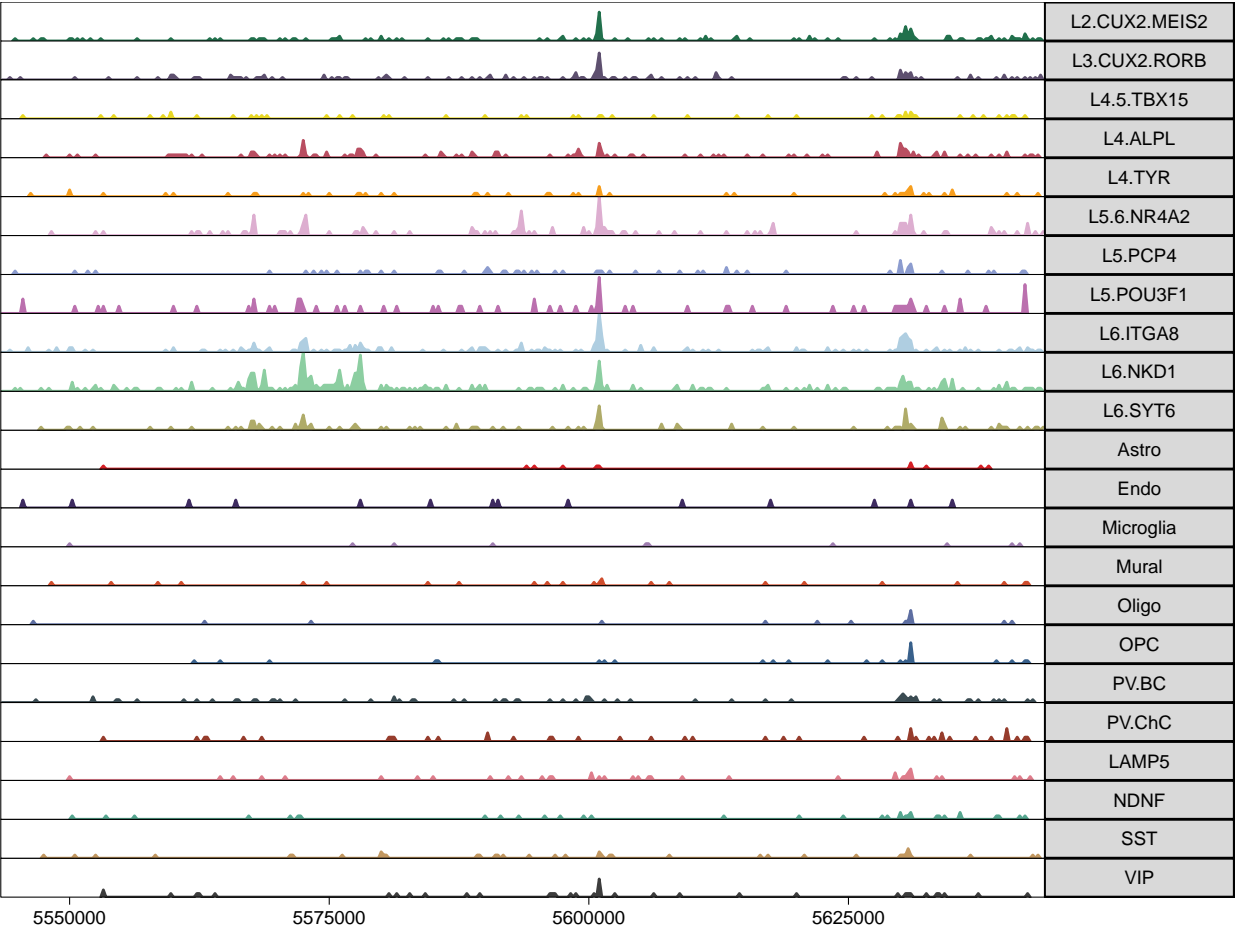
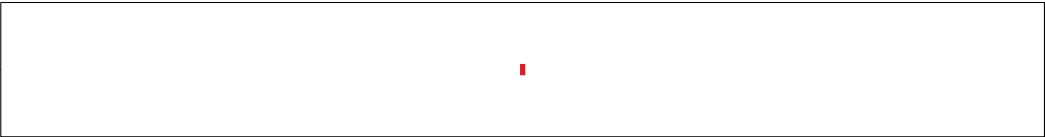


Coverage
(Norm. ATAC Signal Range (0–0.4) by ReadsInTSS)



L5.6.NR4A2



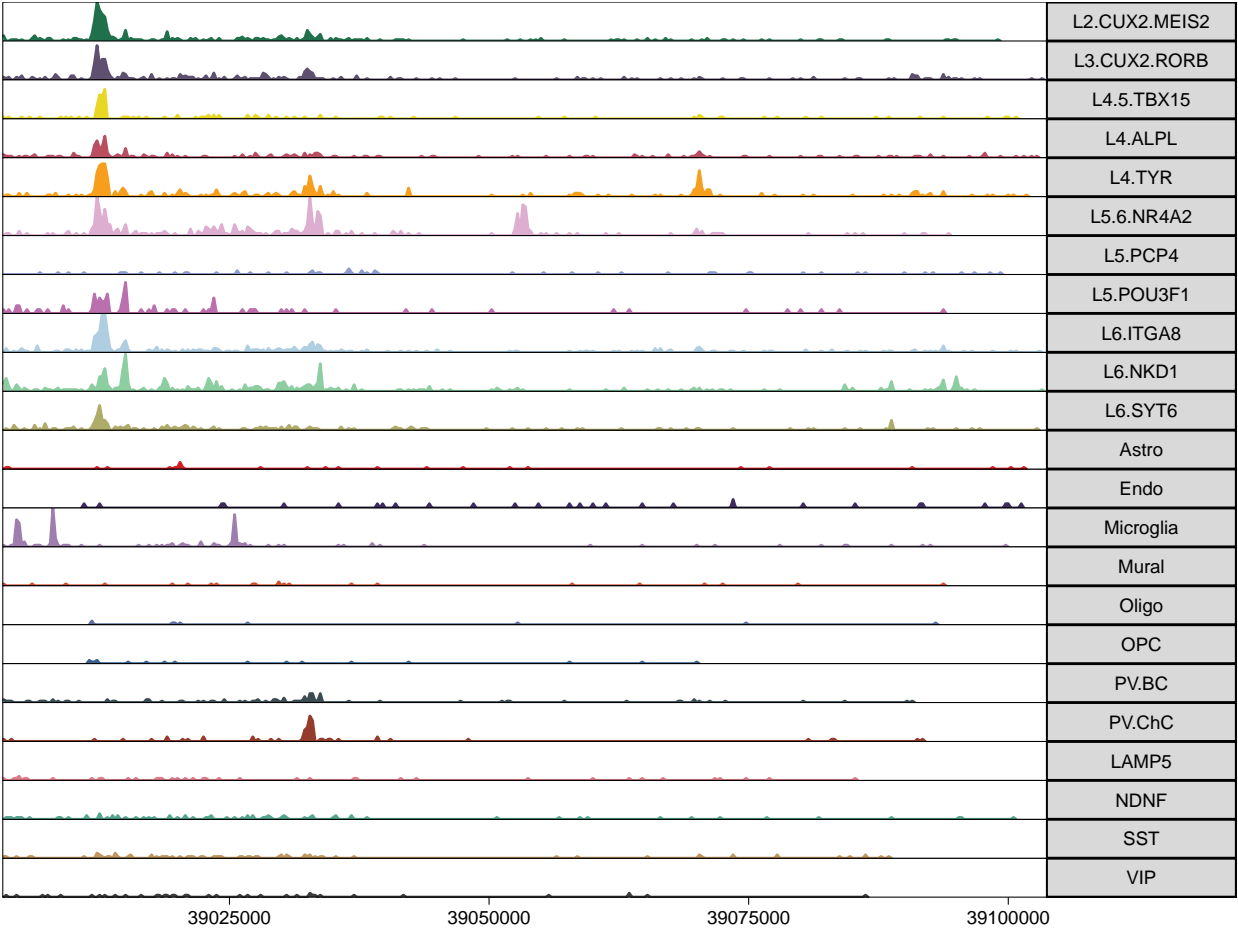
Peaks



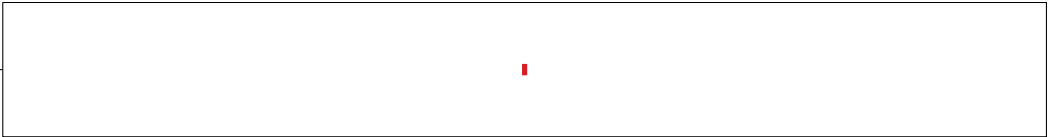
Genes

chr1:39003174–39103675 Marker Peak Tracks for: L5.6.NR4A2

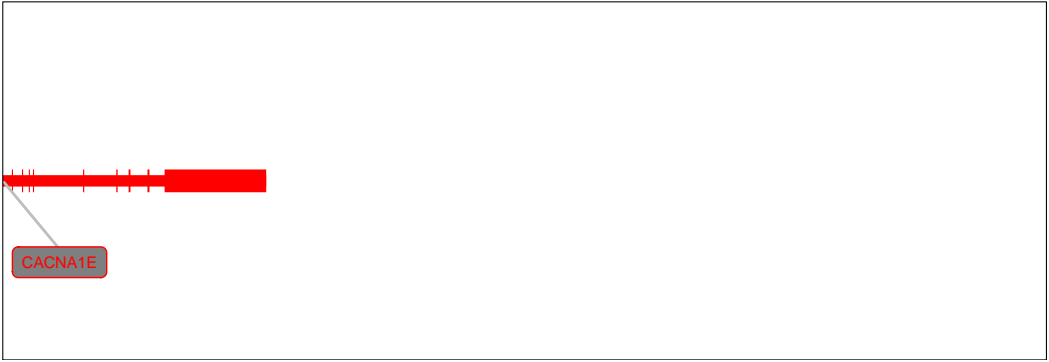
Coverage
(Norm. ATAC Signal Range (0–0.74) by ReadsInTSS)



L5.6.NR4A2



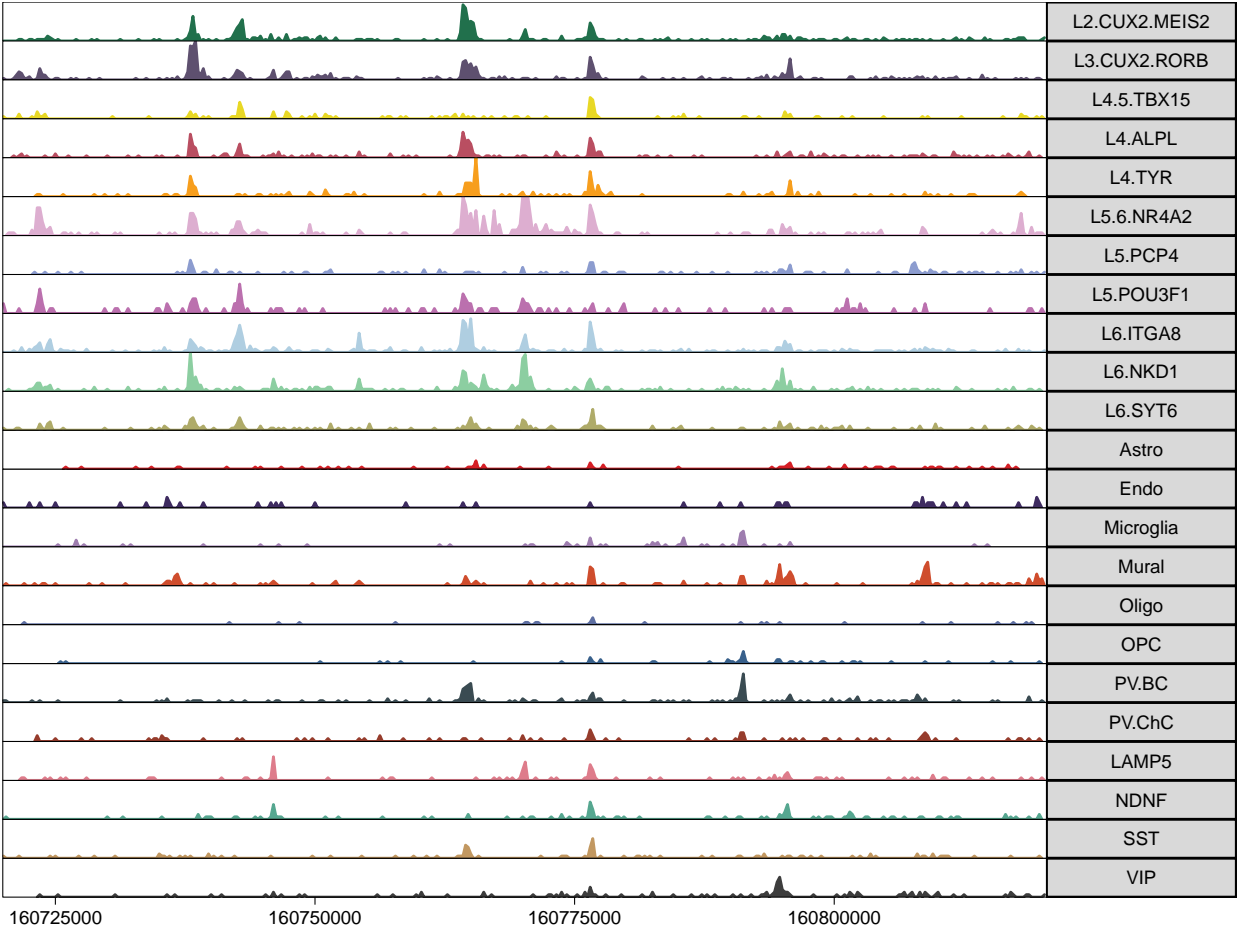
Peaks



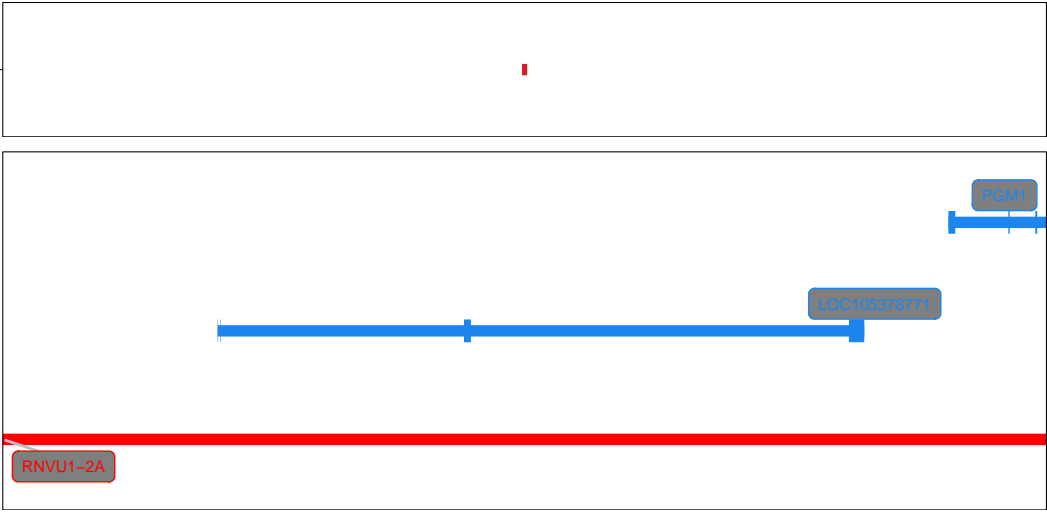
Genes

chr1:160719936–160820437 Marker Peak Tracks for: L5.6.NR4A2

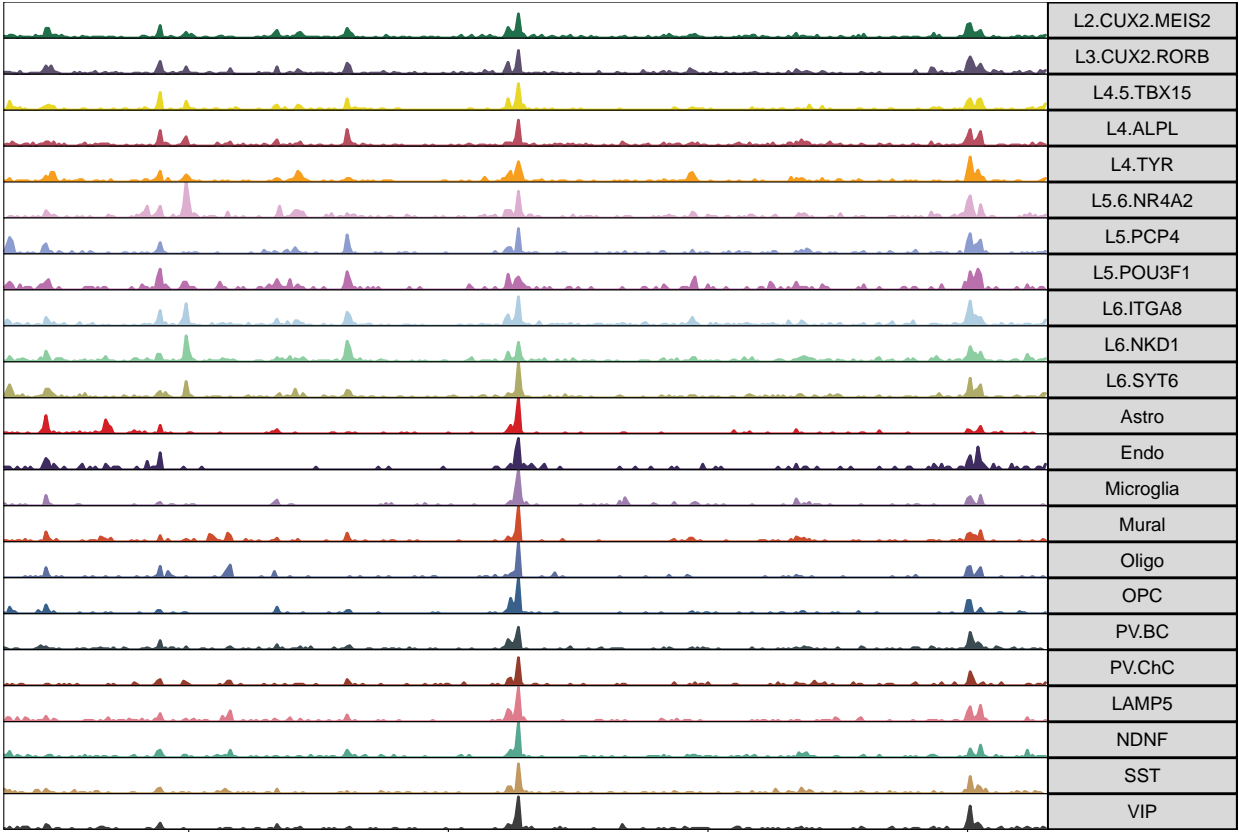
Coverage
(Norm. ATAC Signal Range (0–0.58) by ReadsInTSS)



L5.6.NR4A2



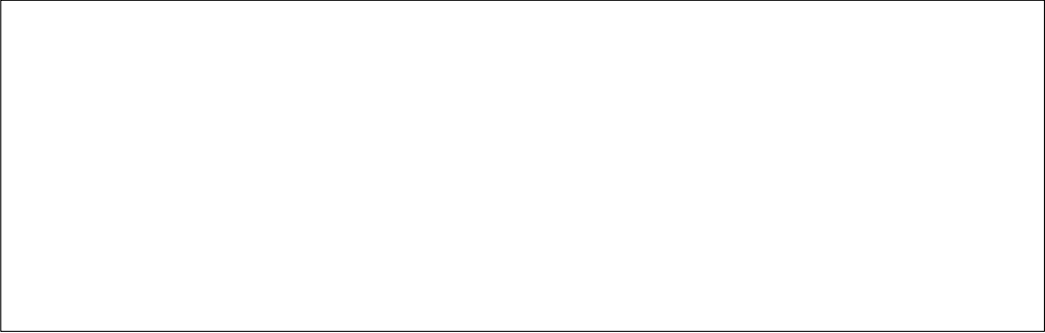
Coverage
(Norm. ATAC Signal Range (0–1.01) by ReadsInTSS)



L5.6.NR4A2



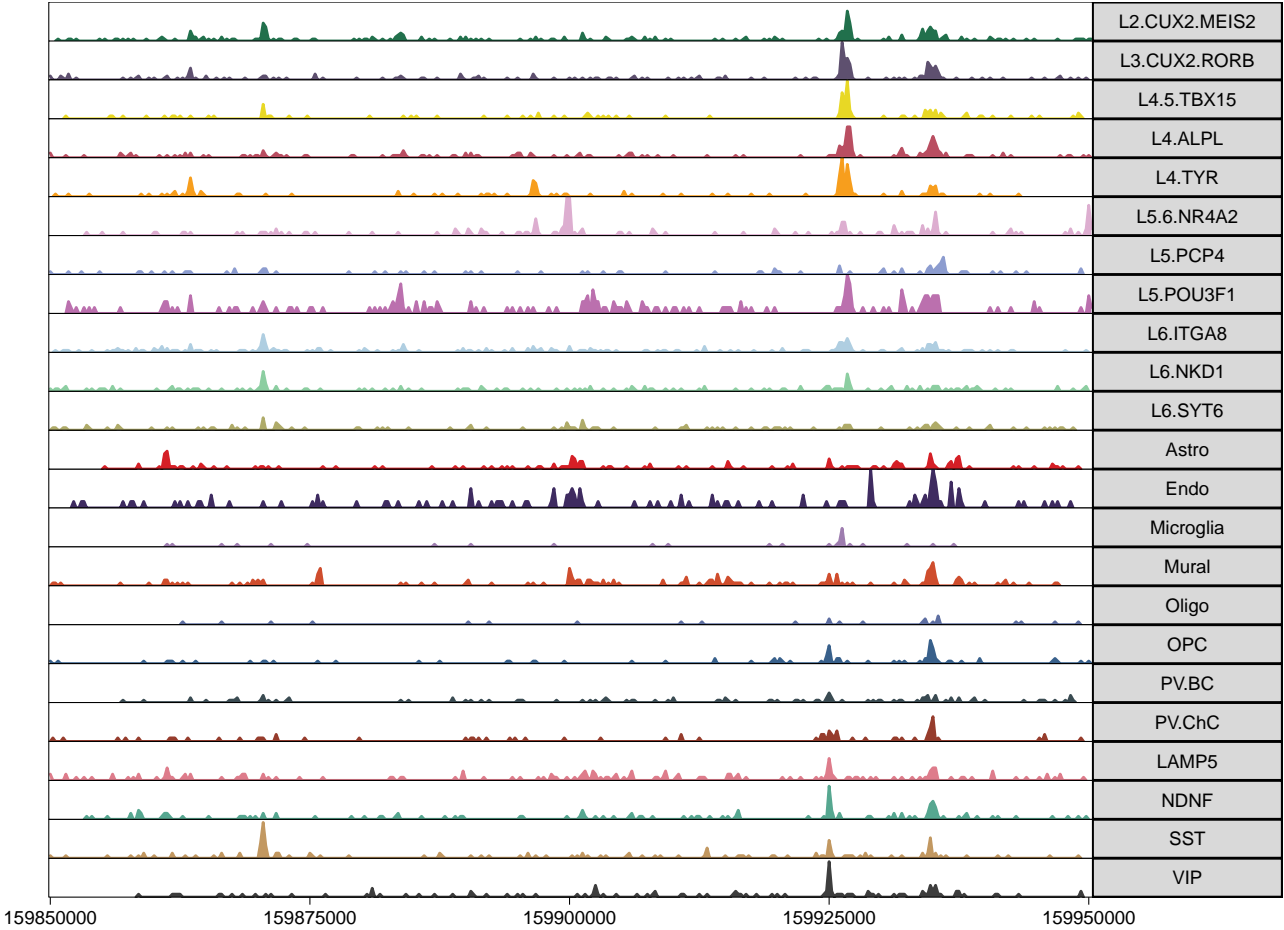
Peaks



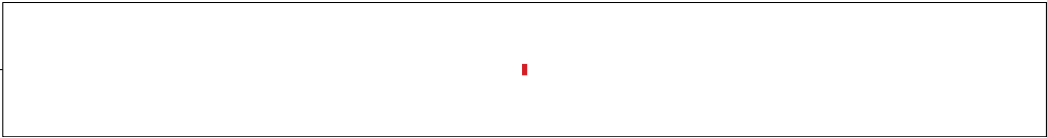
GeneTrack

chr1:159849855–159950356 Marker Peak Tracks for: L5.6.NR4A2

Coverage
(Norm. ATAC Signal Range (0–0.48) by ReadsInTSS)



L5.6.NR4A2



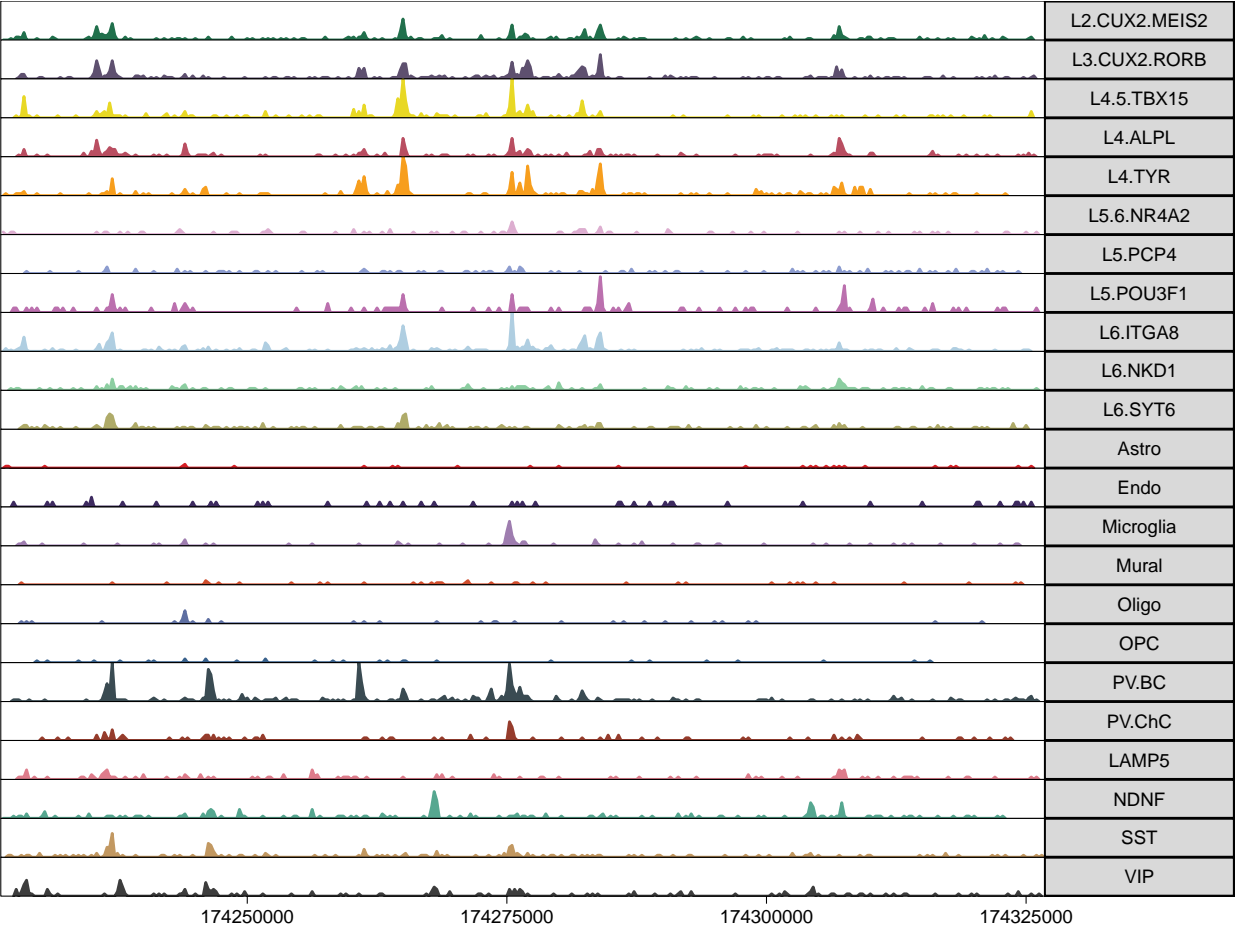
Peaks



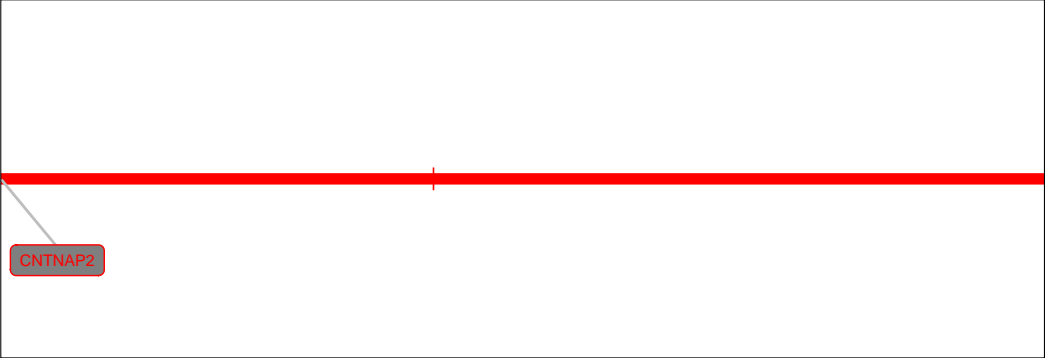
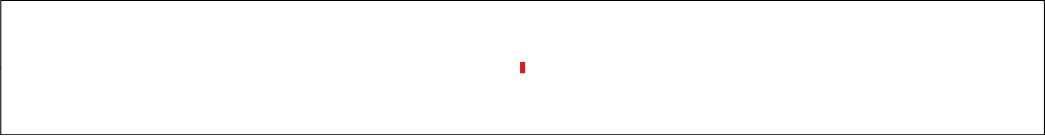
Genes

chr3:174226266–174326767 Marker Peak Tracks for: L5.6.NR4A2

Coverage
(Norm. ATAC Signal Range (0–0.64) by ReadsInTSS)

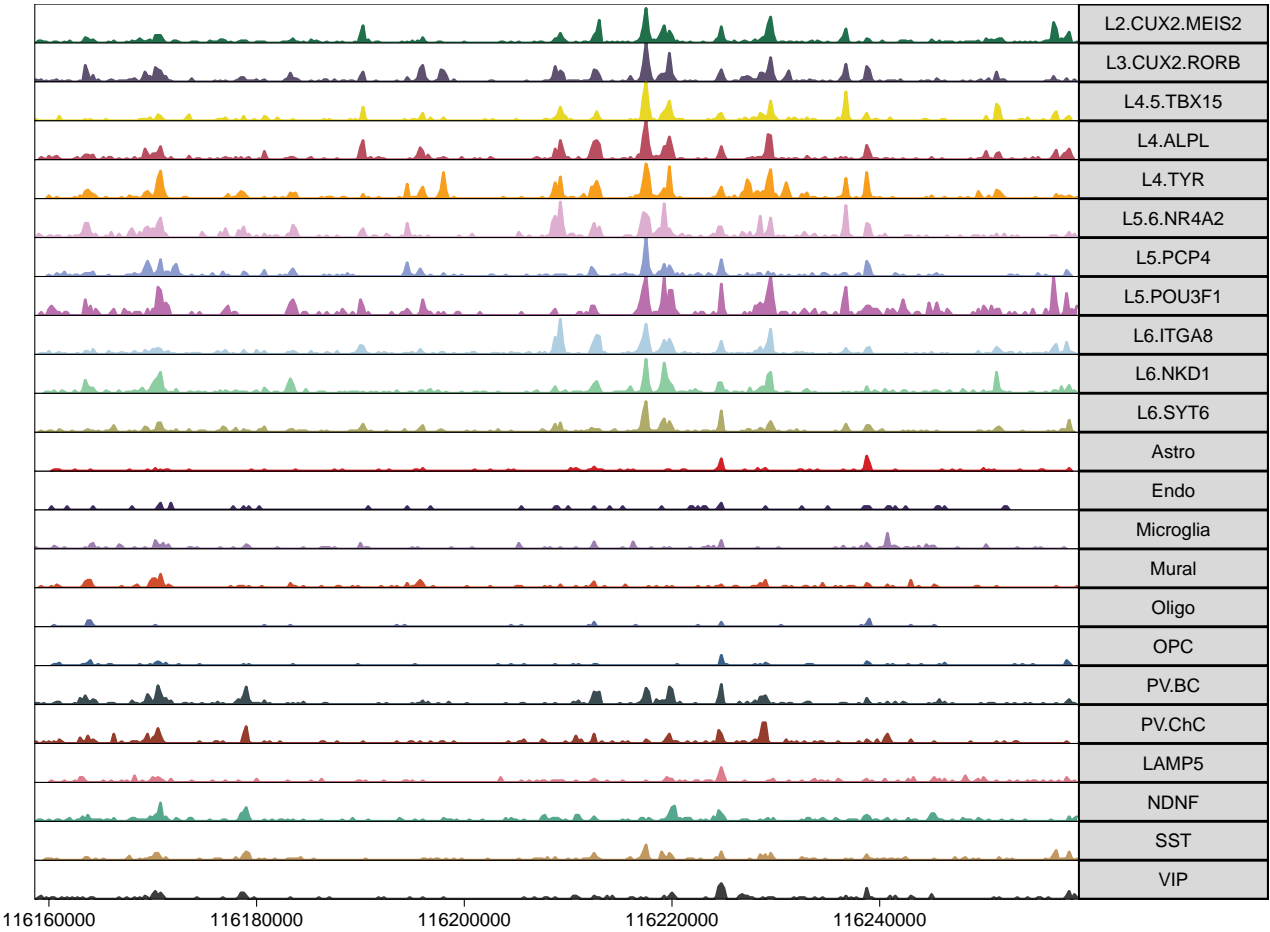


L5.6.NR4A2

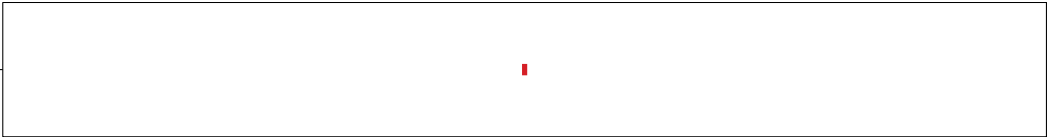


chr14:116158640–116259141 Marker Peak Tracks for: L5.6.NR4A2

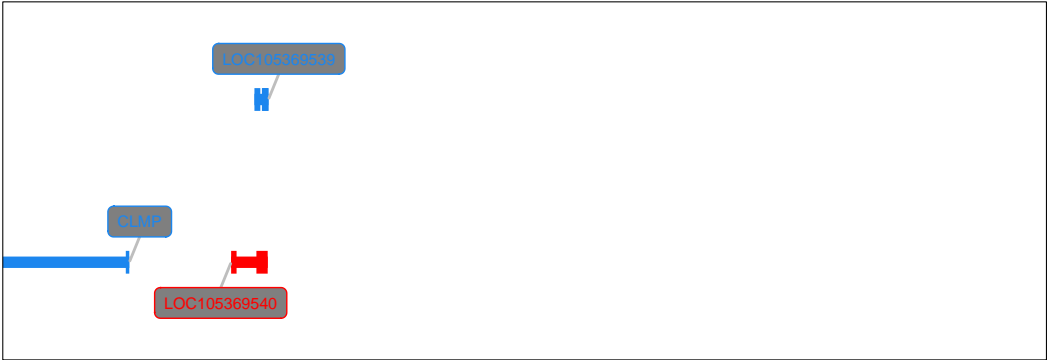
Coverage
(Norm. ATAC Signal Range (0–0.91) by ReadsInTSS)



L5.6.NR4A2

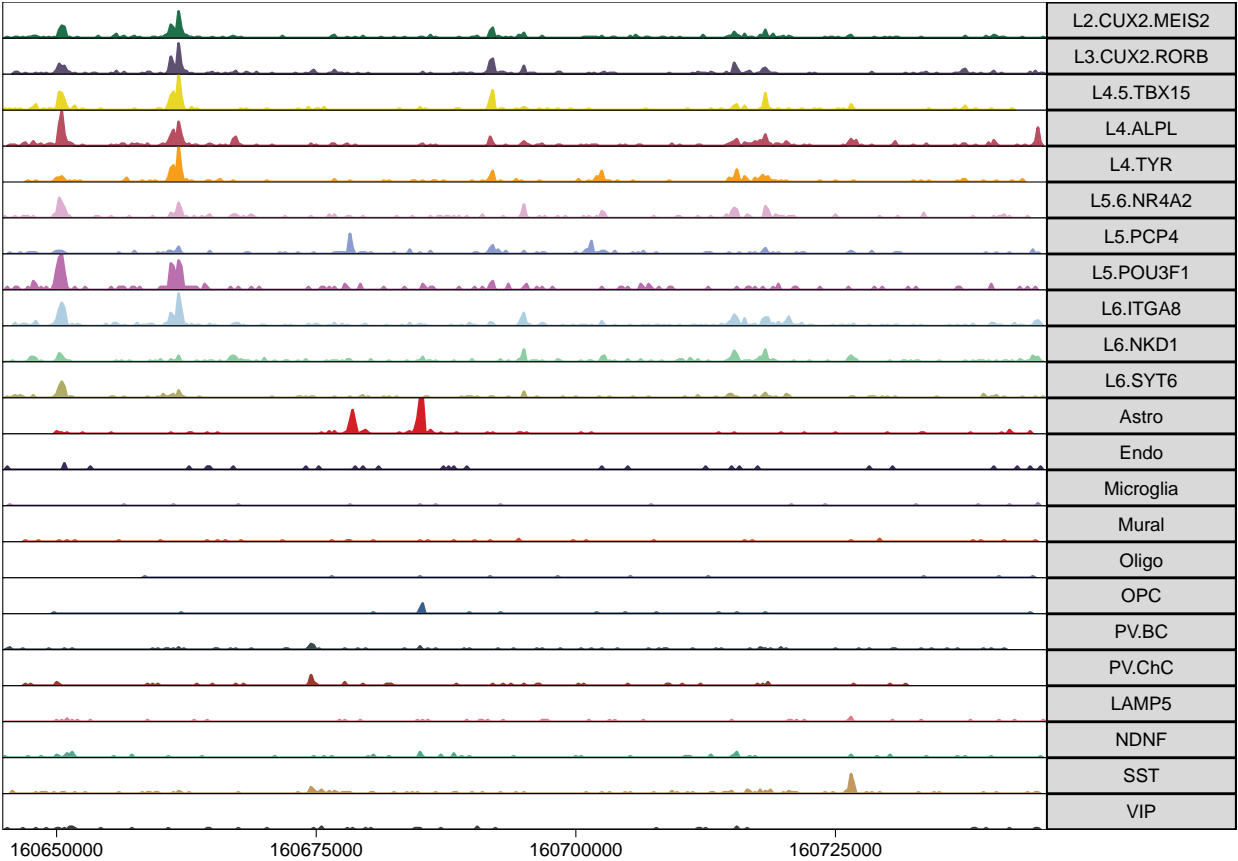


Peaks



Genes

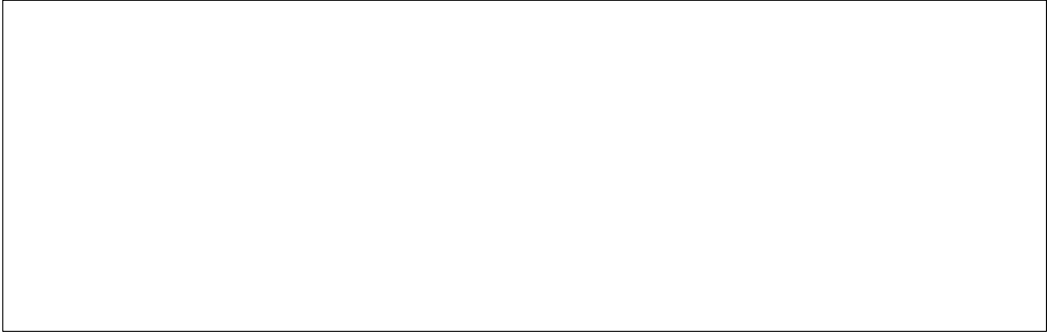
Coverage
(Norm. ATAC Signal Range (0-0.9) by ReadsInTSS)



L5.6.NR4A2



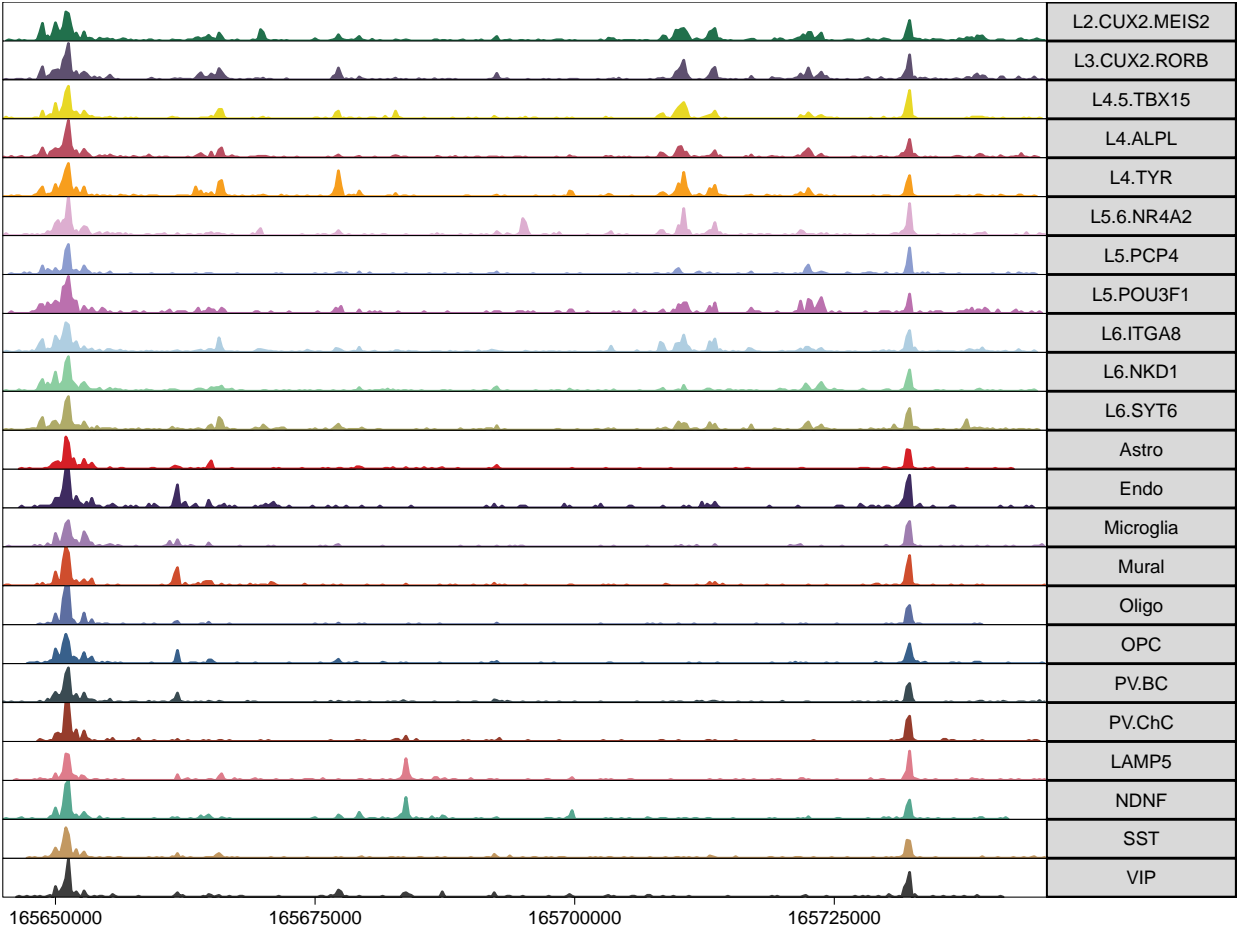
Peaks



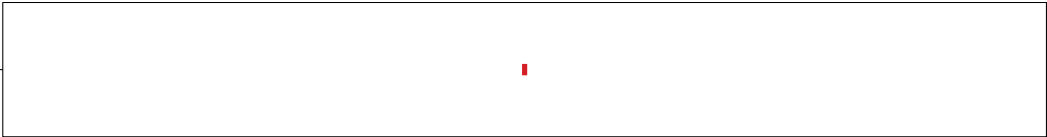
GeneTrack

chr1:165644927-165745428 Marker Peak Tracks for: L5.6.NR4A2

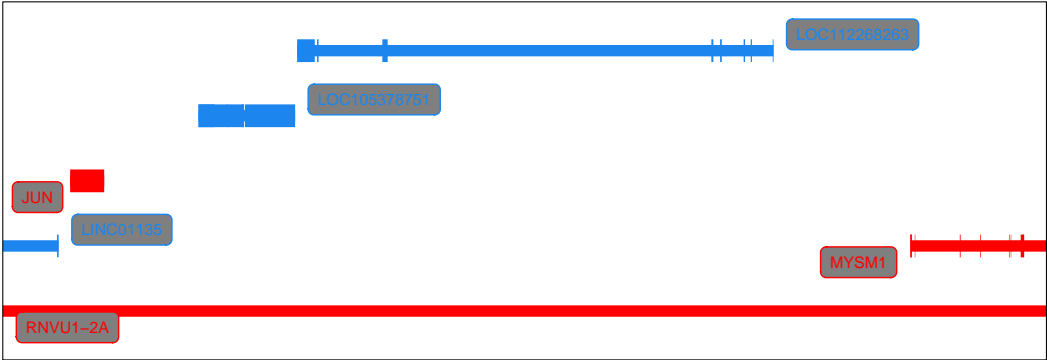
Coverage
(Norm. ATAC Signal Range (0-1.62) by ReadsInTSS)



L5.6.NR4A2

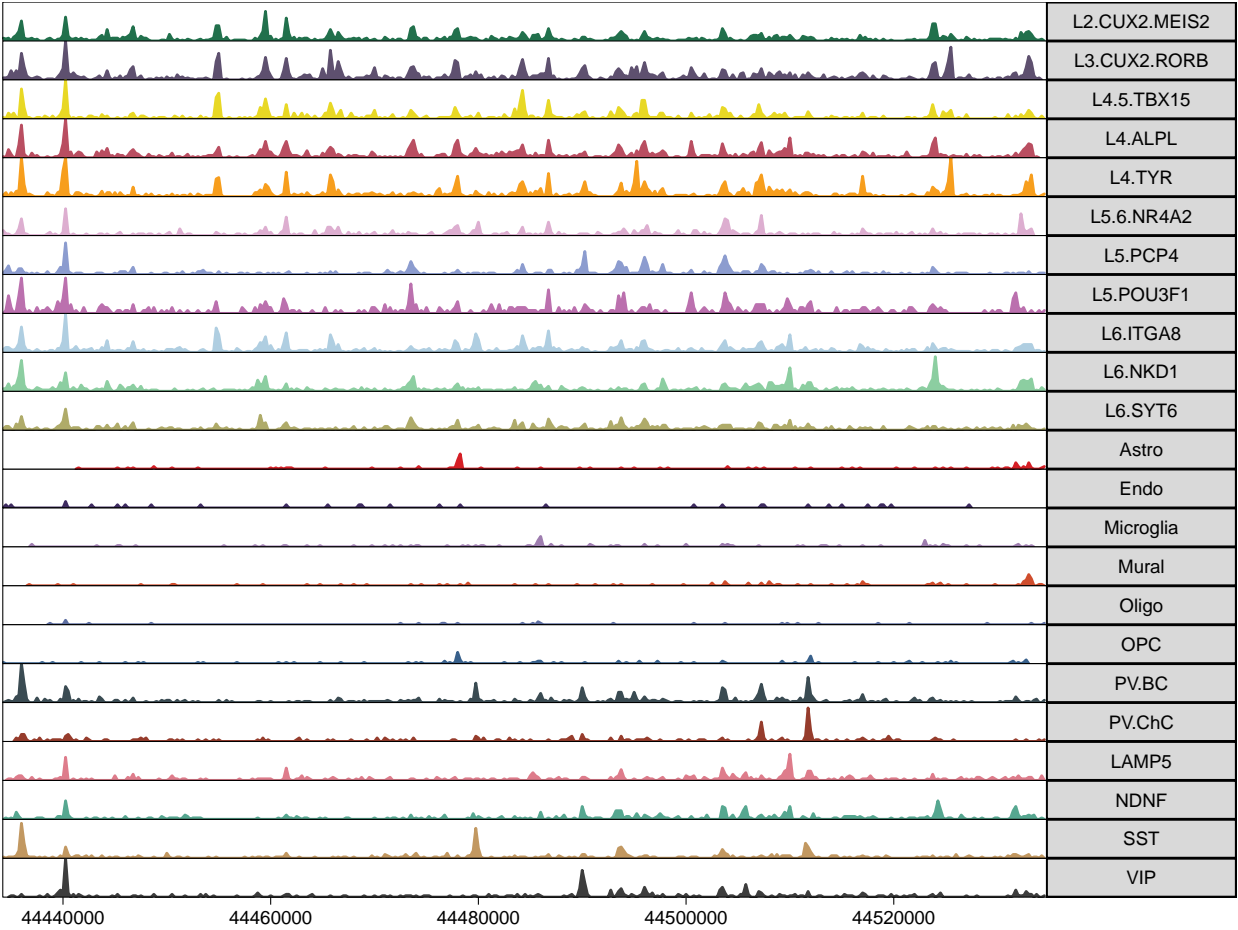


Peaks

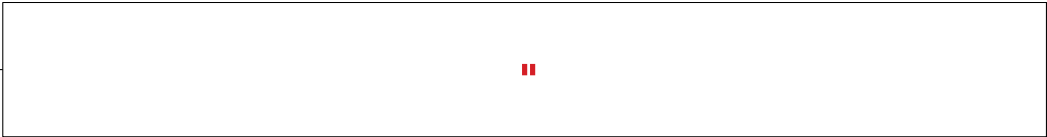


Genes

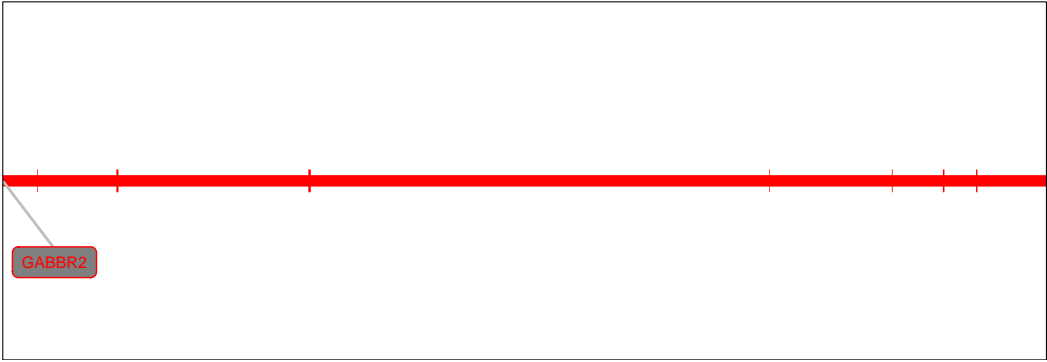
Coverage
(Norm. ATAC Signal Range (0–0.97) by ReadsInTSS)



L5.6.NR4A2

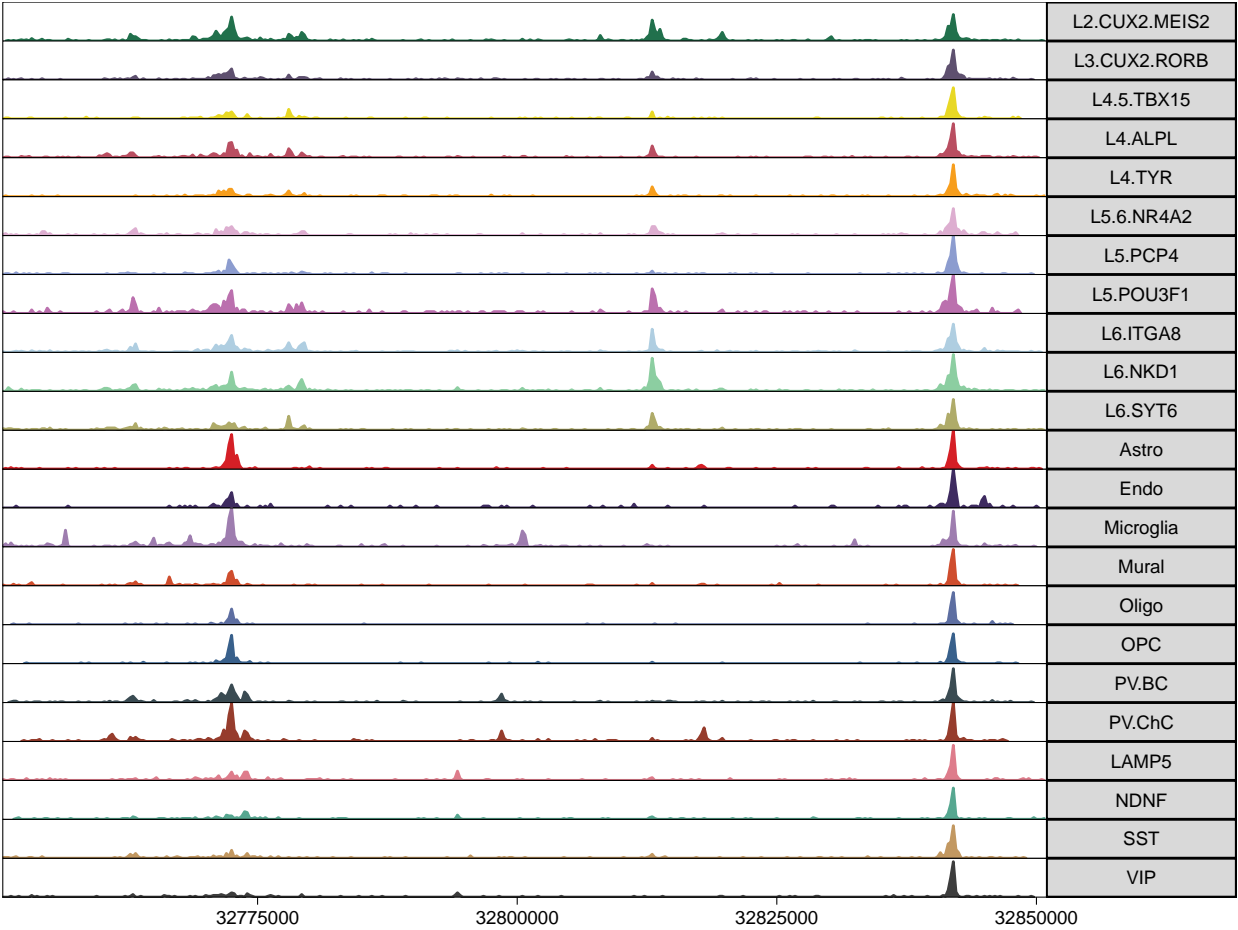


Peaks

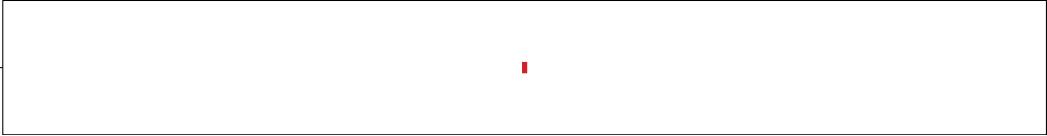


Genes

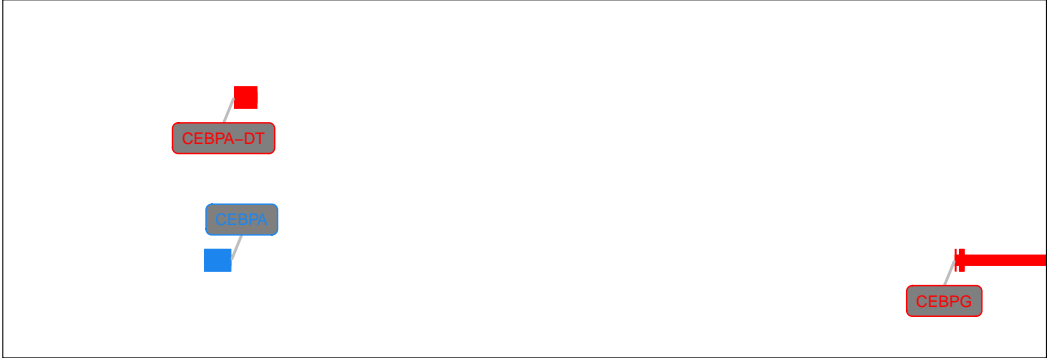
Coverage
(Norm. ATAC Signal Range (0–1.63) by ReadsInTSS)



L5.6.NR4A2

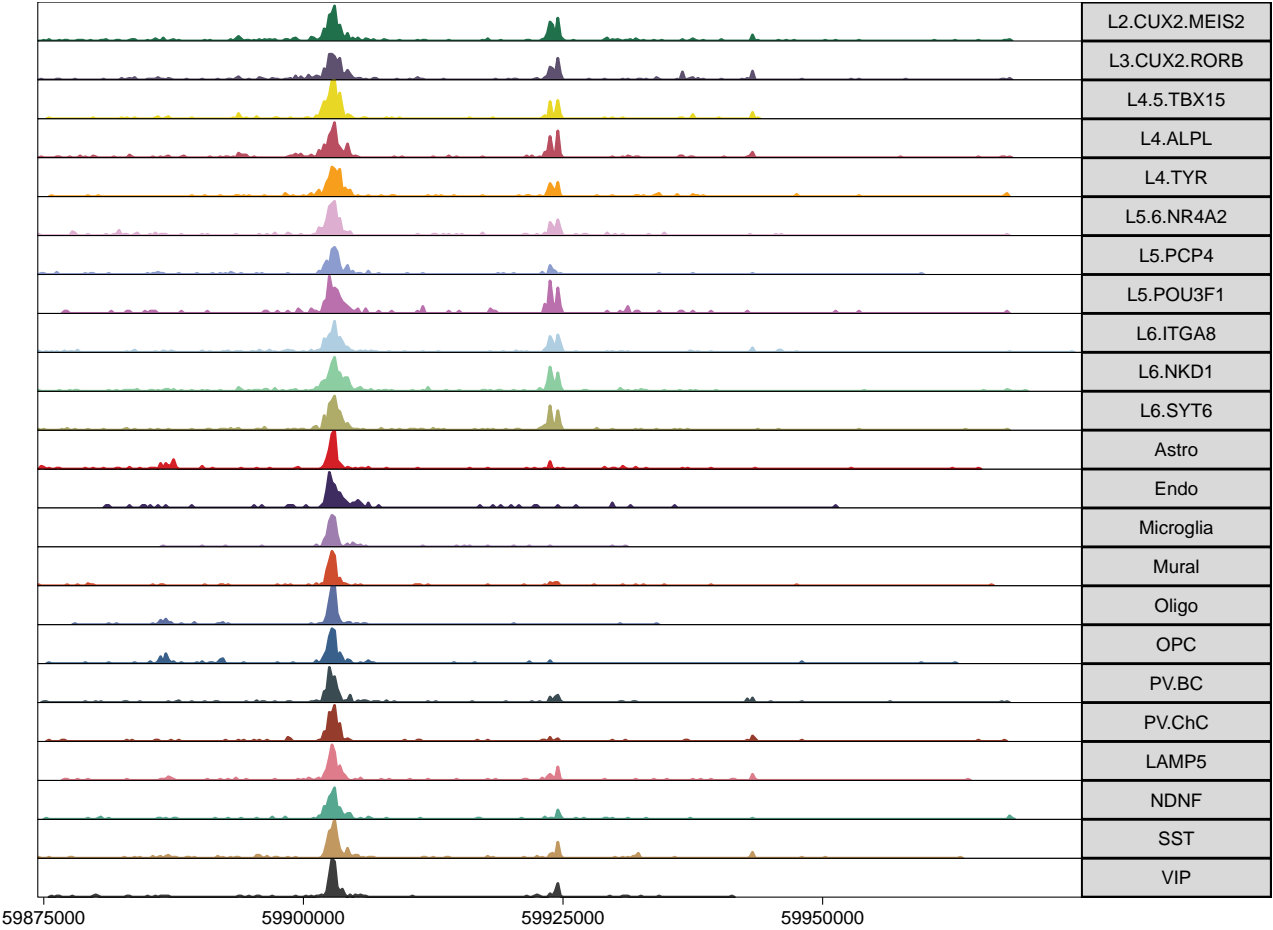


Peaks

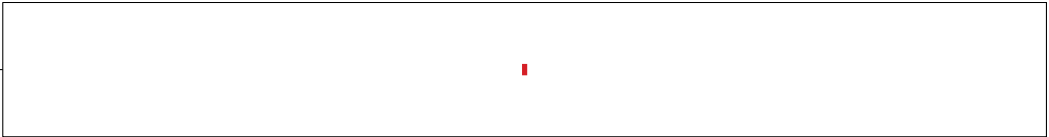


Genes

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



L5.6.NR4A2

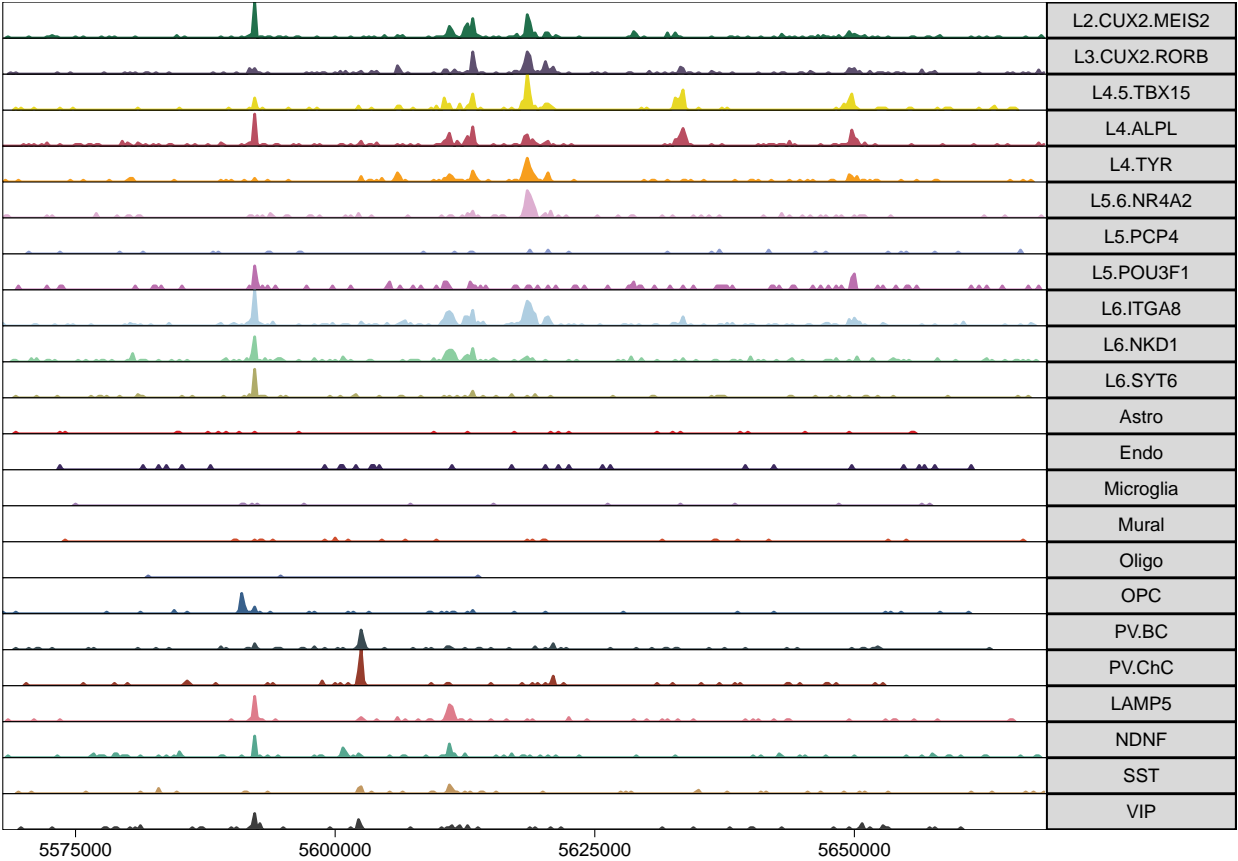


Peaks



Genes

Coverage
(Norm. ATAC Signal Range (0–0.66) by ReadsInTSS)



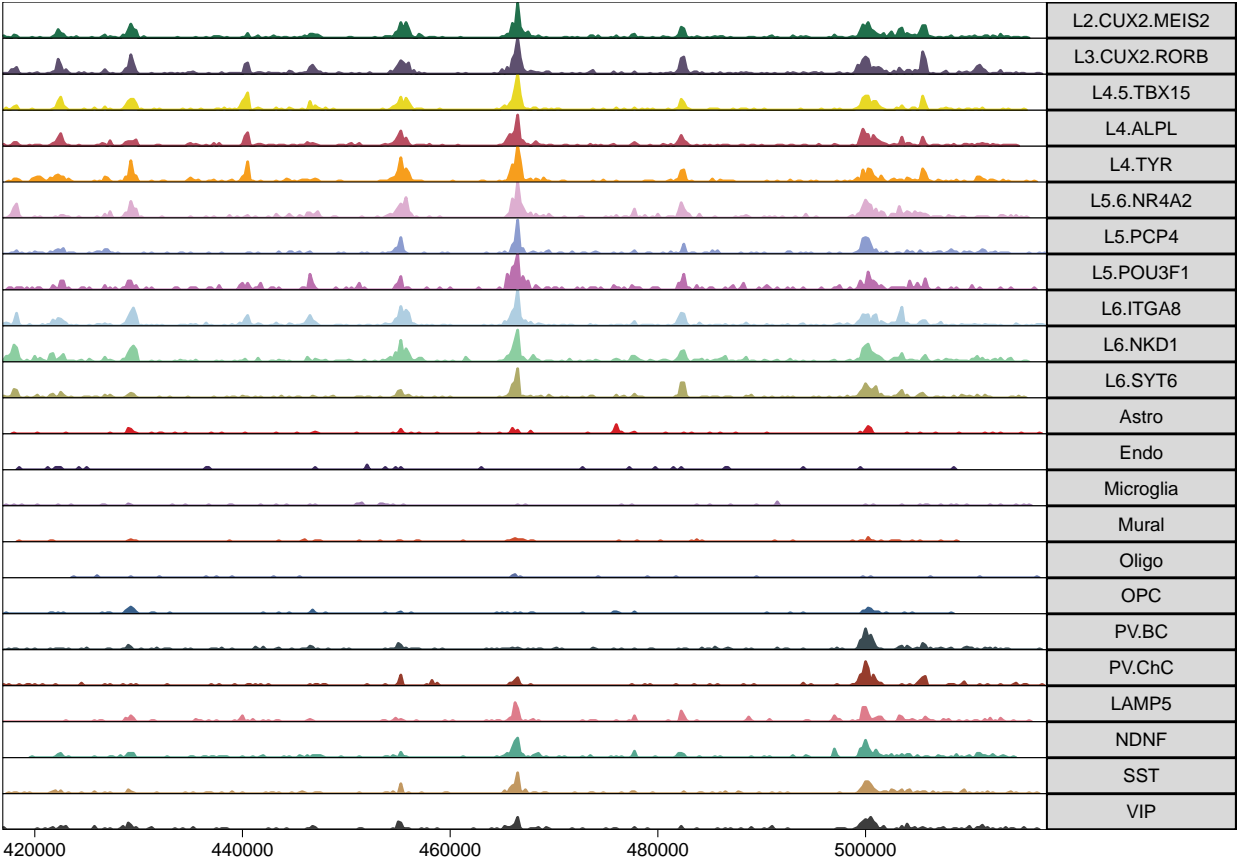
L5.6.NR4A2



Peaks

GeneTrack

Coverage
(Norm. ATAC Signal Range (0–1,18) by ReadsInTSS)



L5.6.NR4A2

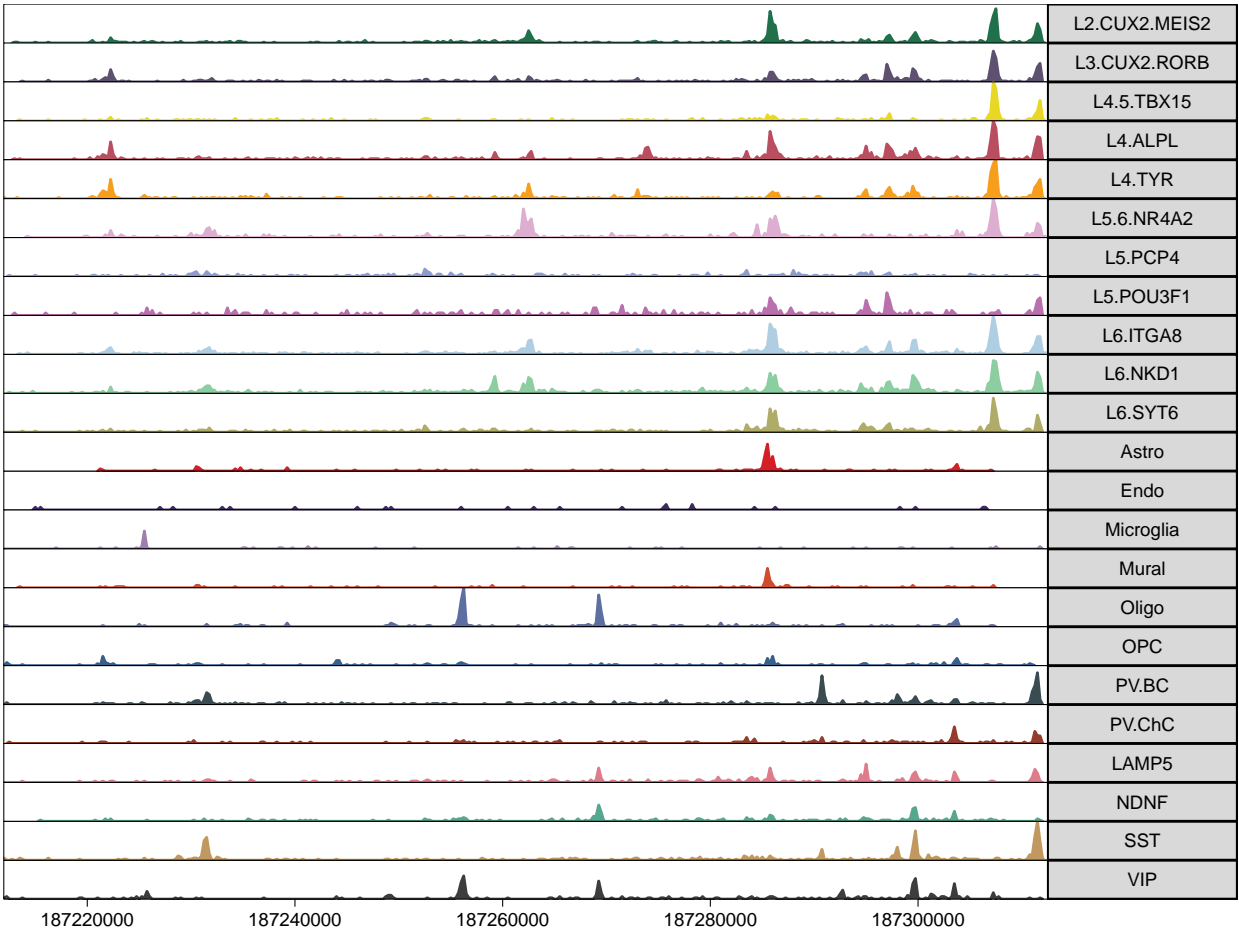


Peaks

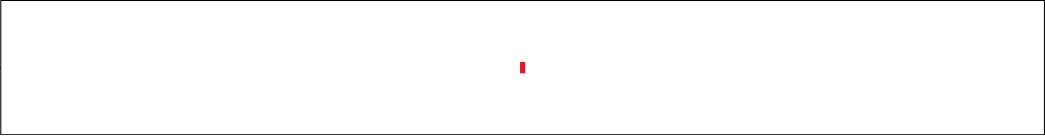
GeneTrack

chr1:187211961–187312462 Marker Peak Tracks for: L5.6.NR4A2

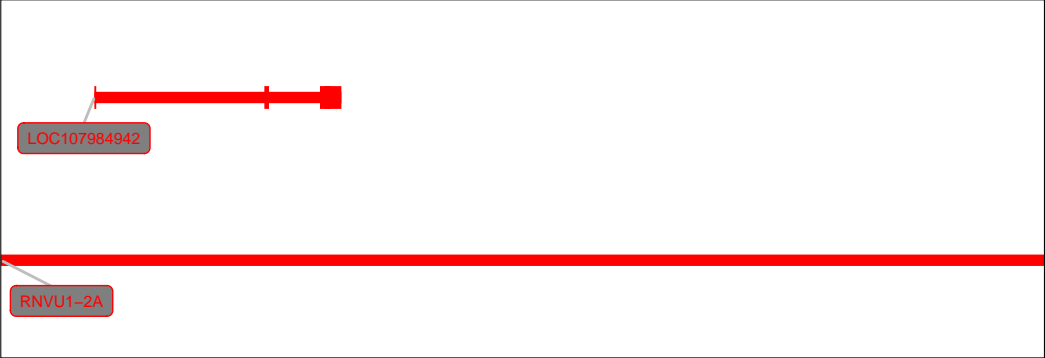
Coverage
(Norm. ATAC Signal Range (0–1.13) by ReadsInTSS)



L5.6.NR4A2

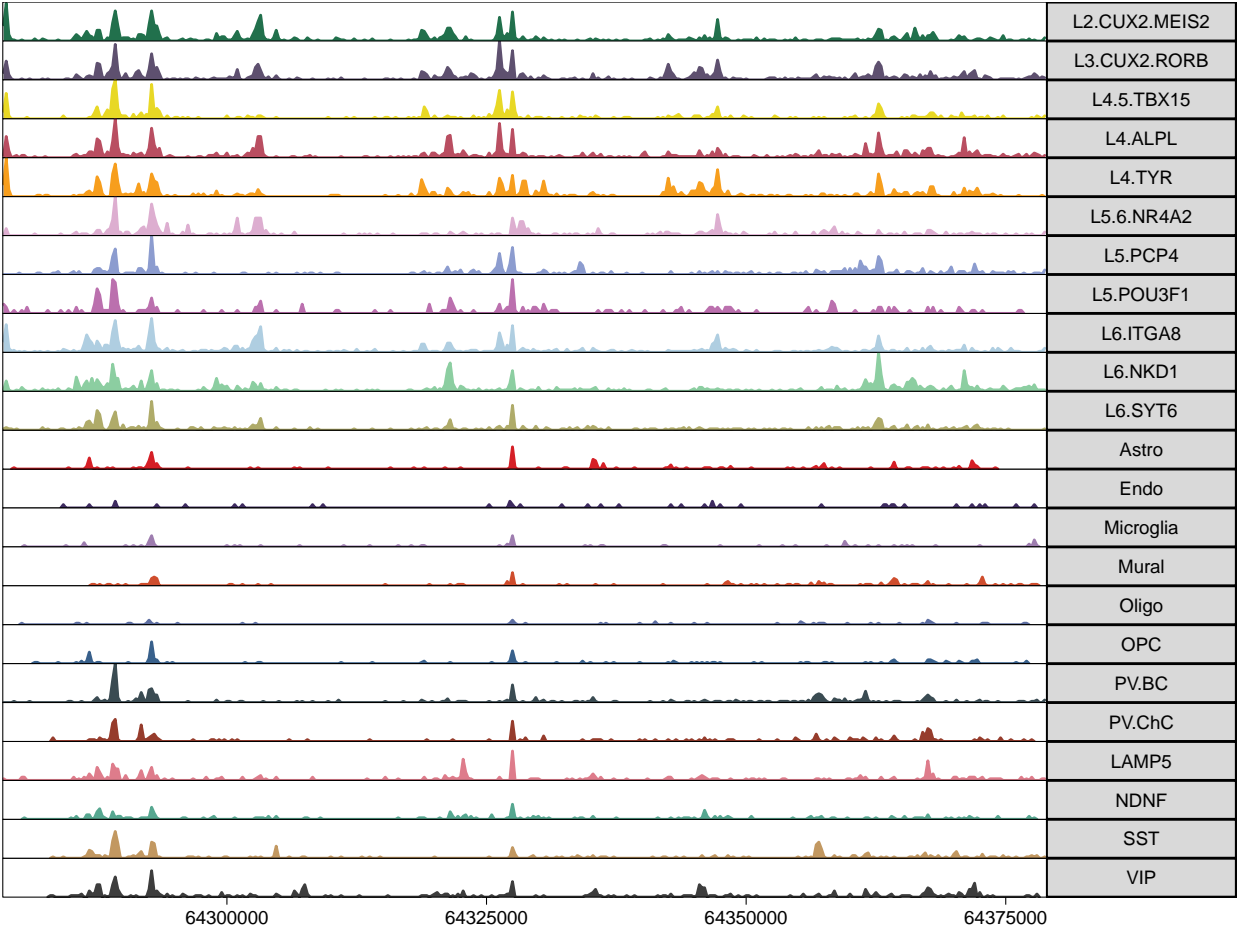


Peaks

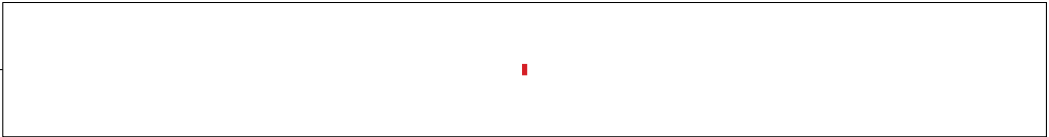


Genes

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



L5.6.NR4A2



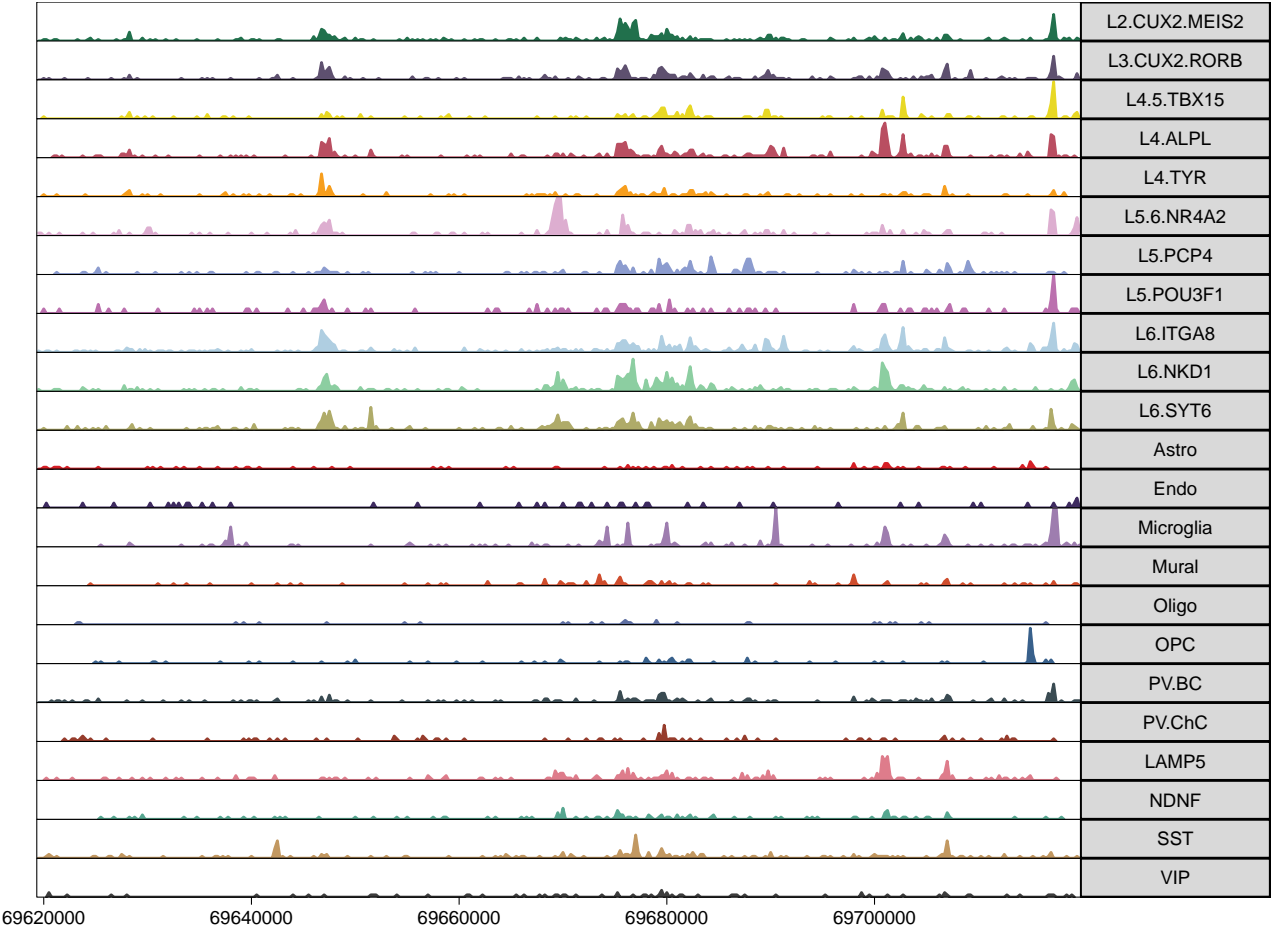
Peaks



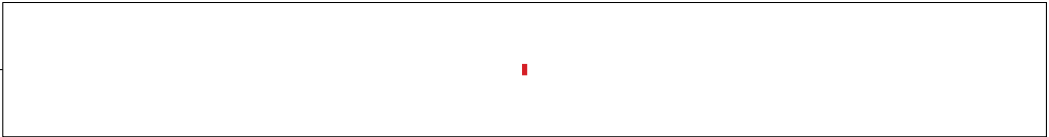
Genes

chr7:69619329–69719830 Marker Peak Tracks for: L5.6.NR4A2

Coverage
(Norm. ATAC Signal Range (0–0.63) by ReadsInTSS)



L5.6.NR4A2



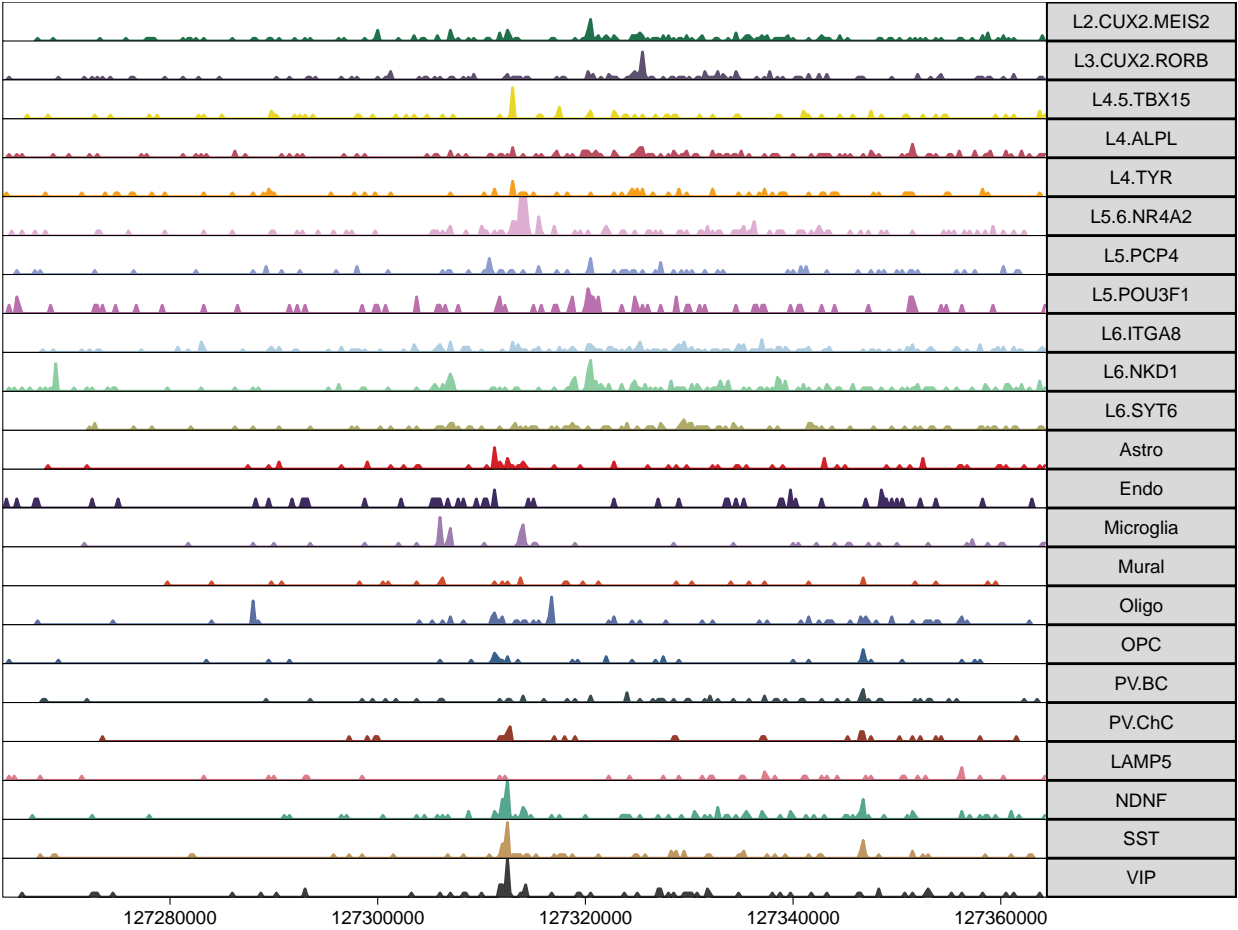
Peaks



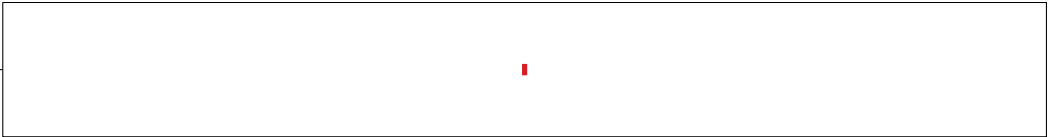
Genes

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

Coverage
(Norm. ATAC Signal Range (0–0.35) by ReadsInTSS)



L5.6.NR4A2



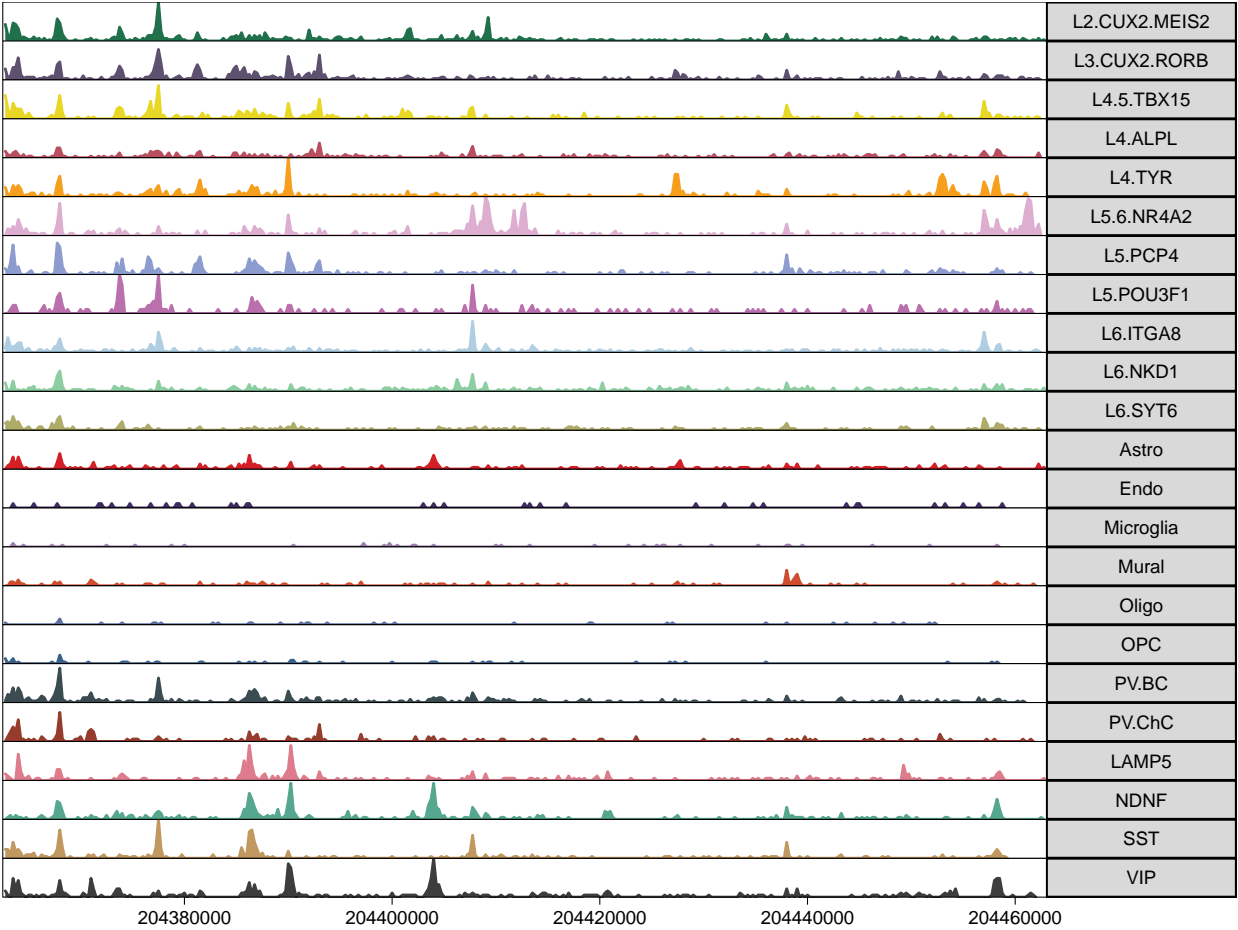
Peaks



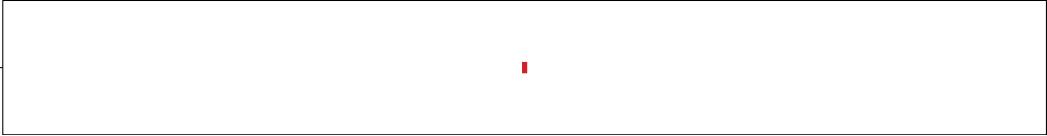
Genes

chr1:204362509-204463010 Marker Peak Tracks for: L5.6.NR4A2

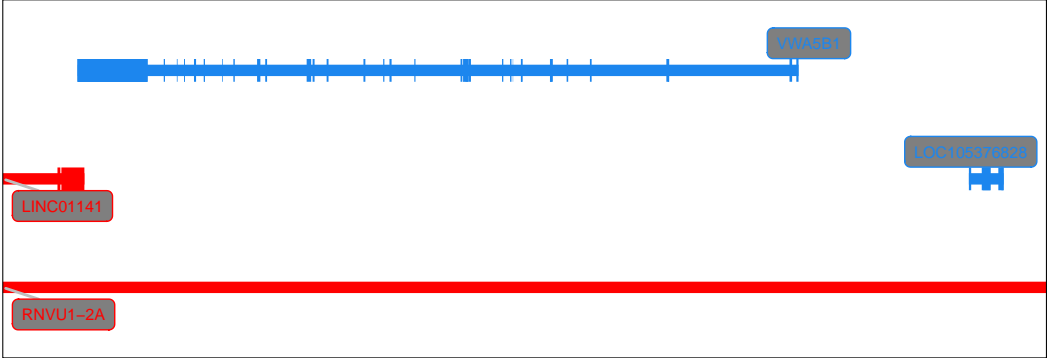
Coverage
(Norm. ATAC Signal Range (0-0.71) by ReadsInTSS)



L5.6.NR4A2

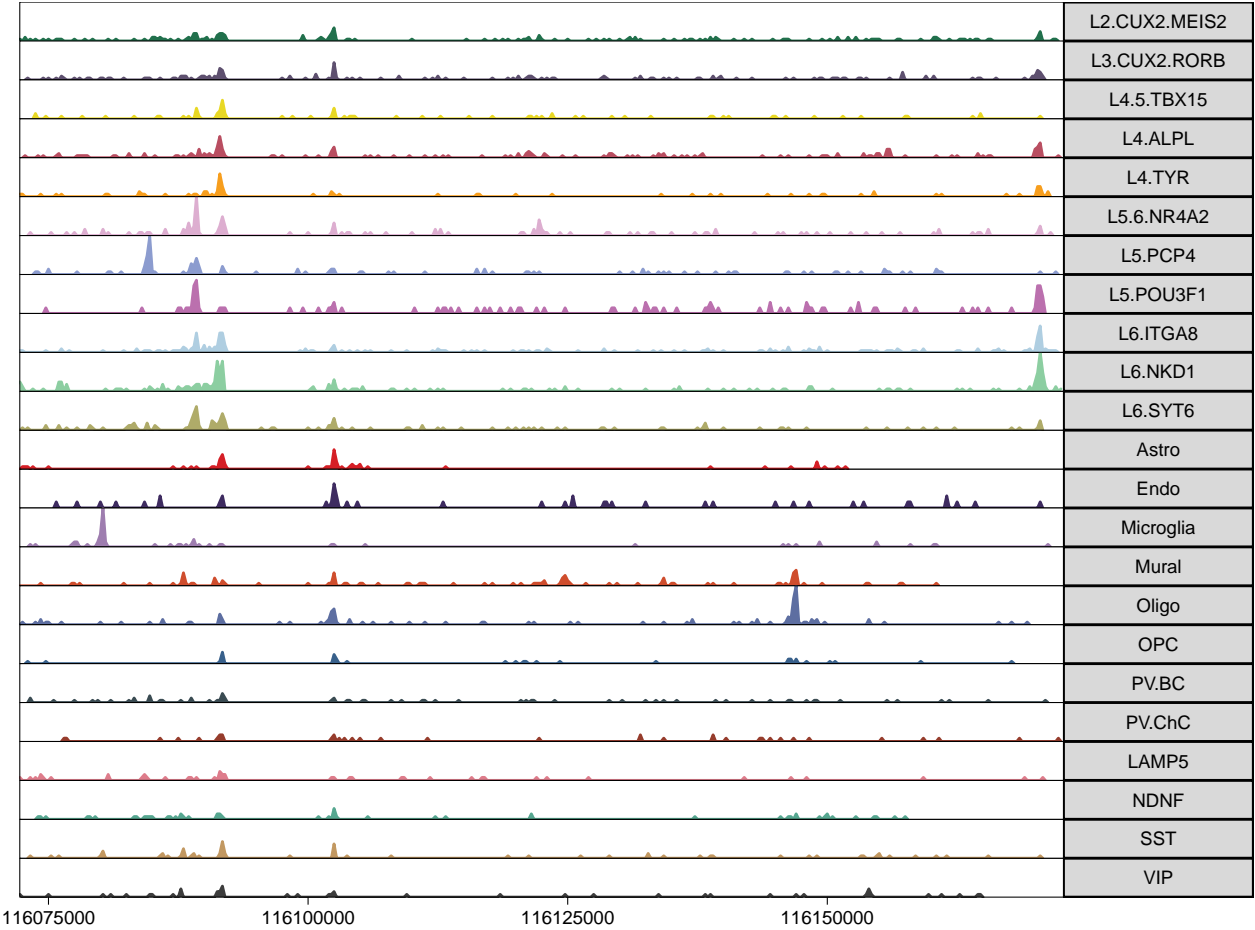


Peaks

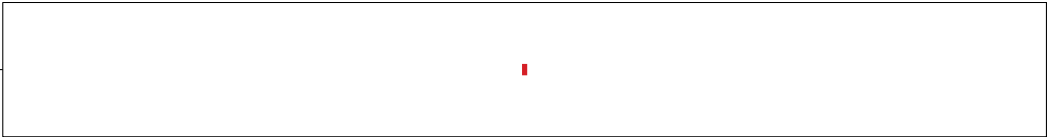


chr4:116072238–116172739 Marker Peak Tracks for: L5.6.NR4A2

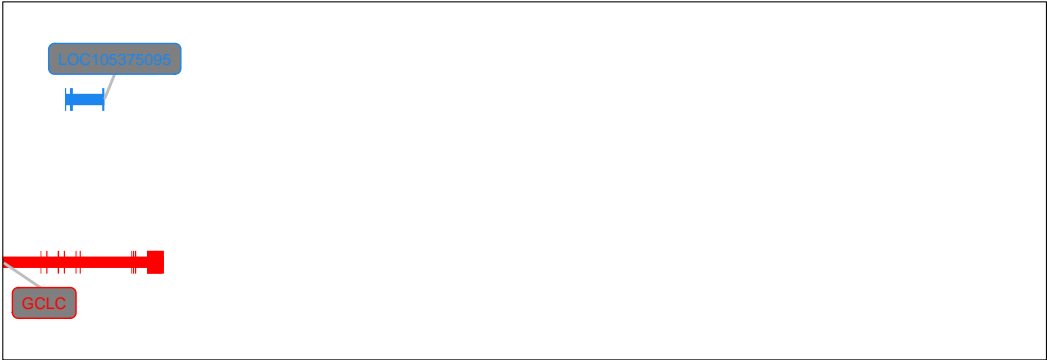
Coverage
(Norm. ATAC Signal Range (0–0.52) by ReadsInTSS)



L5.6.NR4A2

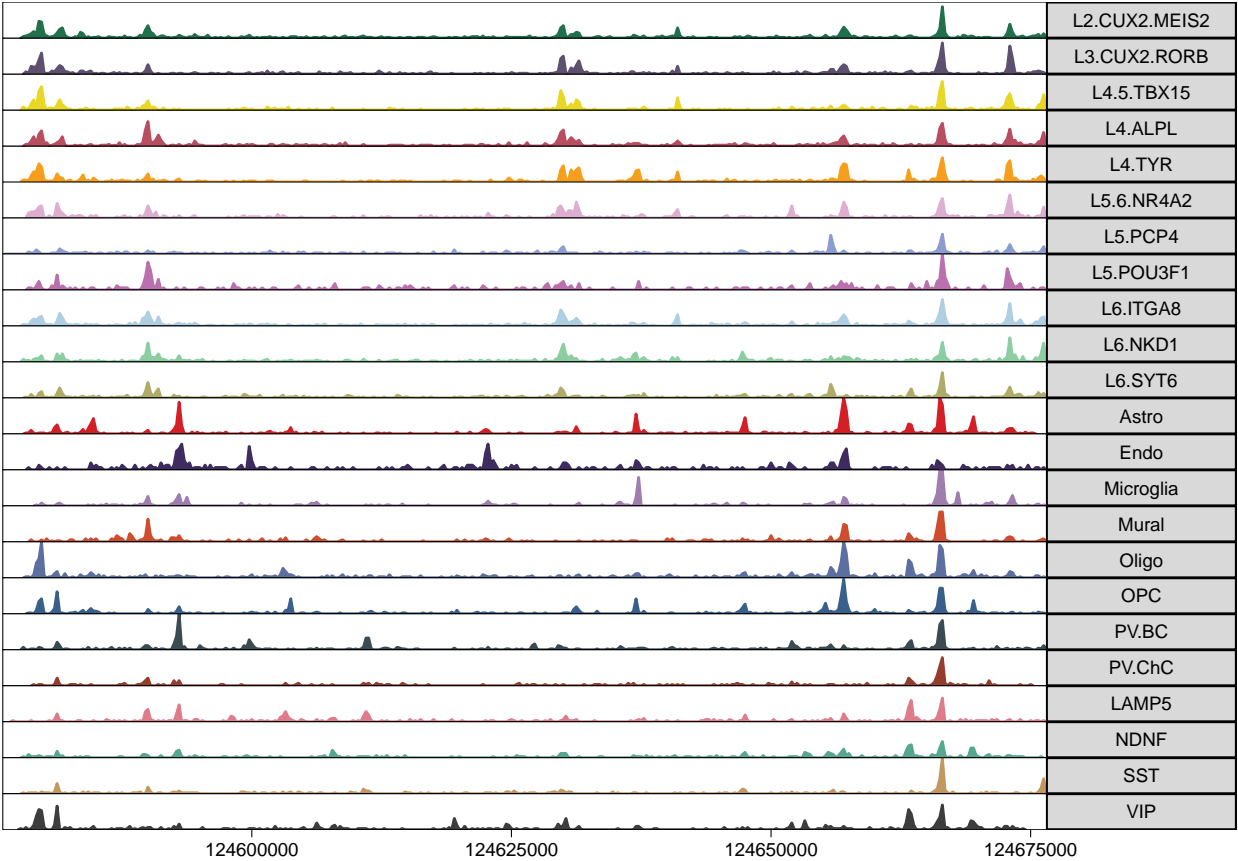


Peaks



Genes

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



L5.6.NR4A2

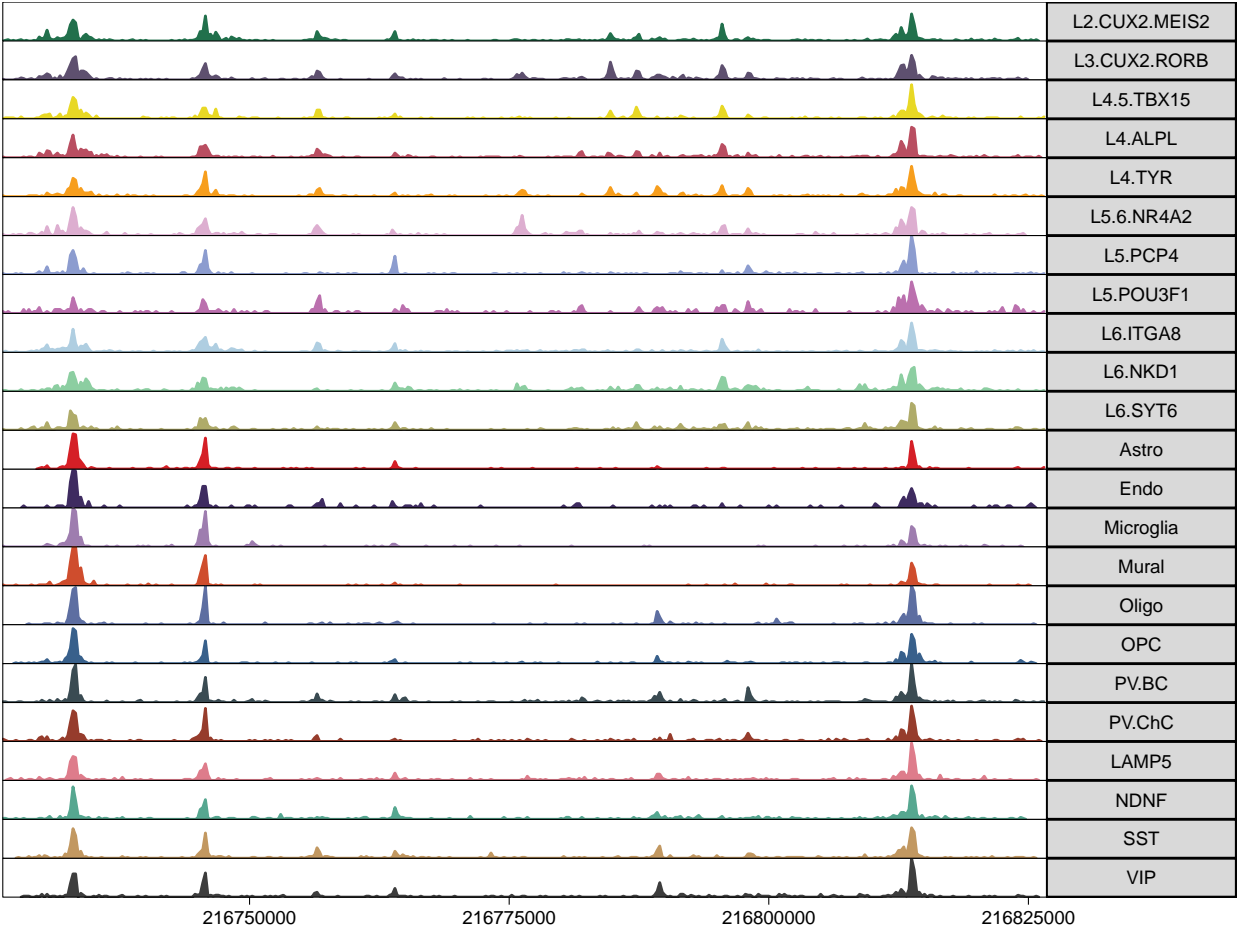


Peaks

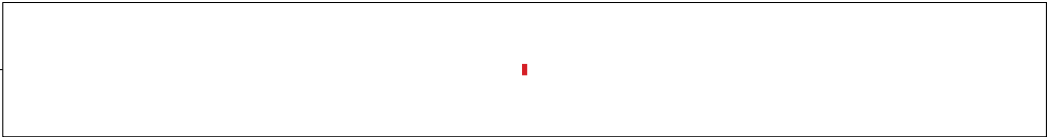
GeneTrack

chr1:216726237-216826738 Marker Peak Tracks for: L5.6.NR4A2

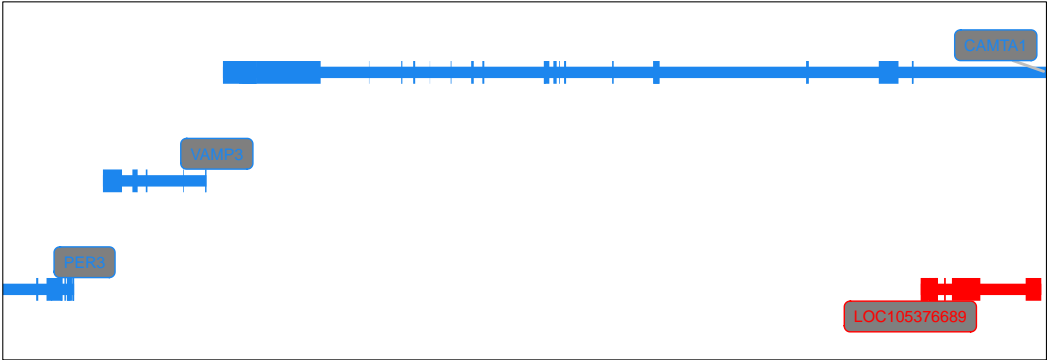
Coverage
(Norm. ATAC Signal Range (0-1.45) by ReadsInTSS)



L5.6.NR4A2



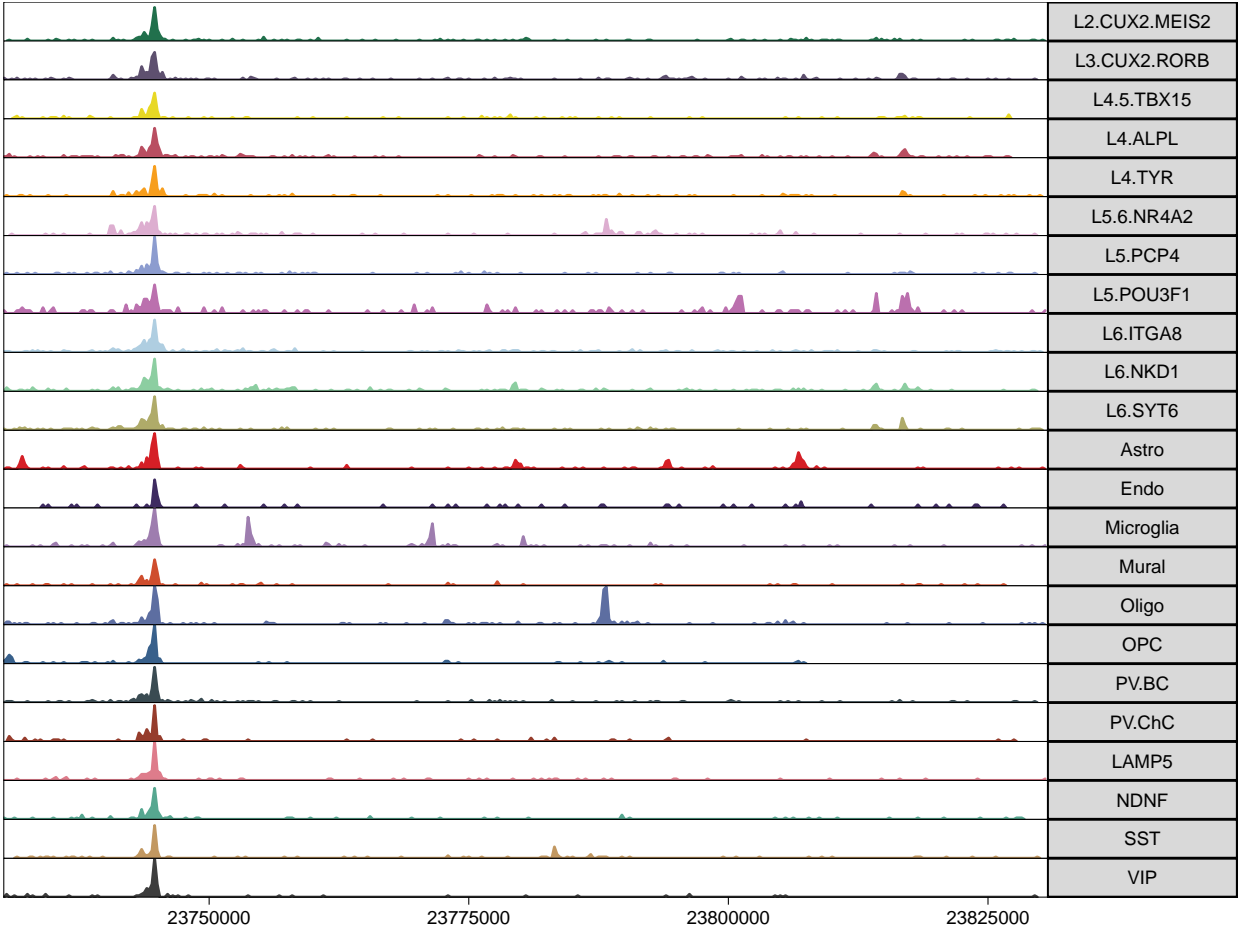
Peaks



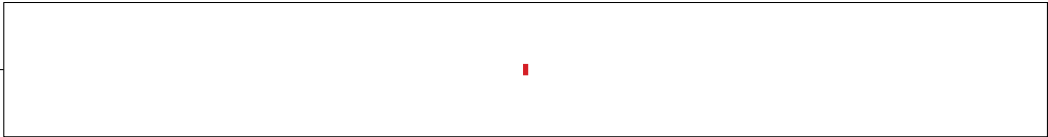
Genes

chr9:23730225–23830726 Marker Peak Tracks for: L5.6.NR4A2

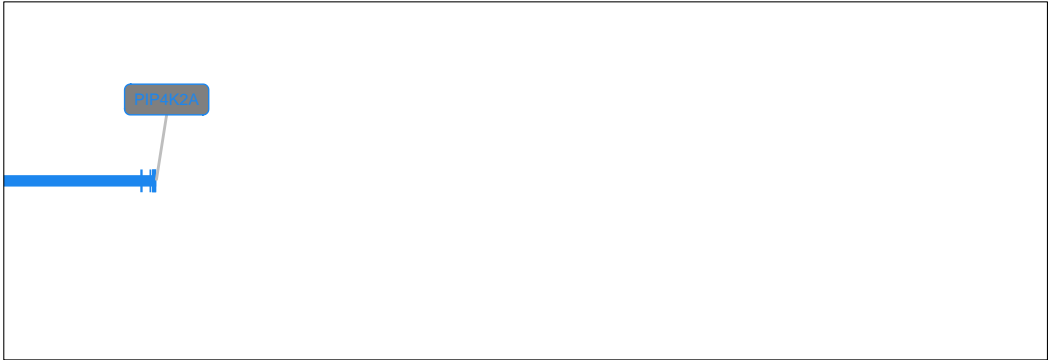
Coverage
(Norm. ATAC Signal Range (0–1) by ReadslnTSS)



L5.6.NR4A2



Peaks



Genes

L5.6.NR4A2

Peaks

Genes

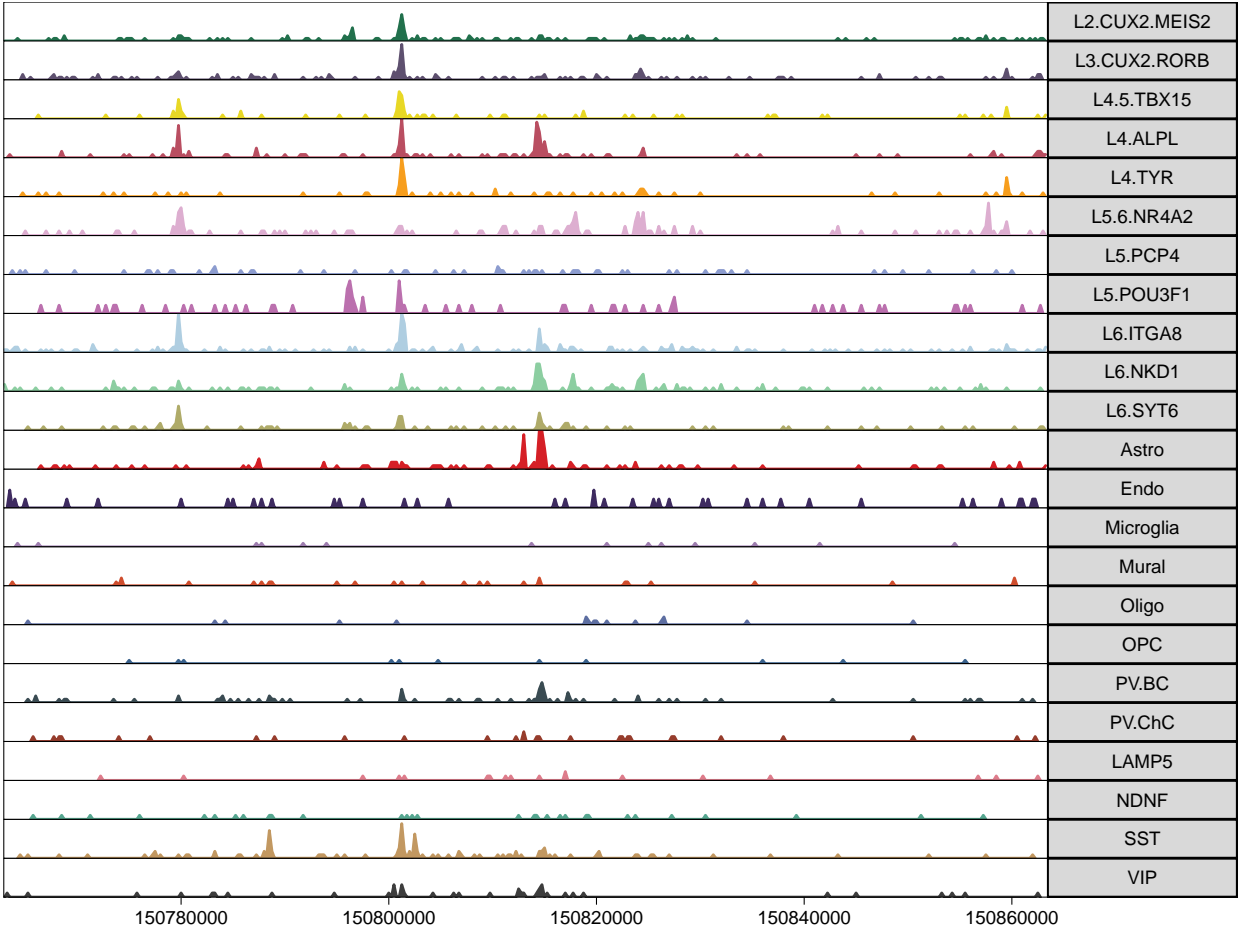
FADS3

FTH1

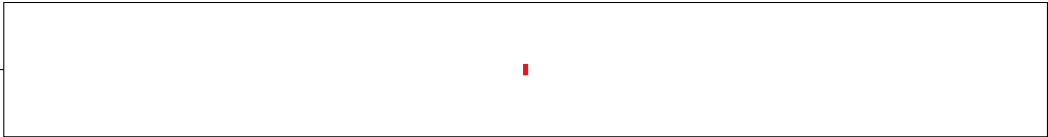
EADS2

chr4:150762926–150863427 Marker Peak Tracks for: L5.6.NR4A2

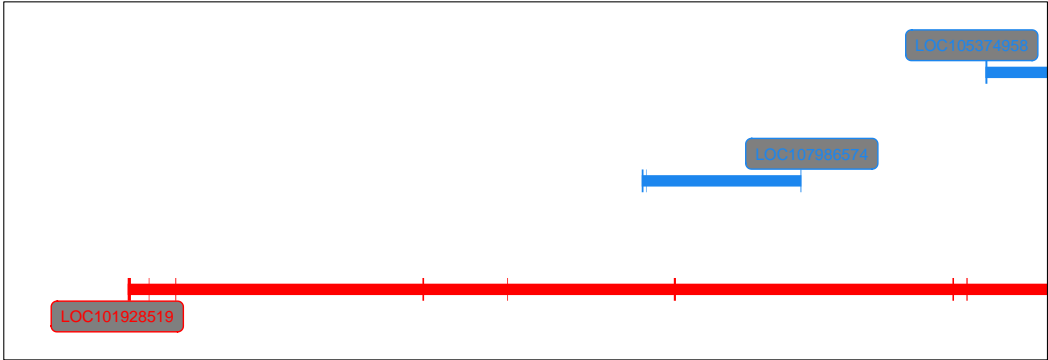
Coverage
(Norm. ATAC Signal Range (0–0.35) by ReadsInTSS)



L5.6.NR4A2

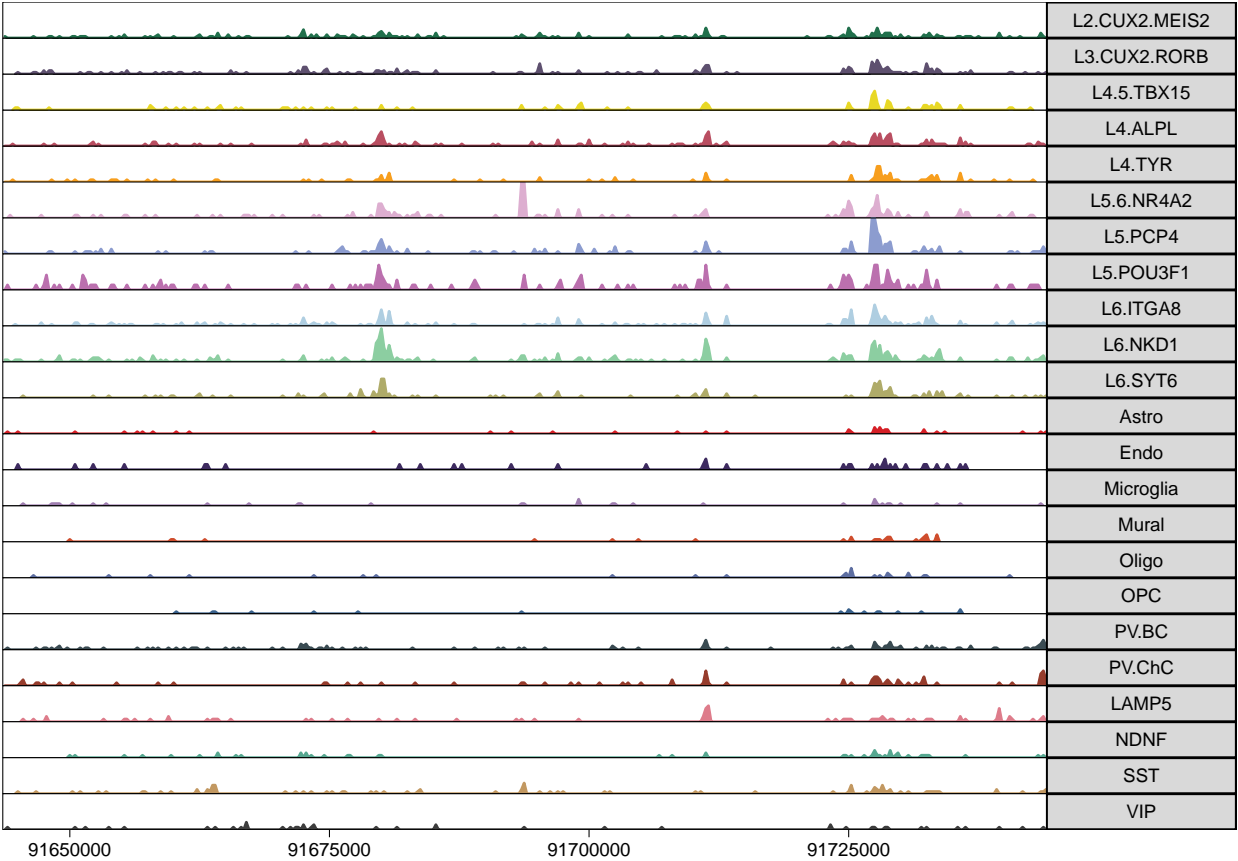


Peaks



Genes

Coverage
(Norm. ATAC Signal Range (0–0.53) by ReadsInTSS)

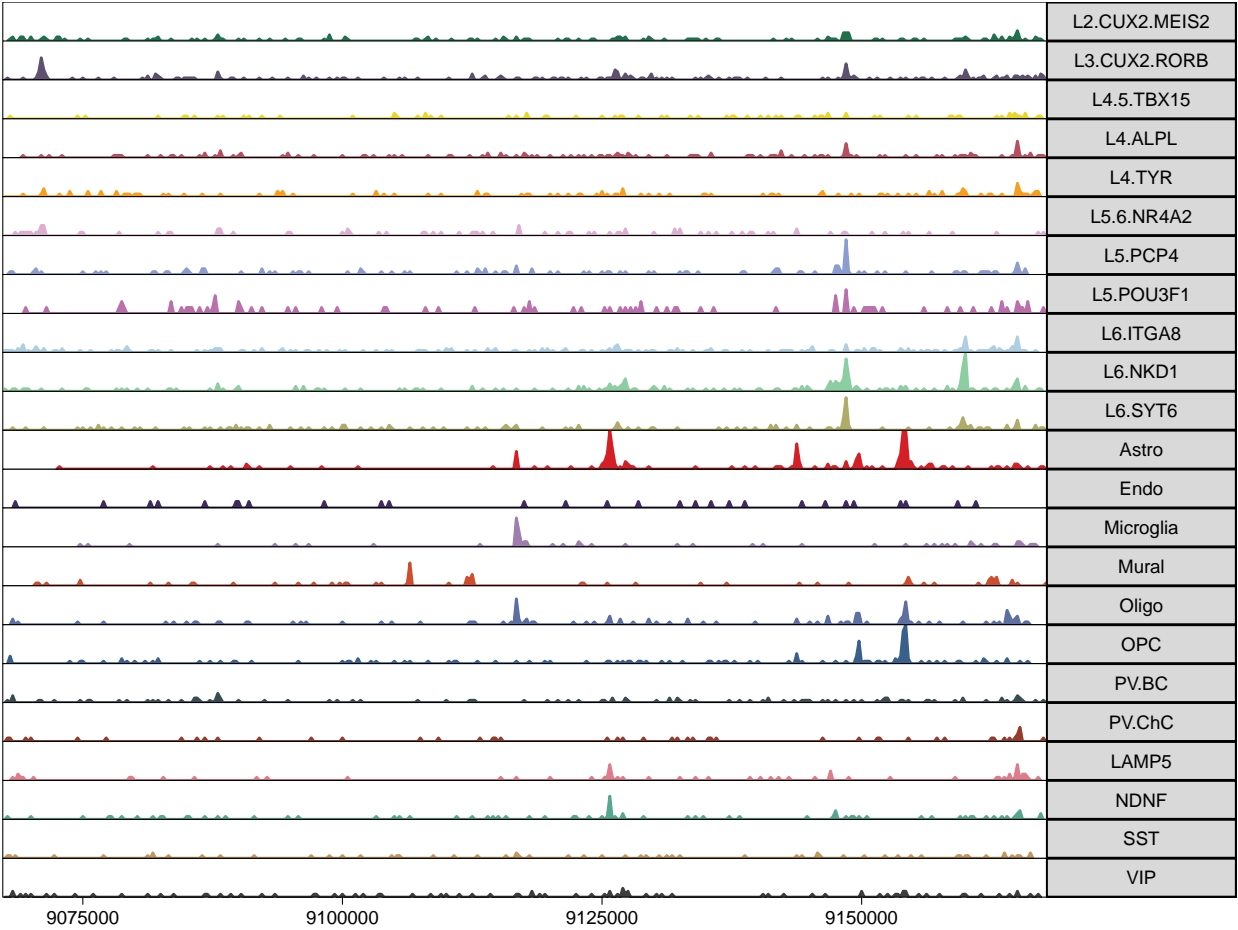


L5.6.NR4A2

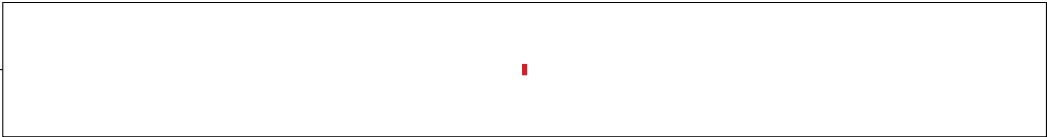


GeneTrack

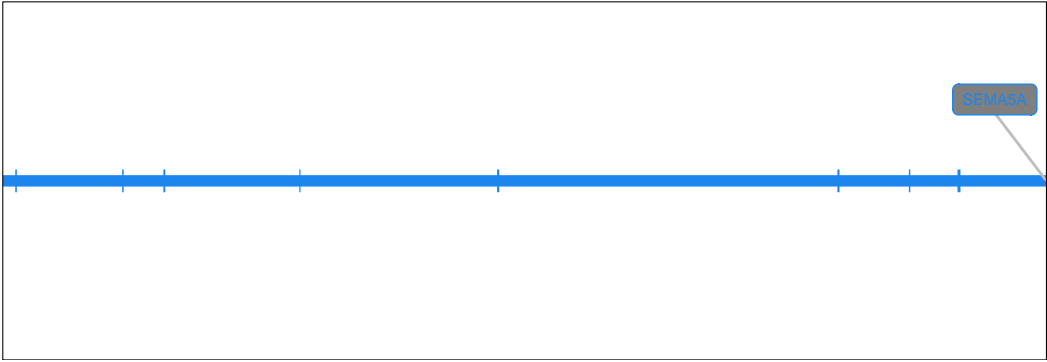
Coverage
(Norm. ATAC Signal Range (0–0.48) by ReadsInTSS)



L5.6.NR4A2

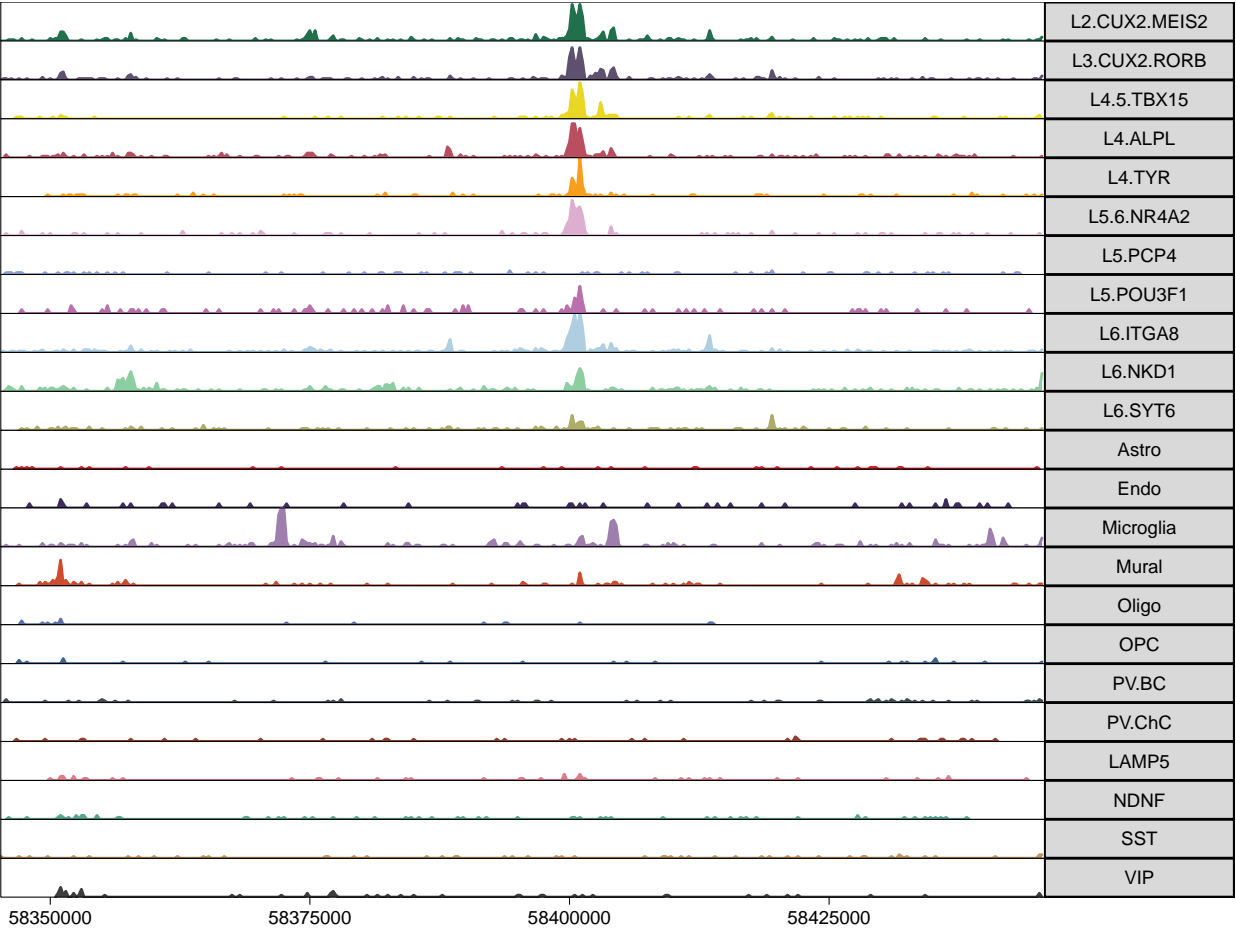


Peaks

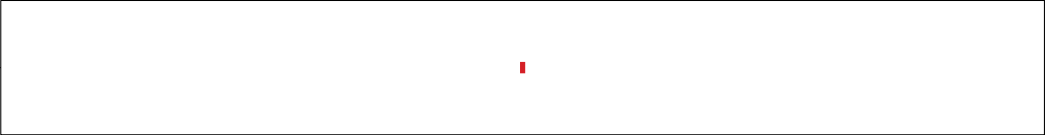


Genes

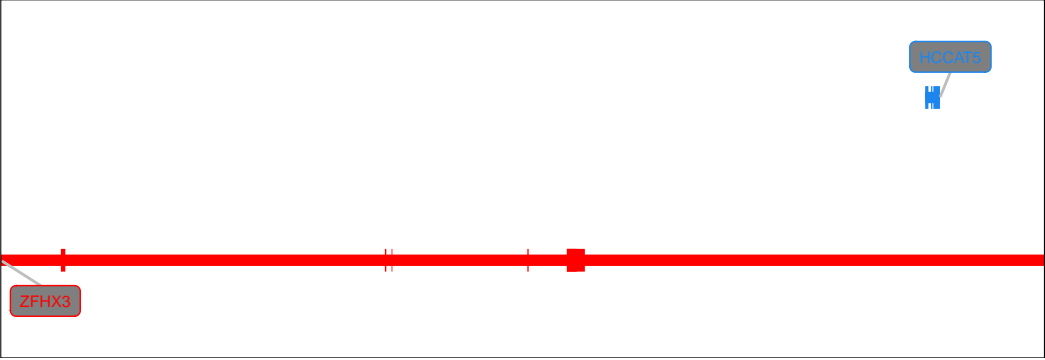
Coverage
(Norm. ATAC Signal Range (0–0.74) by ReadsInTSS)



L5.6.NR4A2



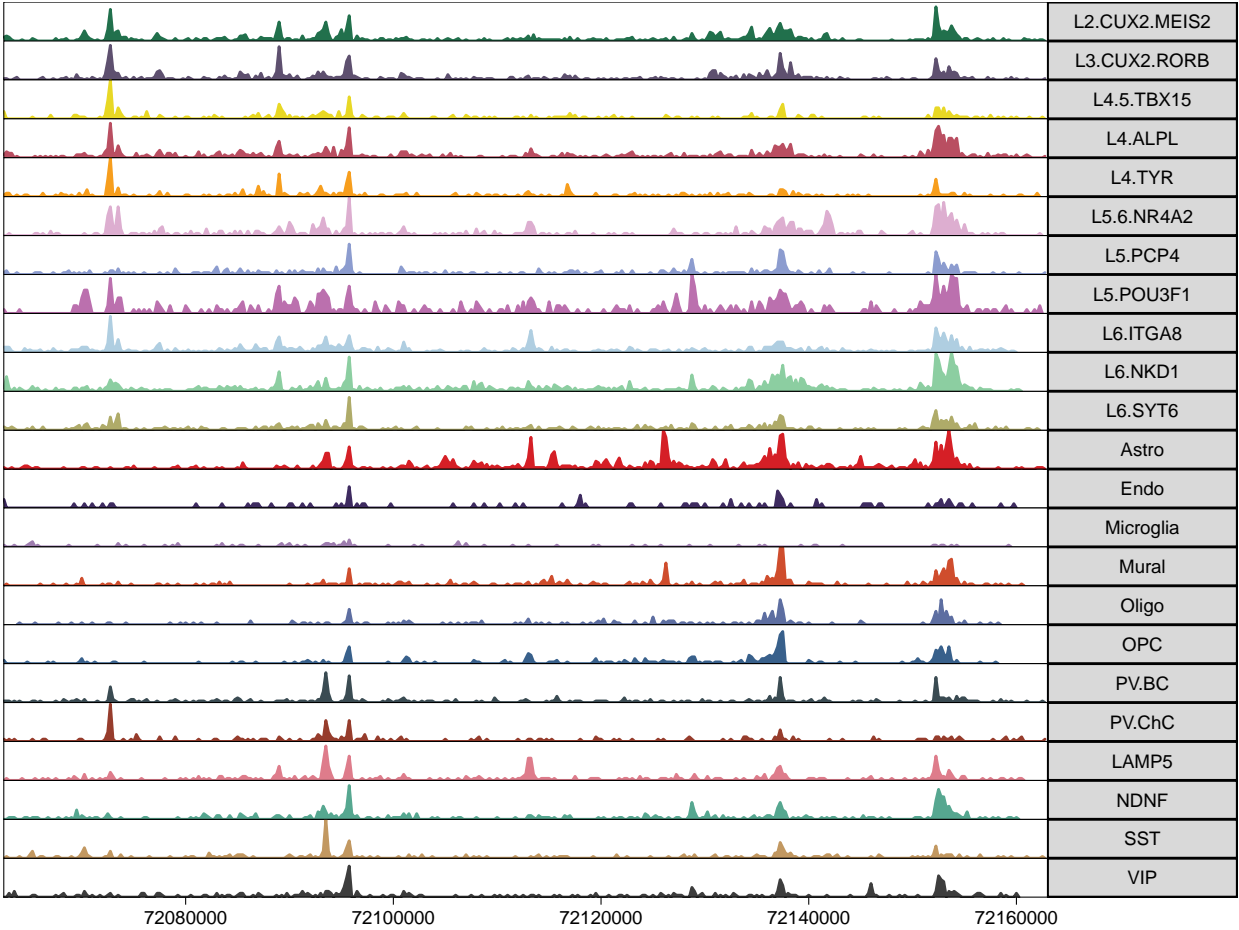
Peaks



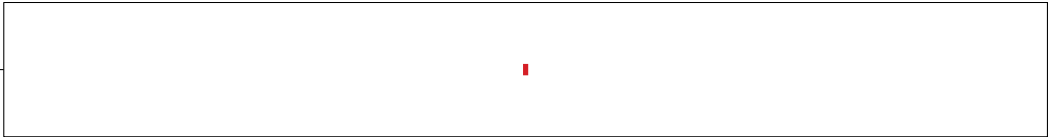
Genes

chr7:72062486–72162987 Marker Peak Tracks for: L5.6.NR4A2

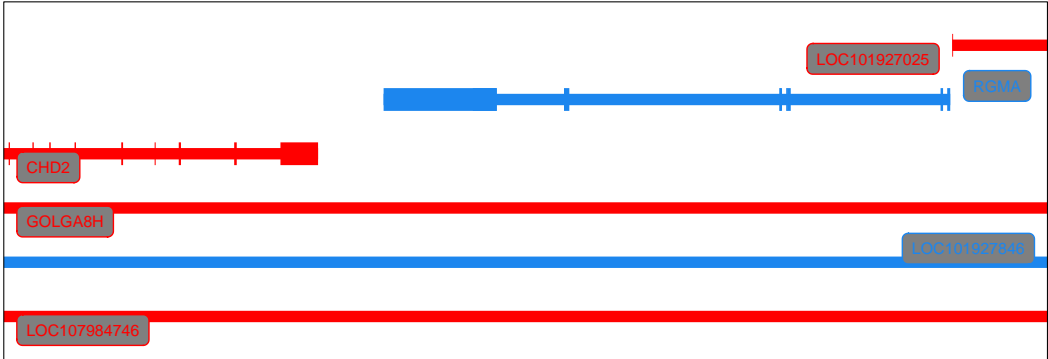
Coverage
(Norm. ATAC Signal Range (0–0.73) by ReadsInTSS)



L5.6.NR4A2

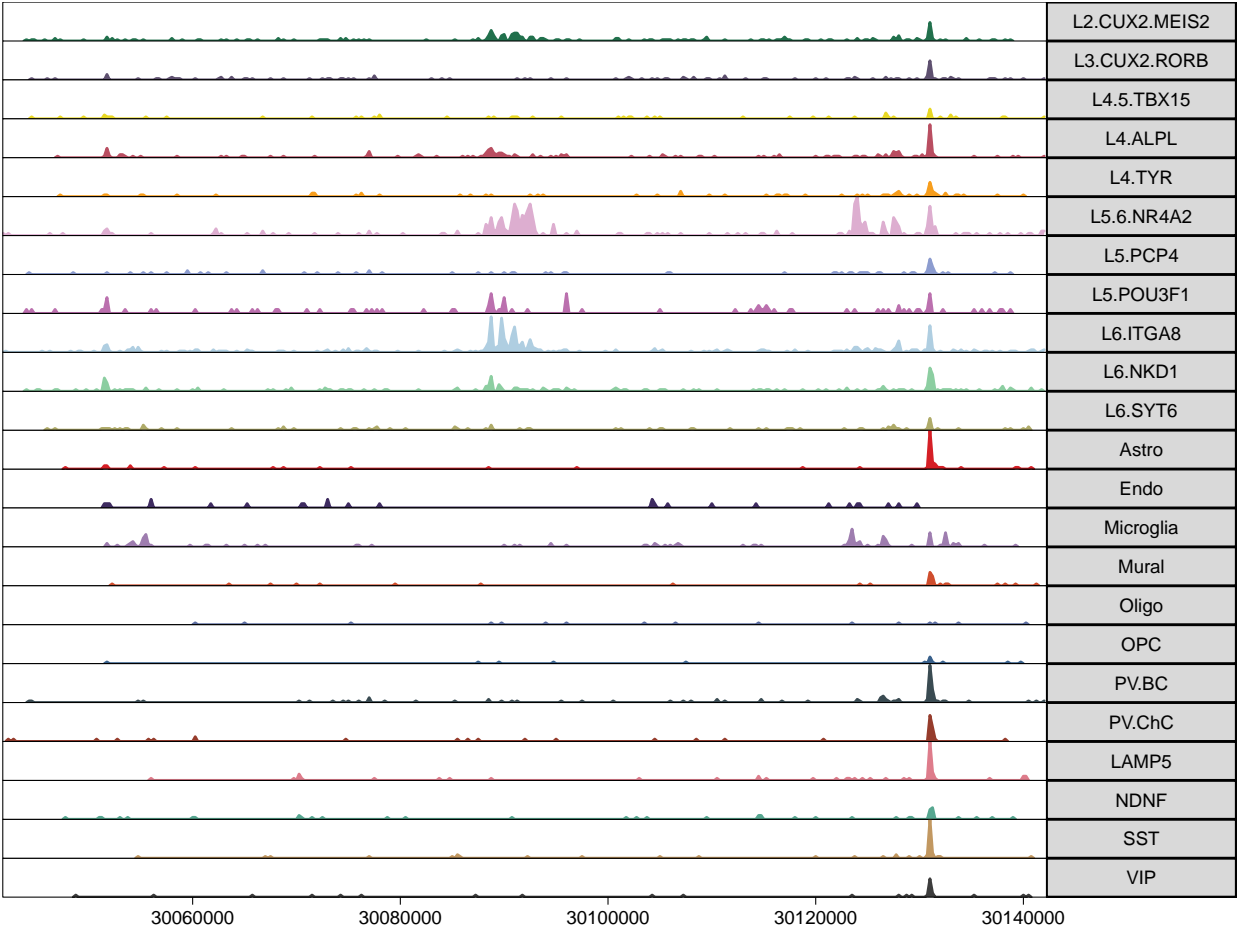


Peaks

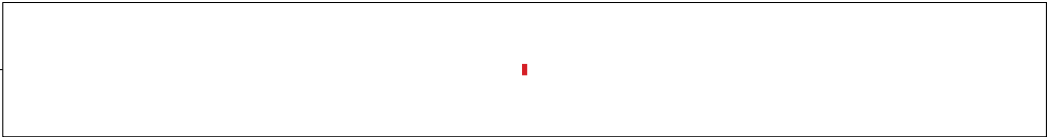


Genes

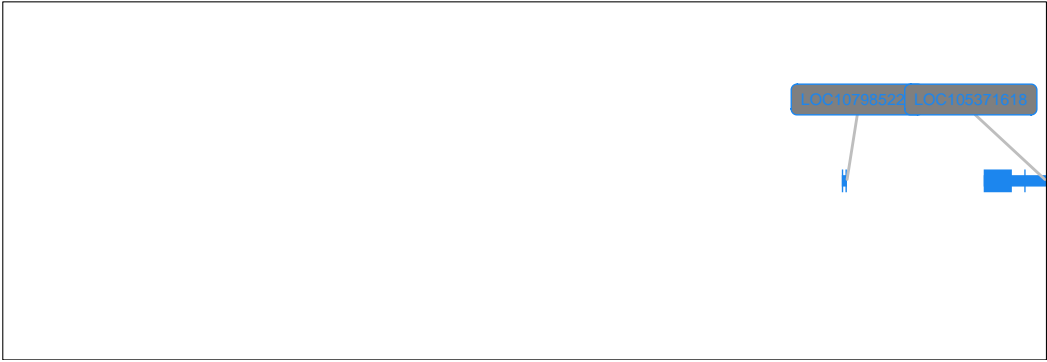
Coverage
(Norm. ATAC Signal Range (0–0.72) by ReadsInTSS)



L5.6.NR4A2



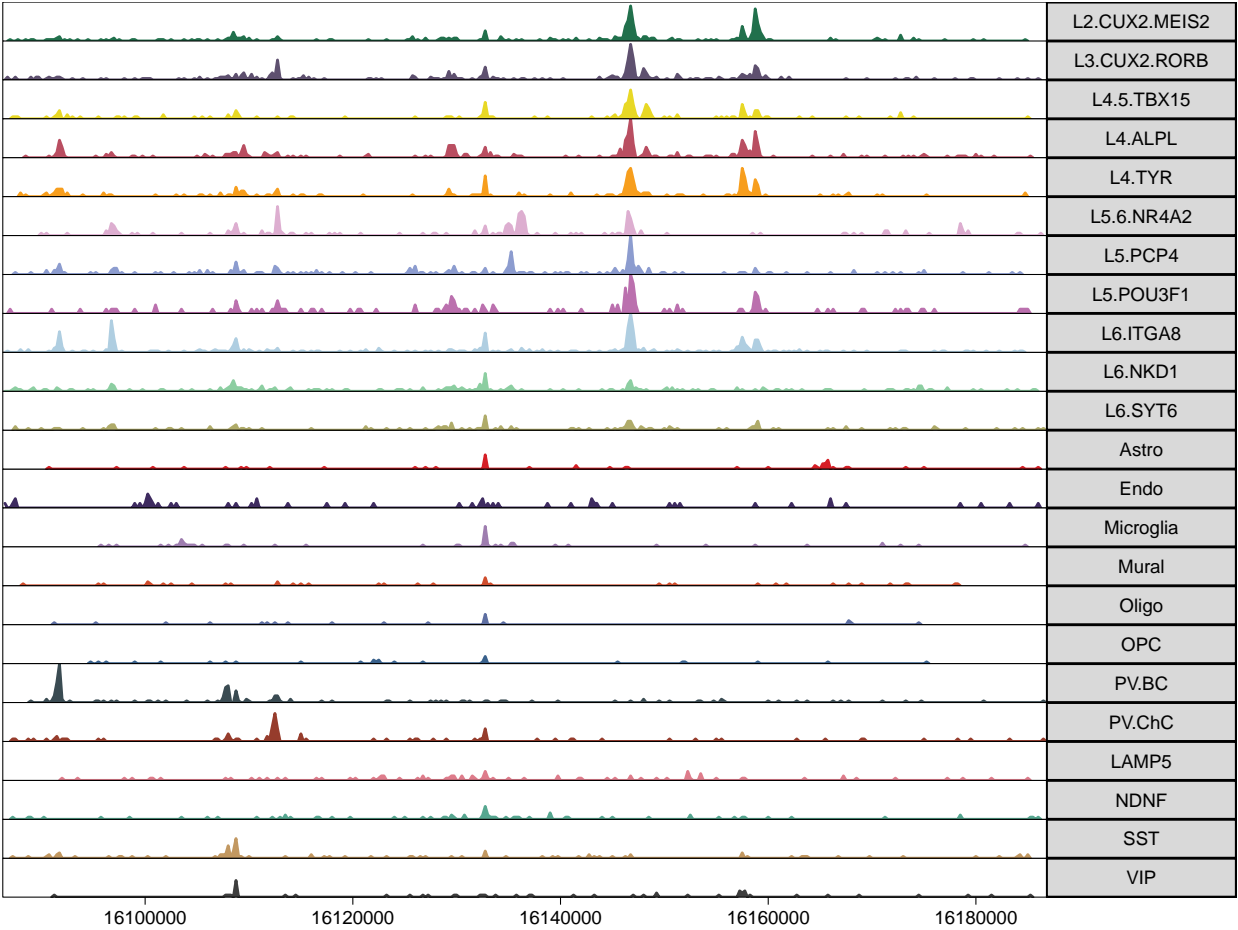
Peaks



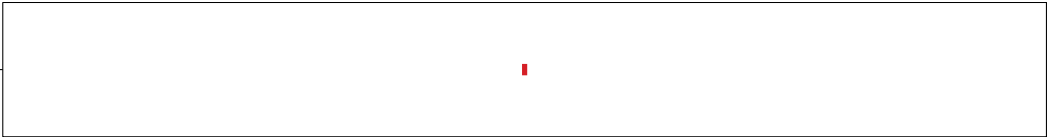
Genes

chr1:16086292–16186793 Marker Peak Tracks for: L5.6.NR4A2

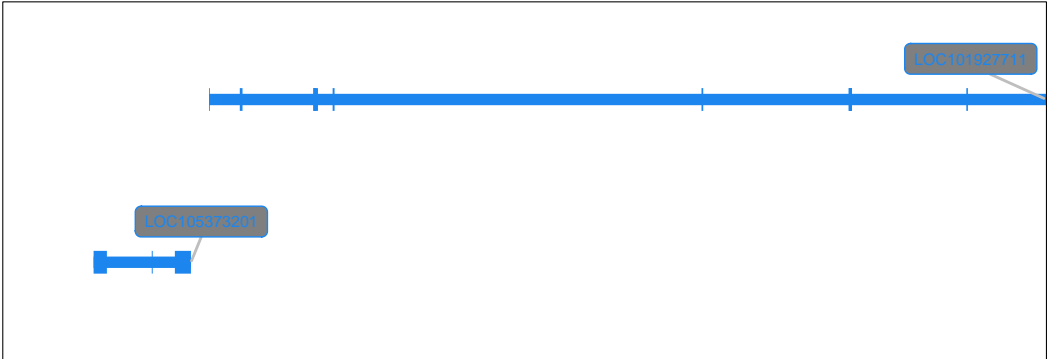
Coverage
(Norm. ATAC Signal Range (0–0.67) by ReadsInTSS)



L5.6.NR4A2

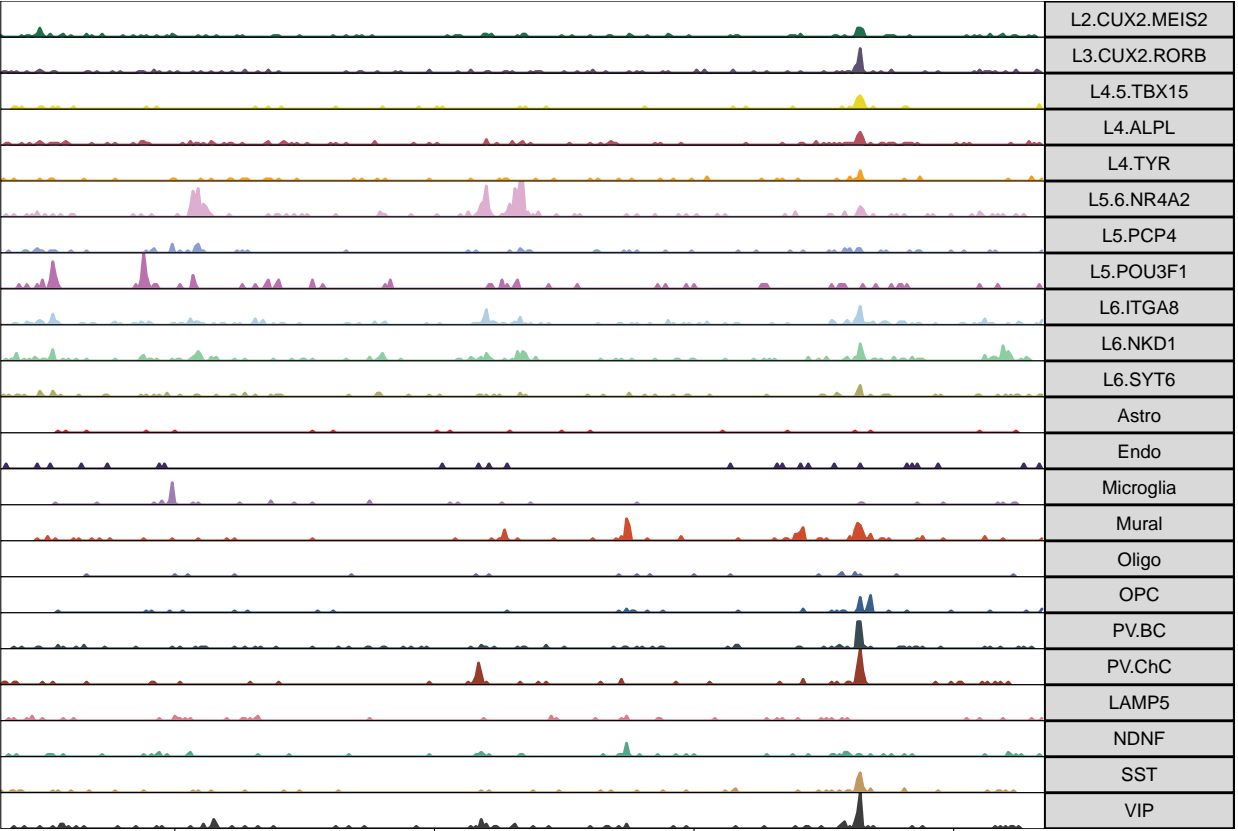


Peaks

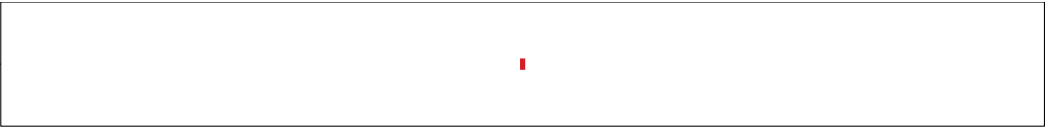


Genes

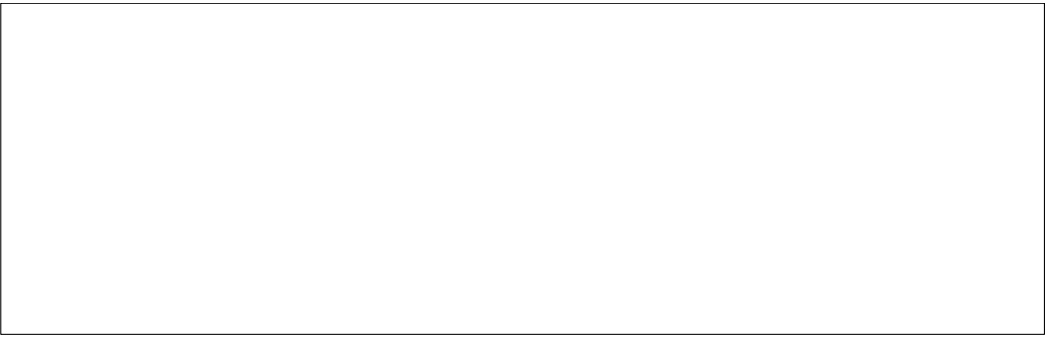
Coverage
(Norm. ATAC Signal Range (0–0.58) by ReadsInTSS)



L5.6.NR4A2



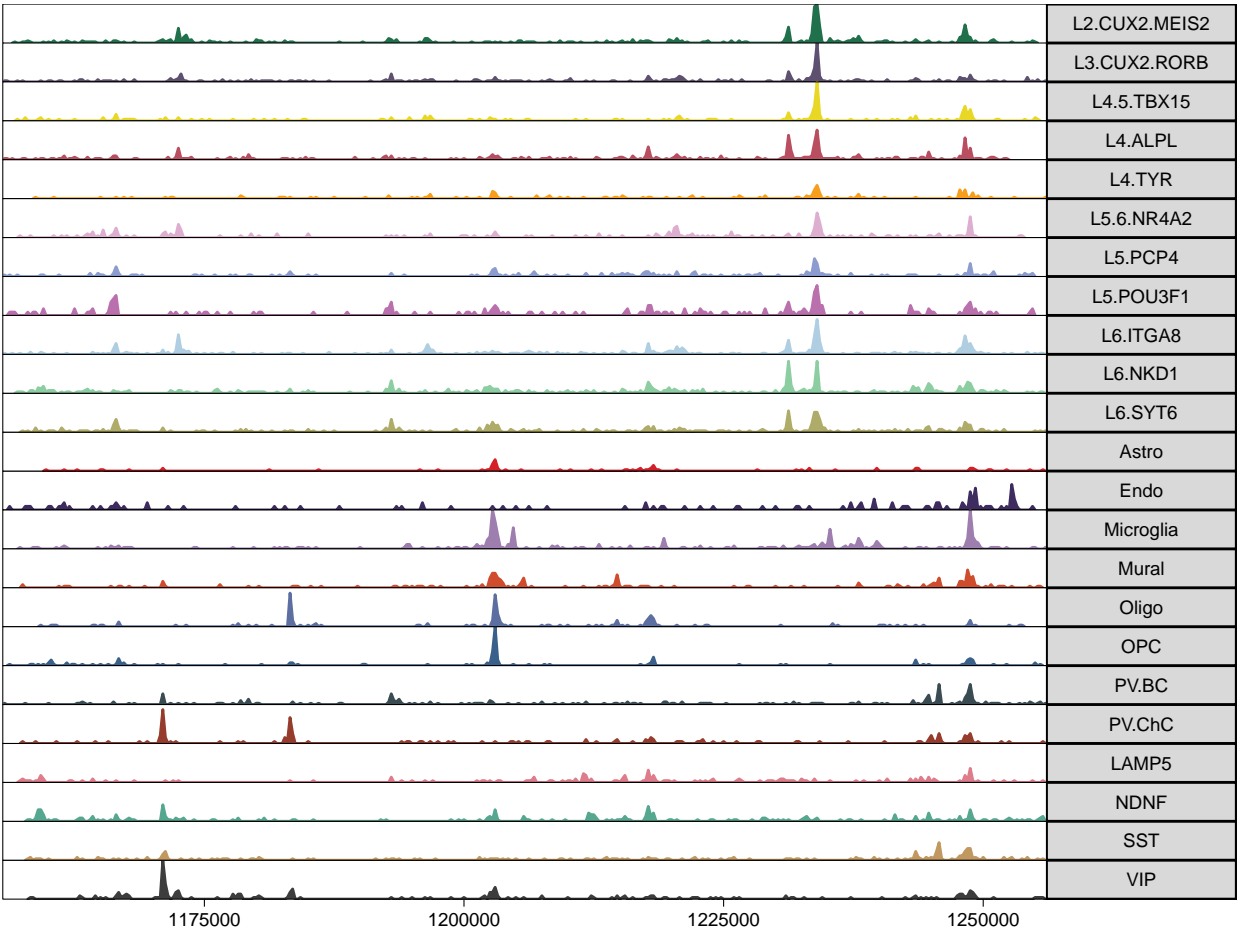
Peaks



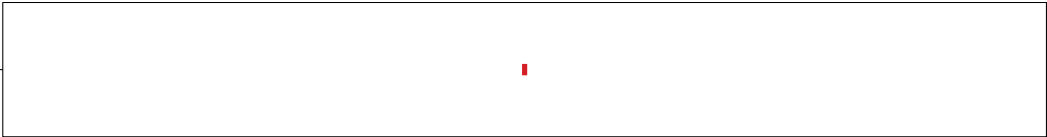
GeneTrack

chr8:1155588–1256089 Marker Peak Tracks for: L5.6.NR4A2

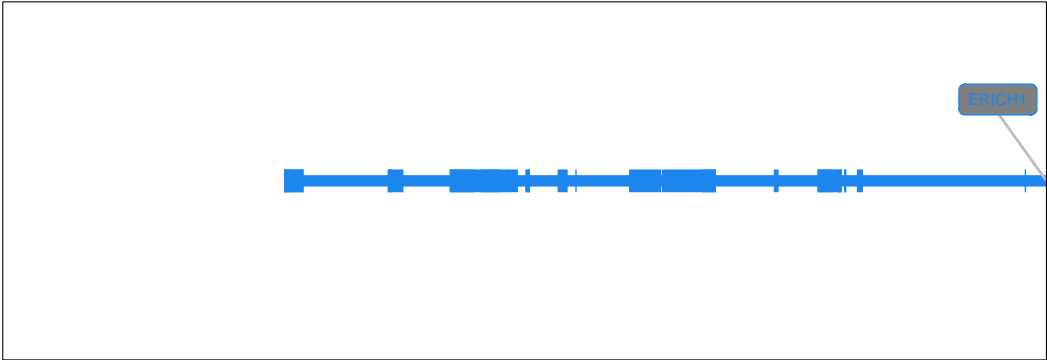
Coverage
(Norm. ATAC Signal Range (0–0.86) by ReadsInTSS)



L5.6.NR4A2

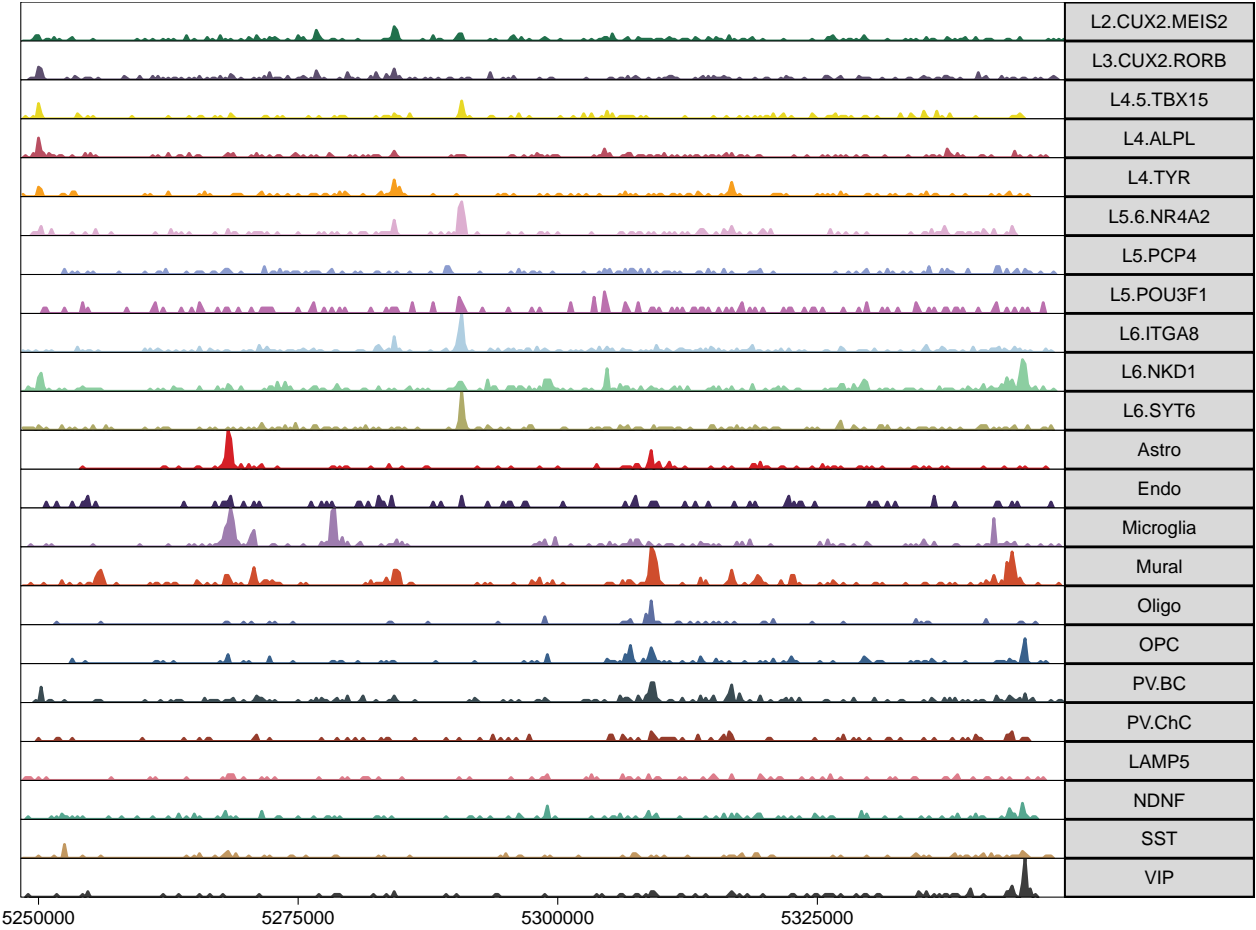


Peaks

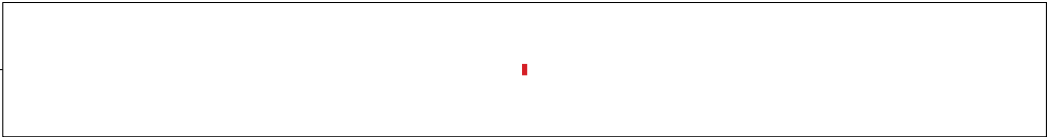


Genes

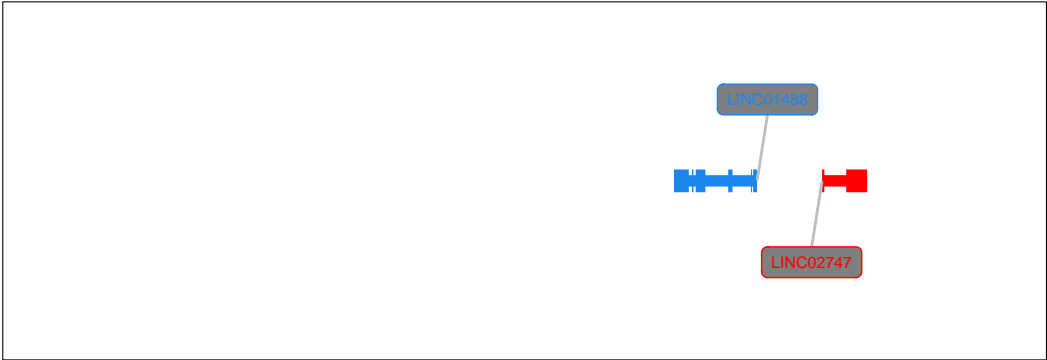
Coverage
(Norm. ATAC Signal Range (0–0.53) by ReadsInTSS)



L5.6.NR4A2

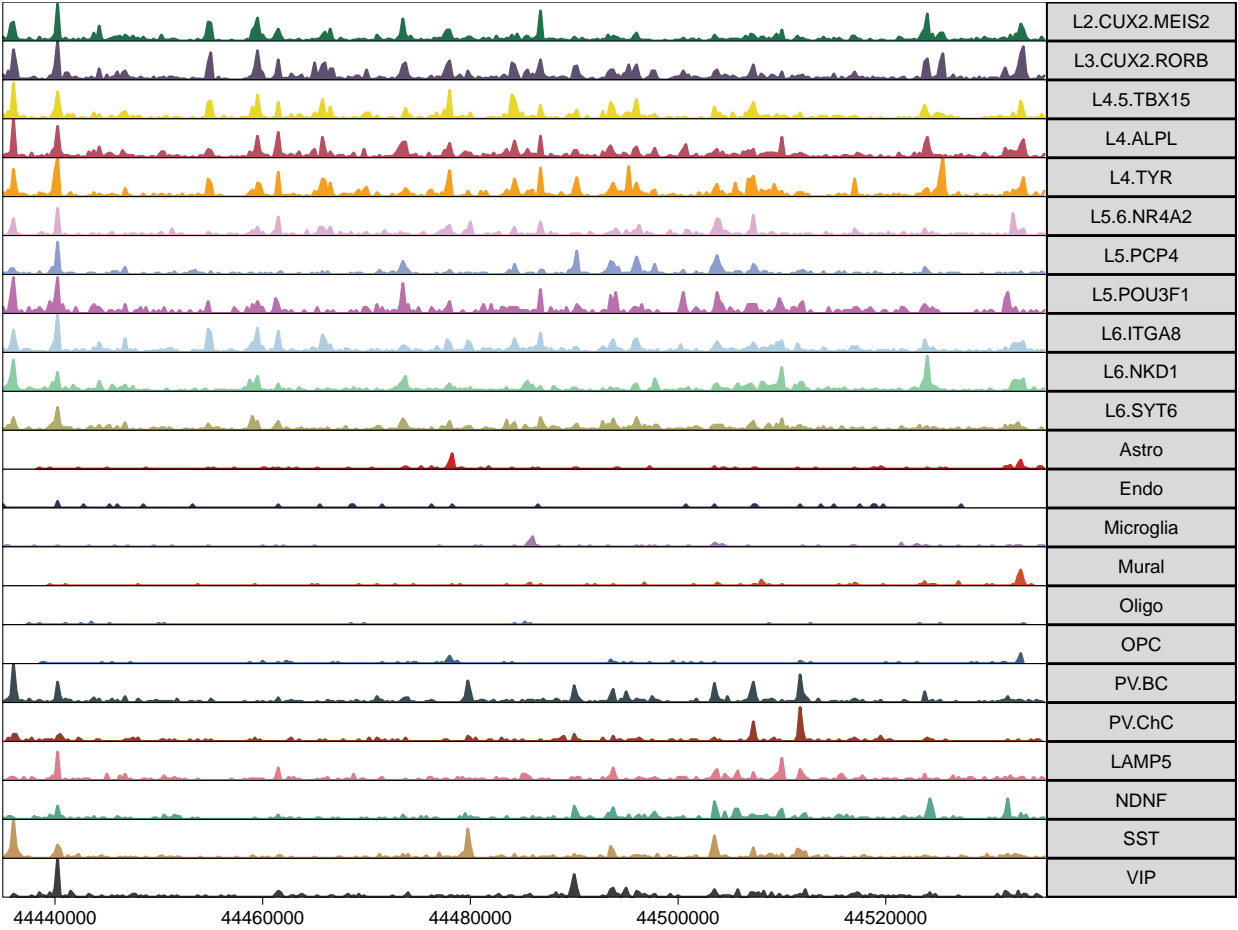


Peaks

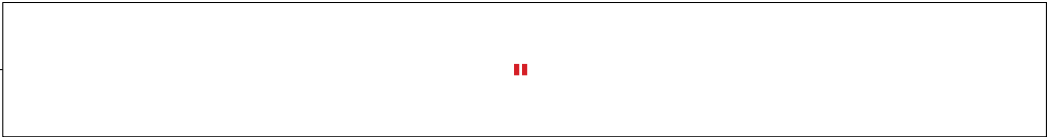


Genes

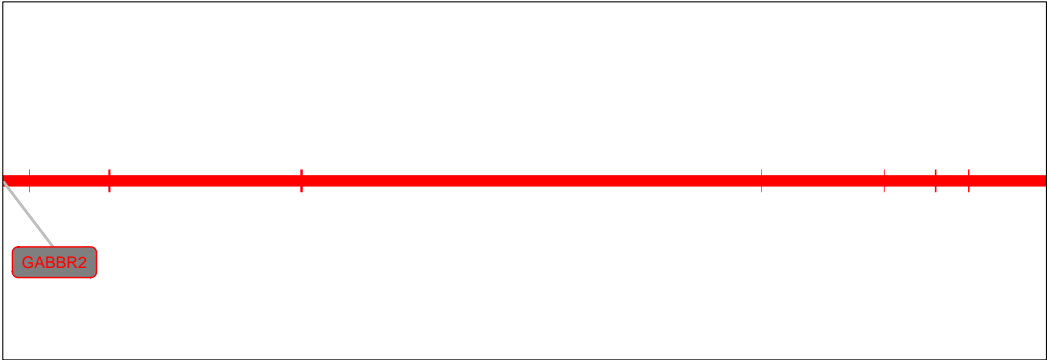
Coverage
(Norm. ATAC Signal Range (0–0.95) by ReadsInTSS)



L5.6.NR4A2

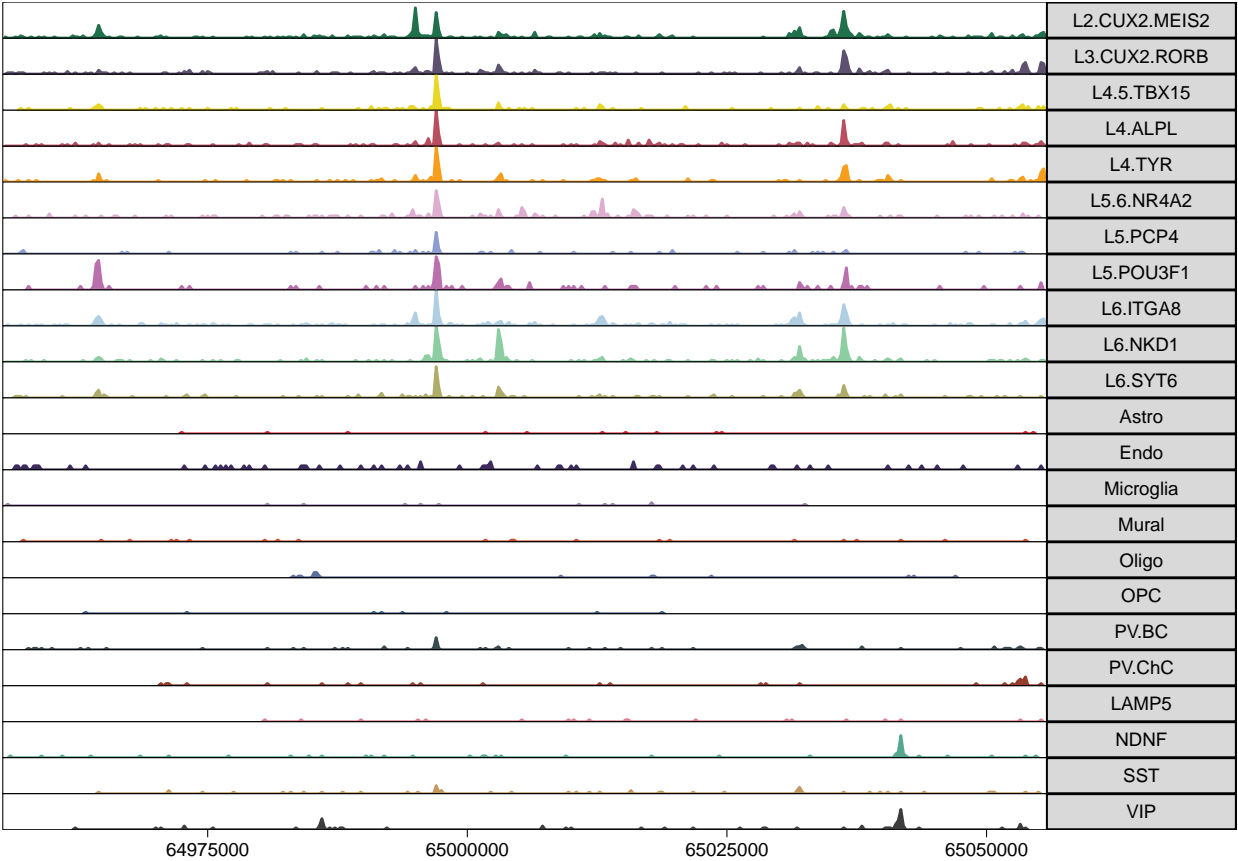


Peaks



Genes

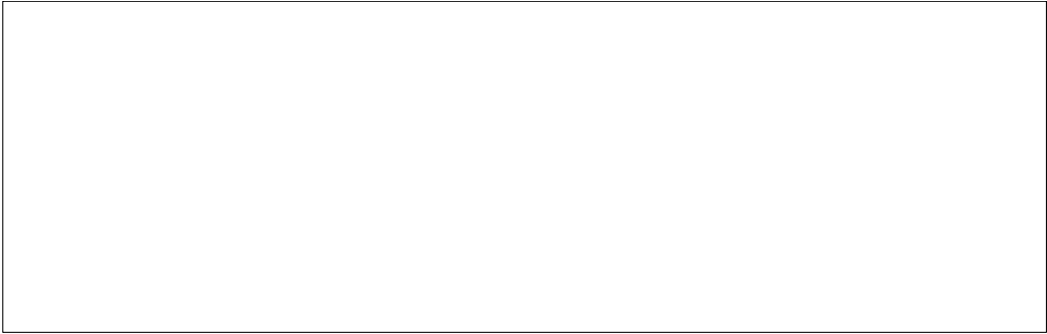
Coverage
(Norm. ATAC Signal Range (0–0.71) by ReadsInTSS)



L5.6.NR4A2



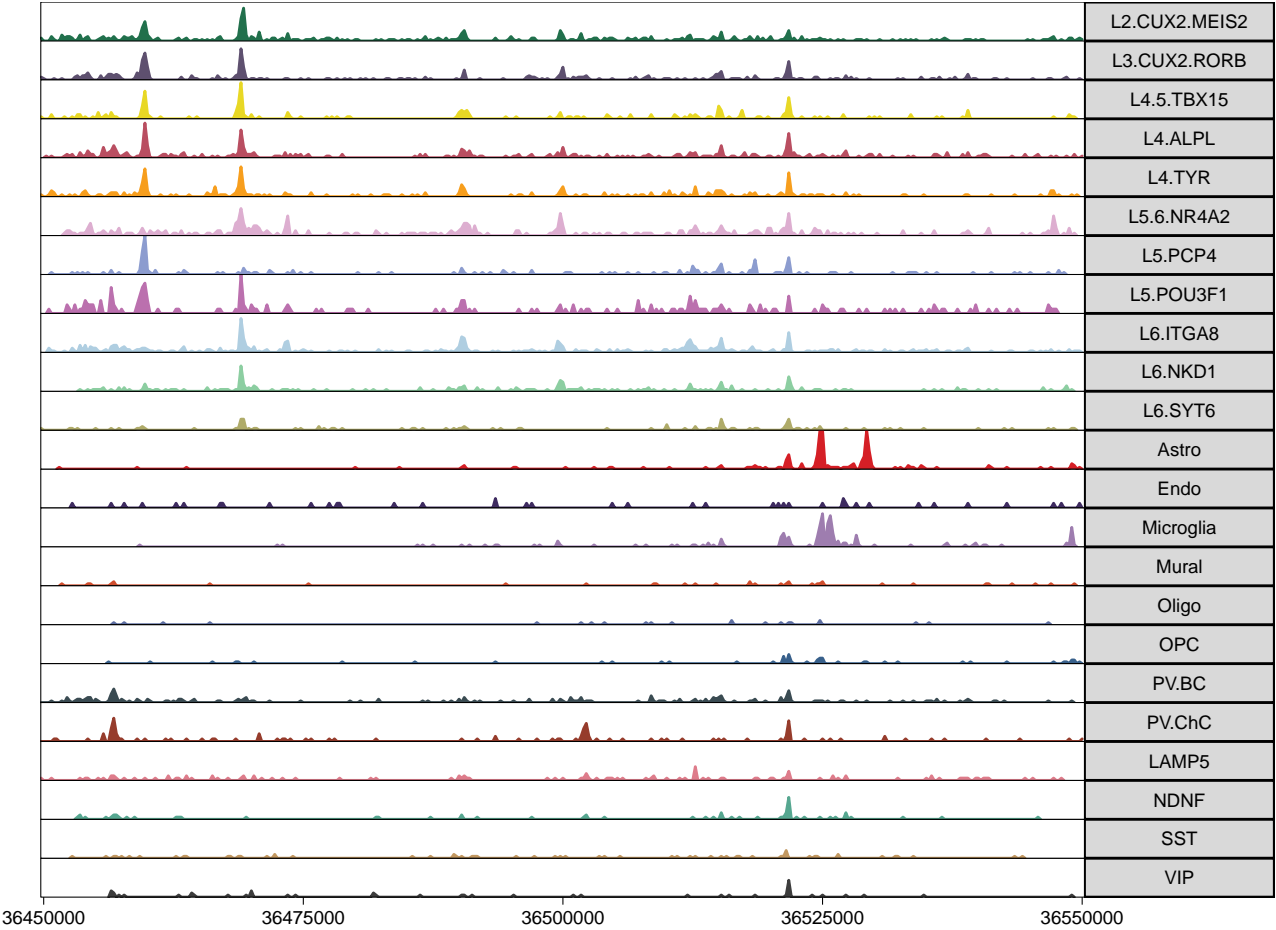
Peaks



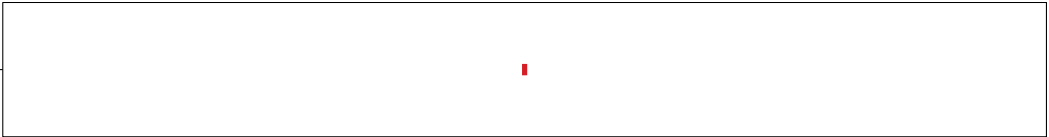
GeneTrack

chr5:36449720–36550221 Marker Peak Tracks for: L5.6.NR4A2

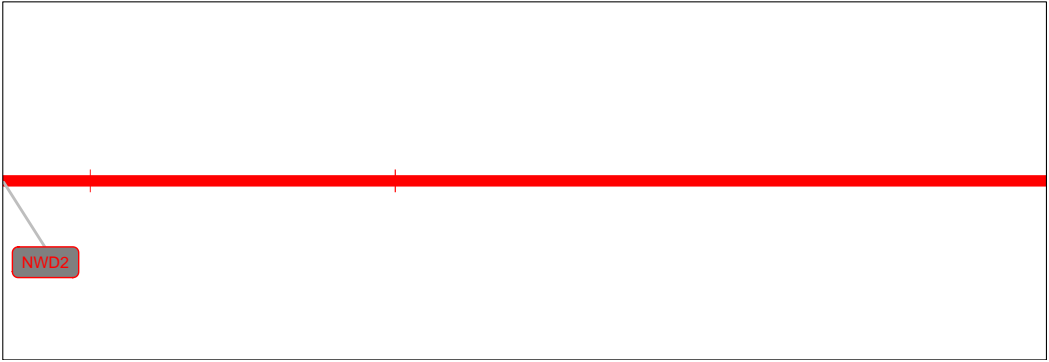
Coverage
(Norm. ATAC Signal Range (0–0.66) by ReadsInTSS)



L5.6.NR4A2

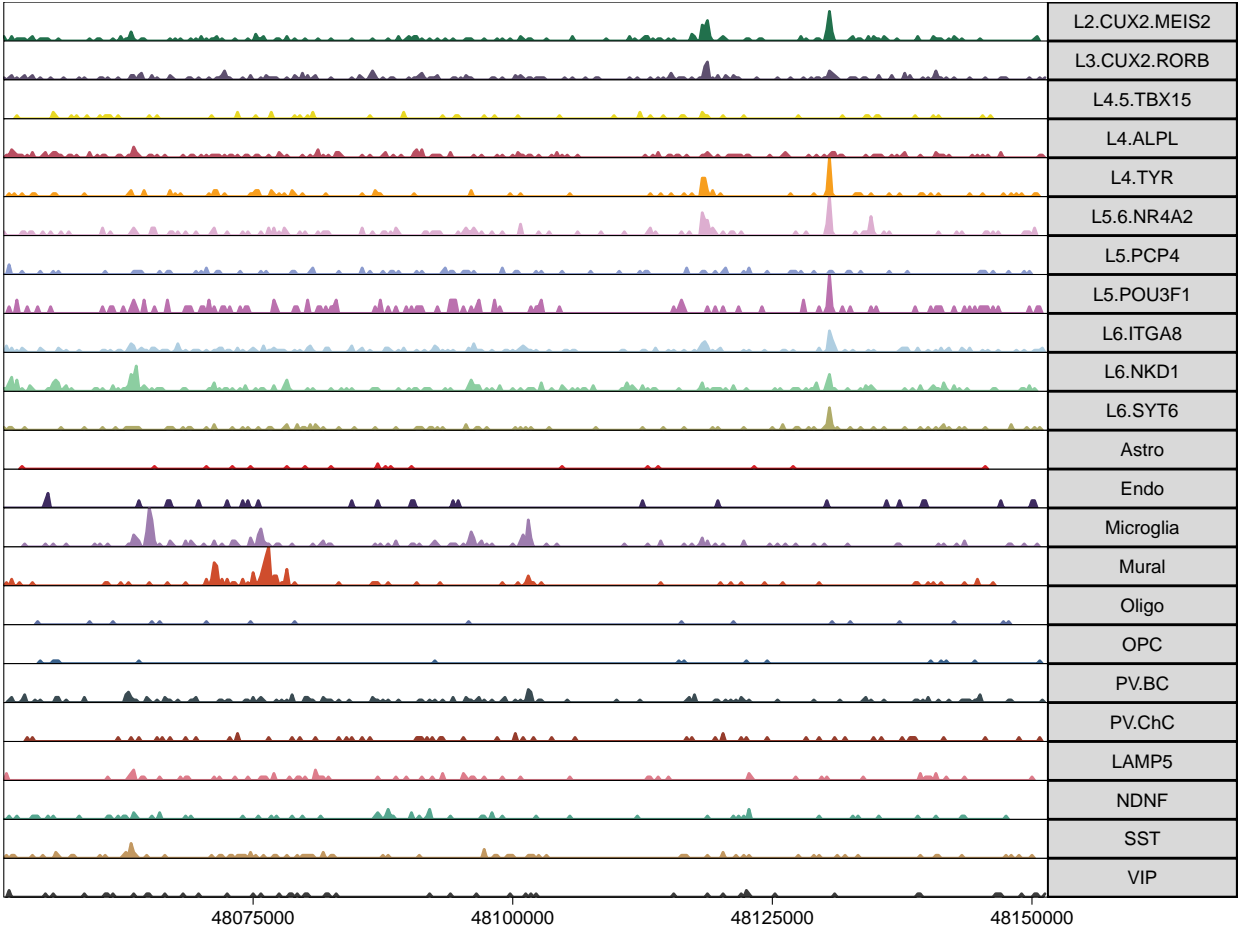


Peaks

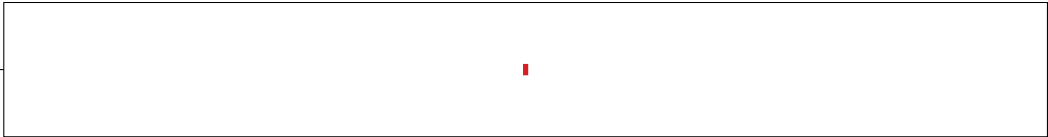


Genes

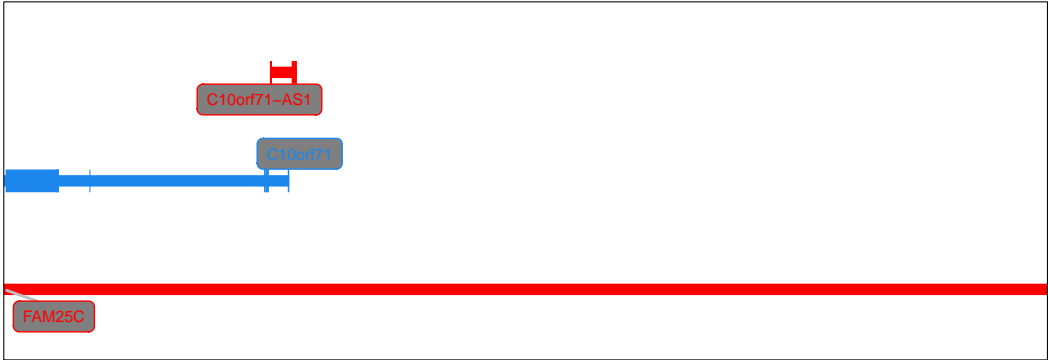
Coverage
(Norm. ATAC Signal Range (0–0.43) by ReadsInTSS)



L5.6.NR4A2

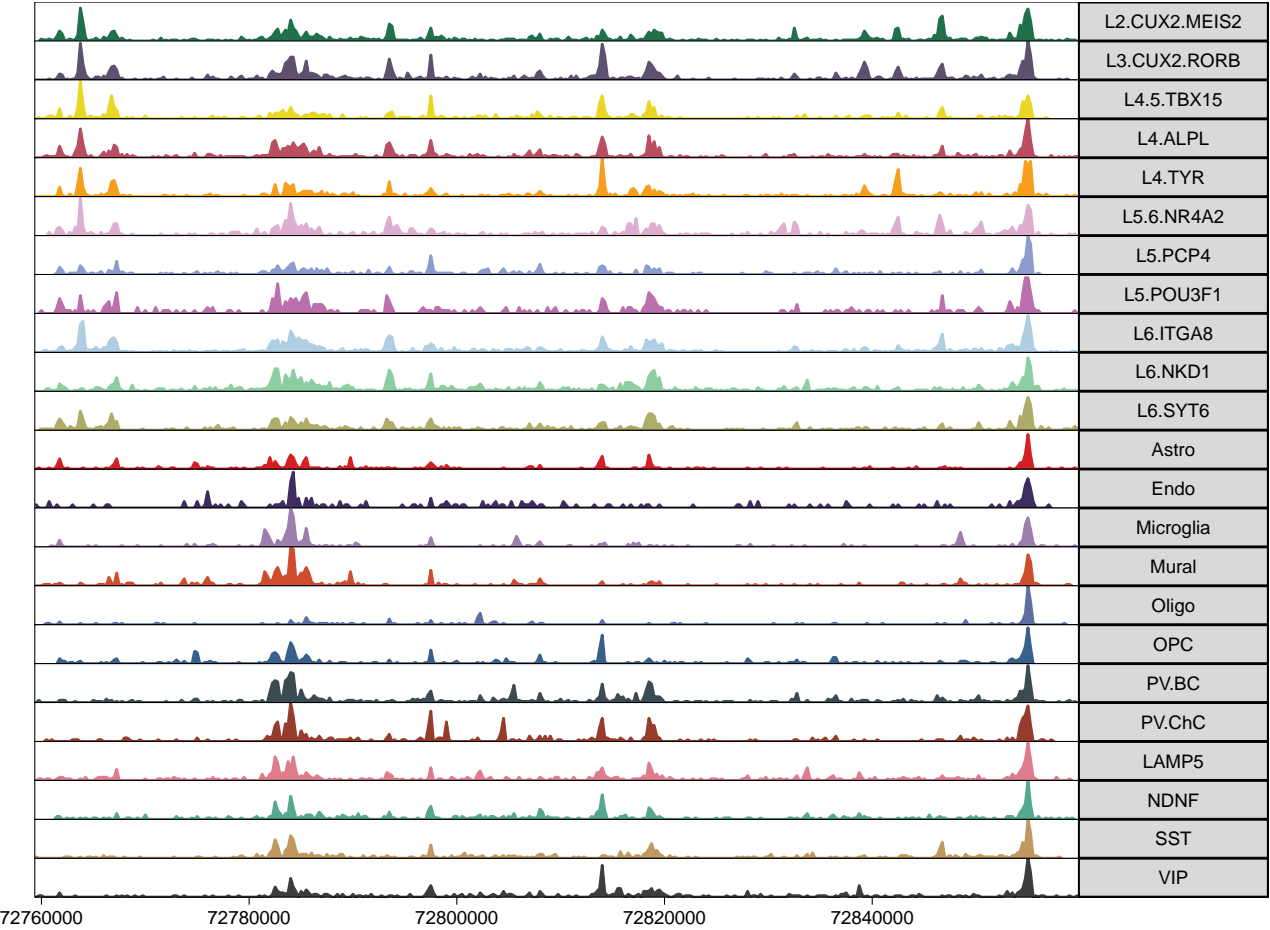


Peaks

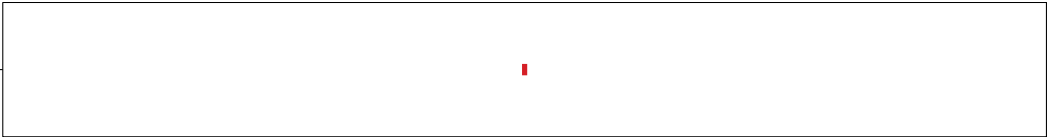


Genes

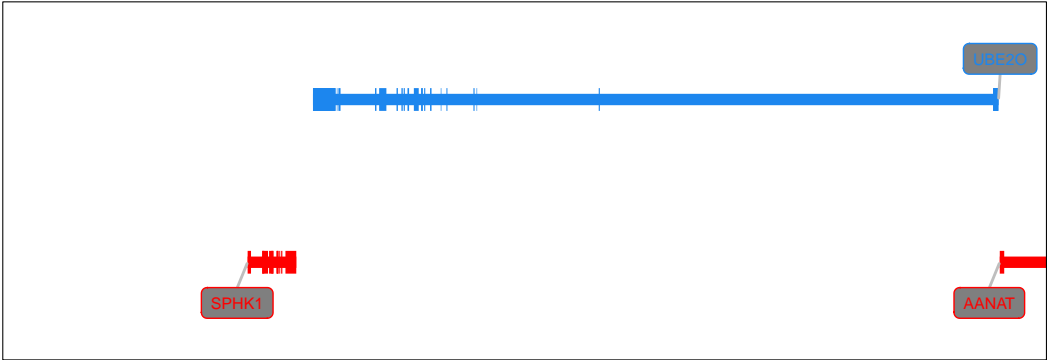
Coverage
(Norm. ATAC Signal Range (0–0.96) by ReadsInTSS)



L5.6.NR4A2



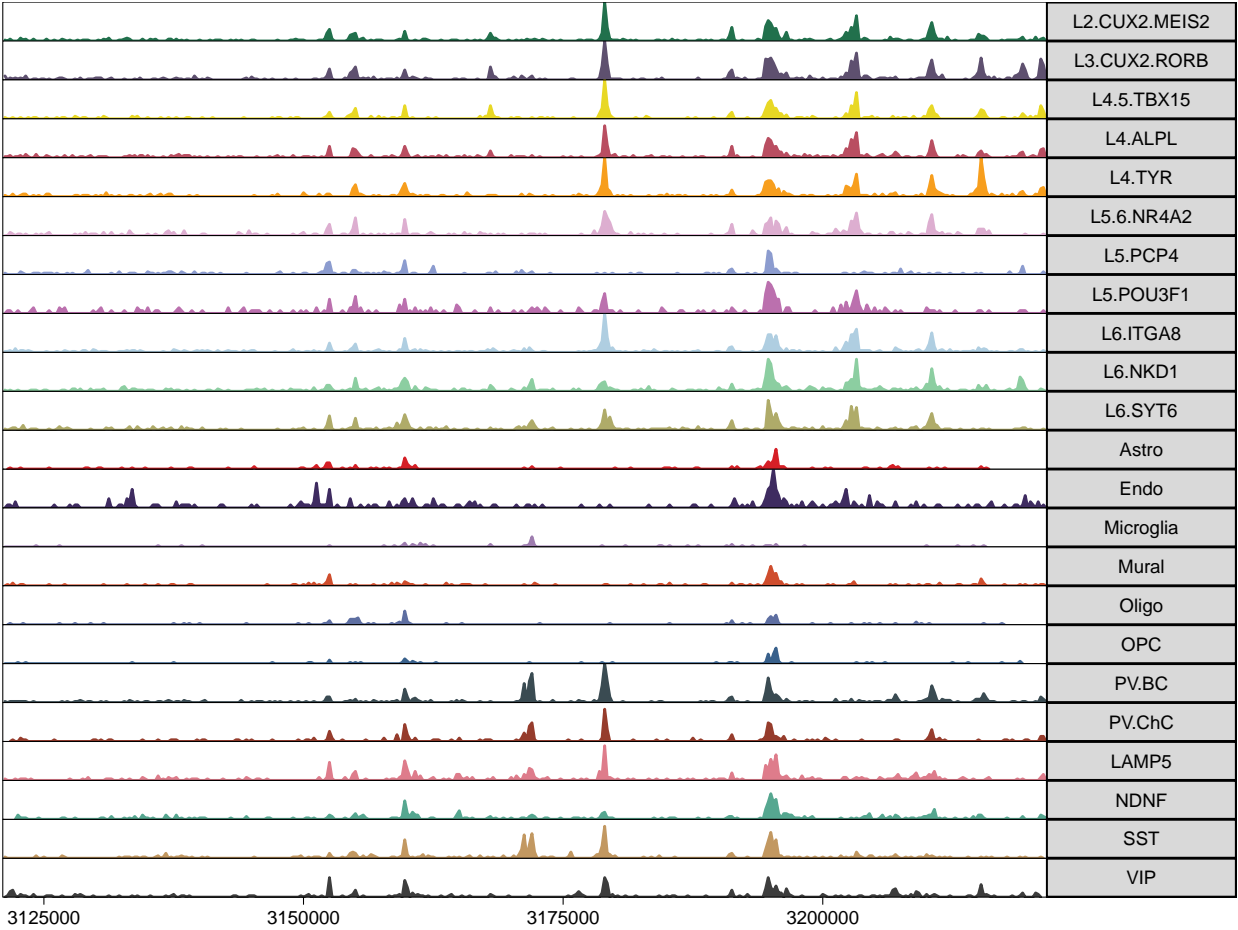
Peaks



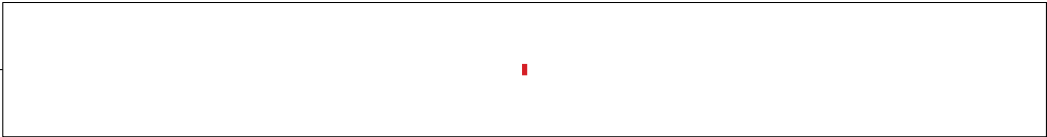
Genes

chr11:3121041–3221542 Marker Peak Tracks for: L5.6.NR4A2

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



L5.6.NR4A2

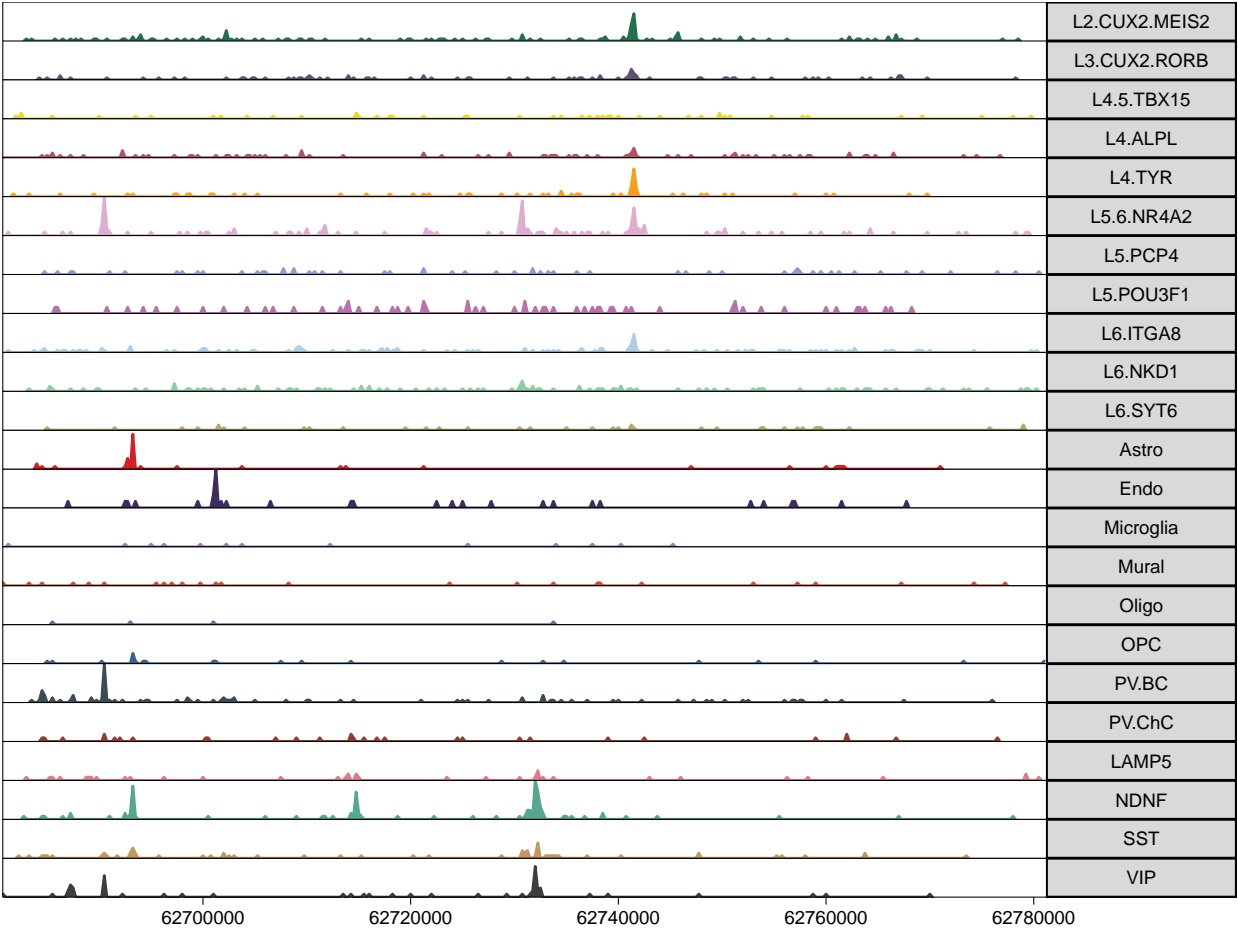


Peaks

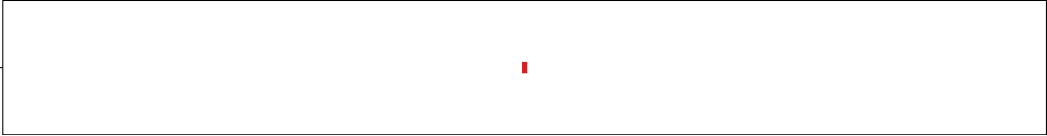


Genes

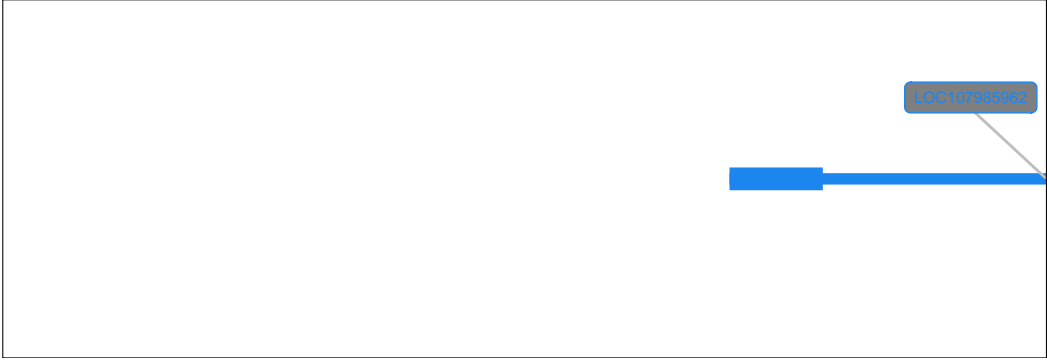
Coverage
(Norm. ATAC Signal Range (0–0.47) by ReadsInTSS)



L5.6.NR4A2

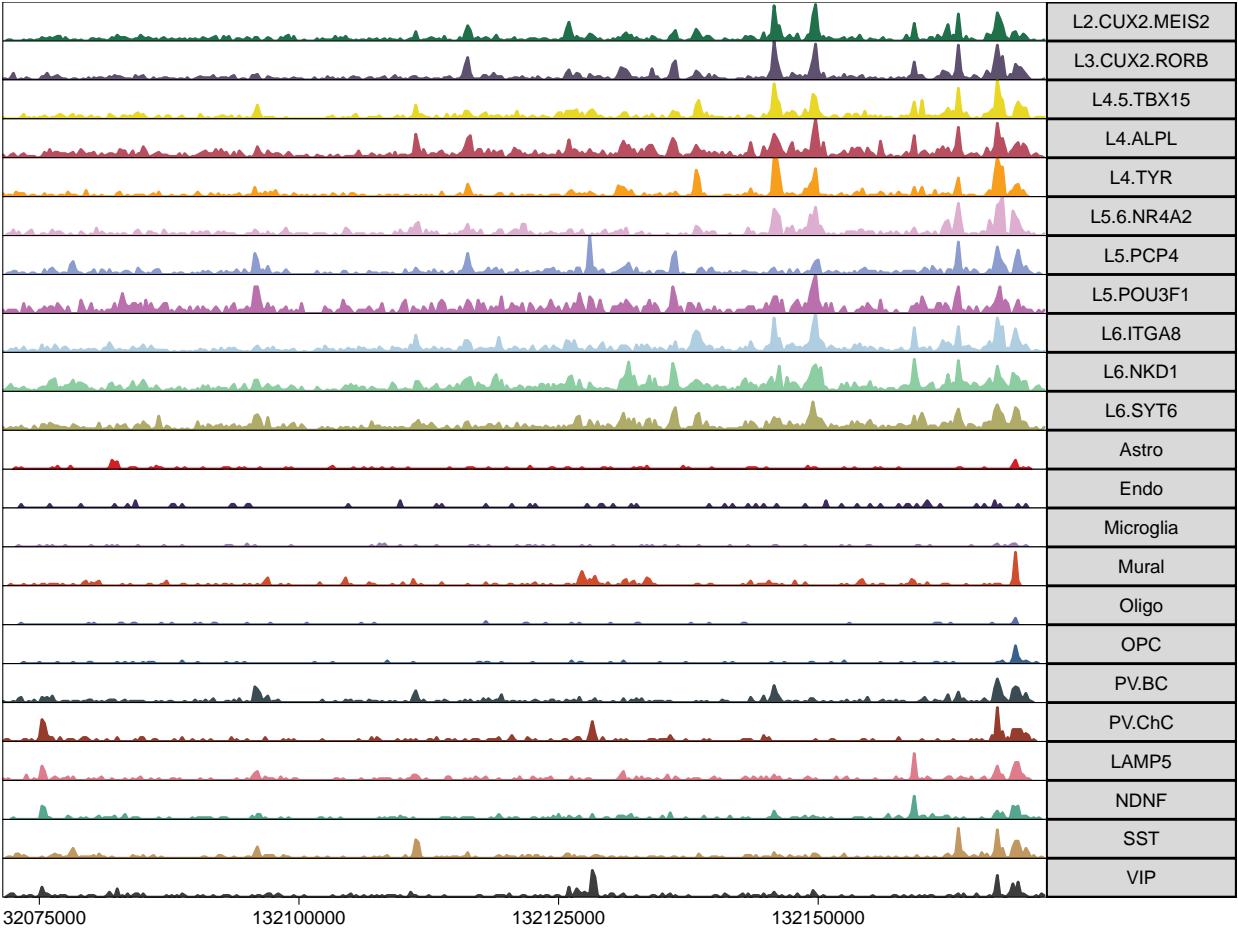


Peaks

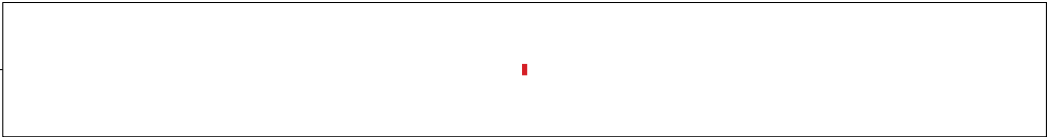


Genes

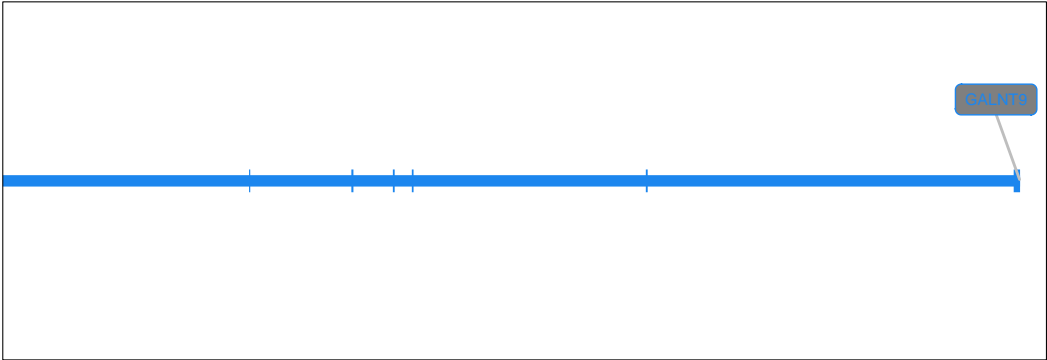
Coverage
(Norm. ATAC Signal Range (0–0.86) by ReadsInTSS)



L5.6.NR4A2



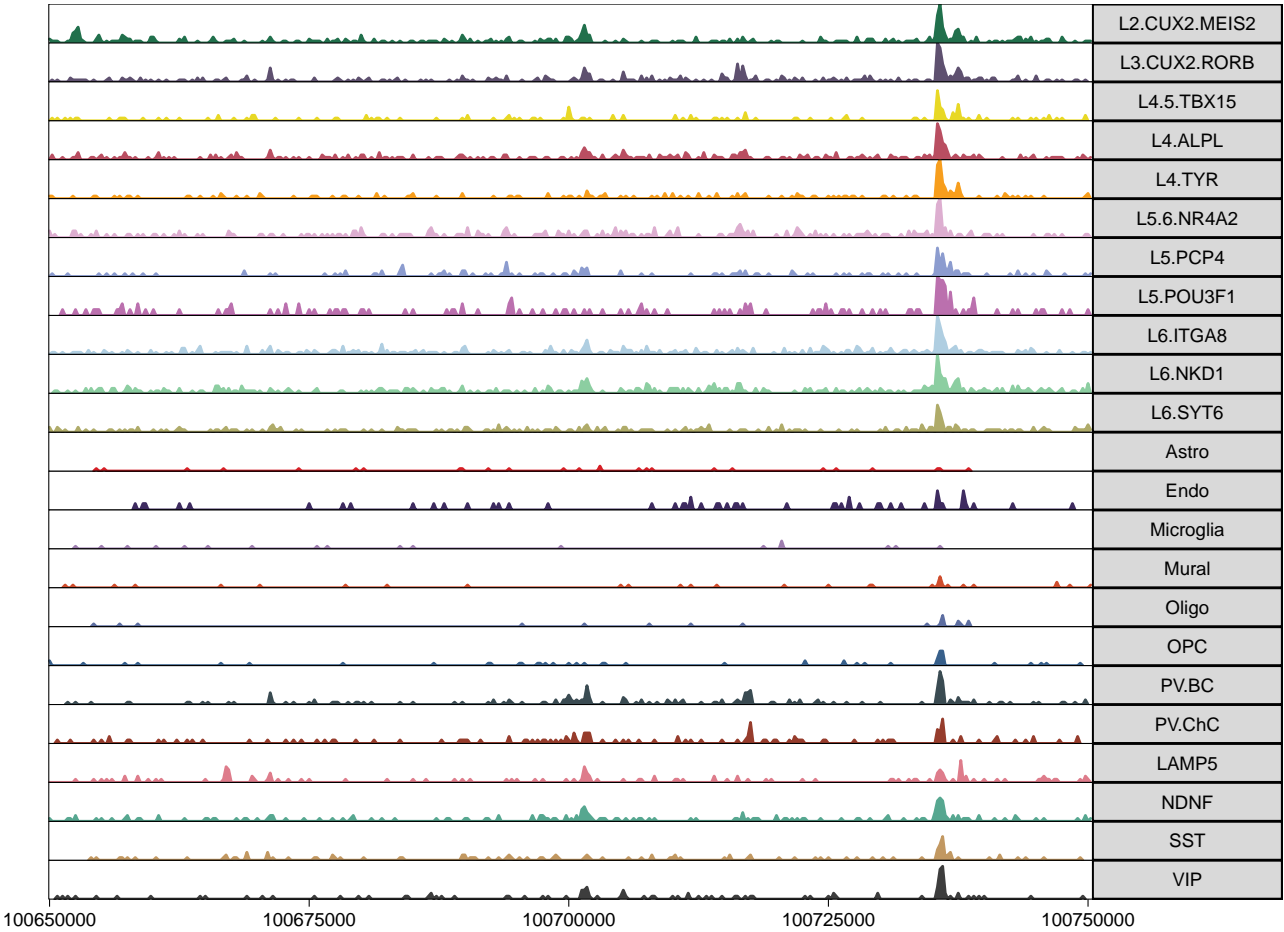
Peaks



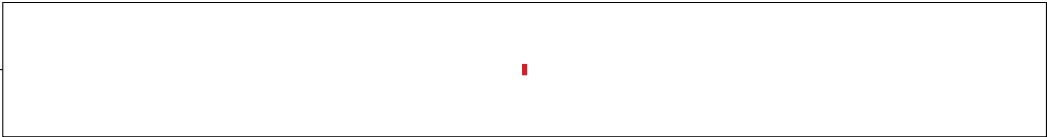
Genes

chr4:100649925–100750426 Marker Peak Tracks for: L5.6.NR4A2

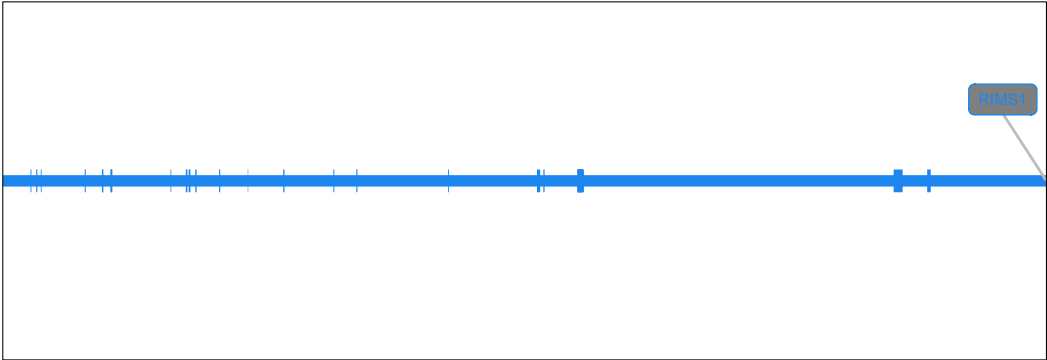
Coverage
(Norm. ATAC Signal Range (0–0.49) by ReadsInTSS)



L5.6.NR4A2

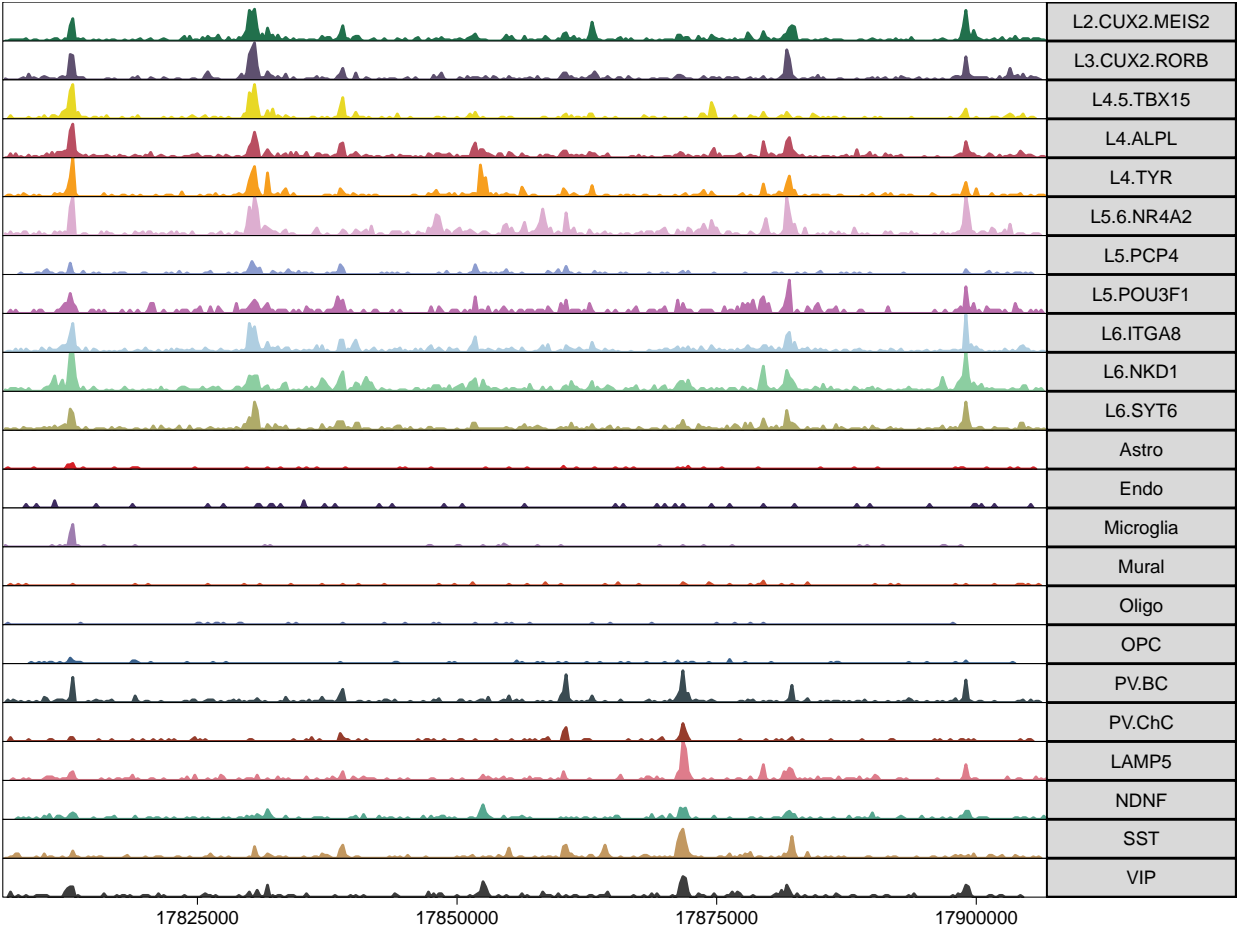


Peaks

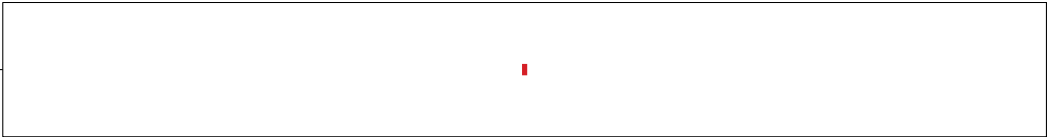


Genes

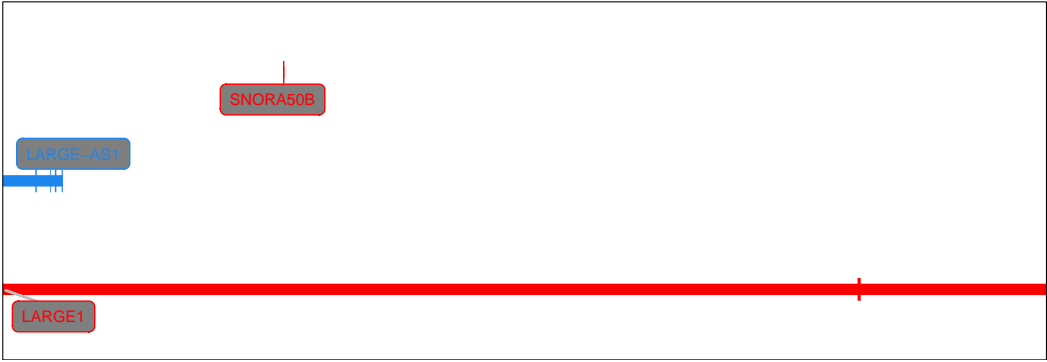
Coverage
(Norm. ATAC Signal Range (0–0.86) by ReadsInTSS)



L5.6.NR4A2

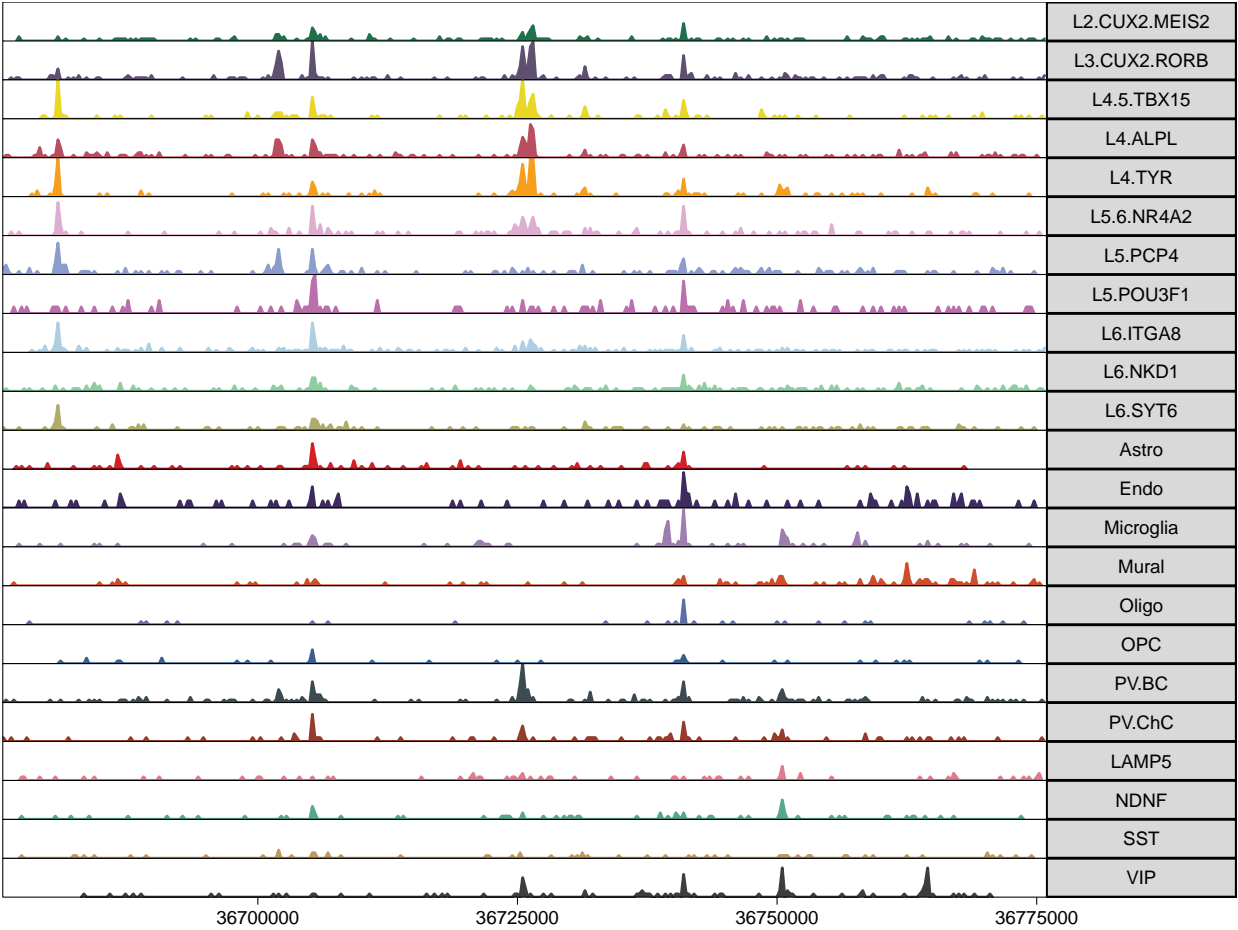


Peaks

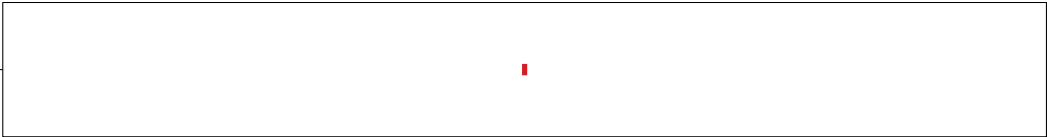


Genes

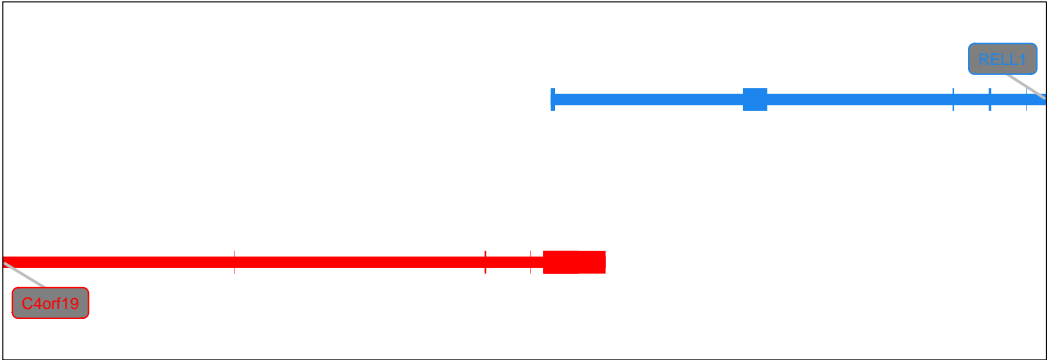
Coverage
(Norm. ATAC Signal Range (0–0.44) by ReadsInTSS)



L5.6.NR4A2



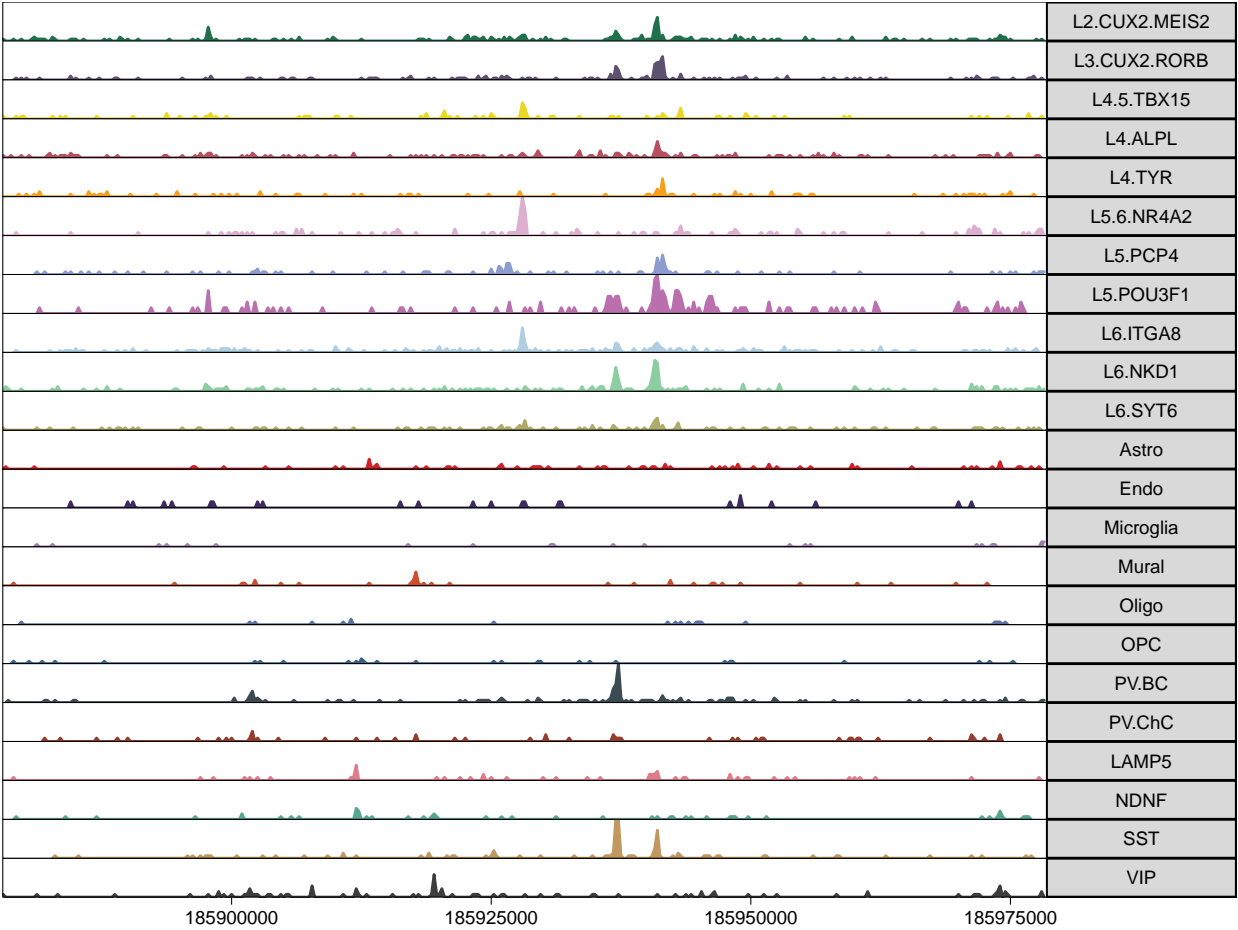
Peaks



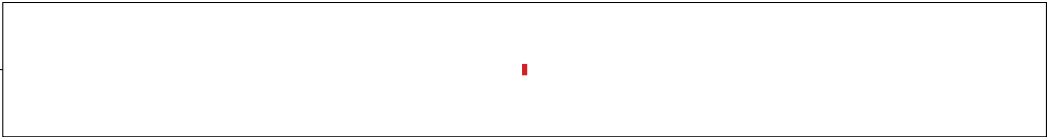
Genes

chr1:185877966–185978467 Marker Peak Tracks for: L5.6.NR4A2

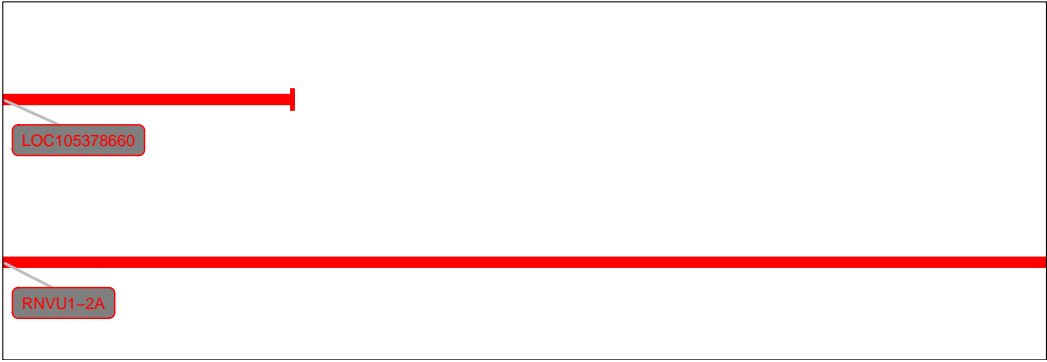
Coverage
(Norm. ATAC Signal Range (0–0.5) by ReadsInTSS)



L5.6.NR4A2



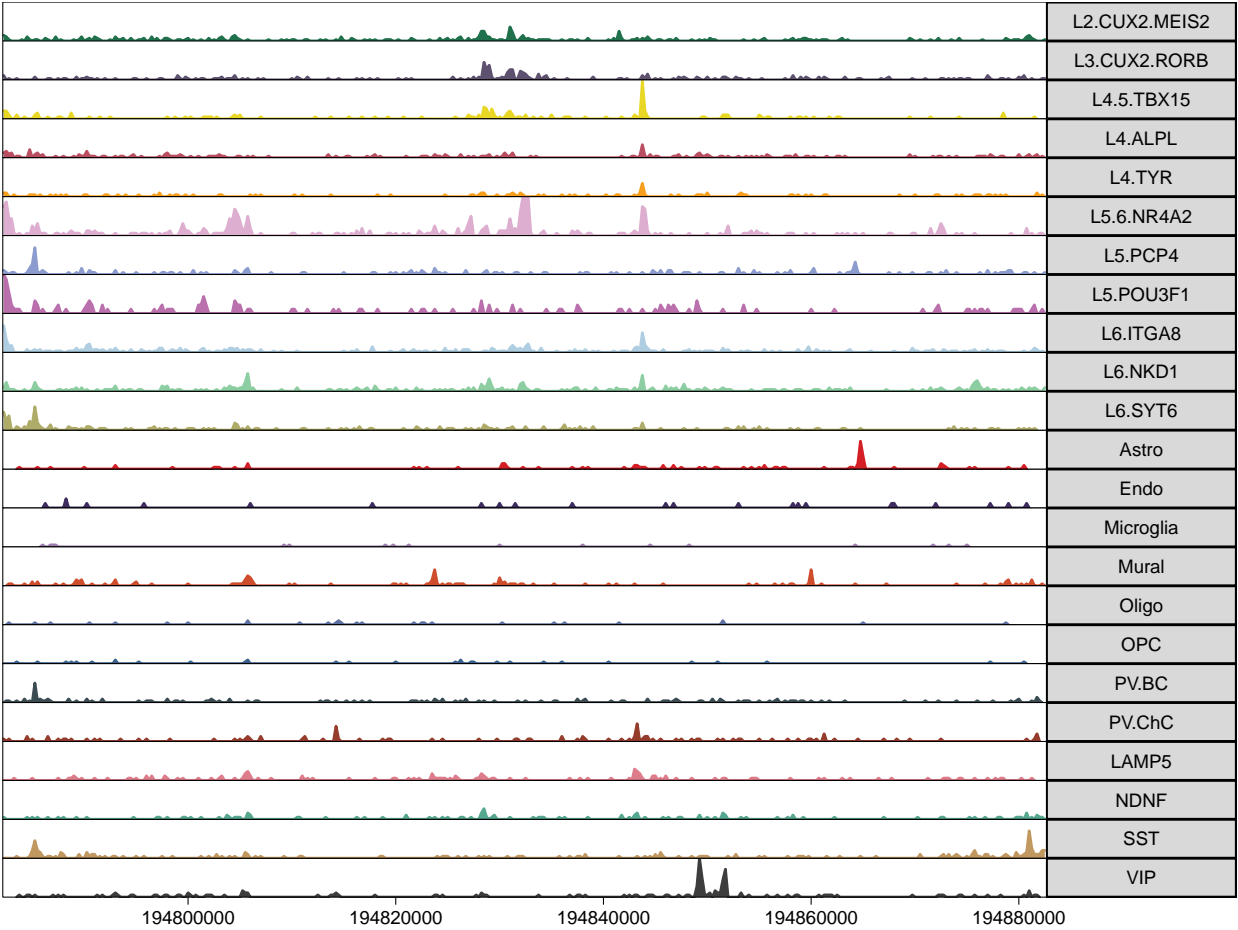
Peaks



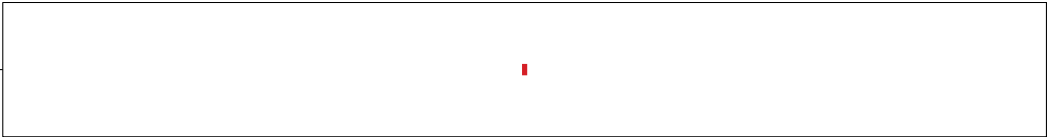
Genes

chr1:194782159–194882660 Marker Peak Tracks for: L5.6.NR4A2

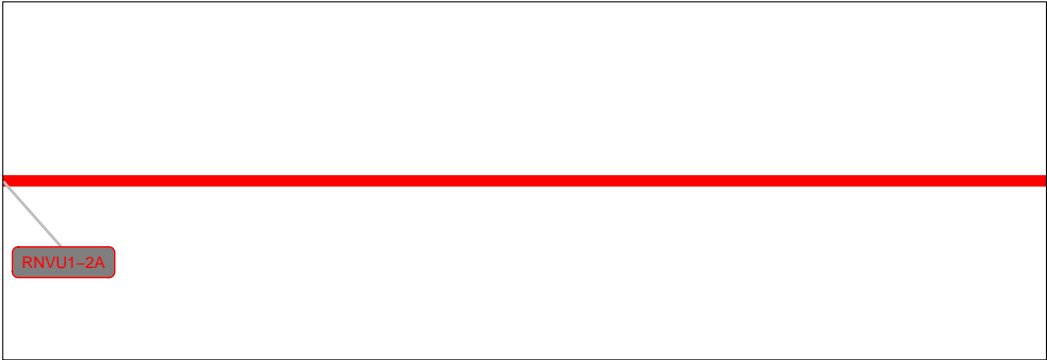
Coverage
(Norm. ATAC Signal Range (0–0.67) by ReadsInTSS)



L5.6.NR4A2



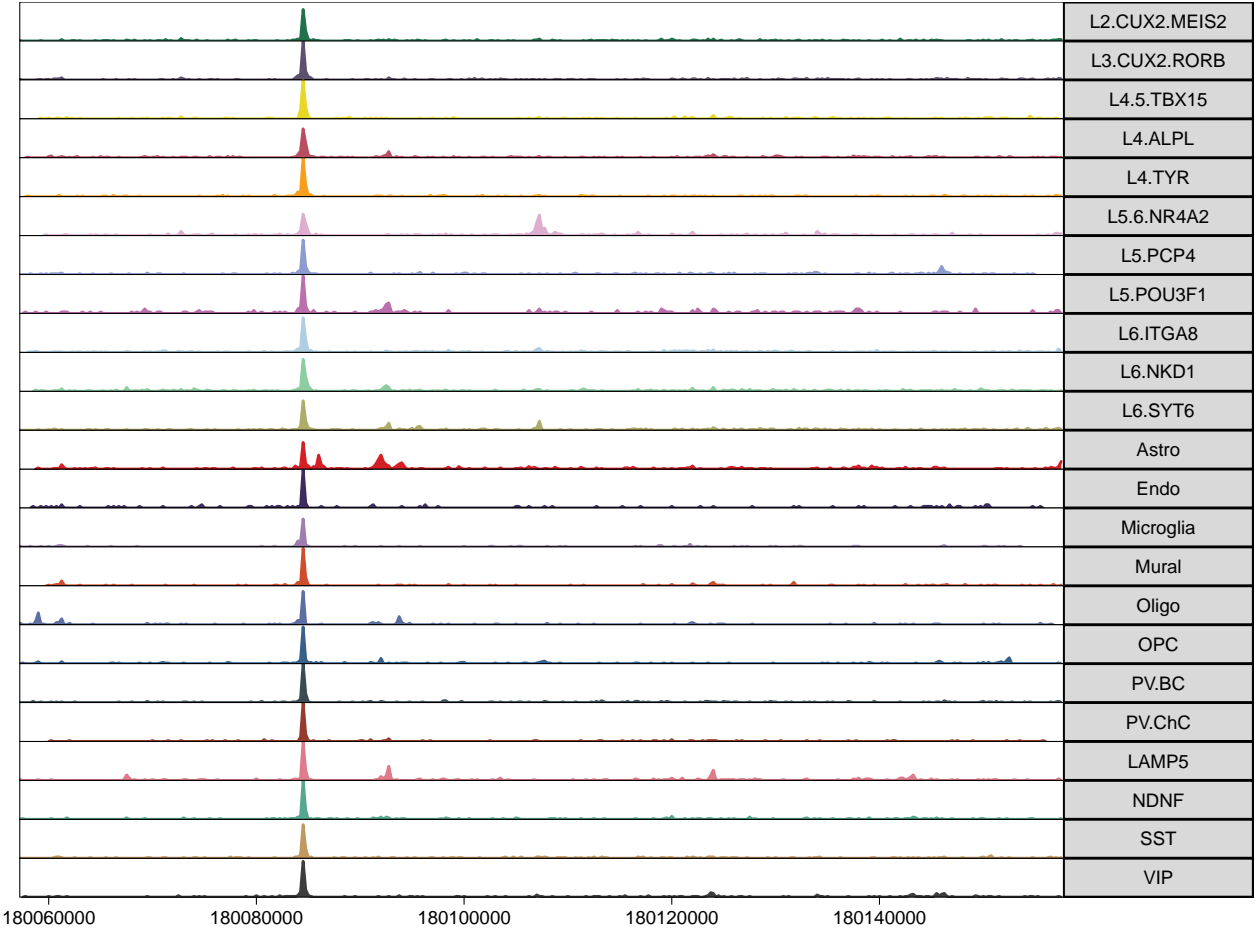
Peaks



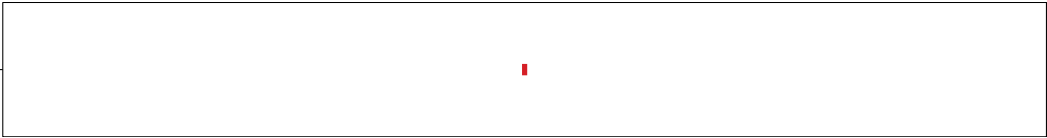
Genes

chr1:180057216–180157717 Marker Peak Tracks for: L5.6.NR4A2

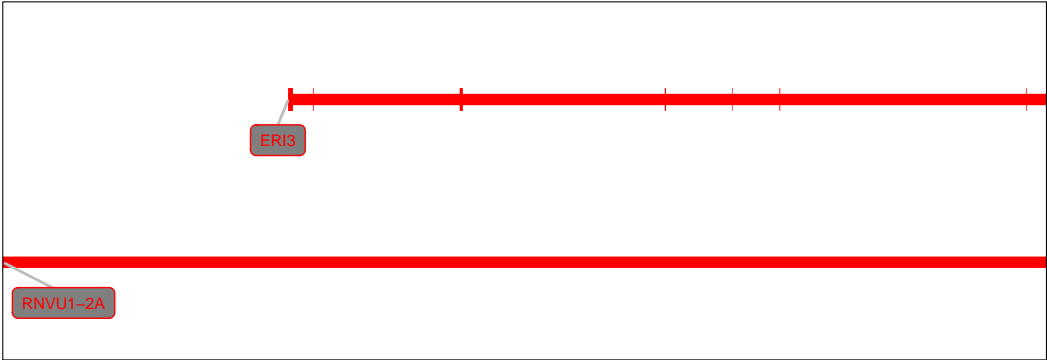
Coverage
(Norm. ATAC Signal Range (0–1.9) by ReadsInTSS)



L5.6.NR4A2



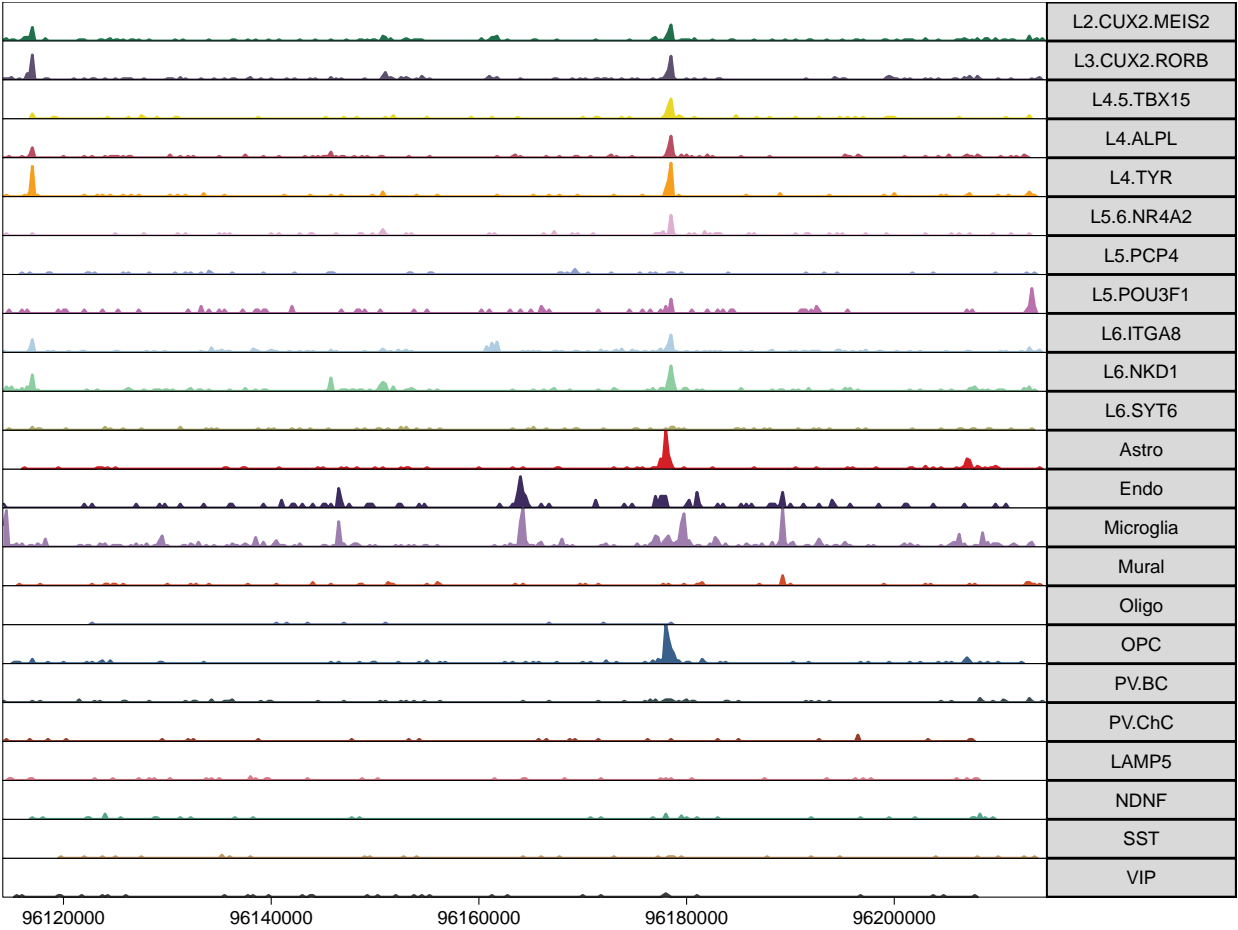
Peaks



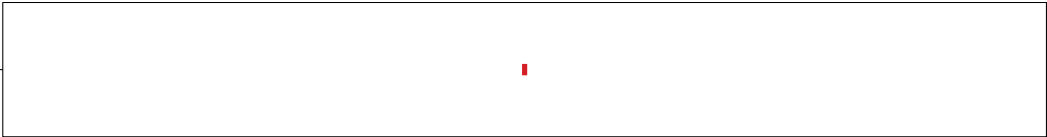
Genes

chr9:96114152–96214653 Marker Peak Tracks for: L5.6.NR4A2

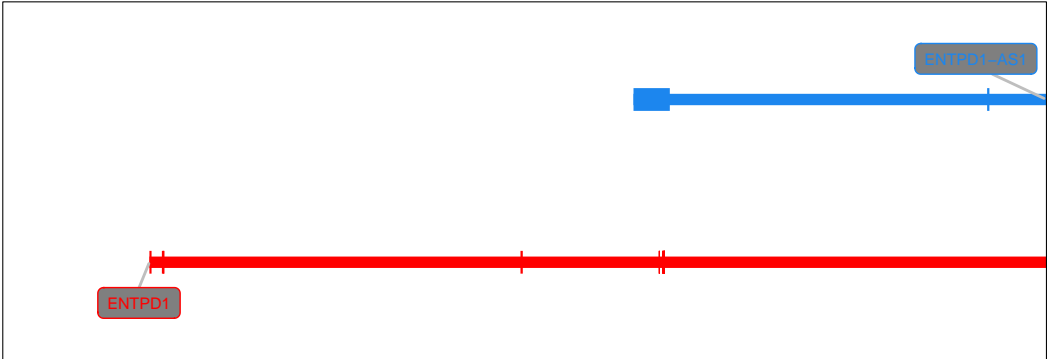
Coverage
(Norm. ATAC Signal Range (0–0.8) by ReadsInTSS)



L5.6.NR4A2

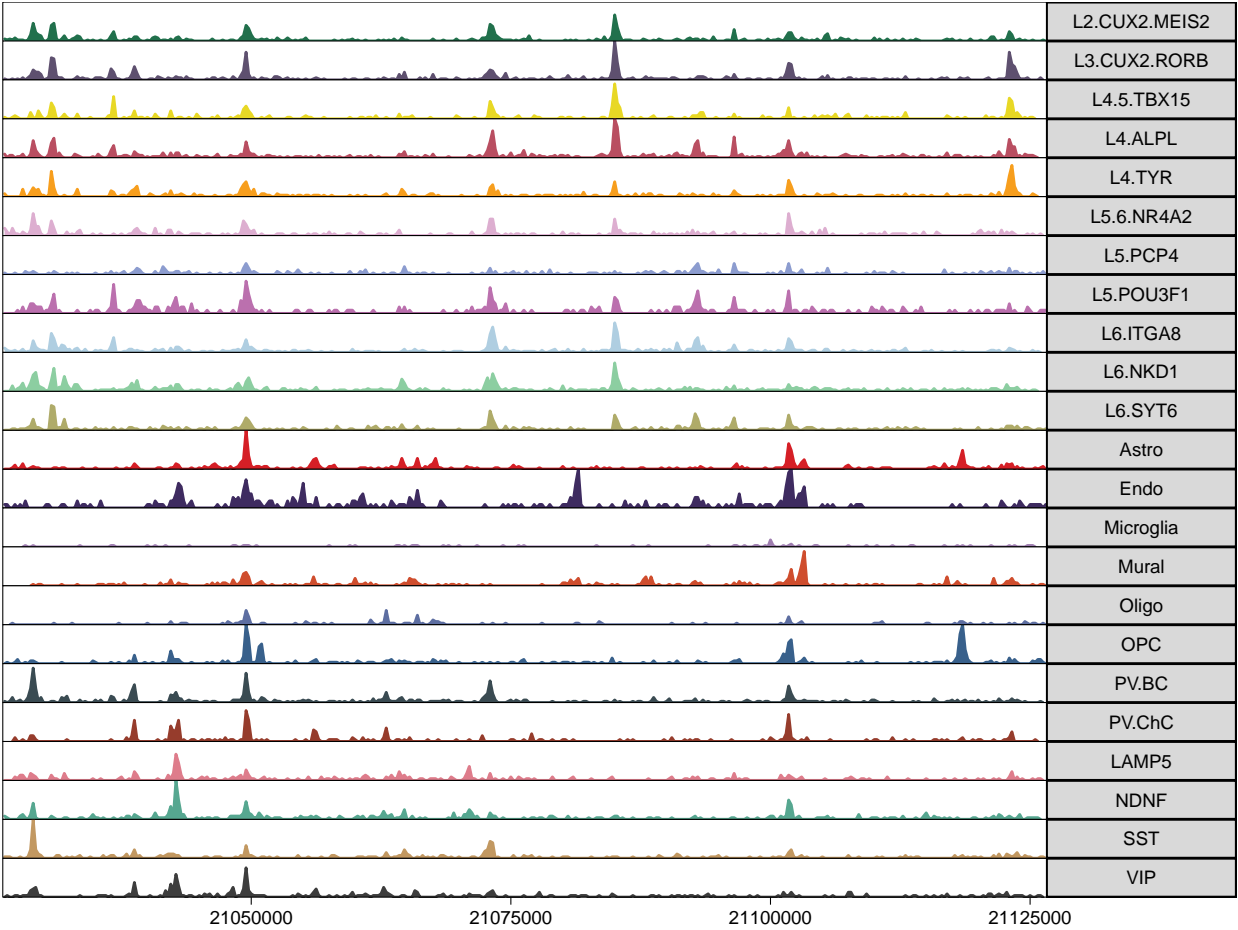


Peaks

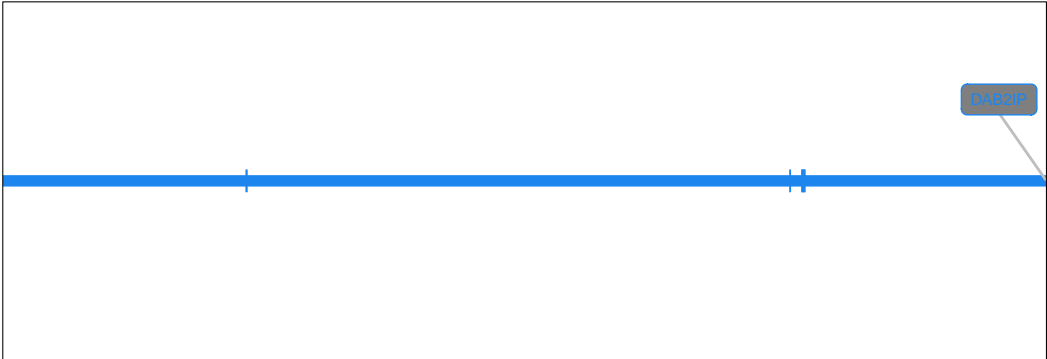
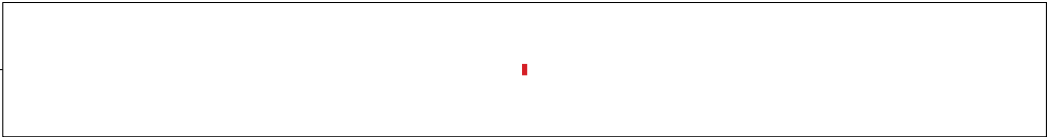


Genes

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

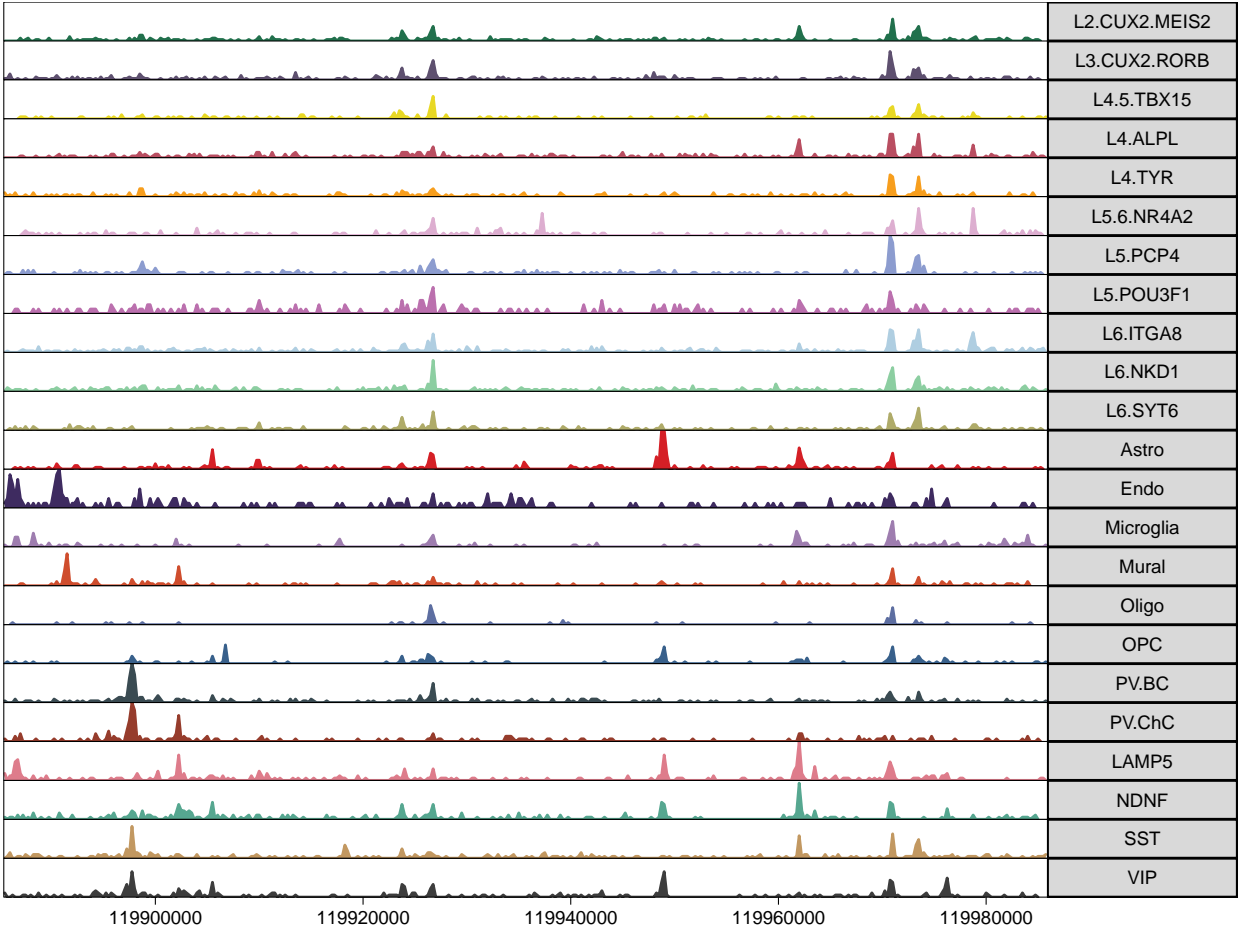


L5.6.NR4A2

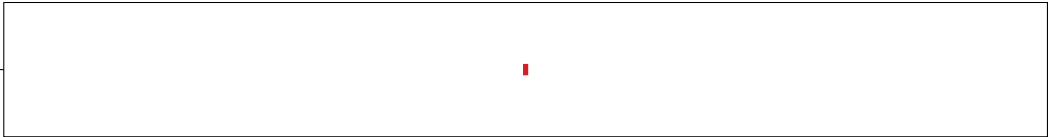


chr9:119885404–119985905 Marker Peak Tracks for: L5.6.NR4A2

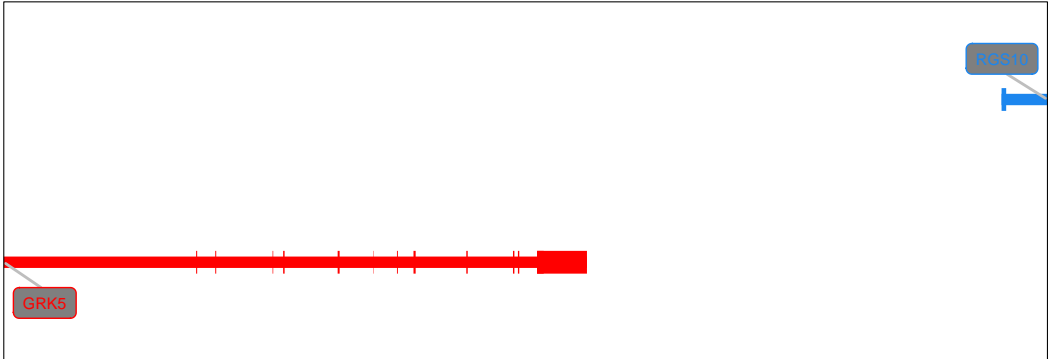
Coverage
(Norm. ATAC Signal Range (0–0.66) by ReadsInTSS)



L5.6.NR4A2



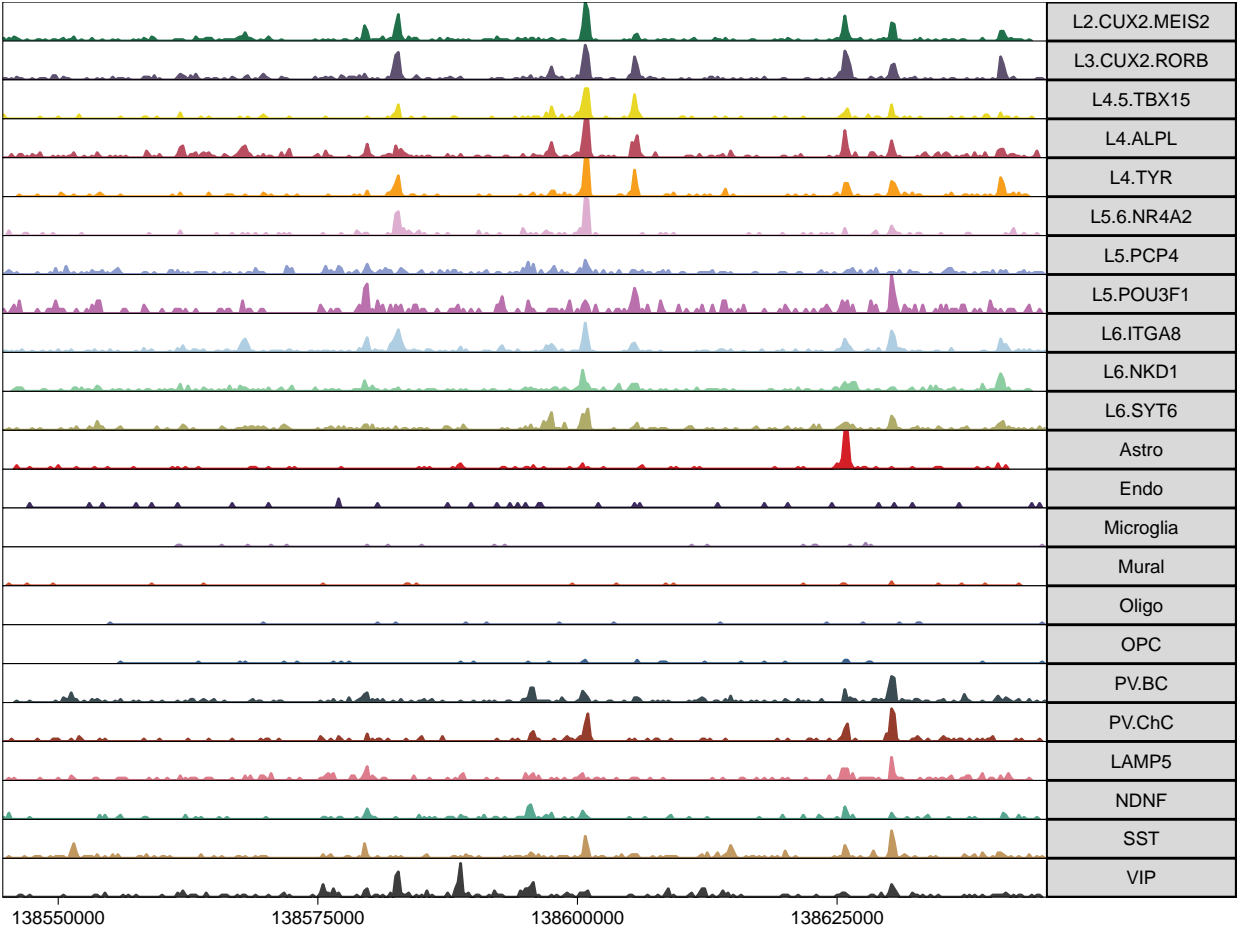
Peaks



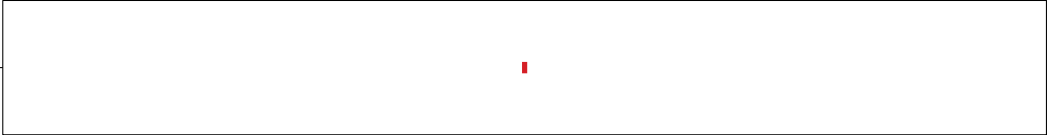
Genes

chr8:138544661–138645162 Marker Peak Tracks for: L5.6.NR4A2

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



L5.6.NR4A2

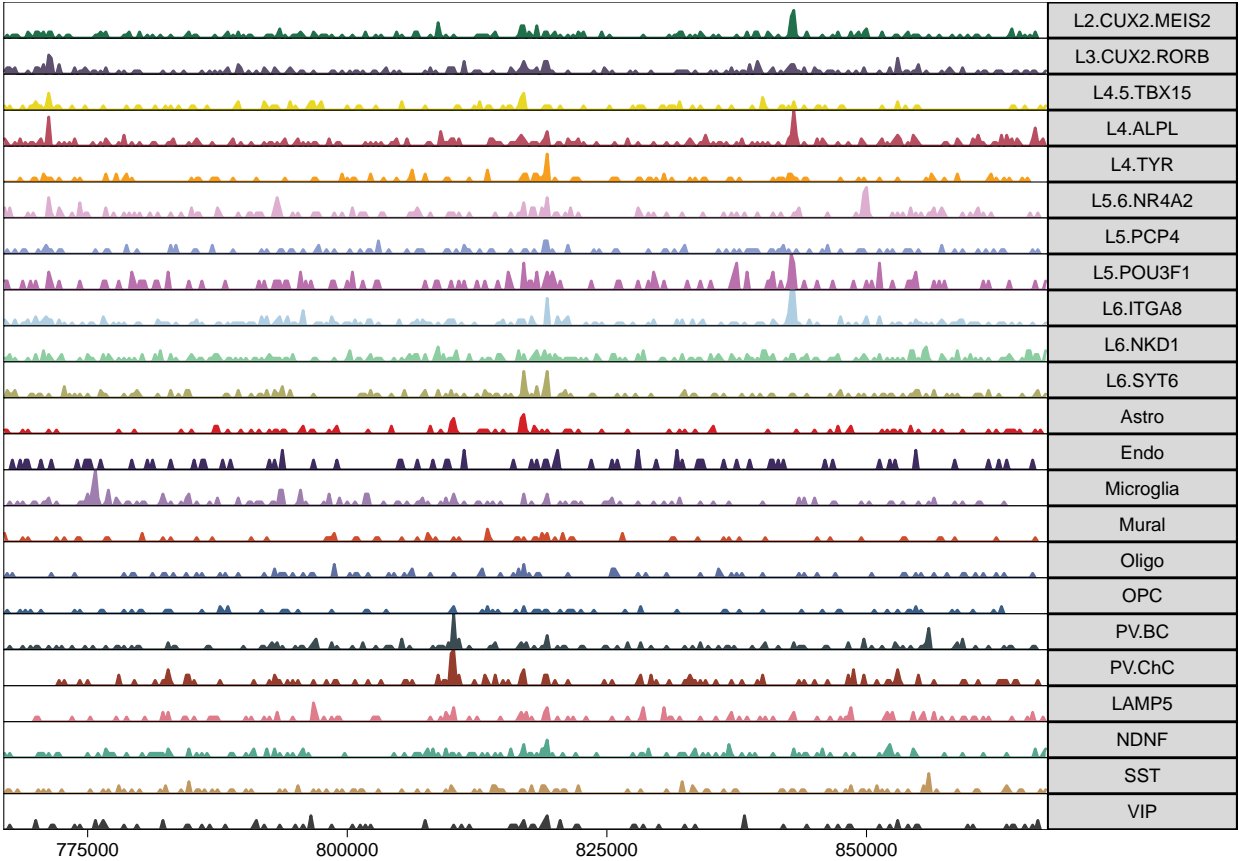


Peaks



Genes

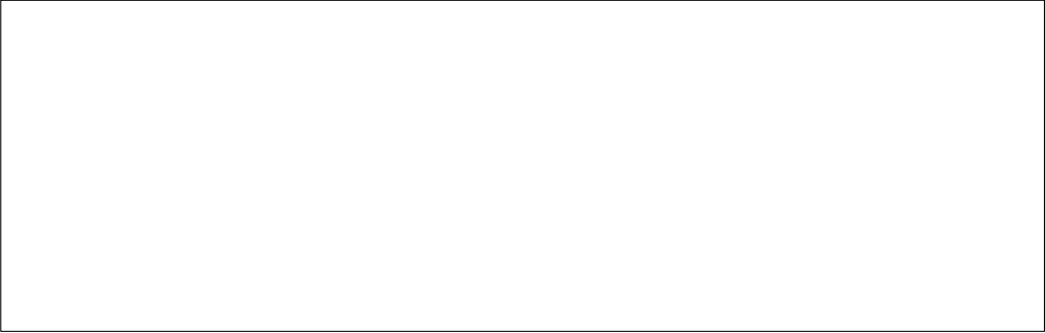
Coverage
(Norm. ATAC Signal Range (0–0.29) by ReadsInTSS)



L5.6.NR4A2

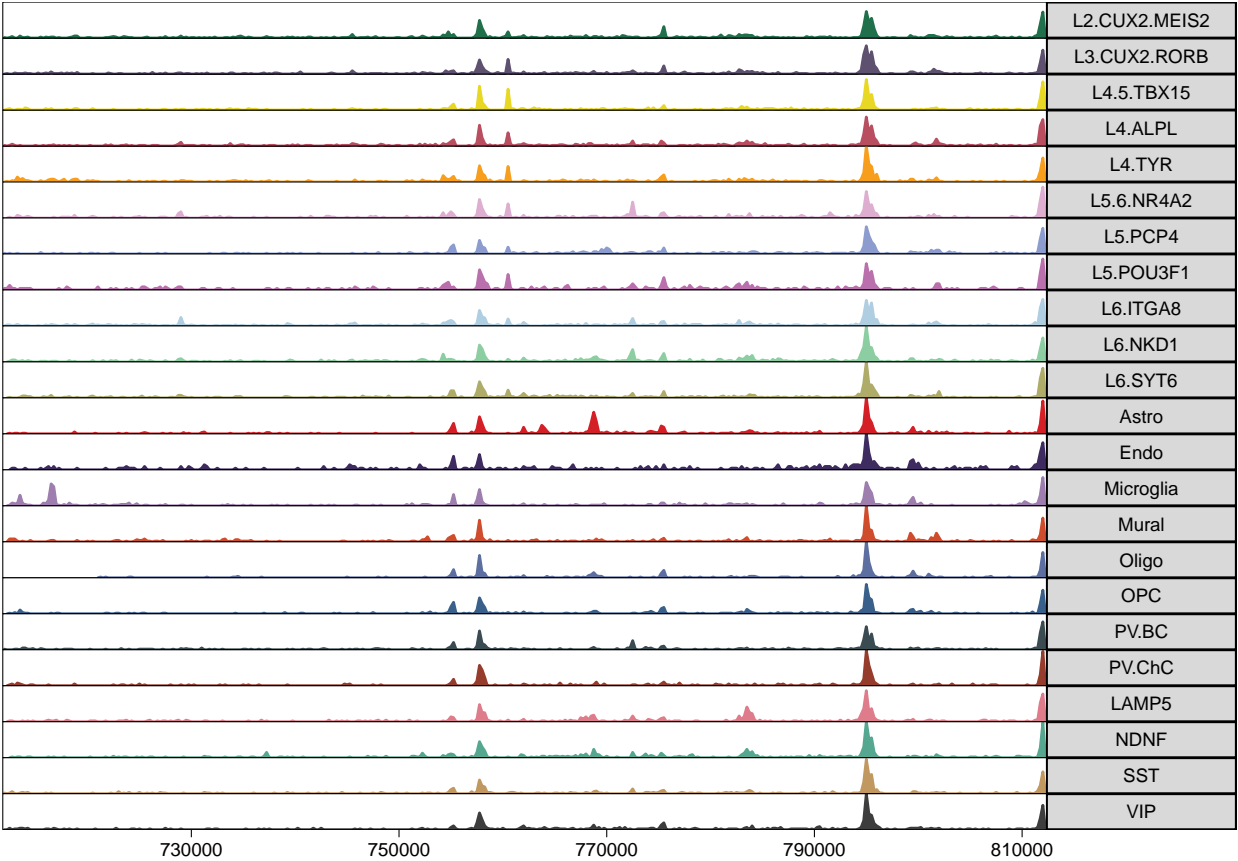


Peaks



GeneTrack

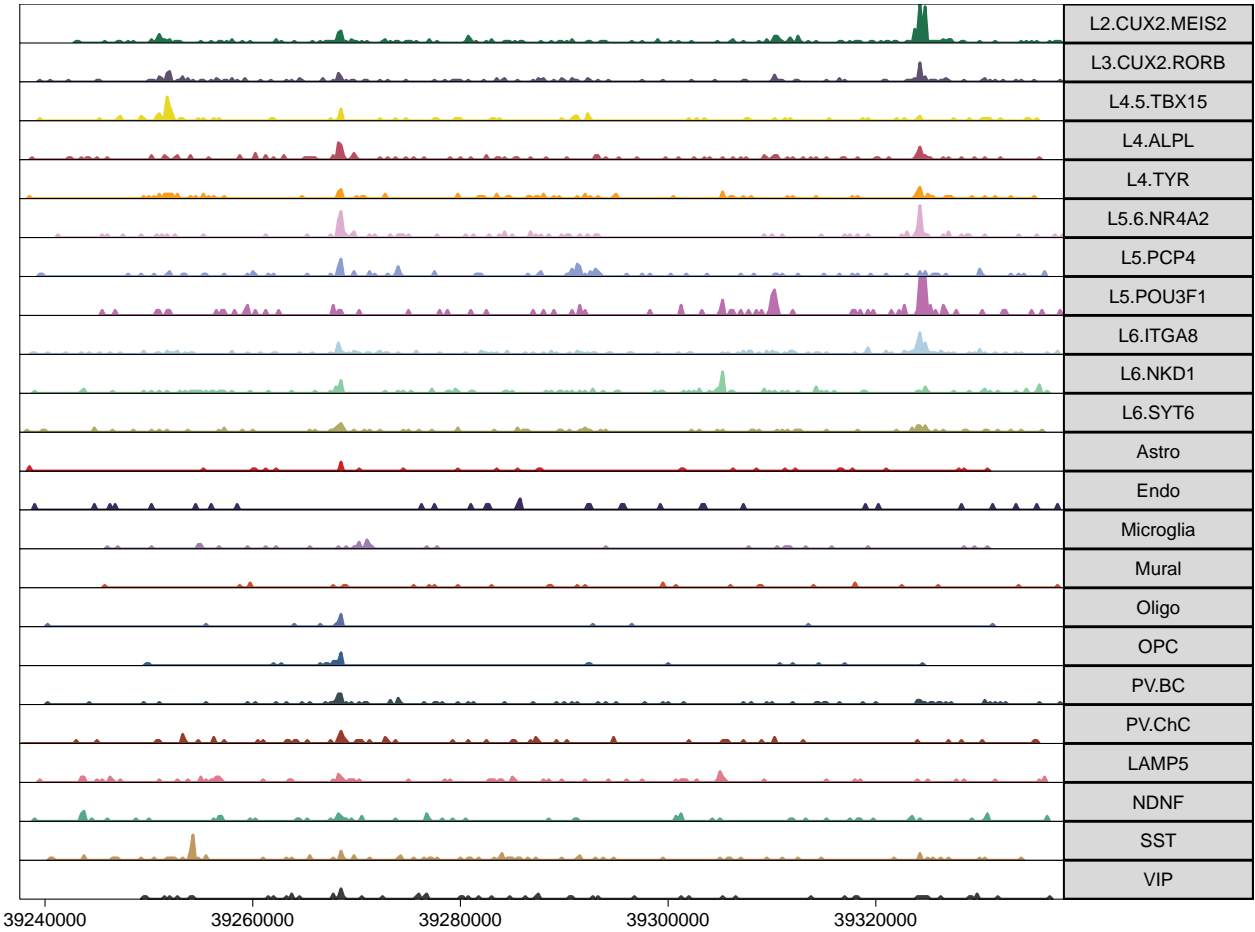
Coverage
(Norm. ATAC Signal Range (0–1.7) by ReadsInTSS)



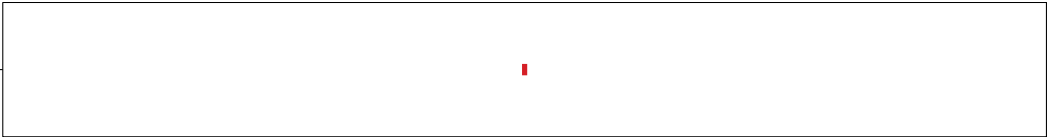
L5.6.NR4A2



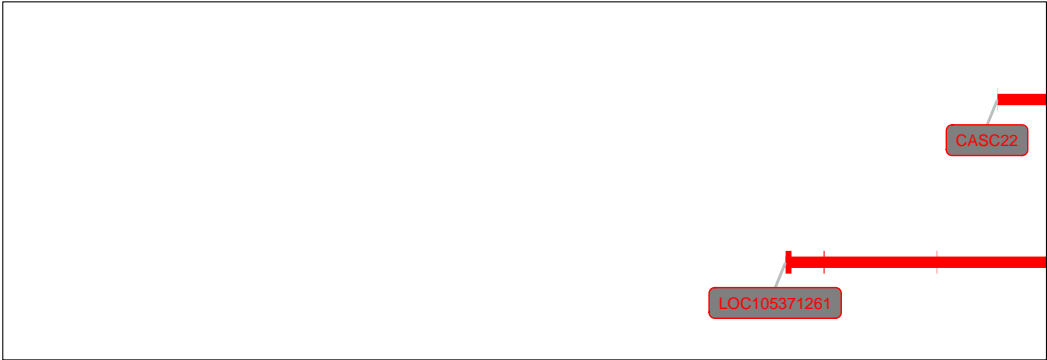
Coverage
(Norm. ATAC Signal Range (0–0.56) by ReadsInTSS)



L5.6.NR4A2

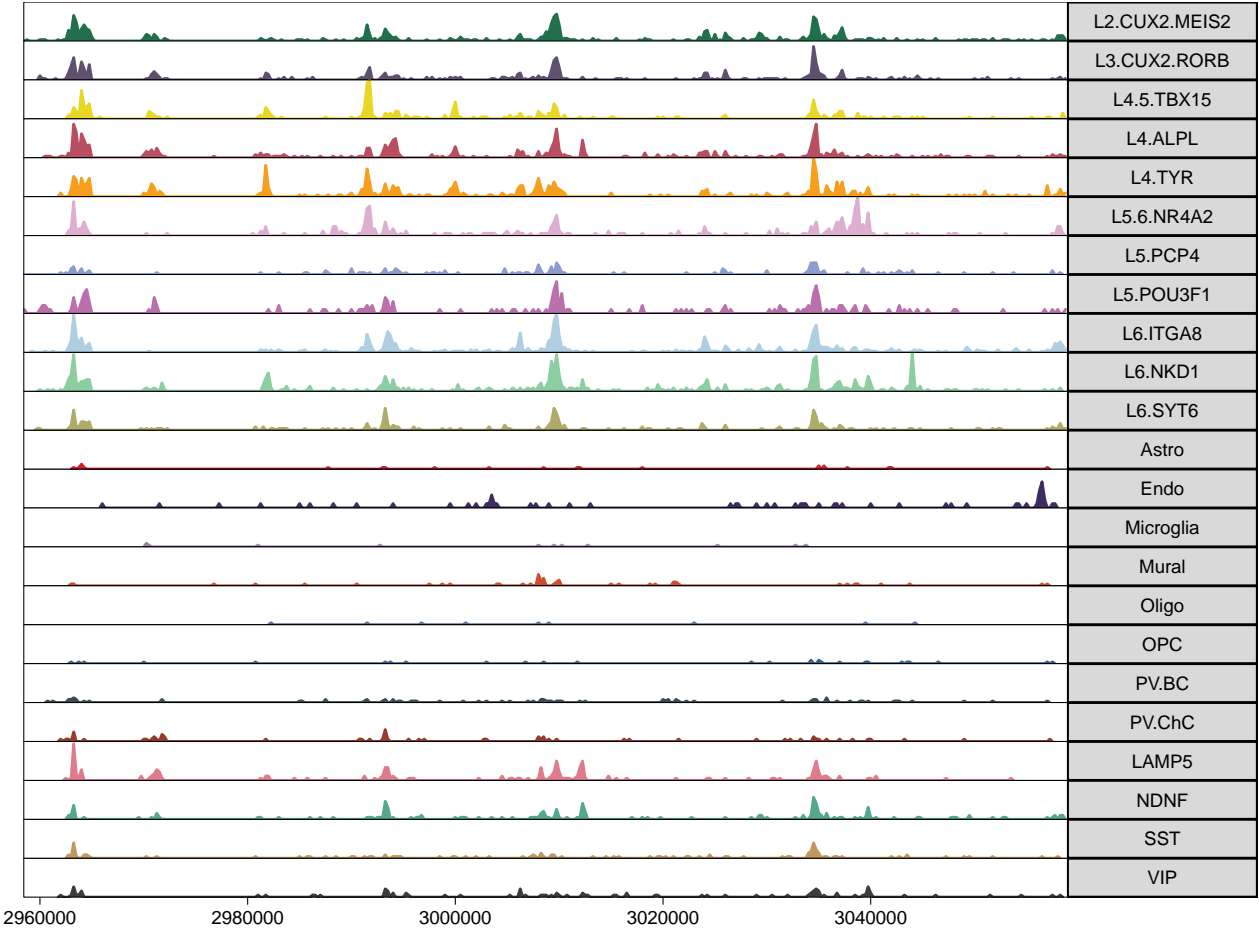


Peaks

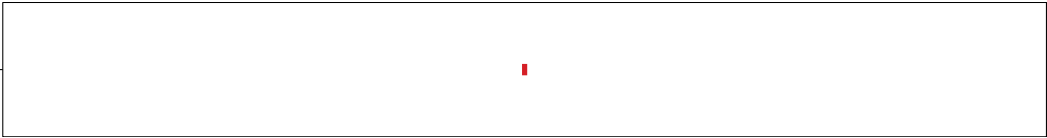


Genes

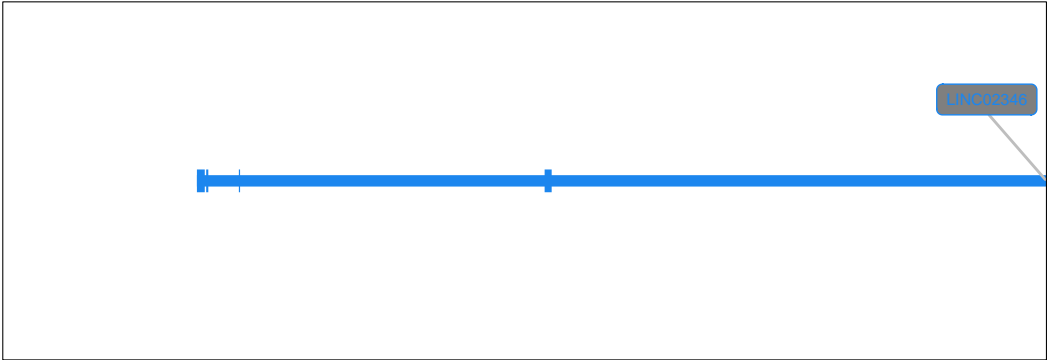
Coverage
(Norm. ATAC Signal Range (0–0.72) by ReadsInTSS)



L5.6.NR4A2

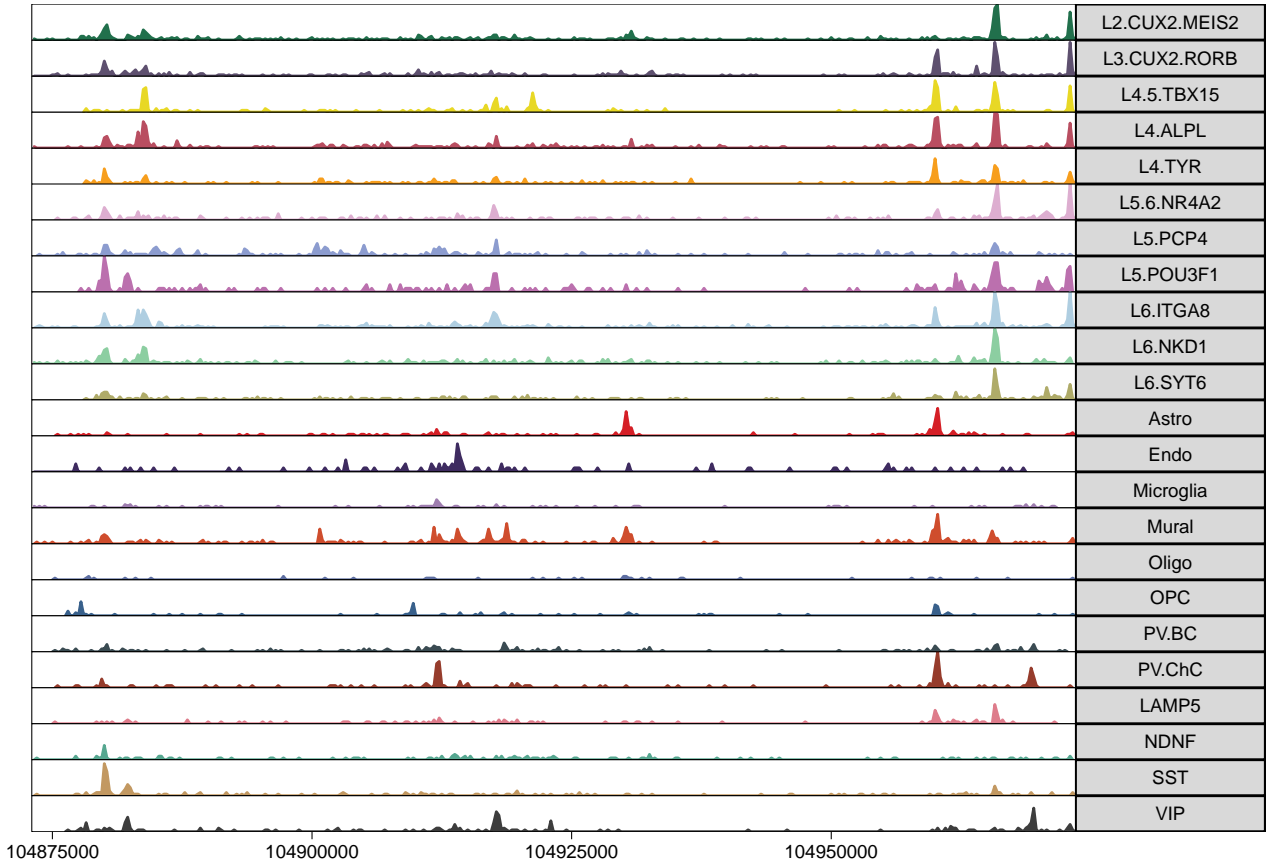


Peaks



Genes

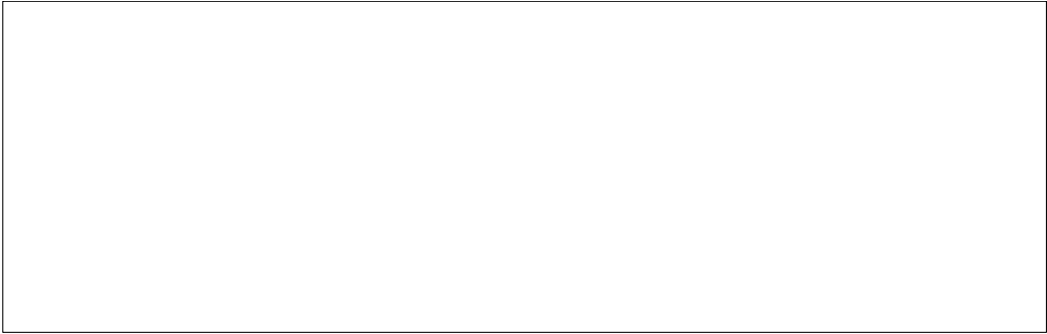
Coverage
(Norm. ATAC Signal Range (0–0.72) by ReadsInTSS)



L5.6.NR4A2

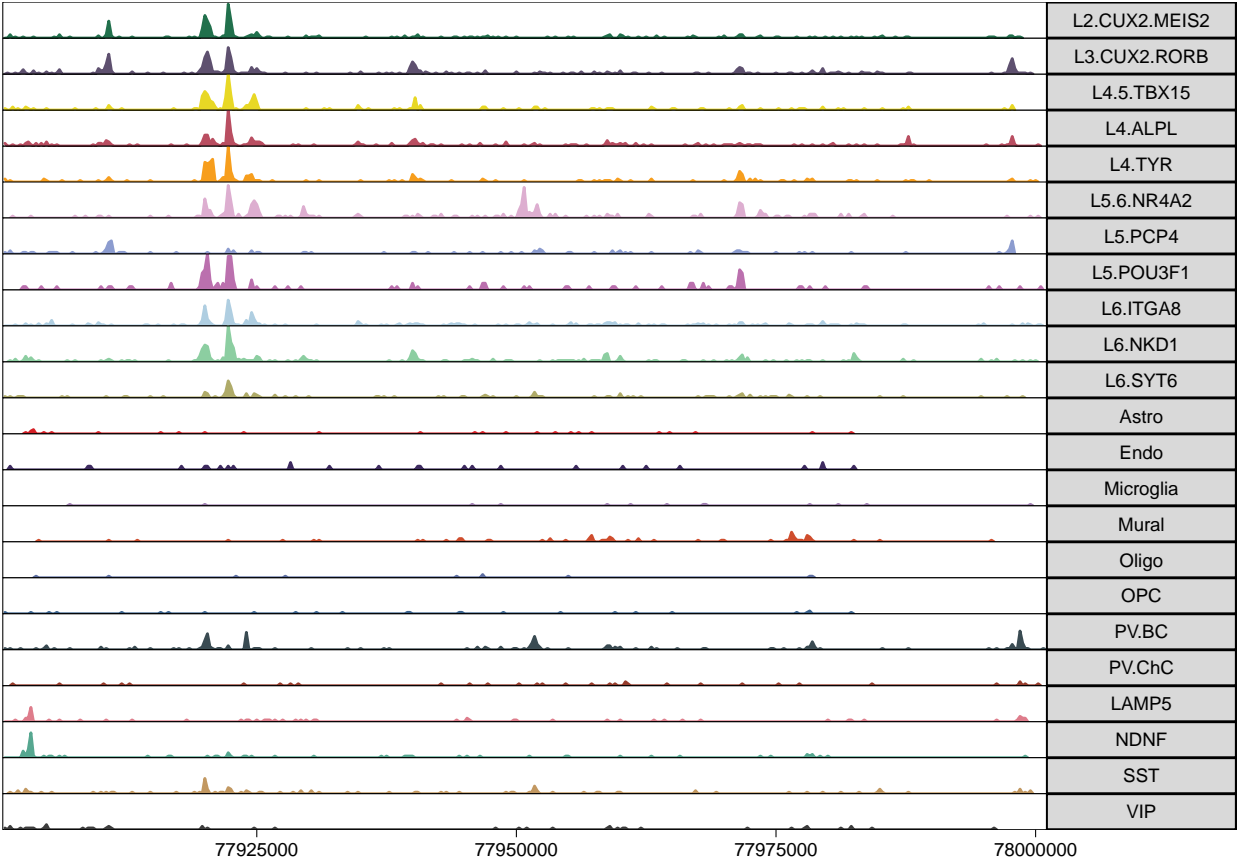


Peaks



GeneTrack

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



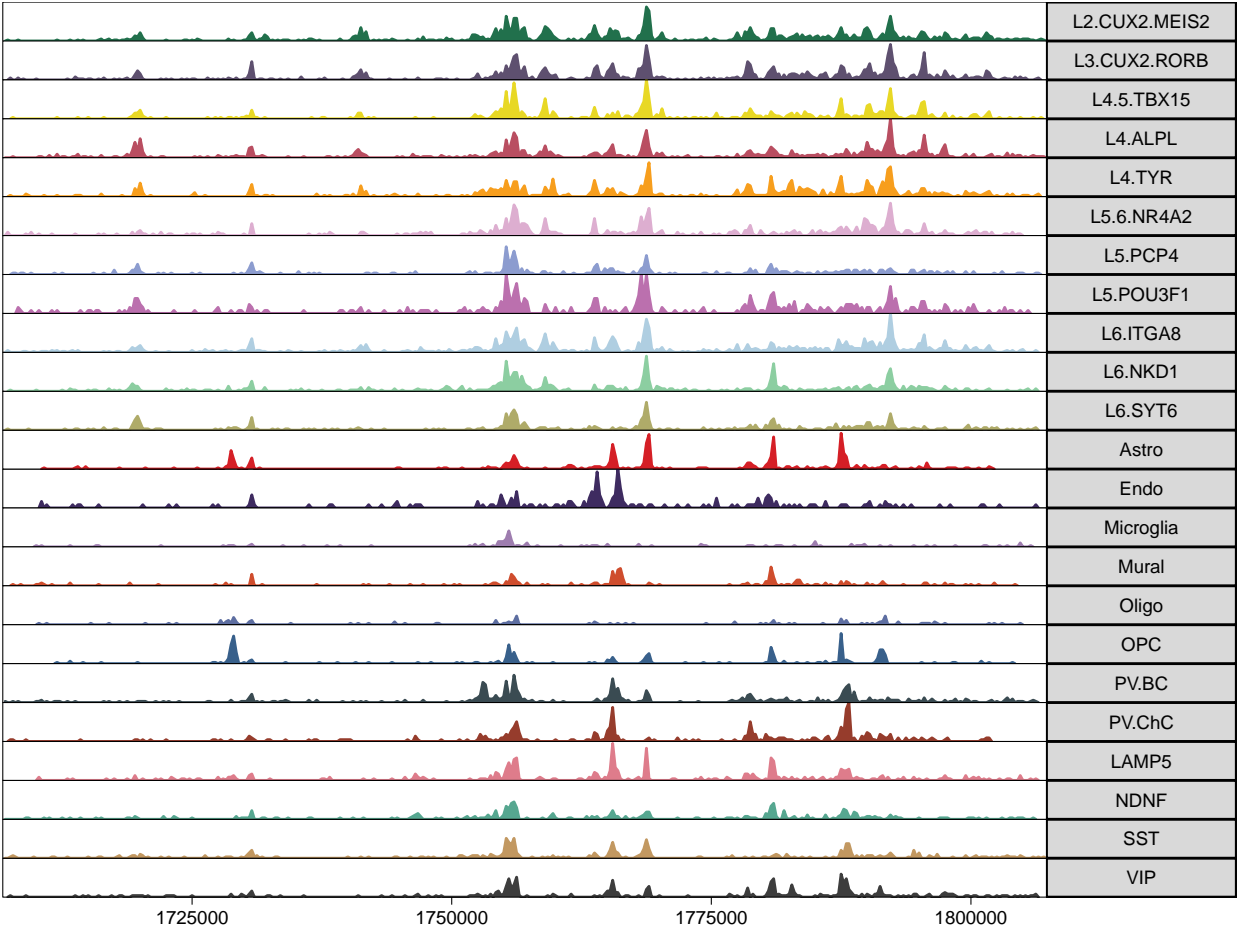
L5.6.NR4A2



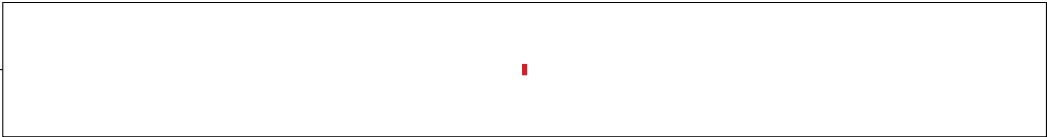
Peaks

GeneTrack

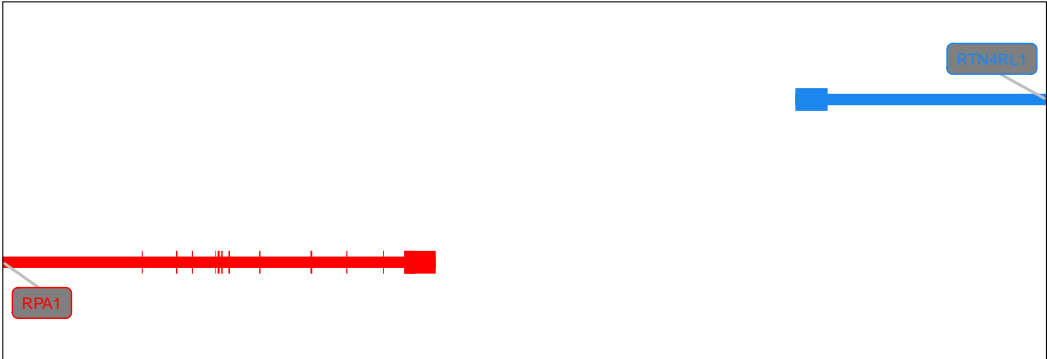
Coverage
(Norm. ATAC Signal Range (0–0.95) by ReadsInTSS)



L5.6.NR4A2

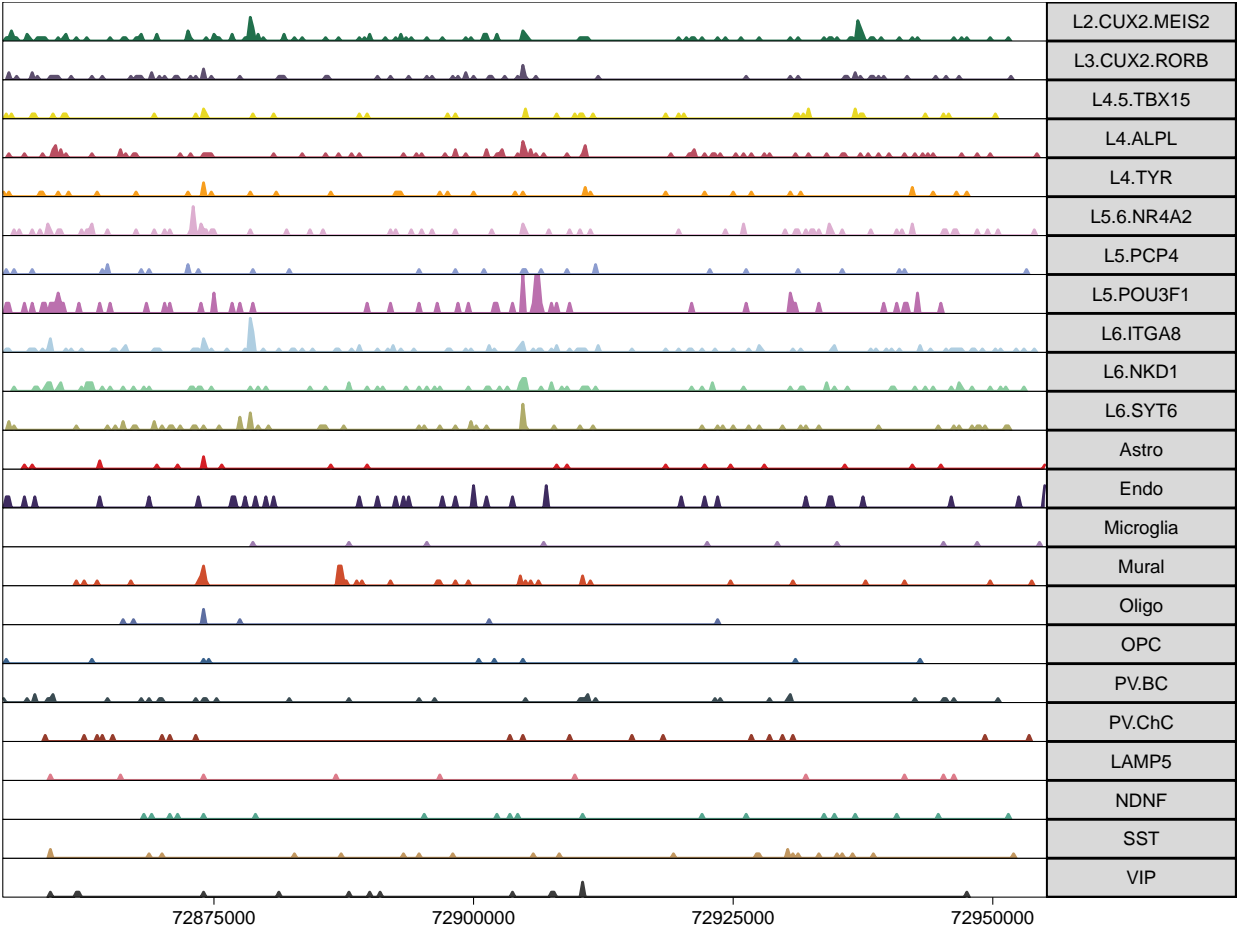


Peaks

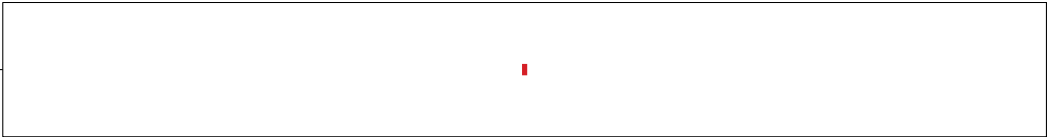


Genes

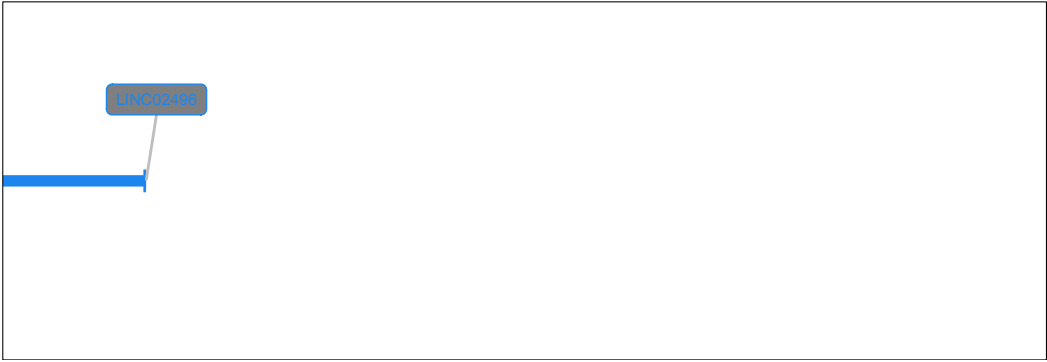
Coverage
(Norm. ATAC Signal Range (0–0.28) by ReadsInTSS)



L5.6.NR4A2

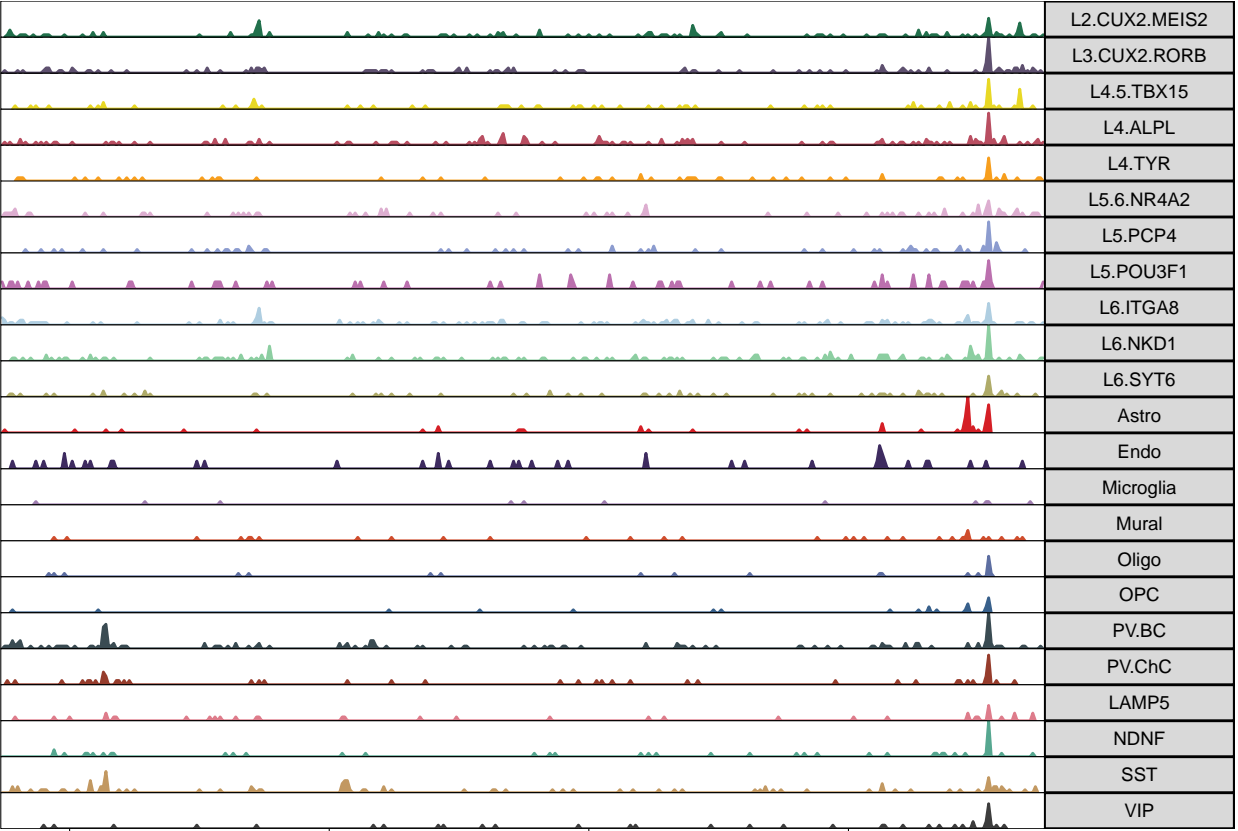


Peaks



Genes

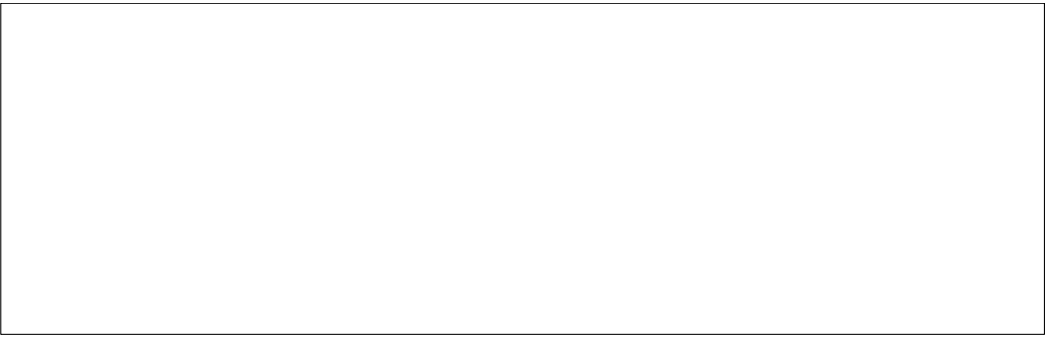
Coverage
(Norm. ATAC Signal Range (0–0.37) by ReadsInTSS)



L5.6.NR4A2

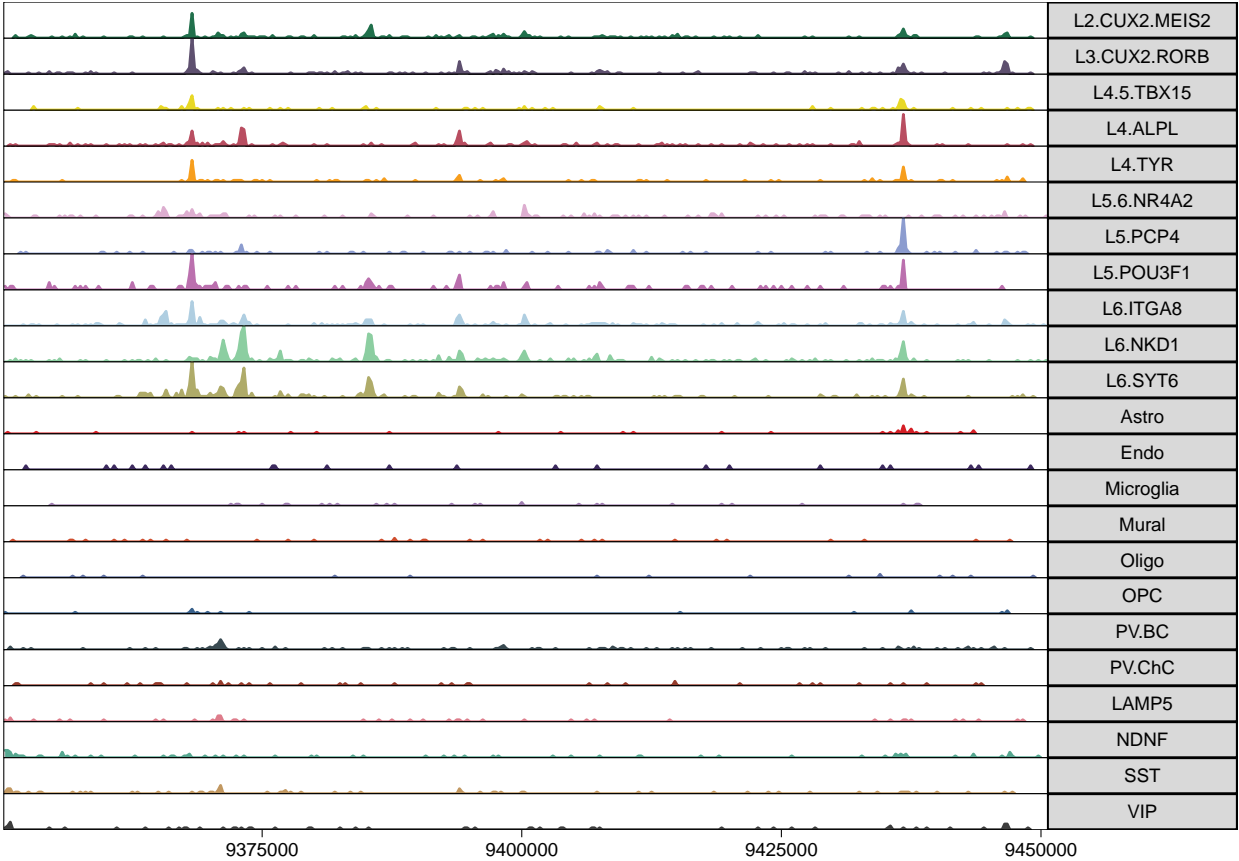


Peaks



GeneTrack

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



L5.6.NR4A2

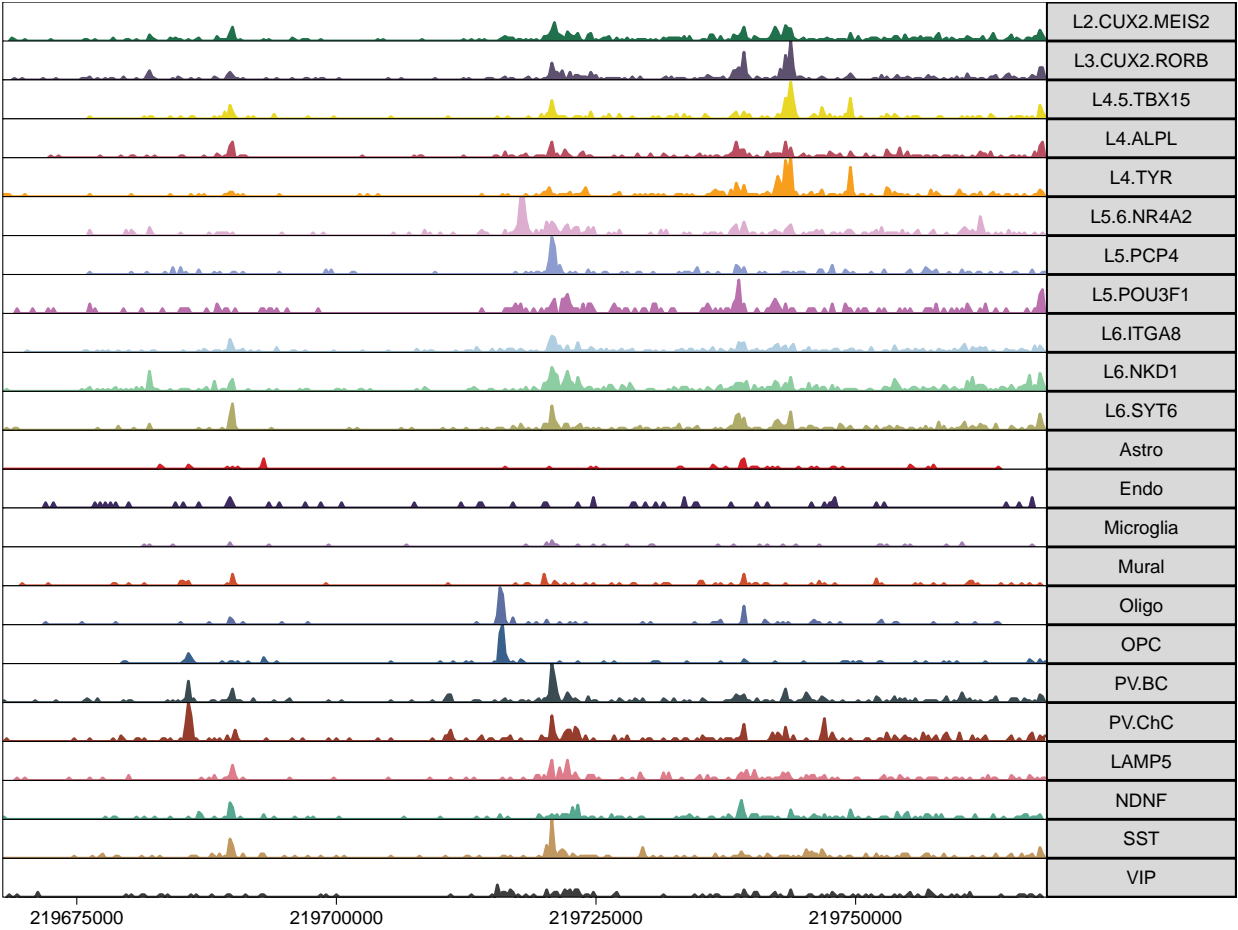


Peaks

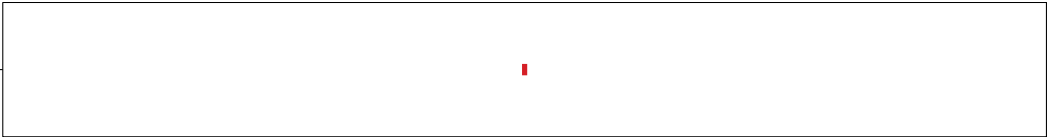
GeneTrack

chr1:219667875–219768376 Marker Peak Tracks for: L5.6.NR4A2

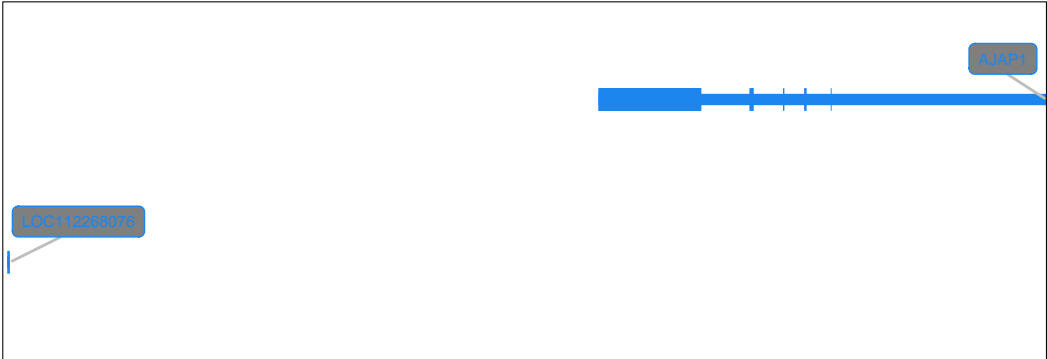
Coverage
(Norm. ATAC Signal Range (0–0.6) by ReadsInTSS)



L5.6.NR4A2



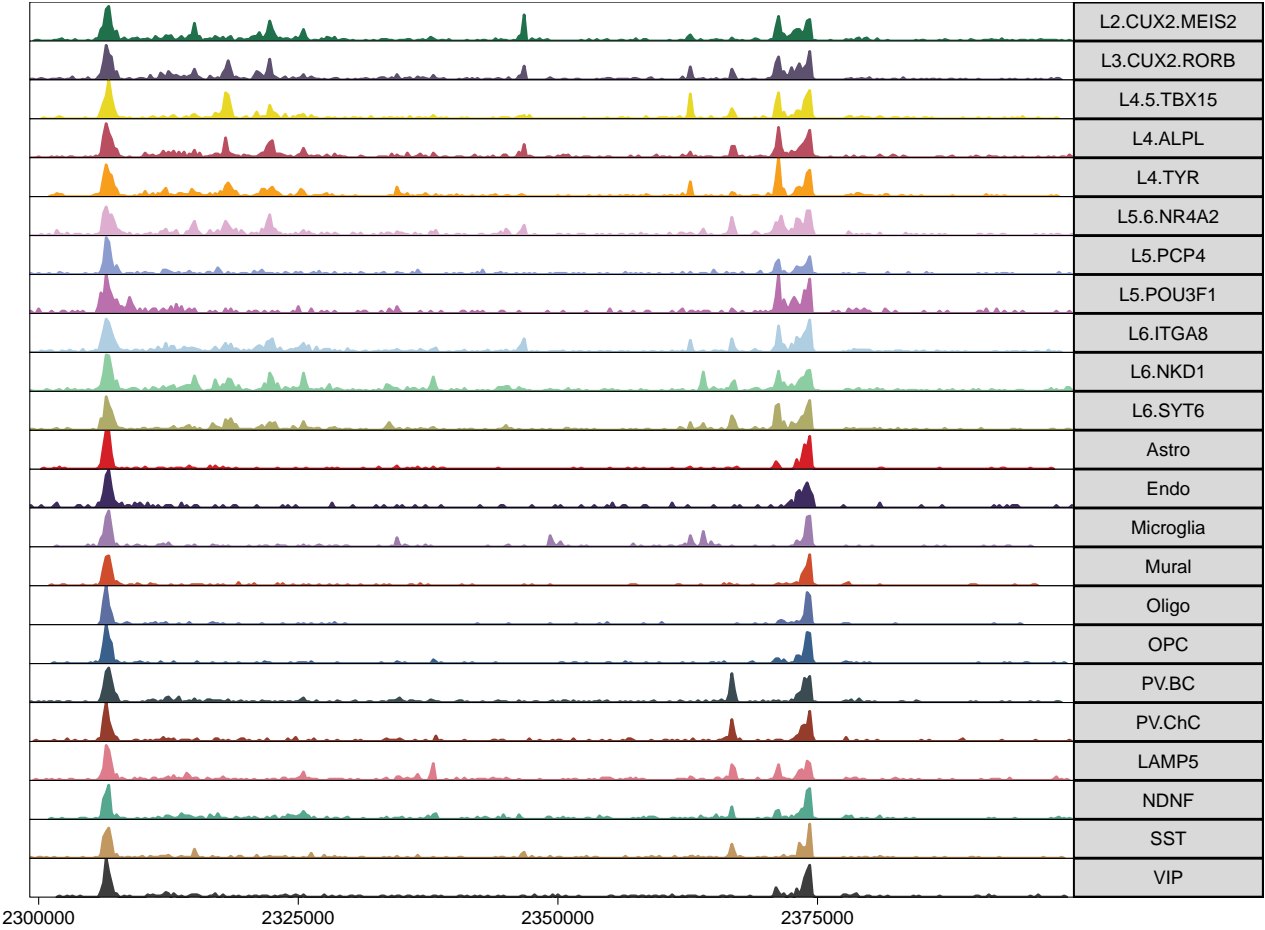
Peaks



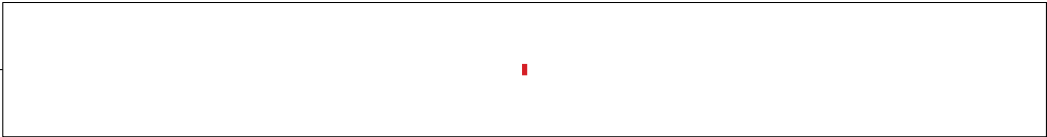
Genes

chr20:2299144–2399645 Marker Peak Tracks for: L5.6.NR4A2

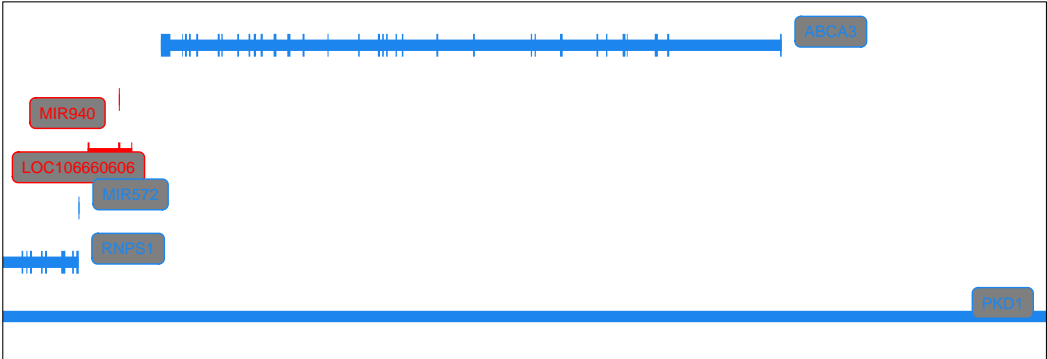
Coverage
(Norm. ATAC Signal Range (0–1.25) by ReadsInTSS)



L5.6.NR4A2



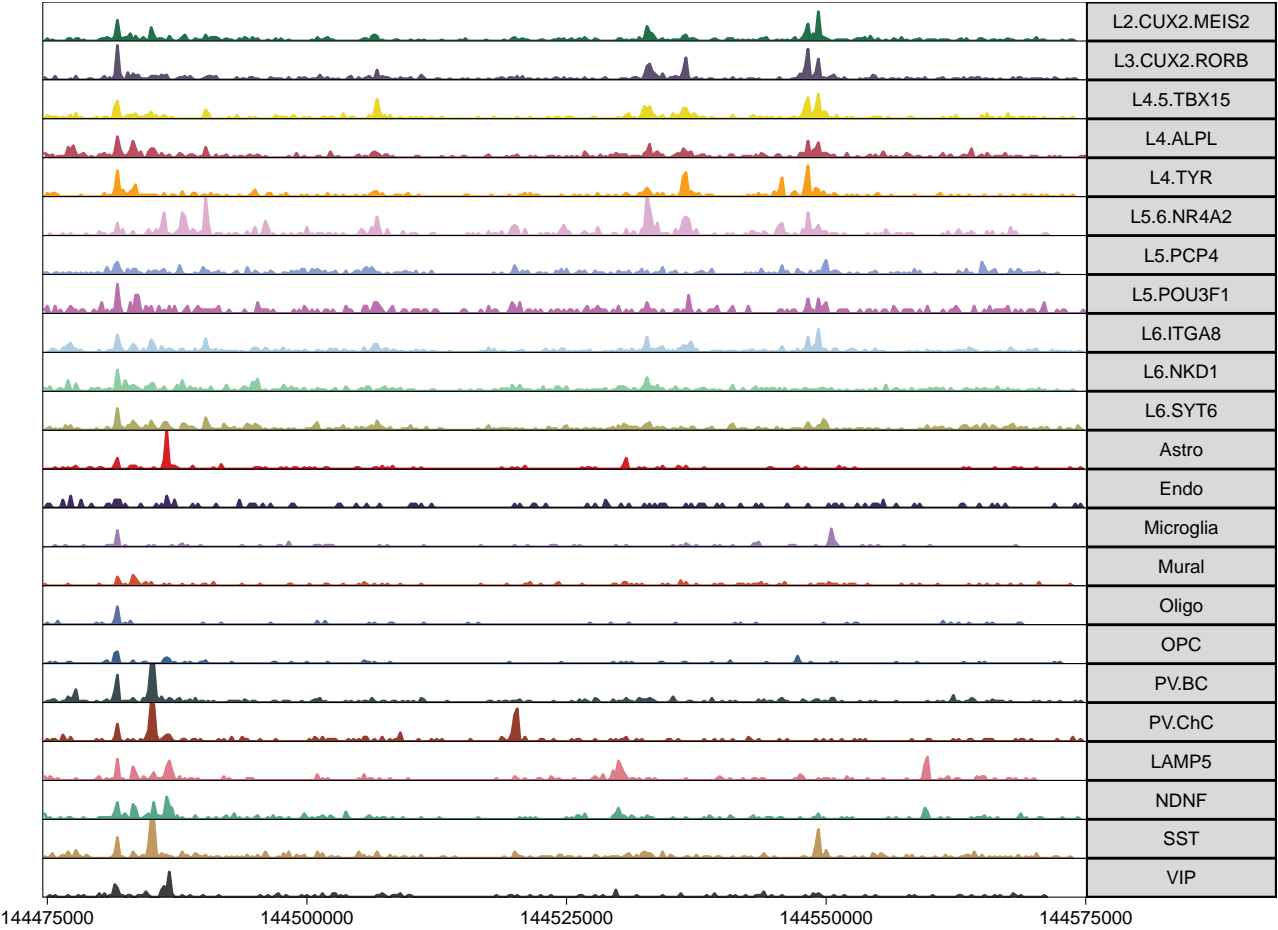
Peaks



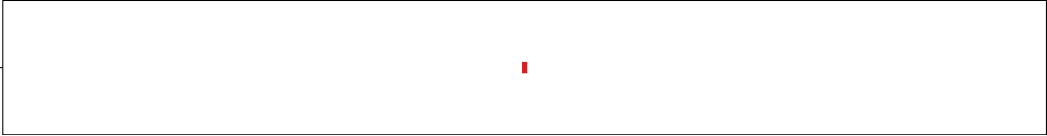
Genes

chr2:144474564–144575065 Marker Peak Tracks for: L5.6.NR4A2

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



L5.6.NR4A2



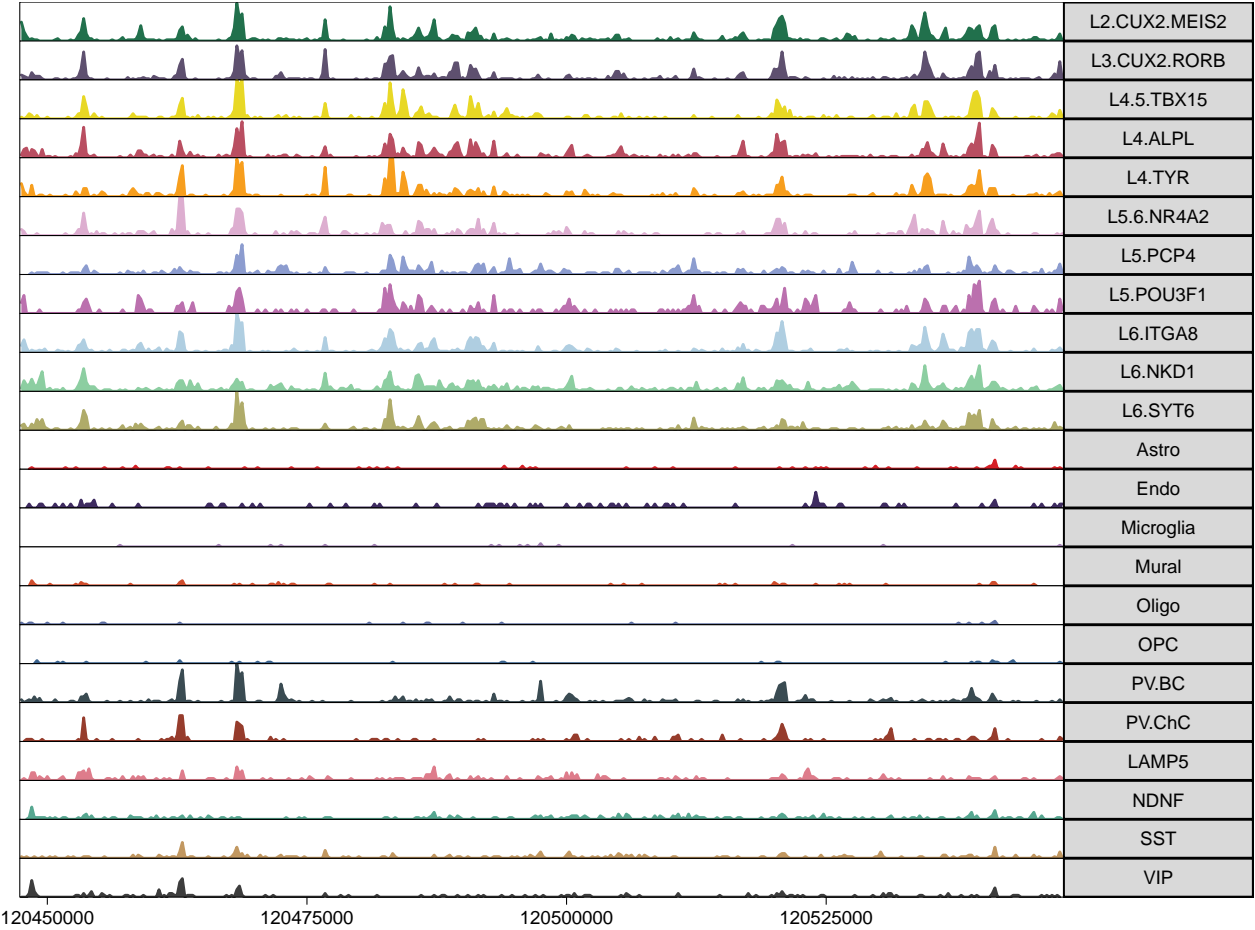
Peaks



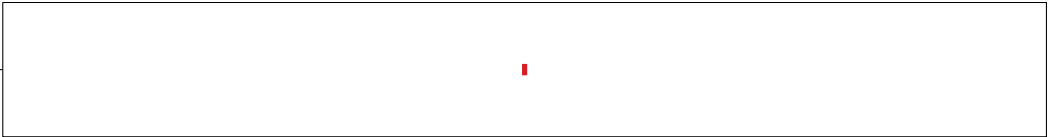
Genes

chr2:120447347–120547848 Marker Peak Tracks for: L5.6.NR4A2

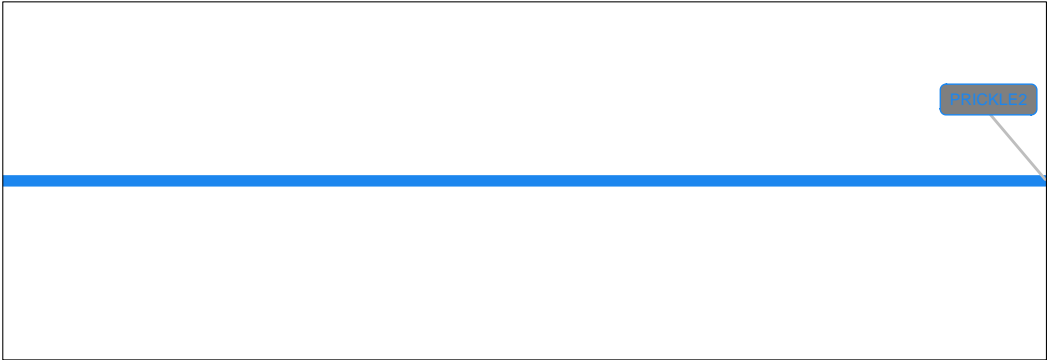
Coverage
(Norm. ATAC Signal Range (0–0.79) by ReadsInTSS)



L5.6.NR4A2

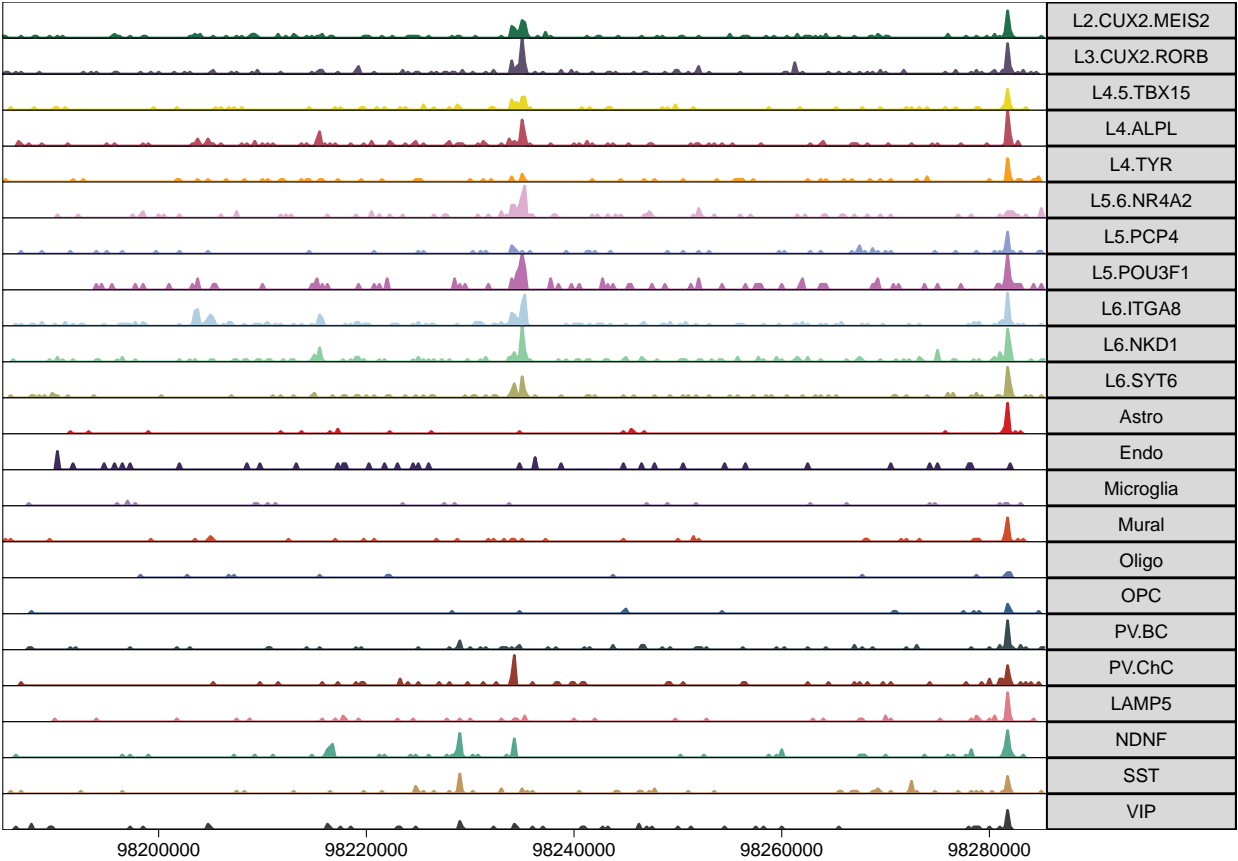


Peaks



Genes

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



L5.6.NR4A2

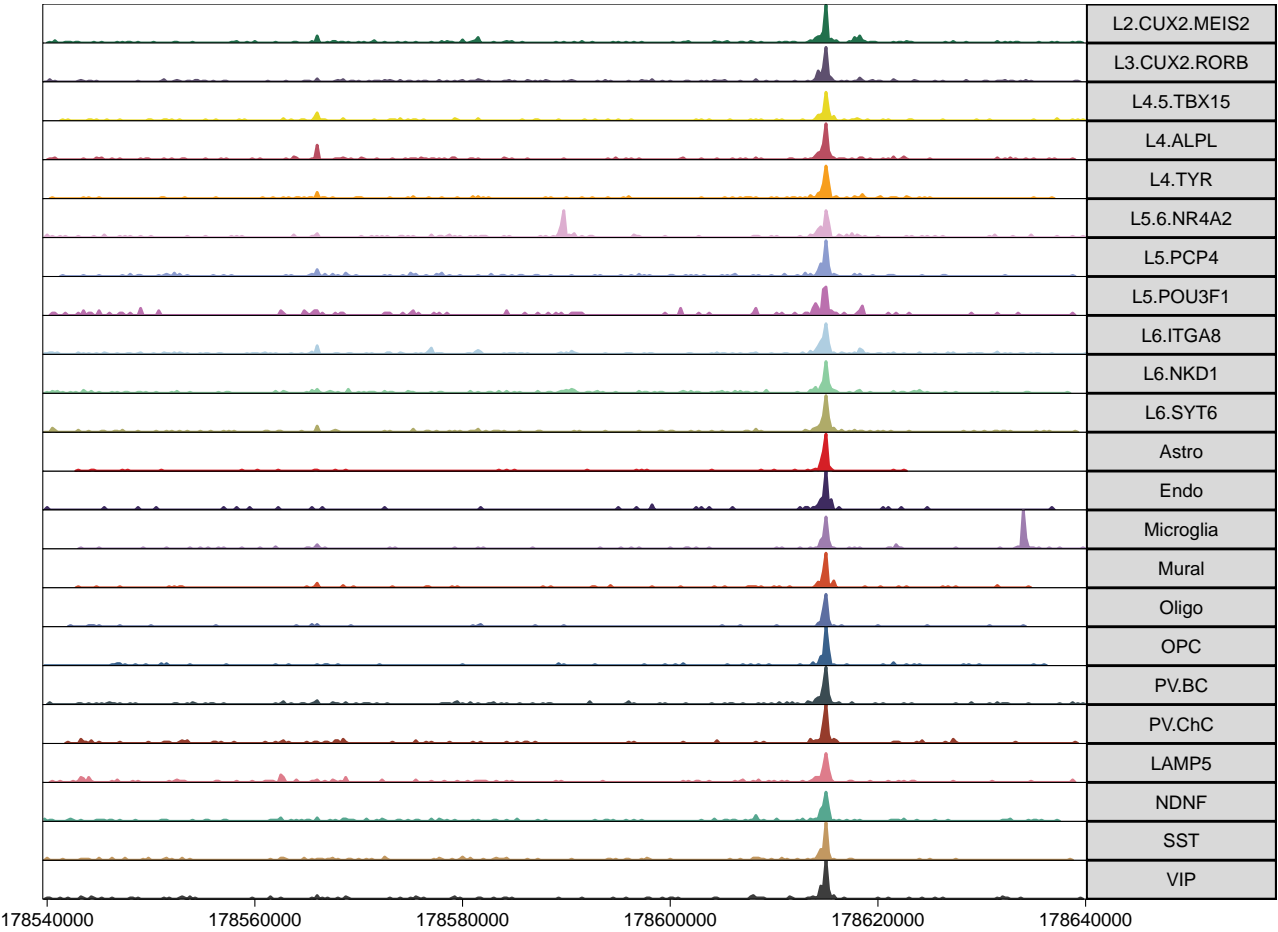


Peaks

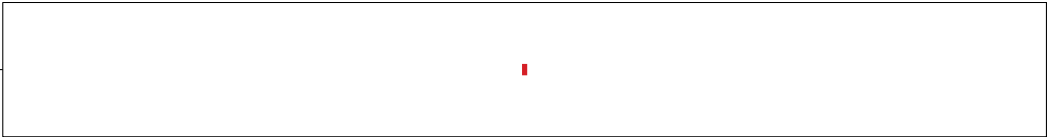
GeneTrack

chr1:178539579–178640080 Marker Peak Tracks for: L5.6.NR4A2

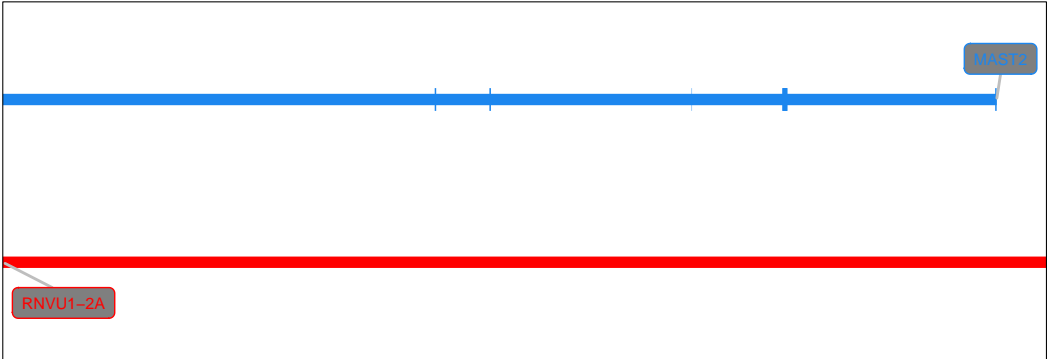
Coverage
(Norm. ATAC Signal Range (0–1.2) by ReadsInTSS)



L5.6.NR4A2

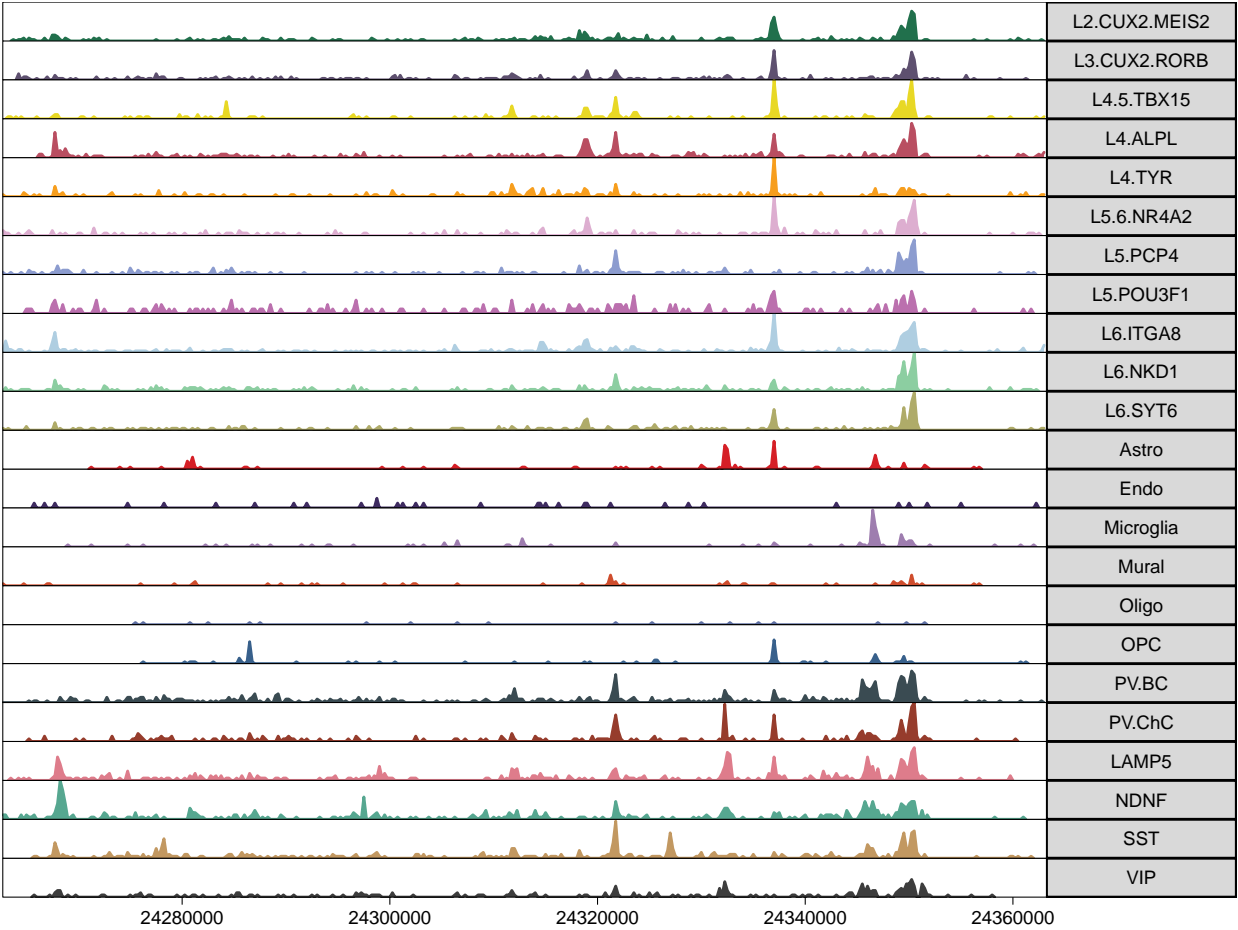


Peaks

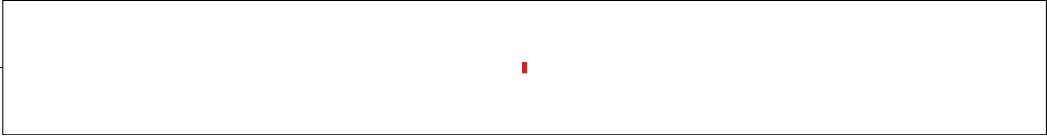


Genes

Coverage
(Norm. ATAC Signal Range (0–0.64) by ReadsInTSS)



L5.6.NR4A2

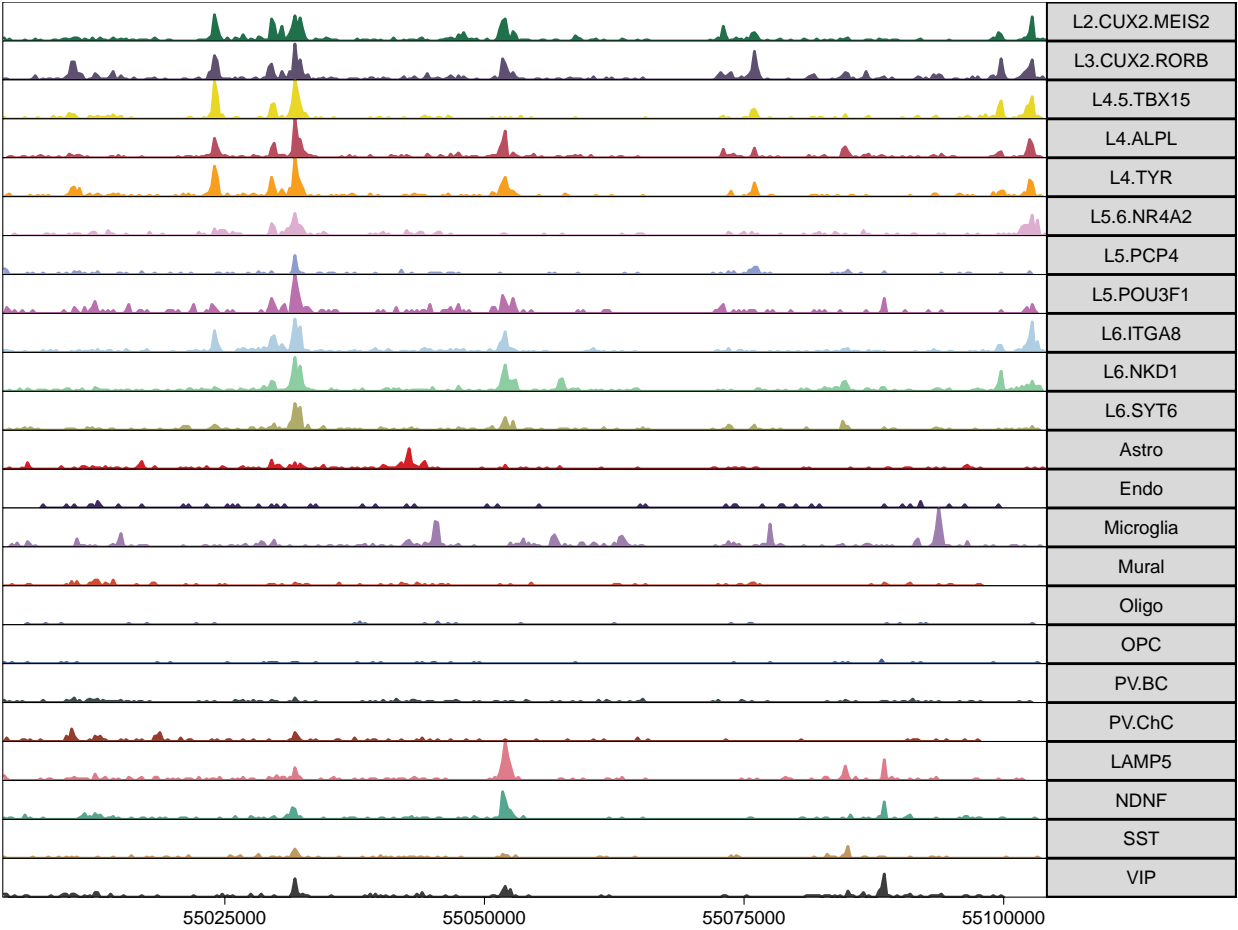


Peaks

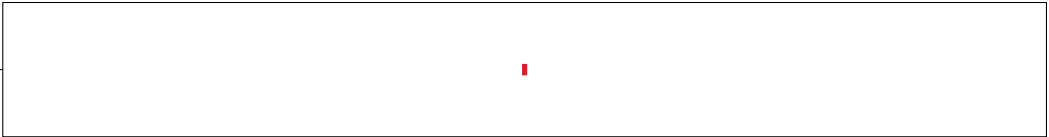


Genes

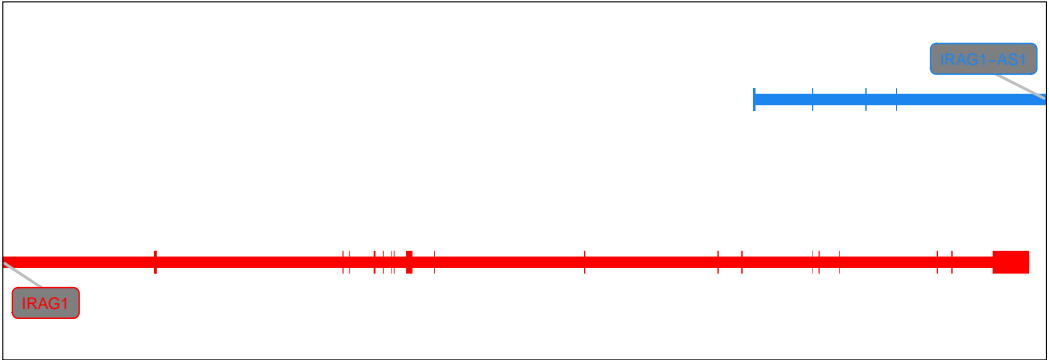
Coverage
(Norm. ATAC Signal Range (0-0.95) by ReadsInTSS)



L5.6.NR4A2



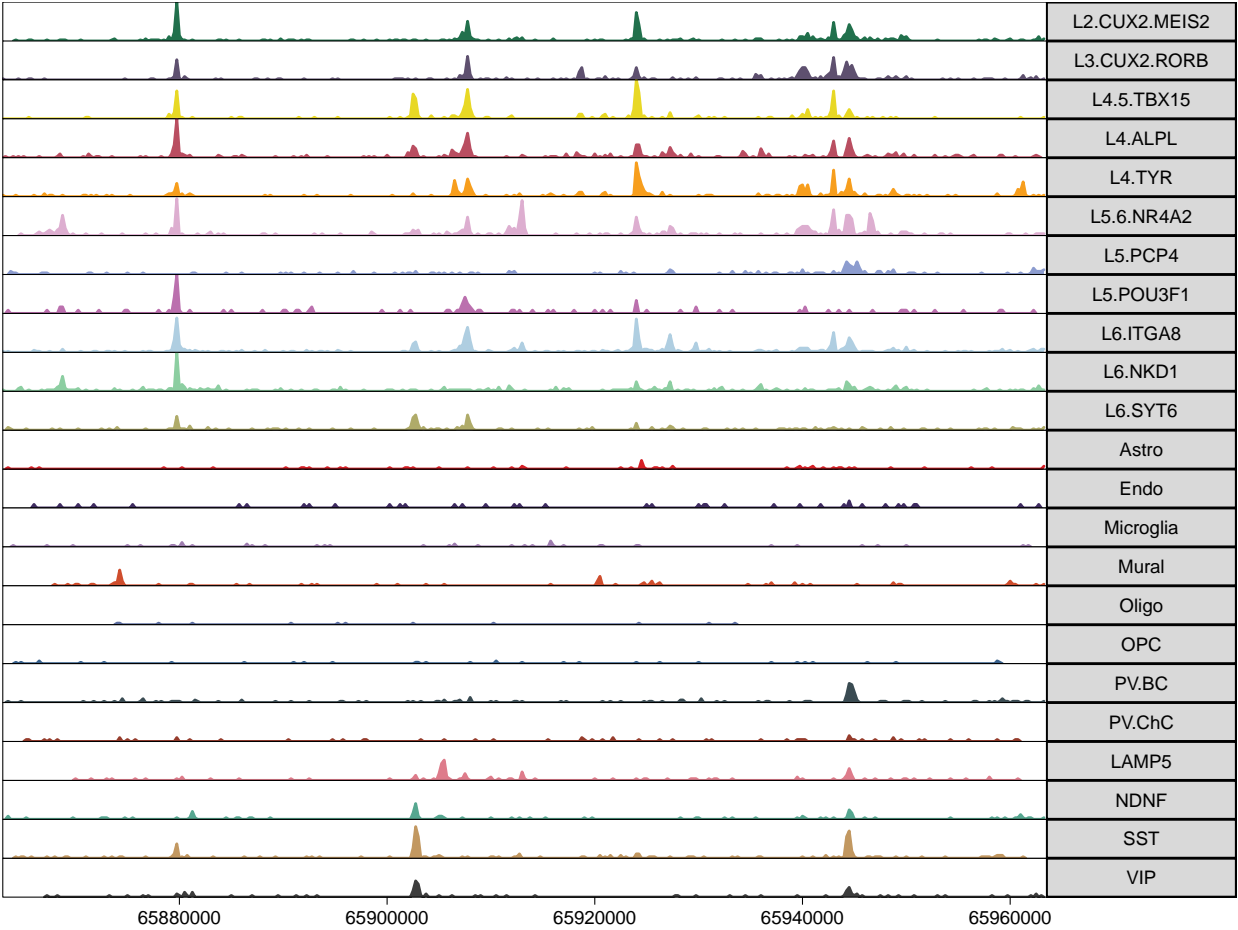
Peaks



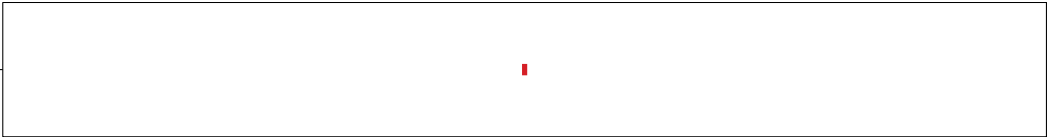
Genes

chr1:65862992–65963493 Marker Peak Tracks for: L5.6.NR4A2

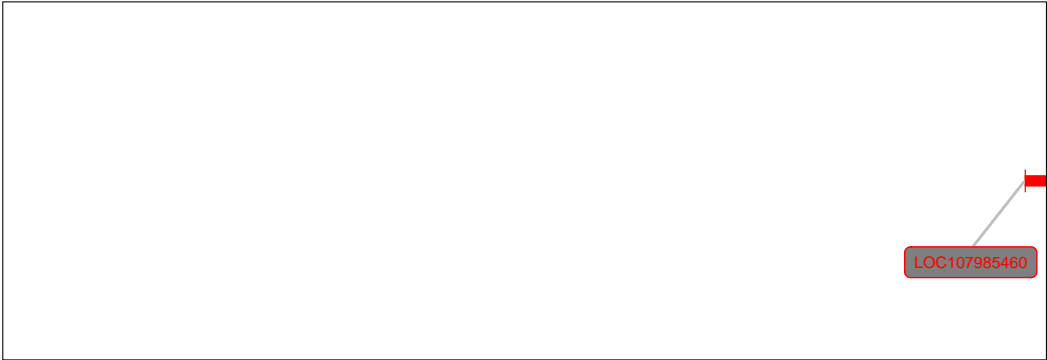
Coverage
(Norm. ATAC Signal Range (0–0.88) by ReadsInTSS)



L5.6.NR4A2

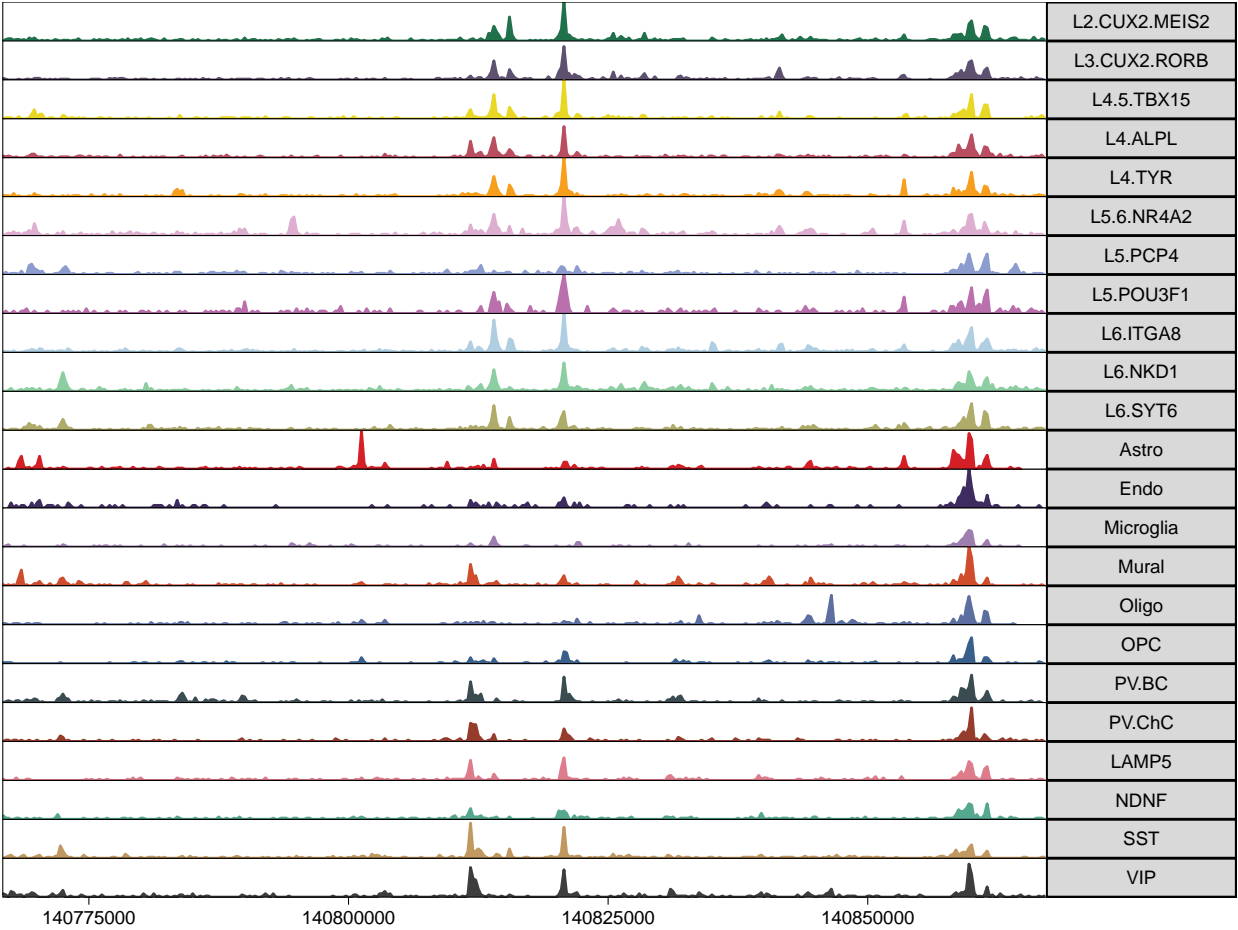


Peaks

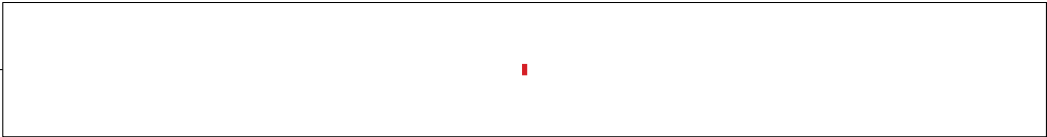


Genes

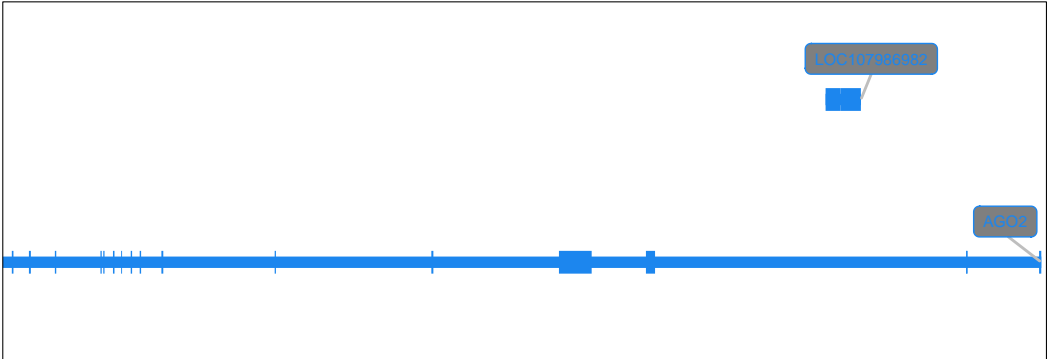
Coverage
(Norm. ATAC Signal Range (0–1.22) by ReadsInTSS)



L5.6.NR4A2

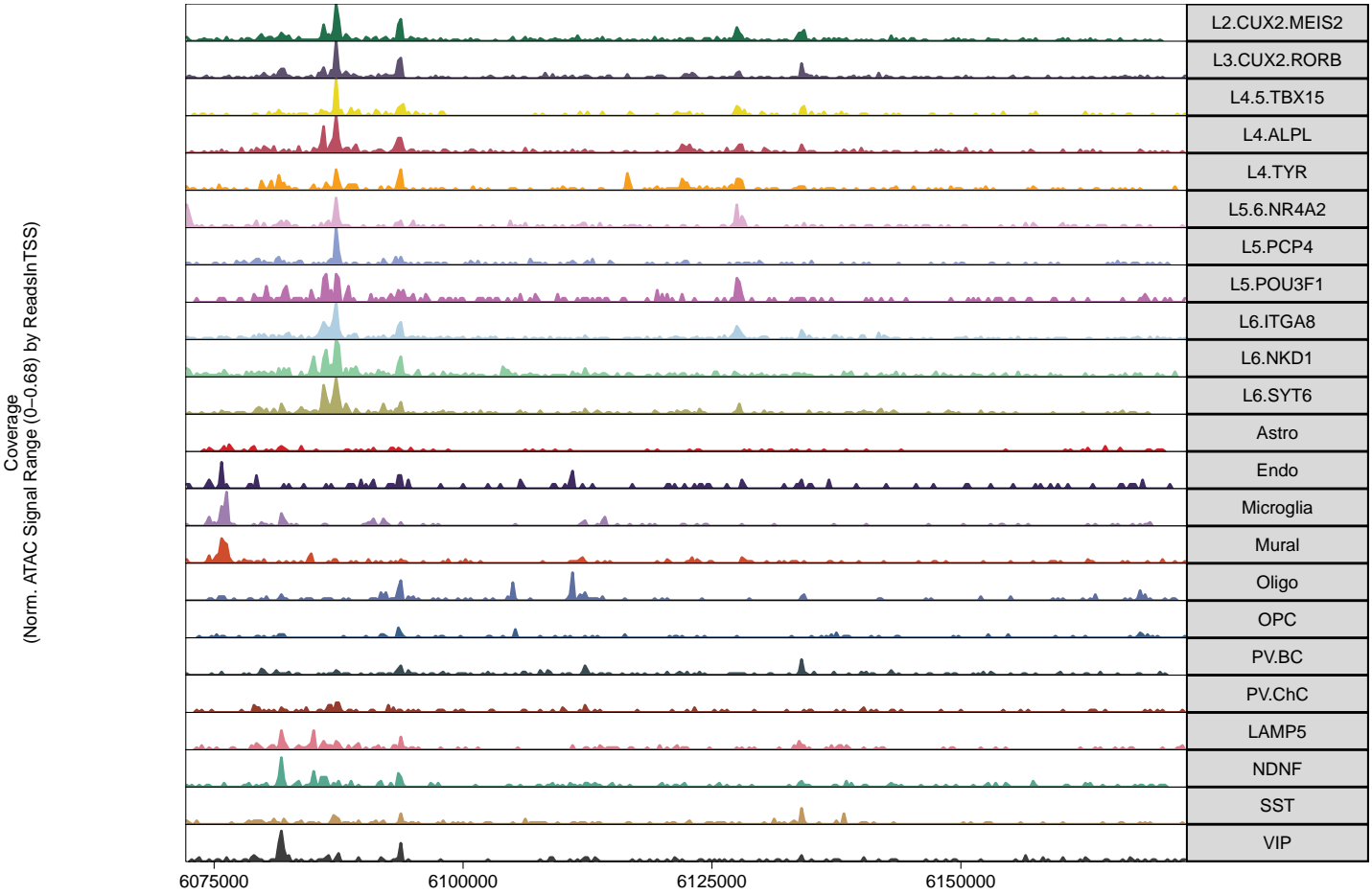


Peaks

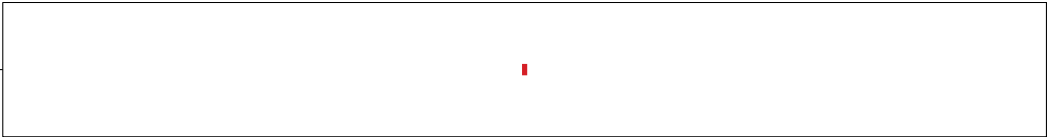


Genes

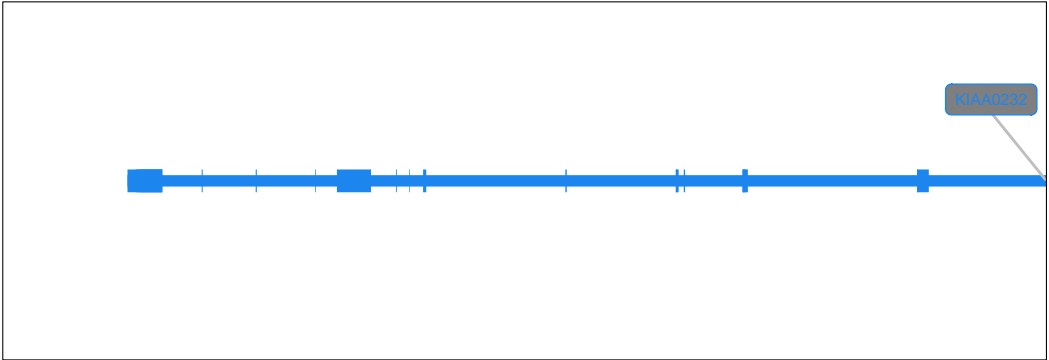
chr5:6072152–6172653 Marker Peak Tracks for: L5.6.NR4A2



L5.6.NR4A2

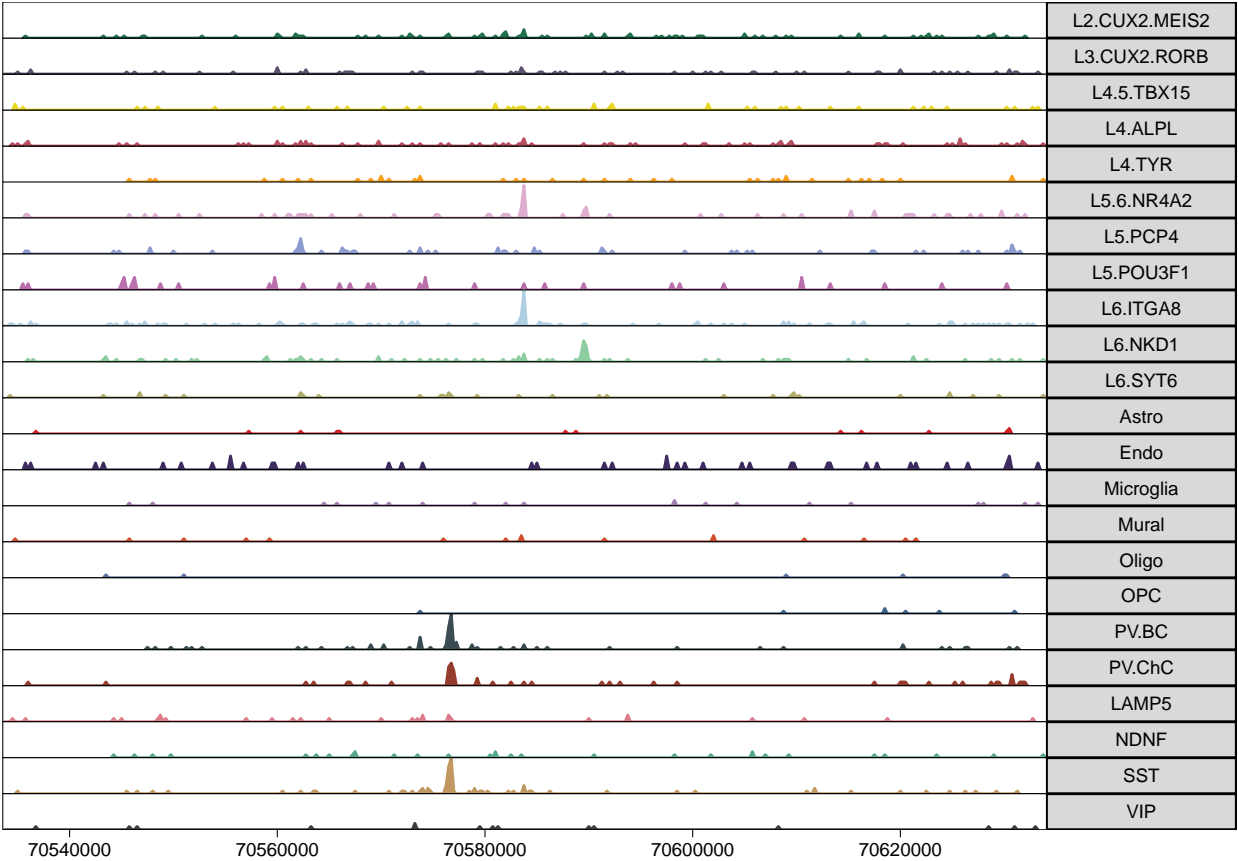


Peaks



Genes

Coverage
(Norm. ATAC Signal Range (0–0.42) by ReadsInTSS)



L5.6.NR4A2

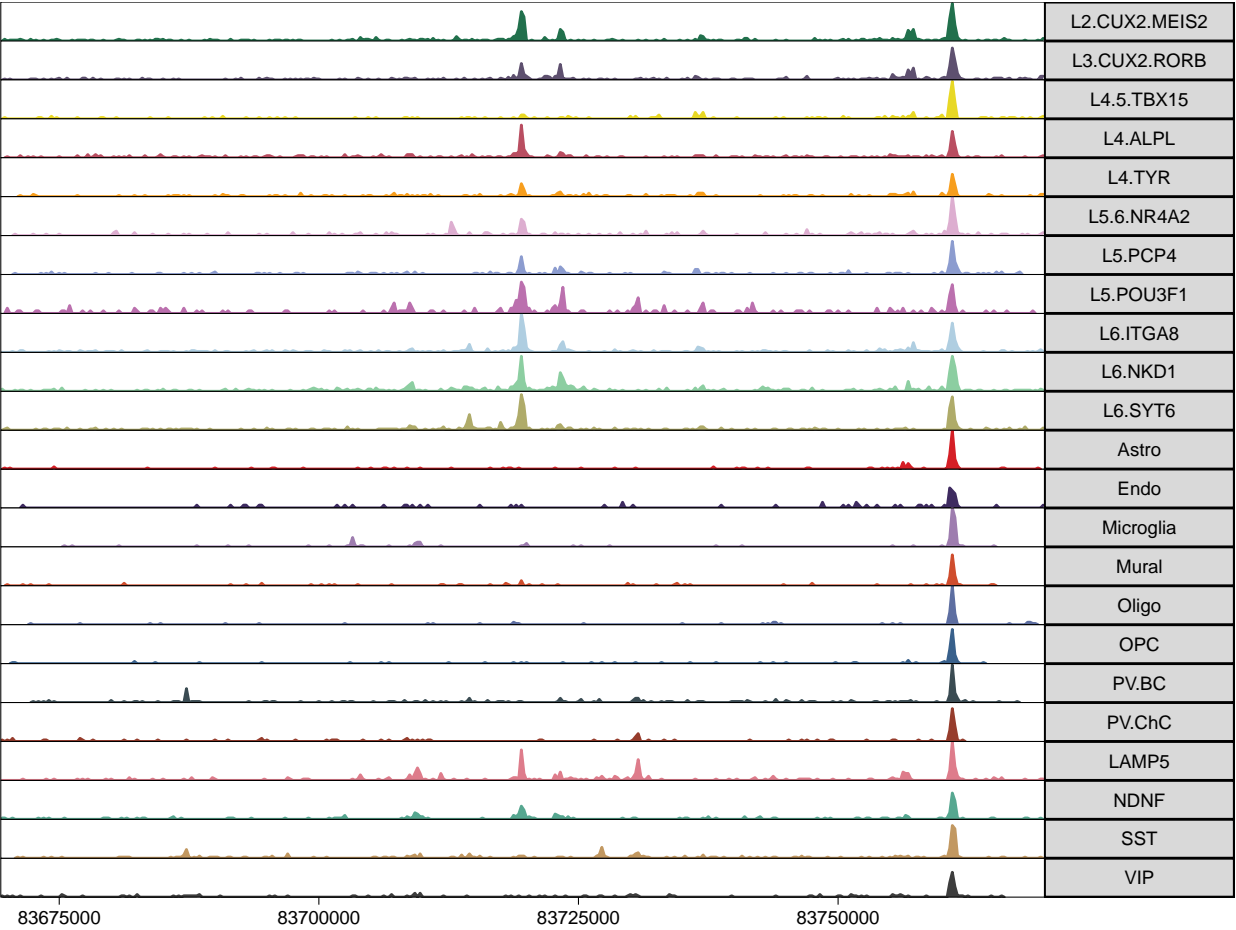


Peaks

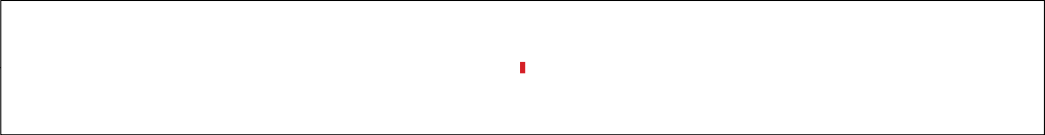
GeneTrack

chr4:83669371–83769872 Marker Peak Tracks for: L5.6.NR4A2

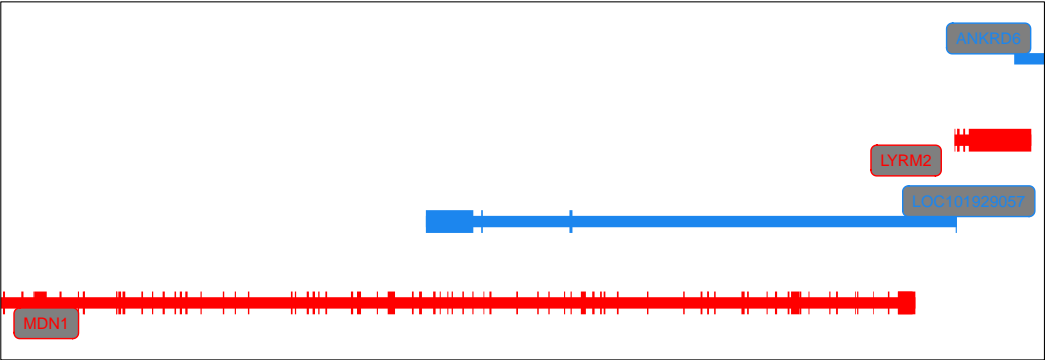
Coverage
(Norm. ATAC Signal Range (0–1.09) by ReadsInTSS)



L5.6.NR4A2

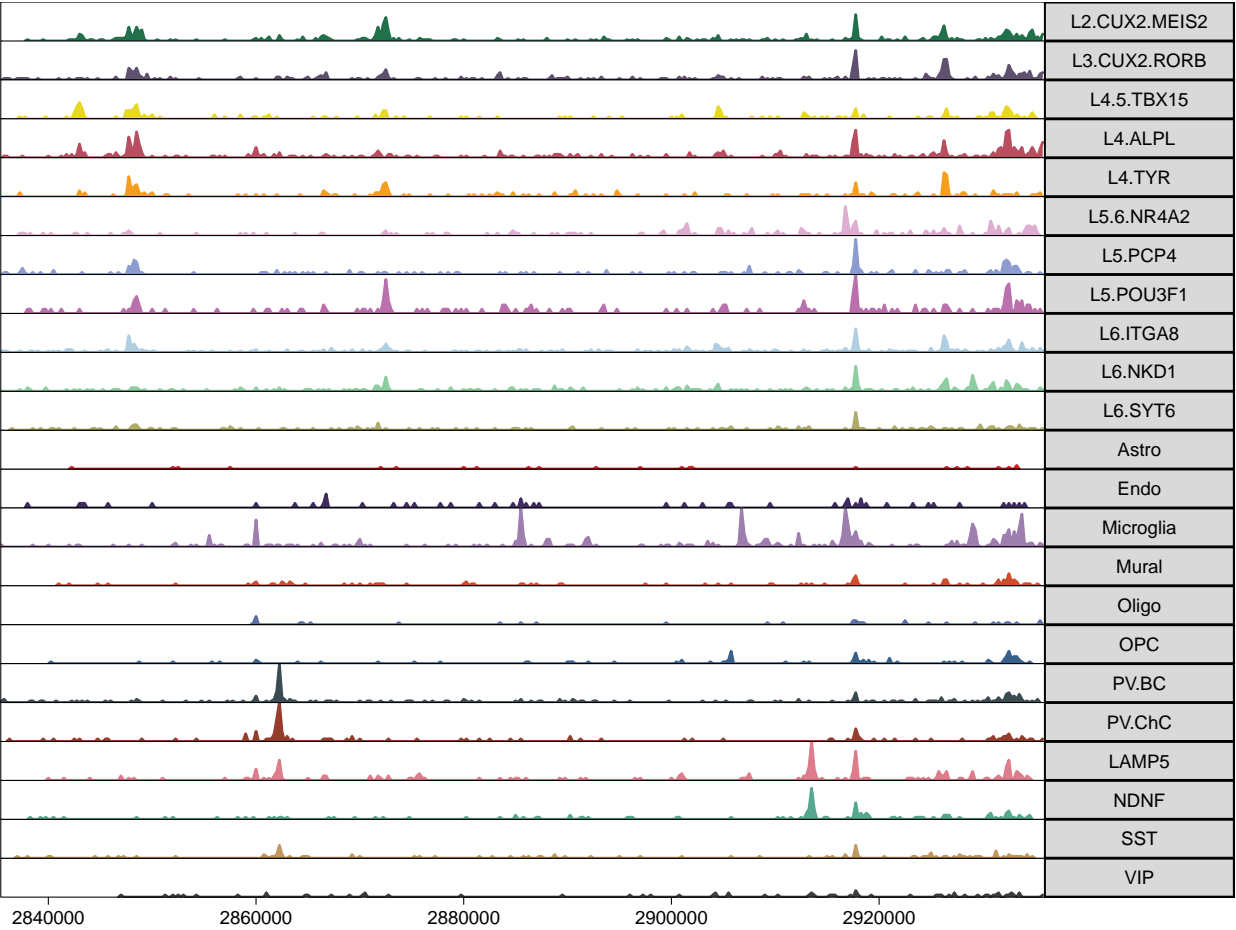


Peaks

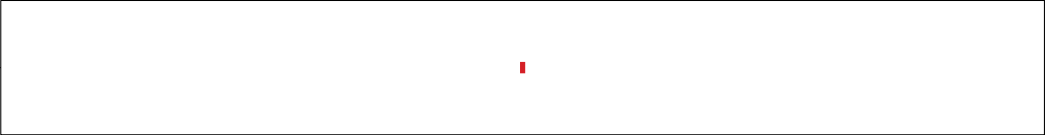


Genes

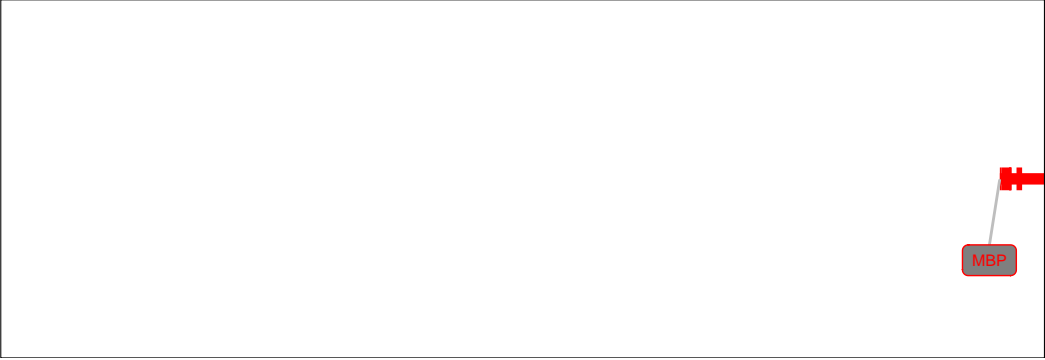
Coverage
(Norm. ATAC Signal Range (0–0.67) by ReadsInTSS)



L5.6.NR4A2

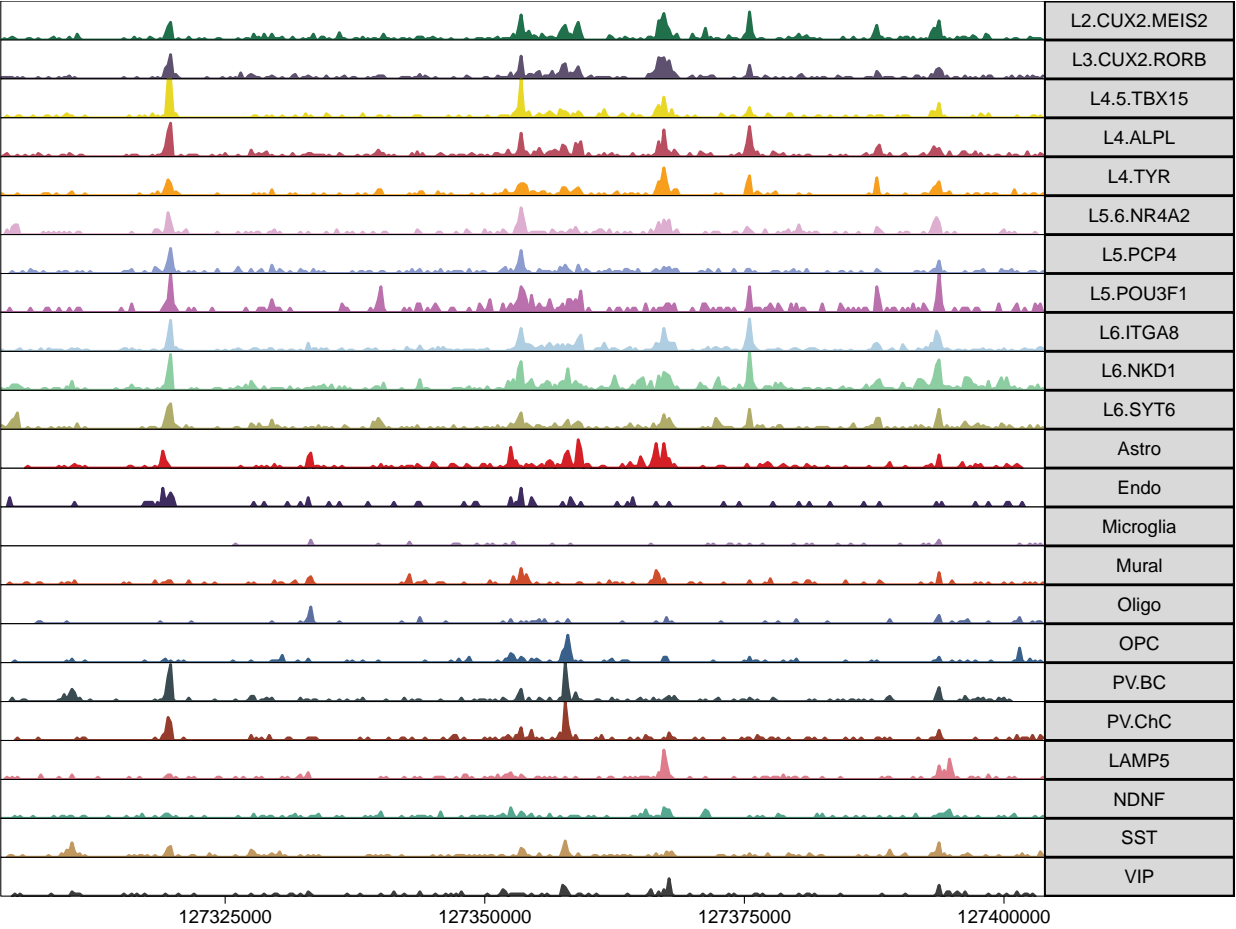


Peaks

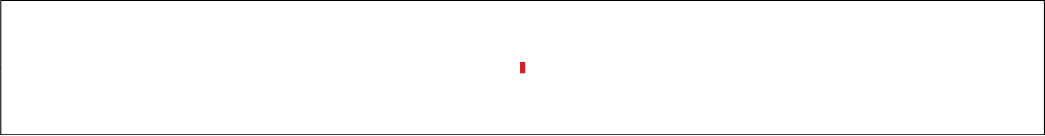


Genes

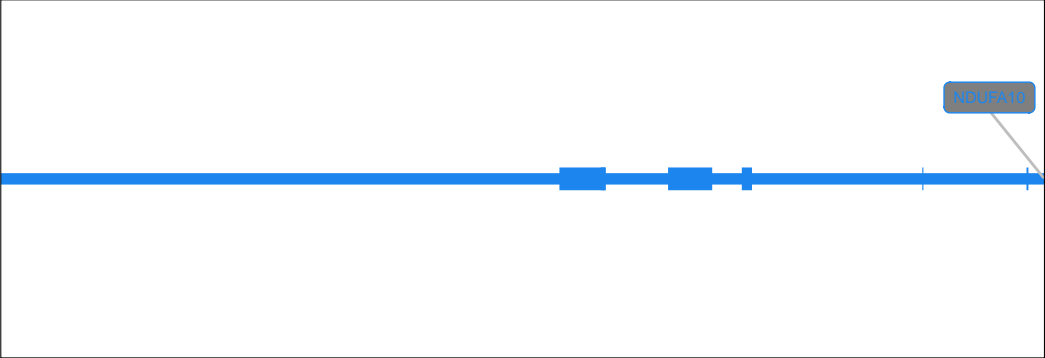
Coverage
(Norm. ATAC Signal Range (0–0.67) by ReadsInTSS)



L5.6.NR4A2

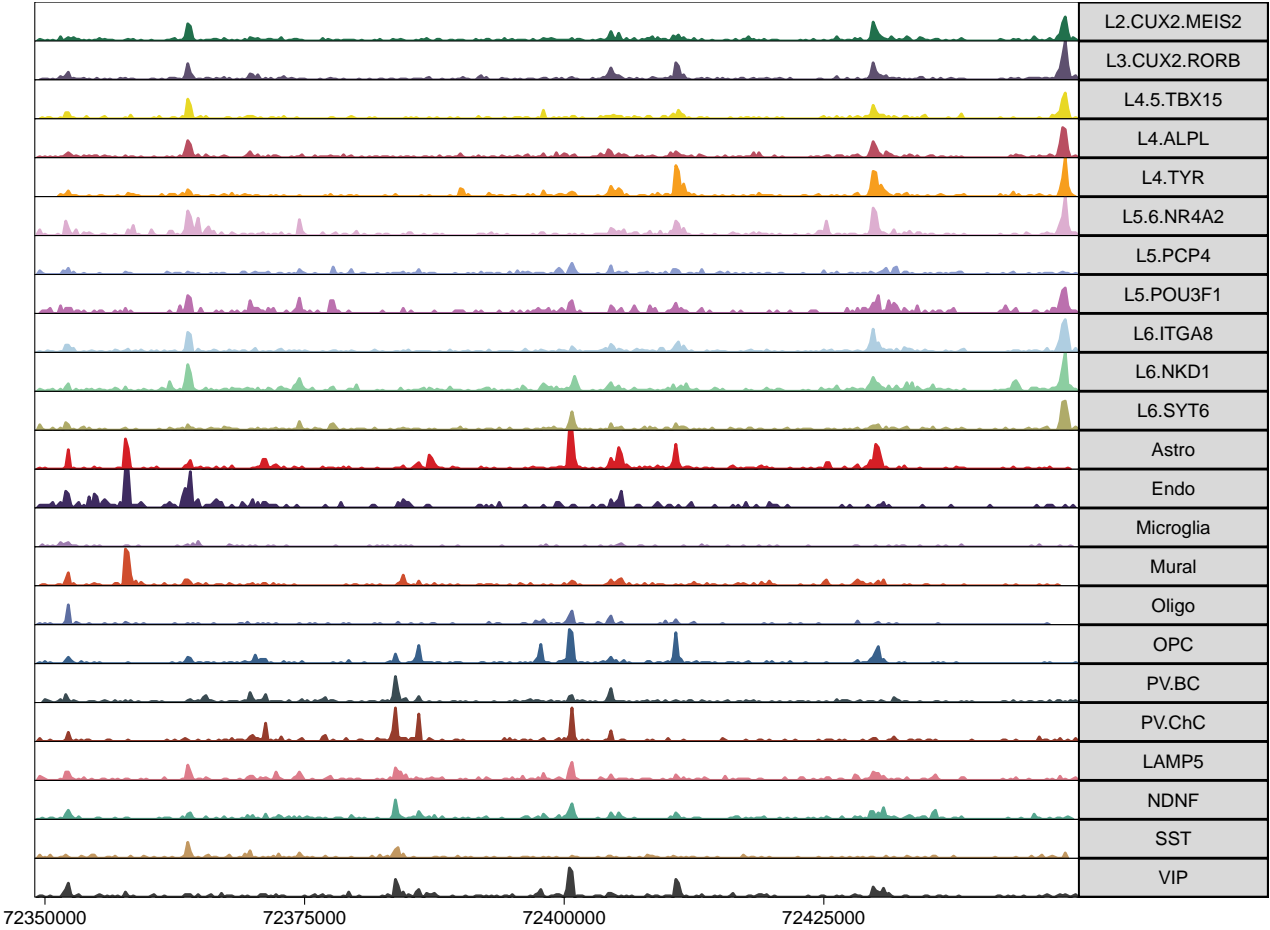


Peaks

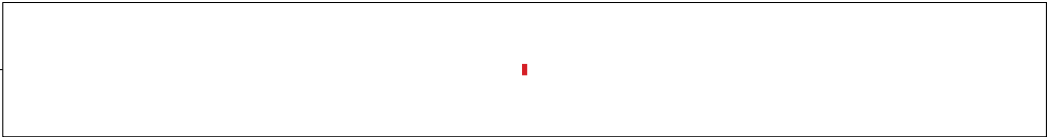


Genes

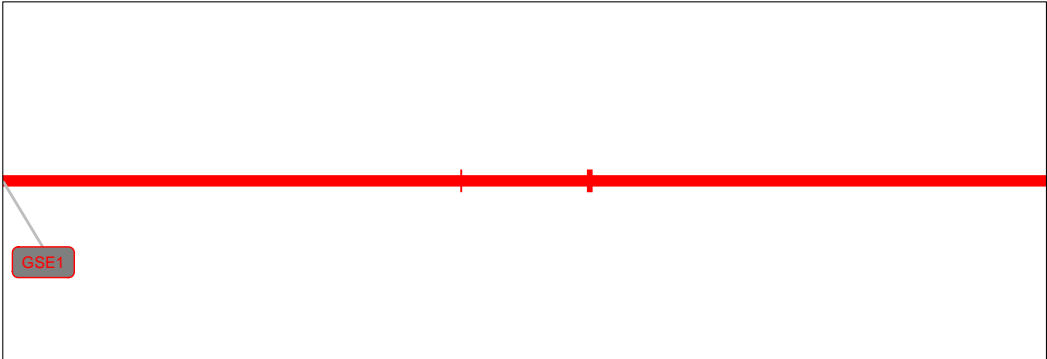
Coverage
(Norm. ATAC Signal Range (0–1.12) by ReadsInTSS)



L5.6.NR4A2

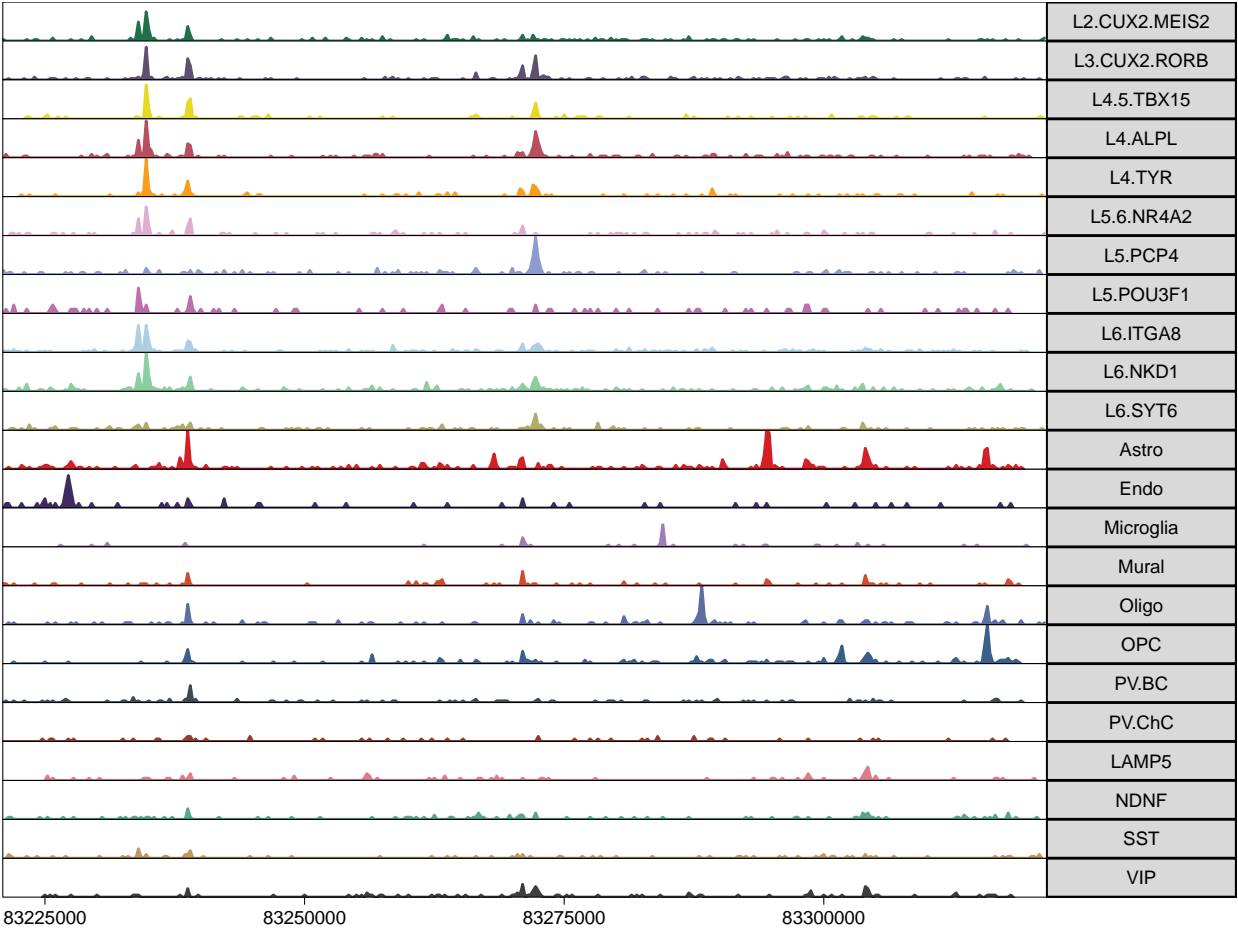


Peaks

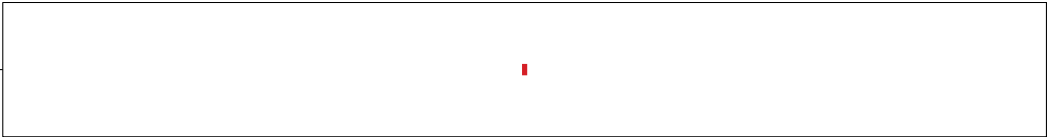


Genes

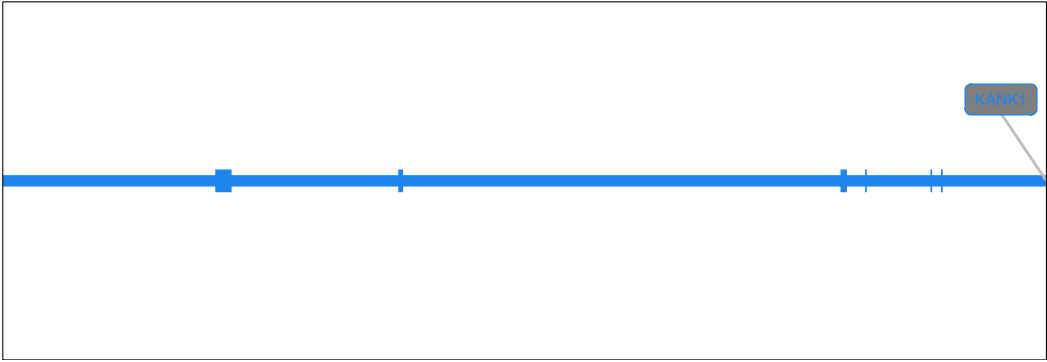
Coverage
(Norm. ATAC Signal Range (0–0.67) by ReadsInTSS)



L5.6.NR4A2



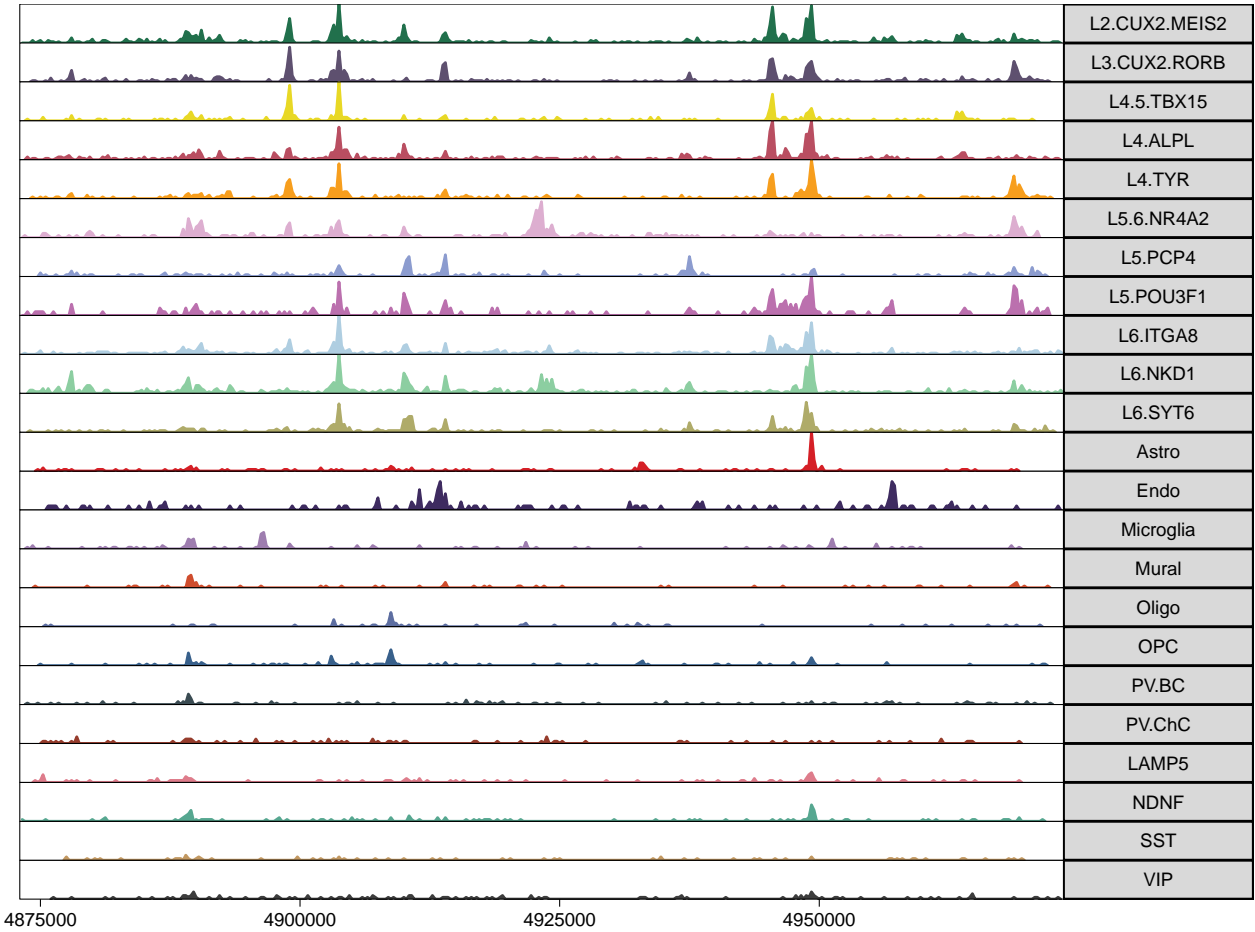
Peaks



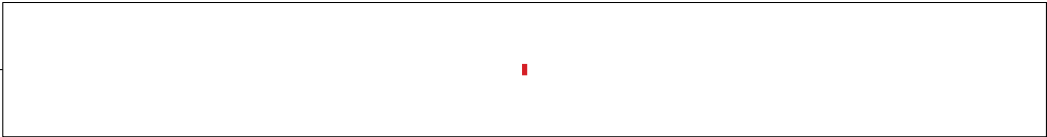
Genes

chr1:4873013–4973514 Marker Peak Tracks for: L5.6.NR4A2

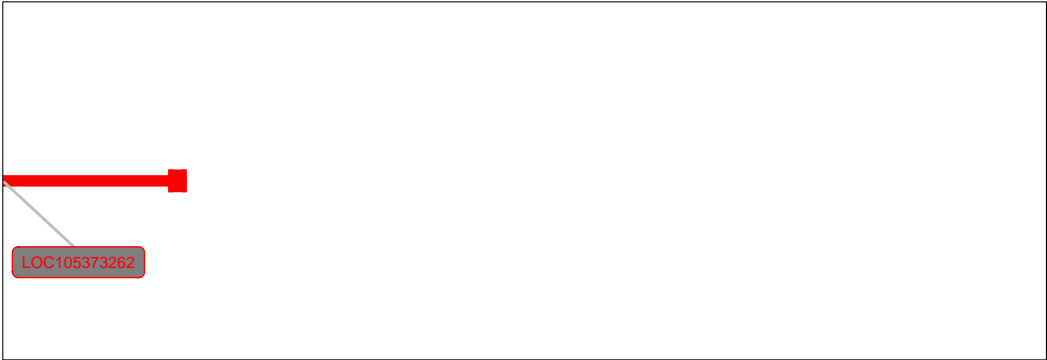
Coverage
(Norm. ATAC Signal Range (0–0.77) by ReadsInTSS)



L5.6.NR4A2

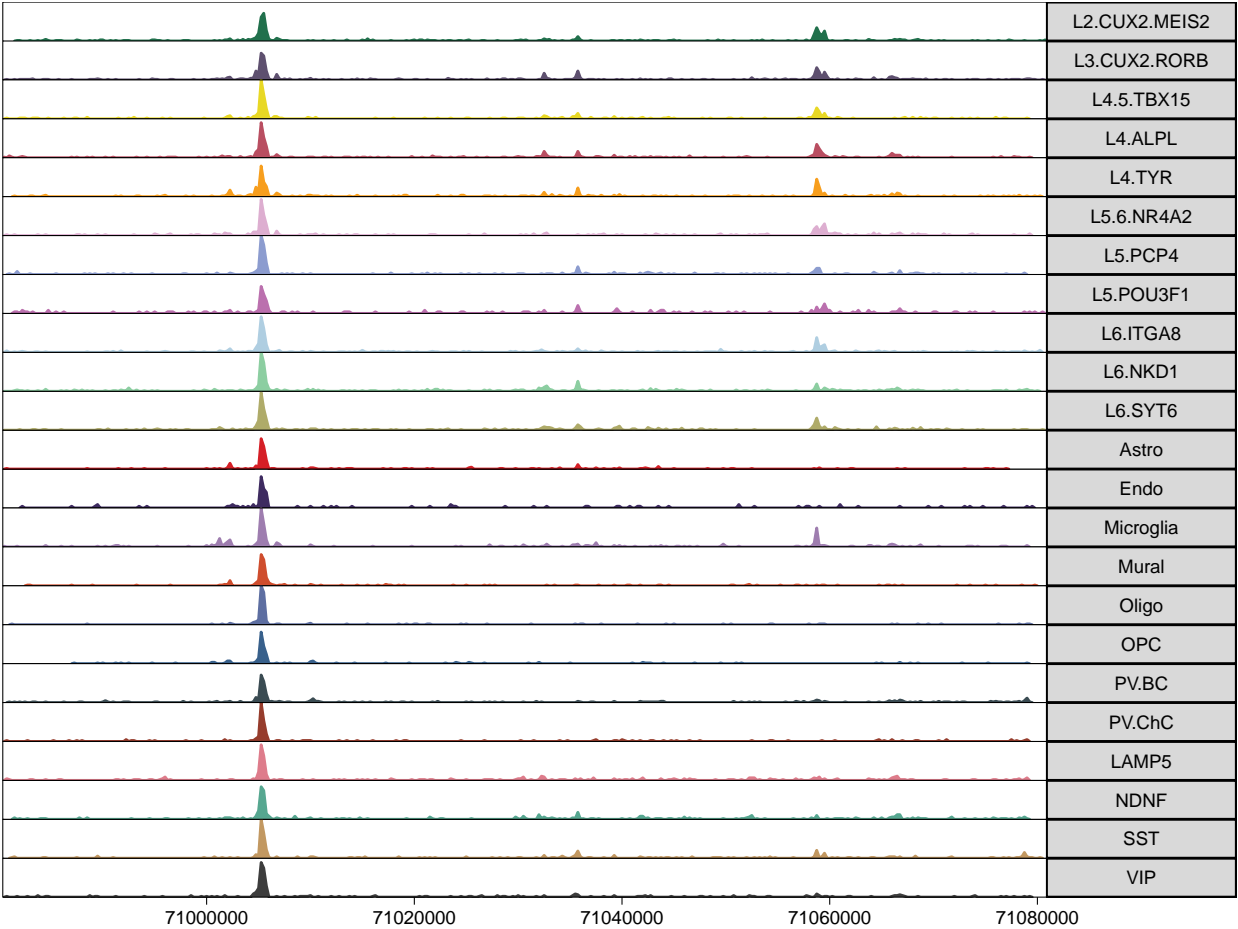


Peaks

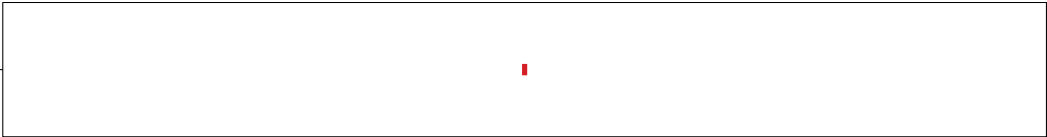


Genes

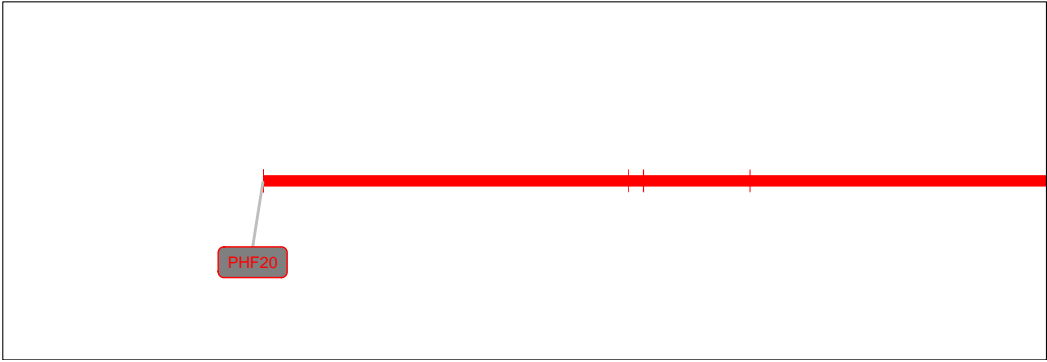
Coverage
(Norm. ATAC Signal Range (0–1.8) by ReadsInTSS)



L5.6.NR4A2



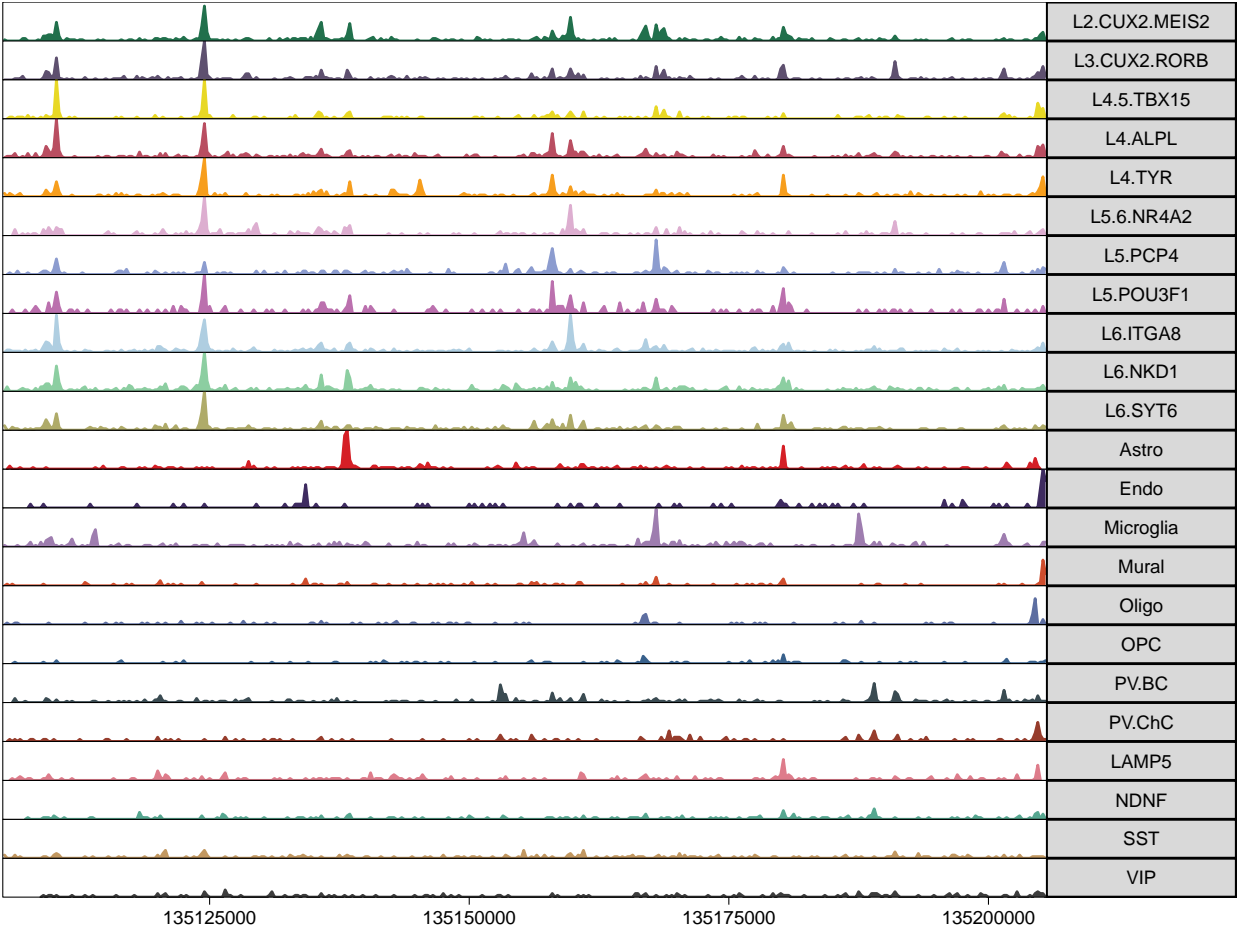
Peaks



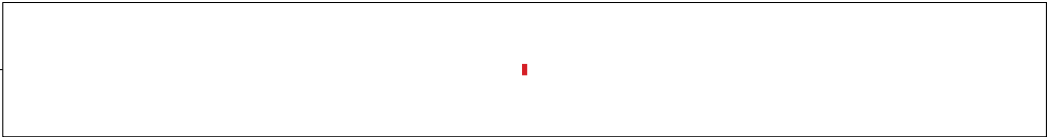
Genes

chr4:135105083–135205584 Marker Peak Tracks for: L5.6.NR4A2

Coverage
(Norm. ATAC Signal Range (0–0.8) by ReadsInTSS)



L5.6.NR4A2



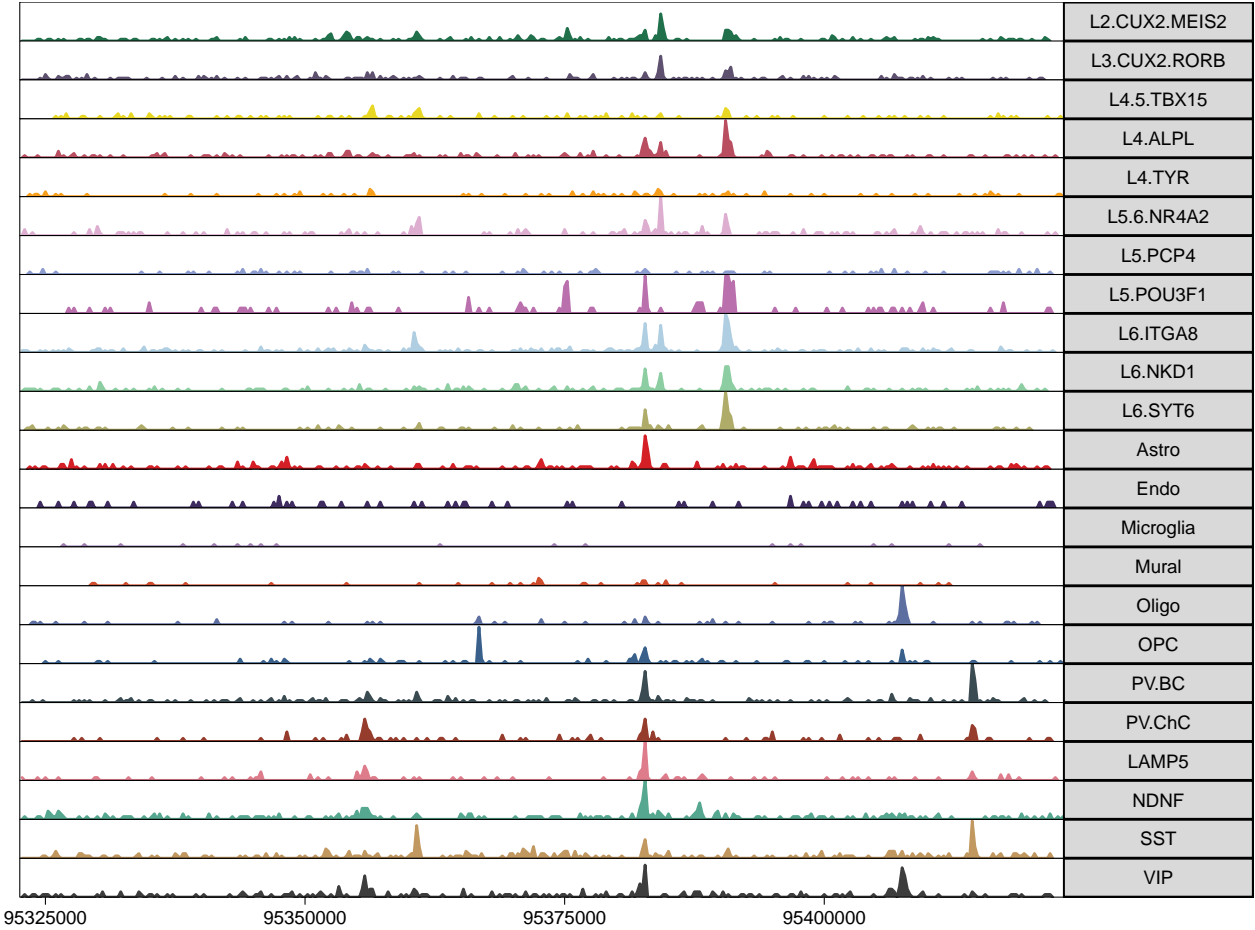
Peaks



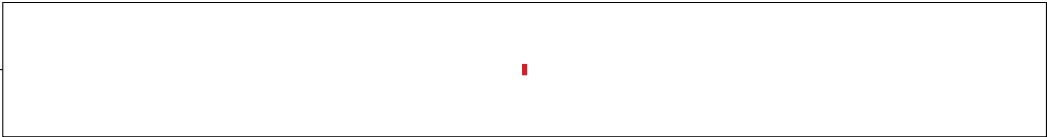
Genes

chr7:95322532–95423033 Marker Peak Tracks for: L5.6.NR4A2

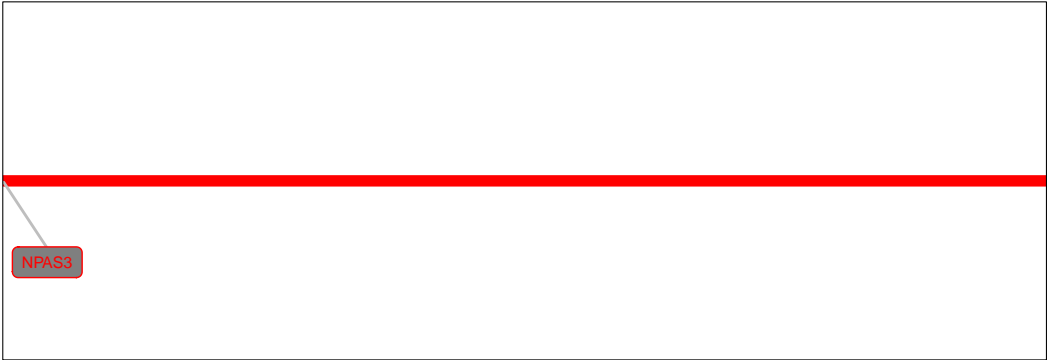
Coverage
(Norm. ATAC Signal Range (0–0.53) by ReadsInTSS)



L5.6.NR4A2

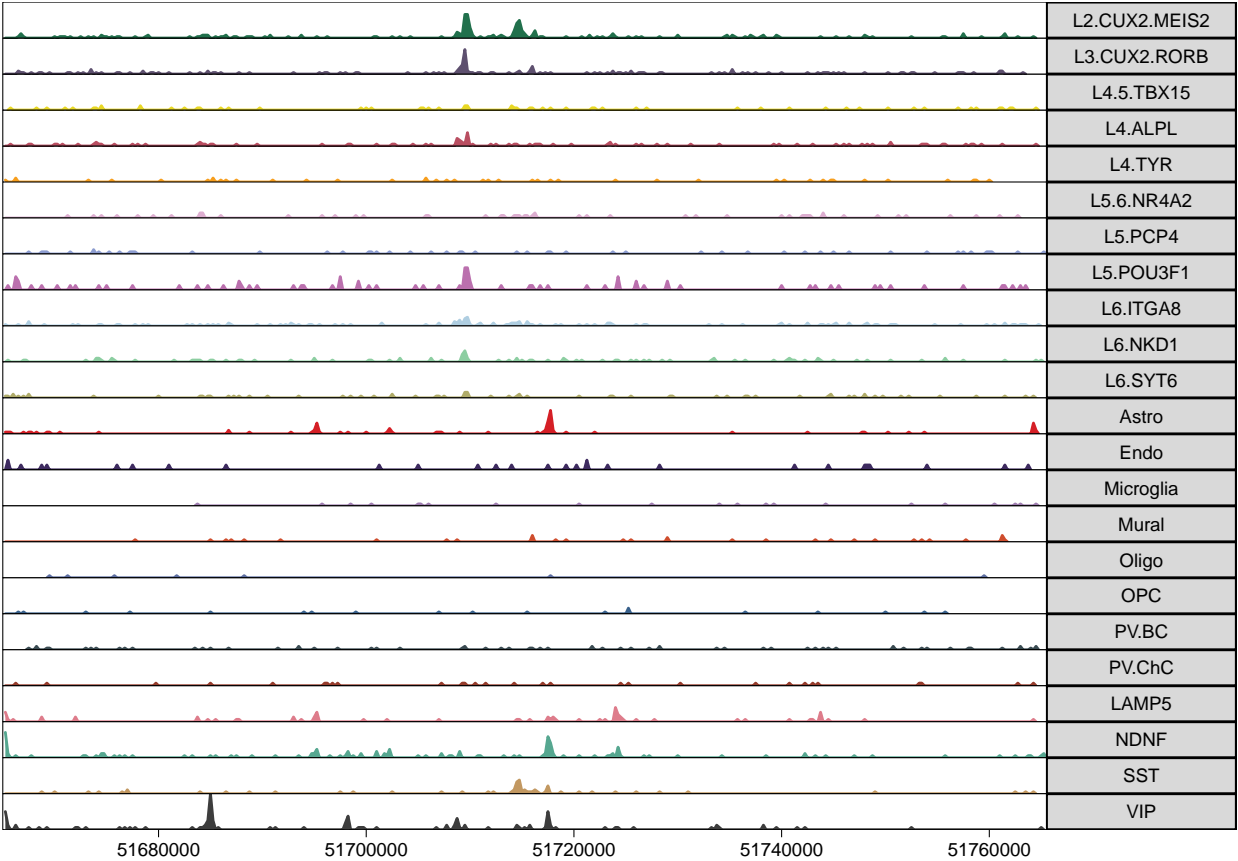


Peaks



Genes

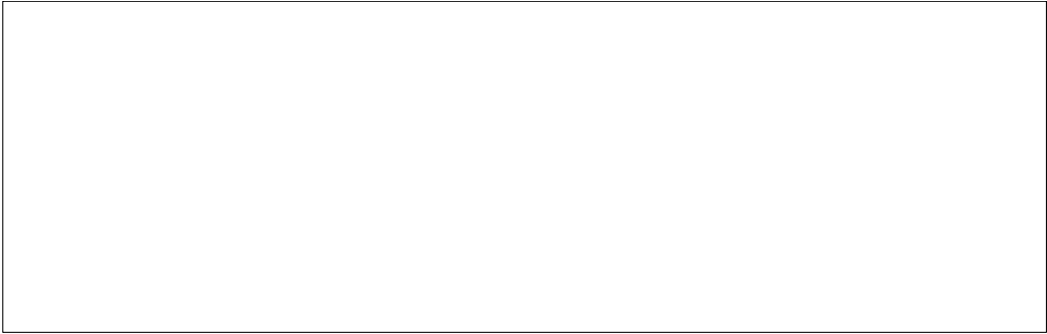
Coverage
(Norm. ATAC Signal Range (0–0.6) by ReadsInTSS)



L5.6.NR4A2

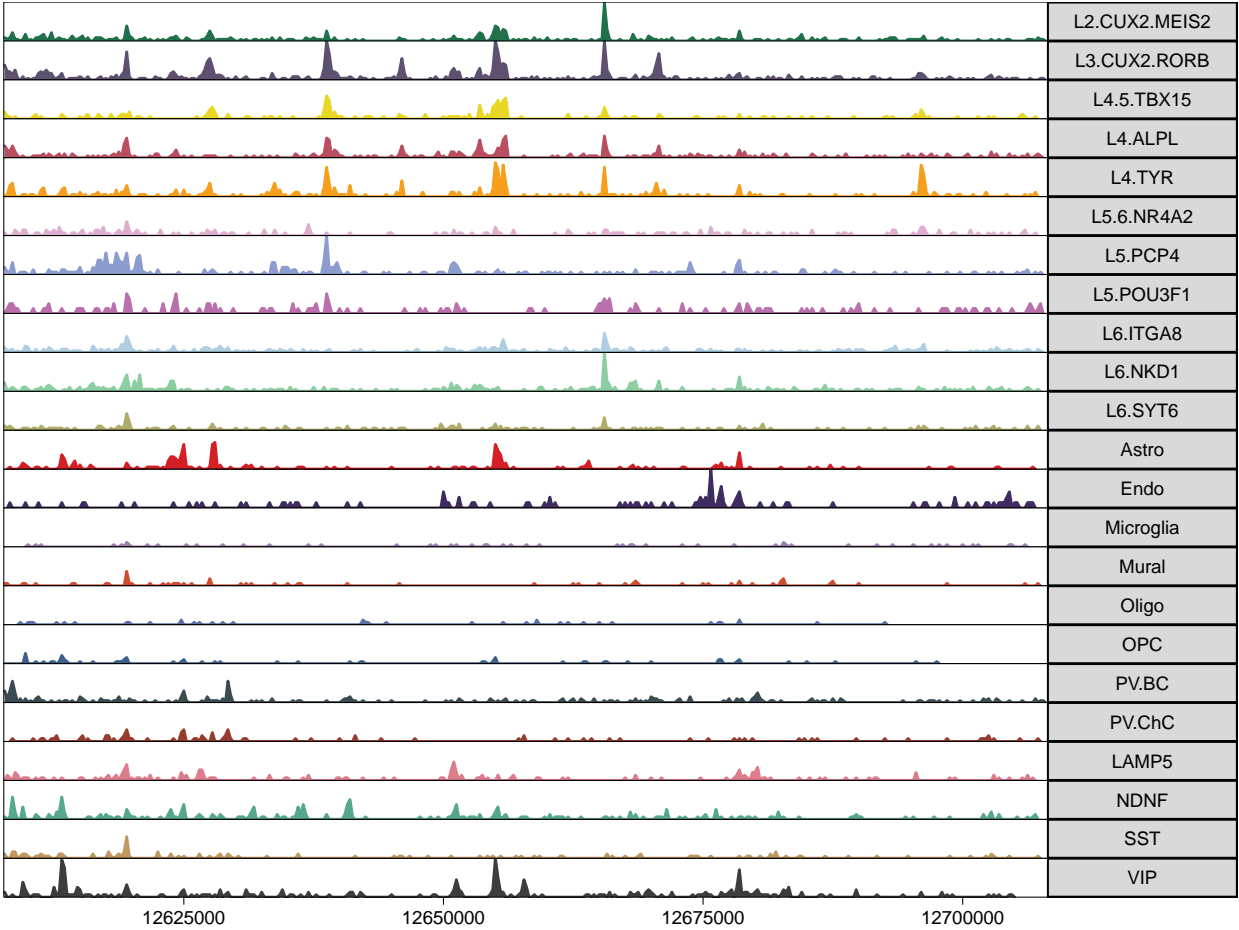


Peaks

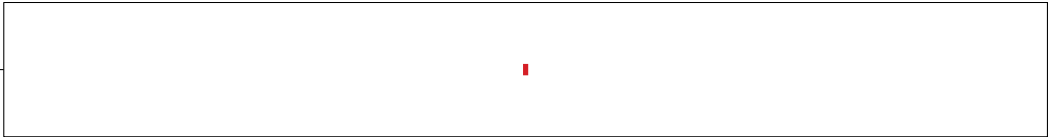


GeneTrack

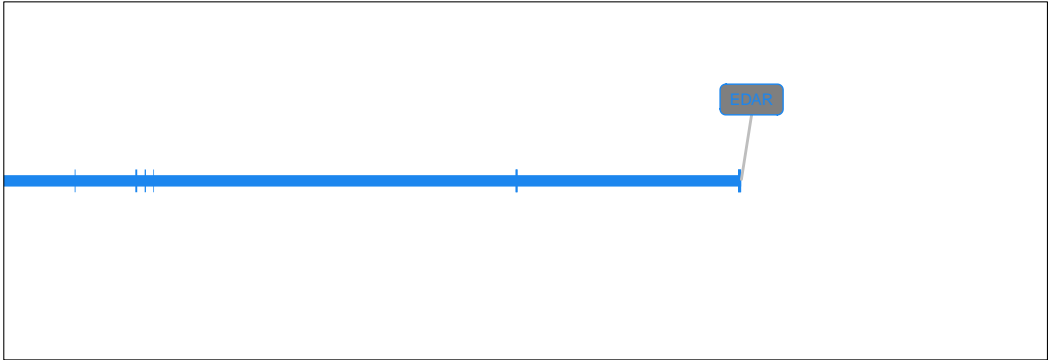
Coverage
(Norm. ATAC Signal Range (0–0.58) by ReadsInTSS)



L5.6.NR4A2



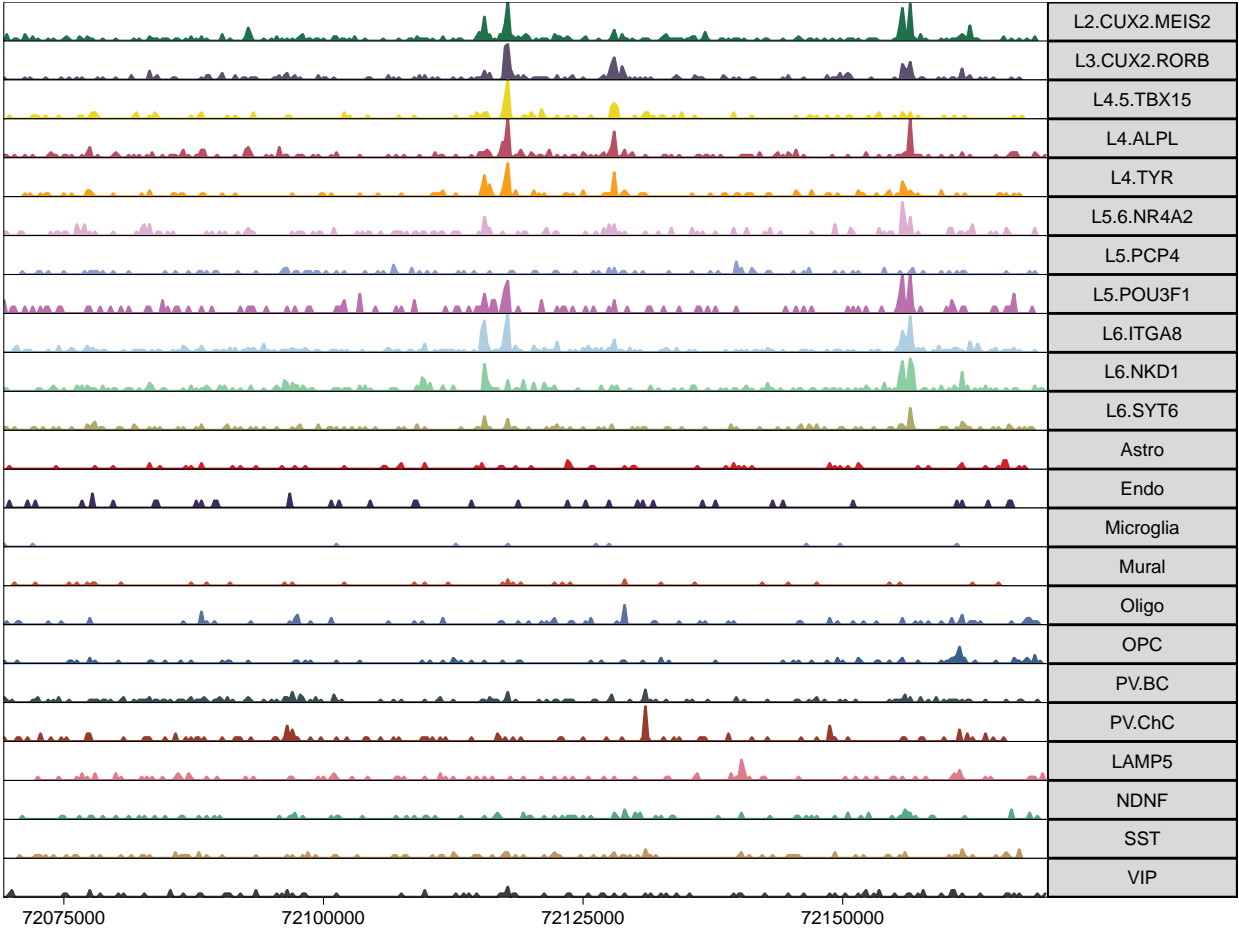
Peaks



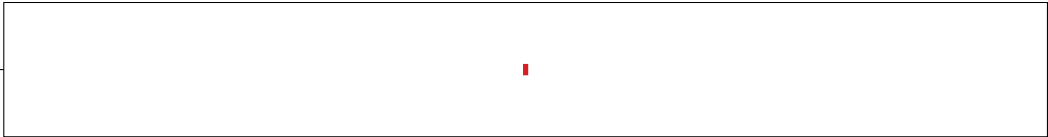
Genes

chr5:72069215–72169716 Marker Peak Tracks for: L5.6.NR4A2

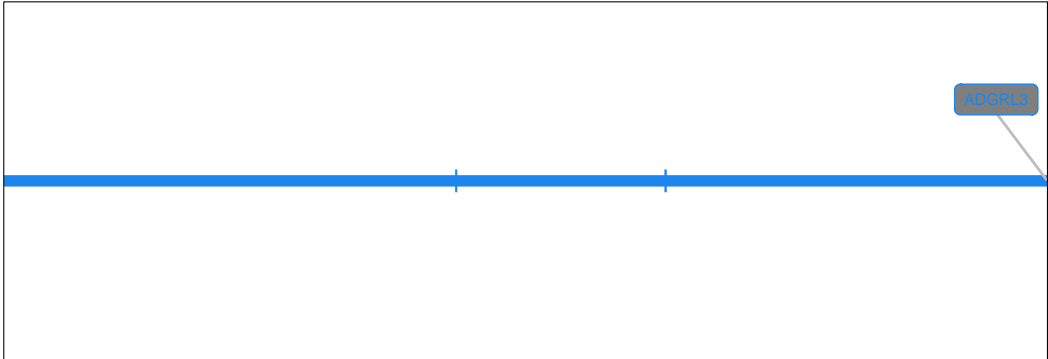
Coverage
(Norm. ATAC Signal Range (0–0.44) by ReadsInTSS)



L5.6.NR4A2



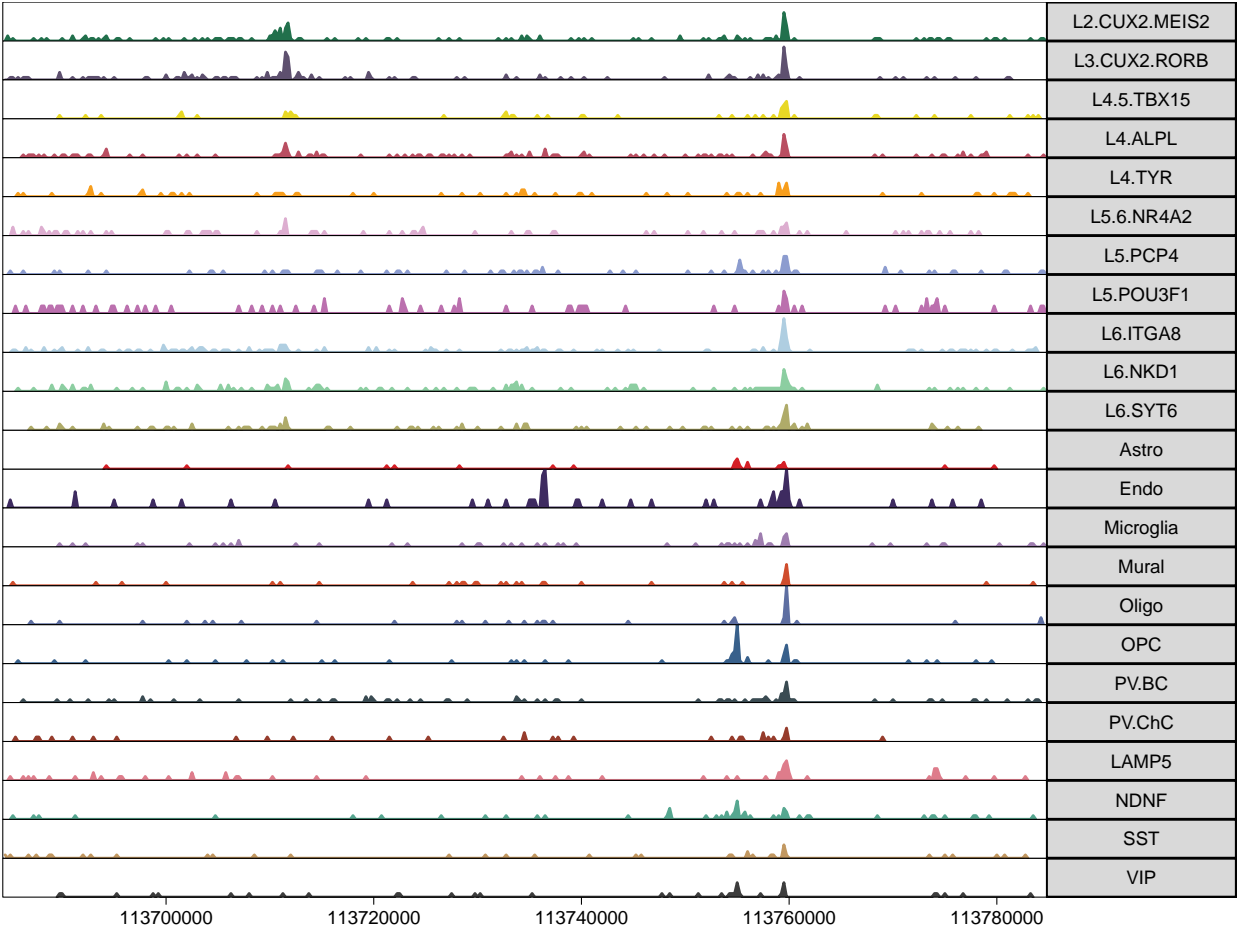
Peaks



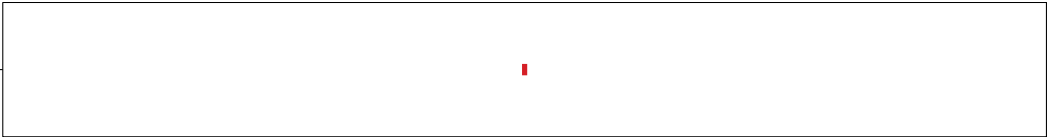
Genes

chr6:113684278–113784779 Marker Peak Tracks for: L5.6.NR4A2

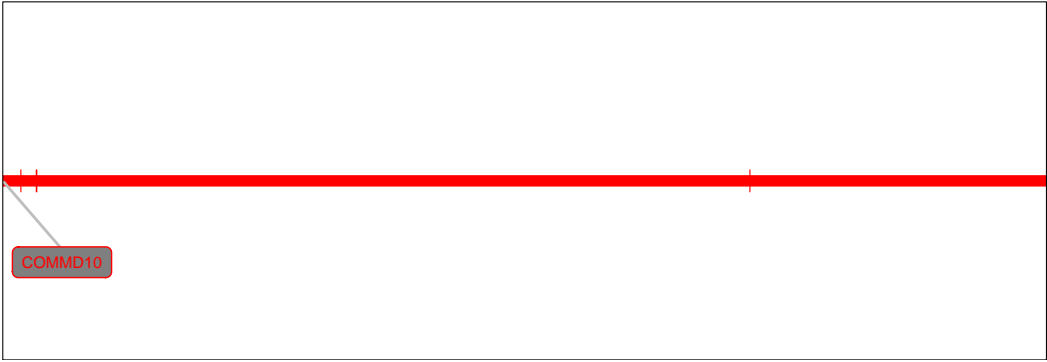
Coverage
(Norm. ATAC Signal Range (0–0.38) by ReadsInTSS)



L5.6.NR4A2



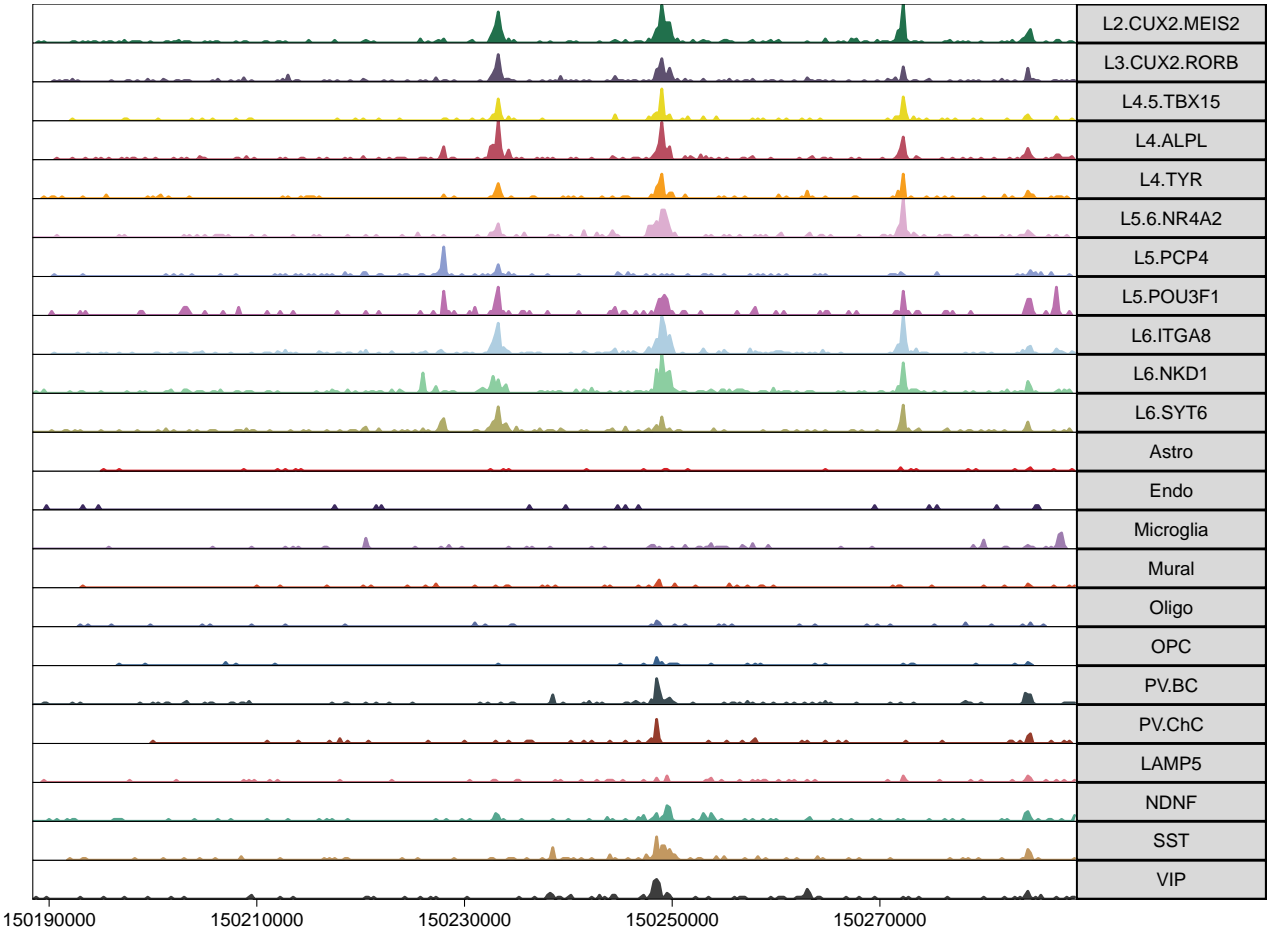
Peaks



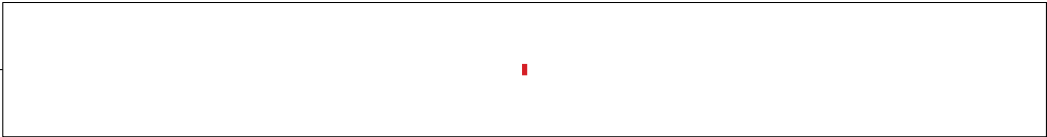
Genes

chr5:150188427–150288928 Marker Peak Tracks for: L5.6.NR4A2

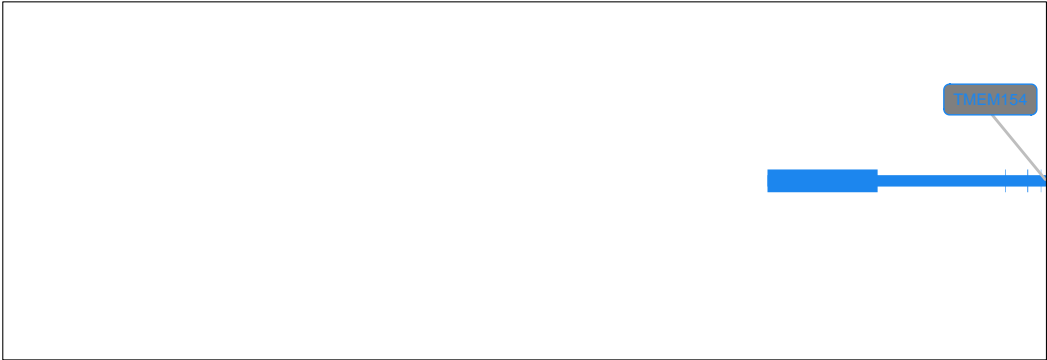
Coverage
(Norm. ATAC Signal Range (0–0.71) by ReadslnTSS)



L5.6.NR4A2

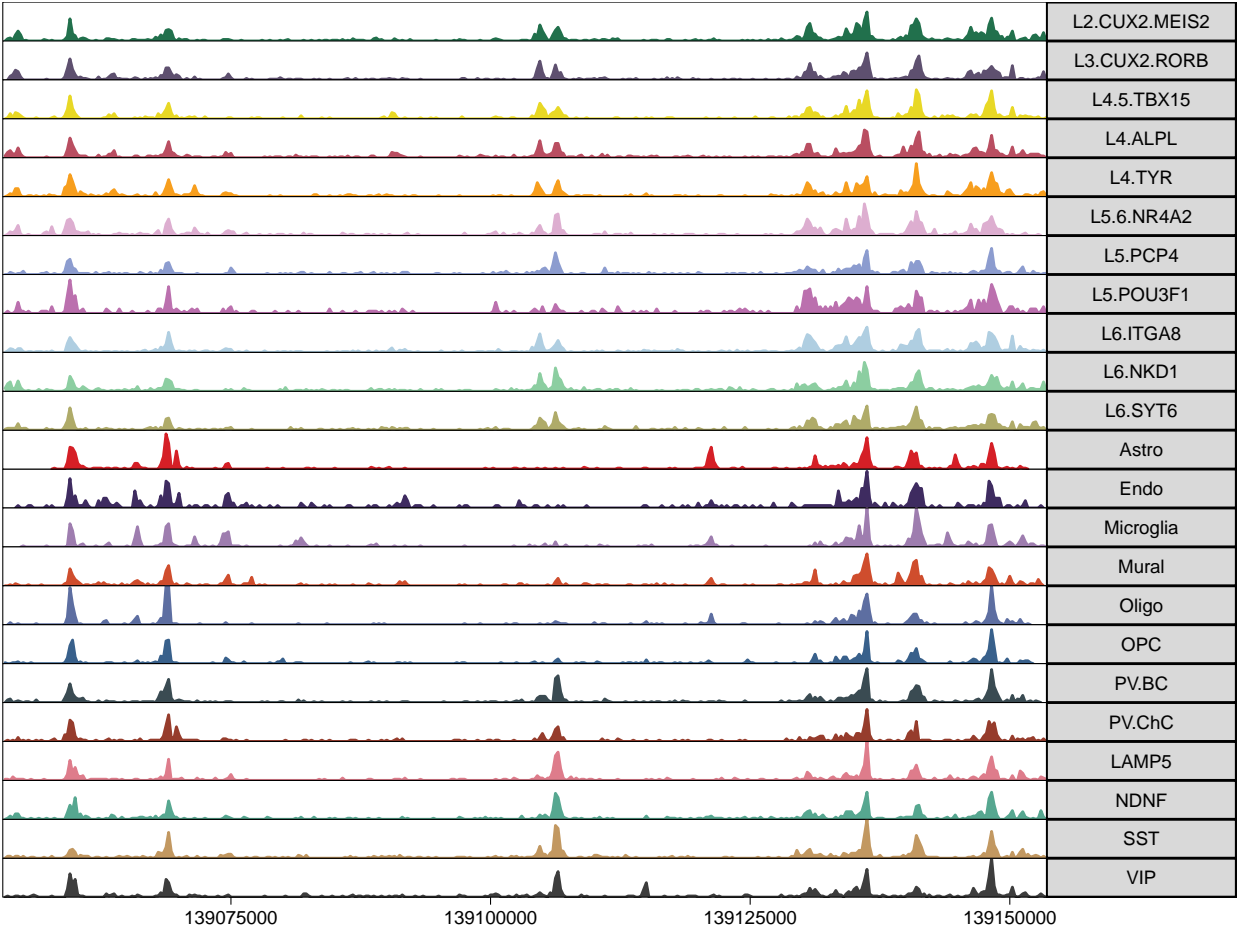


Peaks

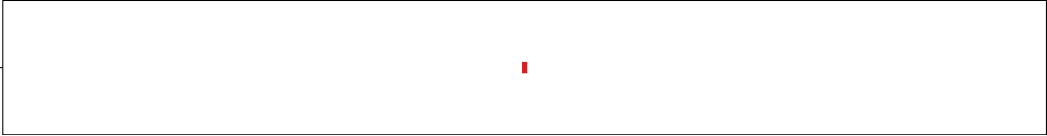


Genes

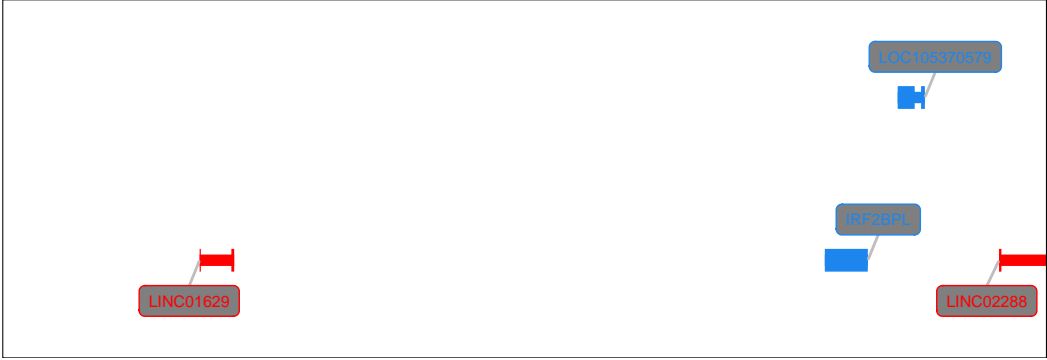
Coverage
(Norm. ATAC Signal Range (0–1.28) by ReadsInTSS)



L5.6.NR4A2



Peaks



Genes