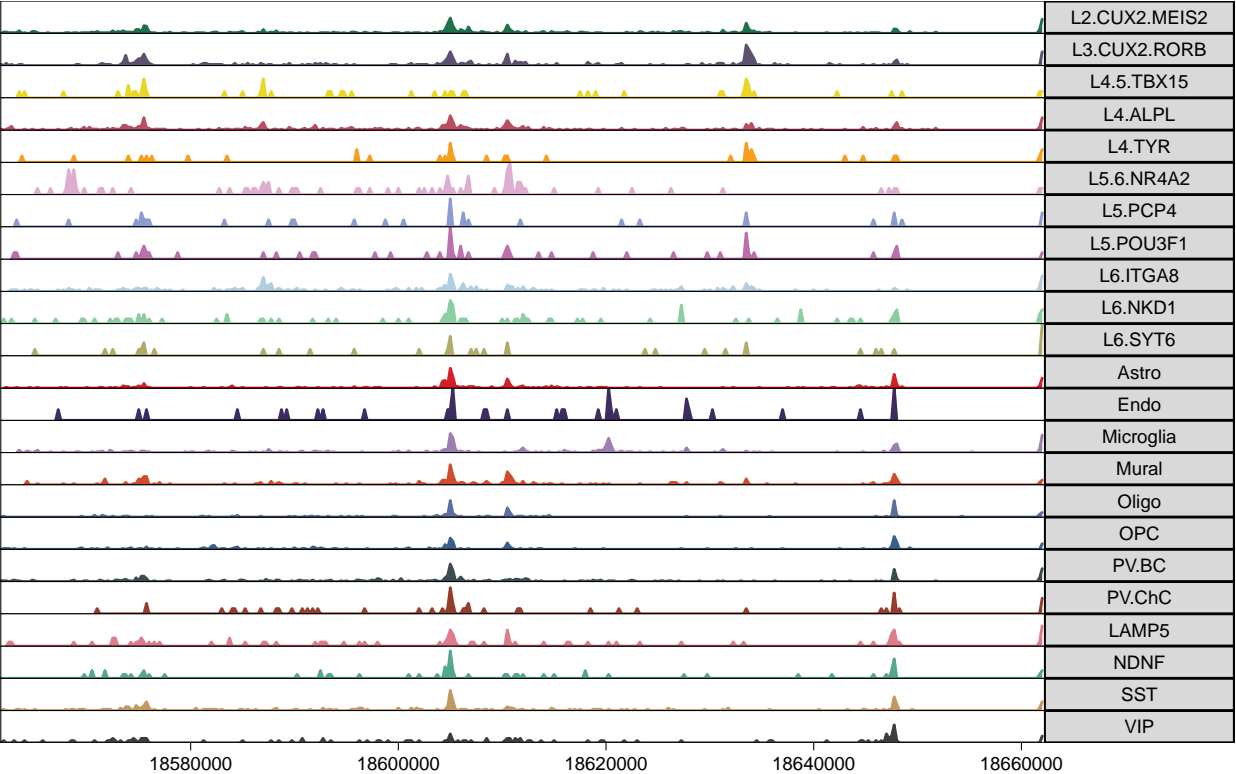
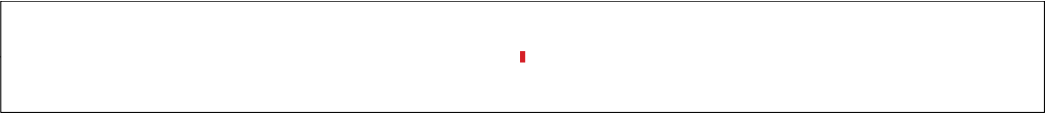


Coverage
(Norm. ATAC Signal Range 0-1.18) by ReadsInTSS)



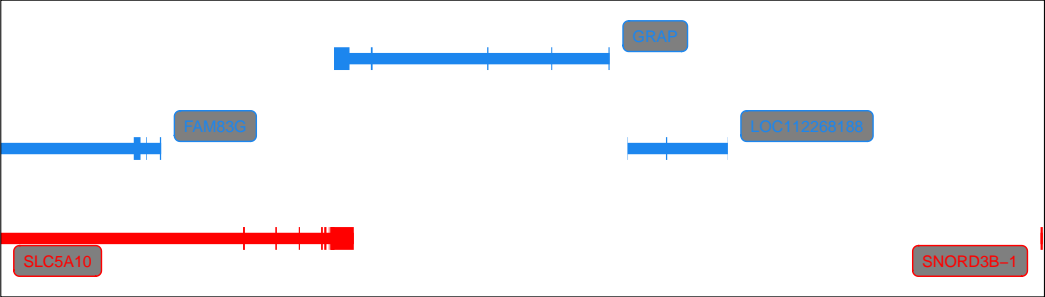
L5.6.NR4A2



Peaks



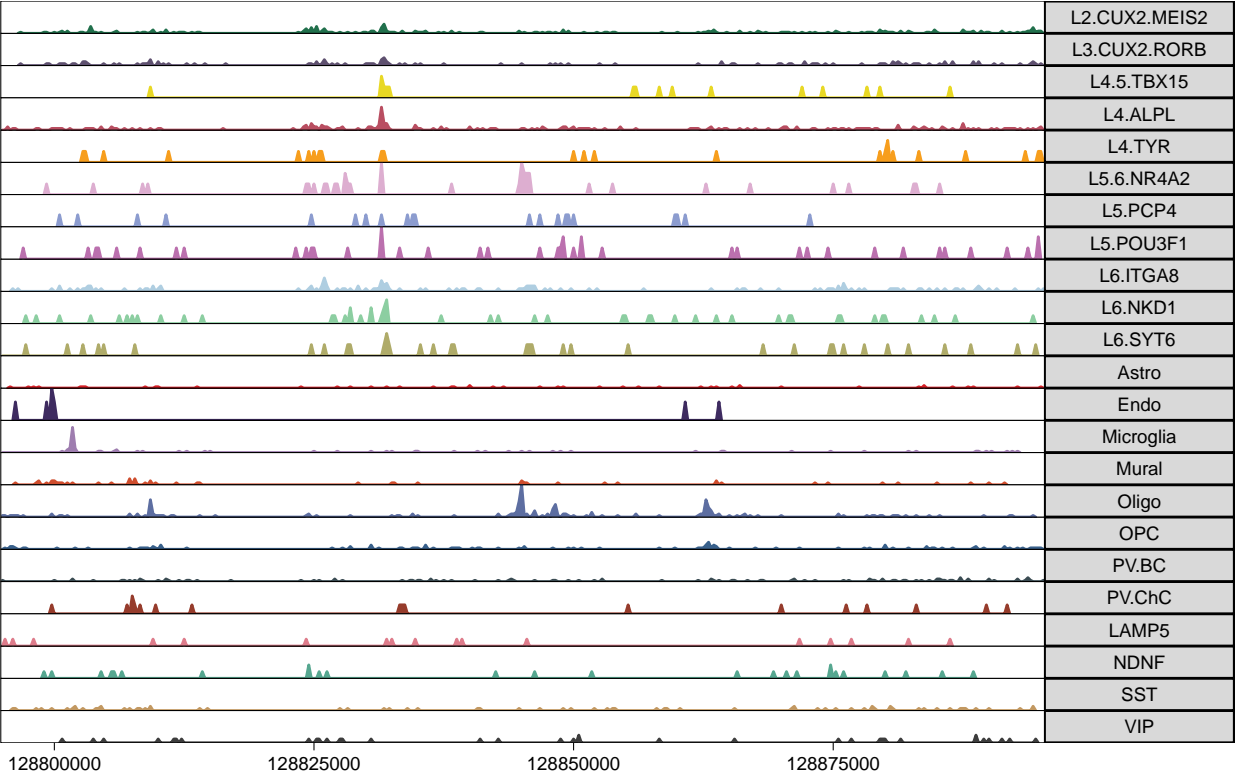
LoopTrack



Genes

chr12:128794841–128895342 Marker Peak Tracks for: L5.6.NR4A2

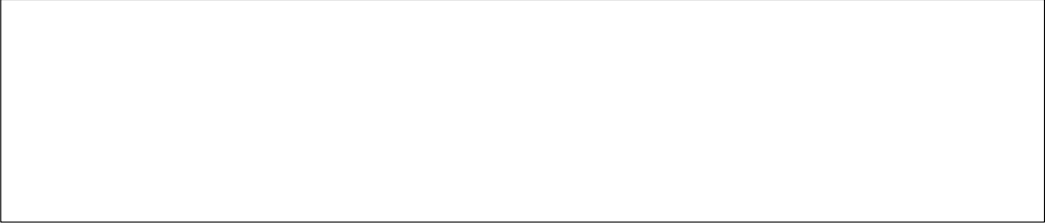
Coverage
(Norm. ATAC Signal Range 0–0.69) by ReadsInTSS)



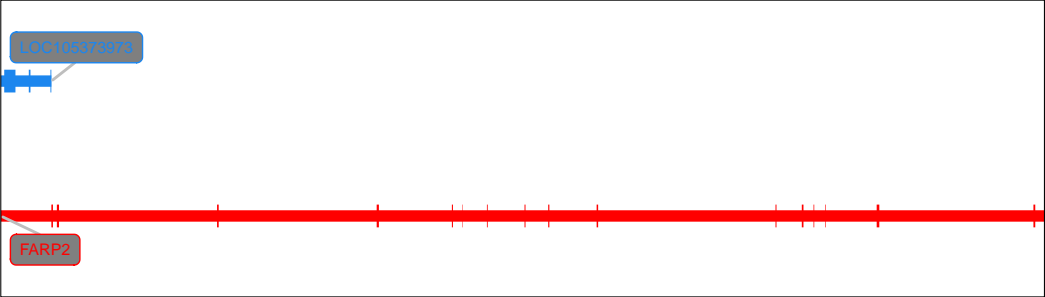
L5.6.NR4A2



Peaks

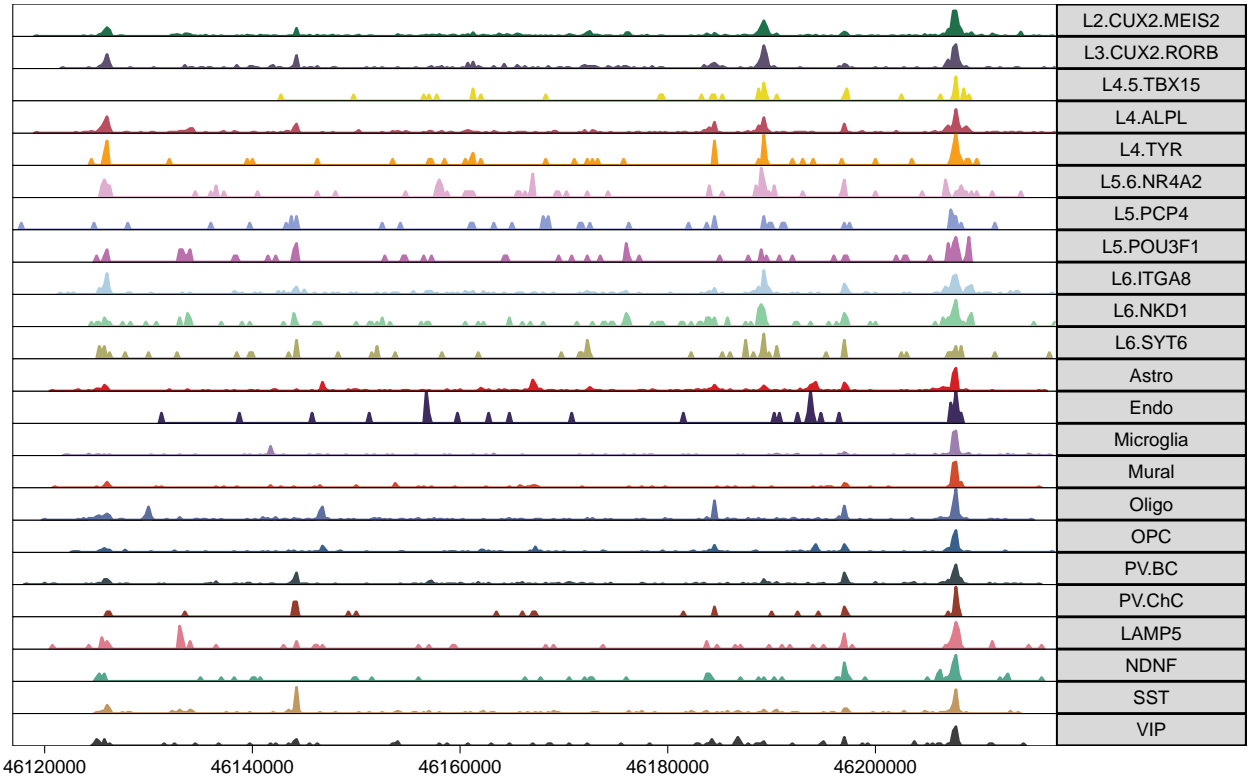


LoopTrack

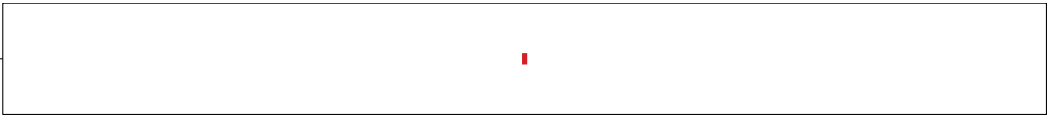


Genes

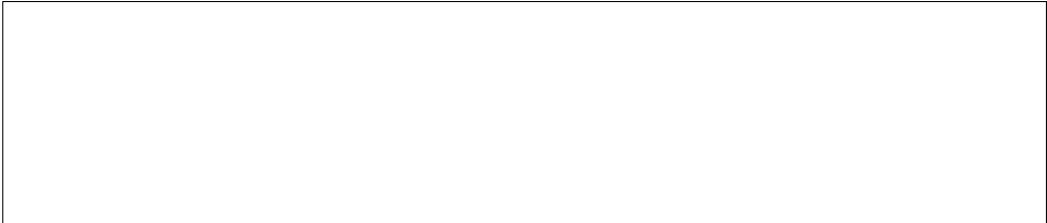
Coverage
(Norm. ATAC Signal Range 0-1.24) by ReadsInTSS)



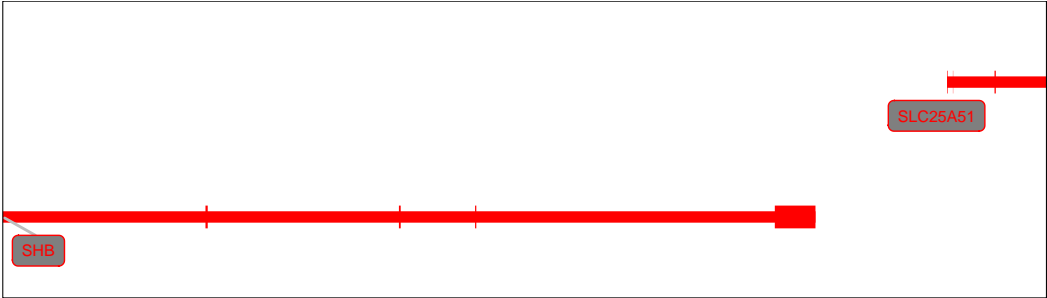
L5.6.NR4A2



Peaks

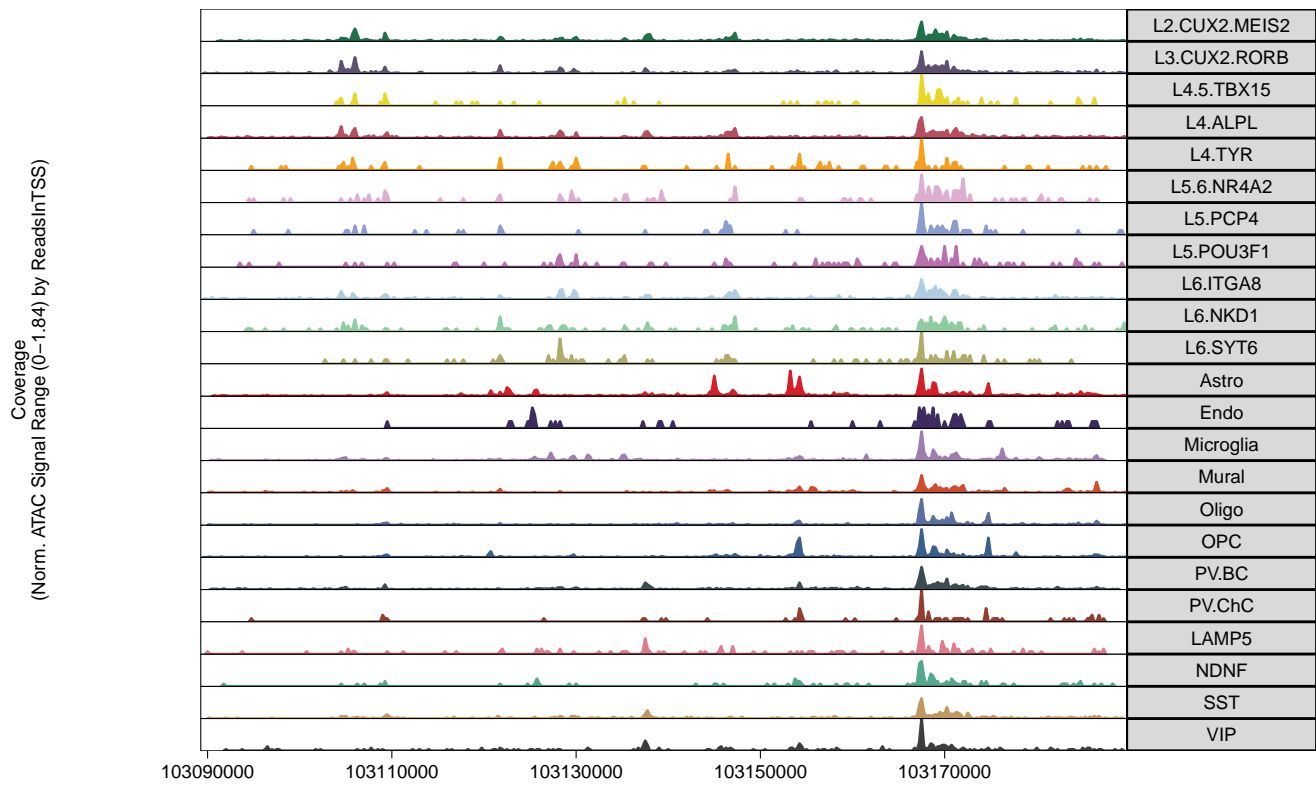


LoopTrack

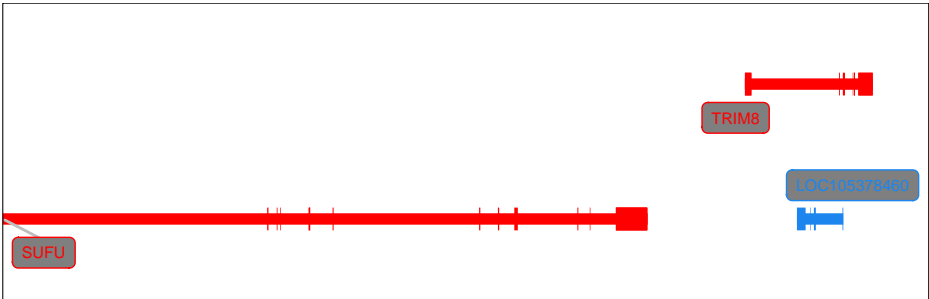
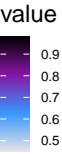
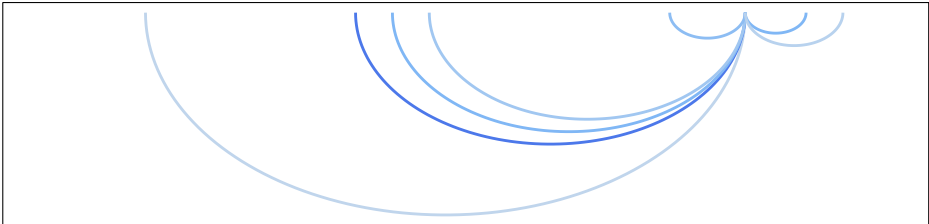


Genes

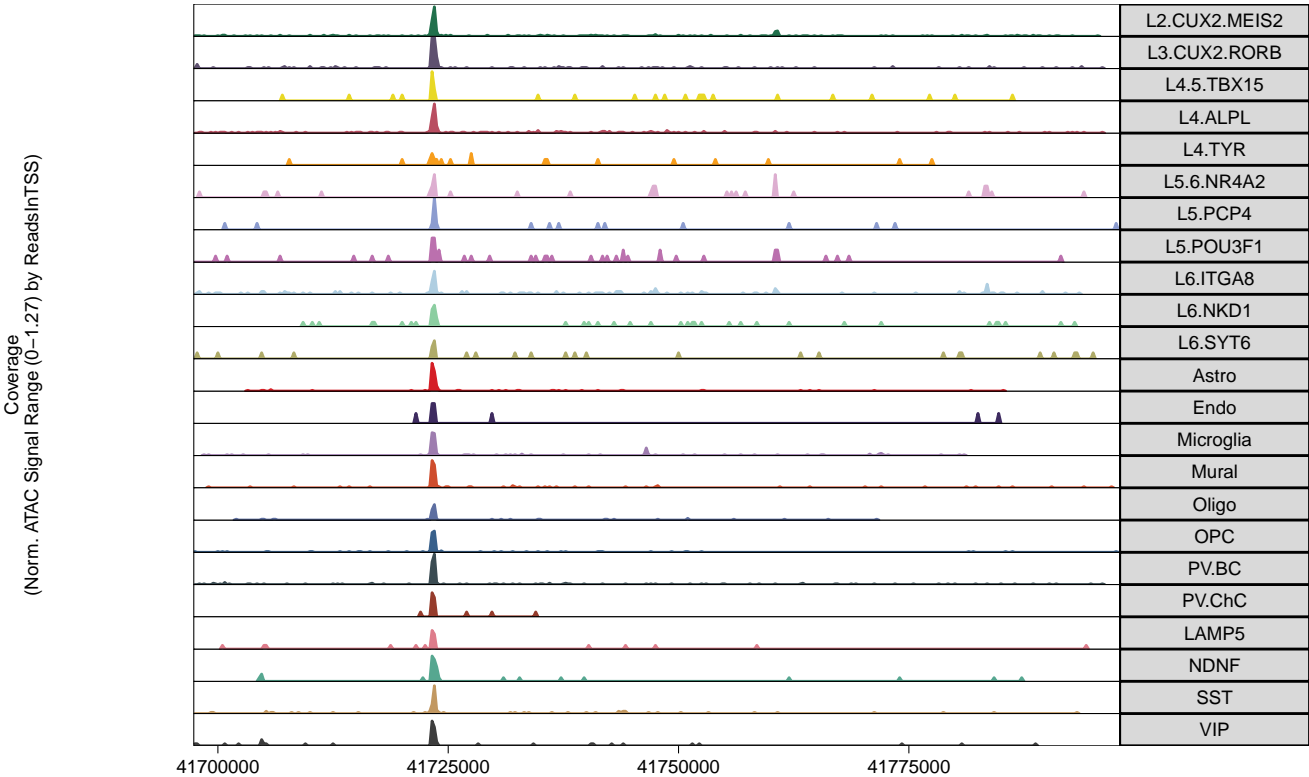
chr9:103089261-103189762 Marker Peak Tracks for: L5.6.NR4A2



L5.6.NR4A2



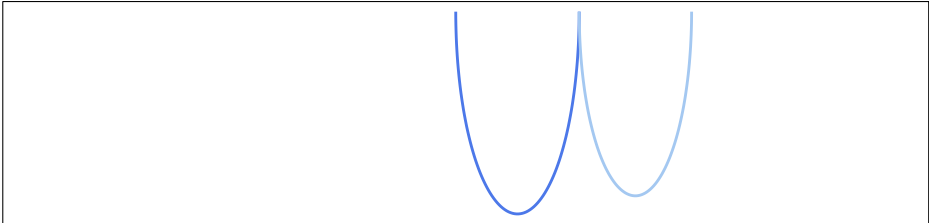
chr3:41697385–41797886 Marker Peak Tracks for: L5.6.NR4A2



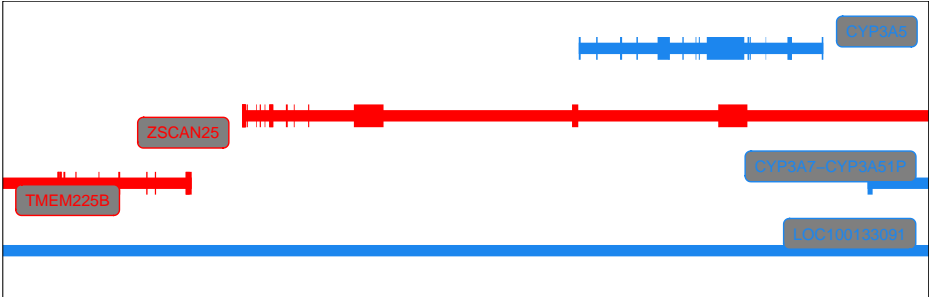
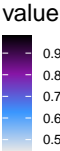
L5.6.NR4A2



Peaks

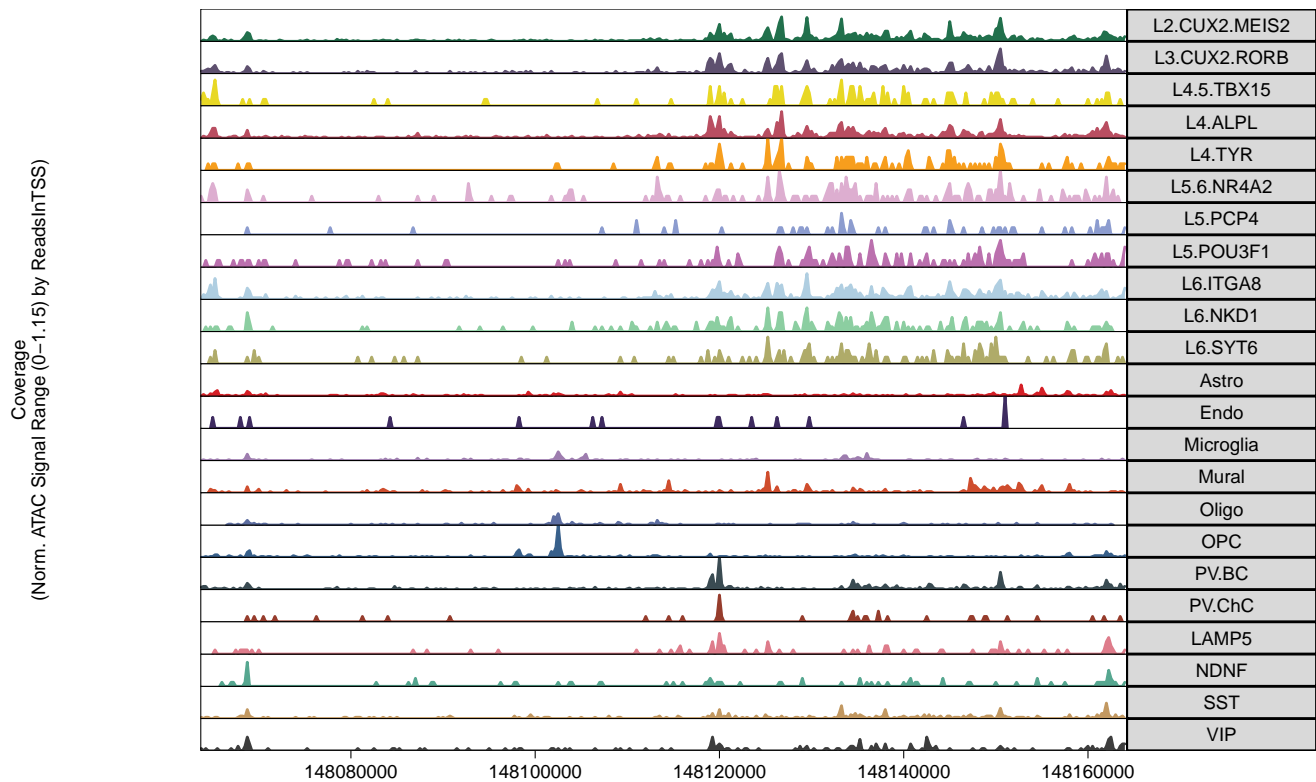


Peak2GeneLinks

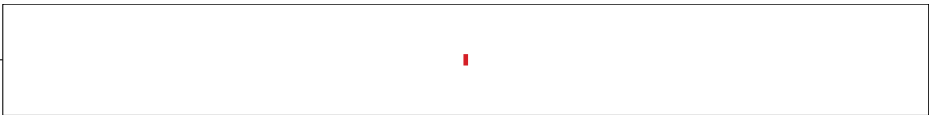


Genes

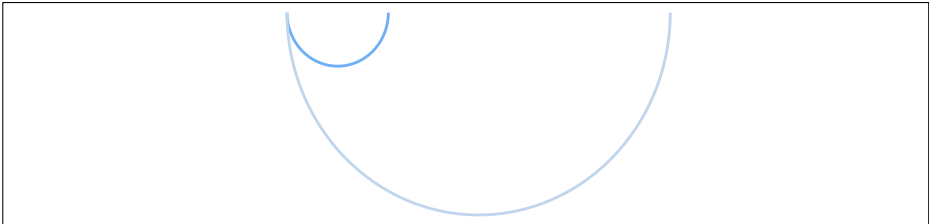
chr6:148063704–148164205 Marker Peak Tracks for: L5.6.NR4A2



L5.6.NR4A2



Peaks



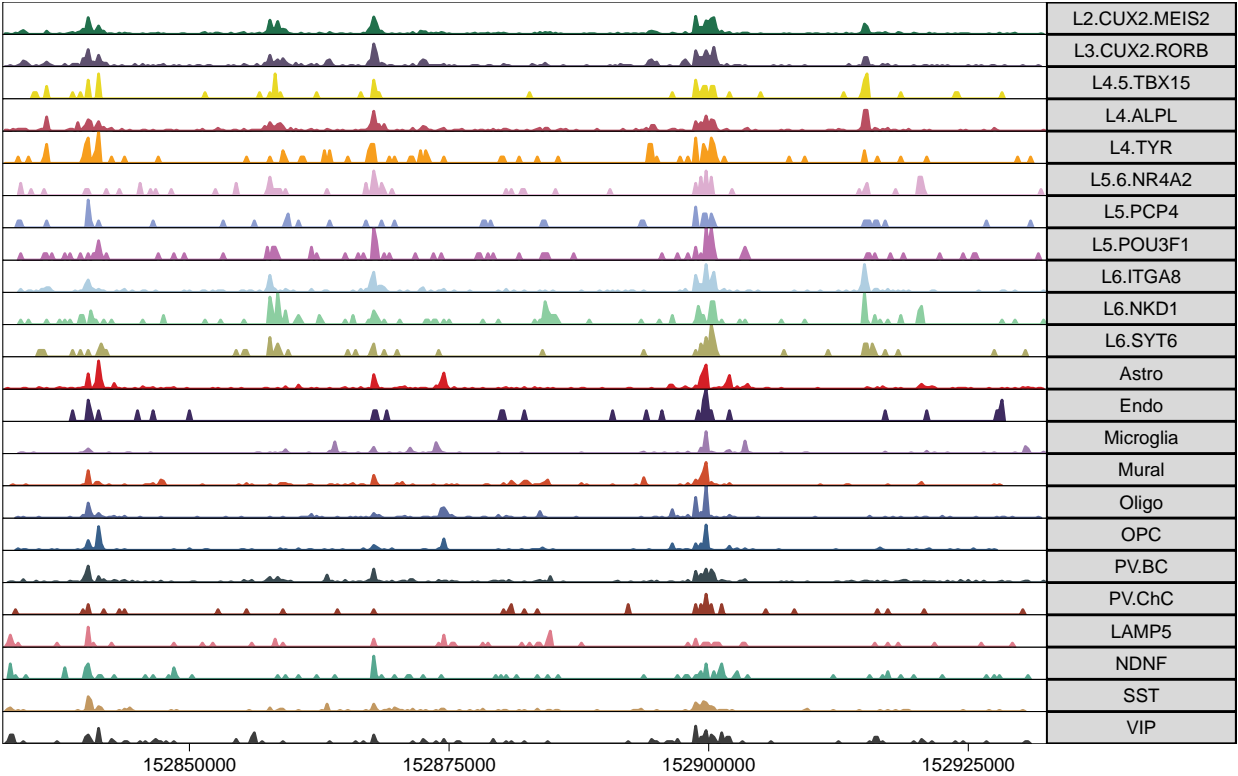
Peak2GeneLinks

value

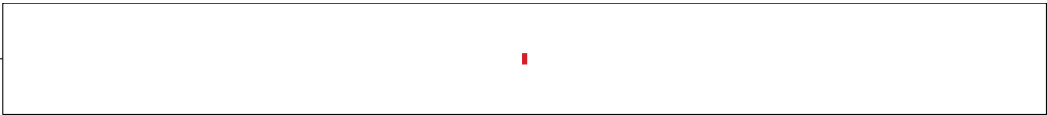


Genes

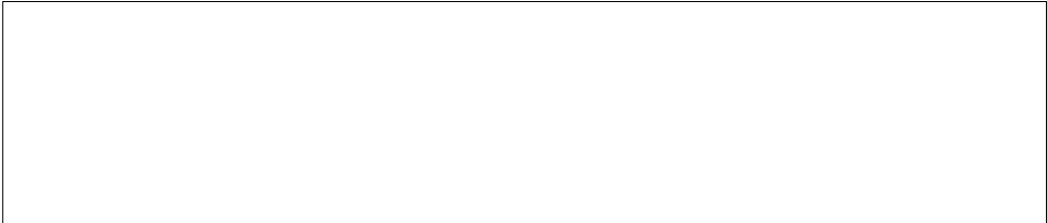
Coverage
(Norm. ATAC Signal Range 0-1.19) by ReadsInTSS)



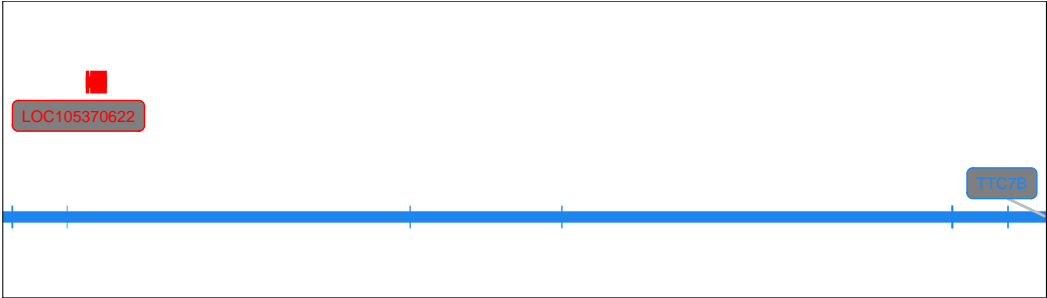
L5.6.NR4A2



Peaks

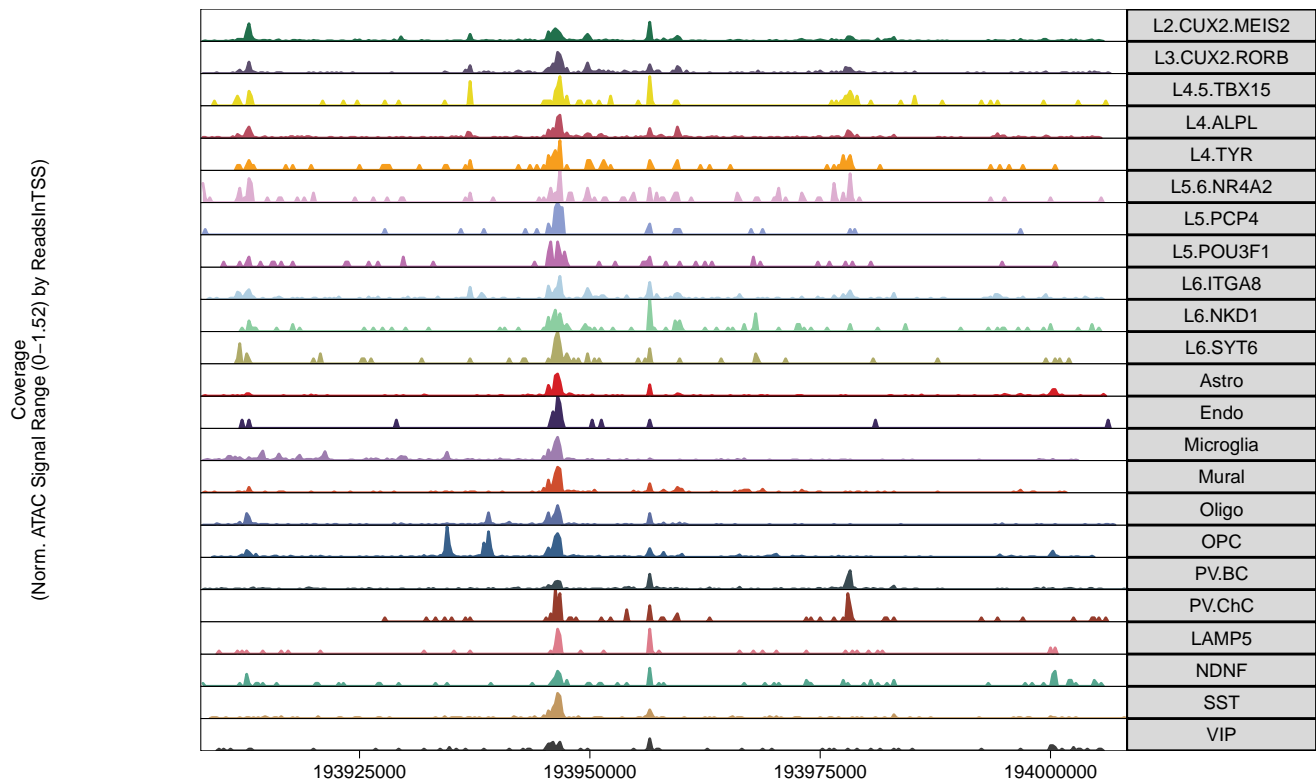


LoopTrack



Genes

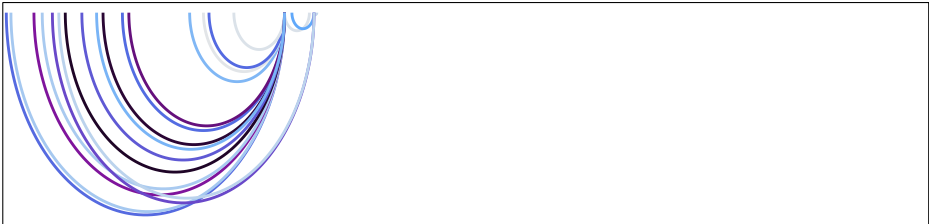
chr1:193907769–194008270 Marker Peak Tracks for: L5.6.NR4A2



L5.6.NR4A2

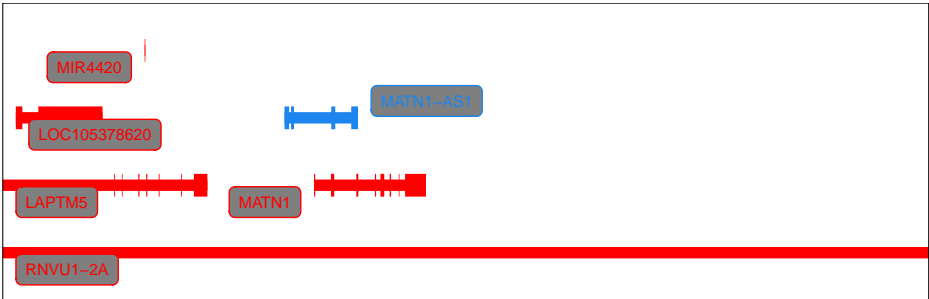
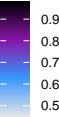


Peaks



Peak2GeneLinks

value



Genes