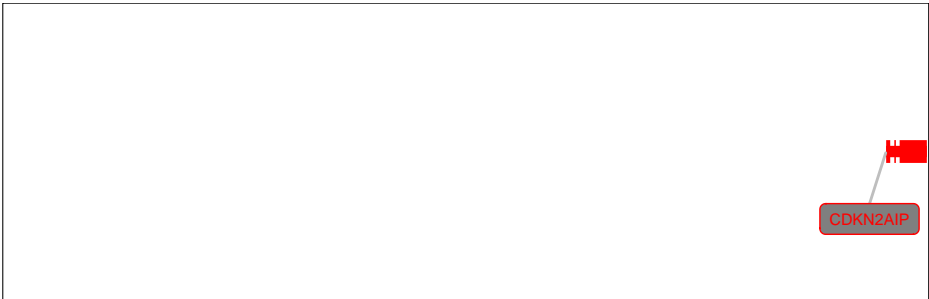
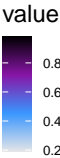
L5.6.NR4A2



Peaks

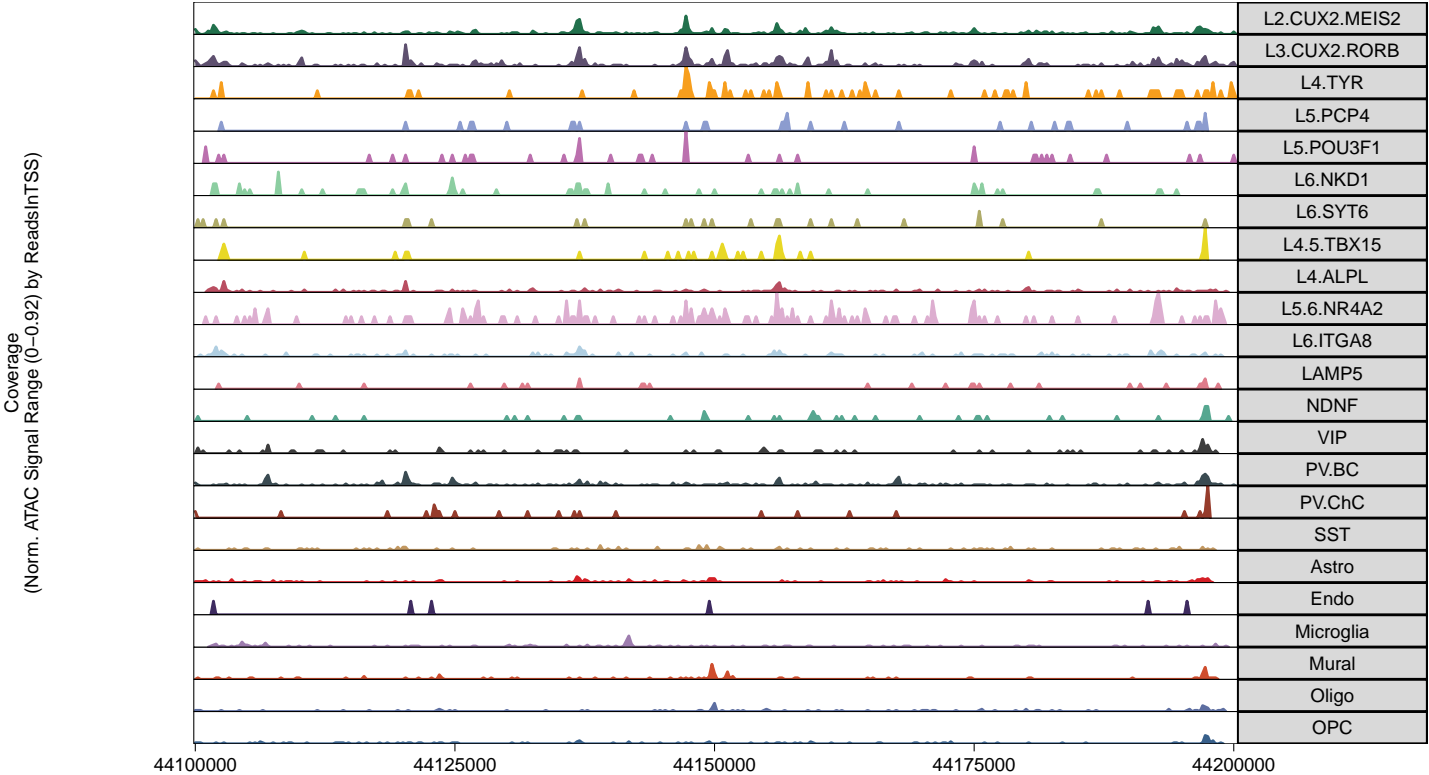


Peak2GeneLinks

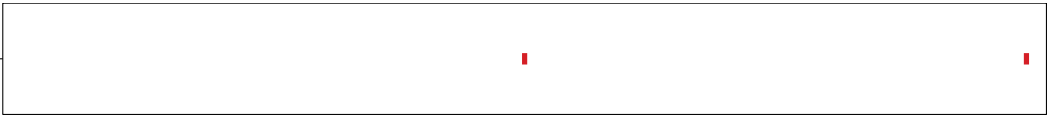


Genes

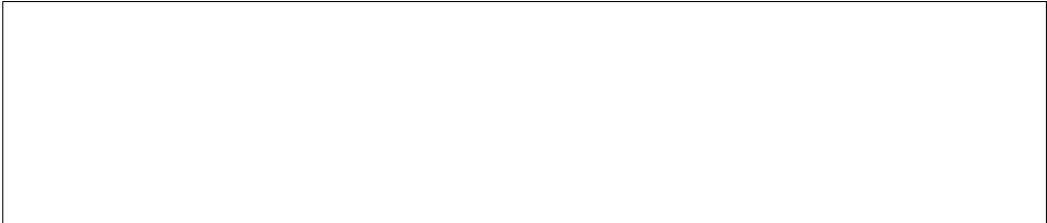
chr3:44099854–44200355 Marker Peak Tracks for: L5.6.NR4A2



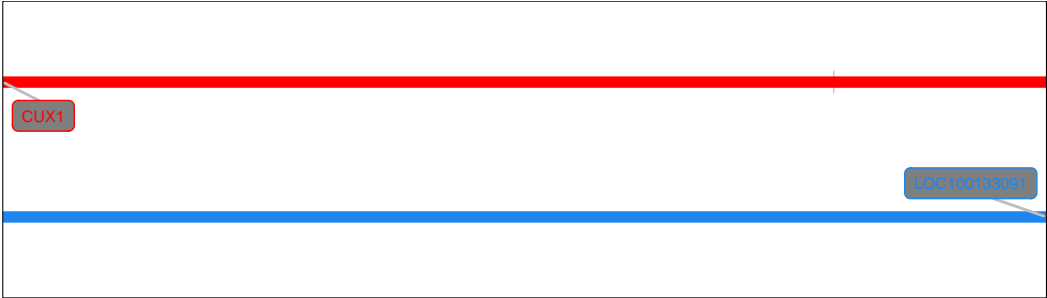
L5.6.NR4A2



Peaks



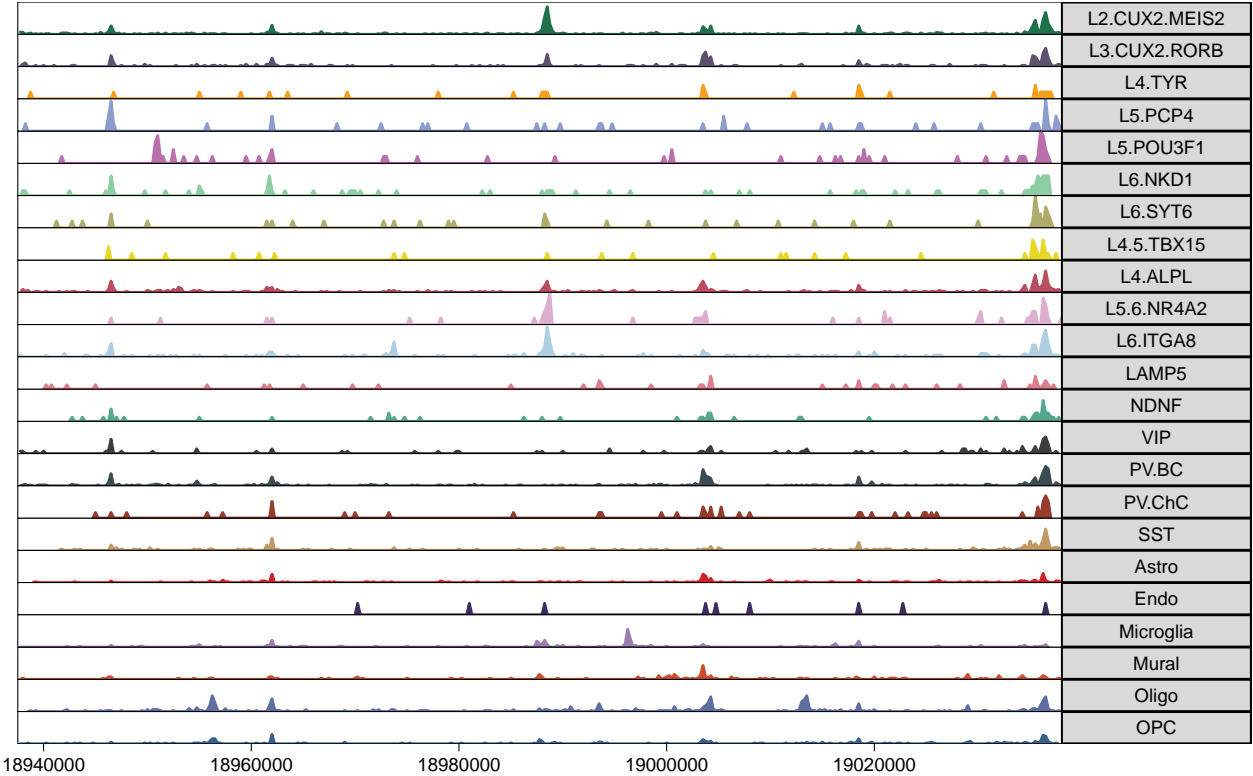
LoopTrack



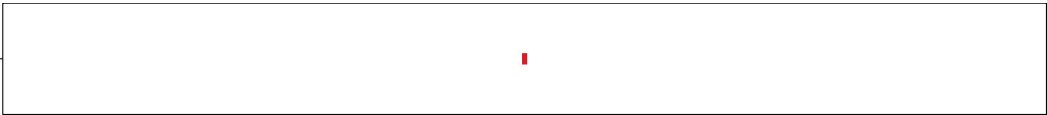
Genes

chr8:18937518–19038019 Marker Peak Tracks for: L5.6.NR4A2

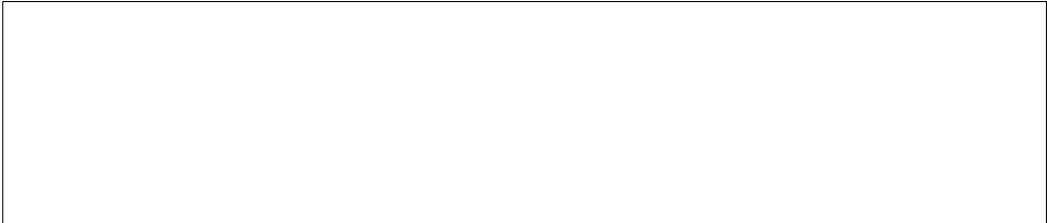
Coverage
(Norm. ATAC Signal Range (0–1.1) by Reads in TSS)



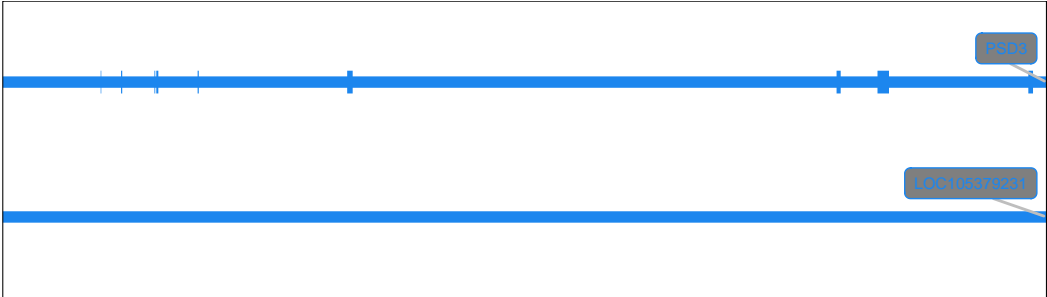
L5.6.NR4A2



Peaks

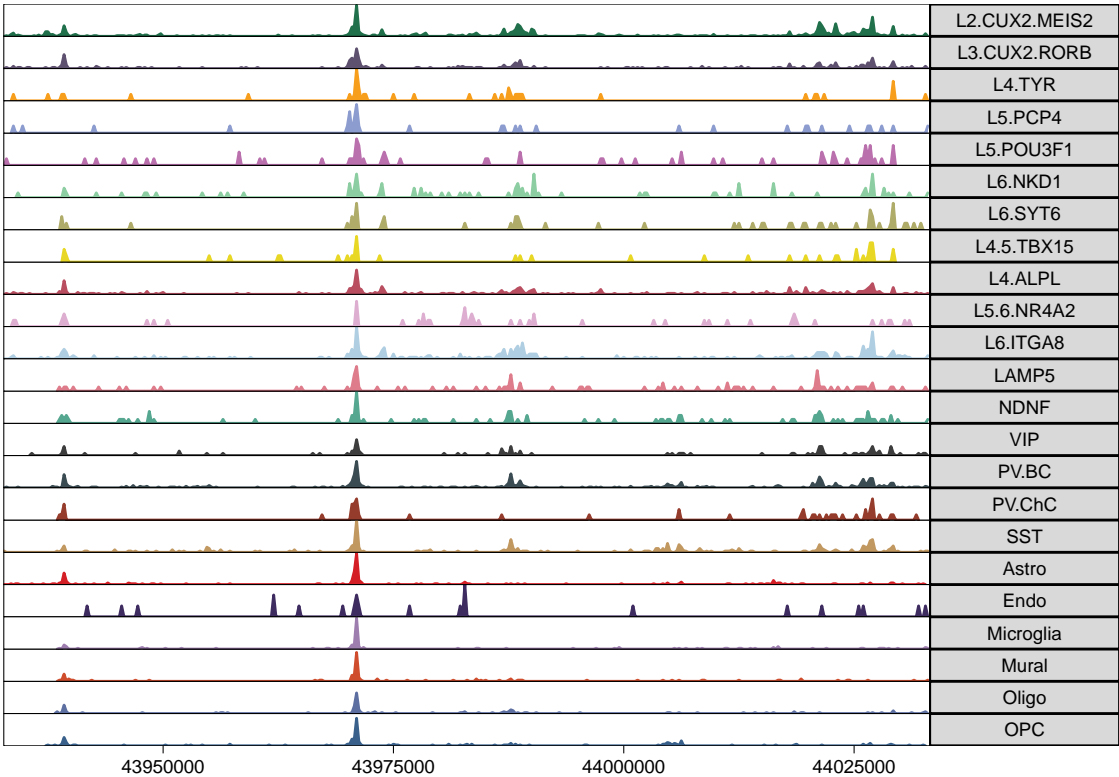


LoopTrack

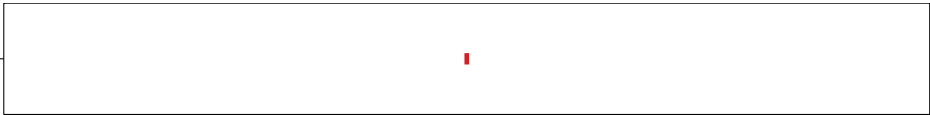


Genes

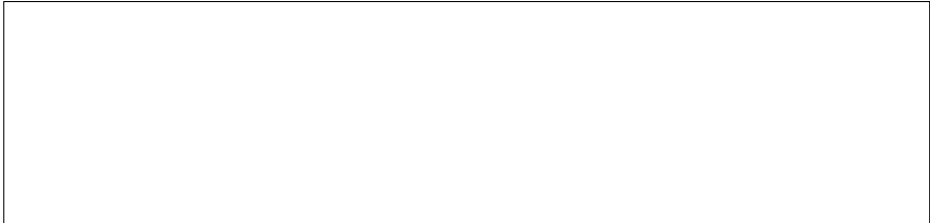
Coverage
(Norm. ATAC Signal Range (0-1.16) by ReadsInTSS)



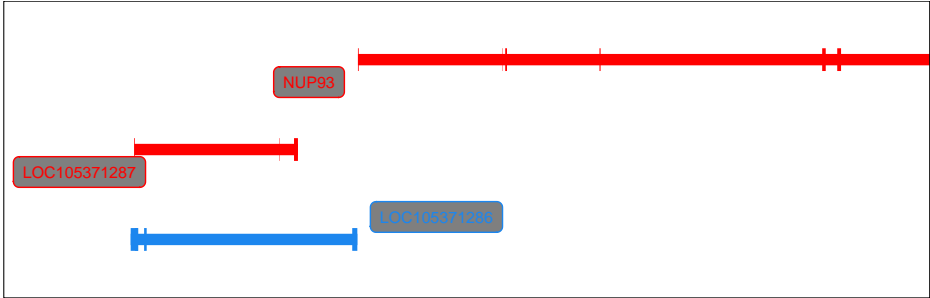
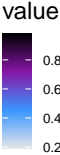
L5.6.NR4A2



Peaks

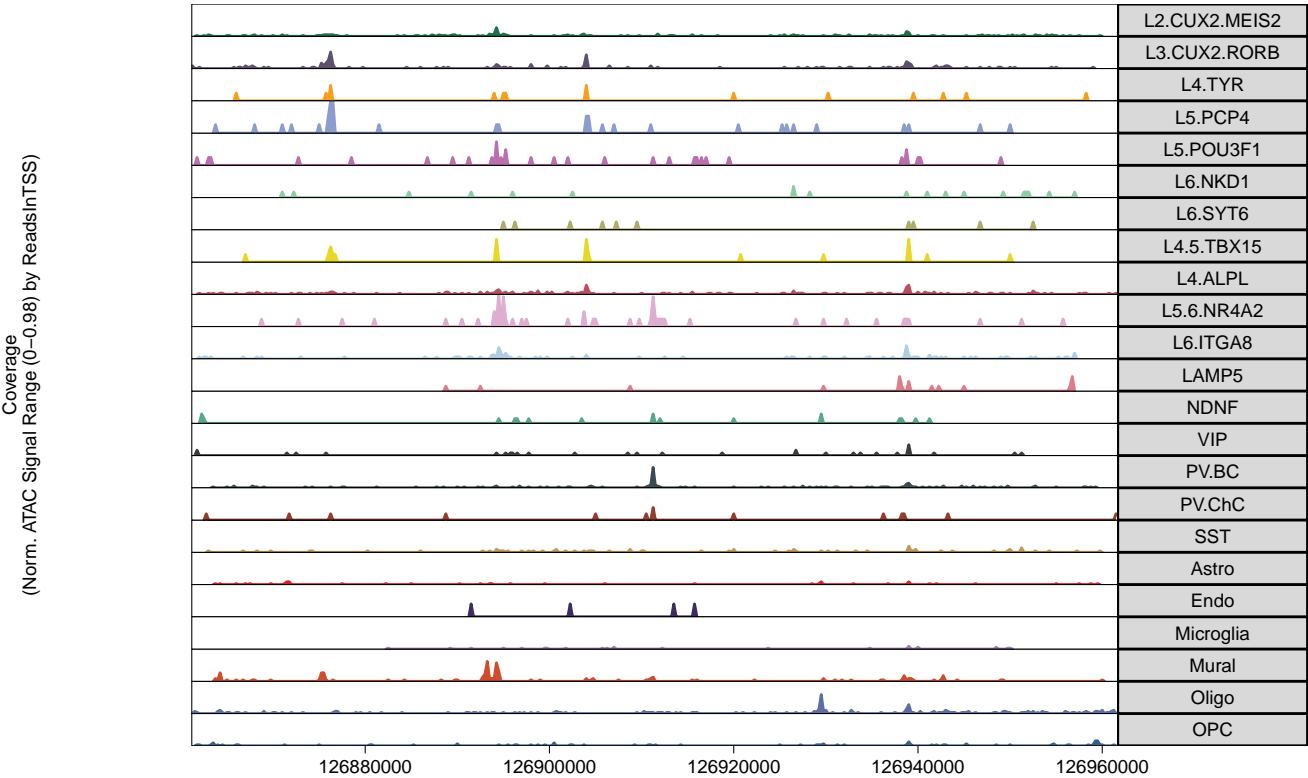


Peak2GeneLinks

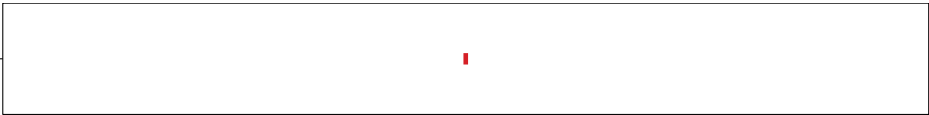


Genes

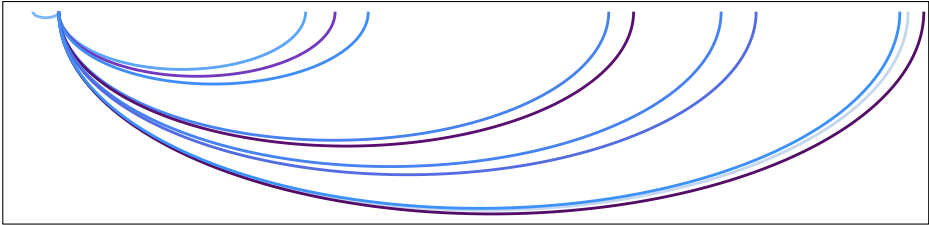
chr9:126861178–126961679 Marker Peak Tracks for: L5.6.NR4A2



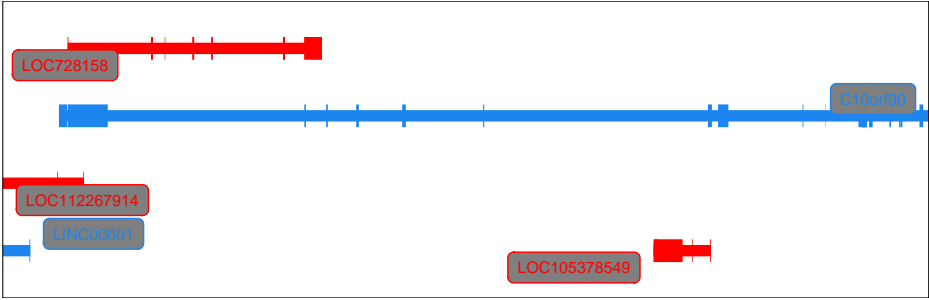
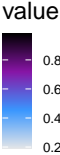
L5.6.NR4A2



Peaks

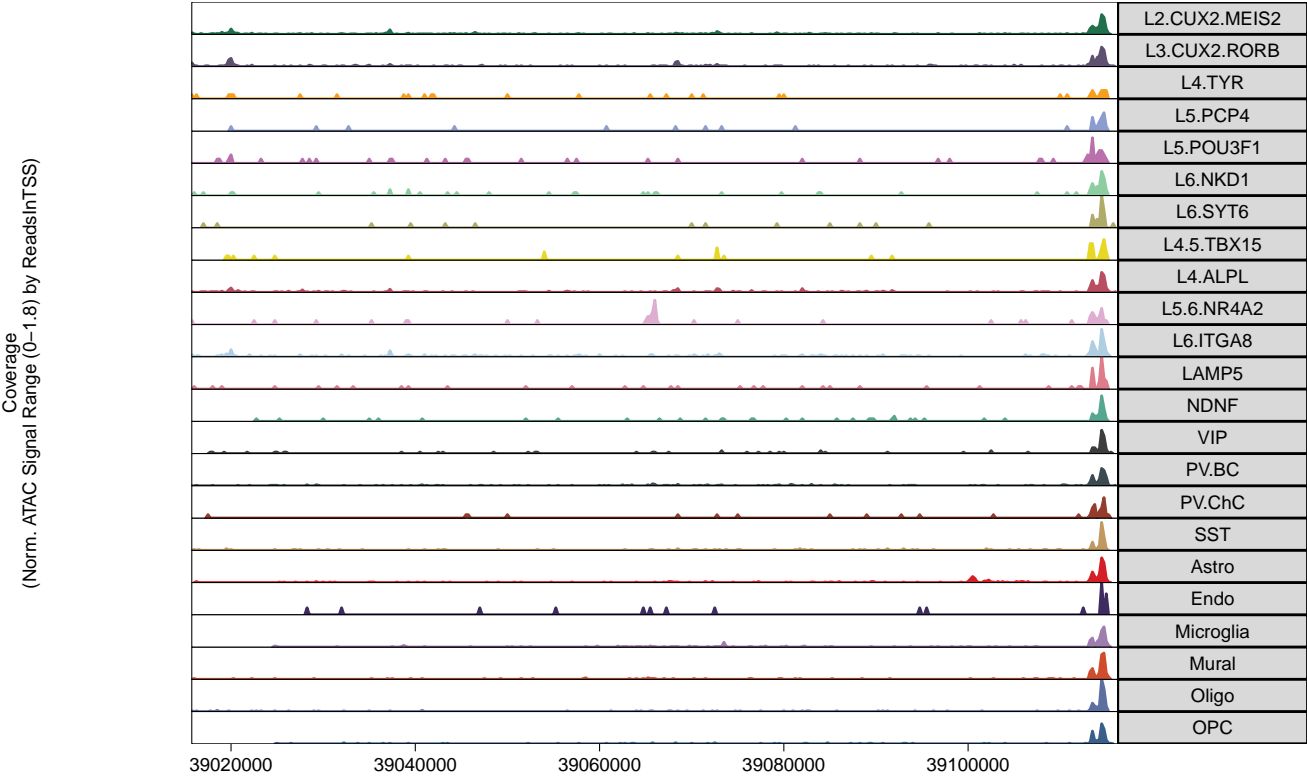


Peak2GeneLinks

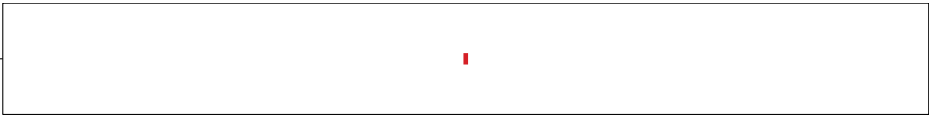


Genes

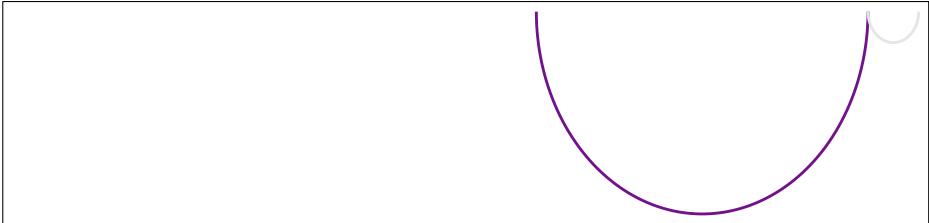
chr5:39015739–39116240 Marker Peak Tracks for: L5.6.NR4A2



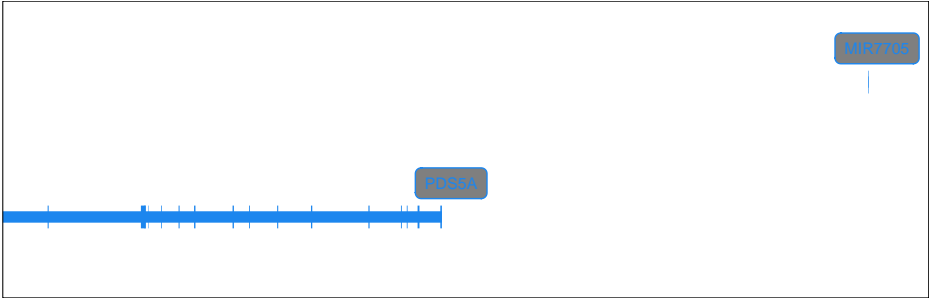
L5.6.NR4A2



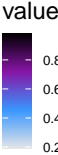
Peaks



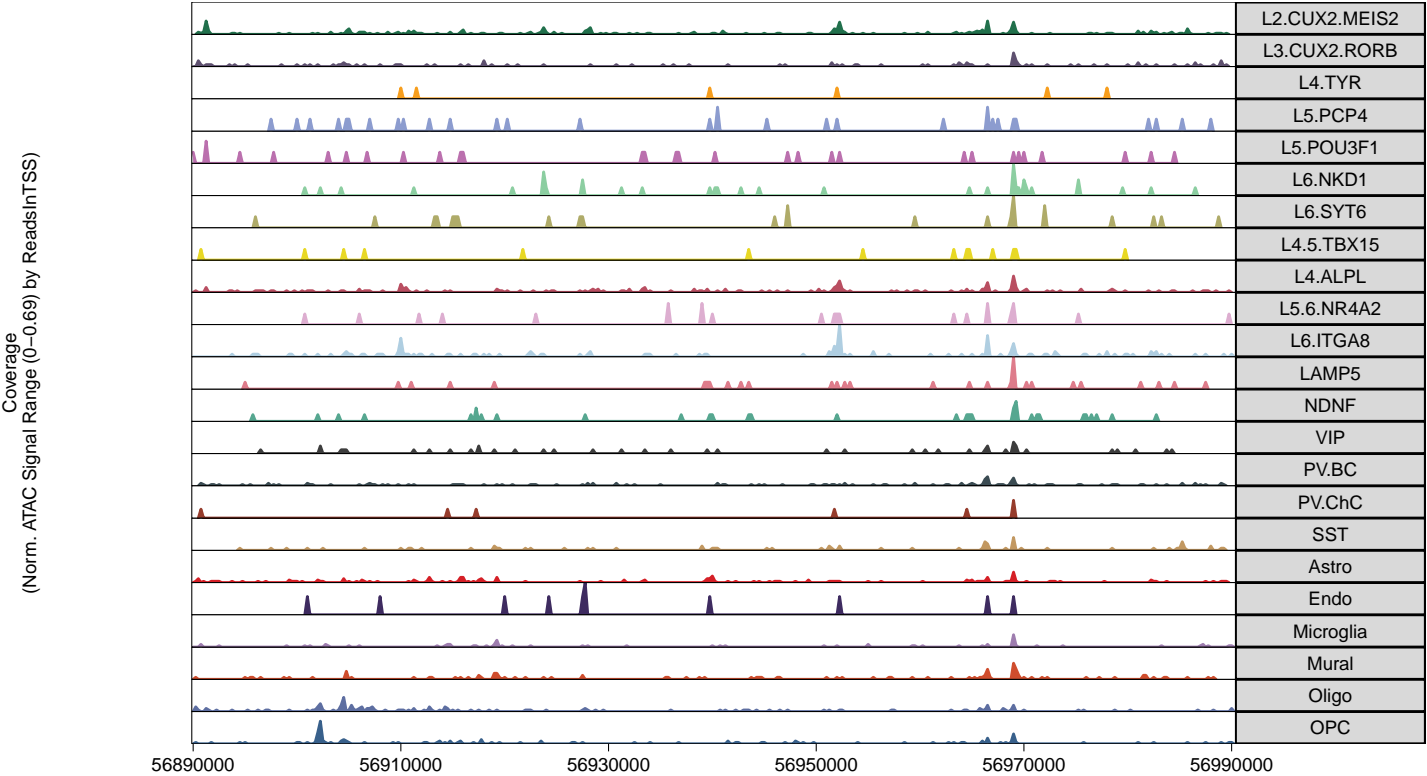
Peak2GeneLinks



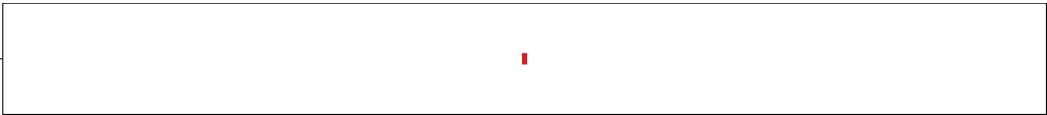
Genes



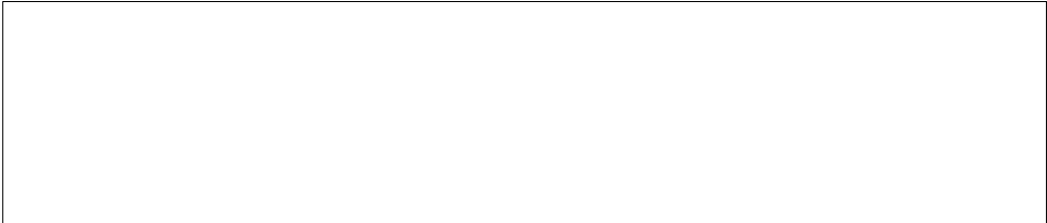
chr4:56889868–56990369 Marker Peak Tracks for: L5.6.NR4A2



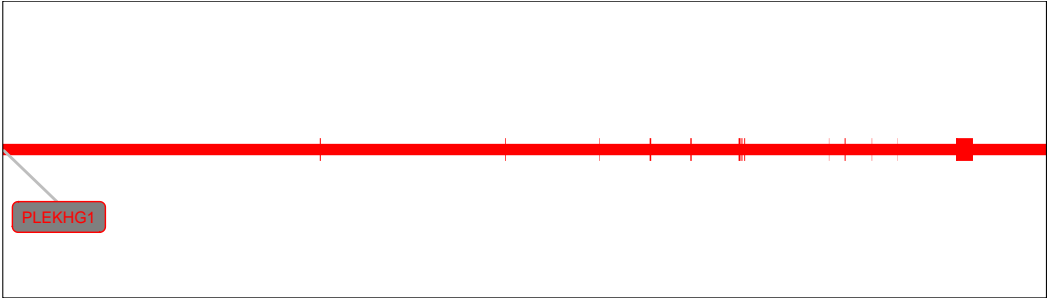
L5.6.NR4A2



Peaks

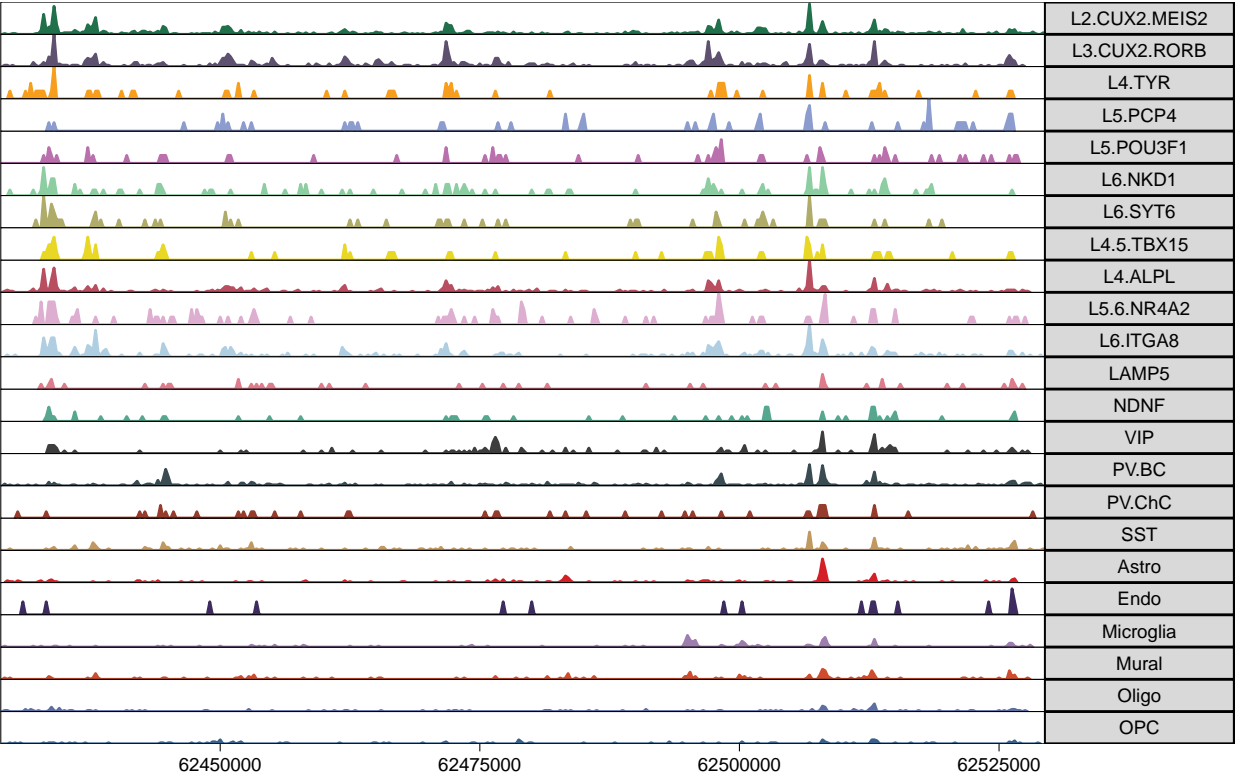


LoopTrack

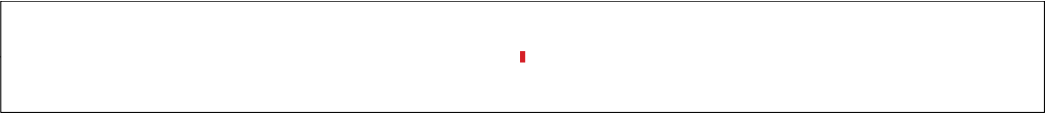


Genes

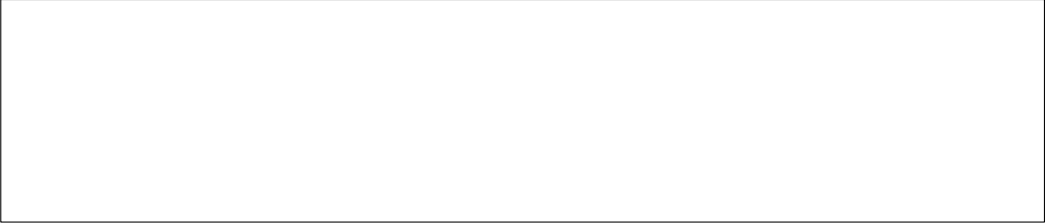
Coverage
(Norm. ATAC Signal Range (0–0.97) by ReadsInTSS)



L5.6.NR4A2



Peaks

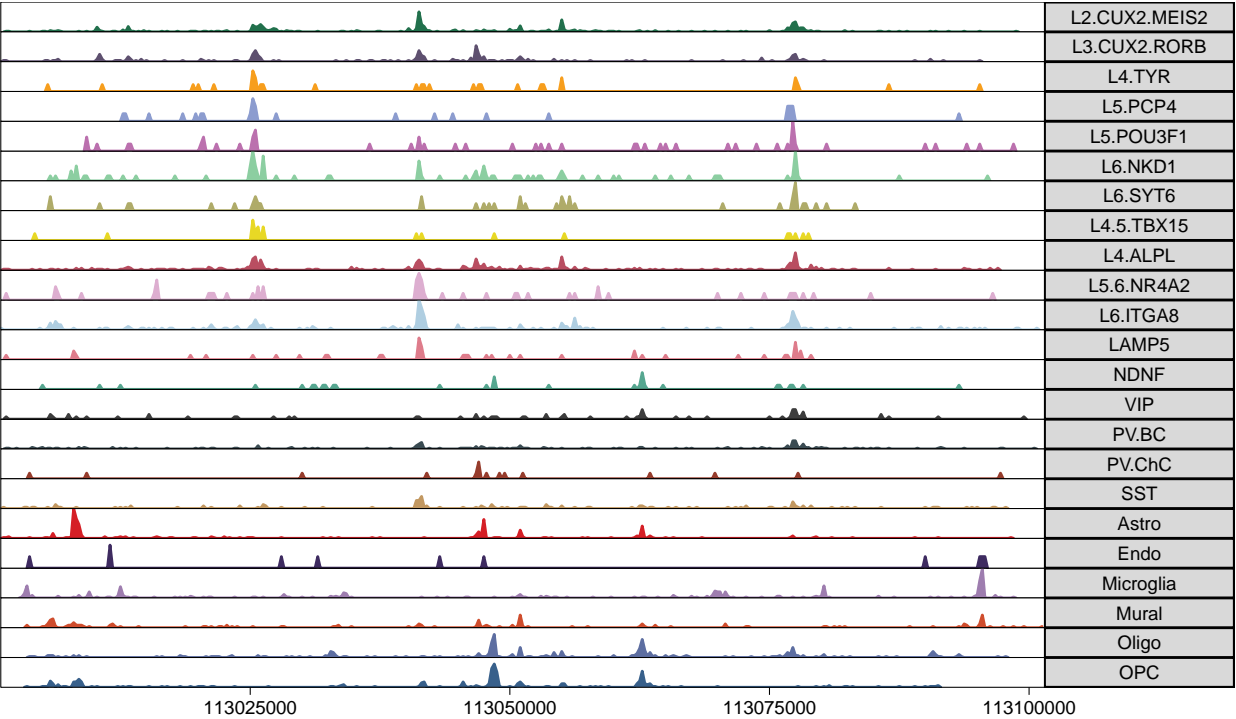


LoopTrack

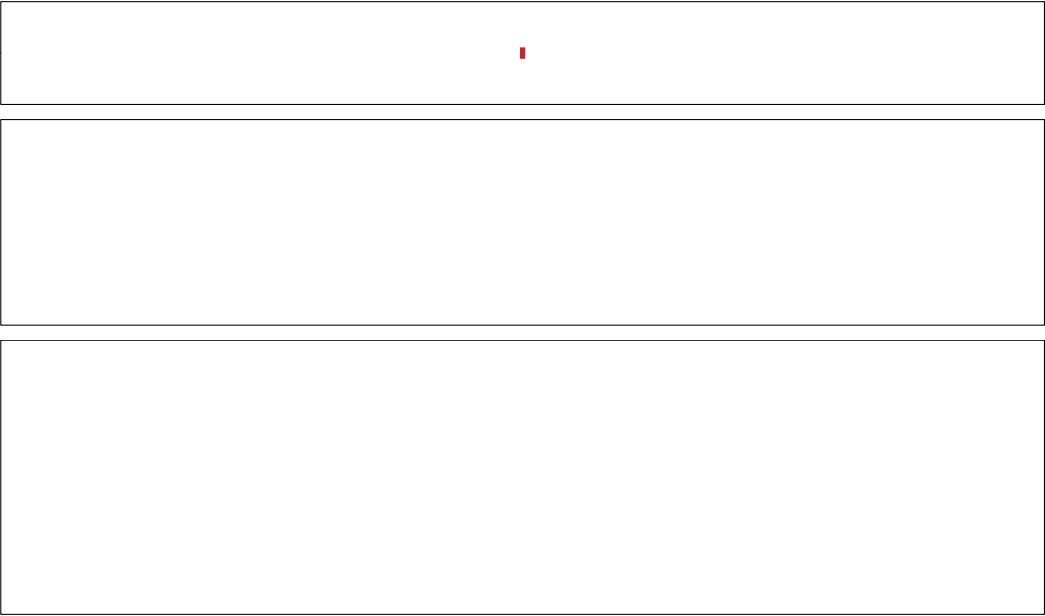


Genes

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



L5.6.NR4A2

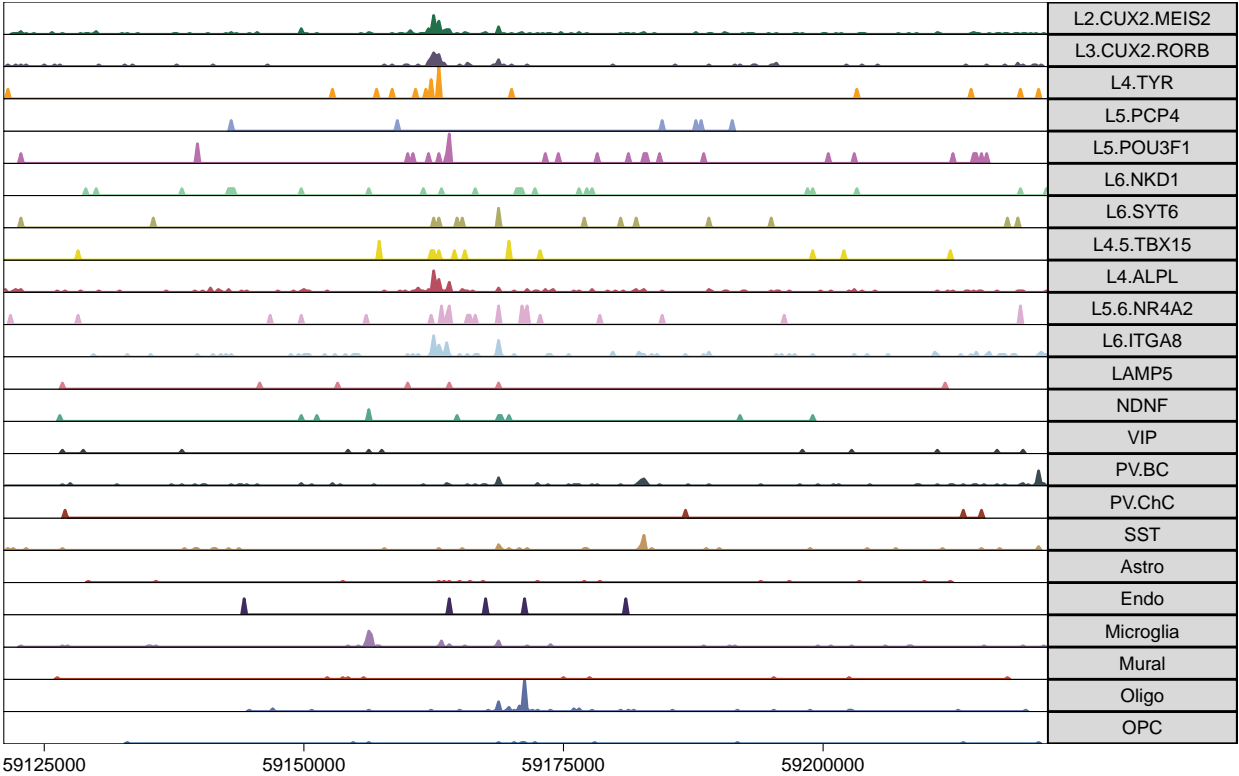


Peaks

LoopTrack

GeneTrack

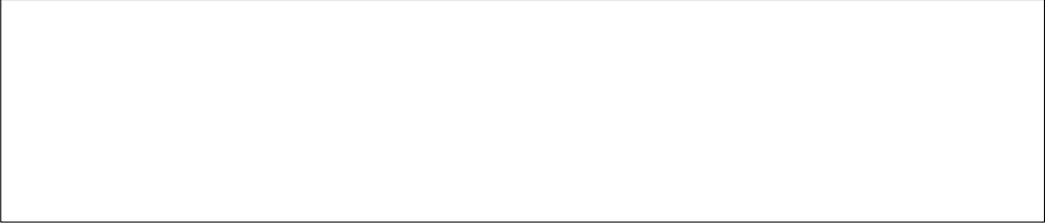
Coverage
(Norm. ATAC Signal Range 0–0.78) by ReadsInTSS)



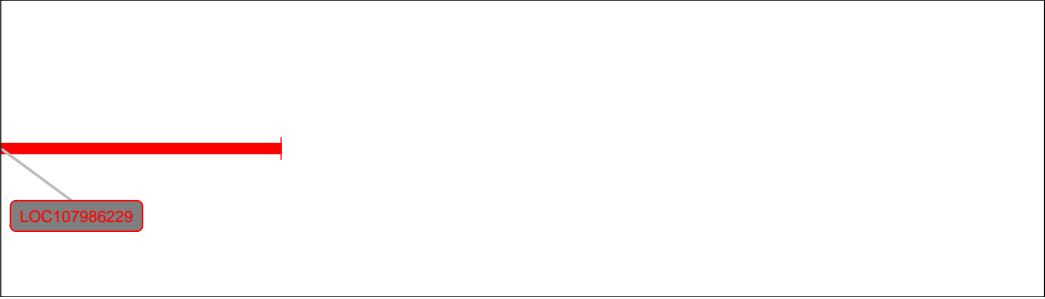
L5.6.NR4A2



Peaks

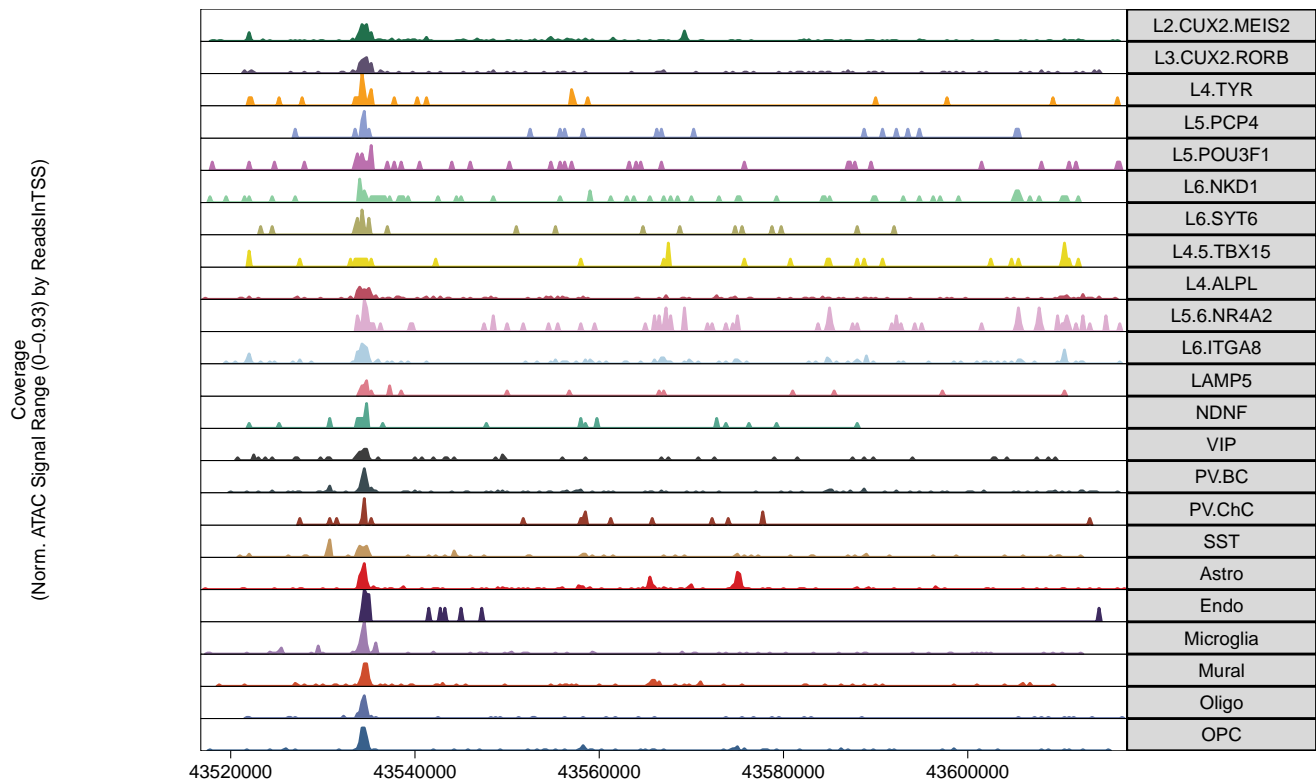


LoopTrack



Genes

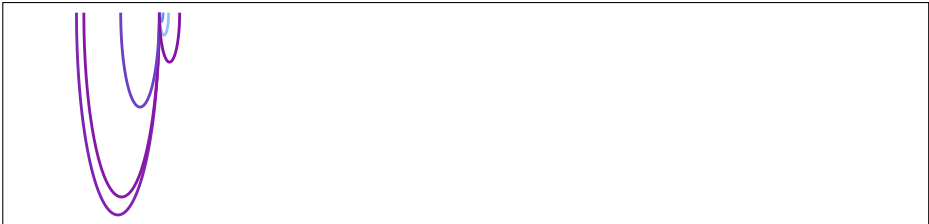
chr12:43516752-43617253 Marker Peak Tracks for: L5.6.NR4A2



L5.6.NR4A2

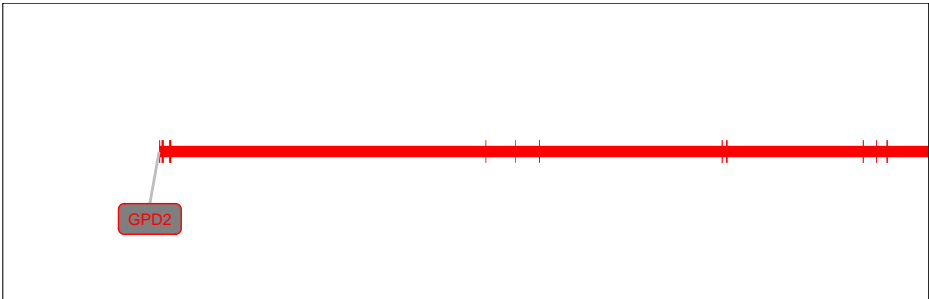
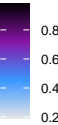


Peaks



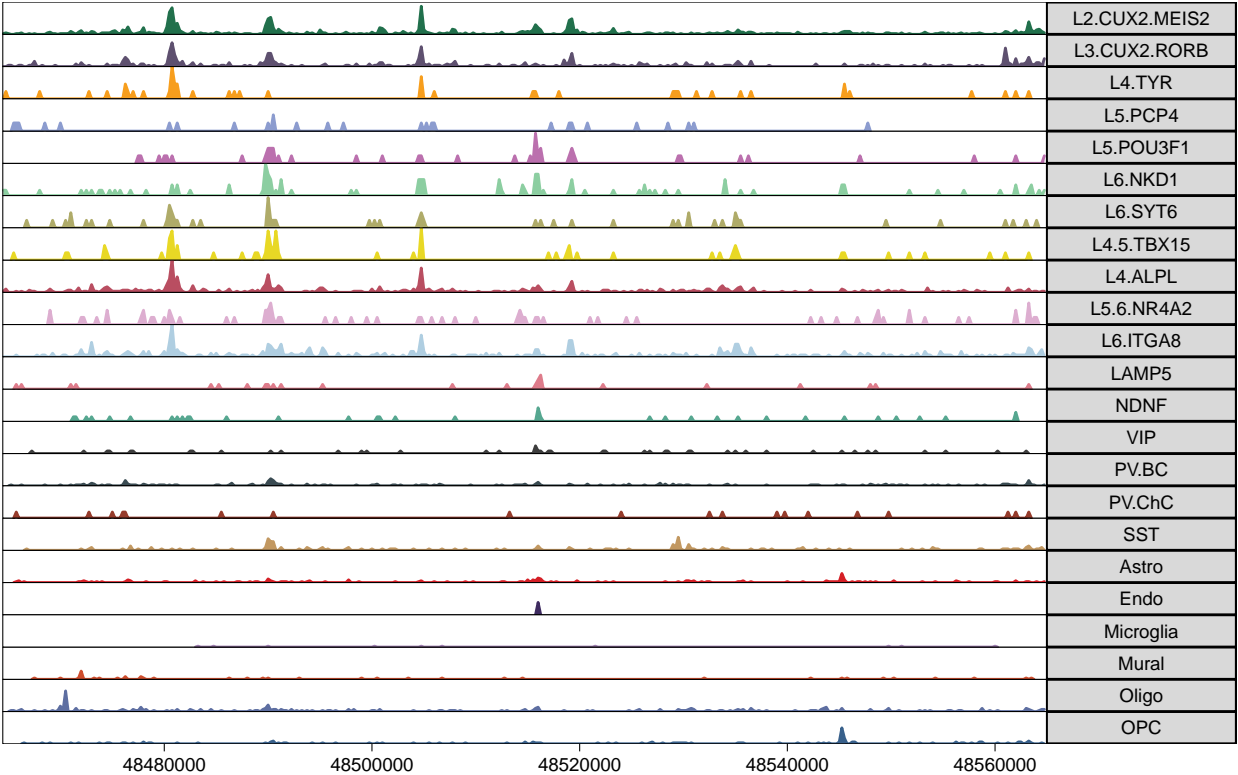
Peak2GeneLinks

value



Genes

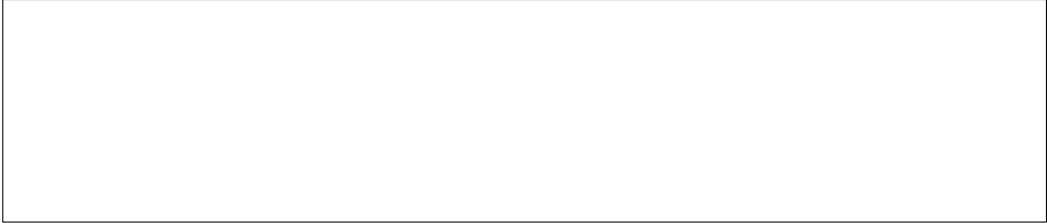
Coverage
(Norm. ATAC Signal Range 0-1.02) by ReadsInTSS)



L5.6.NR4A2



Peaks



LoopTrack



Genes