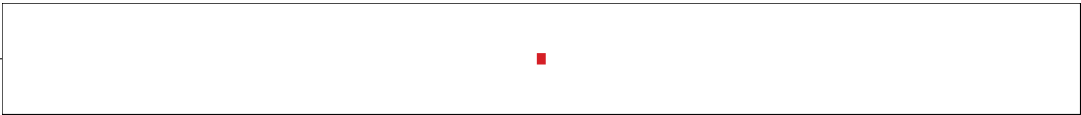
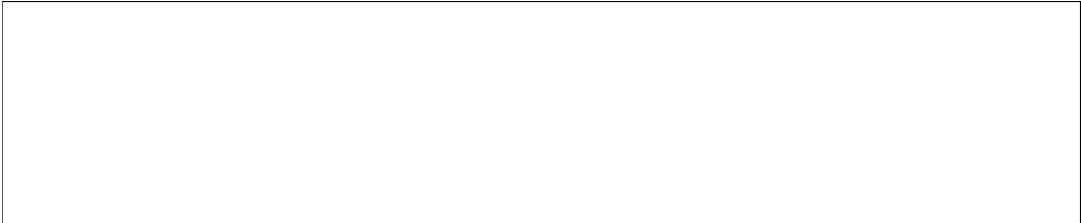


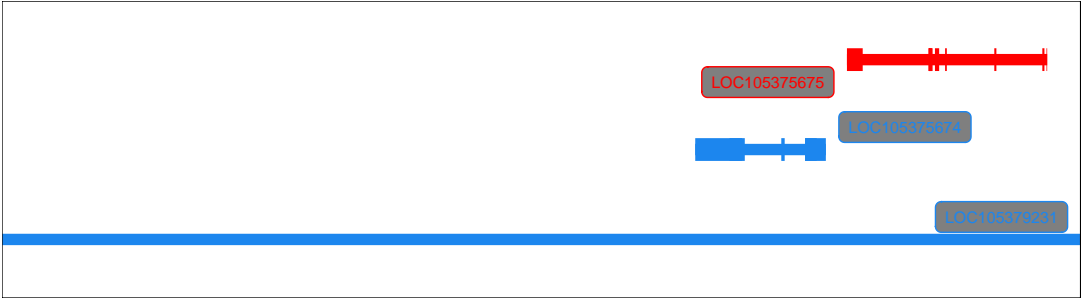
L5.PCP4



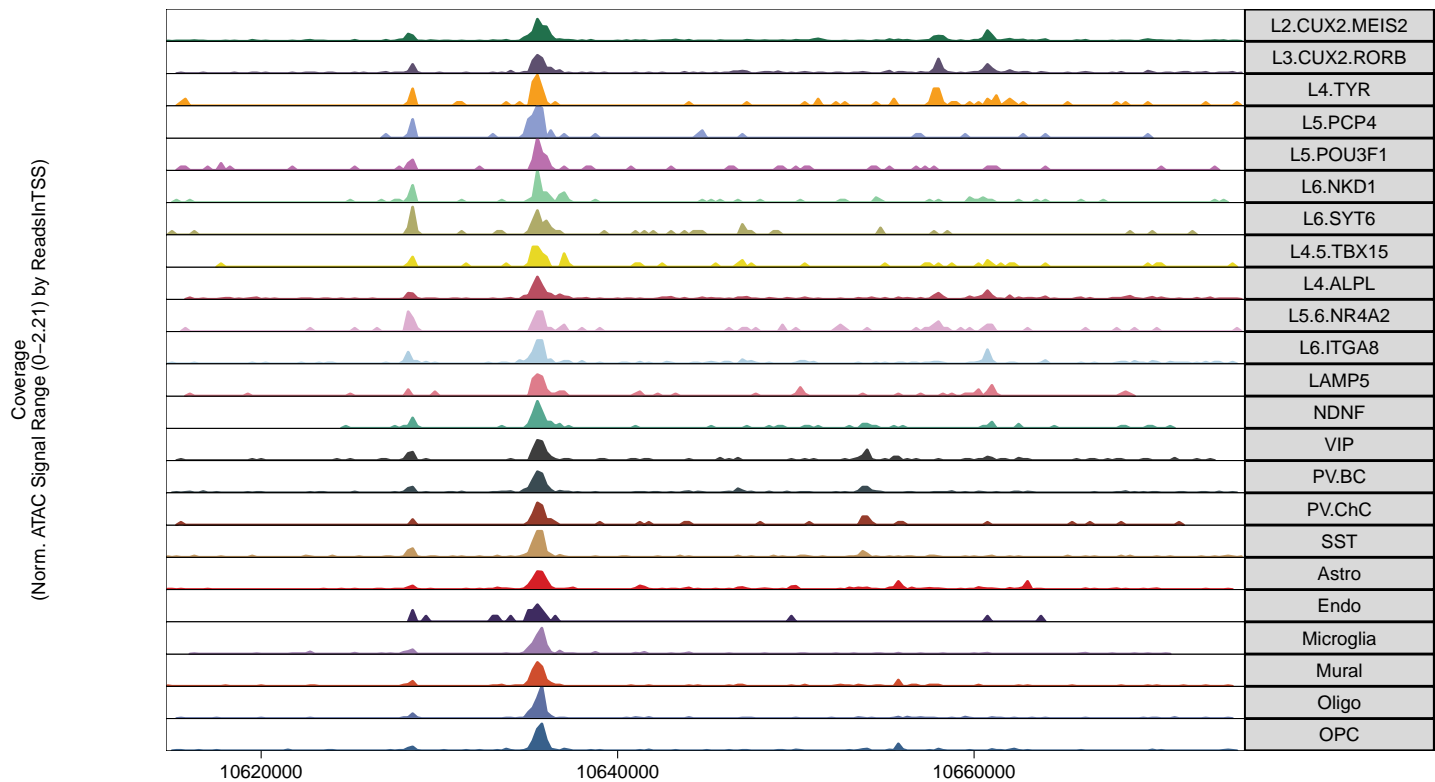
Peaks



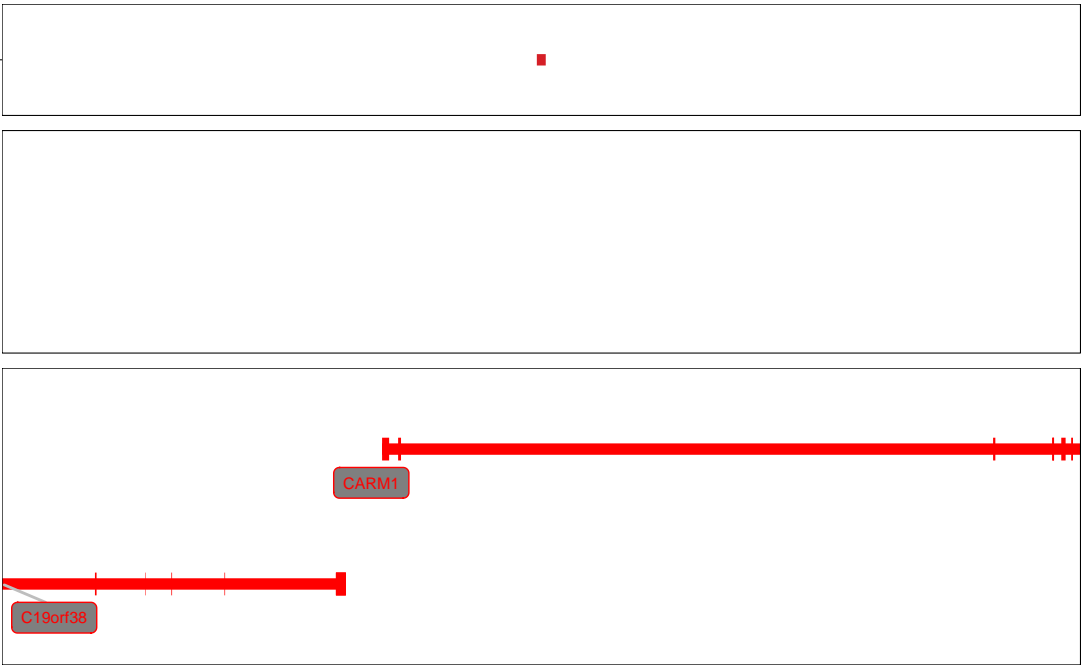
LoopTrack



chr19:10614667-10675168 Marker Peak Tracks for: L5.PCP4



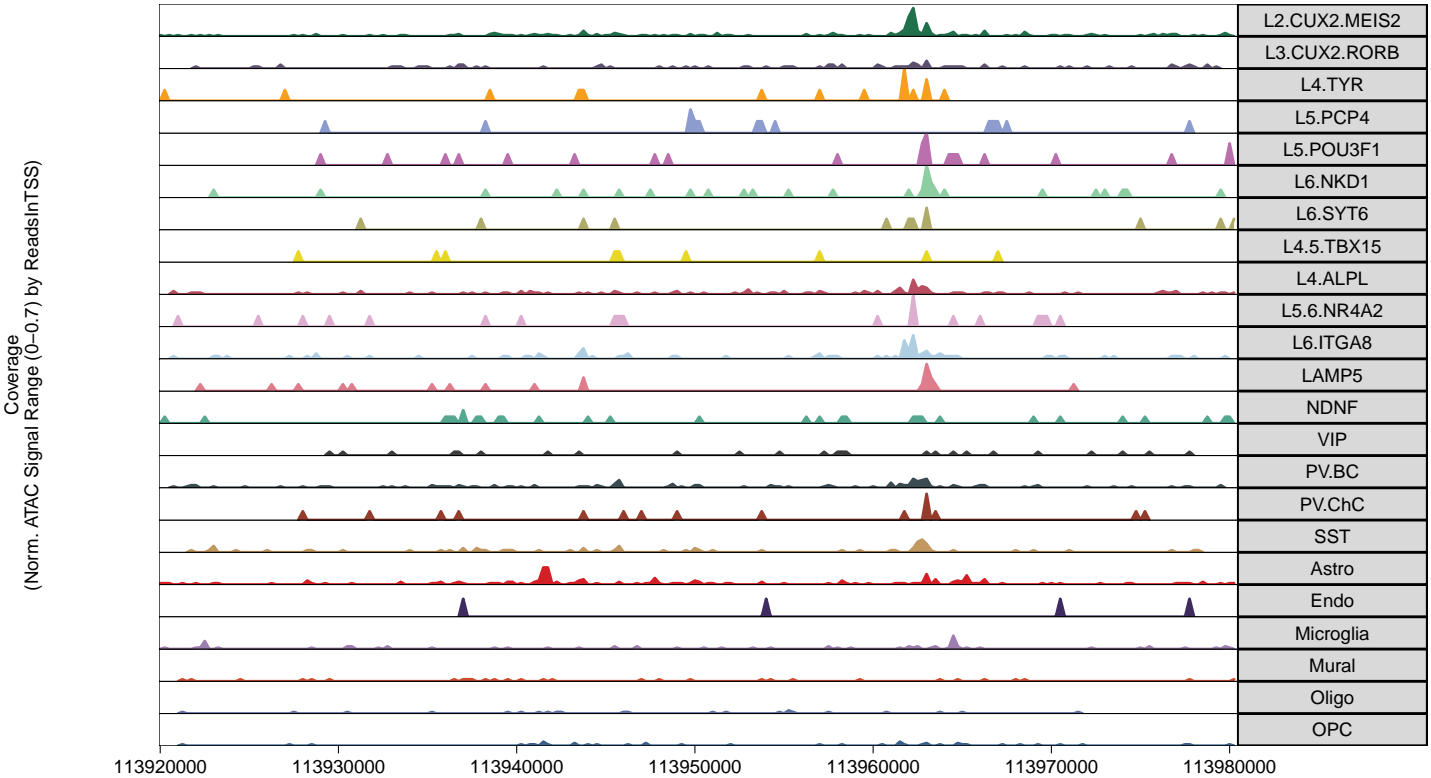
L5.PCP4



Peaks

LoopTrack

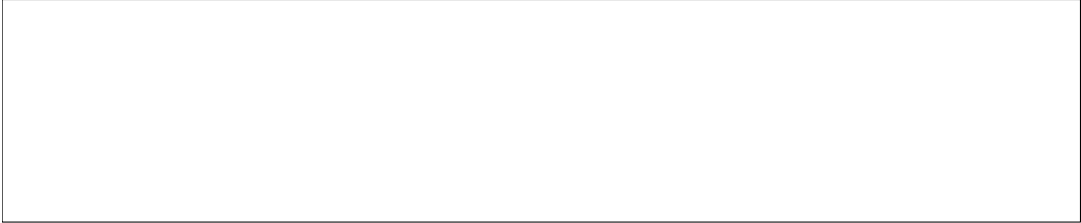
Genes



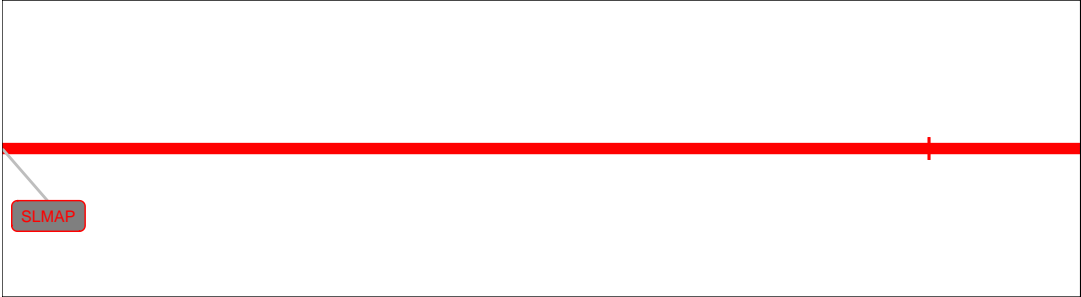
L5.PCP4



Peaks

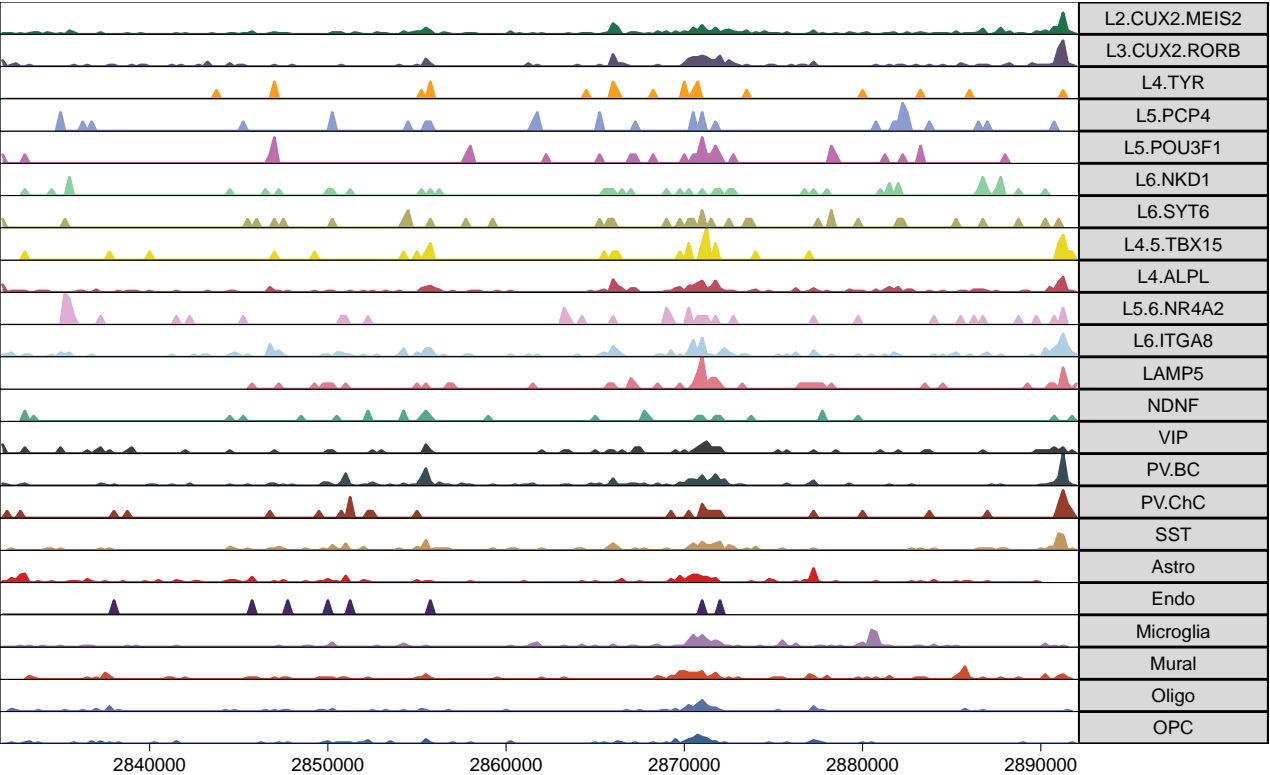


LoopTrack



Genes

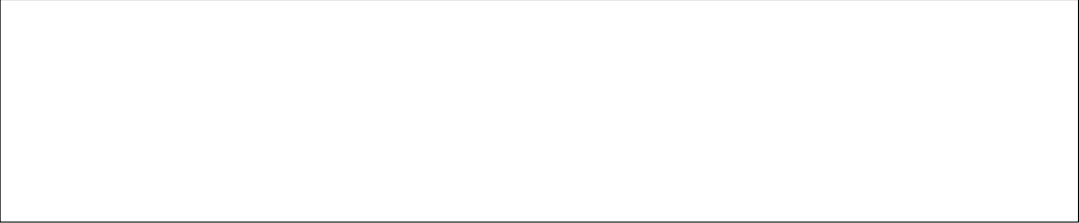
Coverage
(Norm. ATAC Signal Range (0–0.89) by ReadsInTSS)



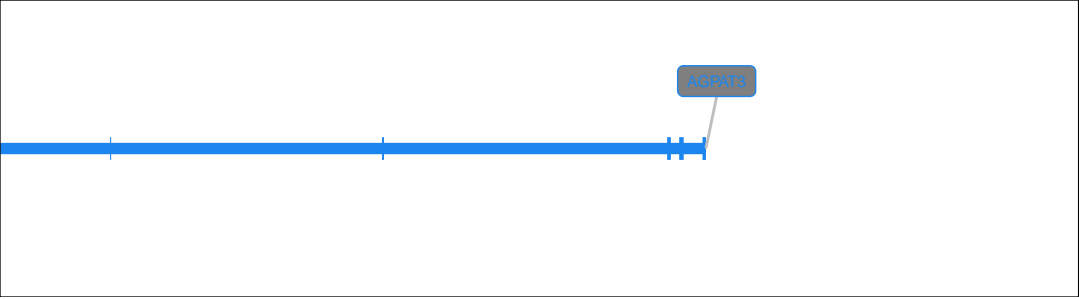
L5.PCP4



Peaks

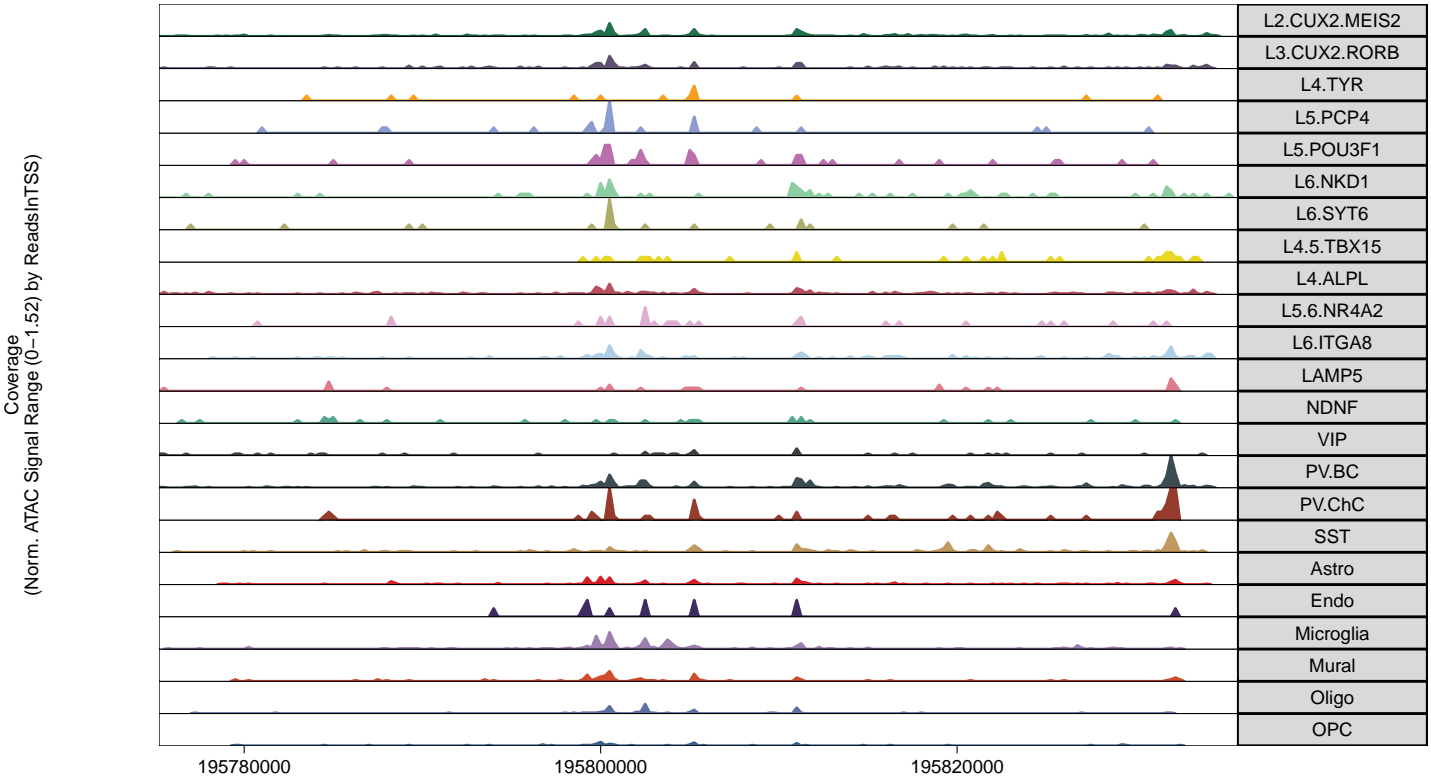


LoopTrack



Genes

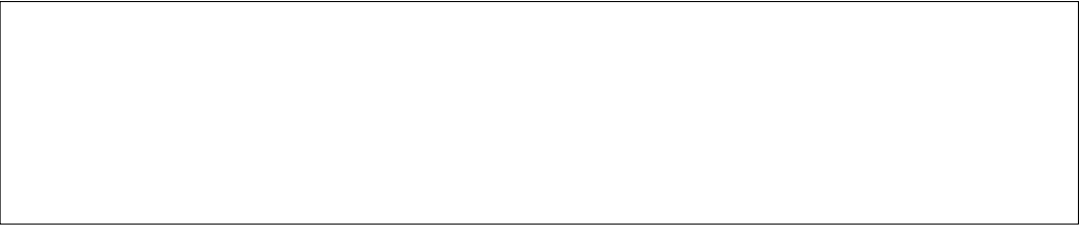
chr1:195775227-195835728 Marker Peak Tracks for: L5.PCP4



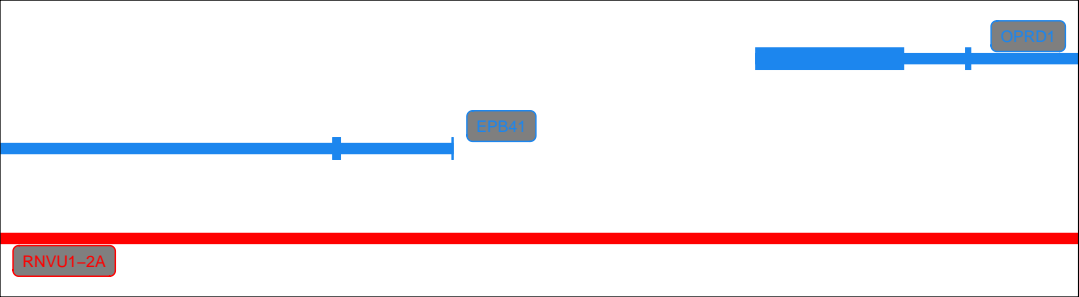
L5.PCP4



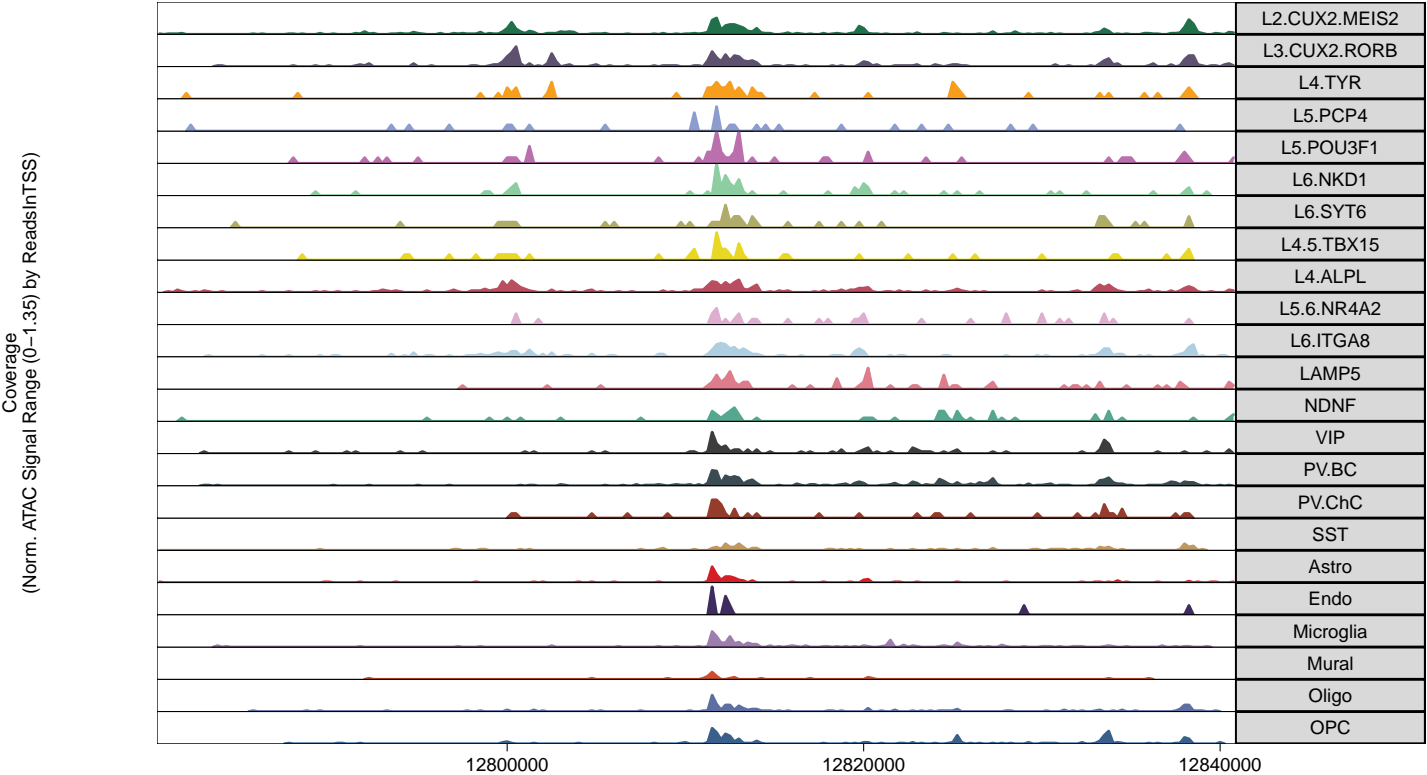
Peaks



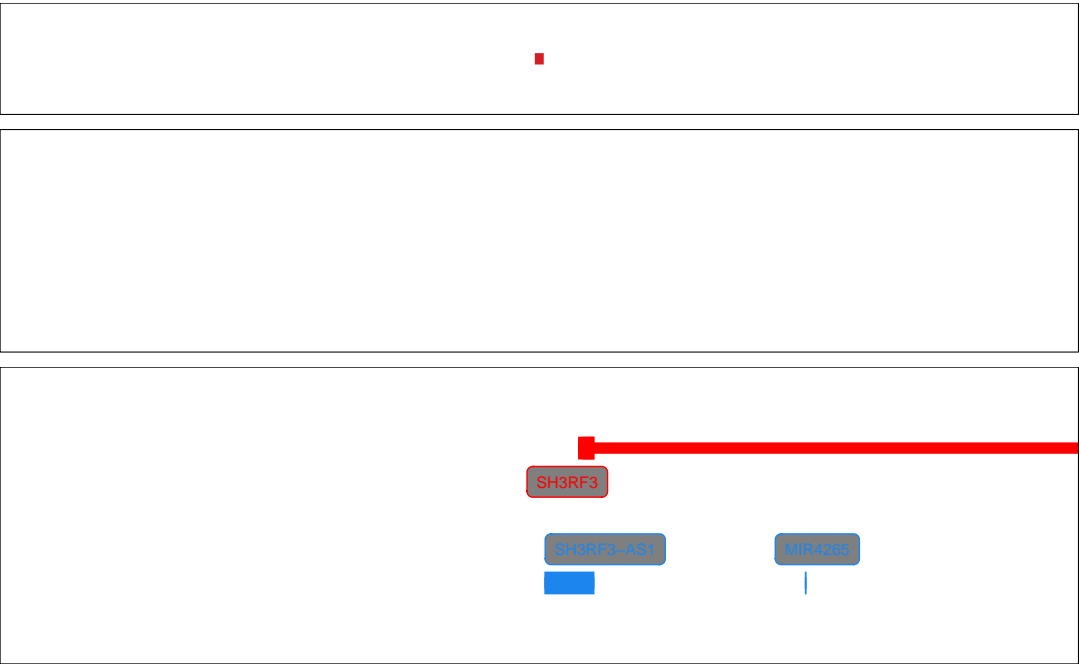
LoopTrack

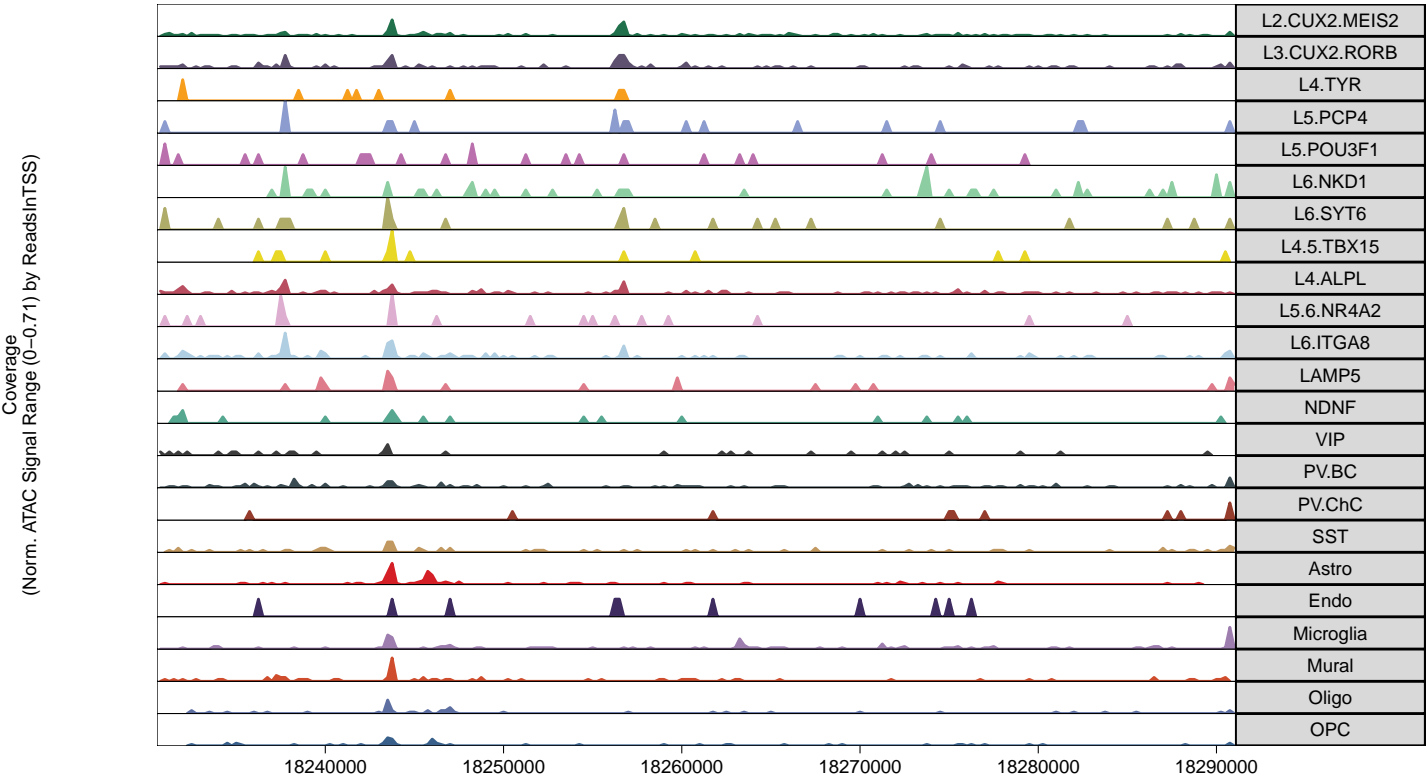


Genes

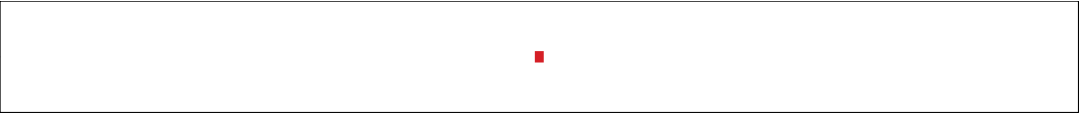


L5.PCP4

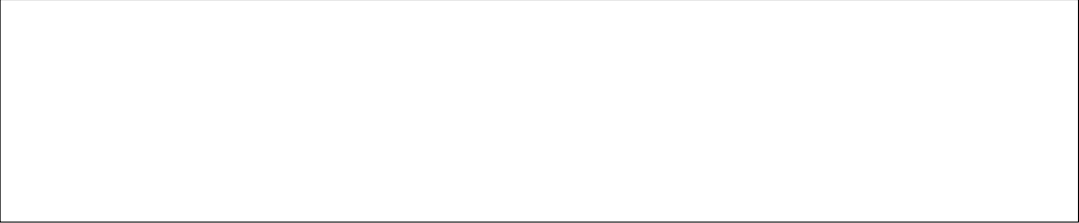




L5.PCP4



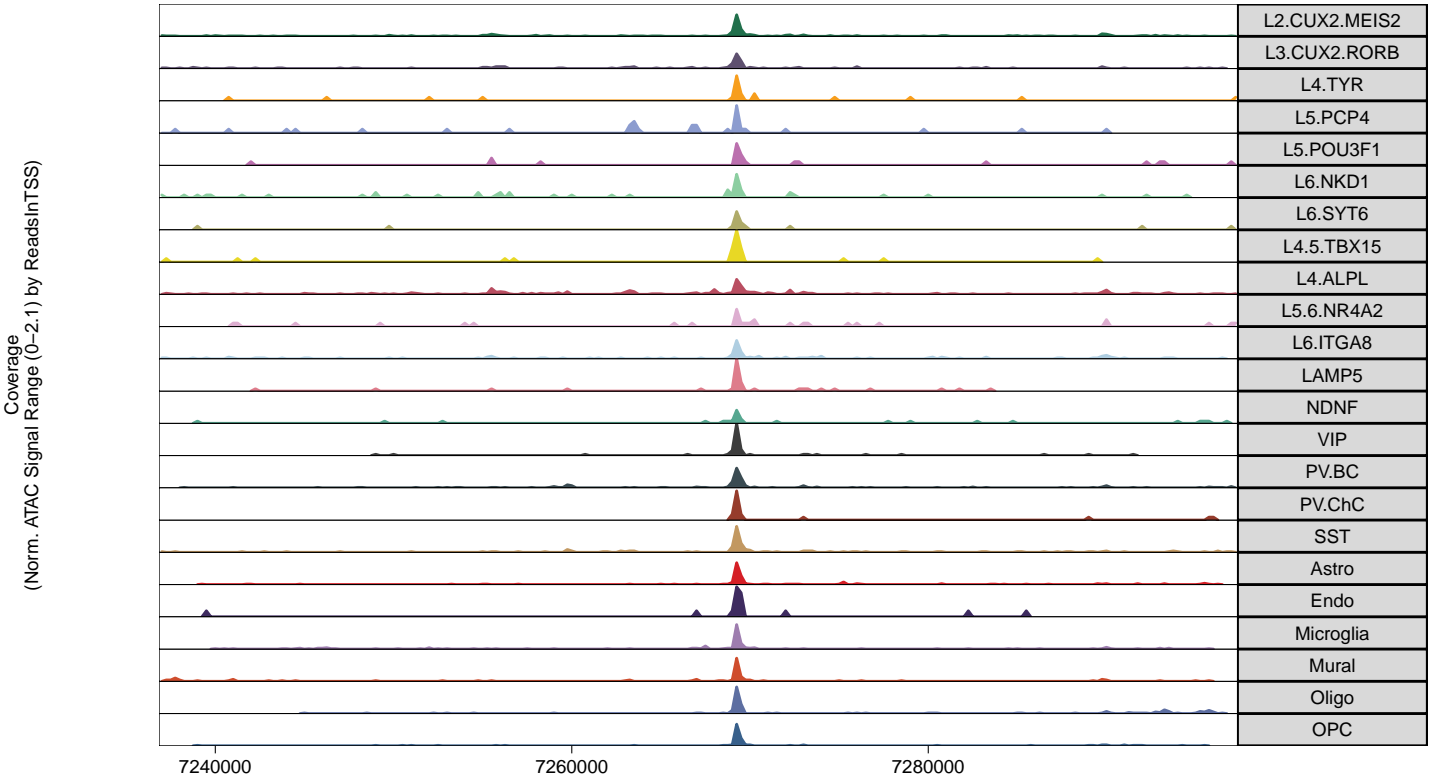
Peaks



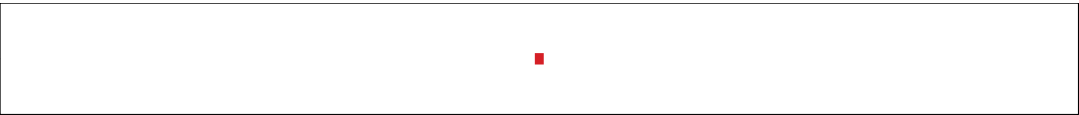
LoopTrack



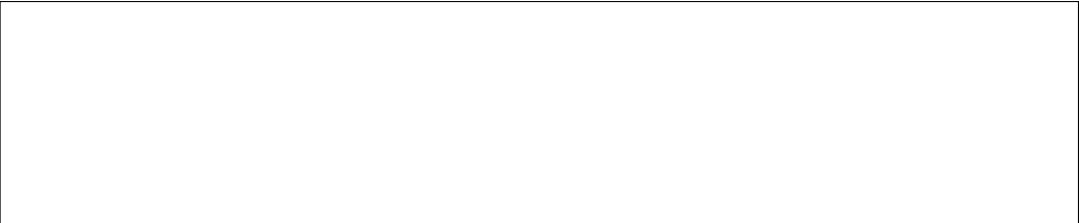
Genes



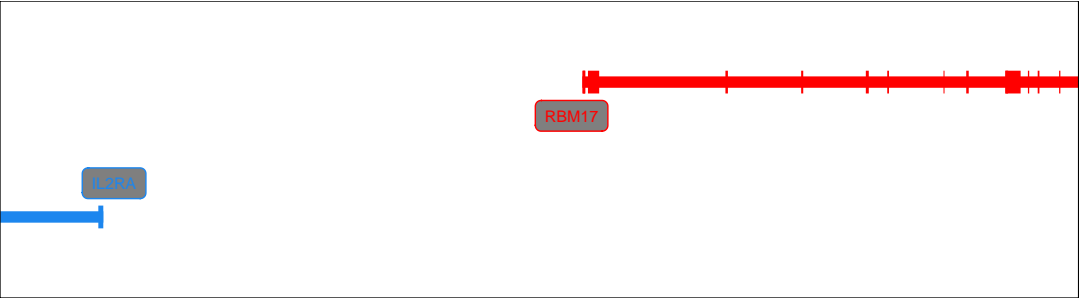
L5.PCP4



Peaks

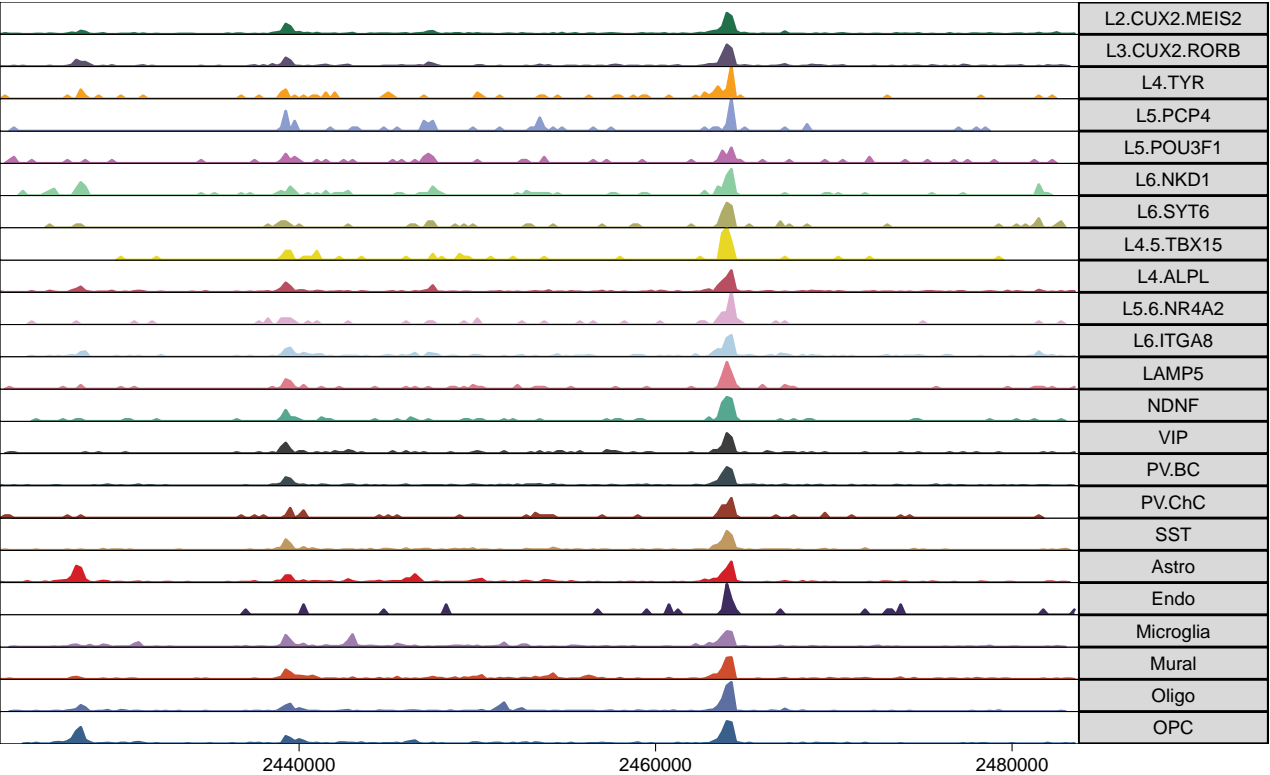


LoopTrack

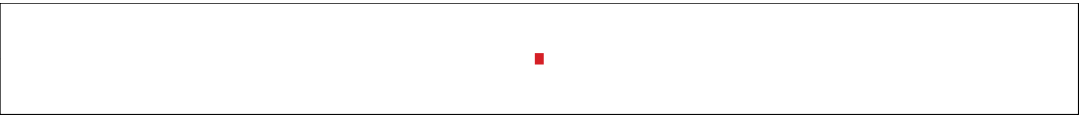


Genes

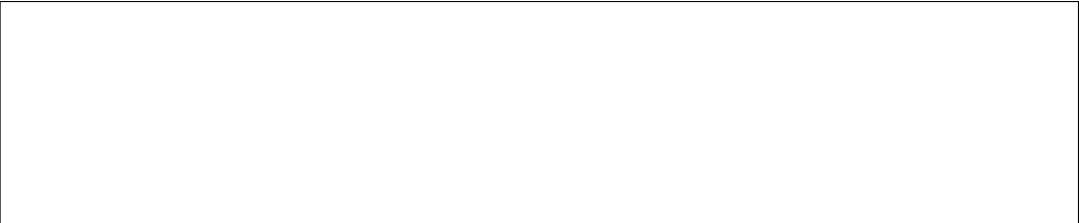
Coverage
(Norm. ATAC Signal Range 0-2.48) by ReadsInTSS)



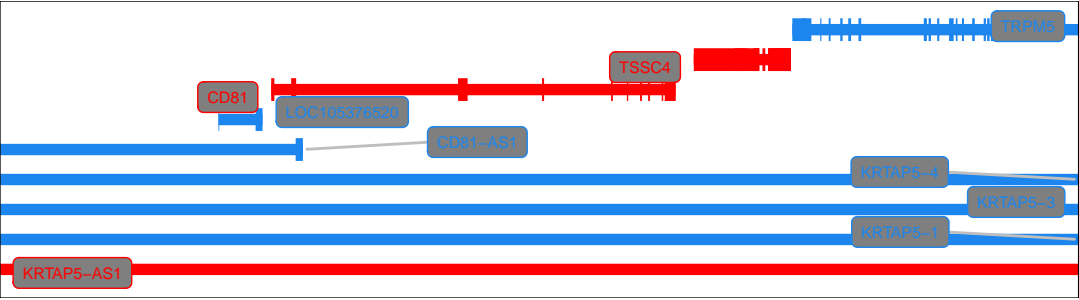
L5.PCP4



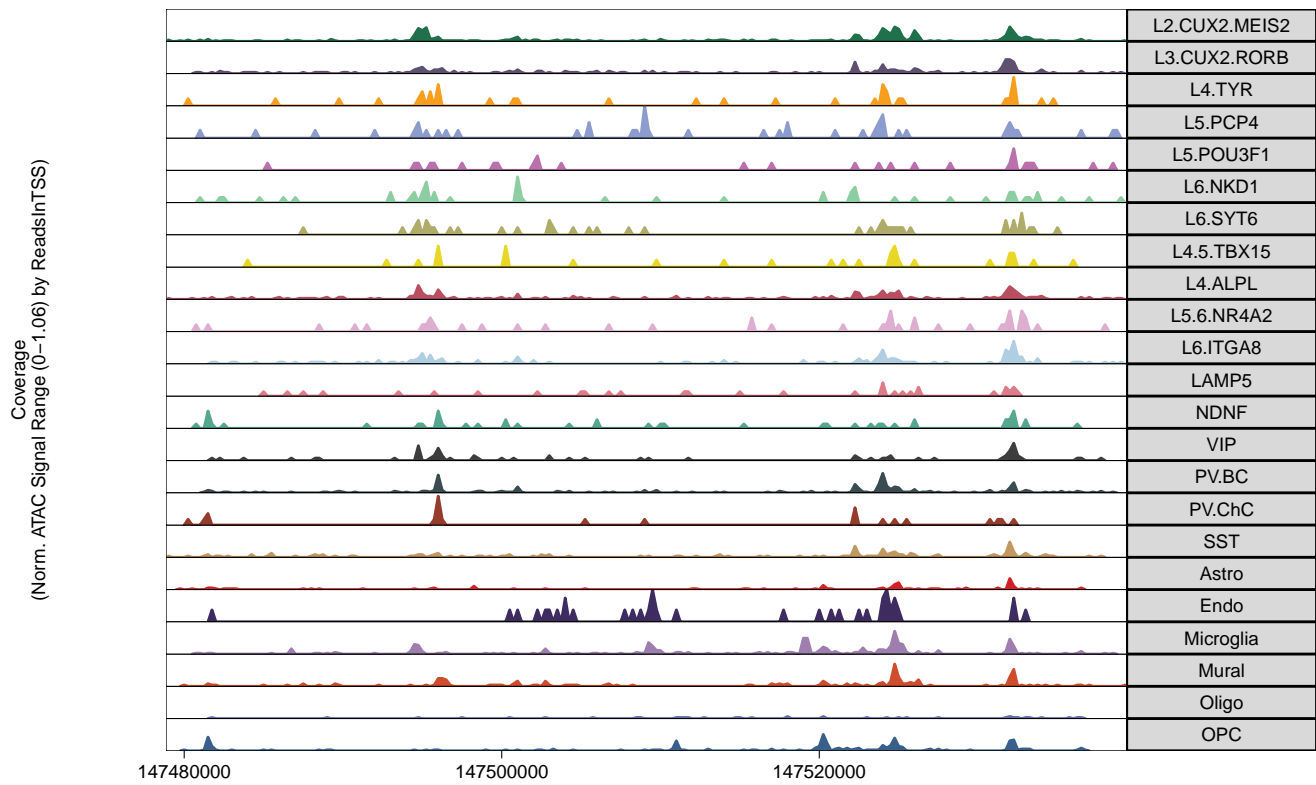
Peaks



LoopTrack



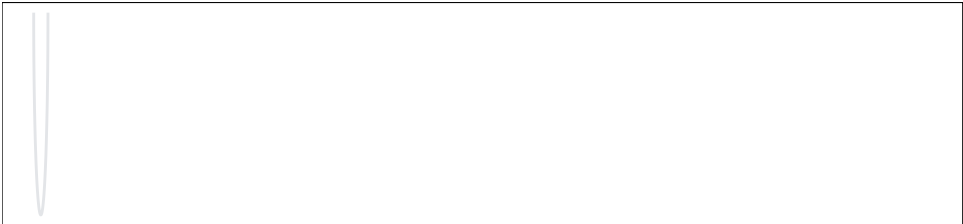
Genes



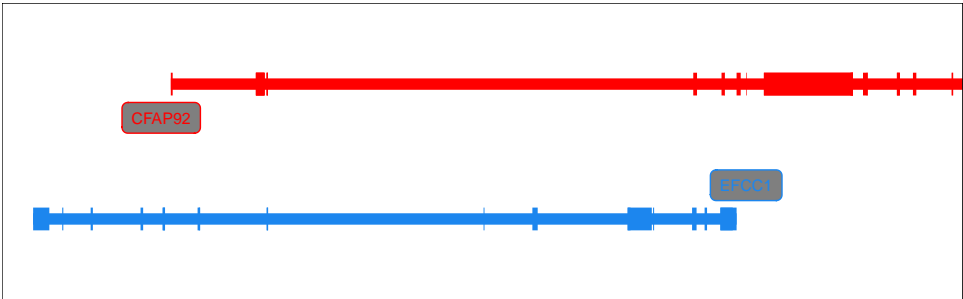
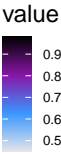
L5.PCP4



Peaks

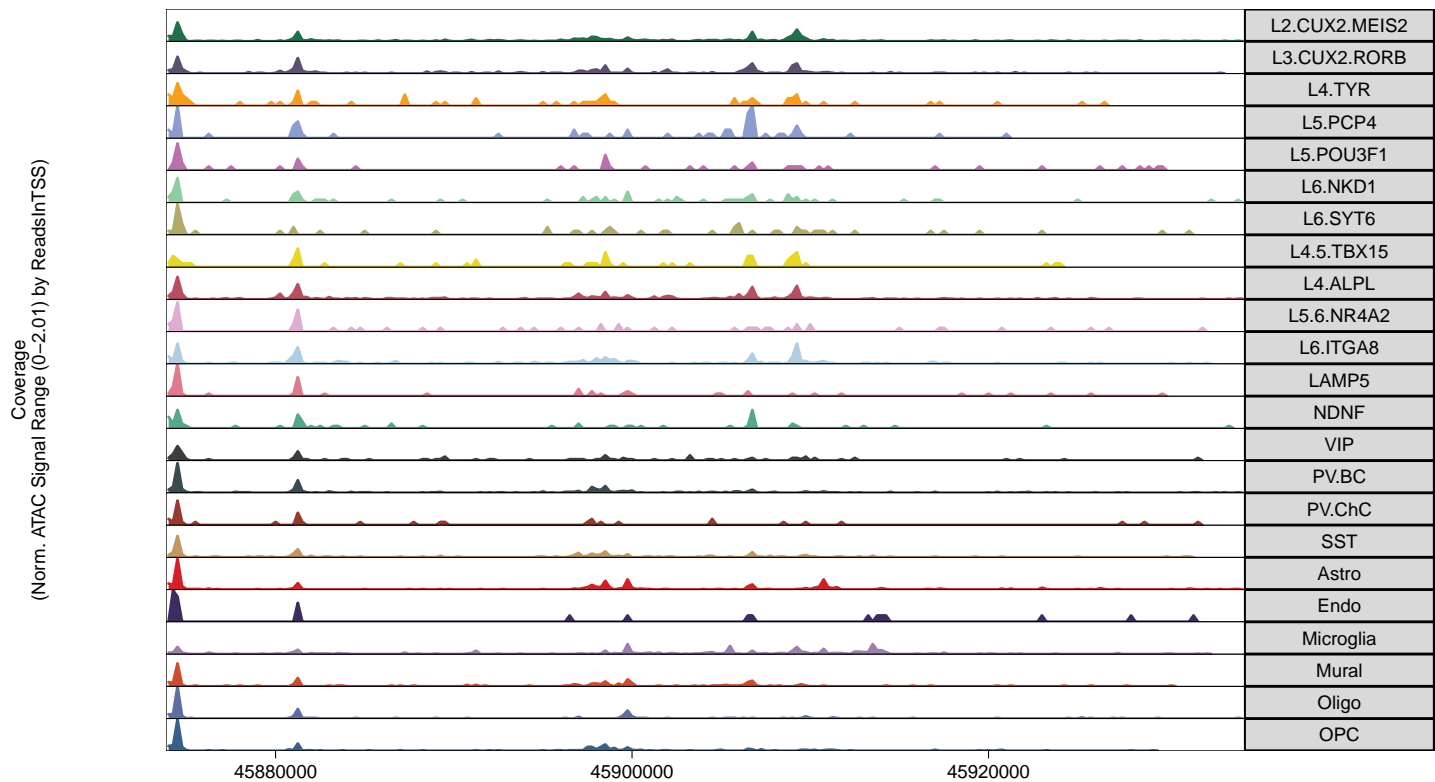


Peak2GeneLinks



Genes

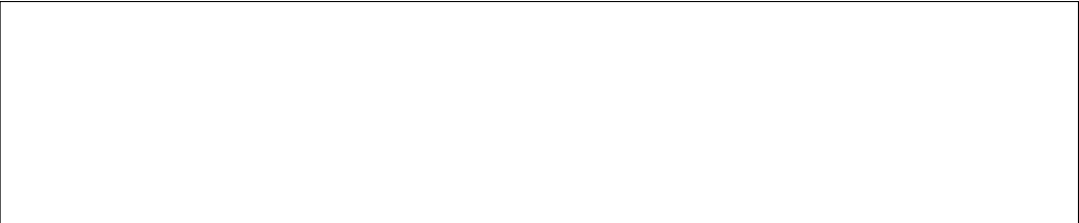
chr19:45873858–45934359 Marker Peak Tracks for: L5.PCP4



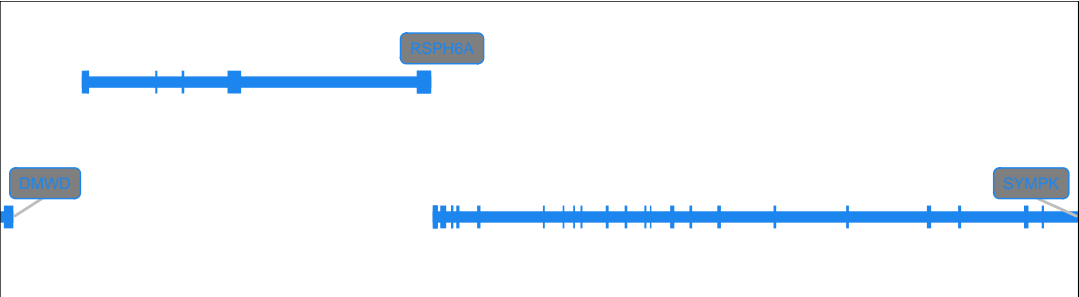
L5.PCP4



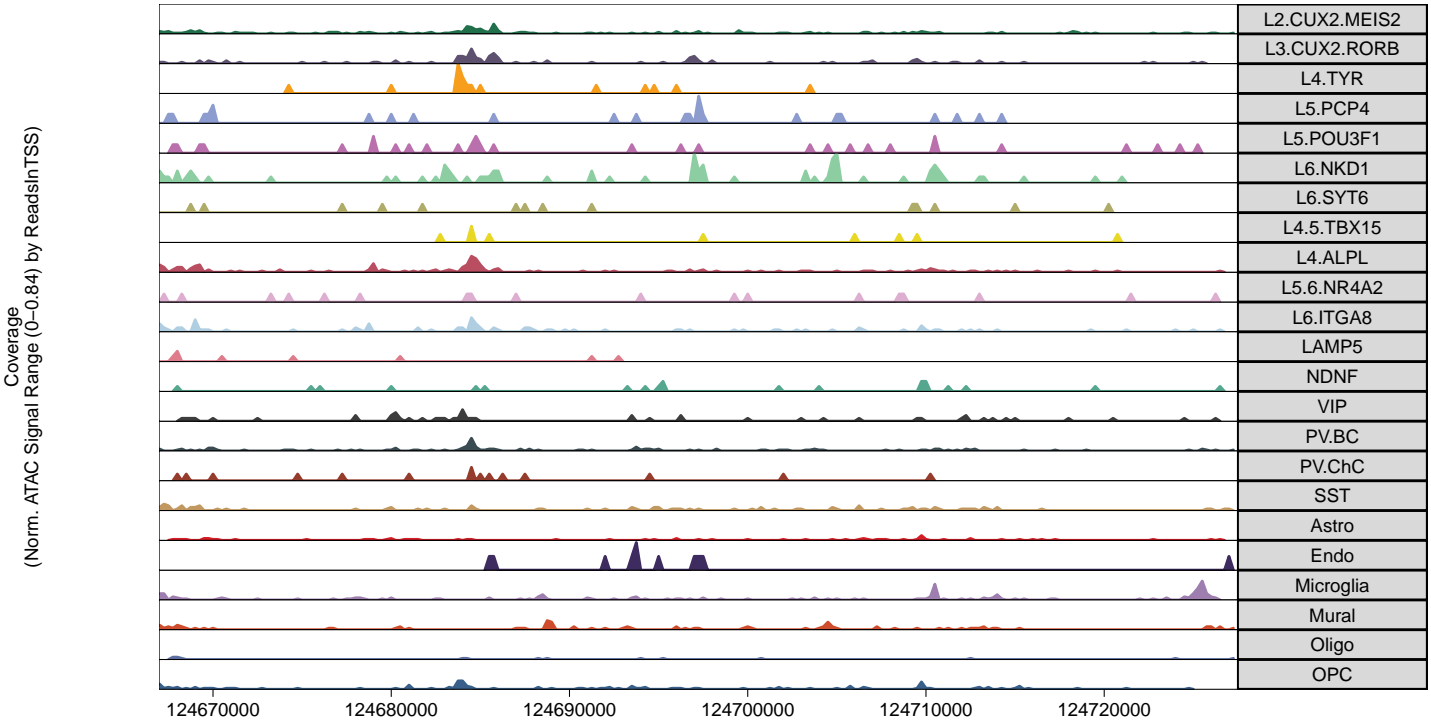
Peaks



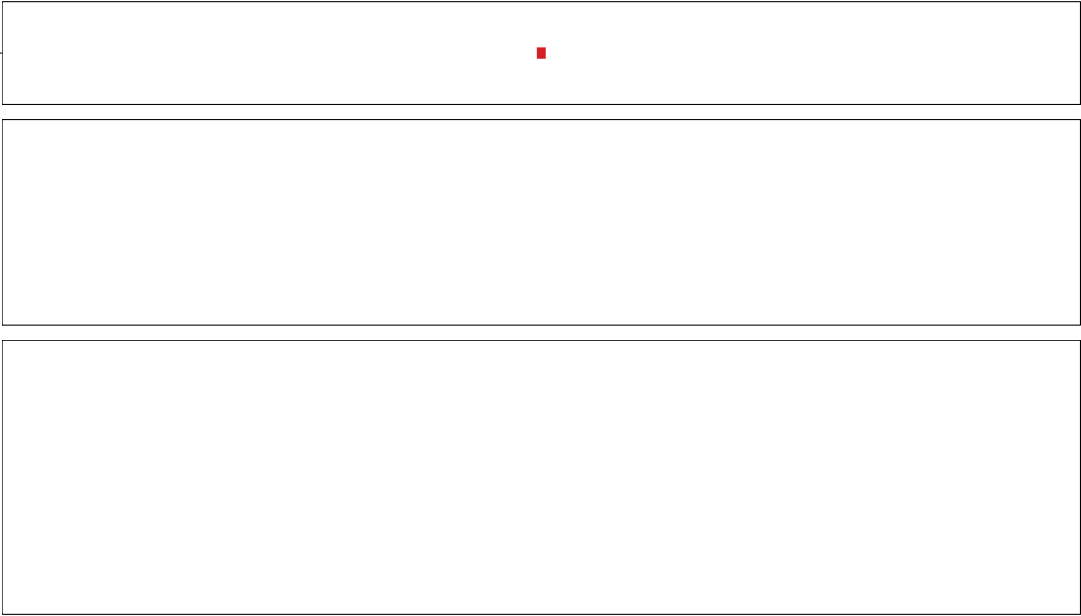
LoopTrack



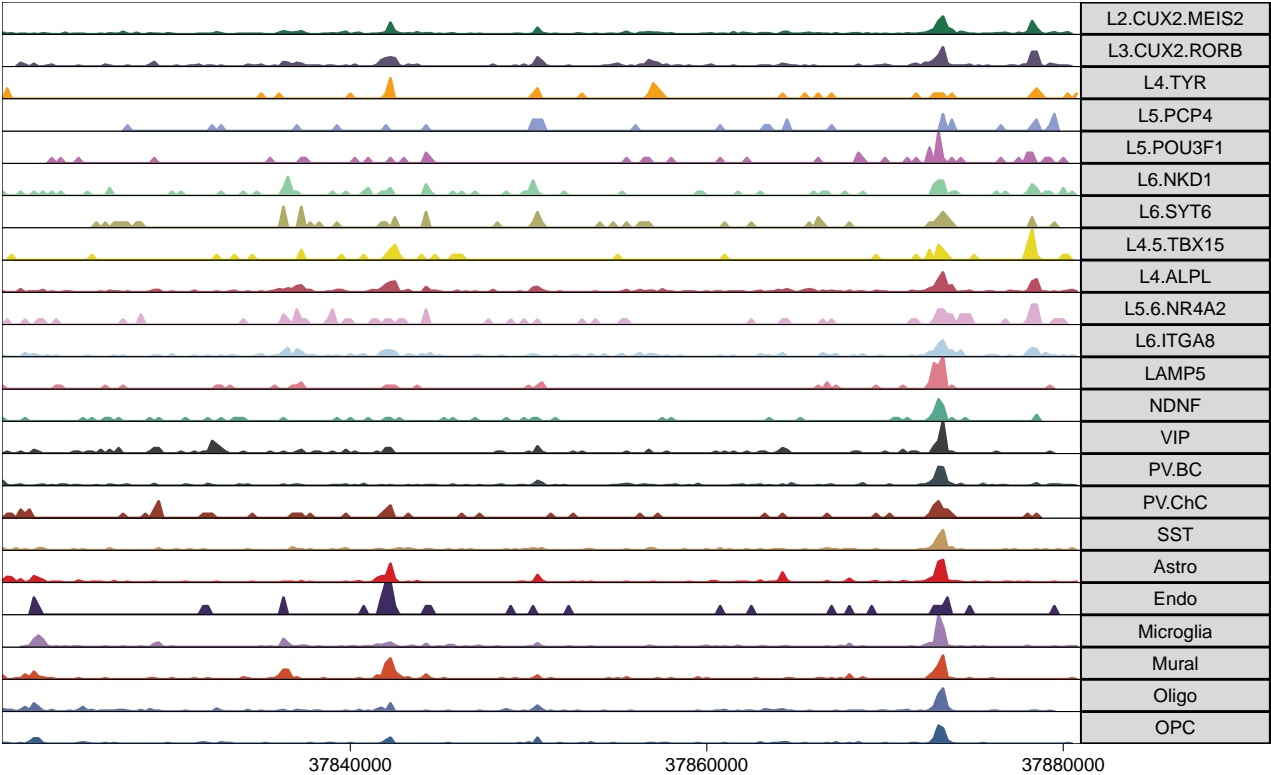
Genes



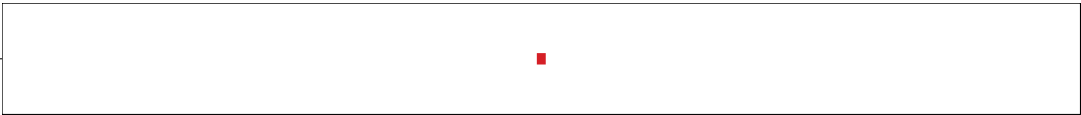
L5.PCP4



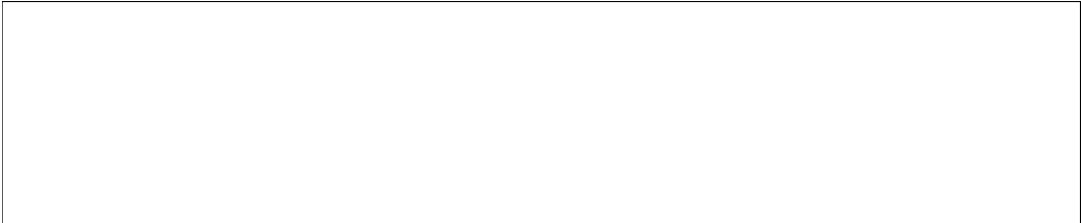
Coverage
(Norm. ATAC Signal Range 0–1.47) by ReadsInTSS)



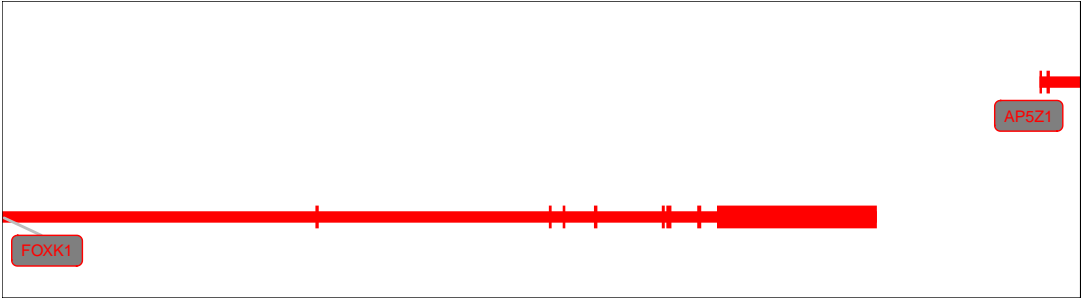
L5.PCP4



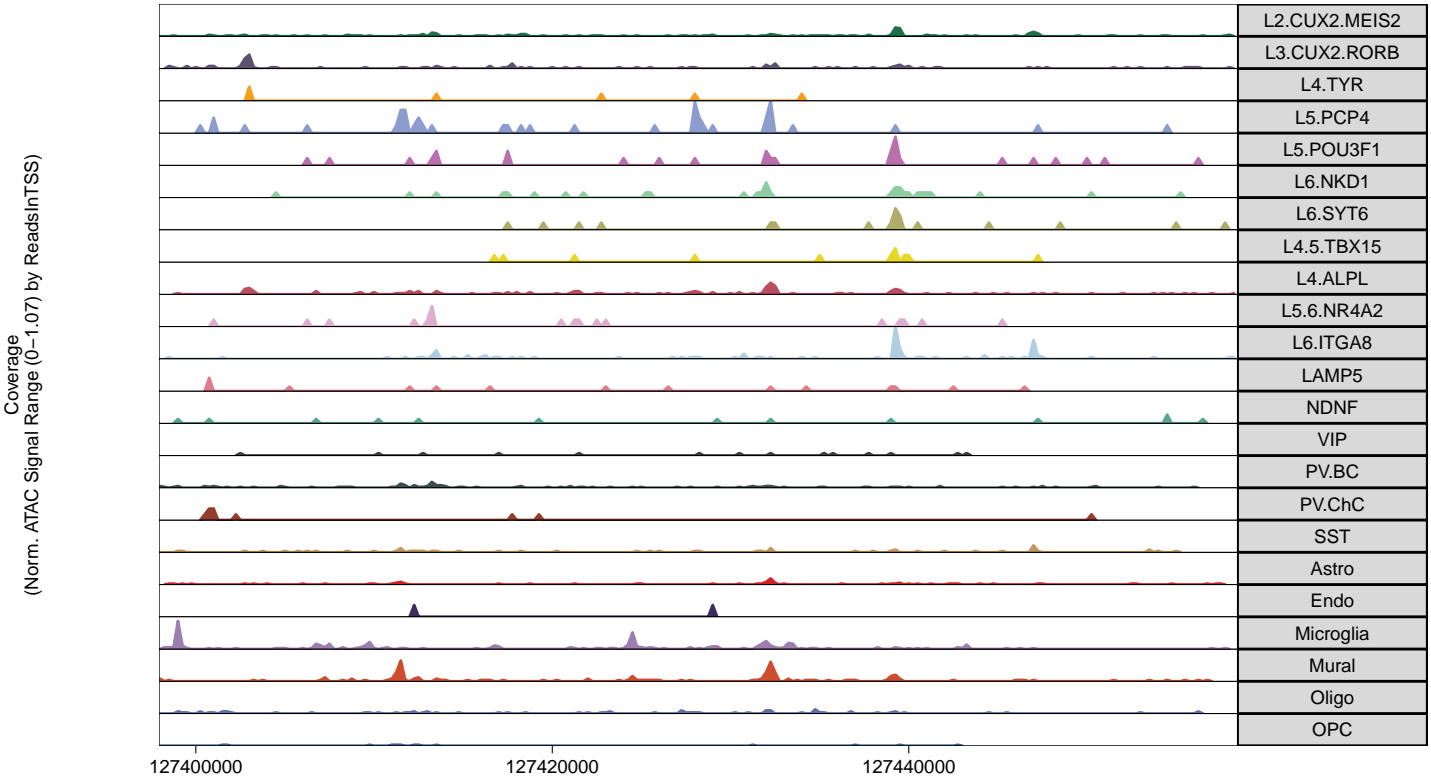
Peaks



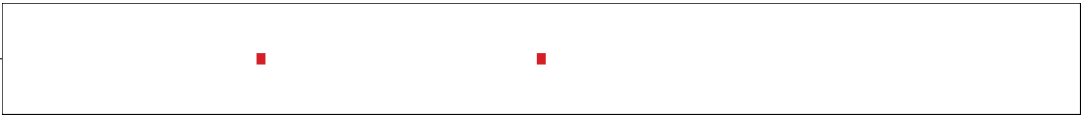
LoopTrack



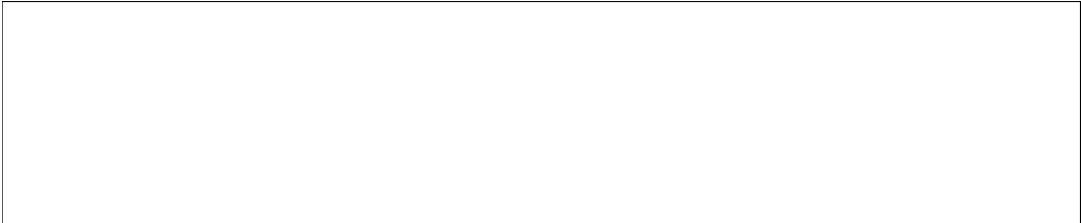
Genes



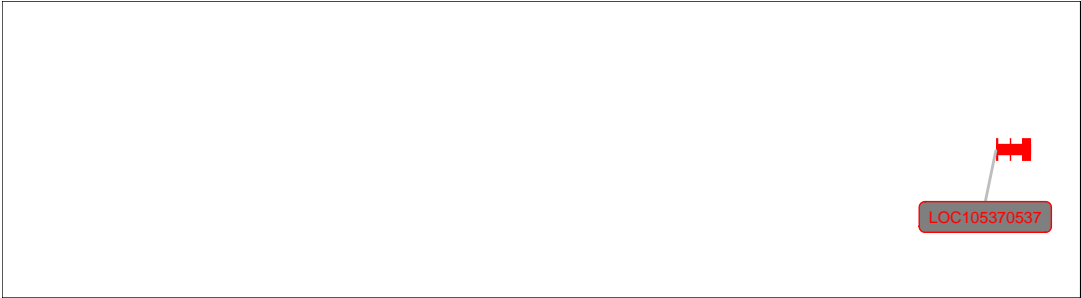
L5.PCP4



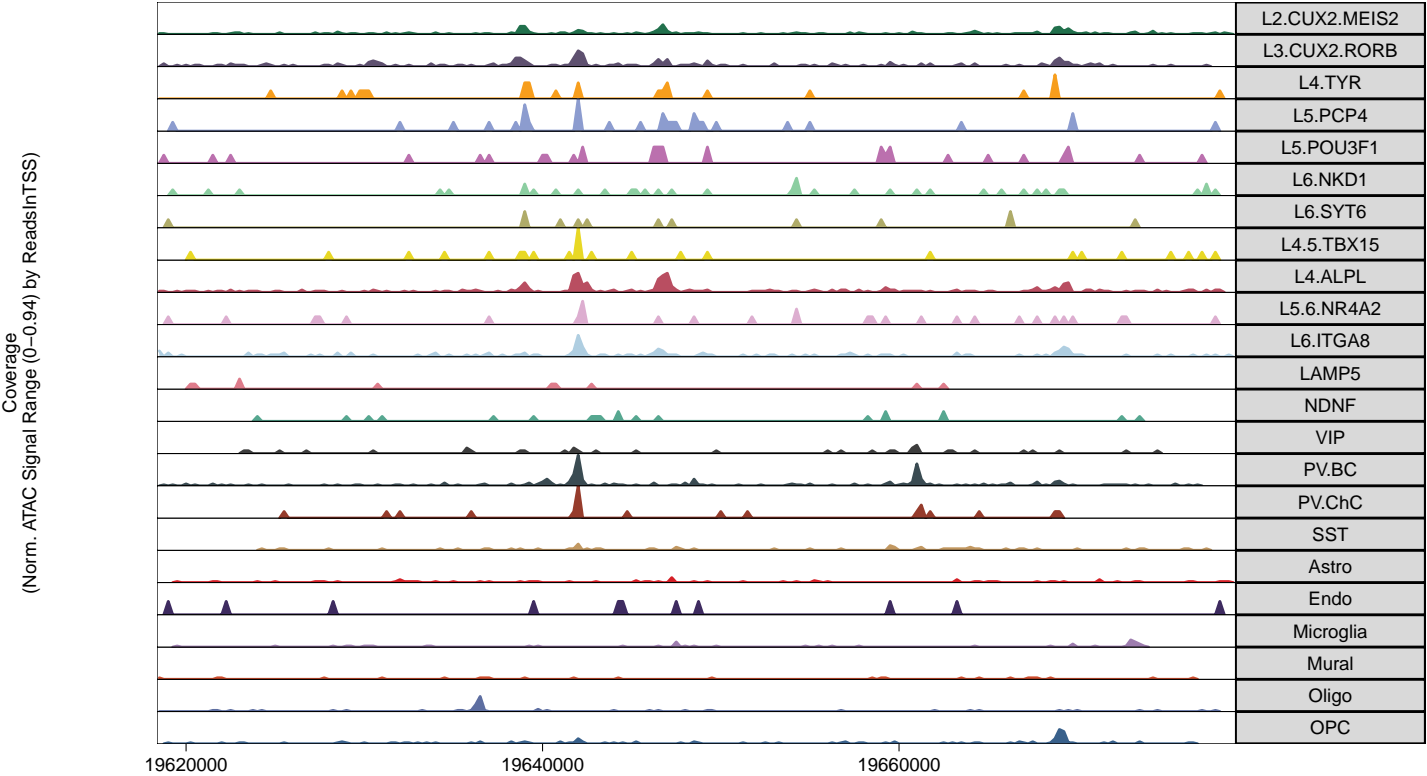
Peaks



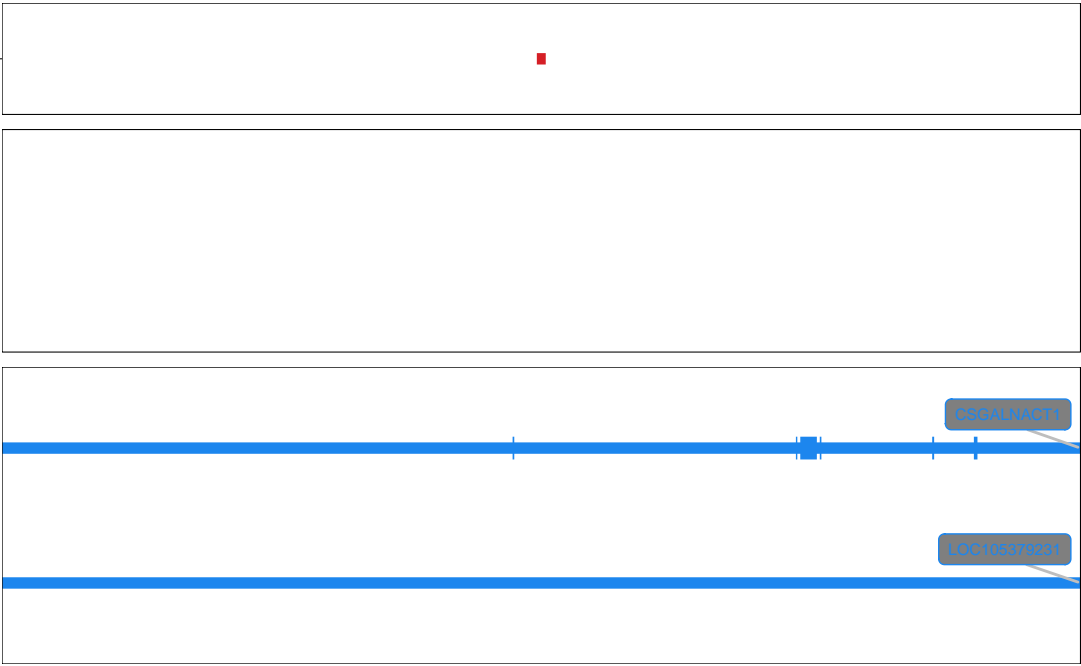
LoopTrack



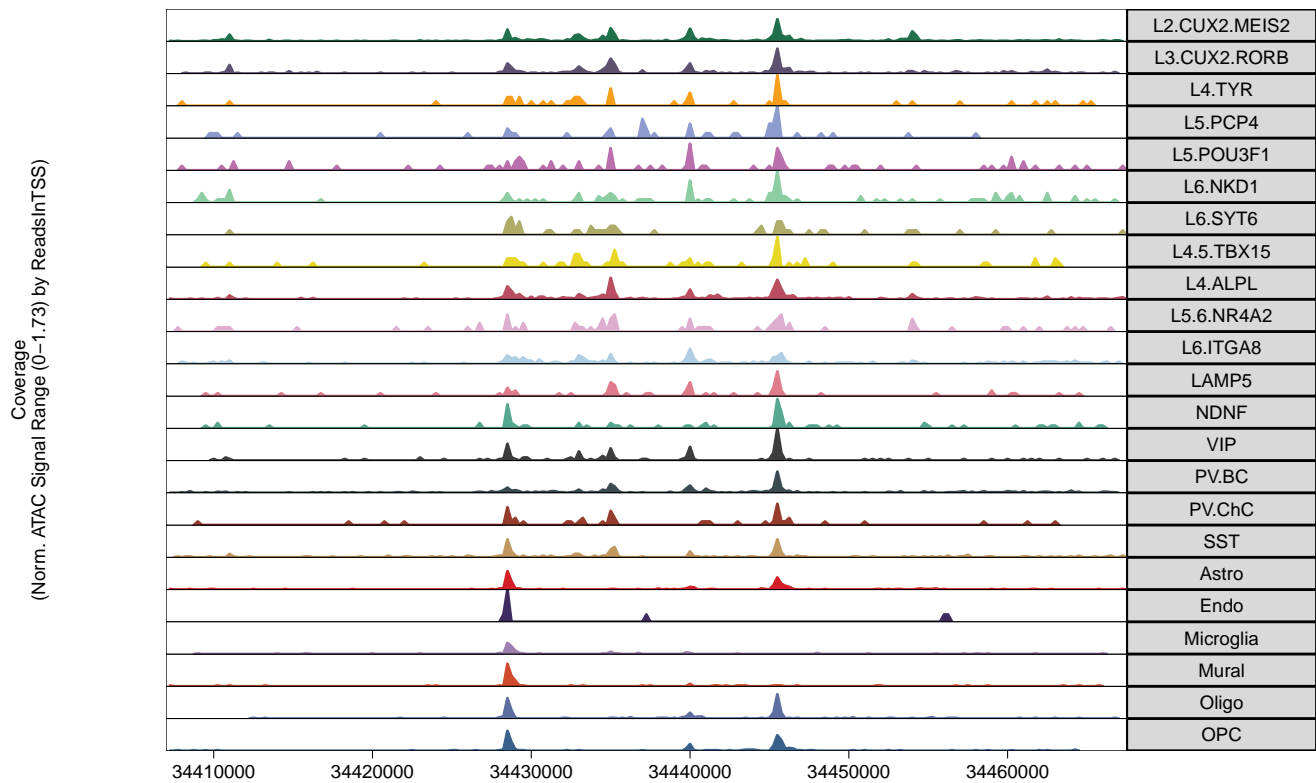
Genes



L5.PCP4



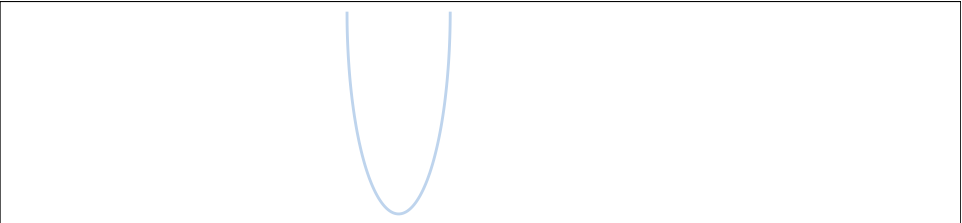
chr13:34407001–34467502 Marker Peak Tracks for: L5.PCP4



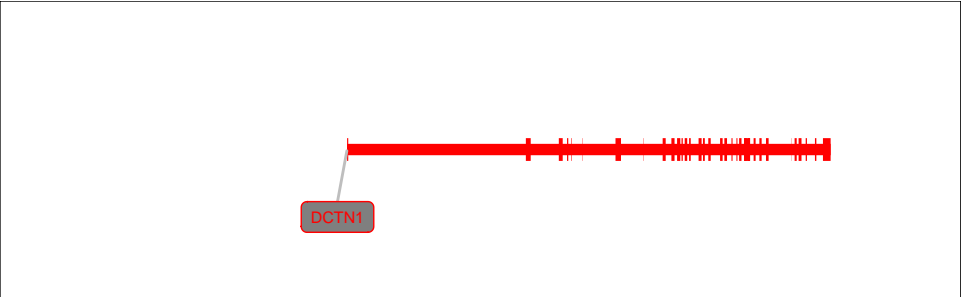
L5.PCP4



Peaks

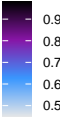


Peak2GeneLinks



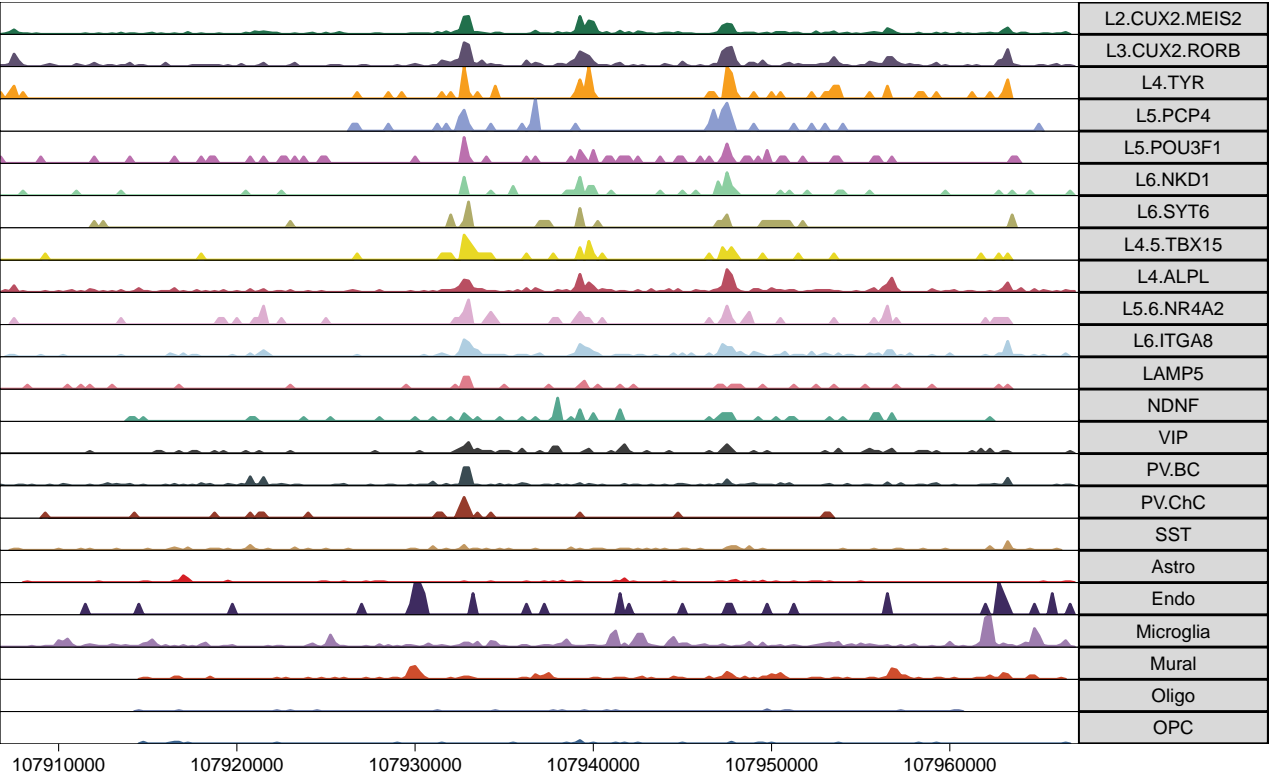
Genes

value

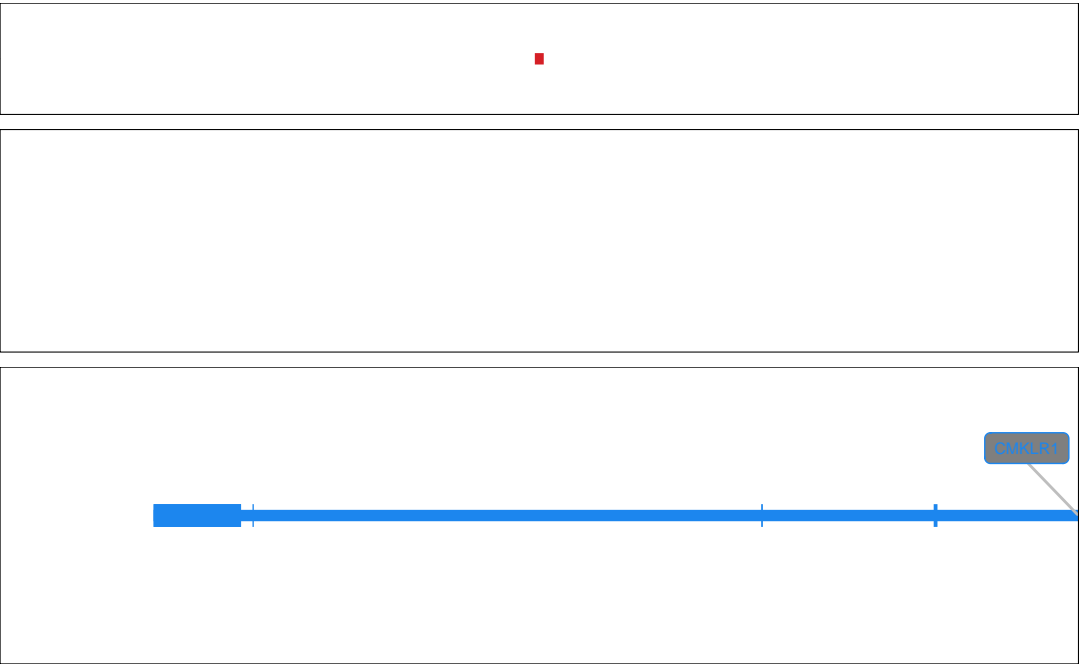


chr11:107906717-107967218 Marker Peak Tracks for: L5.PCP4

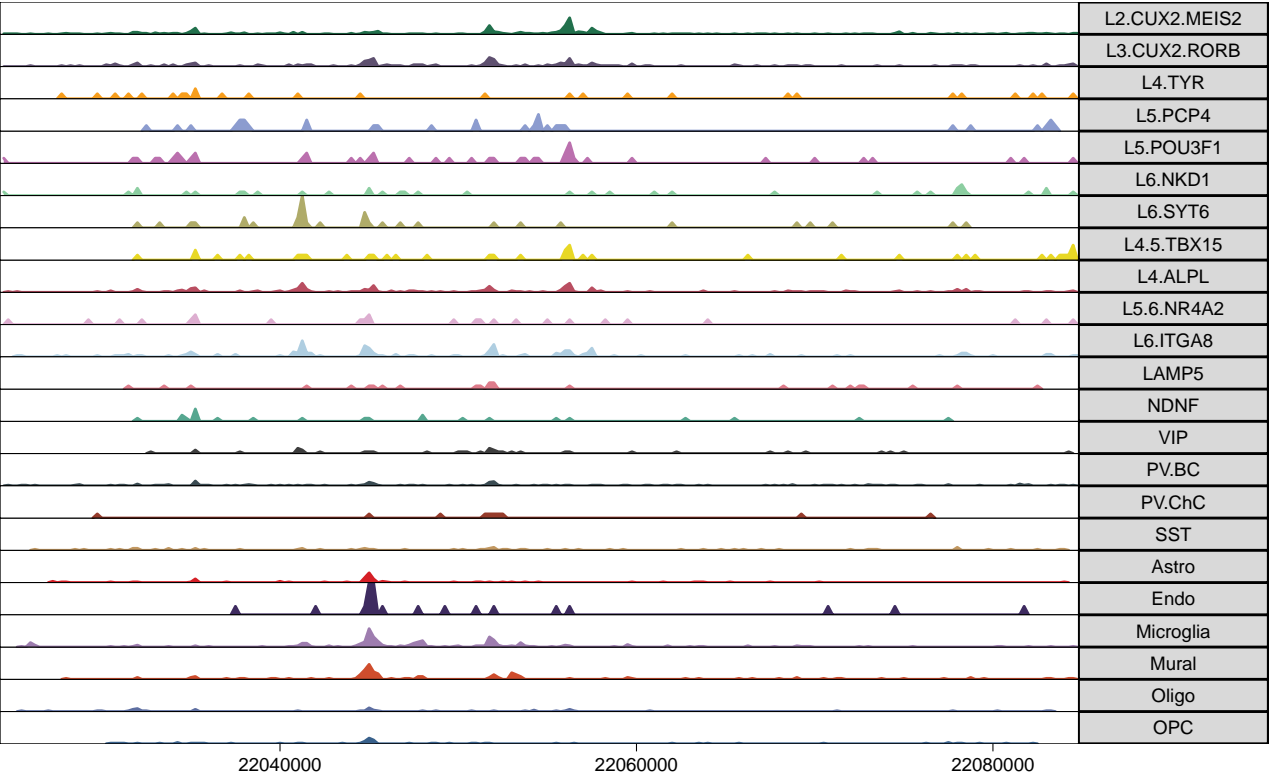
Coverage
(Norm. ATAC Signal Range 0-1.19) by ReadsInTSS)



L5.PCP4



Coverage
(Norm. ATAC Signal Range 0–1.49) by ReadsInTSS)



L5.PCP4



Peaks

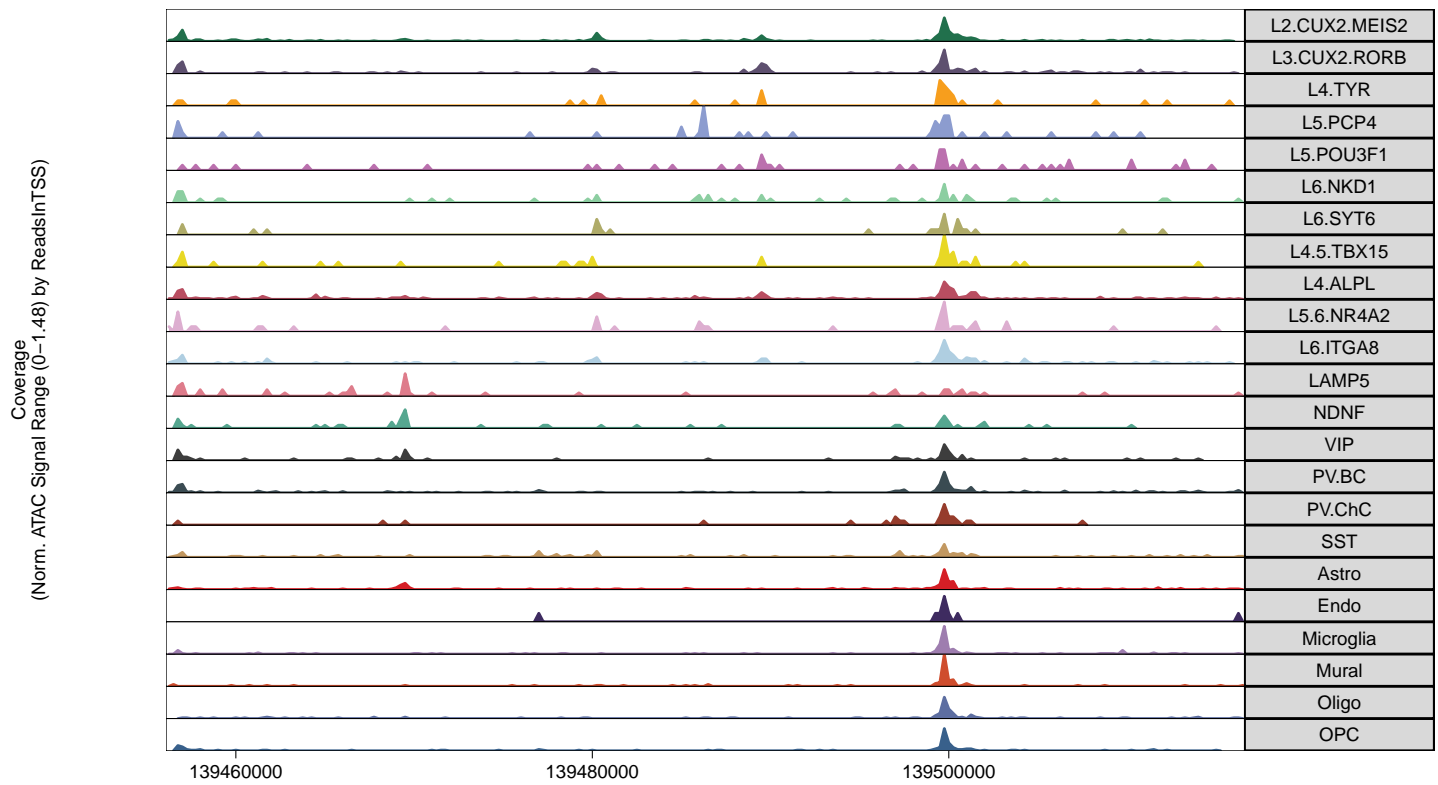


LoopTrack



Genes

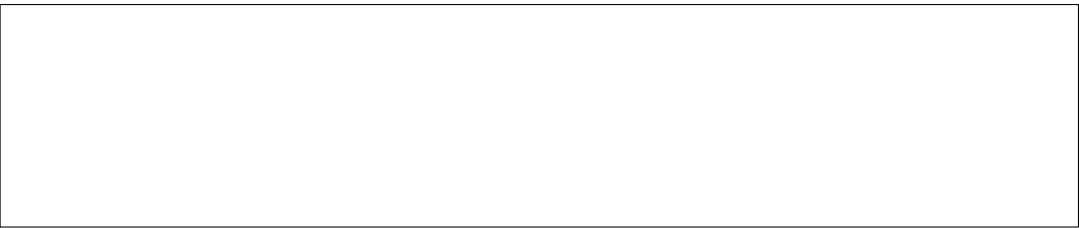
chr6:139456087-139516588 Marker Peak Tracks for: L5.PCP4



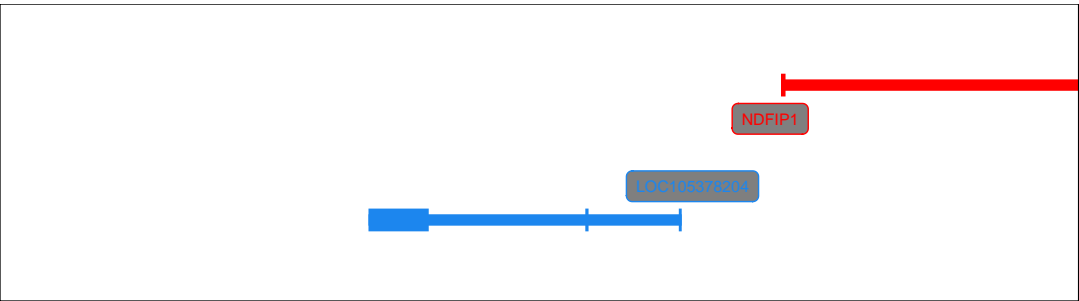
L5.PCP4



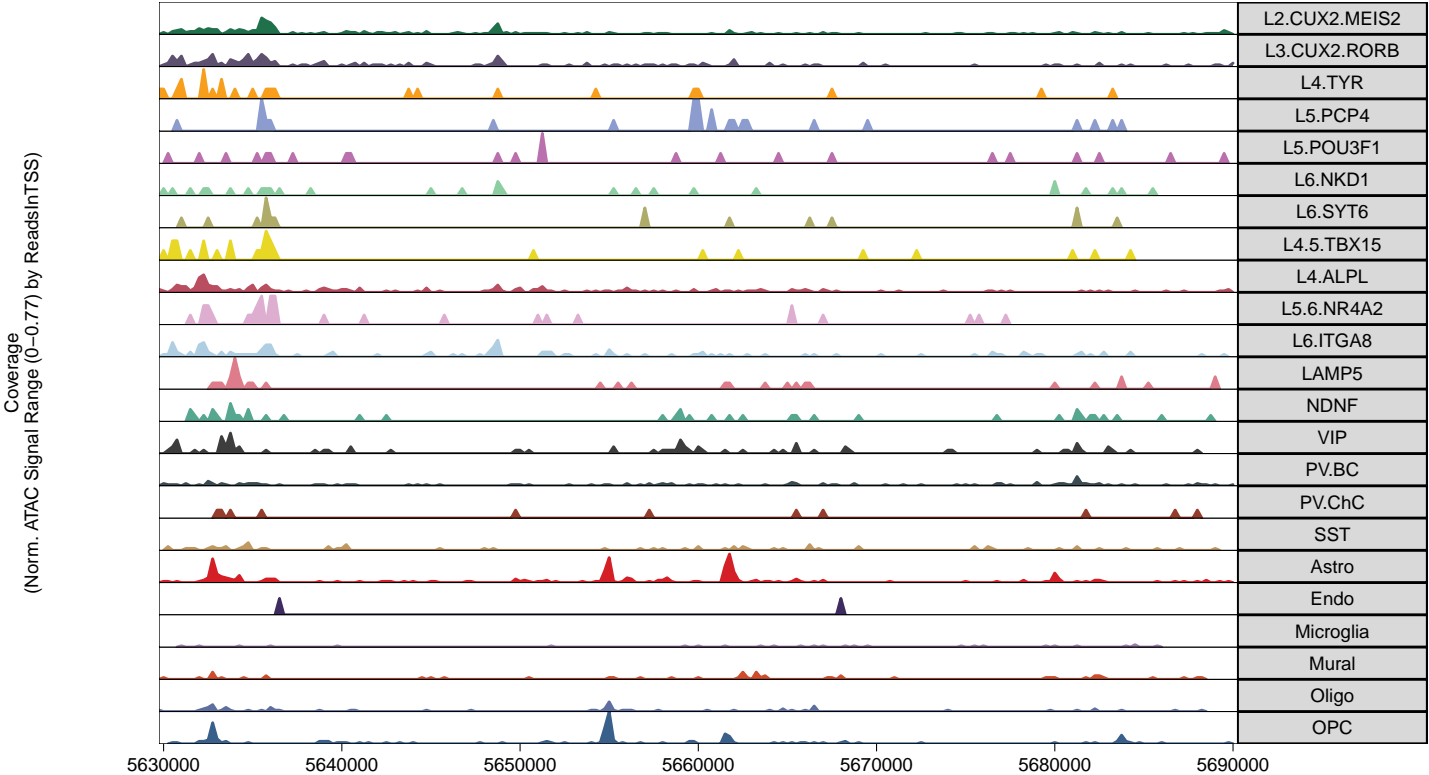
Peaks



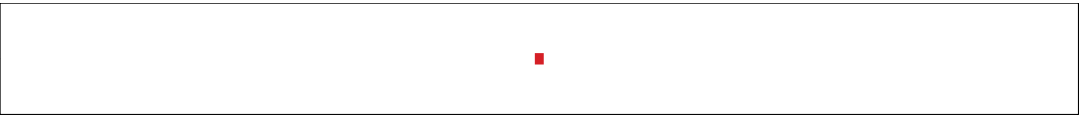
LoopTrack



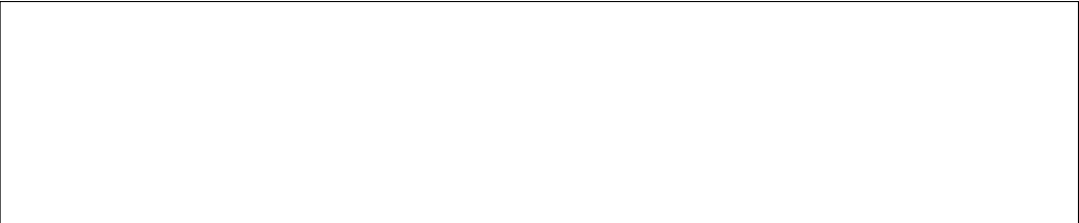
Genes



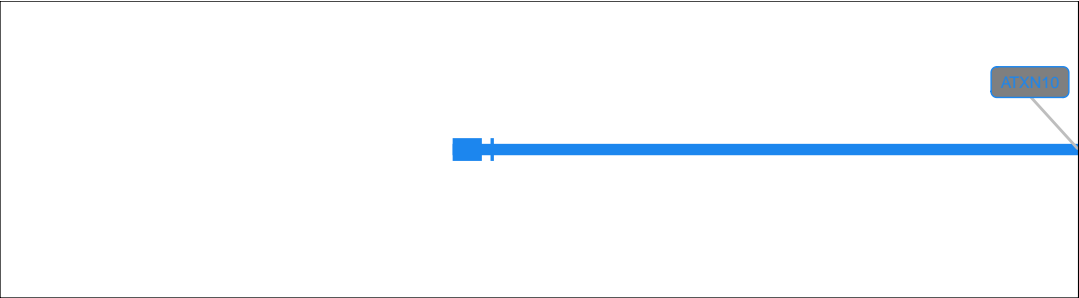
L5.PCP4



Peaks

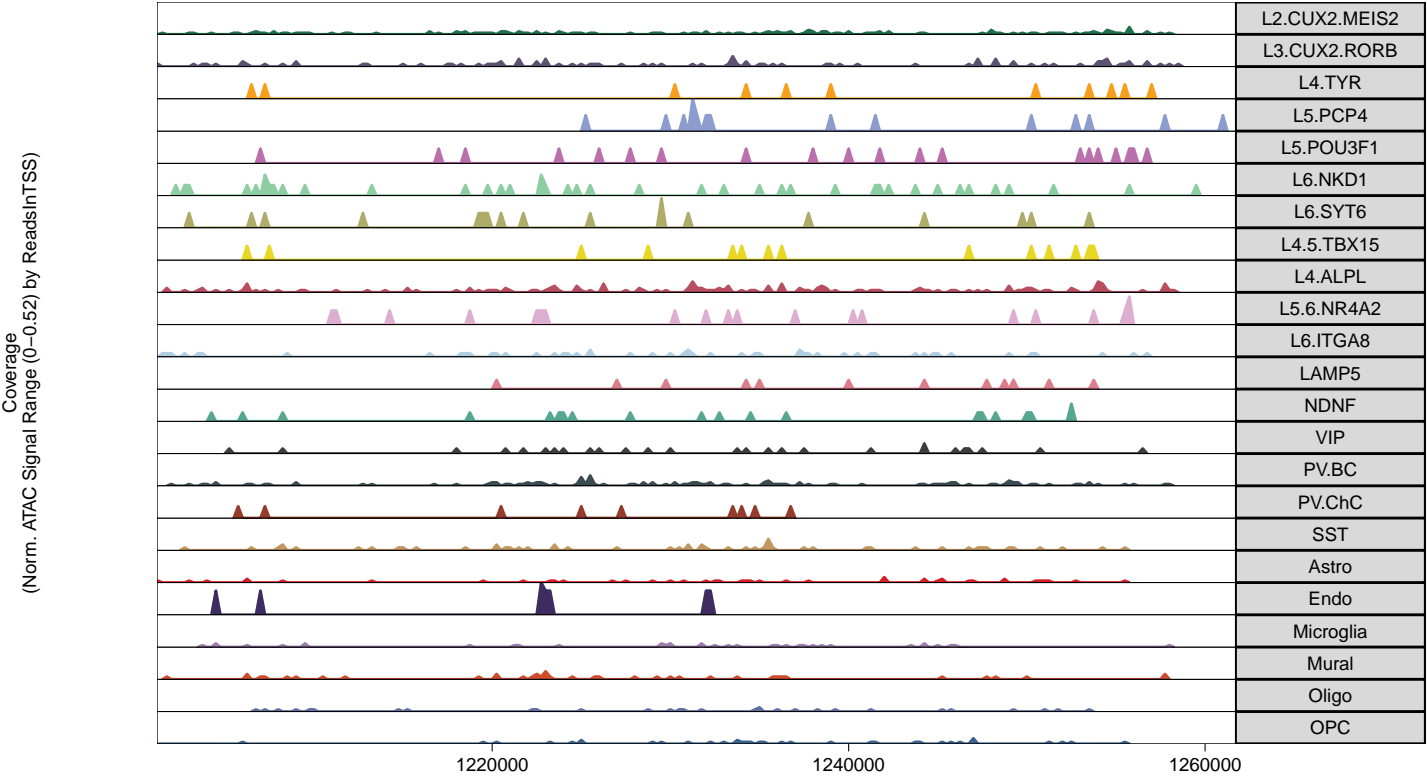


LoopTrack

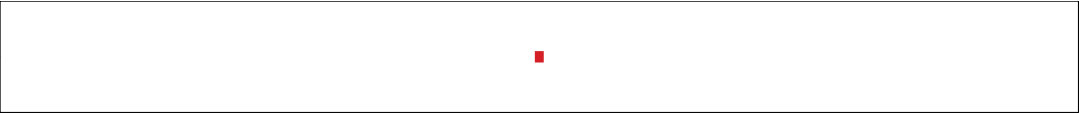


Genes

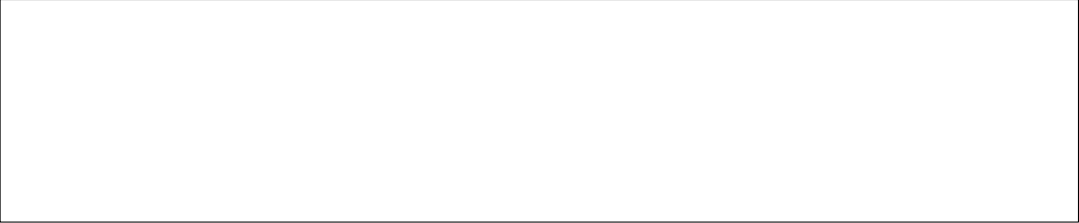
chr14:1201202-1261703 Marker Peak Tracks for: L5.PCP4



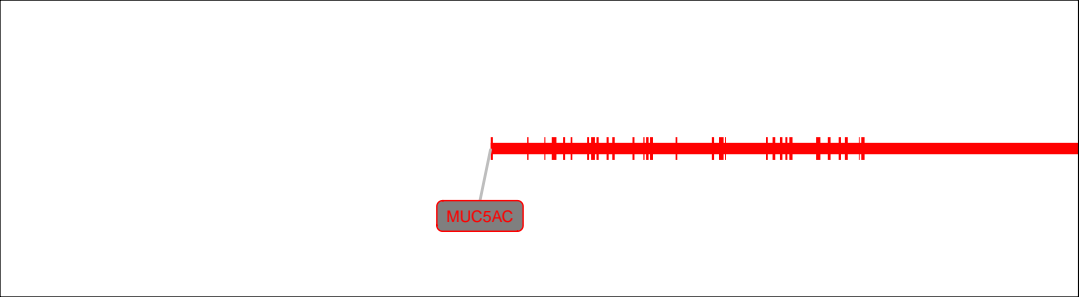
L5.PCP4



Peaks

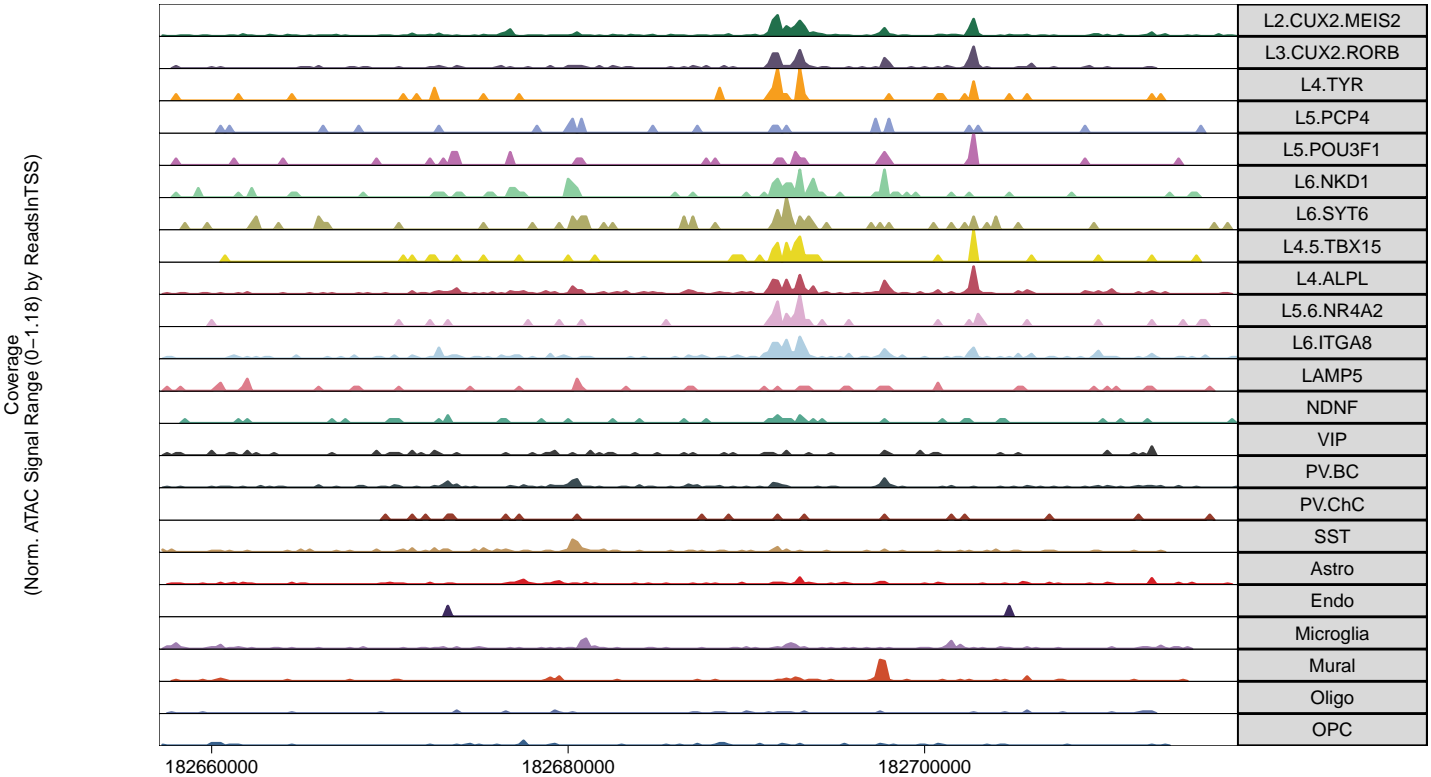


LoopTrack

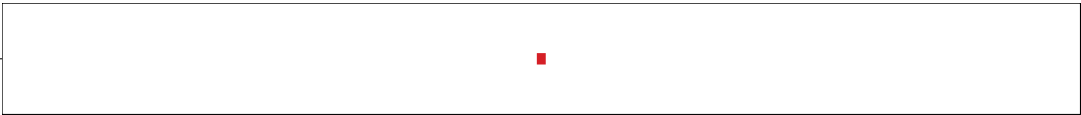


Genes

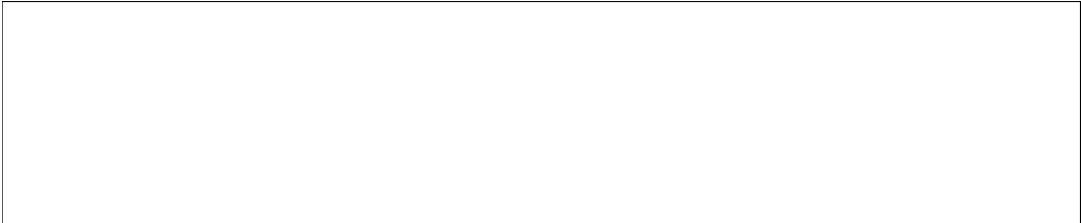
chr1:182657051-182717552 Marker Peak Tracks for: L5.PCP4



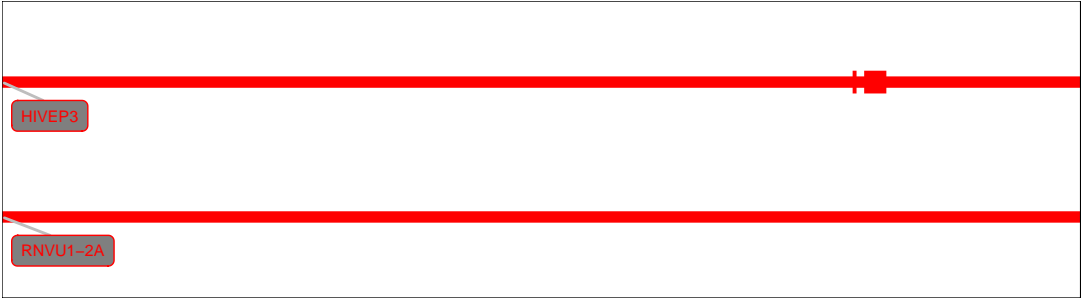
L5.PCP4



Peaks

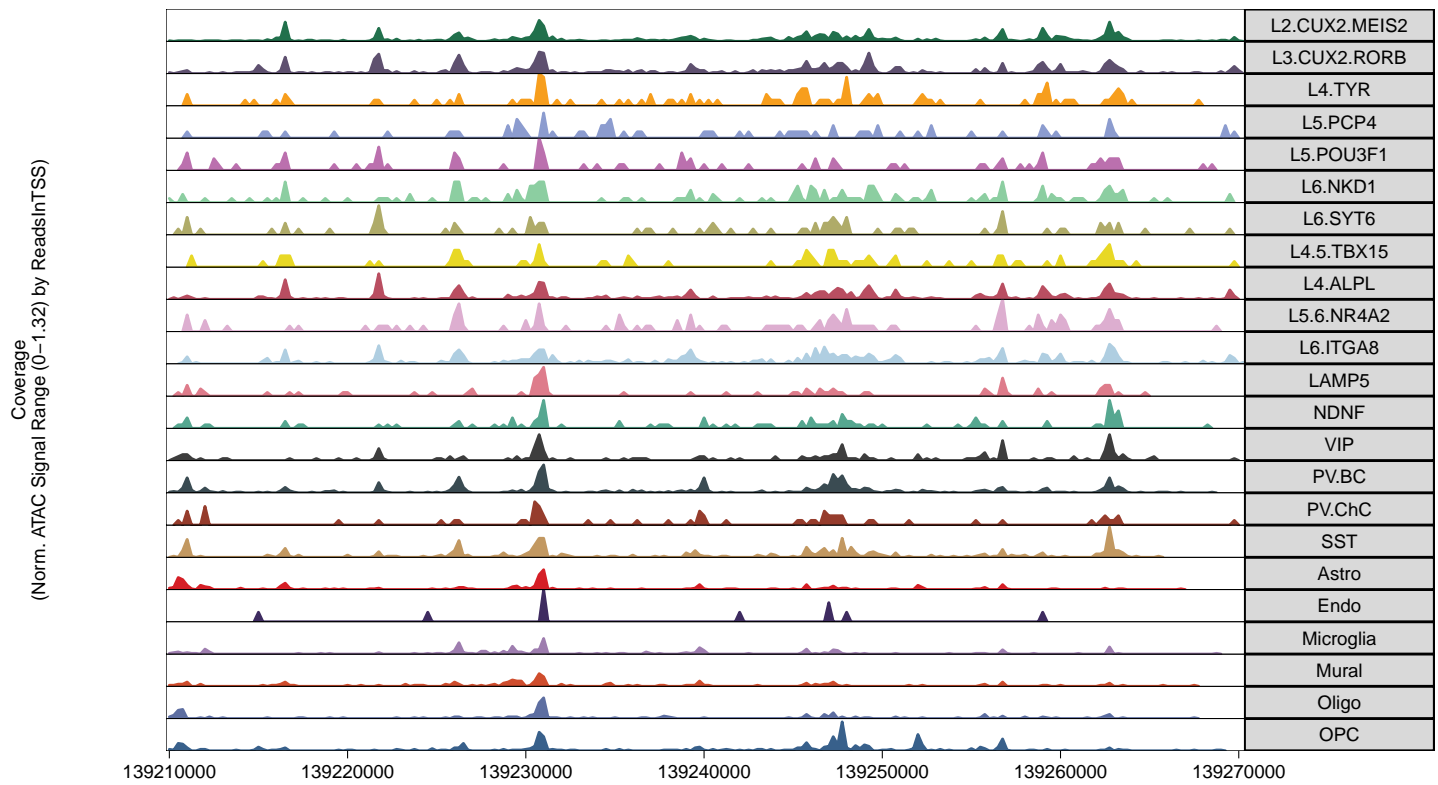


LoopTrack



Genes

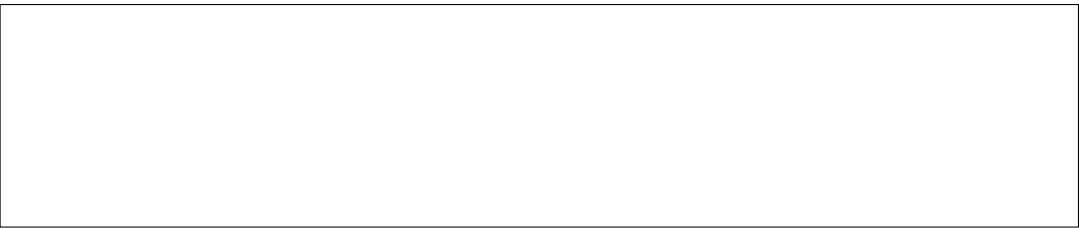
chr7:139209818–139270319 Marker Peak Tracks for: L5.PCP4



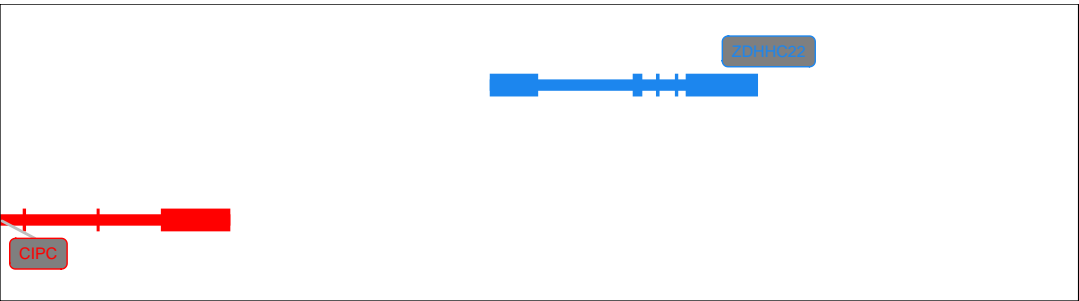
L5.PCP4



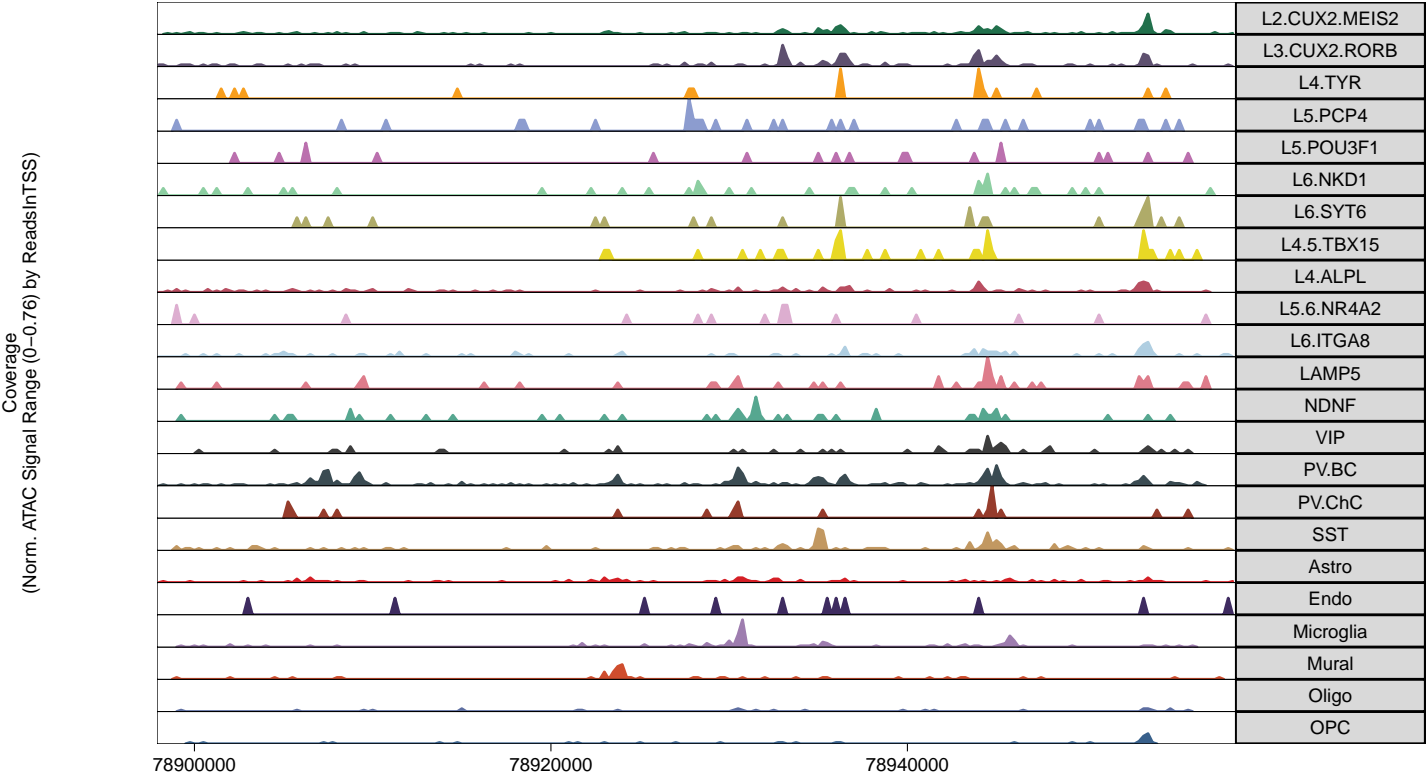
Peaks



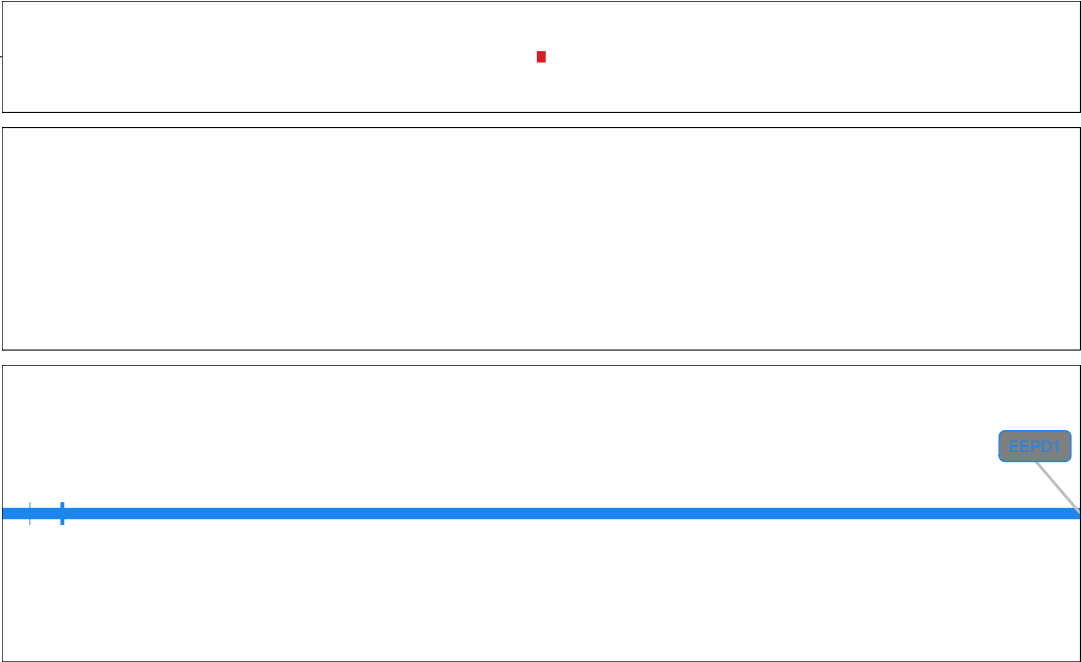
LoopTrack

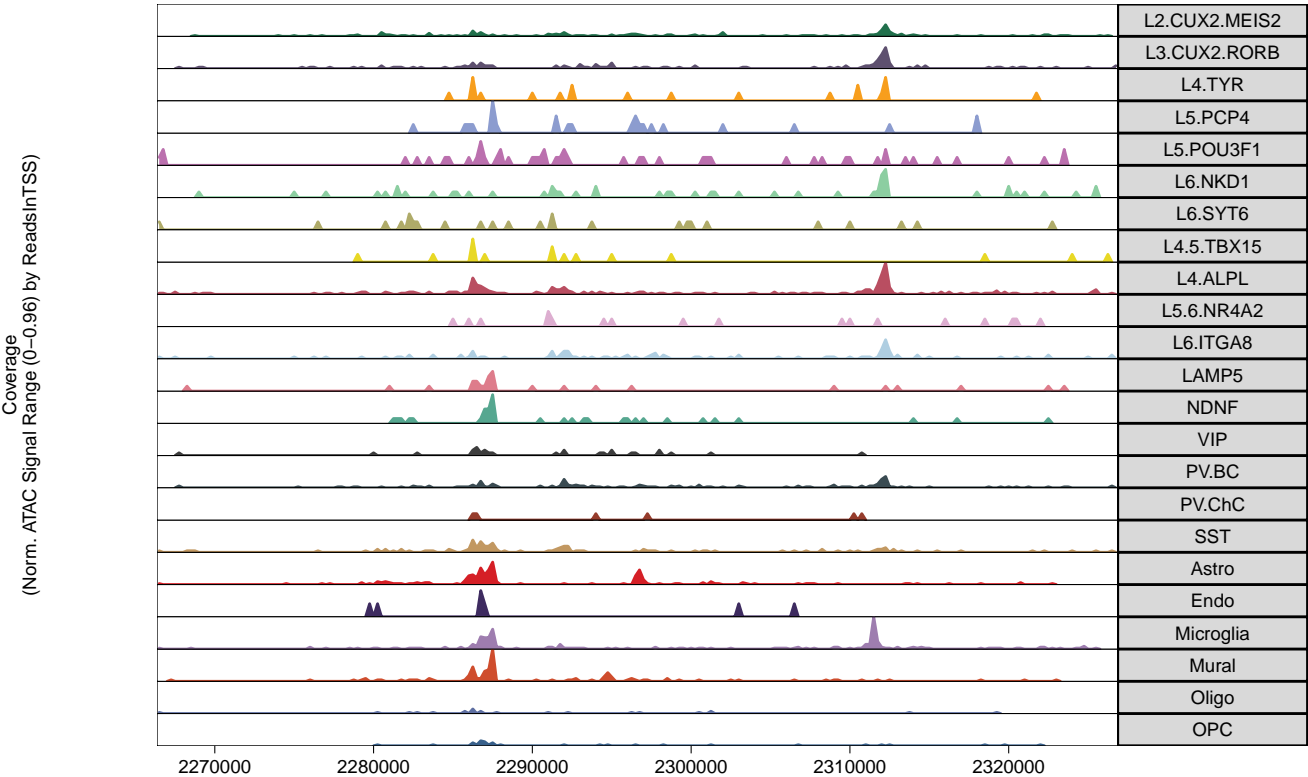


Genes



L5.PCP4

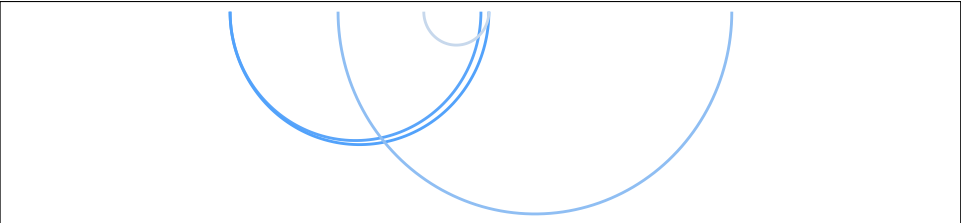




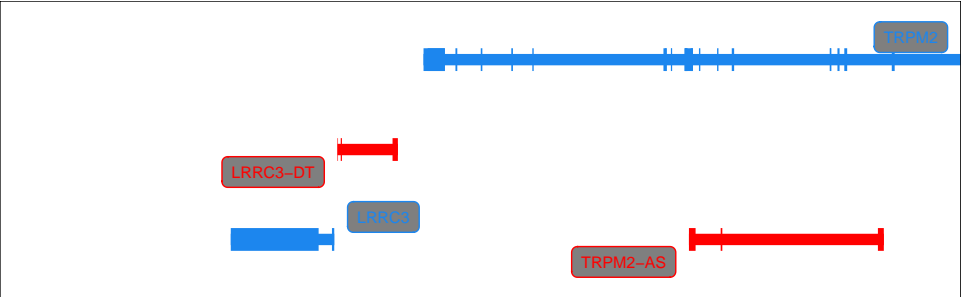
L5.PCP4



Peaks

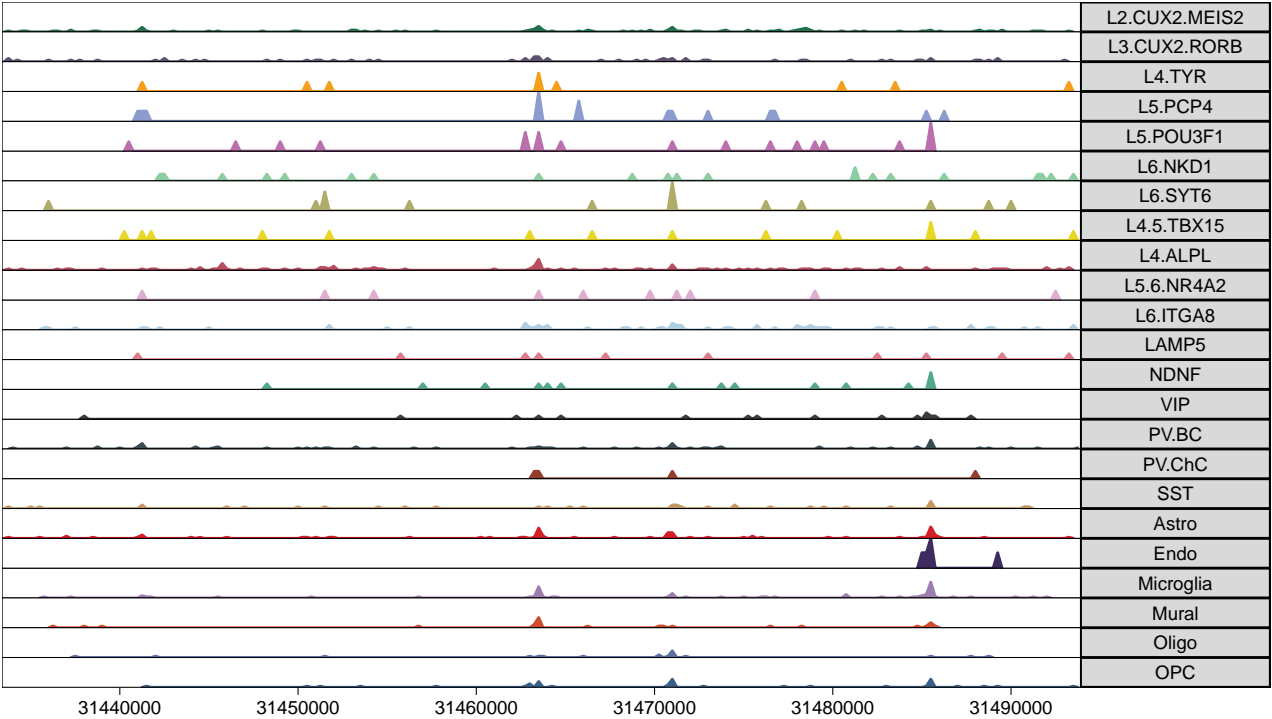


Peak2GeneLinks

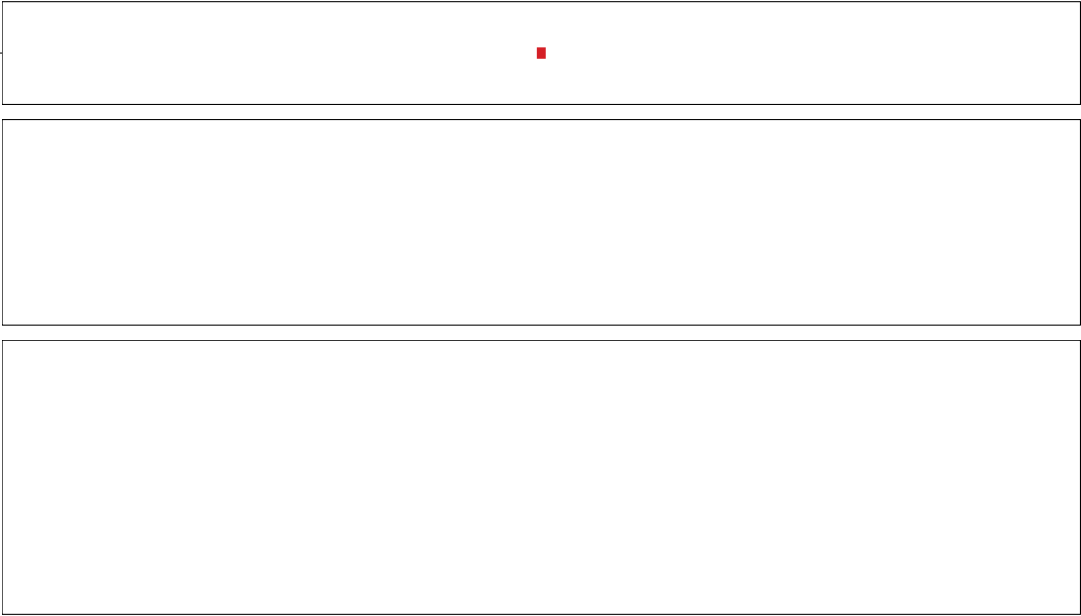


Genes

Coverage
(Norm. ATAC Signal Range (0–0.75) by Reads in TSS)



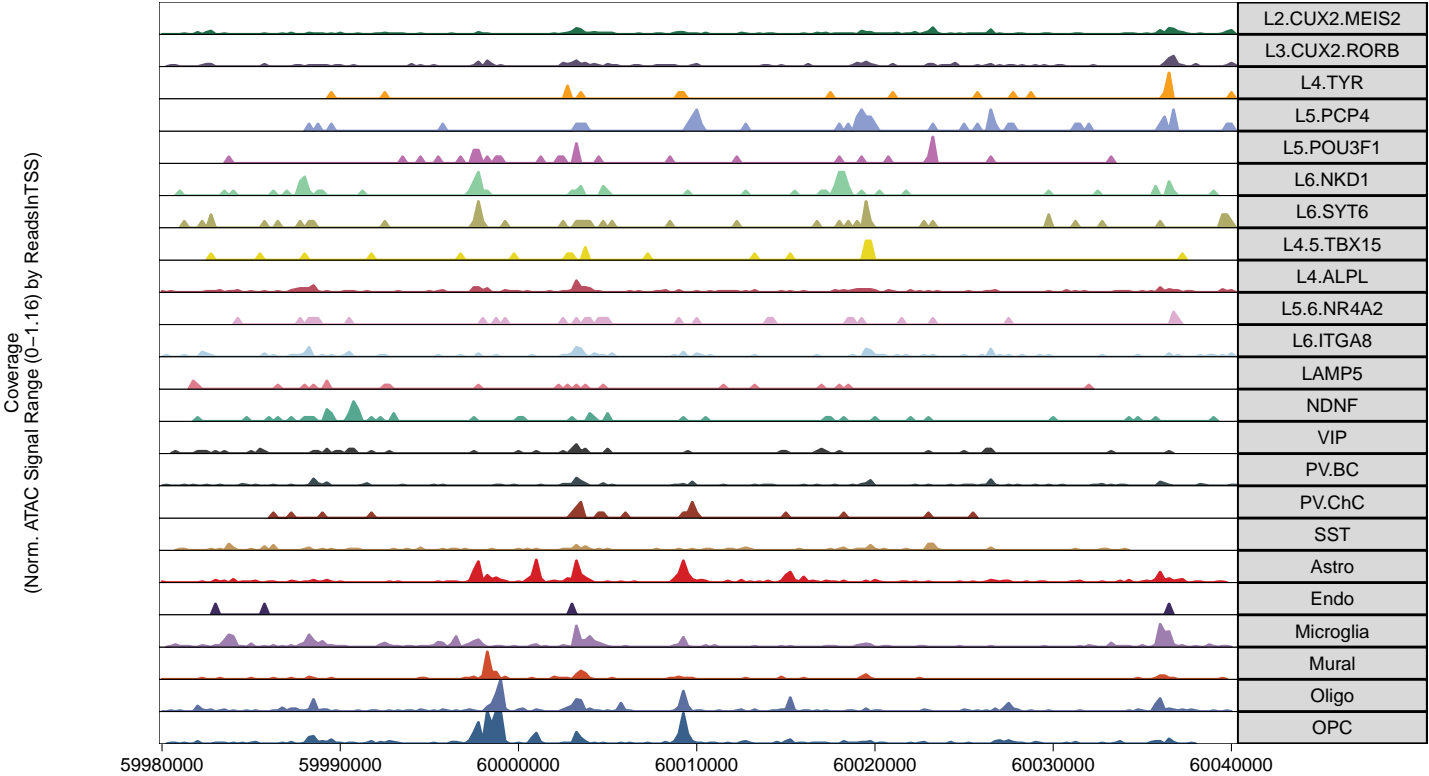
L5.PCP4



Peaks

LoopTrack

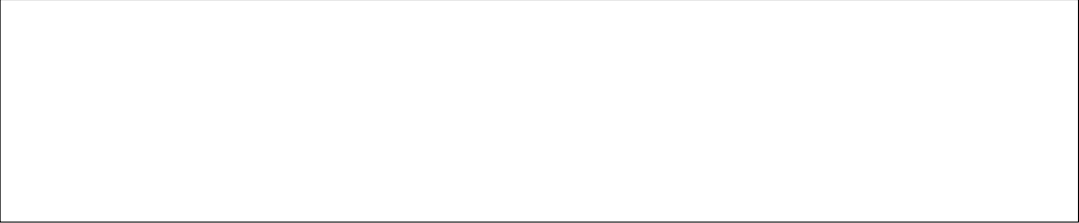
GeneTrack



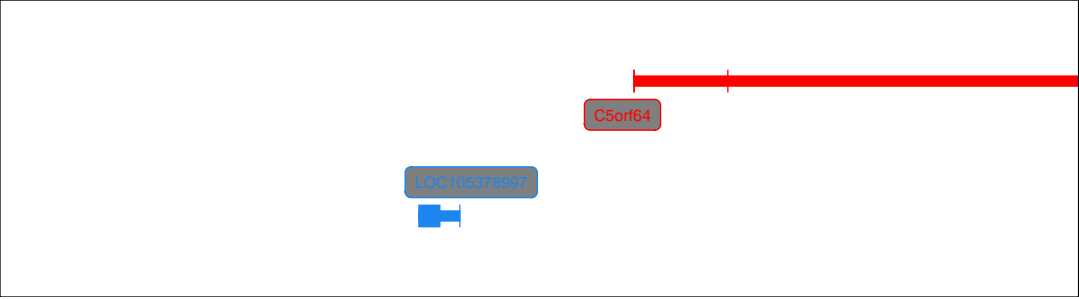
L5.PCP4



Peaks

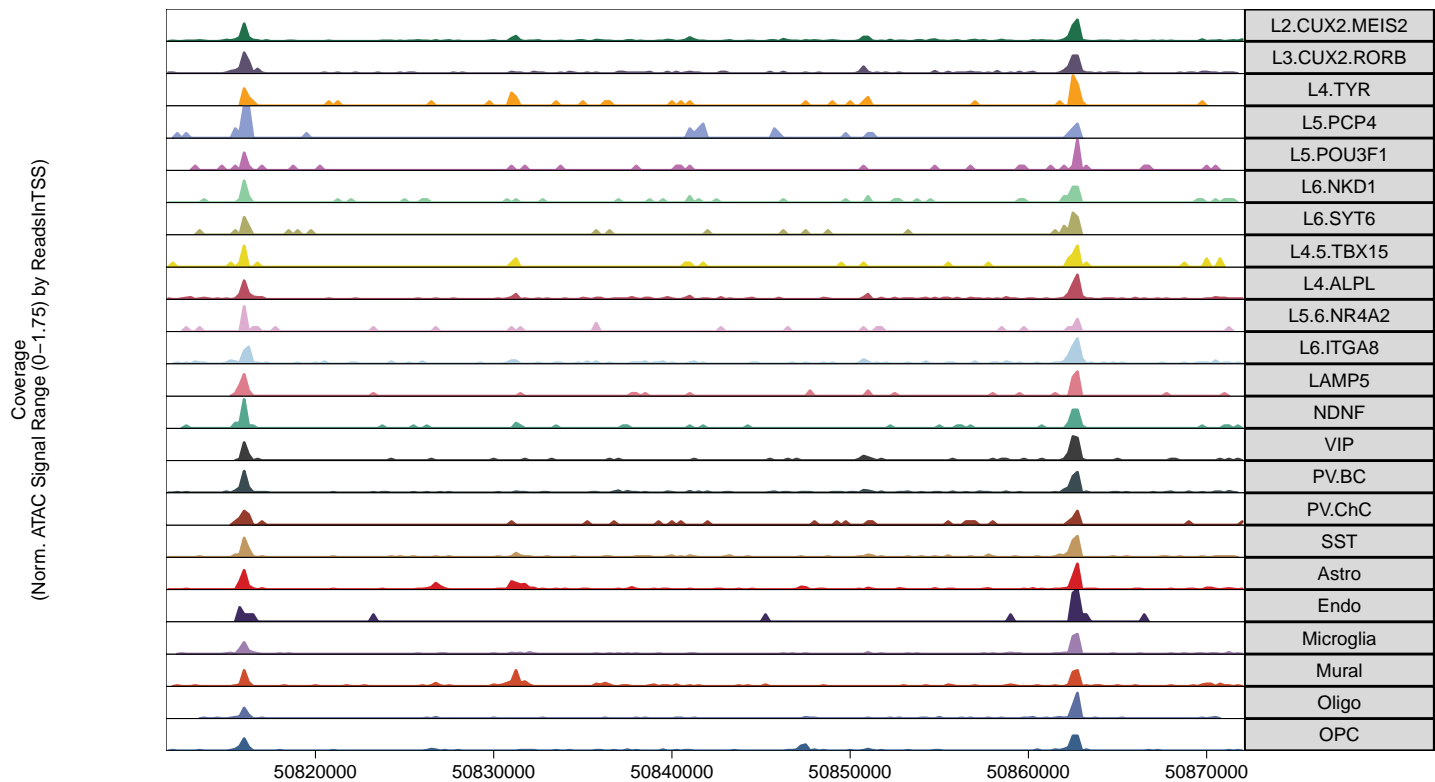


LoopTrack



Genes

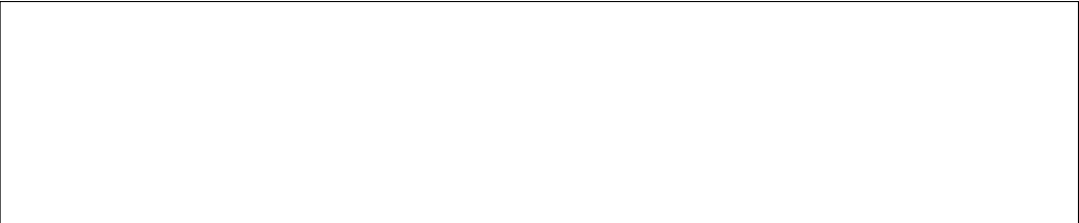
chr16:50811620–50872121 Marker Peak Tracks for: L5.PCP4



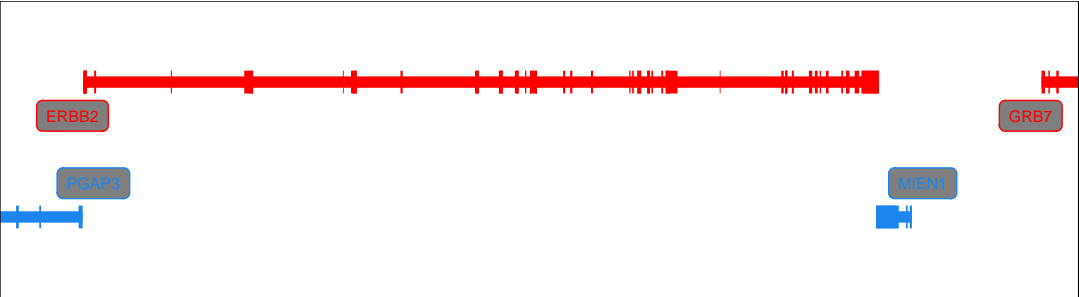
L5.PCP4



Peaks



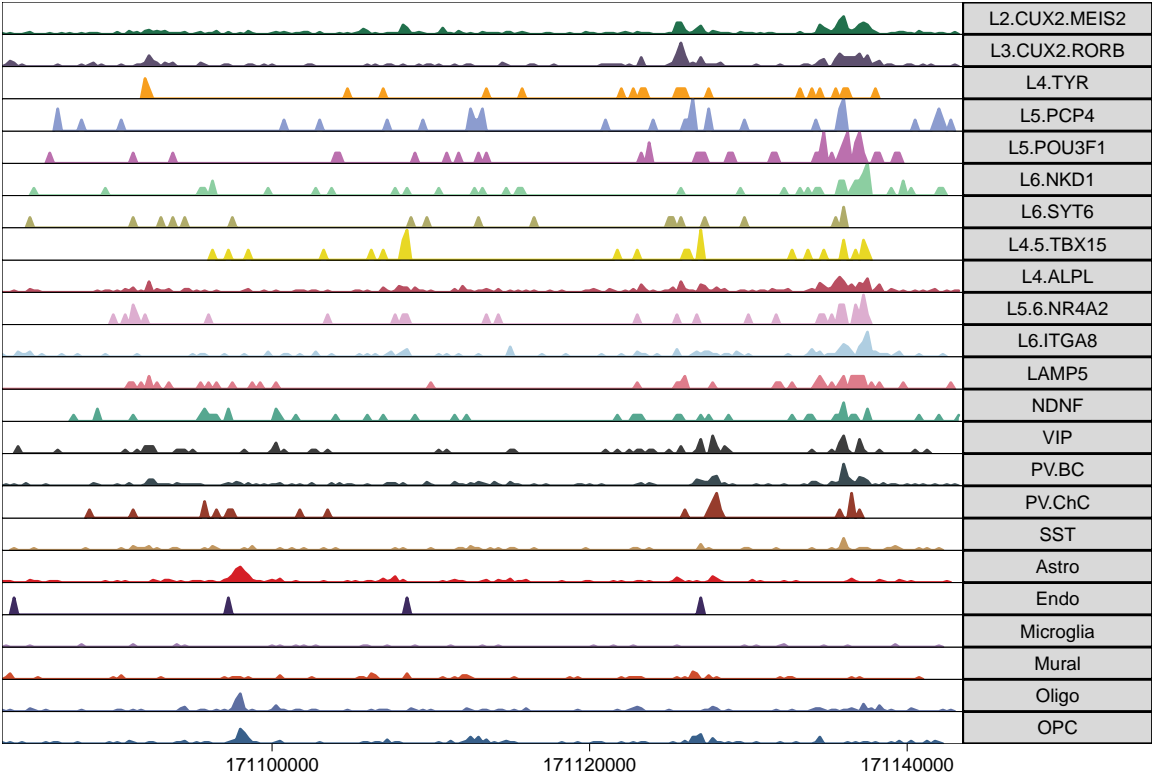
LoopTrack



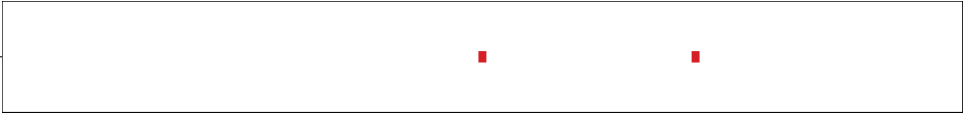
Genes

chr6:171082997-171143498 Marker Peak Tracks for: L5.PCP4

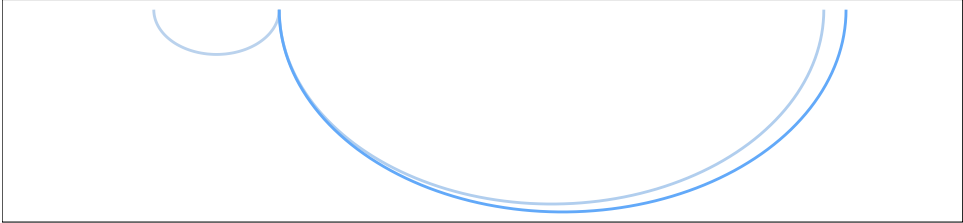
Coverage
(Norm. ATAC Signal Range 0-0.74) by ReadsInTSS)



L5.PCP4

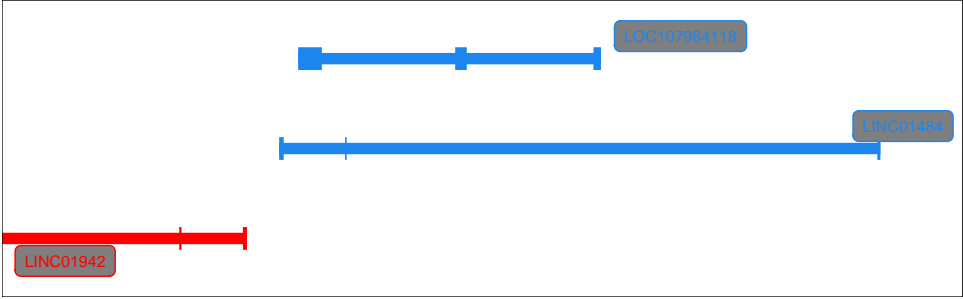


Peaks



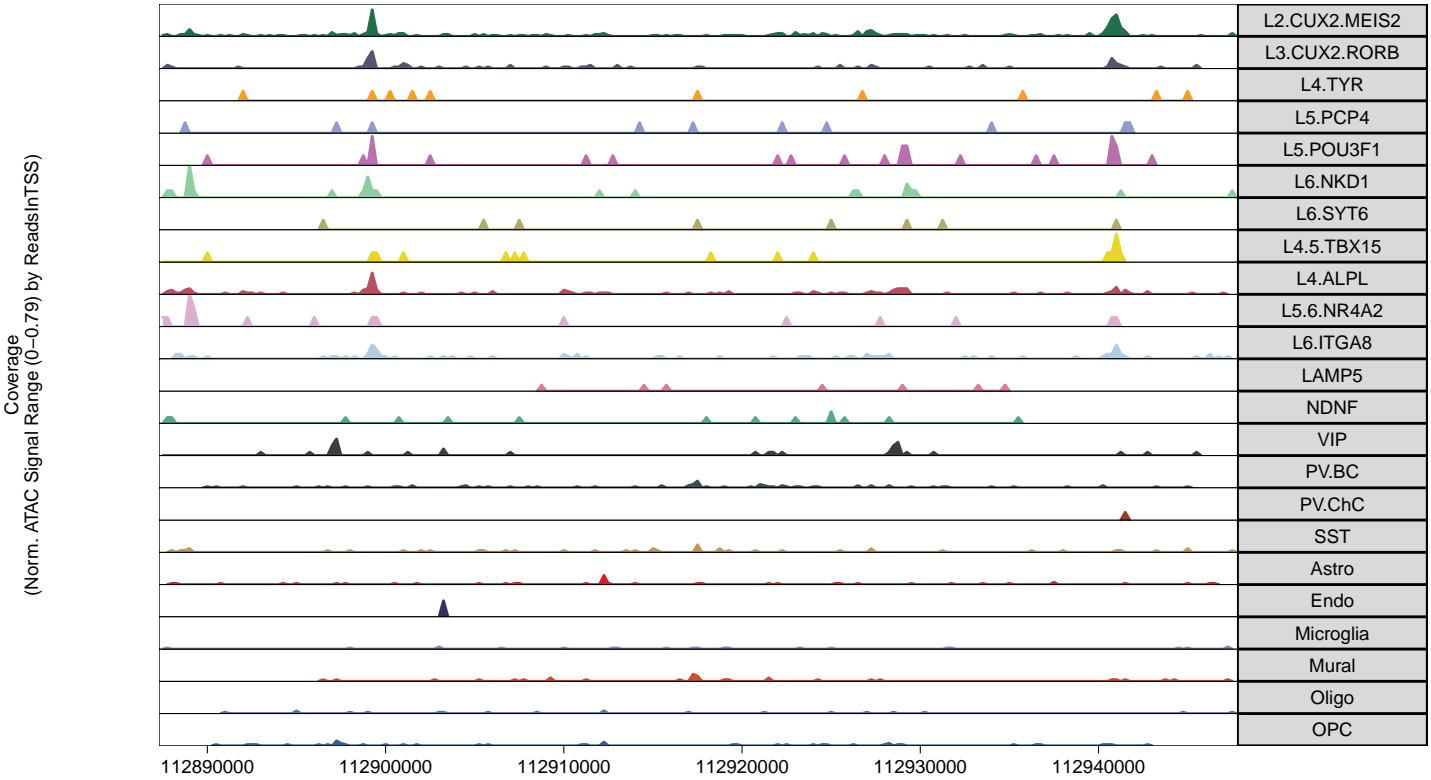
Peak2GeneLinks

value



Genes

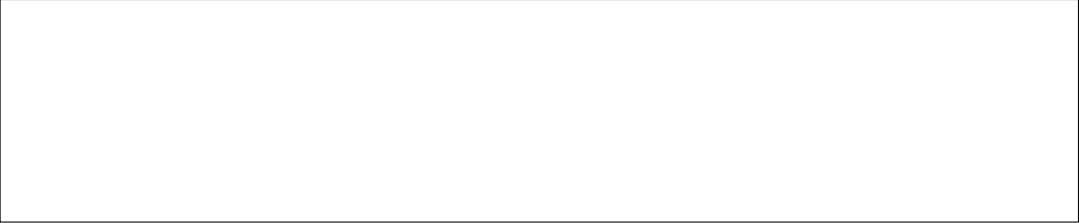
chr6:112887293–112947794 Marker Peak Tracks for: L5.PCP4



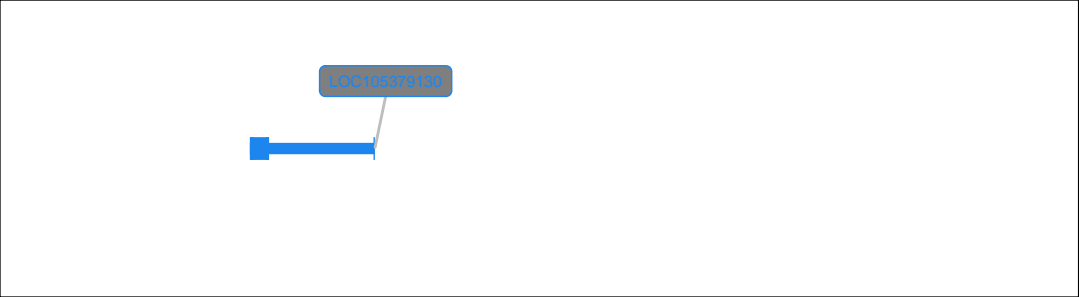
L5.PCP4



Peaks

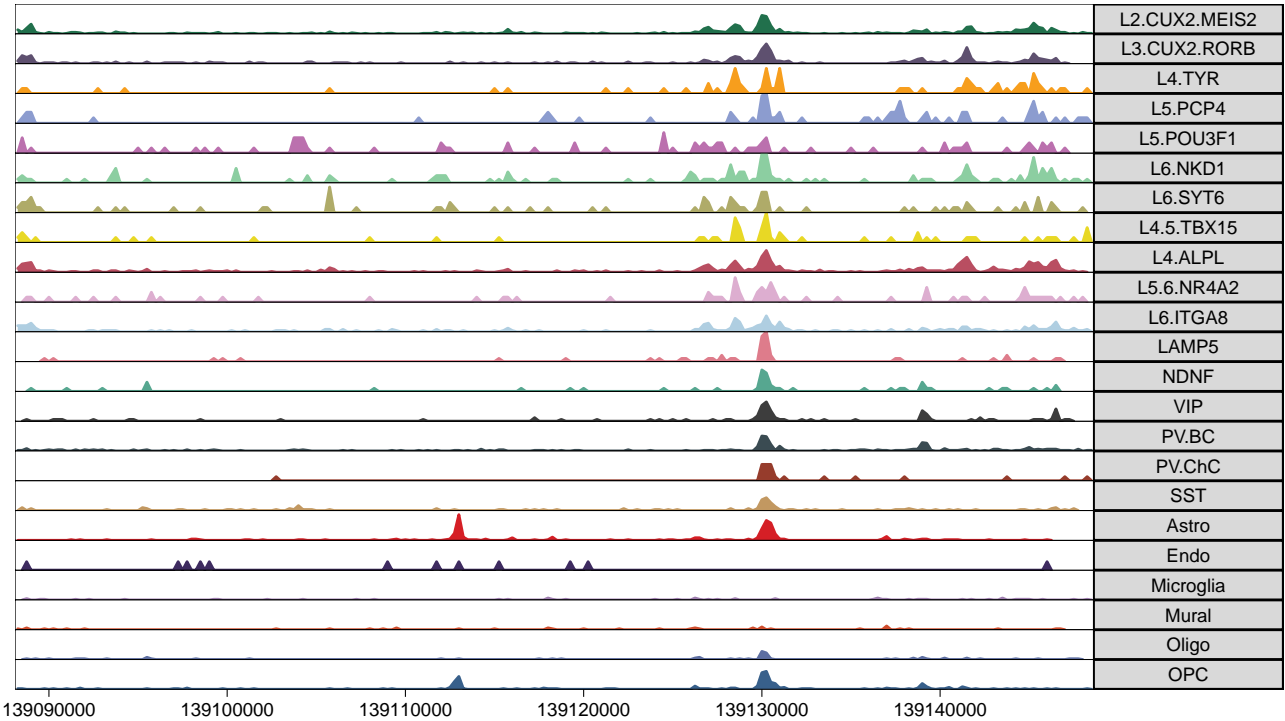


LoopTrack

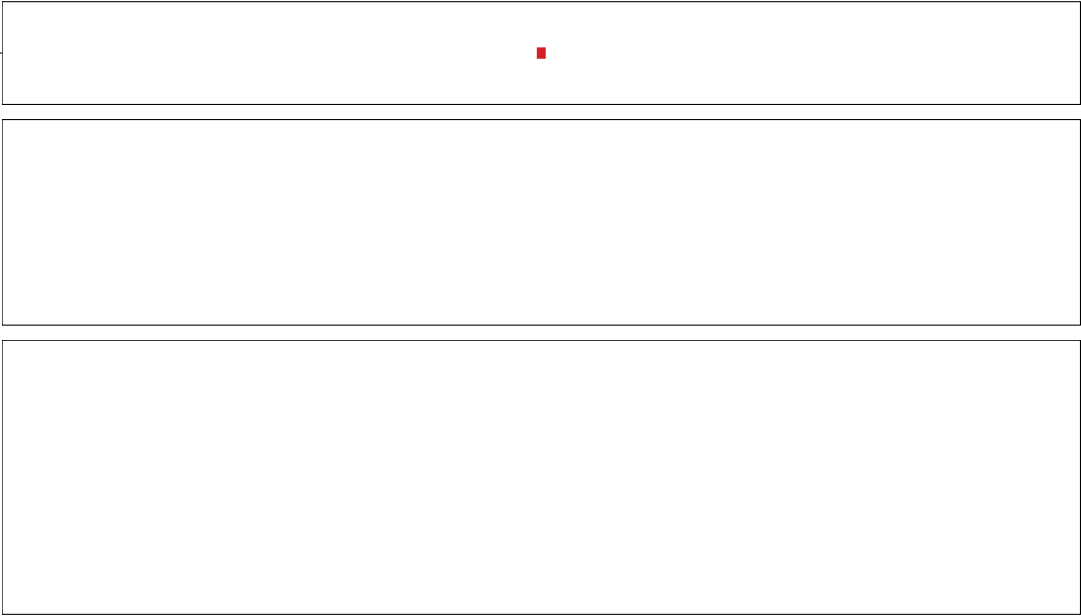


Genes

Coverage
(Norm. ATAC Signal Range (0–1.39) by Reads in TSS)



L5.PCP4

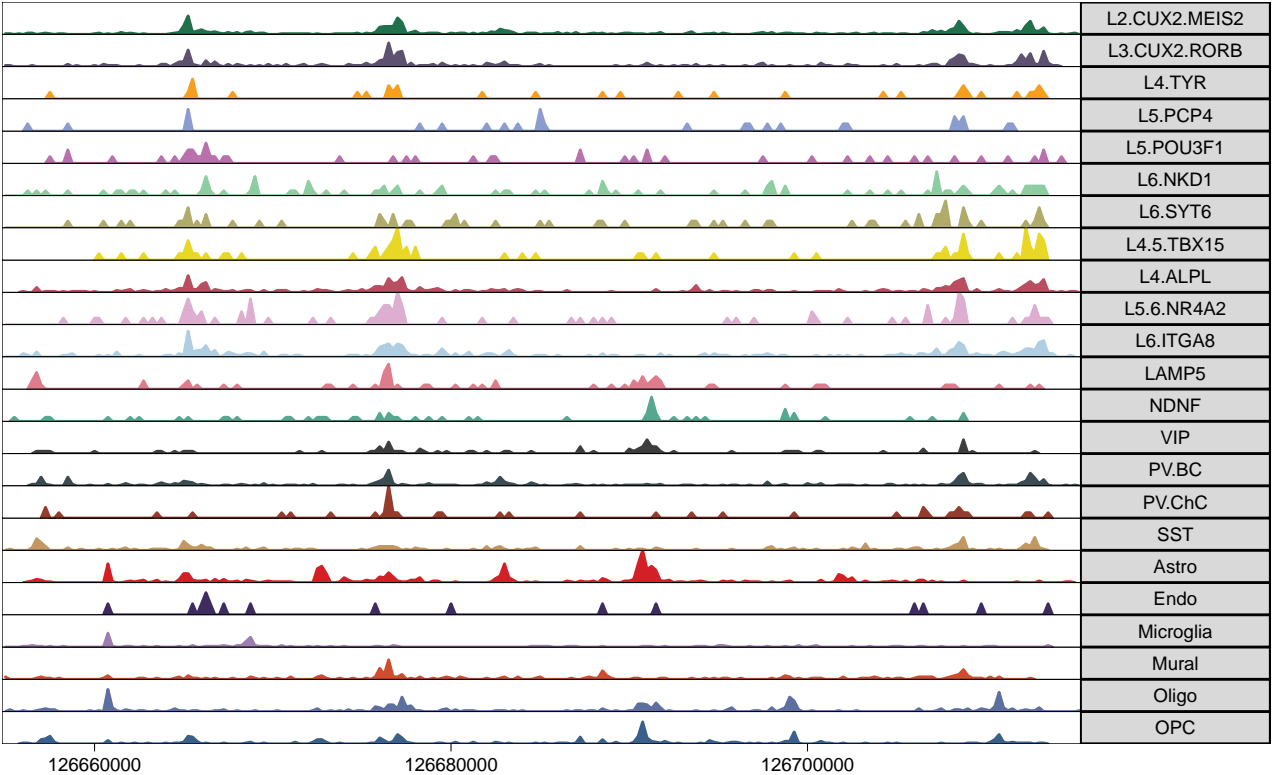


Peaks

LoopTrack

GeneTrack

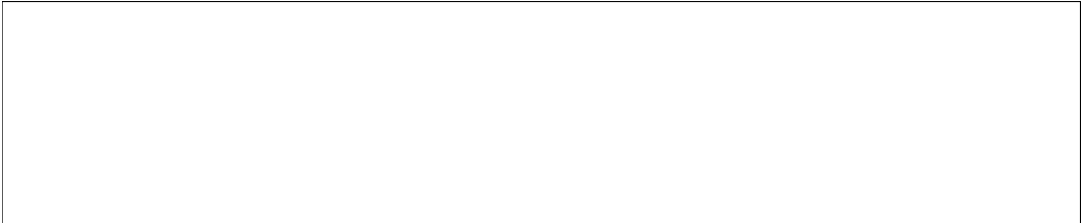
Coverage
(Norm. ATAC Signal Range 0–1.14) by ReadsInTSS)



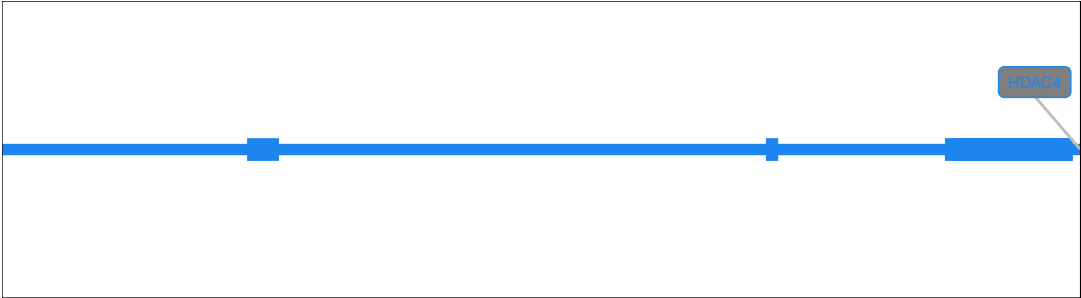
L5.PCP4



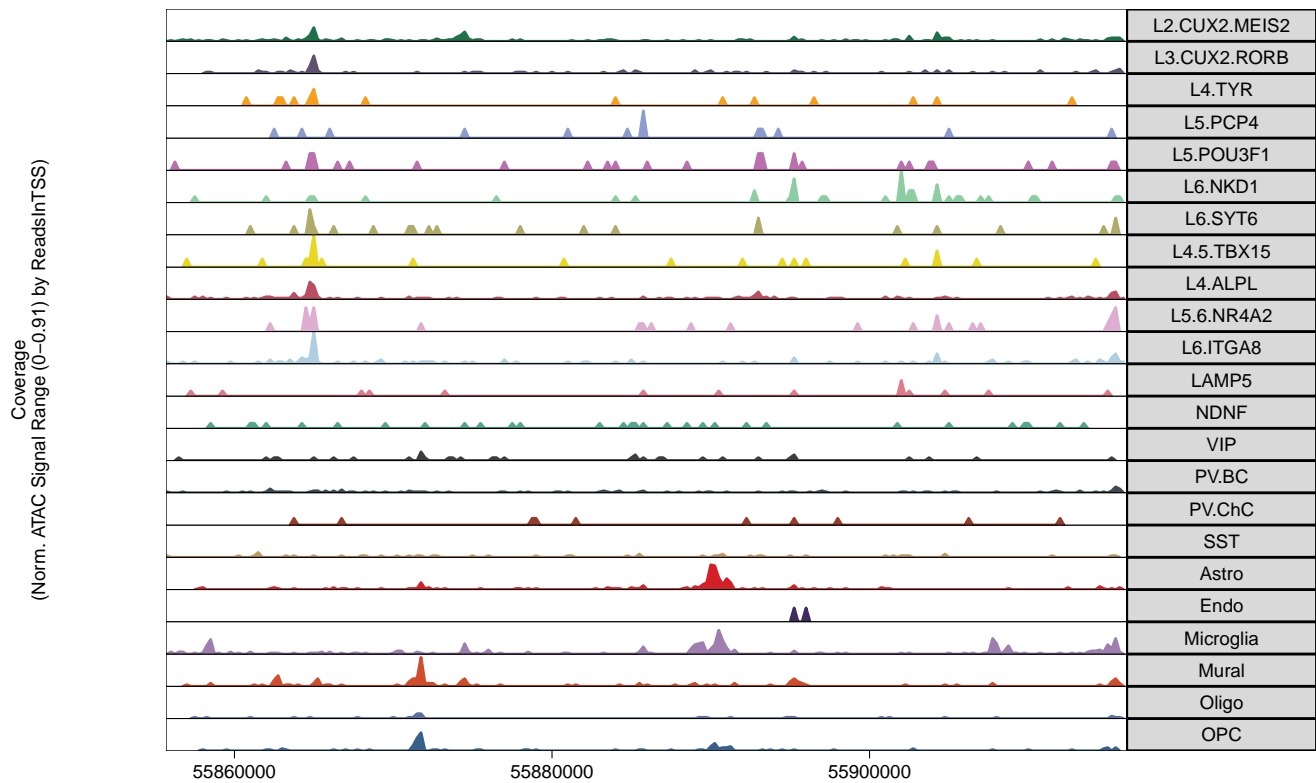
Peaks



LoopTrack



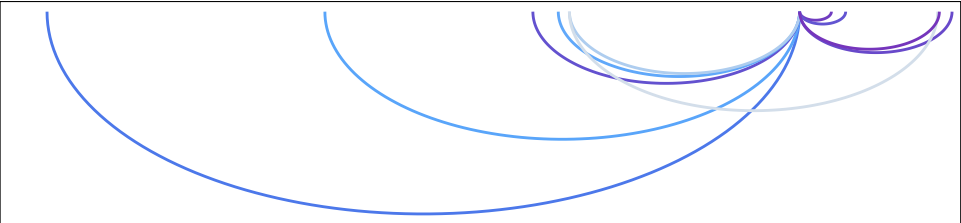
Genes



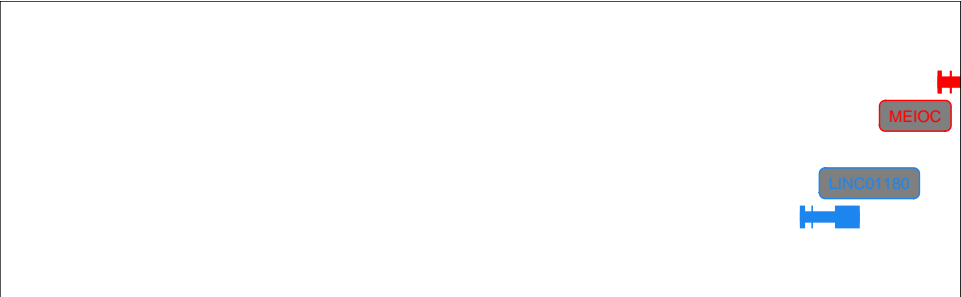
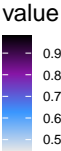
L5.PCP4



Peaks

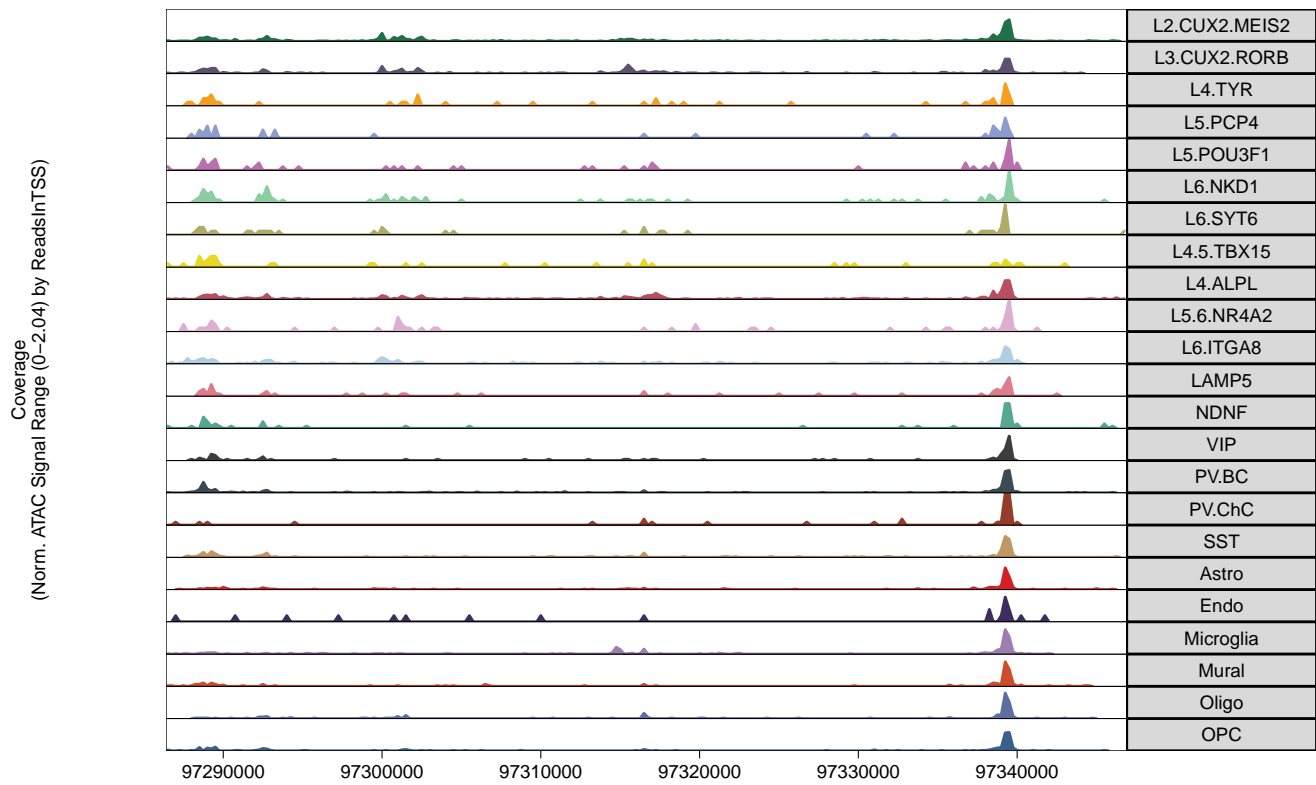


Peak2GeneLinks



Genes

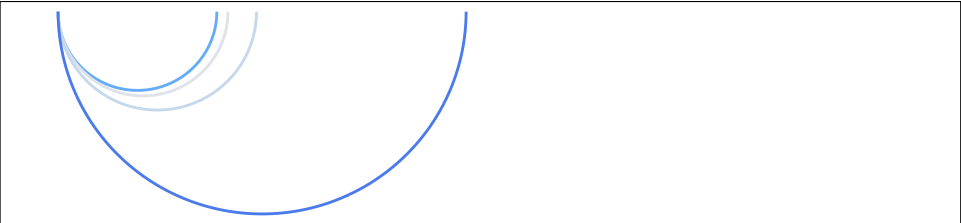
chr1:97286400-97346901 Marker Peak Tracks for: L5.PCP4



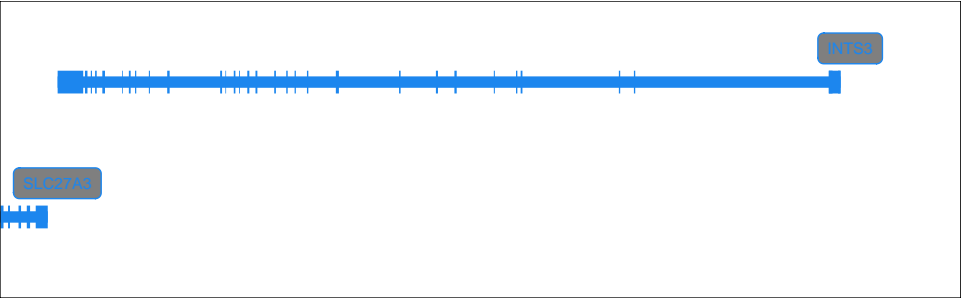
L5.PCP4



Peaks

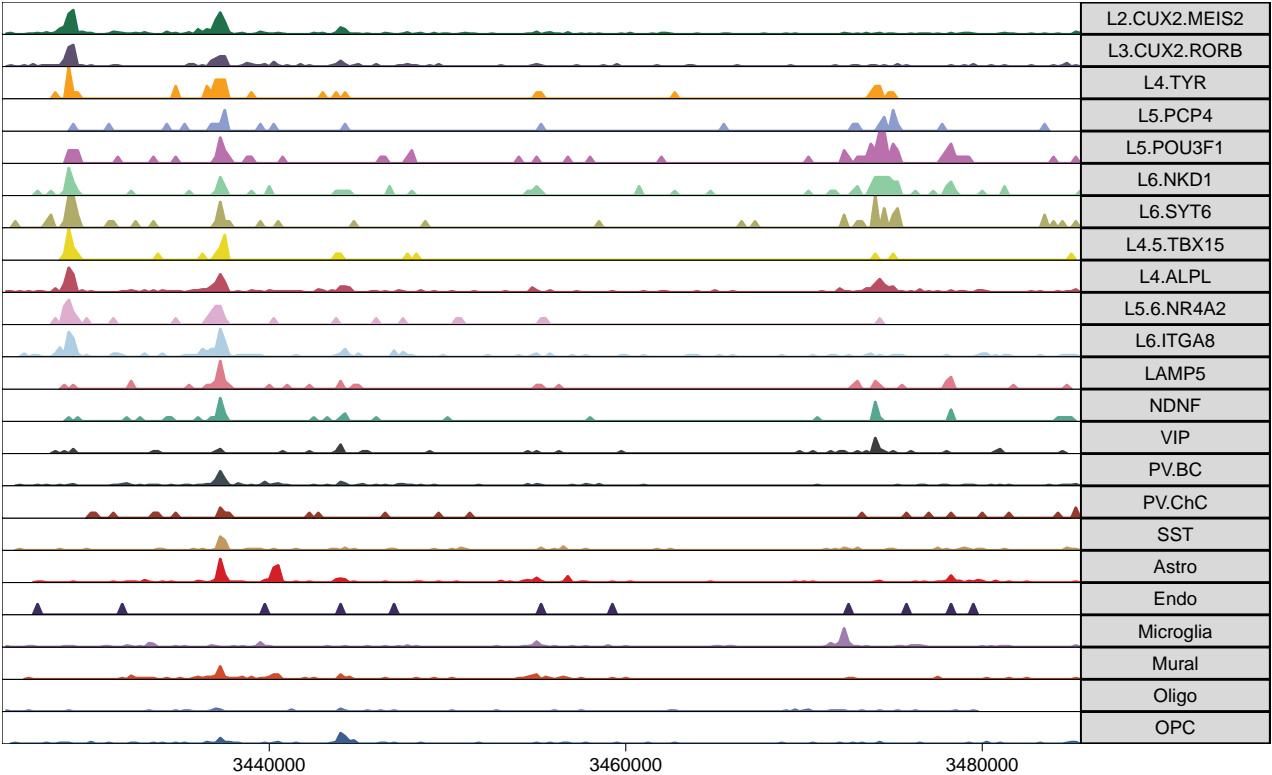


Peak2GeneLinks



Genes

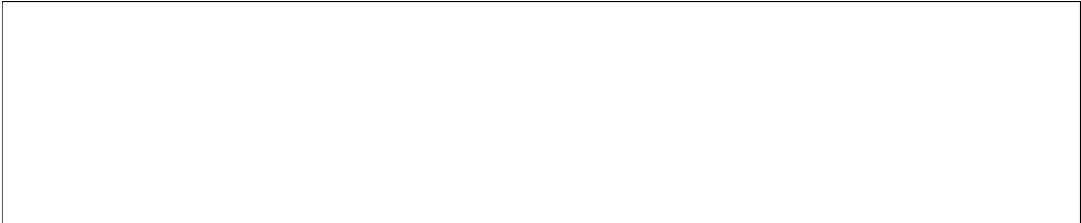
Coverage
(Norm. ATAC Signal Range 0–1.19) by ReadsInTSS)



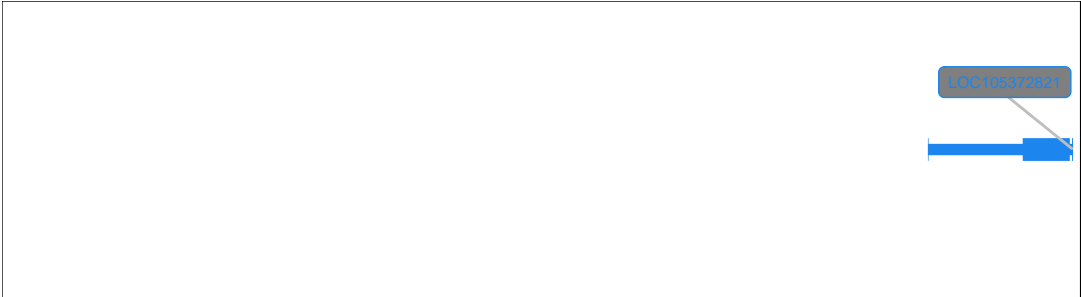
L5.PCP4



Peaks

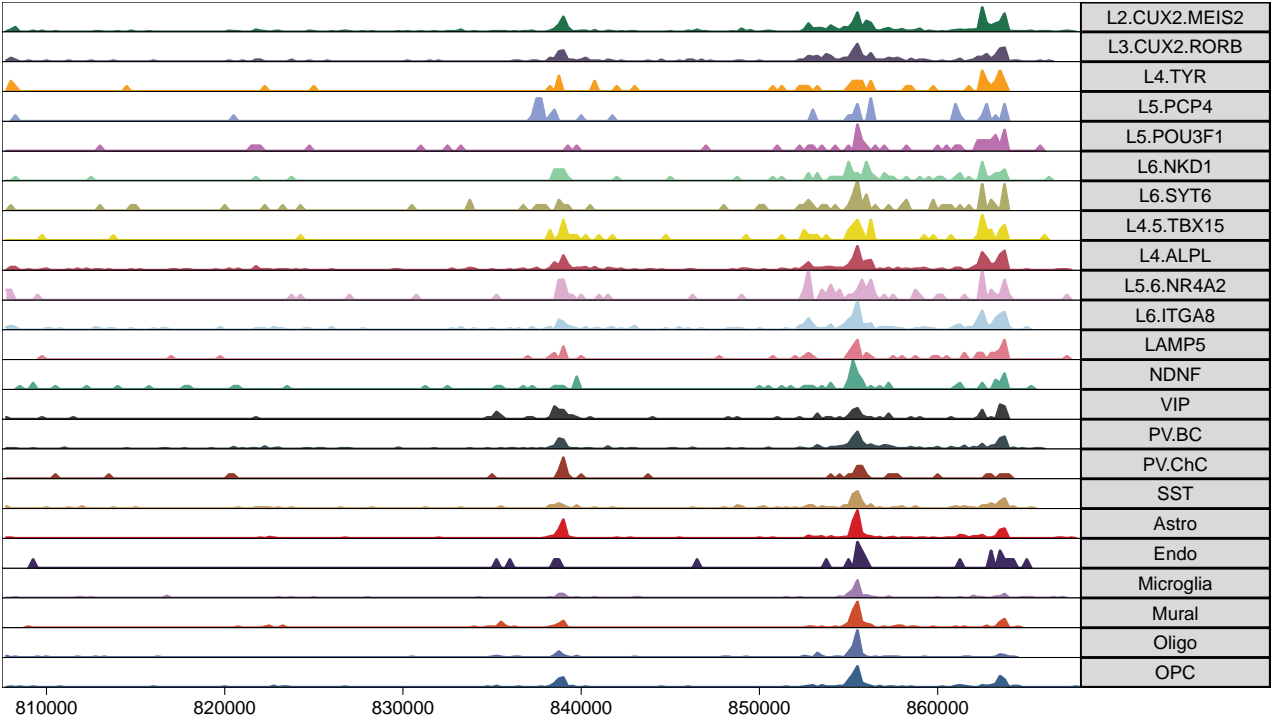


LoopTrack

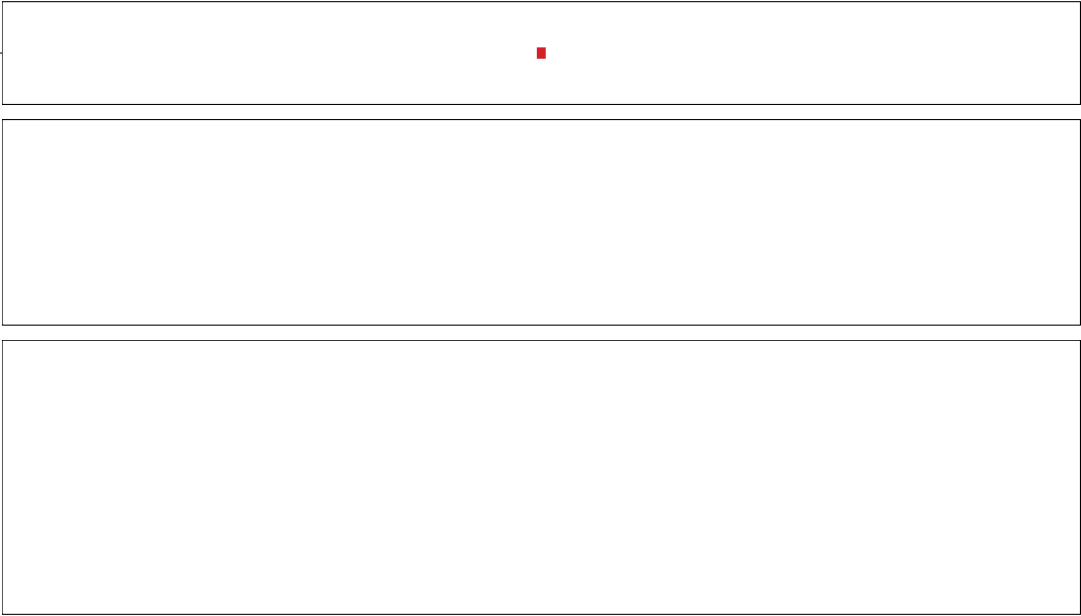


Genes

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



L5.PCP4

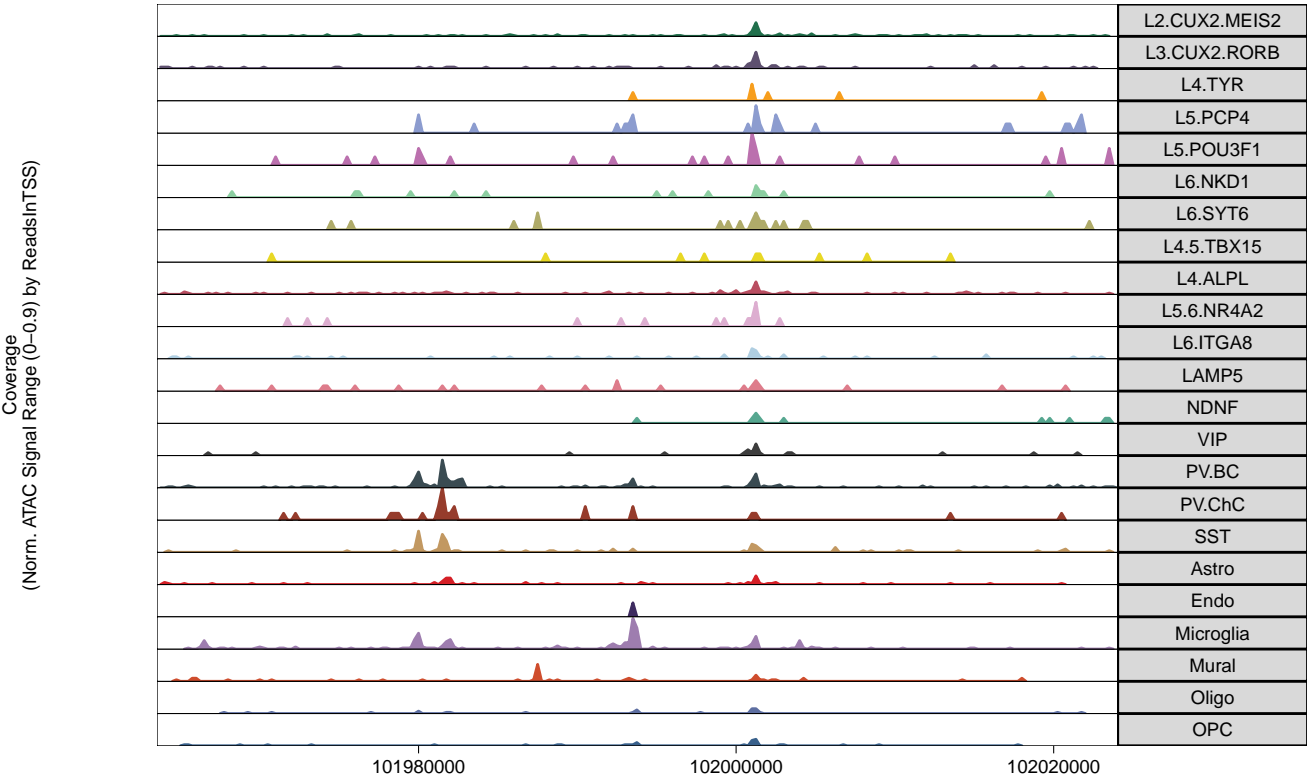


Peaks

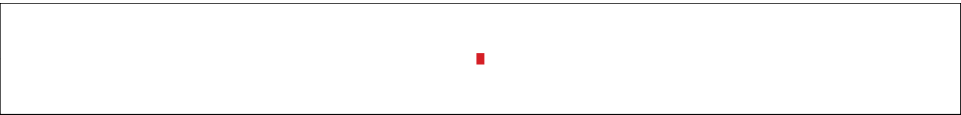
LoopTrack

GeneTrack

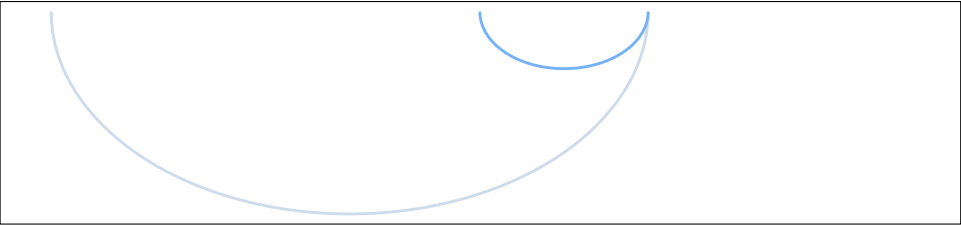
chr13:101963533–102024034 Marker Peak Tracks for: L5.PCP4



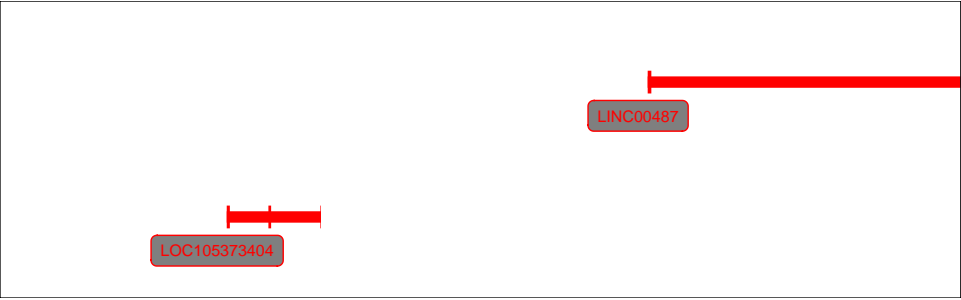
L5.PCP4



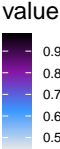
Peaks



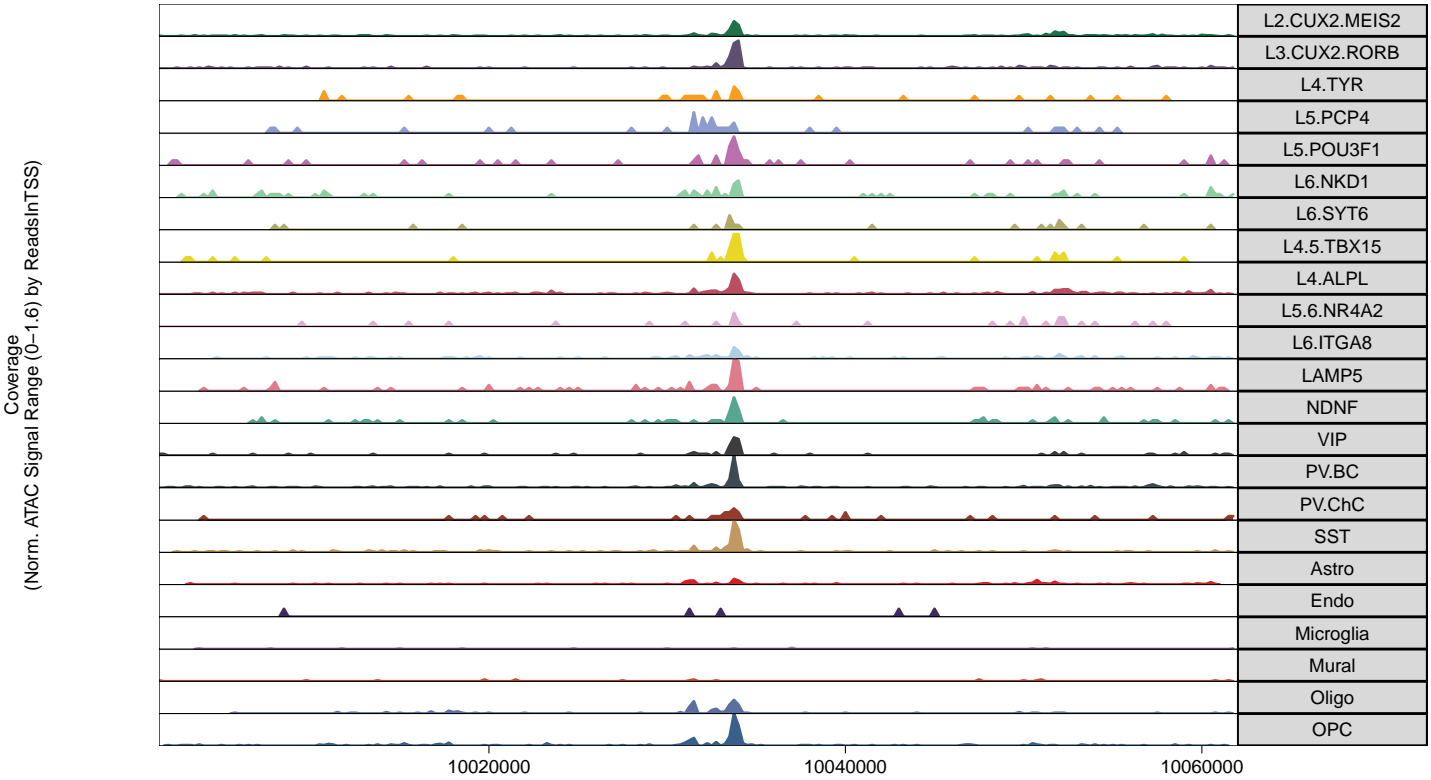
Peak2GeneLinks



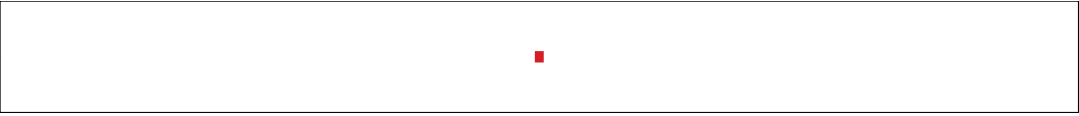
Genes



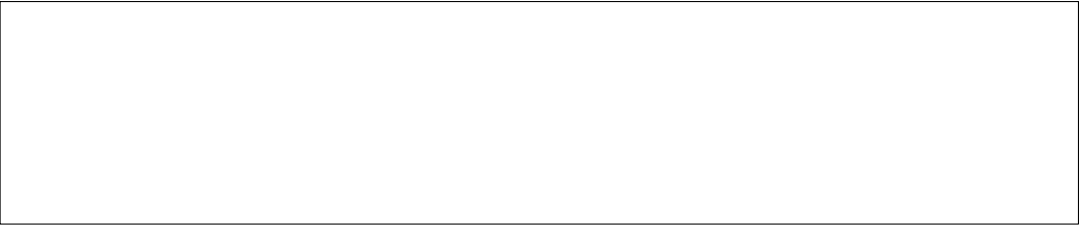
chr8:10001495–10061996 Marker Peak Tracks for: L5.PCP4



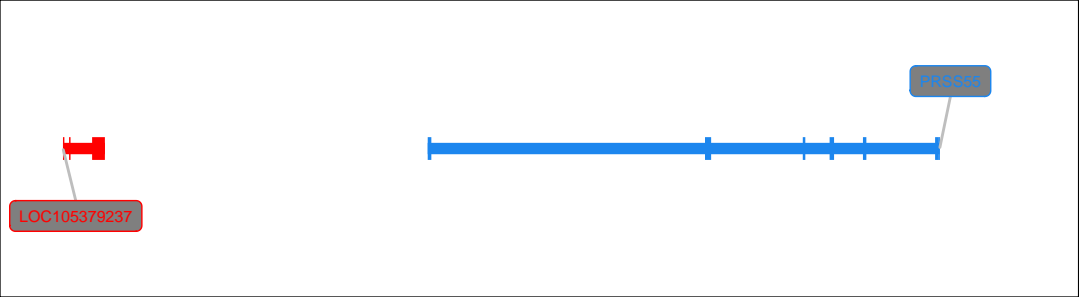
L5.PCP4



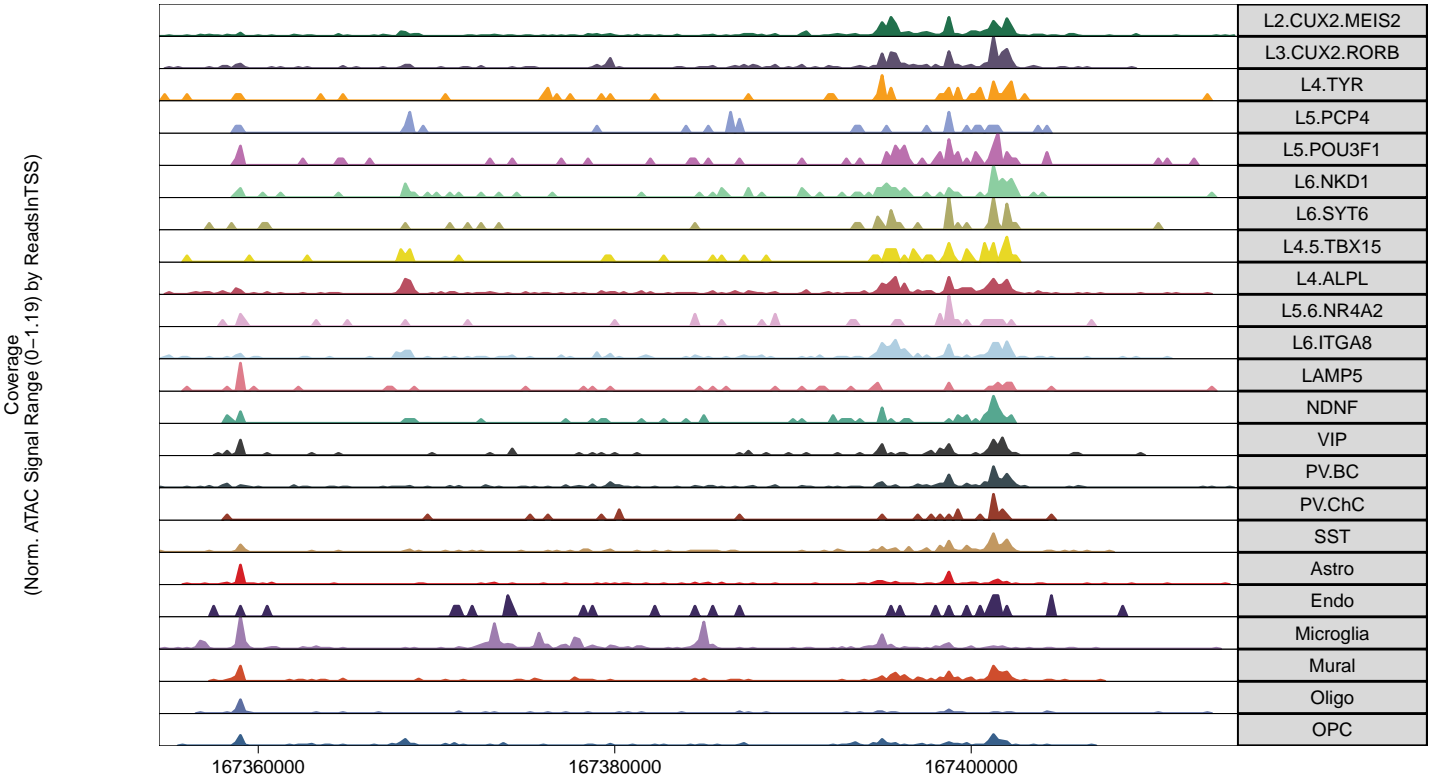
Peaks



LoopTrack



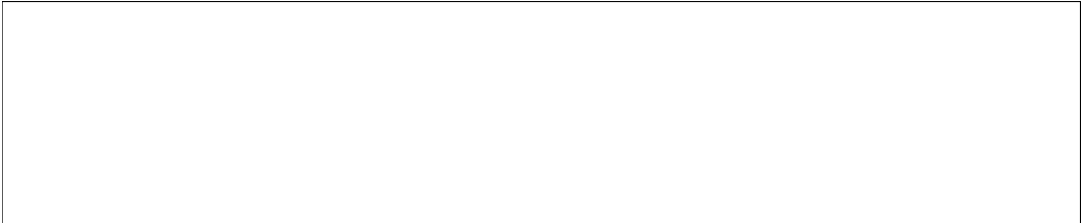
Genes



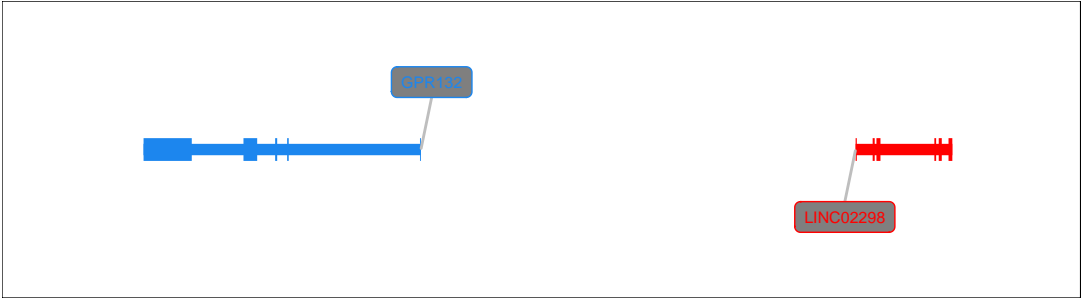
L5.PCP4



Peaks

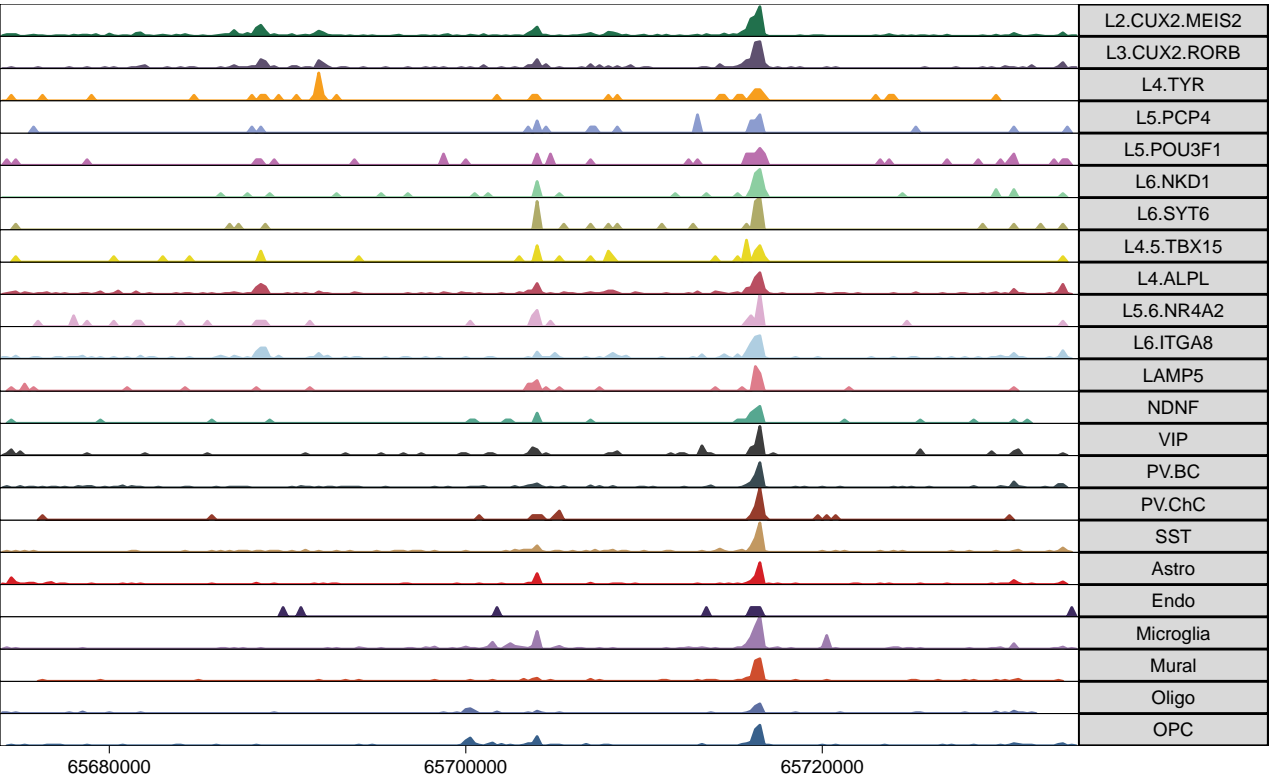


LoopTrack



Genes

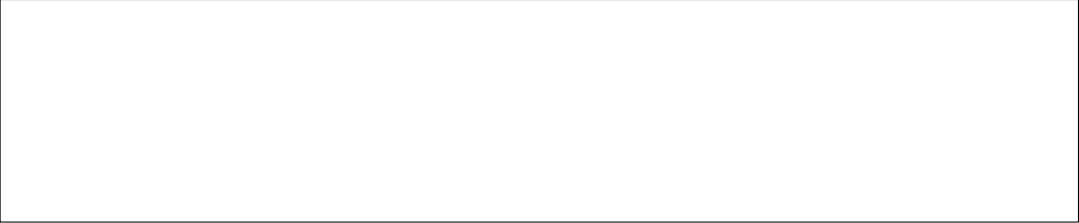
Coverage
(Norm. ATAC Signal Range 0–1.35) by ReadsInTSS)



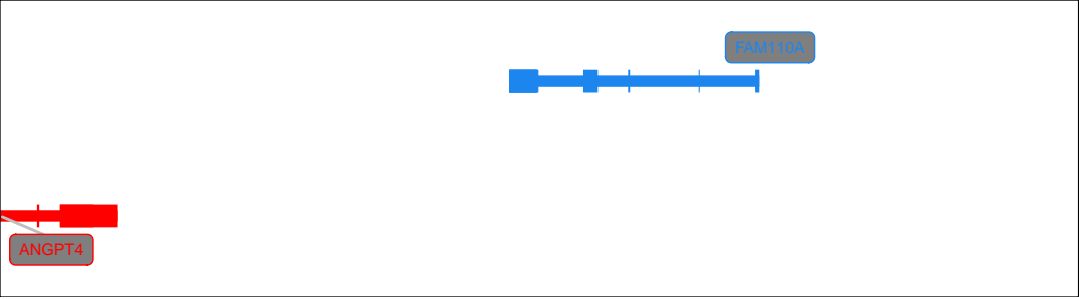
L5.PCP4



Peaks



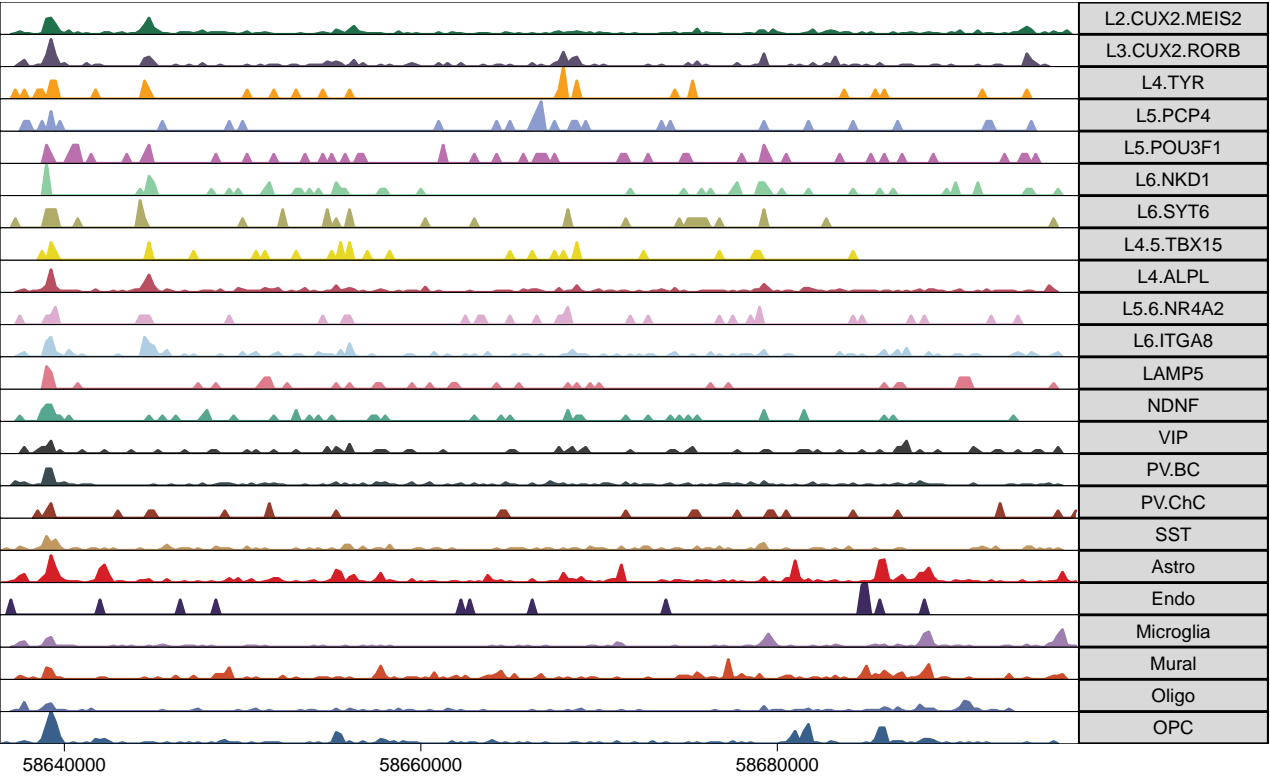
LoopTrack



Genes

chr9:58636394–58696895 Marker Peak Tracks for: L5.PCP4

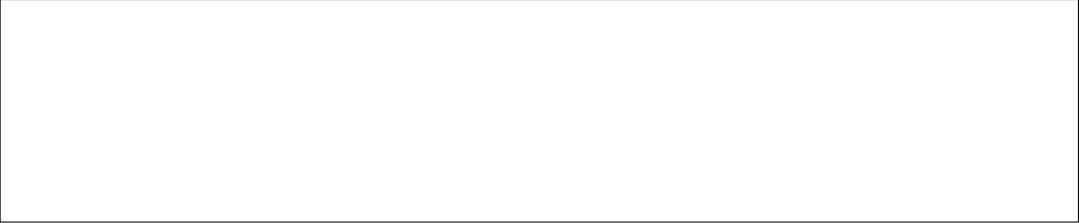
Coverage
(Norm. ATAC Signal Range 0–0.84) by ReadsInTSS)



L5.PCP4



Peaks

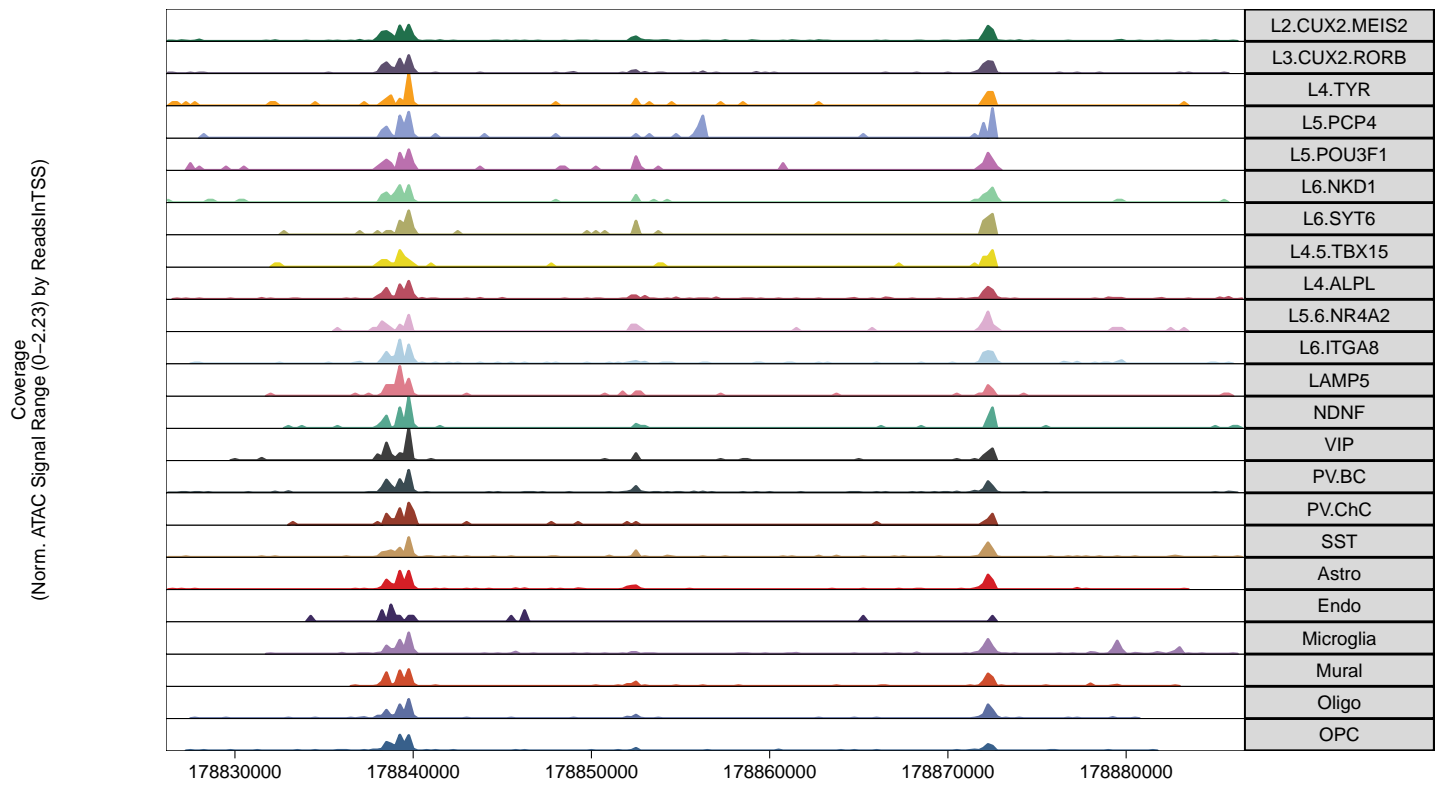


LoopTrack



Genes

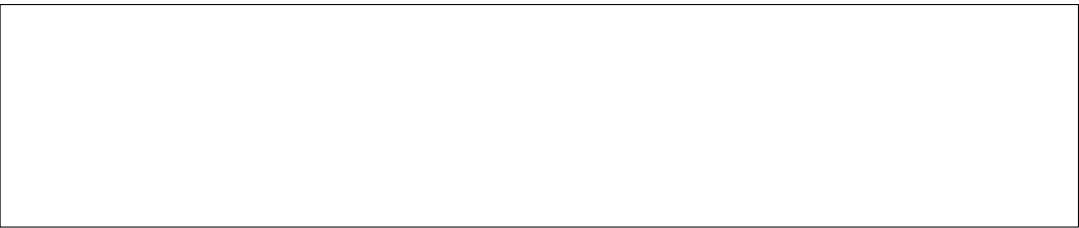
chr1:178826137-178886638 Marker Peak Tracks for: L5.PCP4



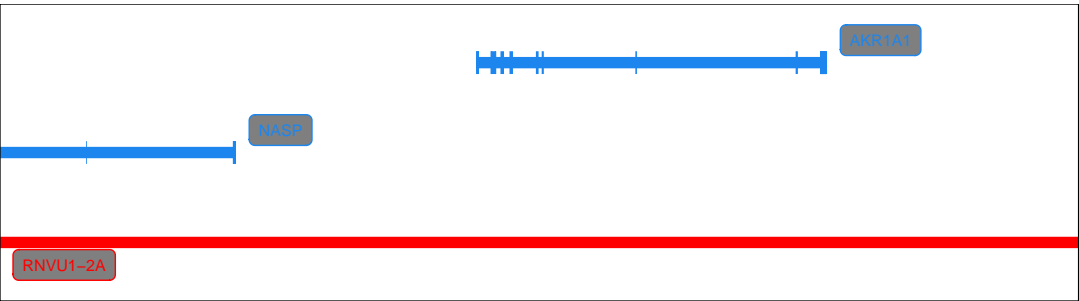
L5.PCP4



Peaks

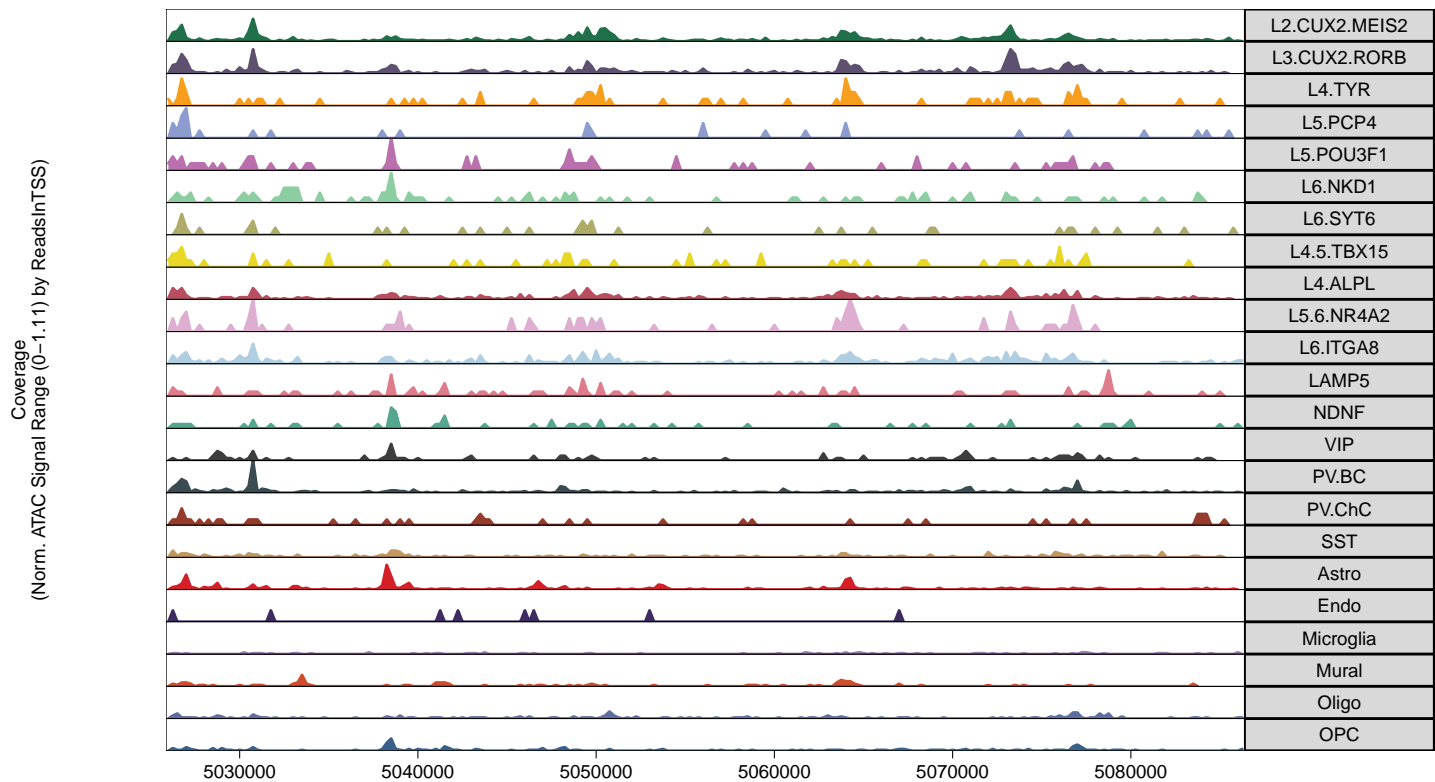


LoopTrack



Genes

chr19:5025872–5086373 Marker Peak Tracks for: L5.PCP4



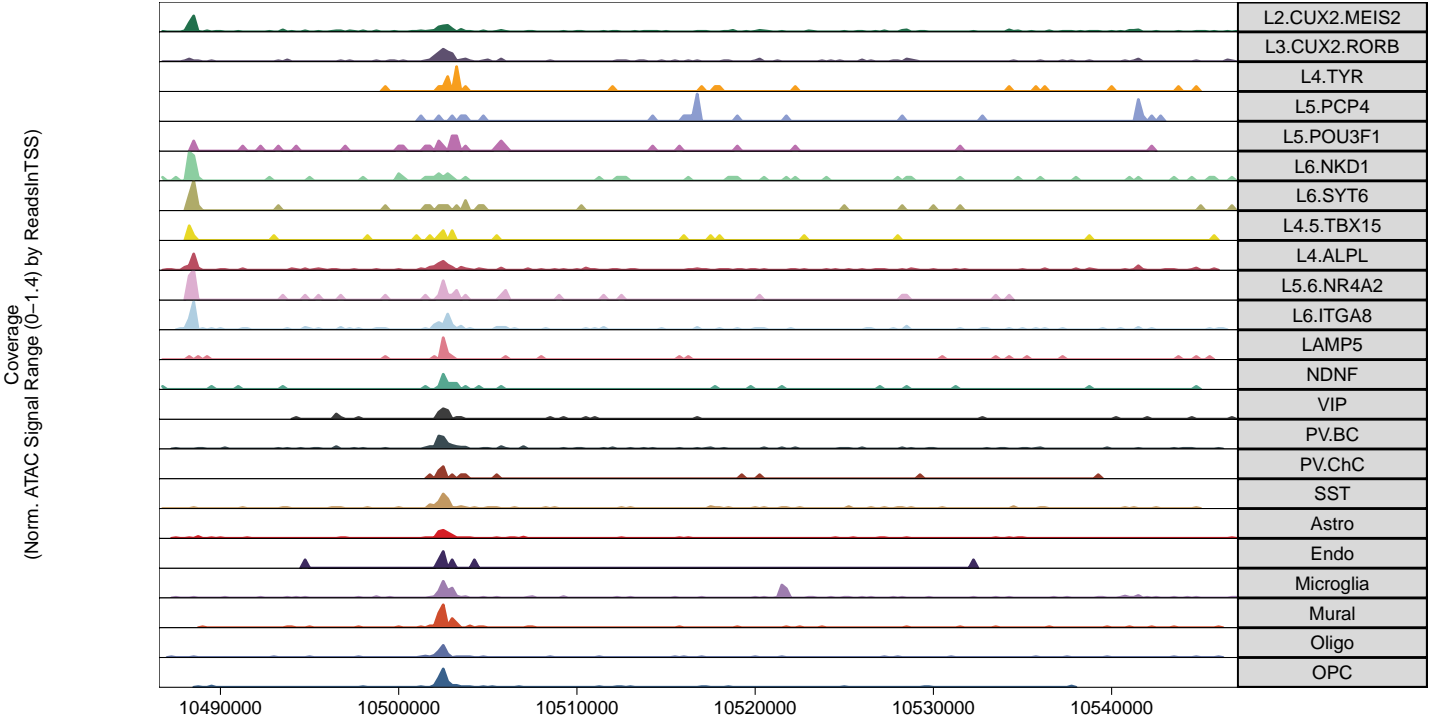
L5.PCP4



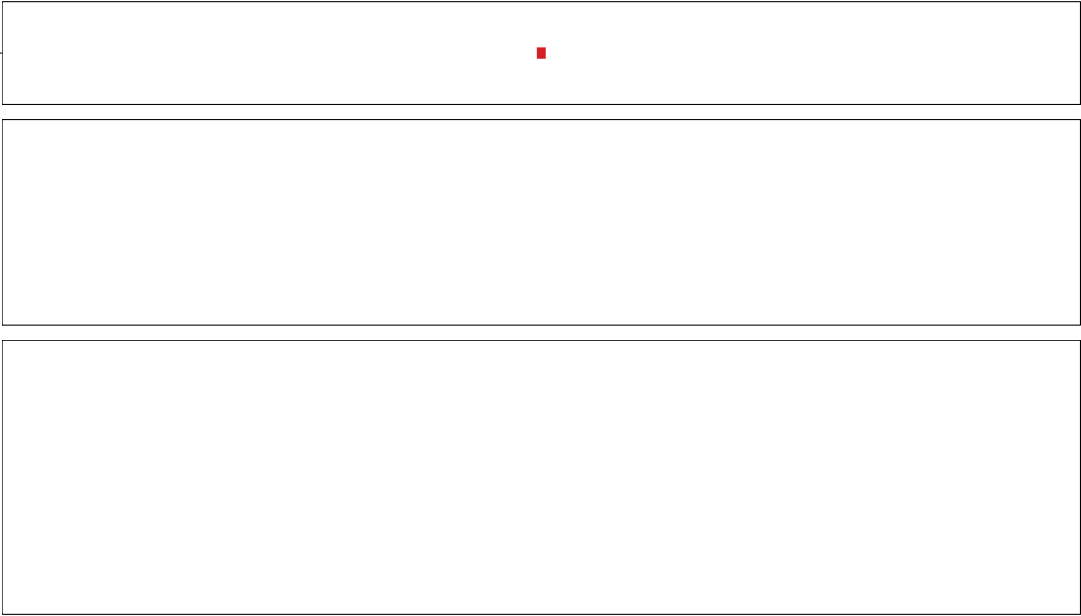
Peaks

LoopTrack

Genes



L5.PCP4

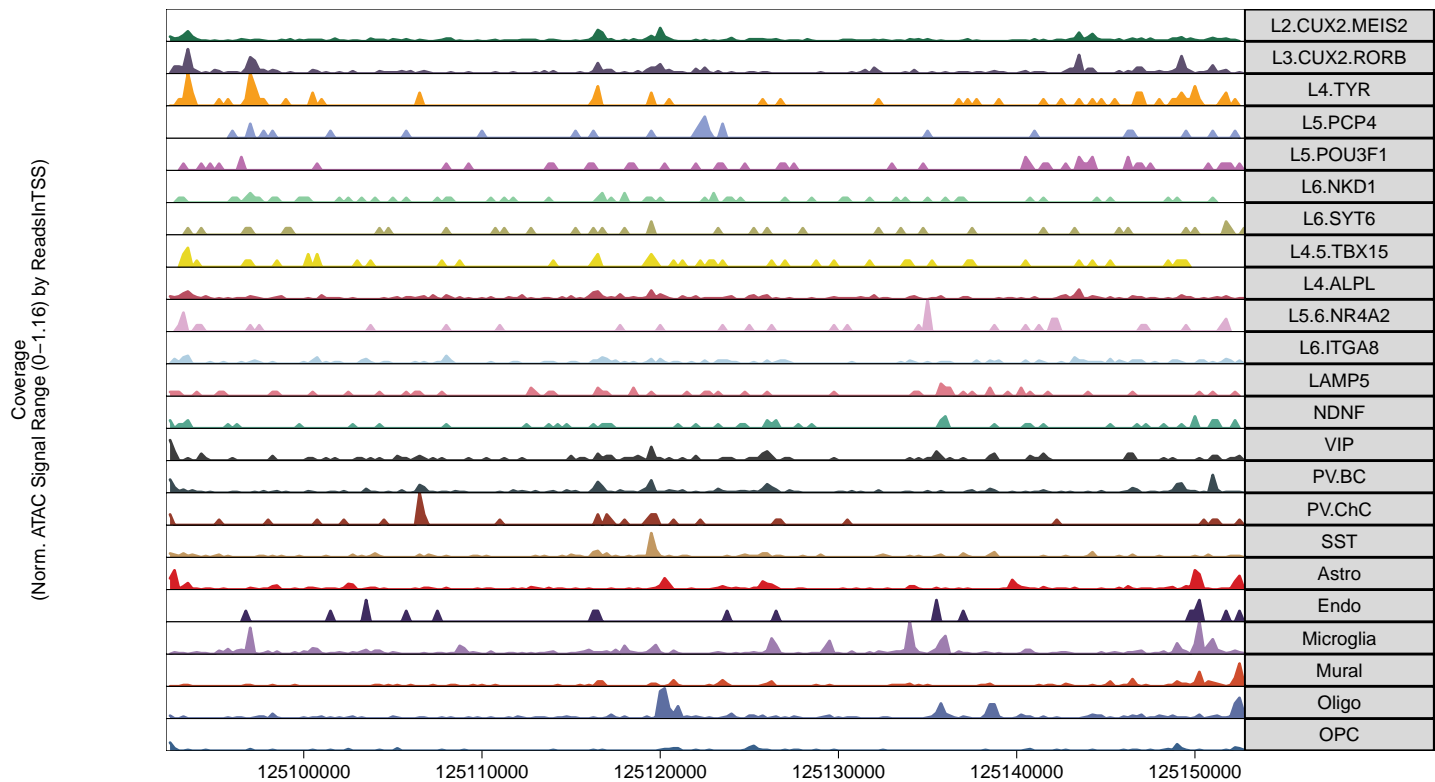


Peaks

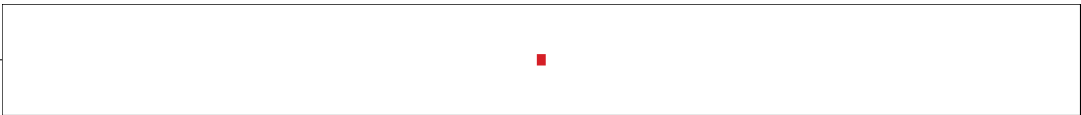
LoopTrack

GeneTrack

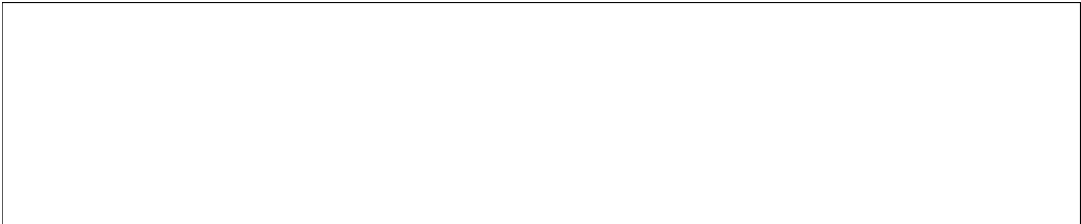
chr9:125092277-125152778 Marker Peak Tracks for: L5.PCP4



L5.PCP4



Peaks

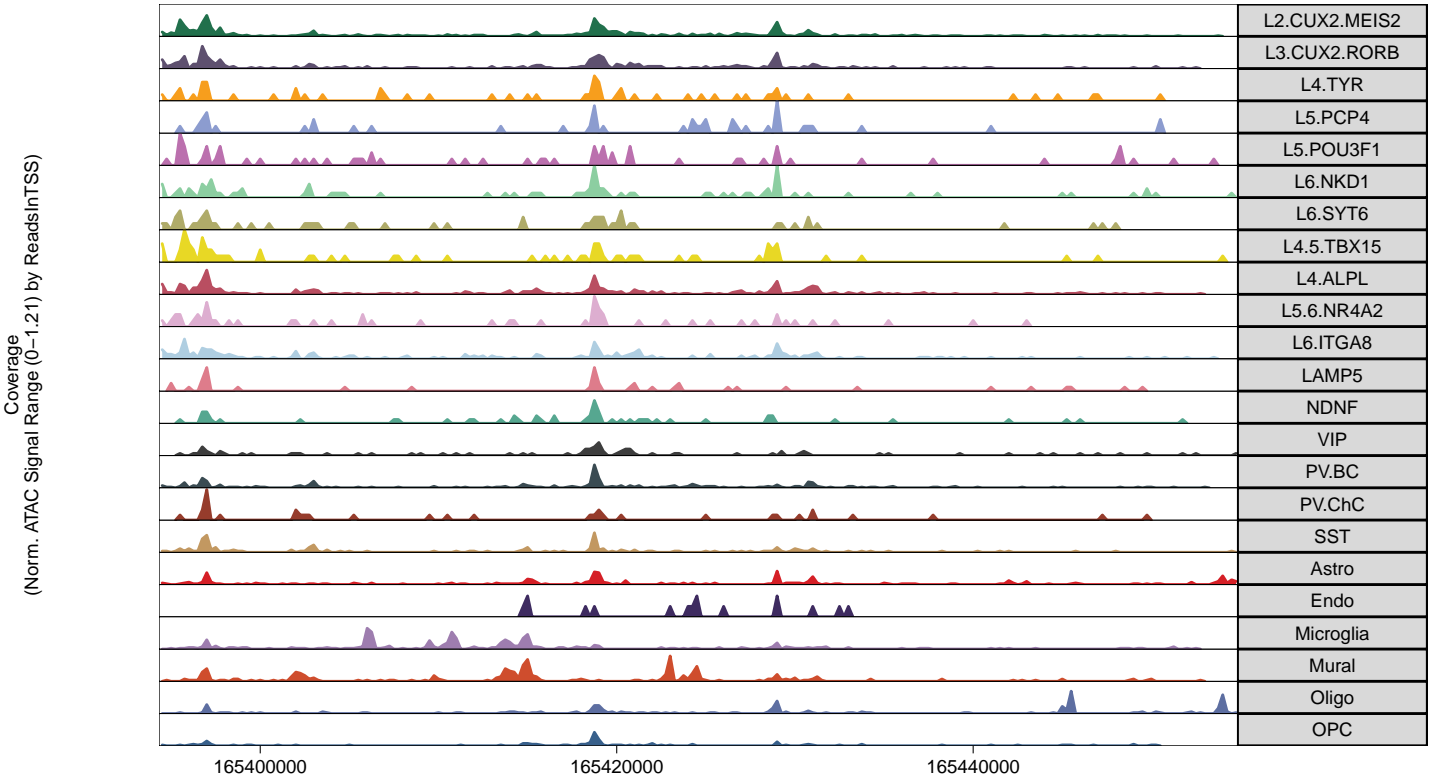


LoopTrack



Genes

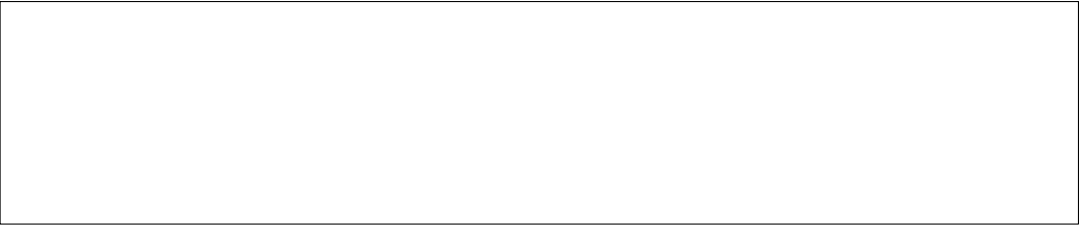
chr7:165394326-165454827 Marker Peak Tracks for: L5.PCP4



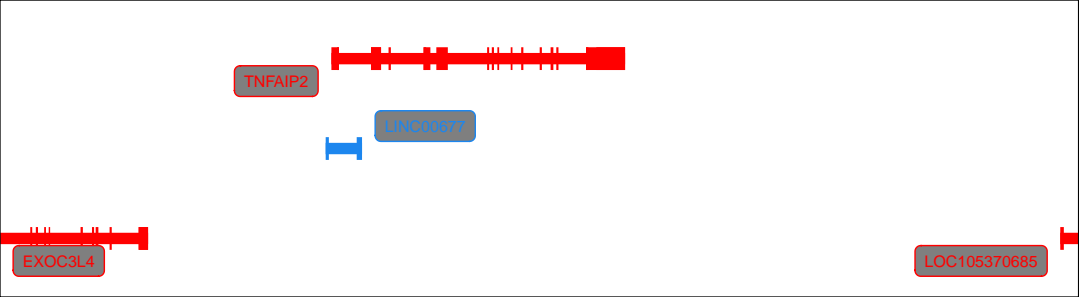
L5.PCP4



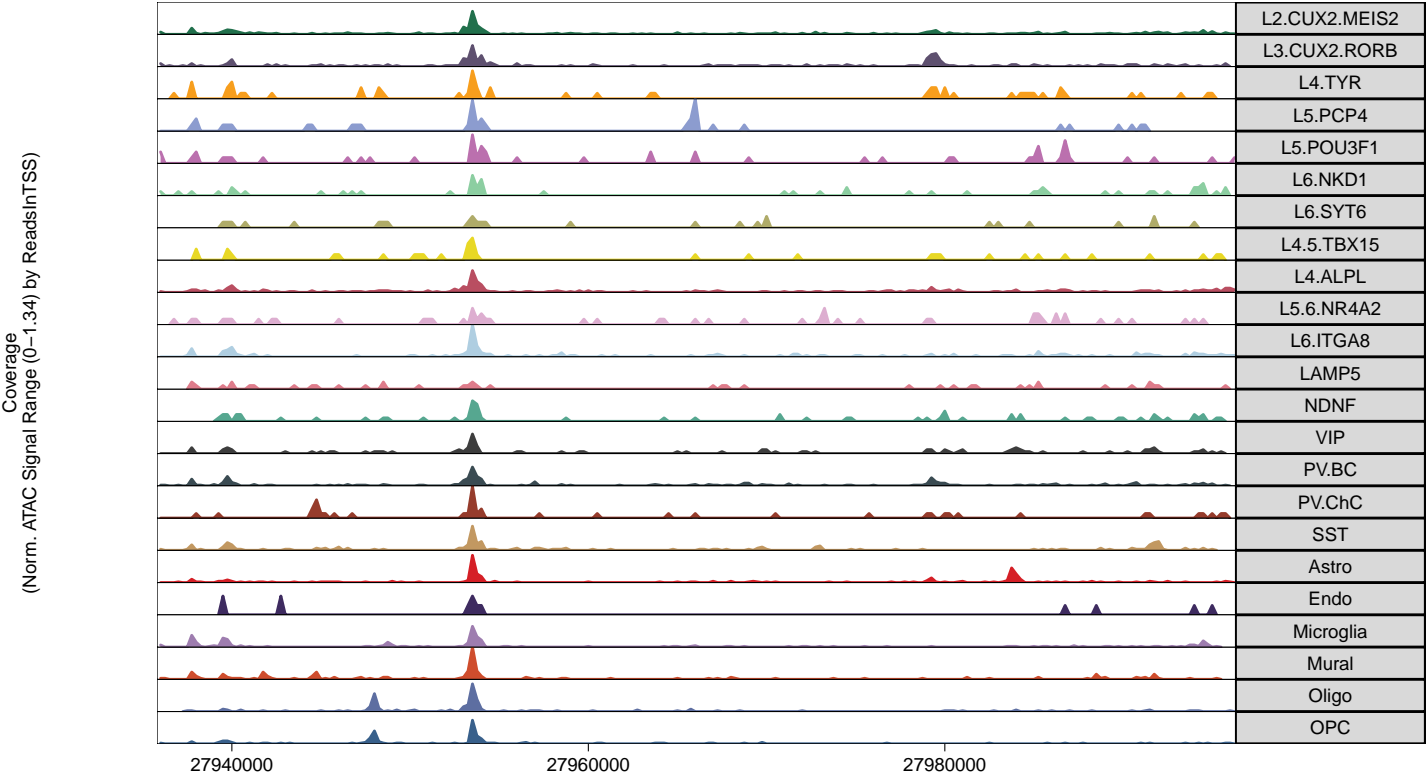
Peaks



LoopTrack



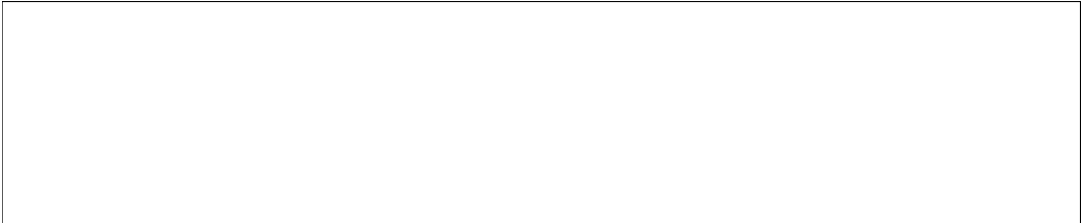
Genes



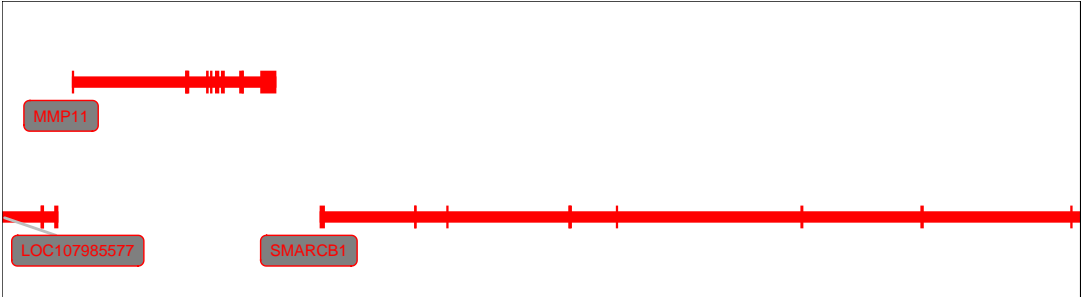
L5.PCP4



Peaks

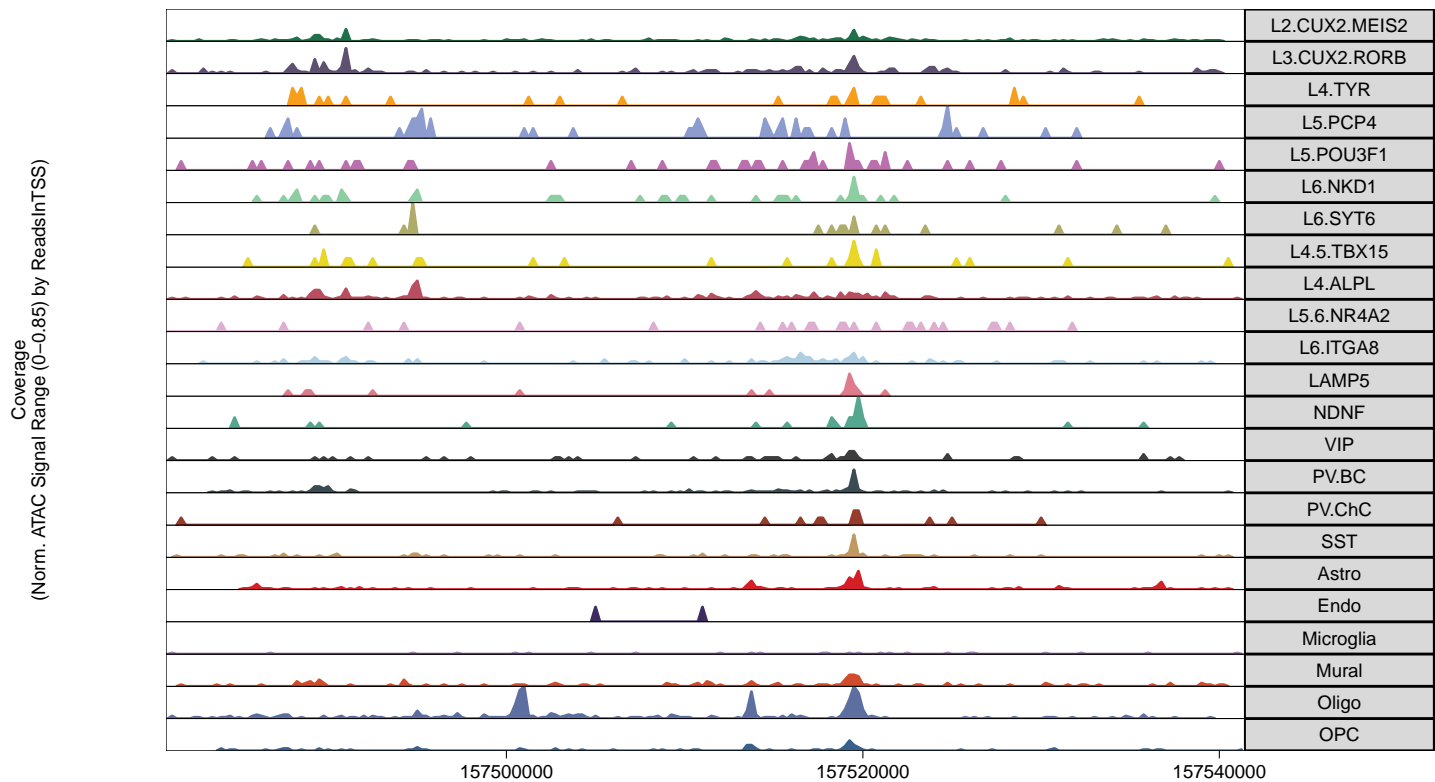


LoopTrack



Genes

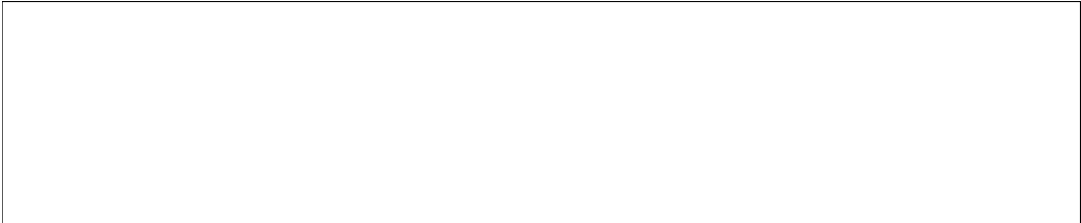
chr7:157480907–157541408 Marker Peak Tracks for: L5.PCP4



L5.PCP4



Peaks

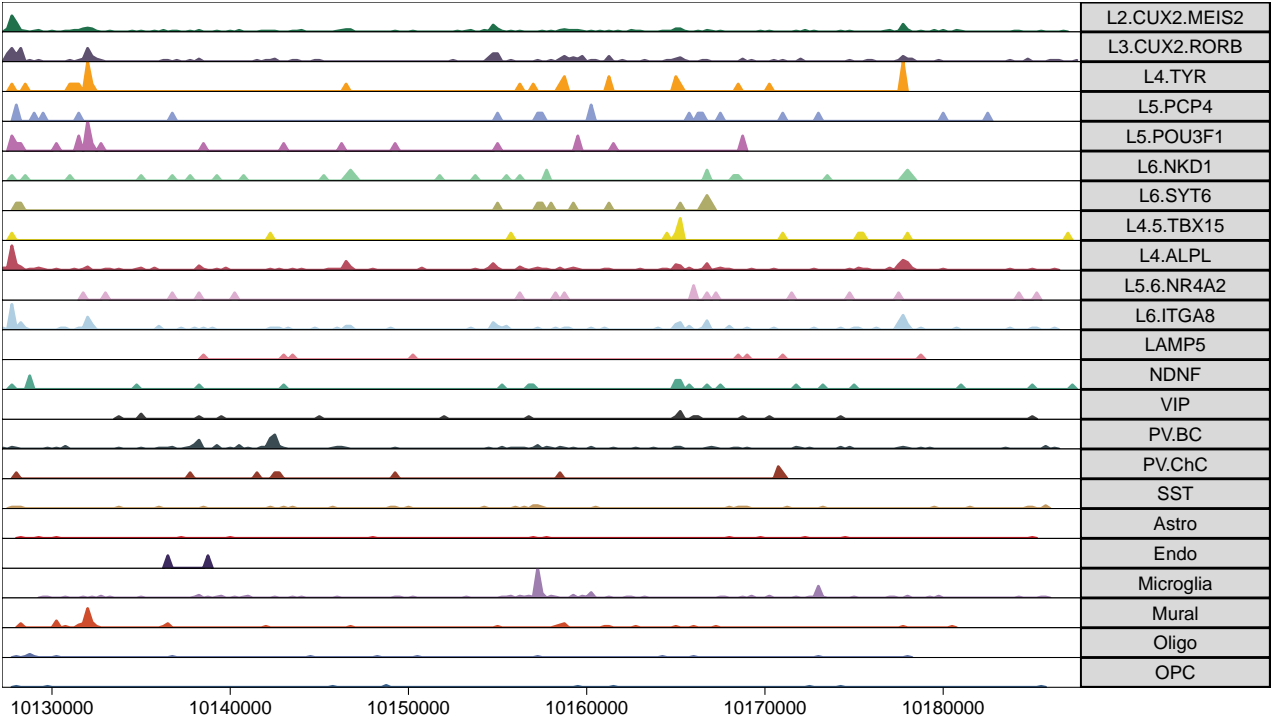


LoopTrack

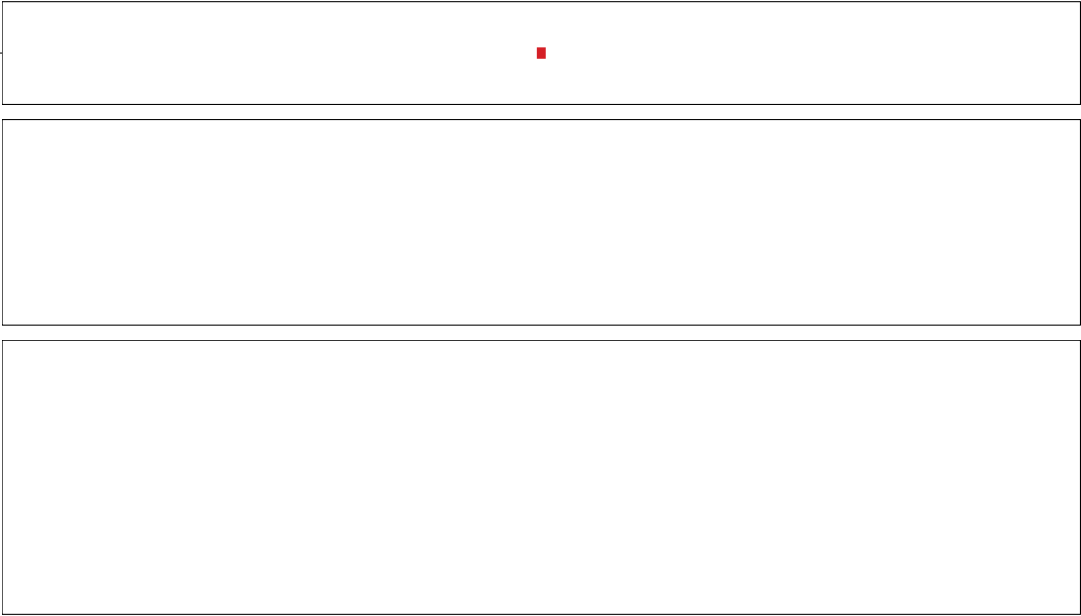


Genes

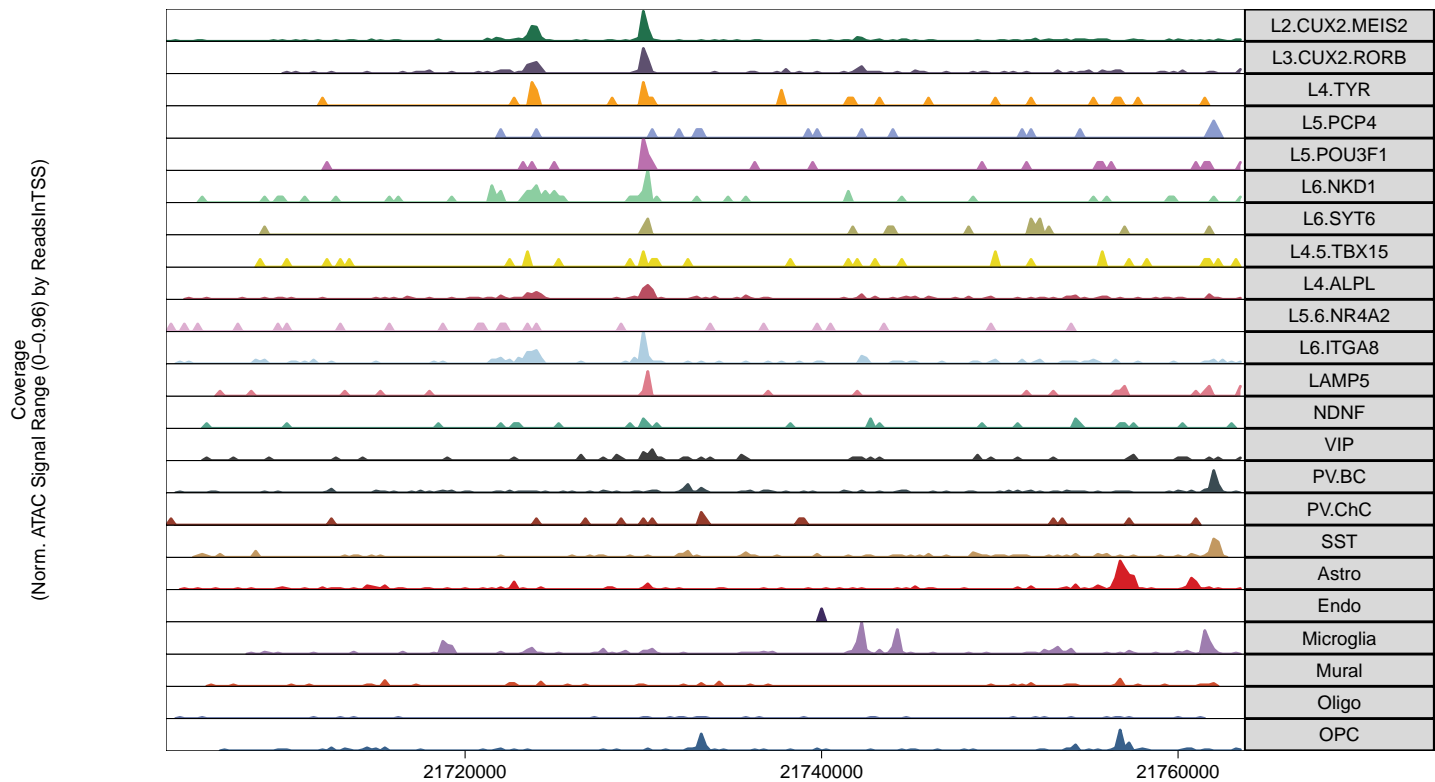
Coverage
(Norm. ATAC Signal Range (0-0.93) by Reads in TSS)



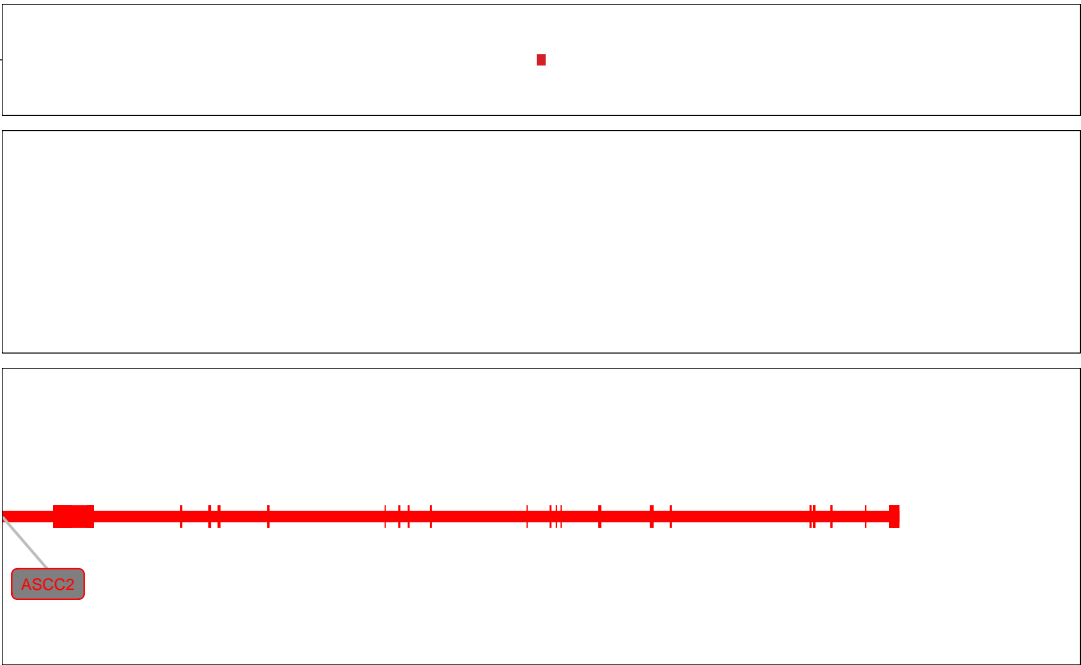
L5.PCP4



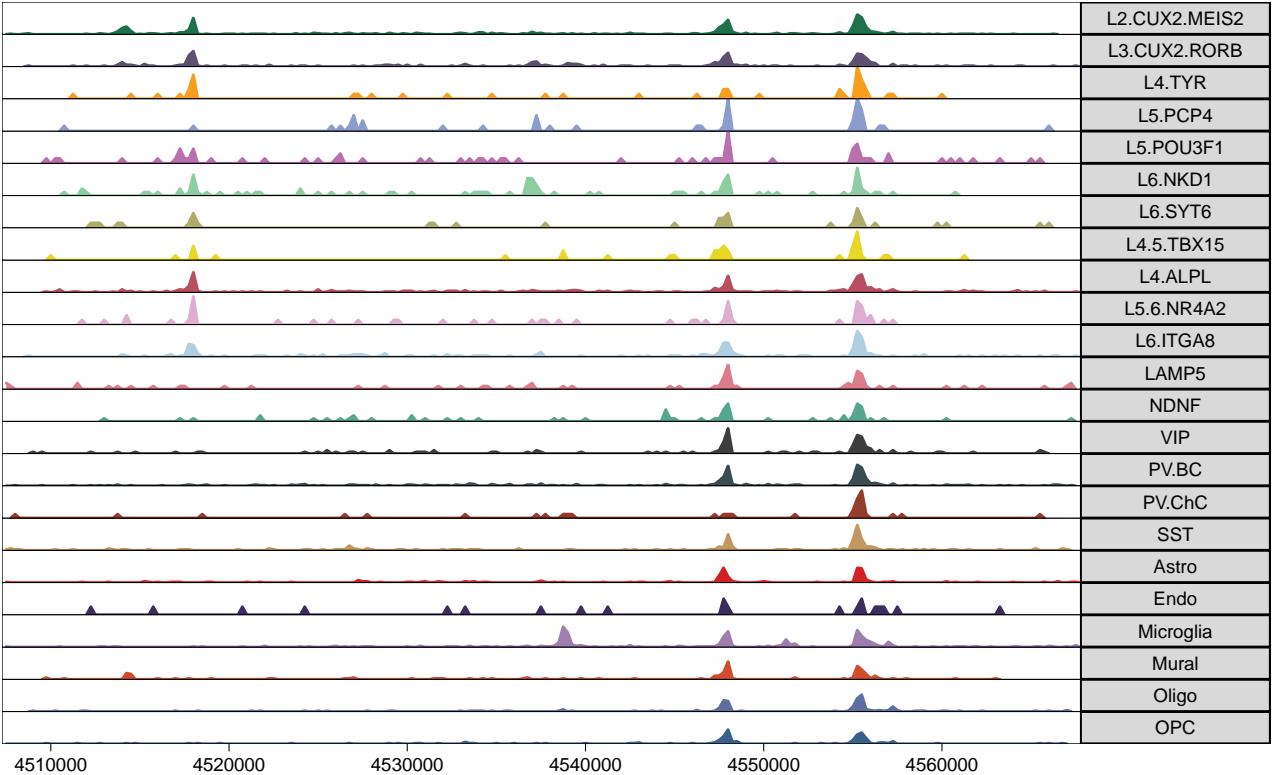
chr10:21703224–21763725 Marker Peak Tracks for: L5.PCP4



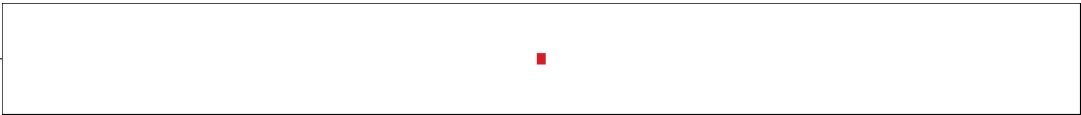
L5.PCP4



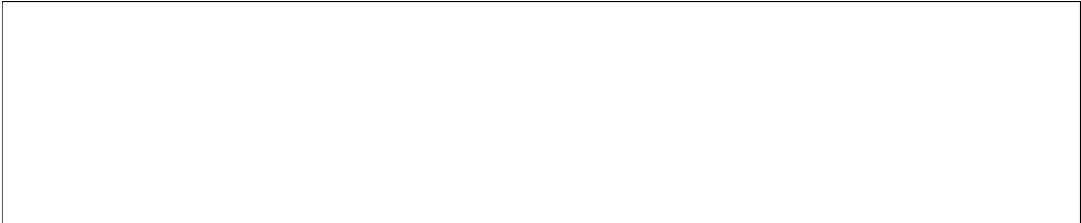
Coverage
(Norm. ATAC Signal Range 0–1.57) by ReadsInTSS)



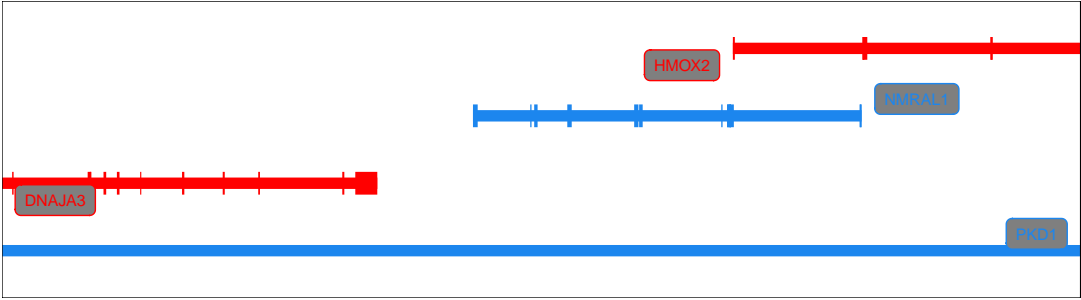
L5.PCP4



Peaks

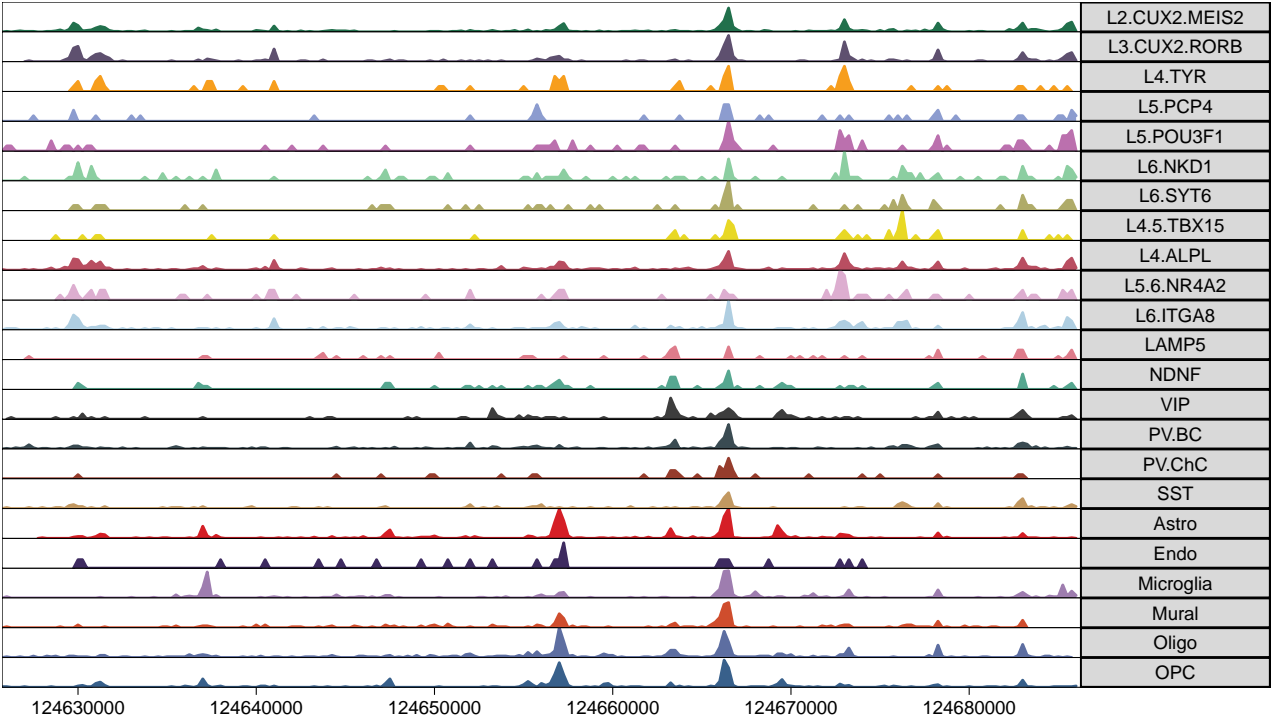


LoopTrack

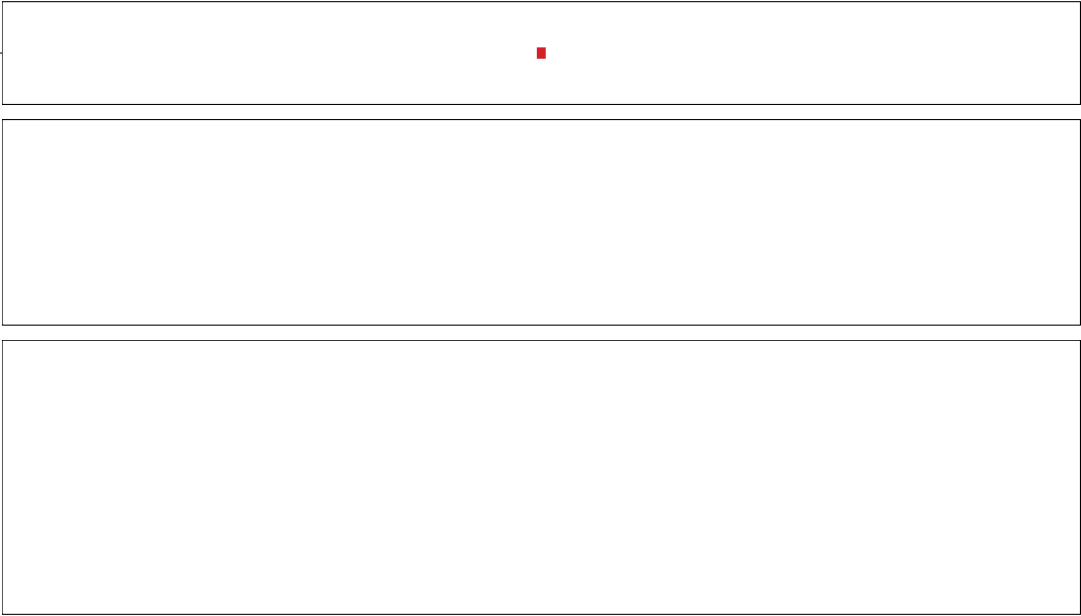


Genes

Coverage
(Norm. ATAC Signal Range (0–1.38) by Reads in TSS)



L5.PCP4

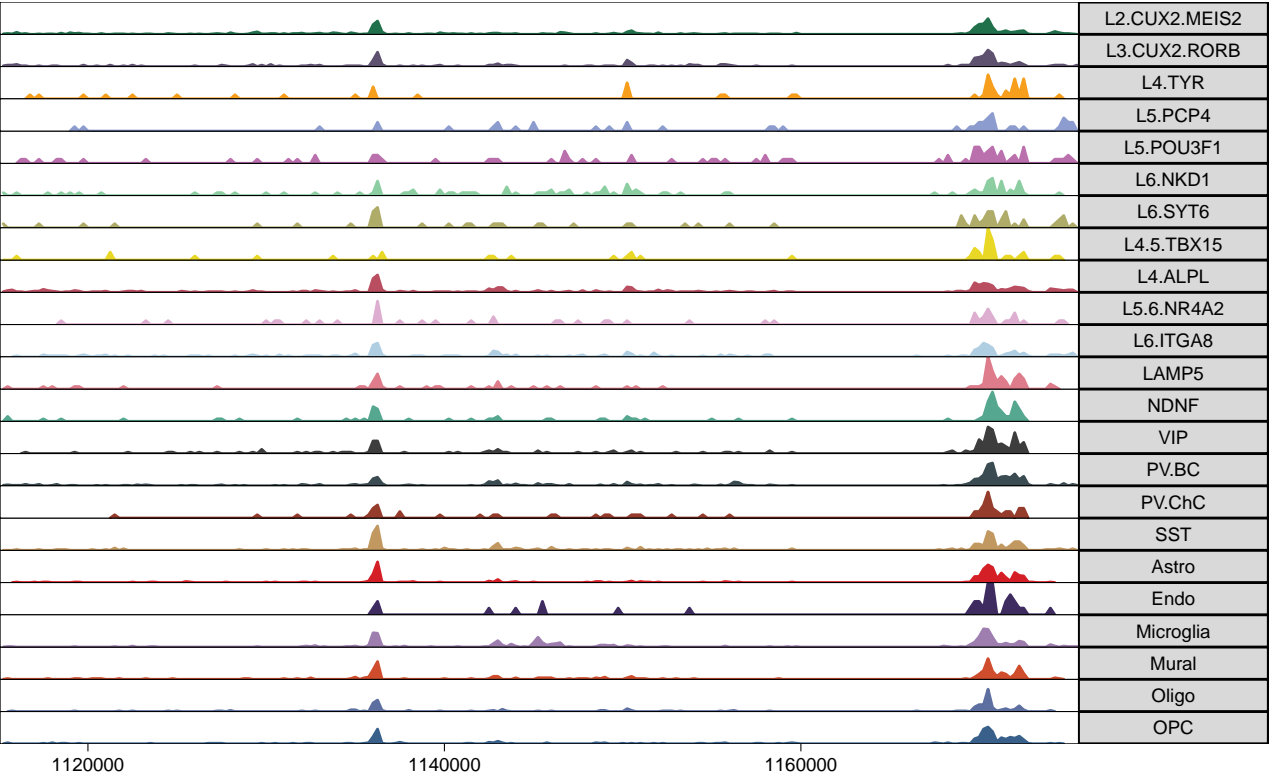


Peaks

LoopTrack

GeneTrack

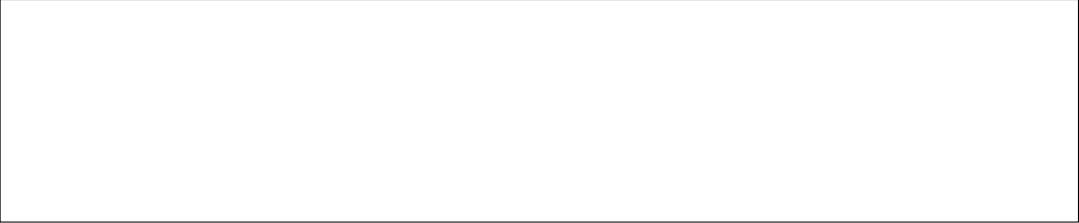
Coverage
(Norm. ATAC Signal Range 0-1.88) by ReadsInTSS)



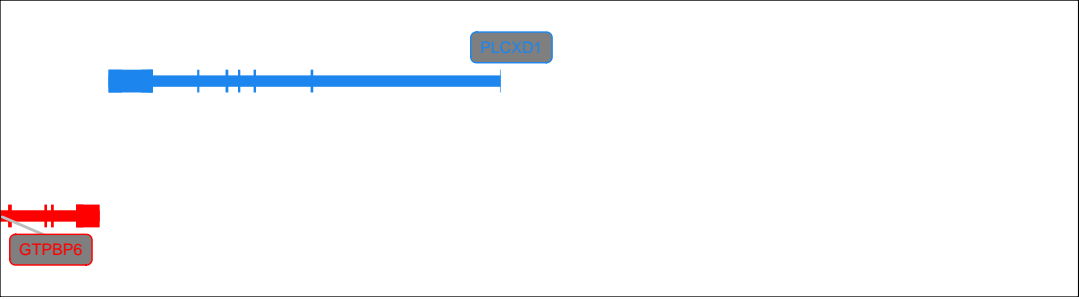
L5.PCP4



Peaks

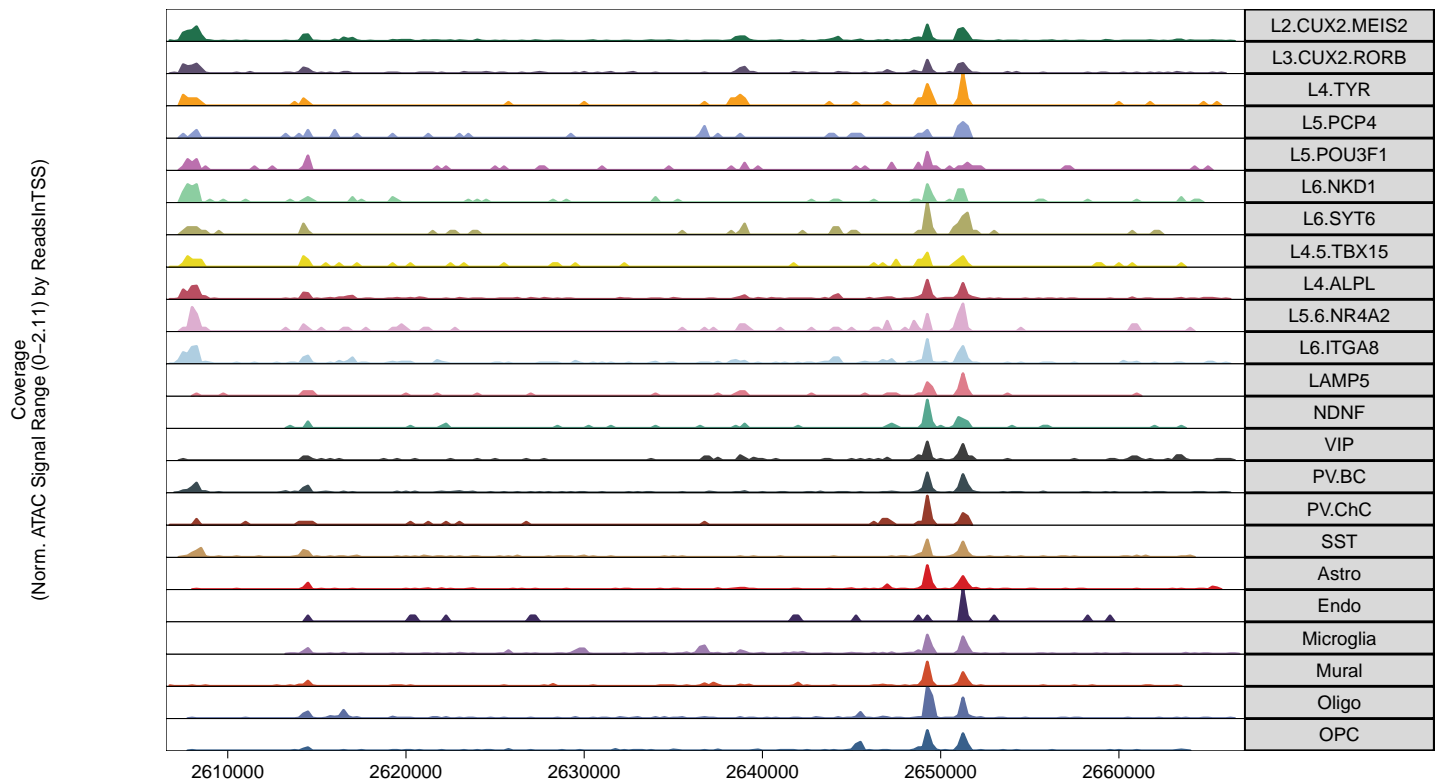


LoopTrack

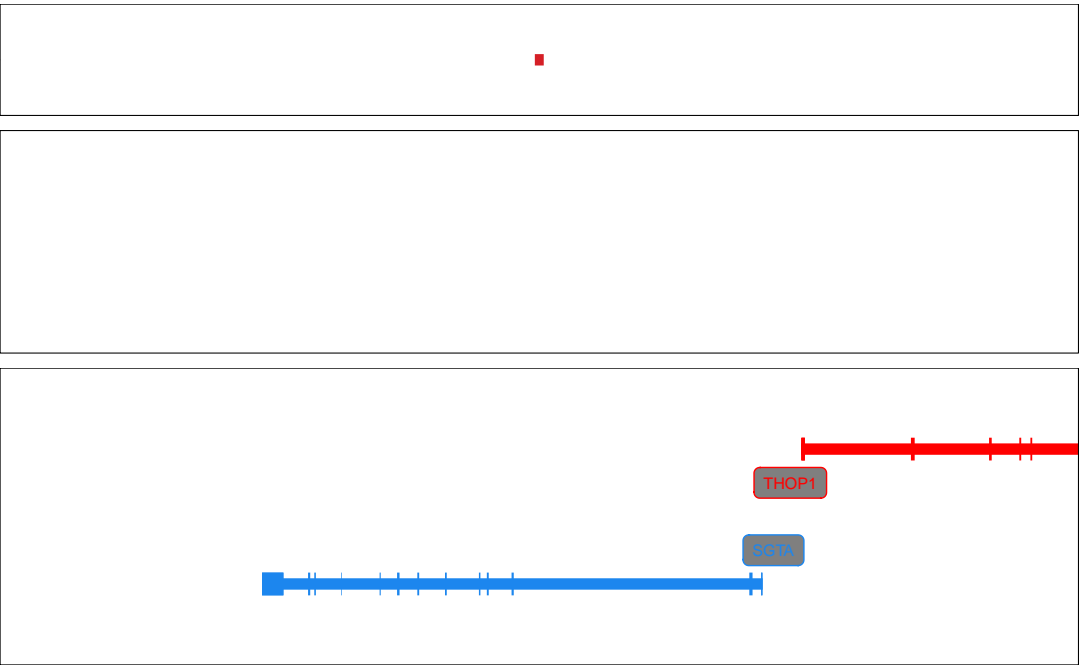


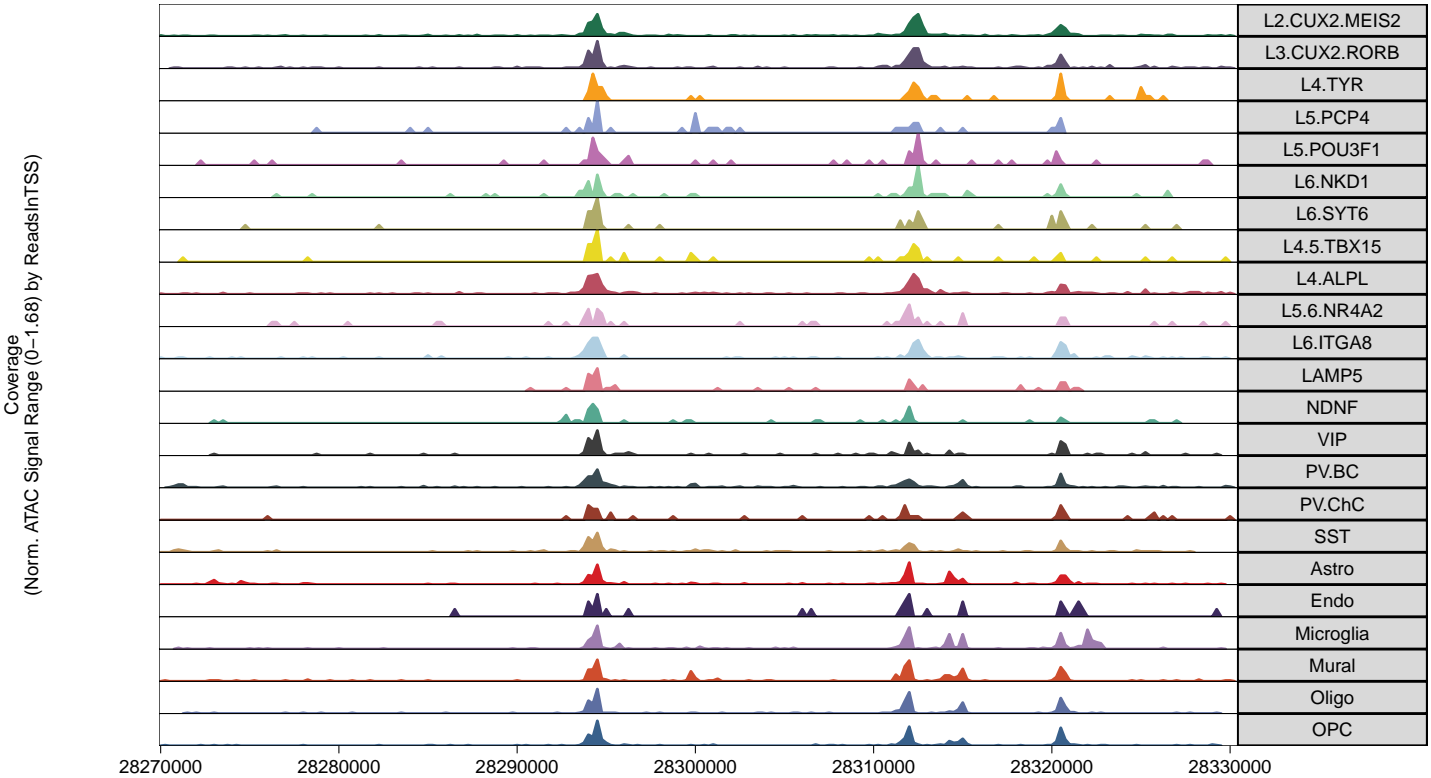
Genes

chr19:2606542-2667043 Marker Peak Tracks for: L5.PCP4



L5.PCP4

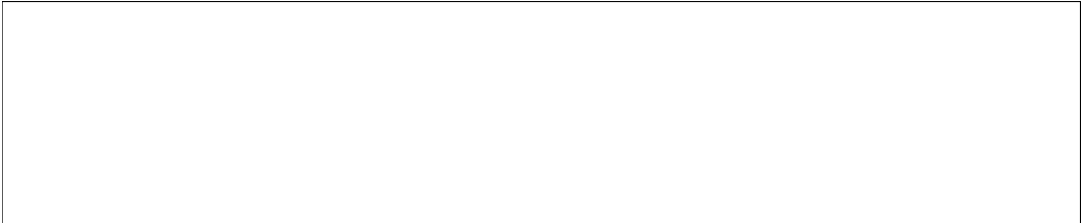




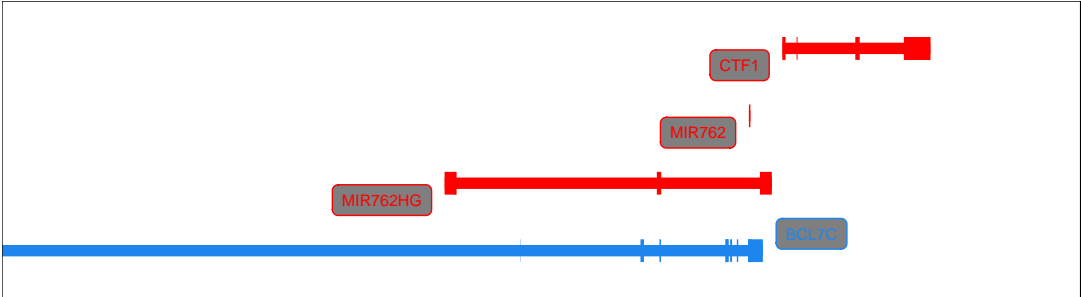
L5.PCP4



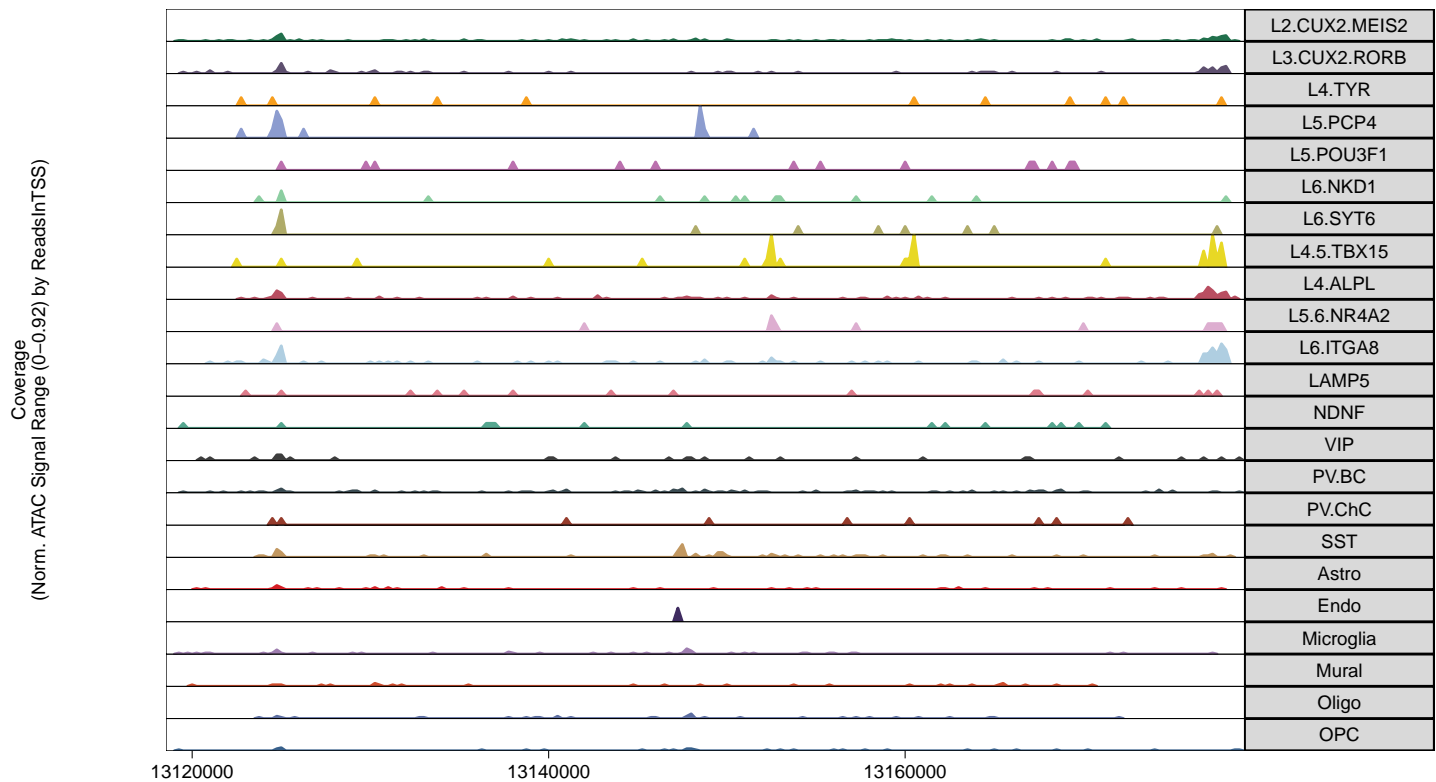
Peaks



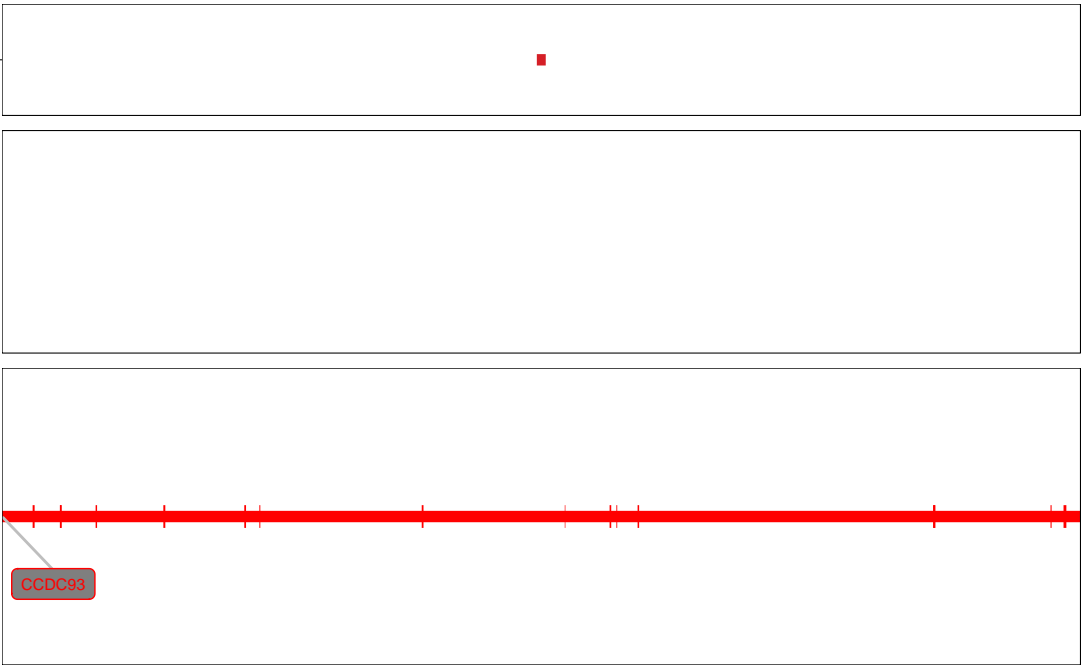
LoopTrack



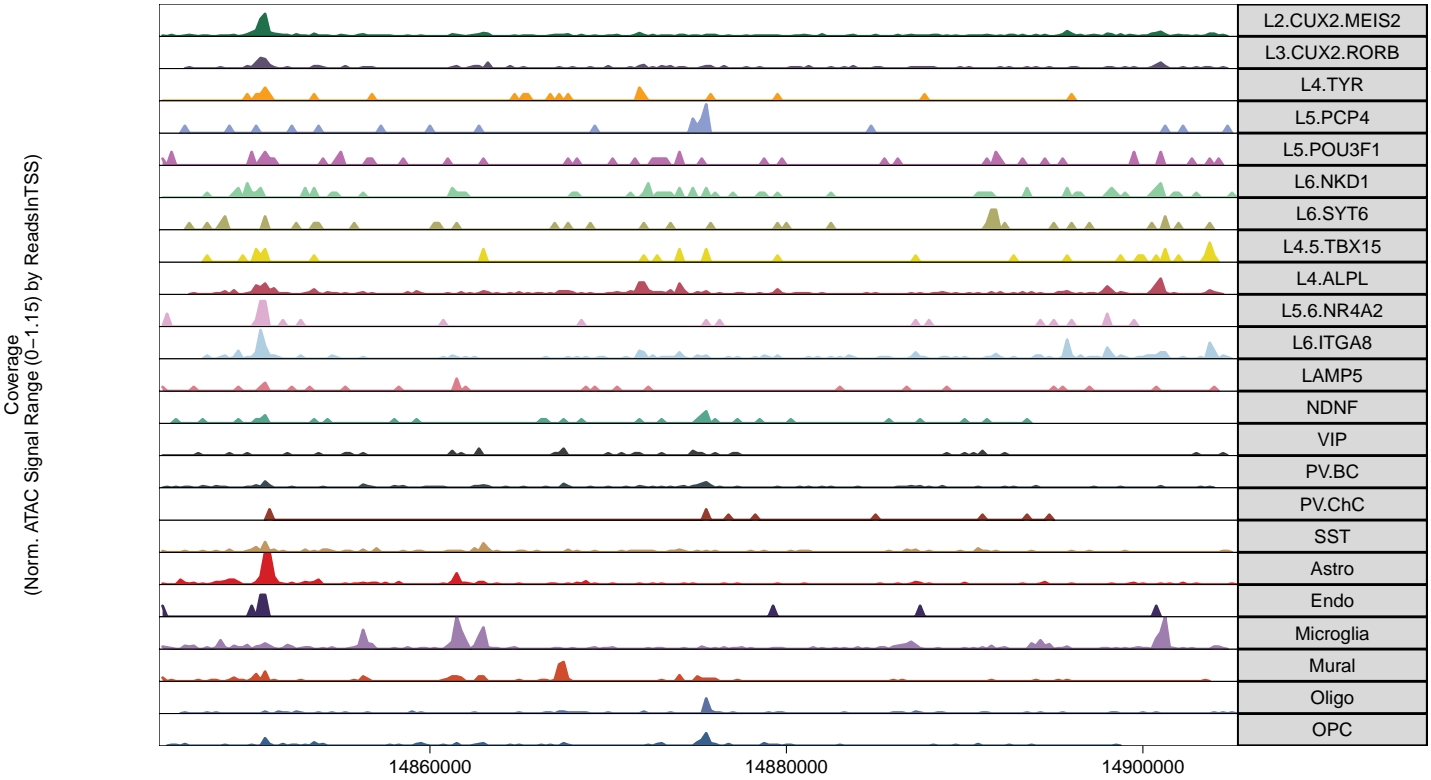
Genes



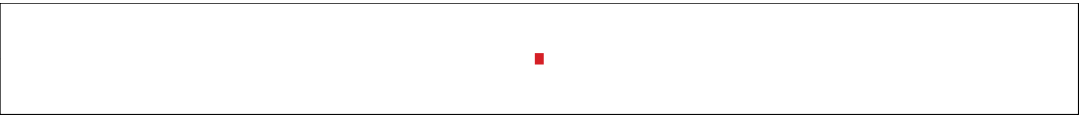
L5.PCP4



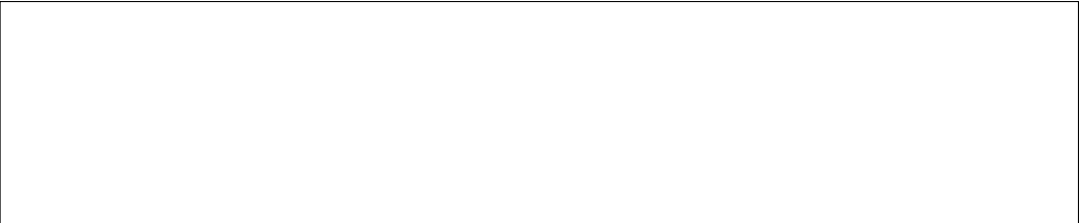
chr9:14844804–14905305 Marker Peak Tracks for: L5.PCP4



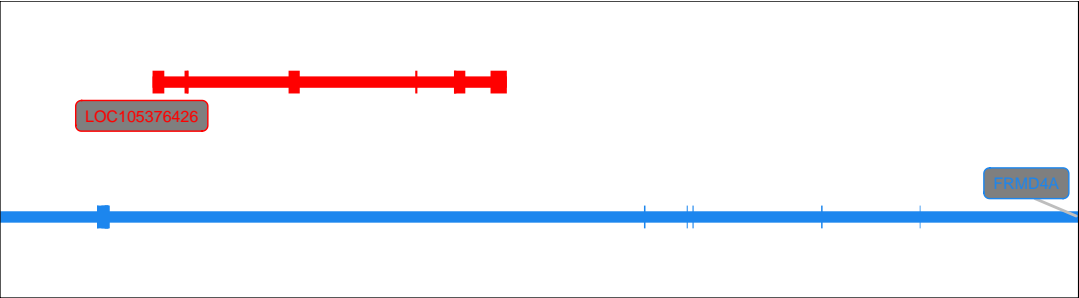
L5.PCP4



Peaks

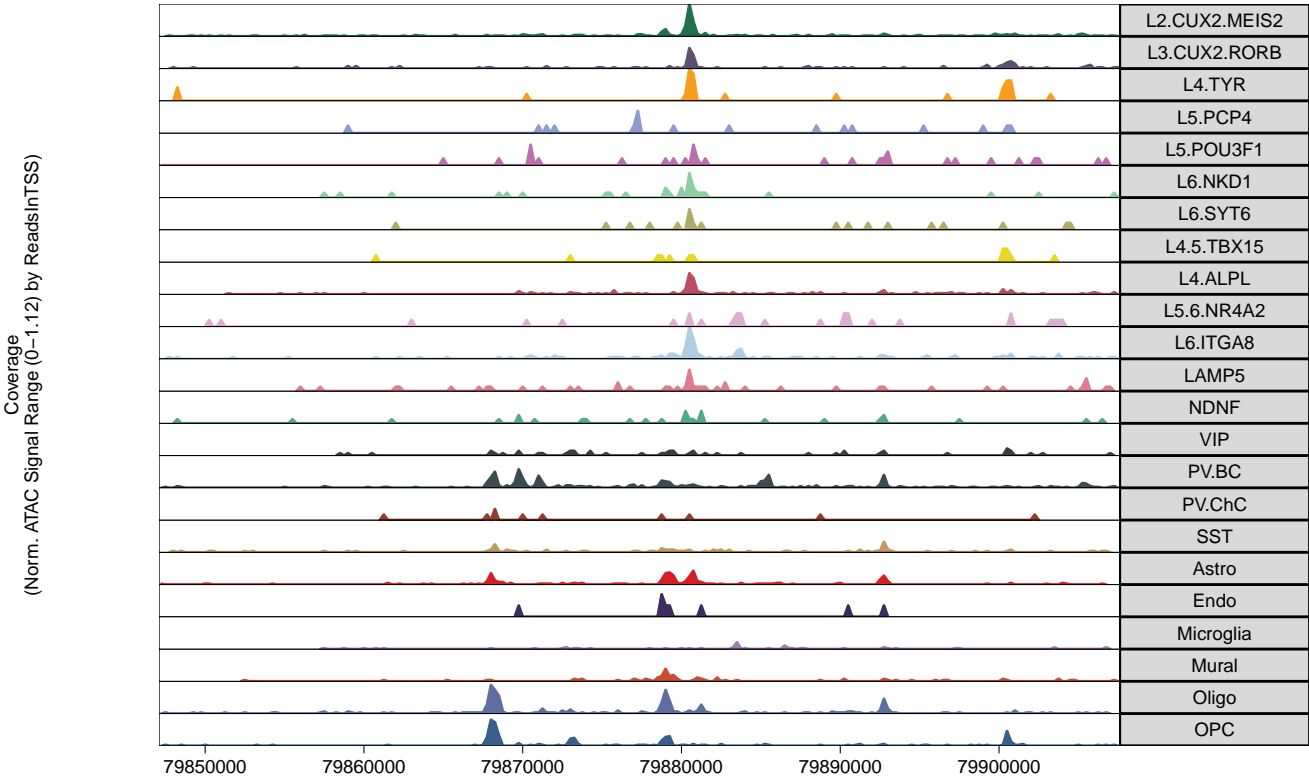


LoopTrack

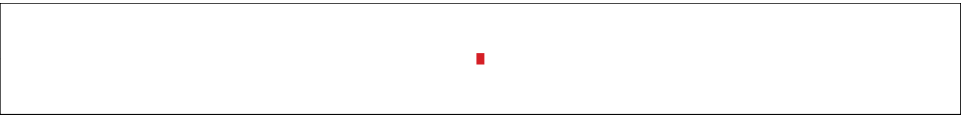


Genes

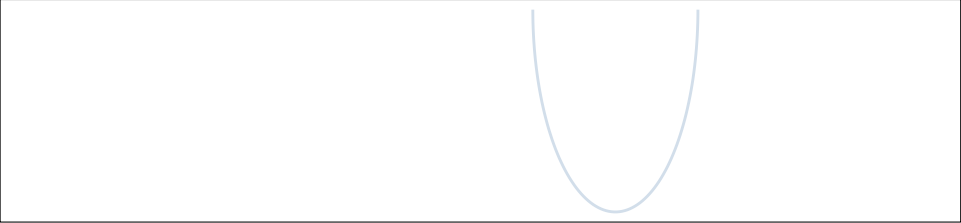
chr13:79847099–79907600 Marker Peak Tracks for: L5.PCP4



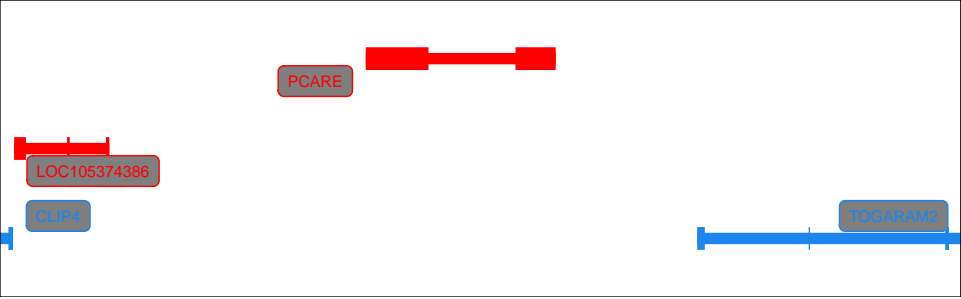
L5.PCP4



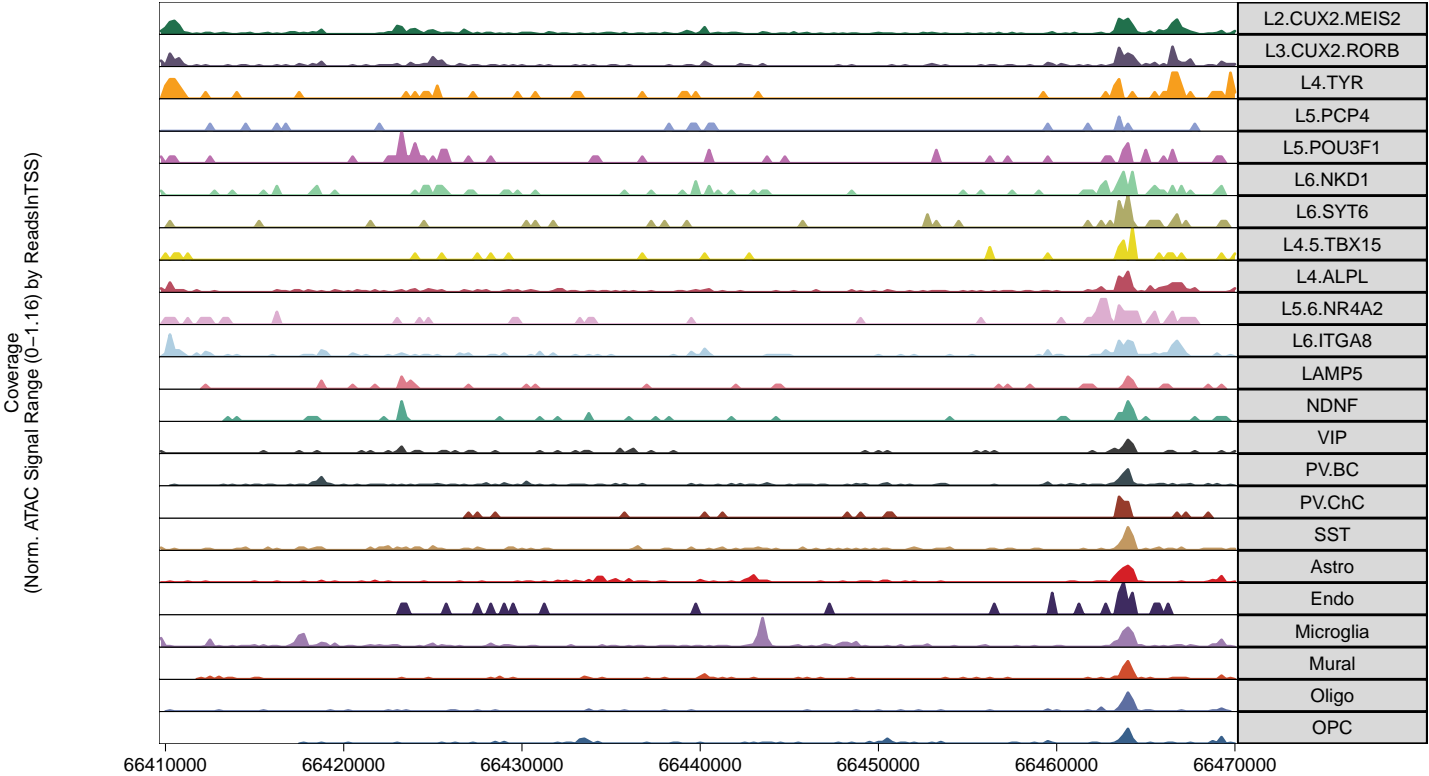
Peaks



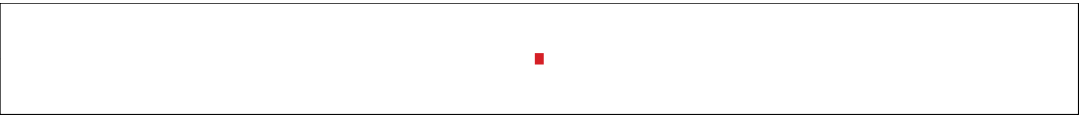
Peak2GeneLinks



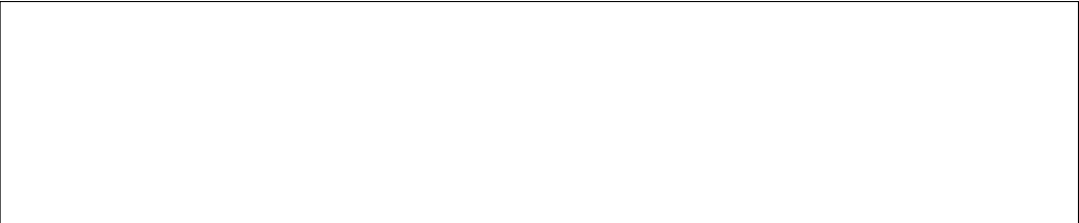
Genes



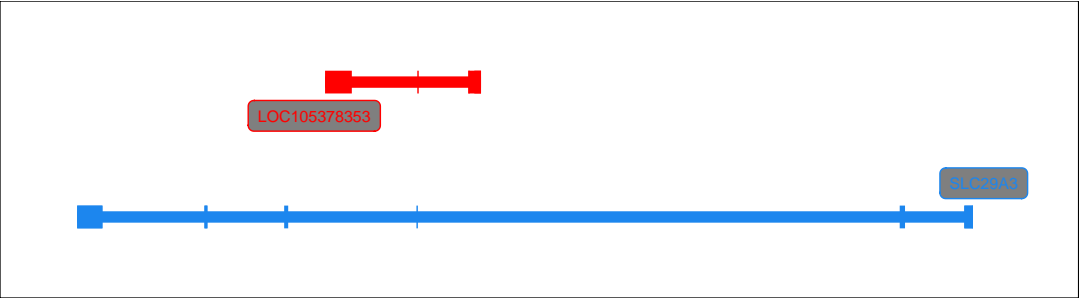
L5.PCP4



Peaks

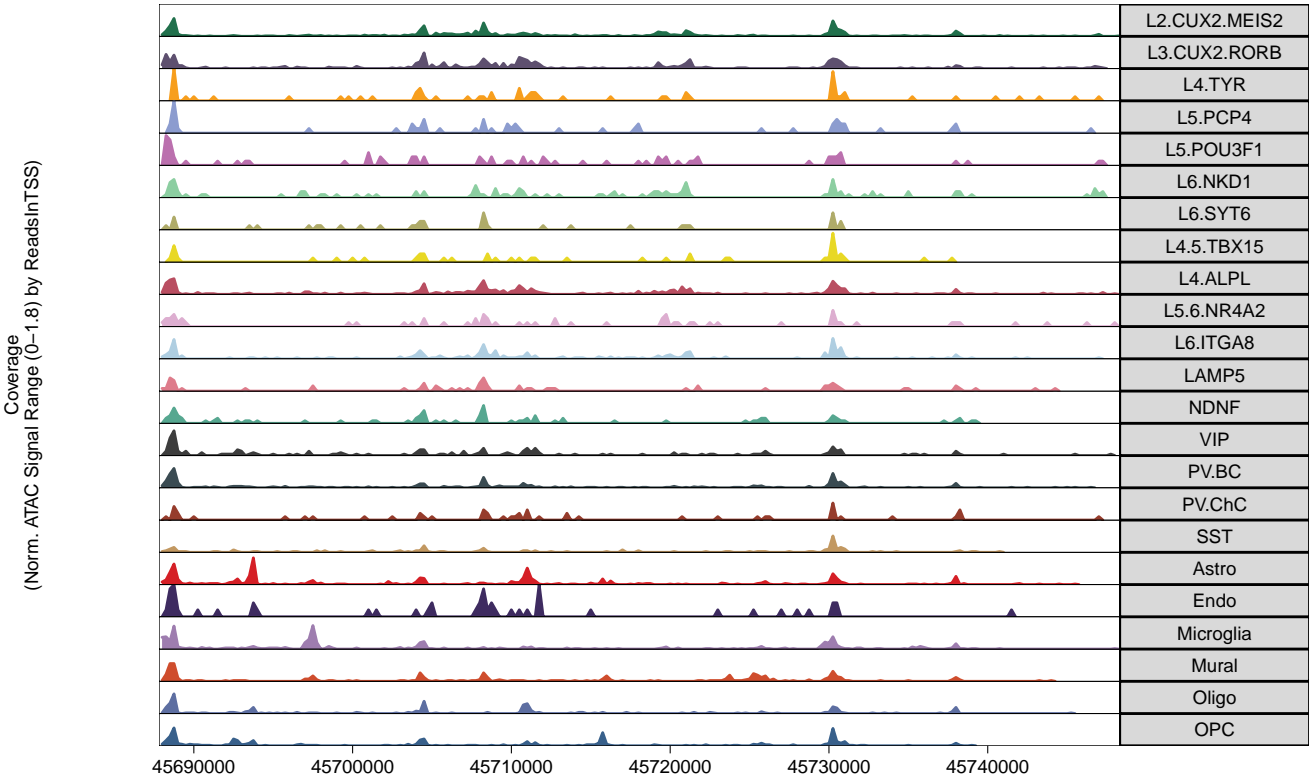


LoopTrack



Genes

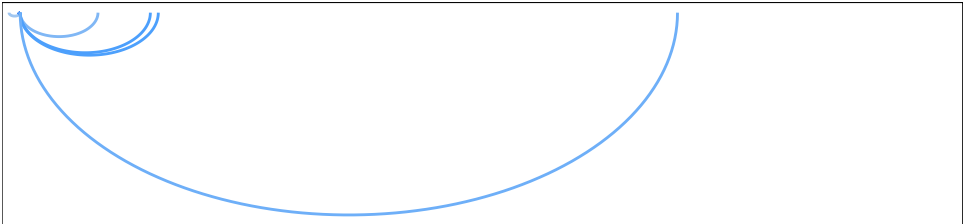
chr9:45687812-45748313 Marker Peak Tracks for: L5.PCP4



L5.PCP4

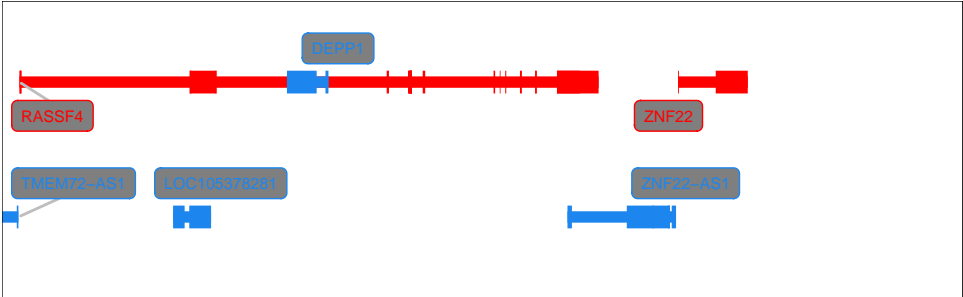


Peaks



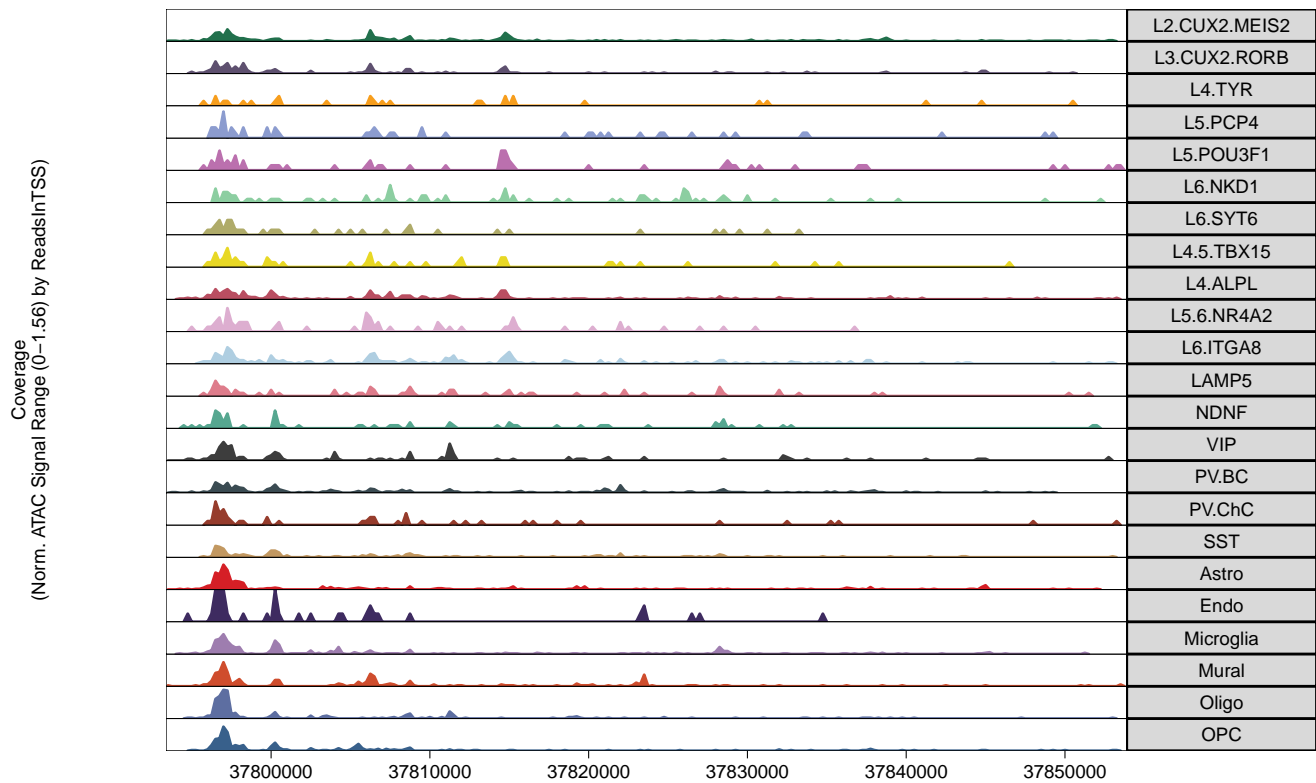
Peak2GeneLinks

value



Genes

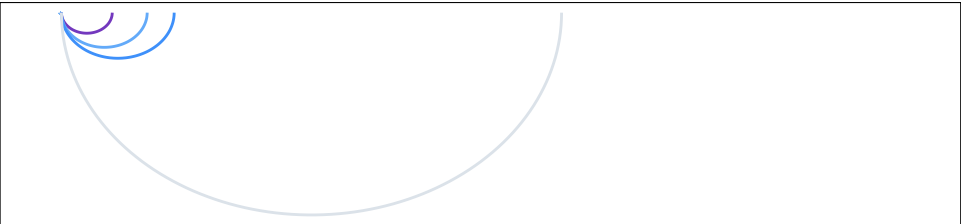
chr5:37793387-37853888 Marker Peak Tracks for: L5.PCP4



L5.PCP4

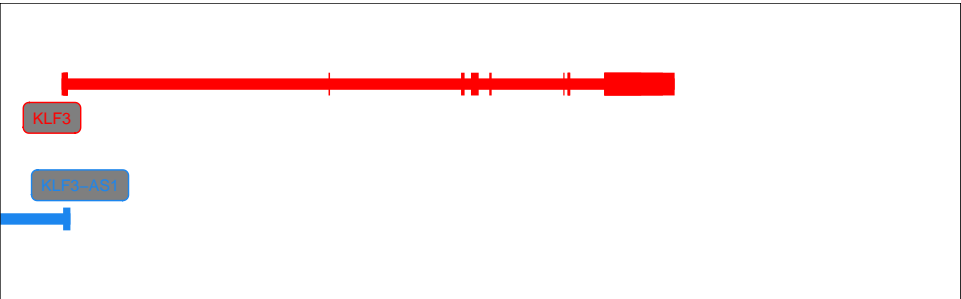


Peaks



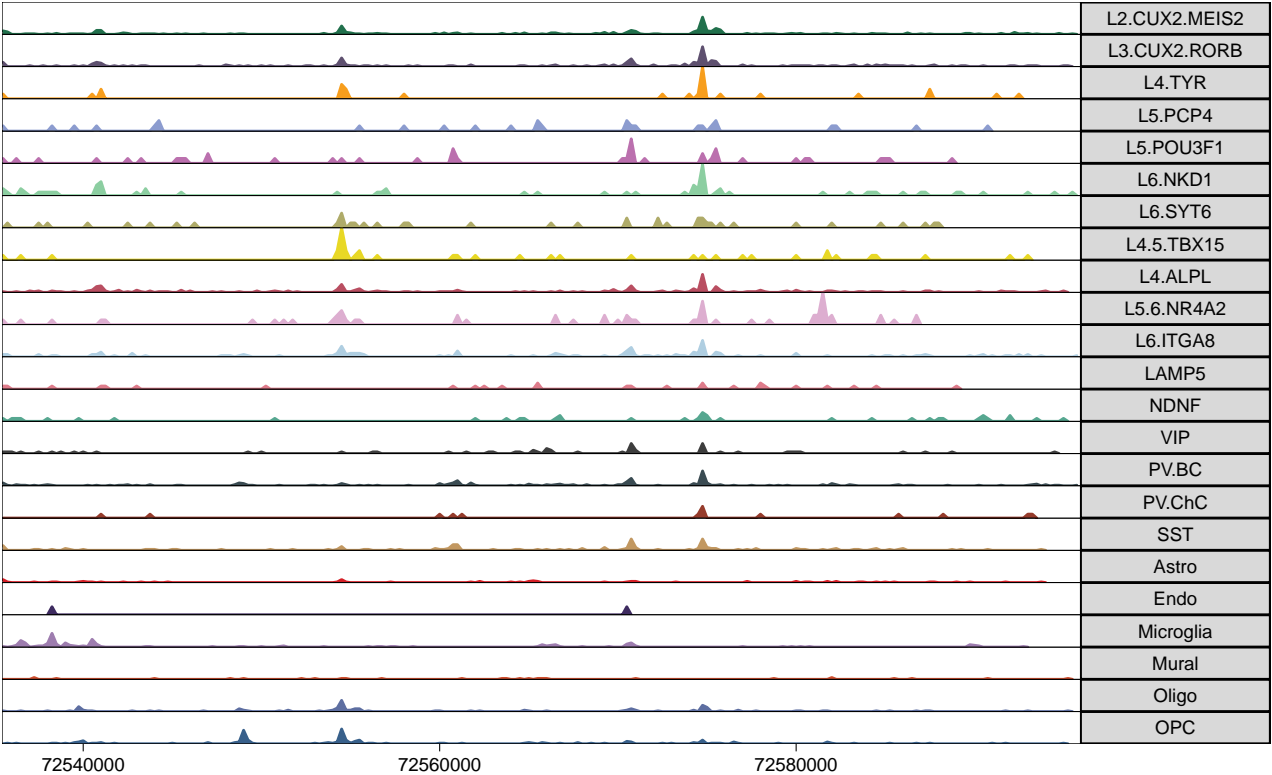
Peak2GeneLinks

value



Genes

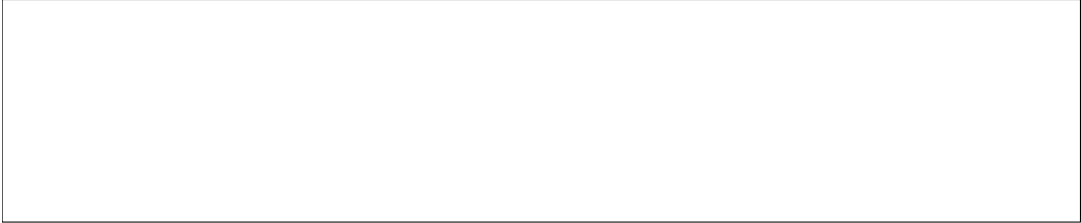
Coverage
(Norm. ATAC Signal Range 0–1.57) by ReadsInTSS)



L5.PCP4



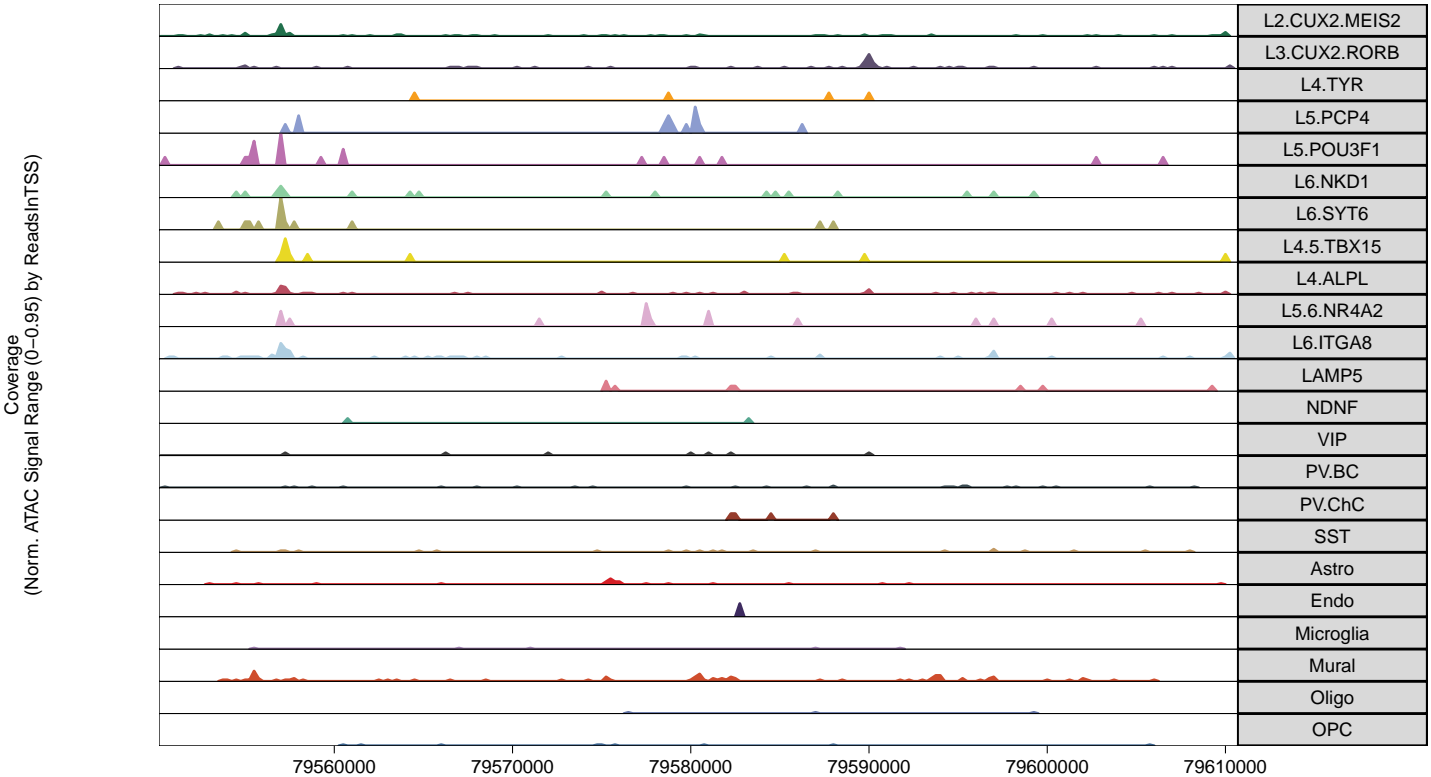
Peaks



LoopTrack



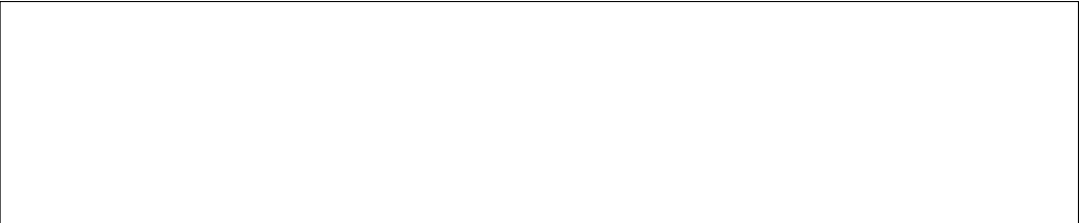
Genes



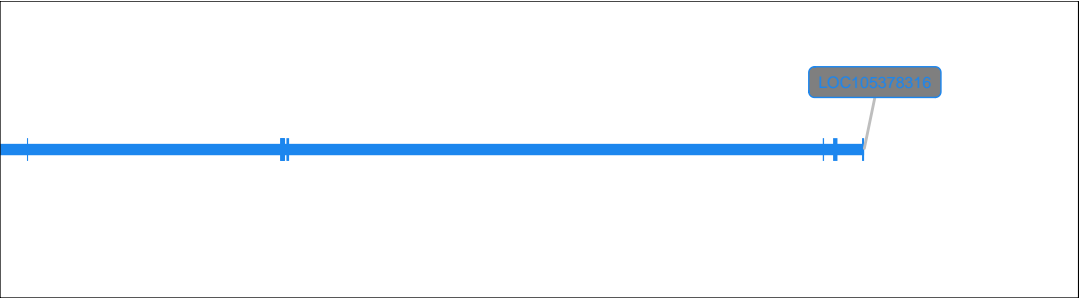
L5.PCP4



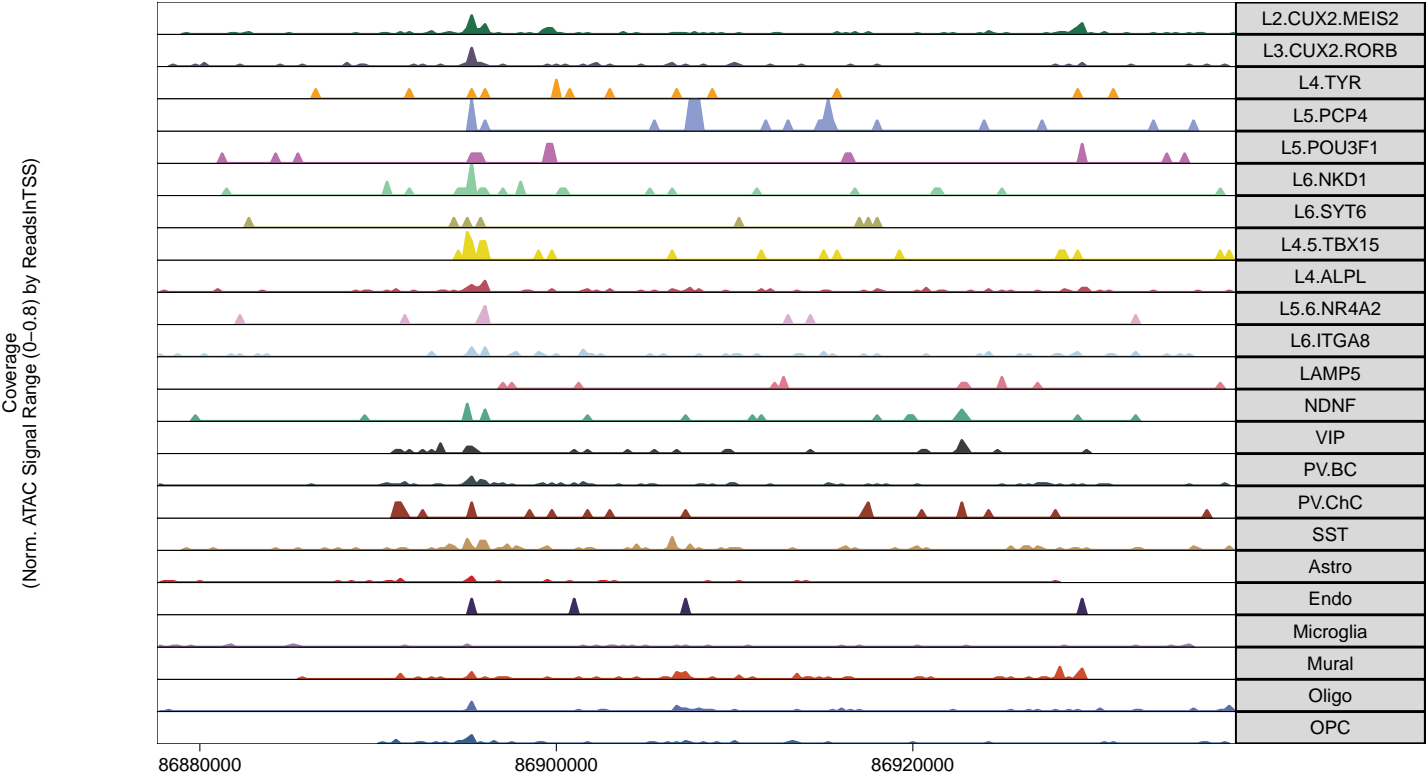
Peaks



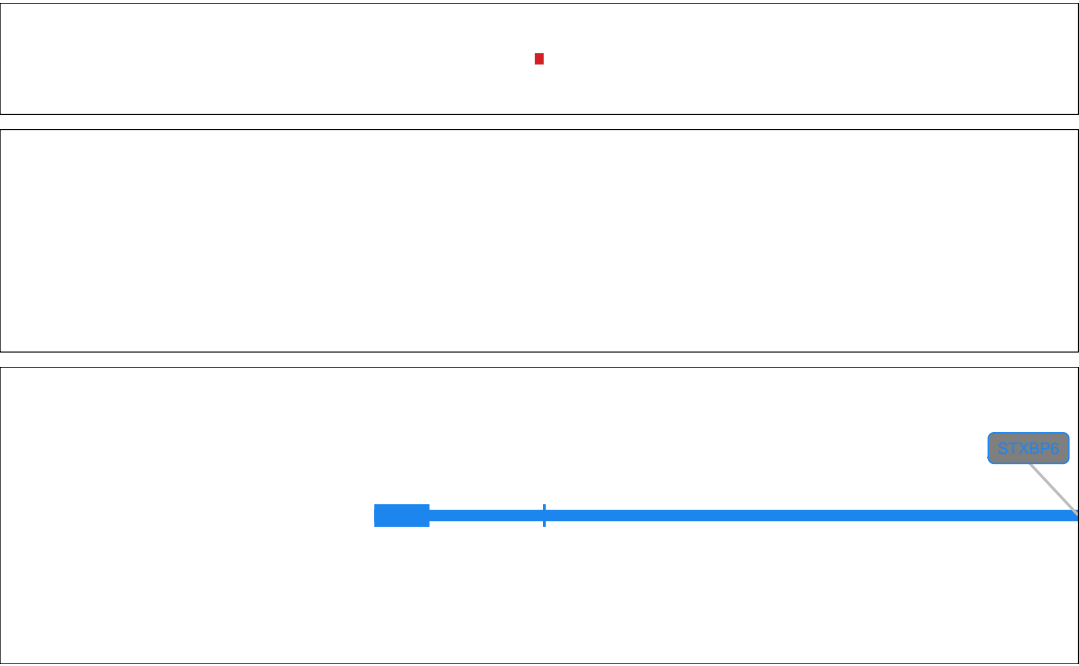
LoopTrack



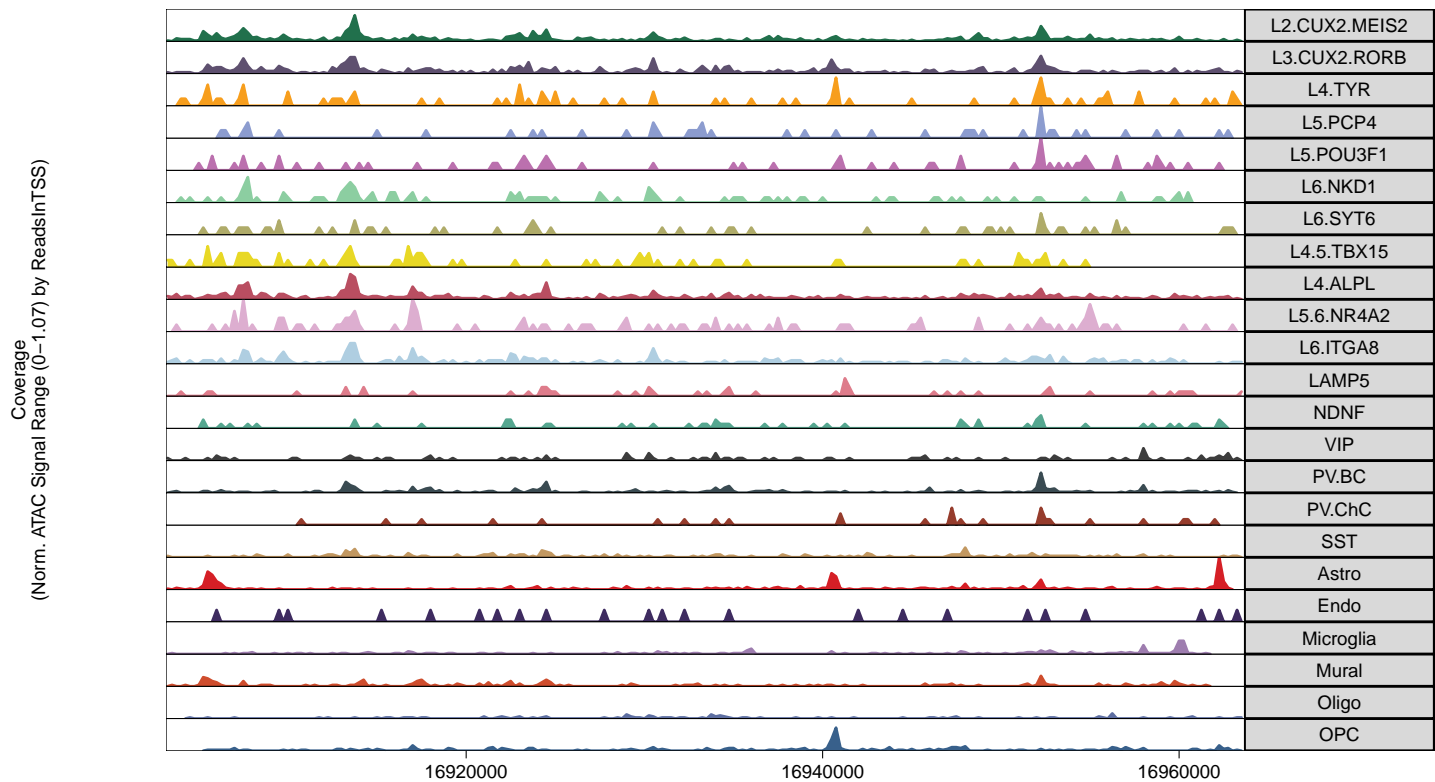
Genes



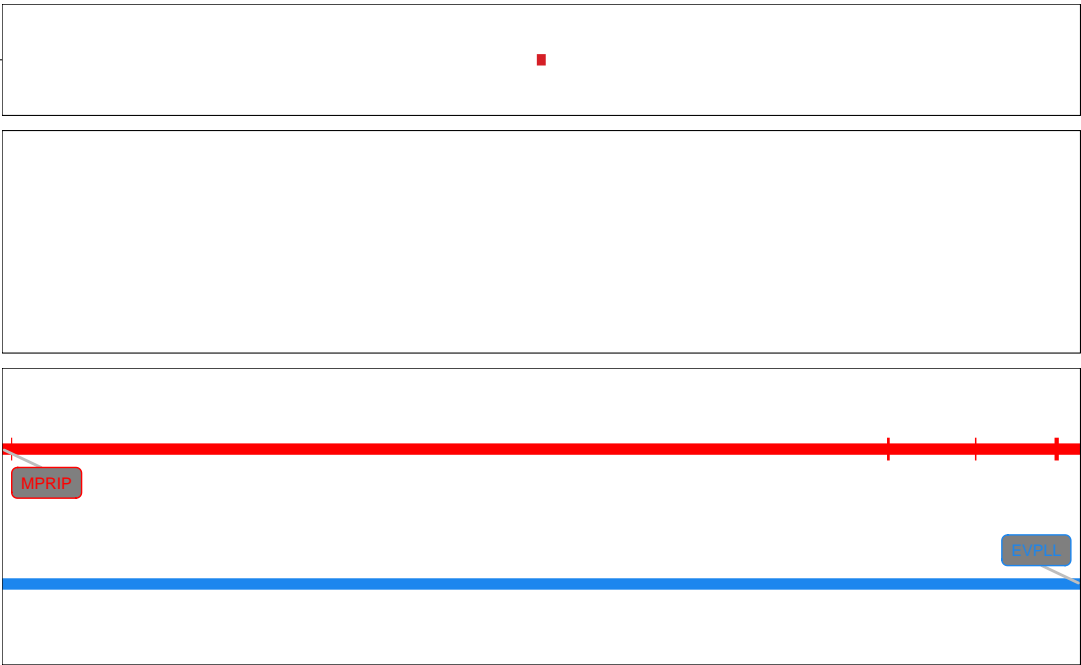
L5.PCP4



chr16:16903165–16963666 Marker Peak Tracks for: L5.PCP4



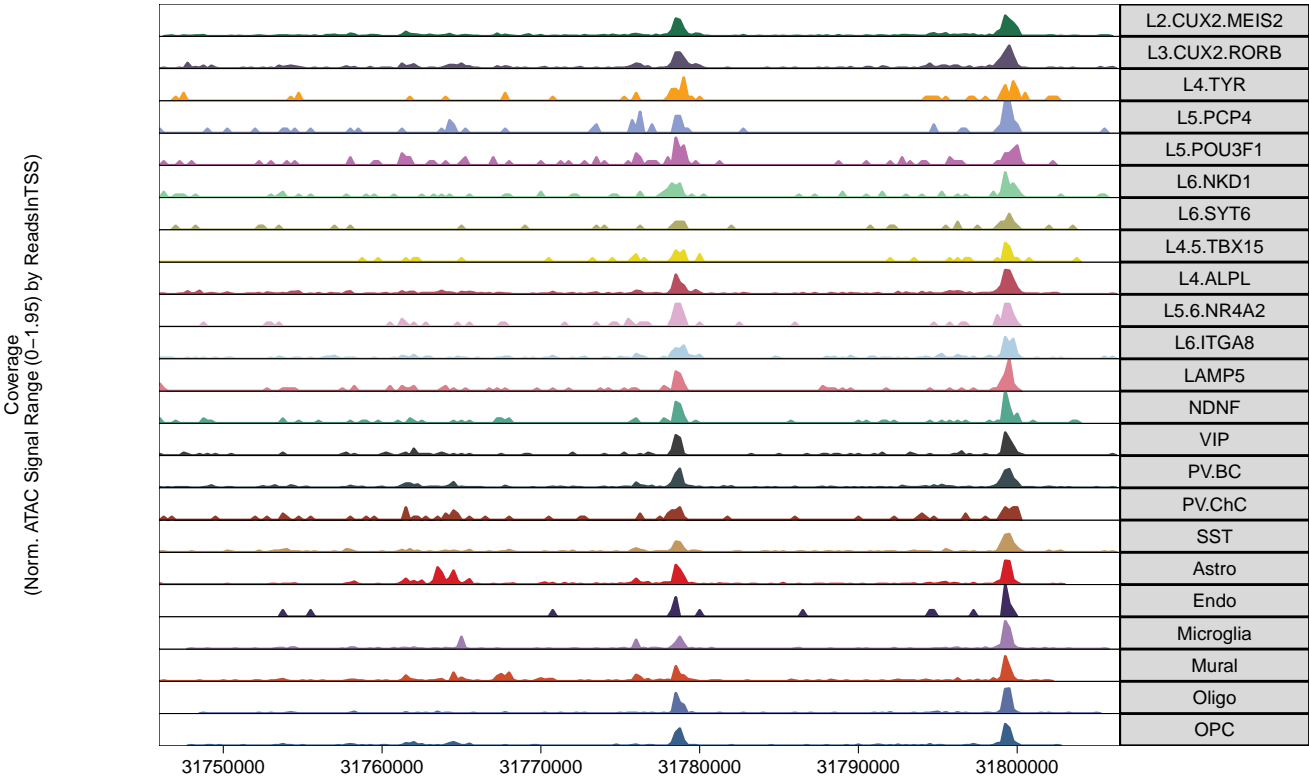
L5.PCP4



Peaks

LoopTrack

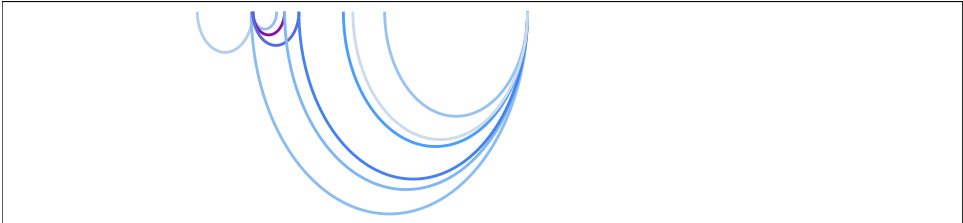
Genes



L5.PCP4

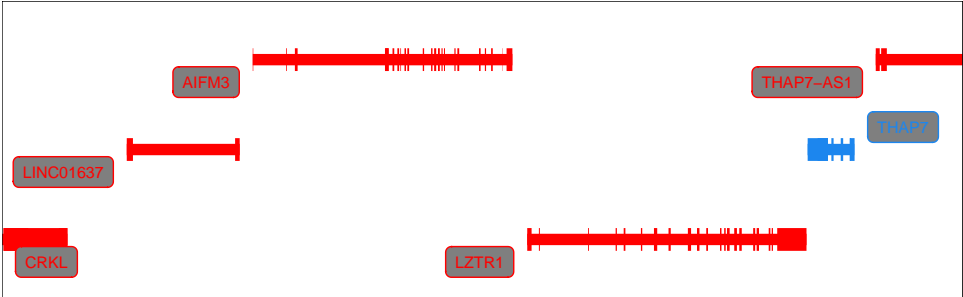


Peaks



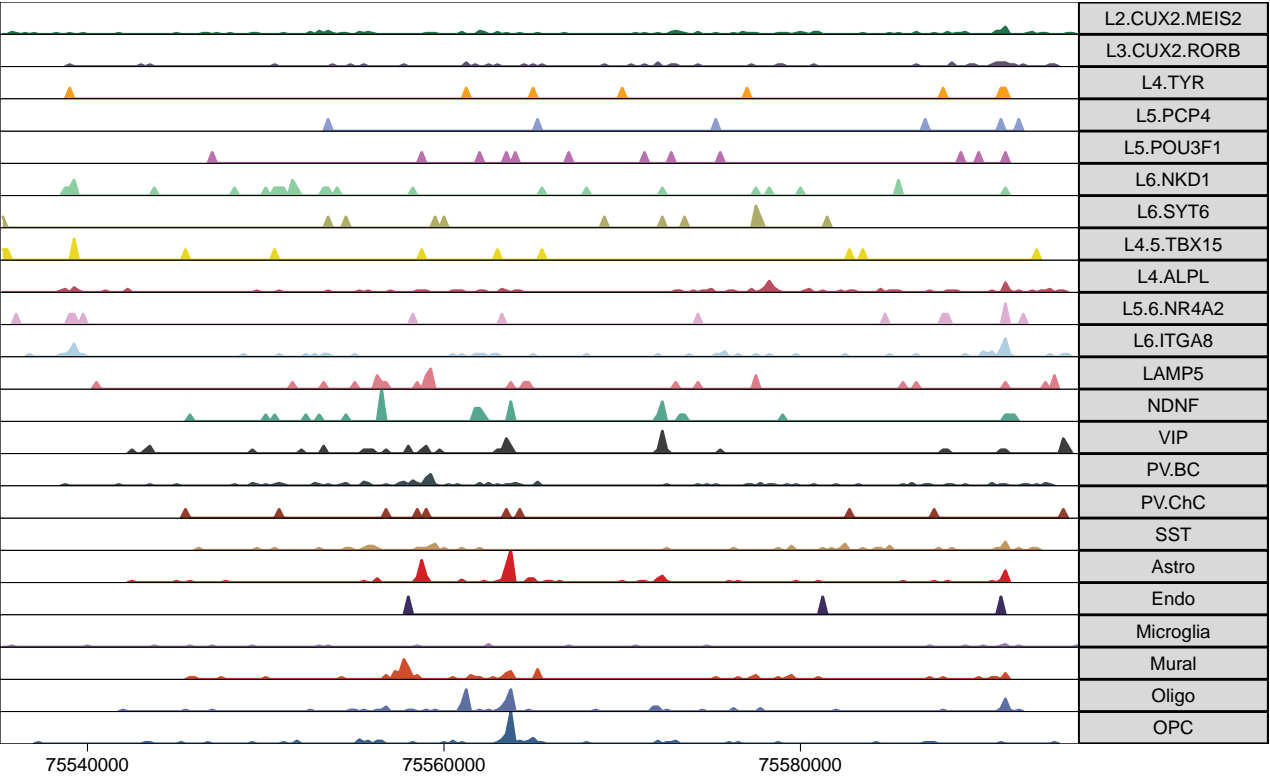
Peak2GeneLinks

value

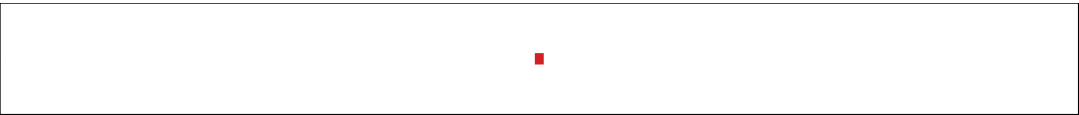


Genes

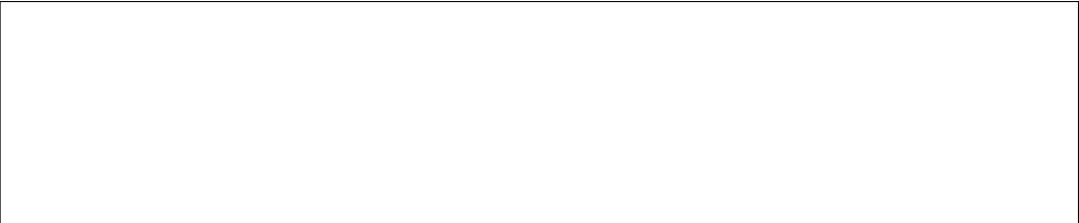
Coverage
(Norm. ATAC Signal Range (0-0.7) by Reads in TSS)



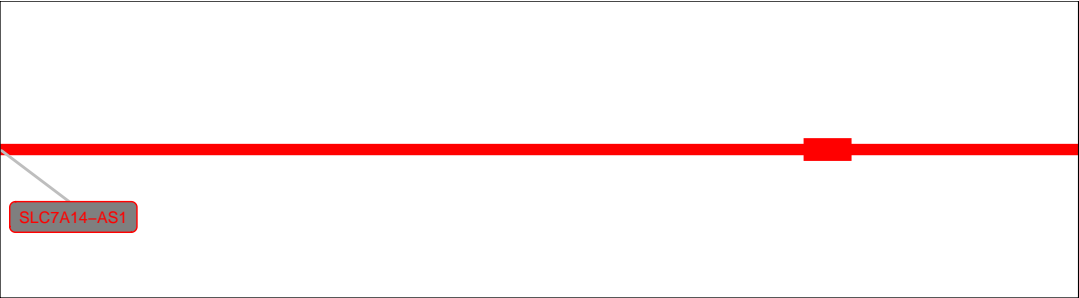
L5.PCP4



Peaks

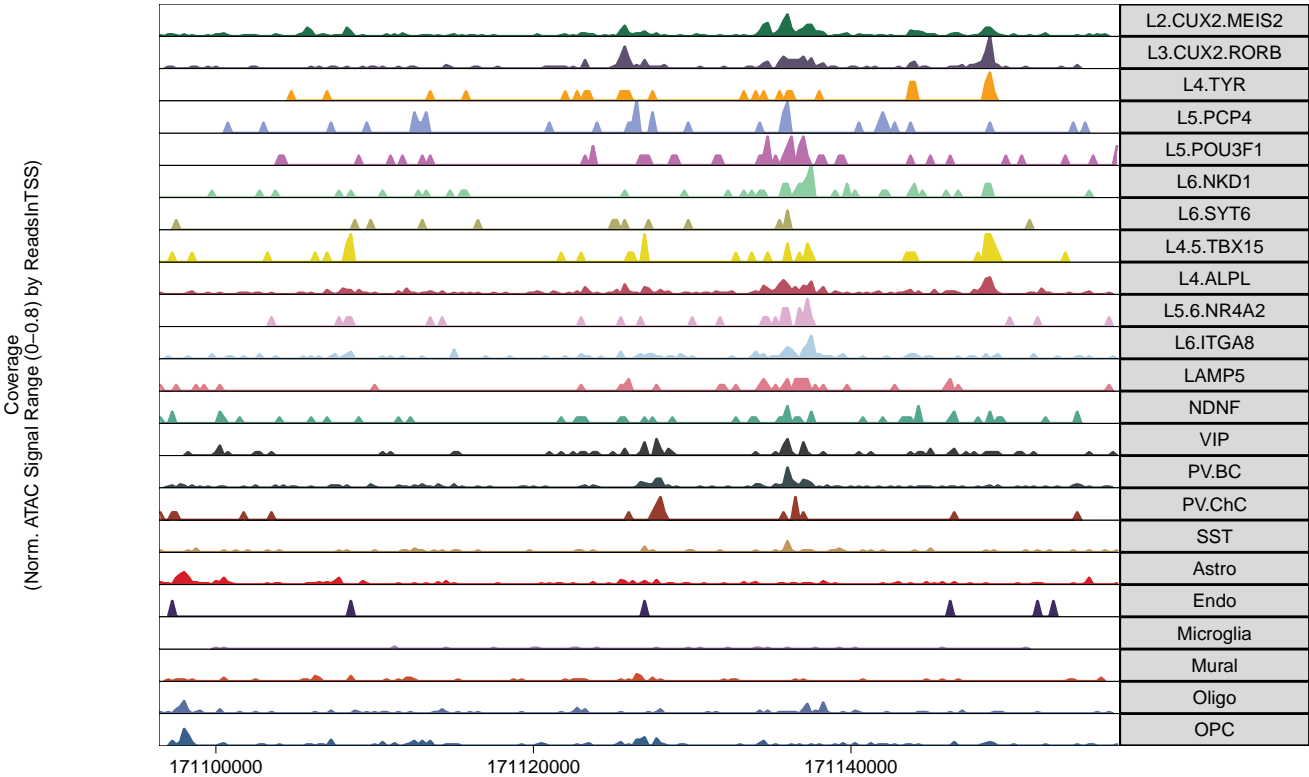


LoopTrack



Genes

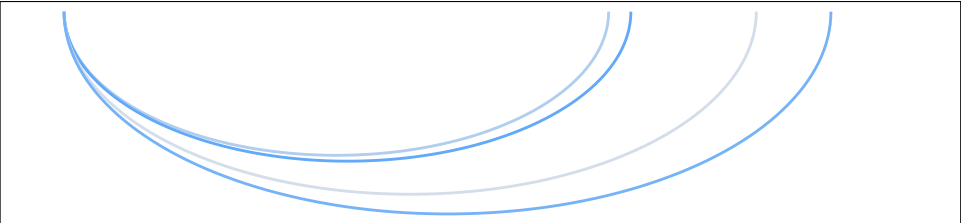
chr6:171096423-171156924 Marker Peak Tracks for: L5.PCP4



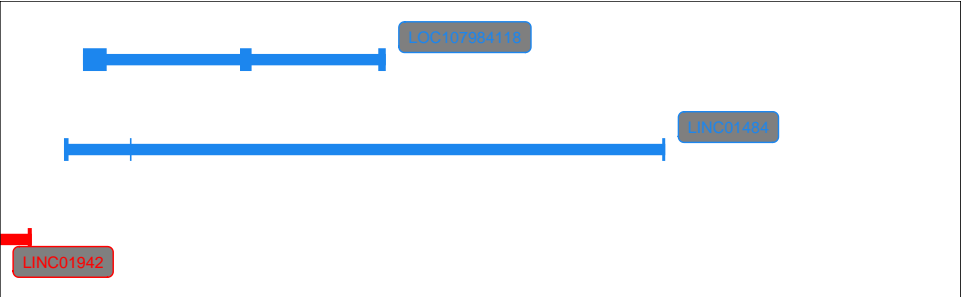
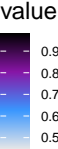
L5.PCP4



Peaks

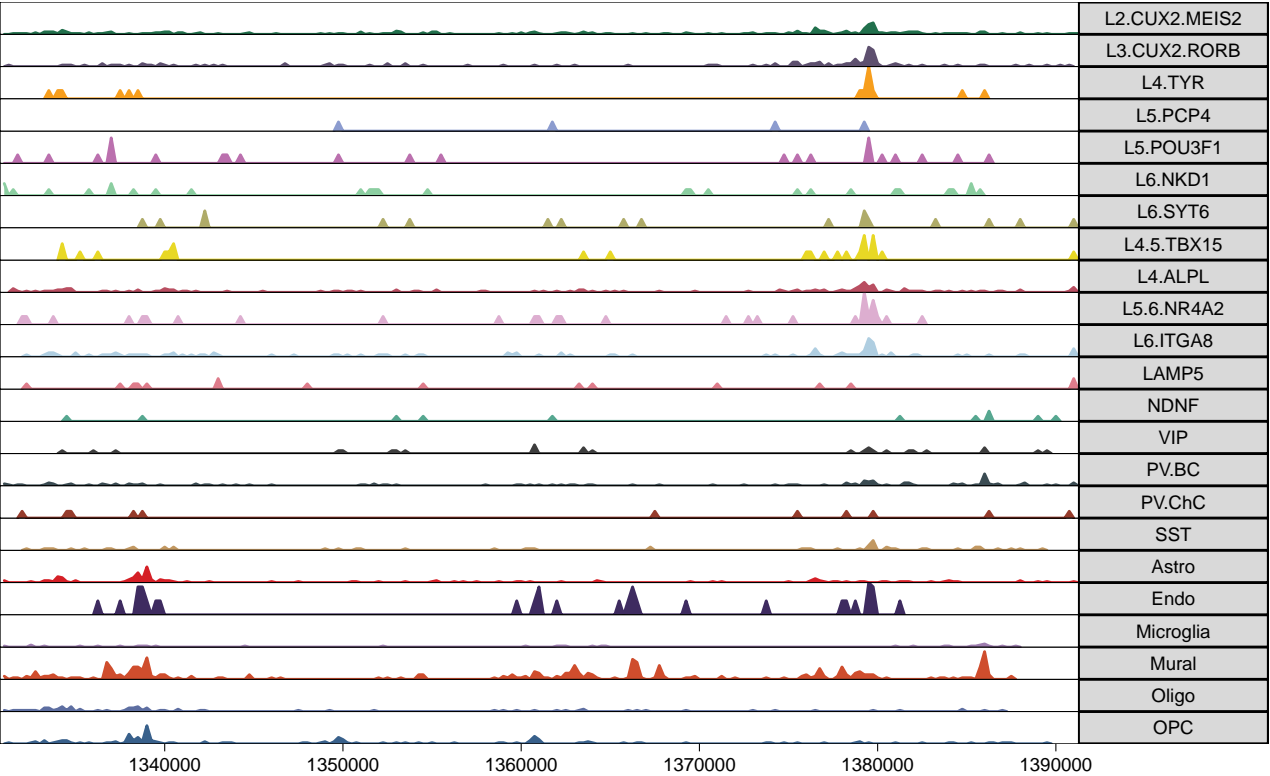


Peak2GeneLinks



Genes

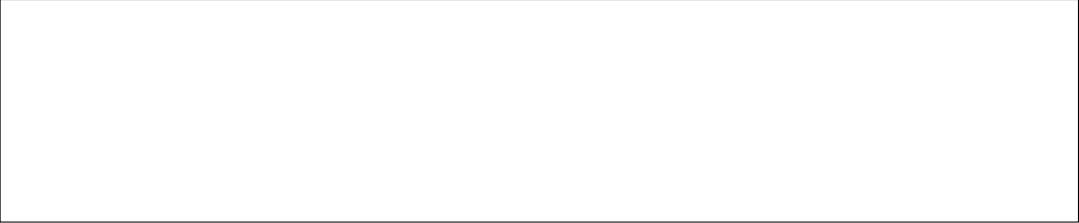
Coverage
(Norm. ATAC Signal Range (0-0.9) by Reads in TSS)



L5.PCP4



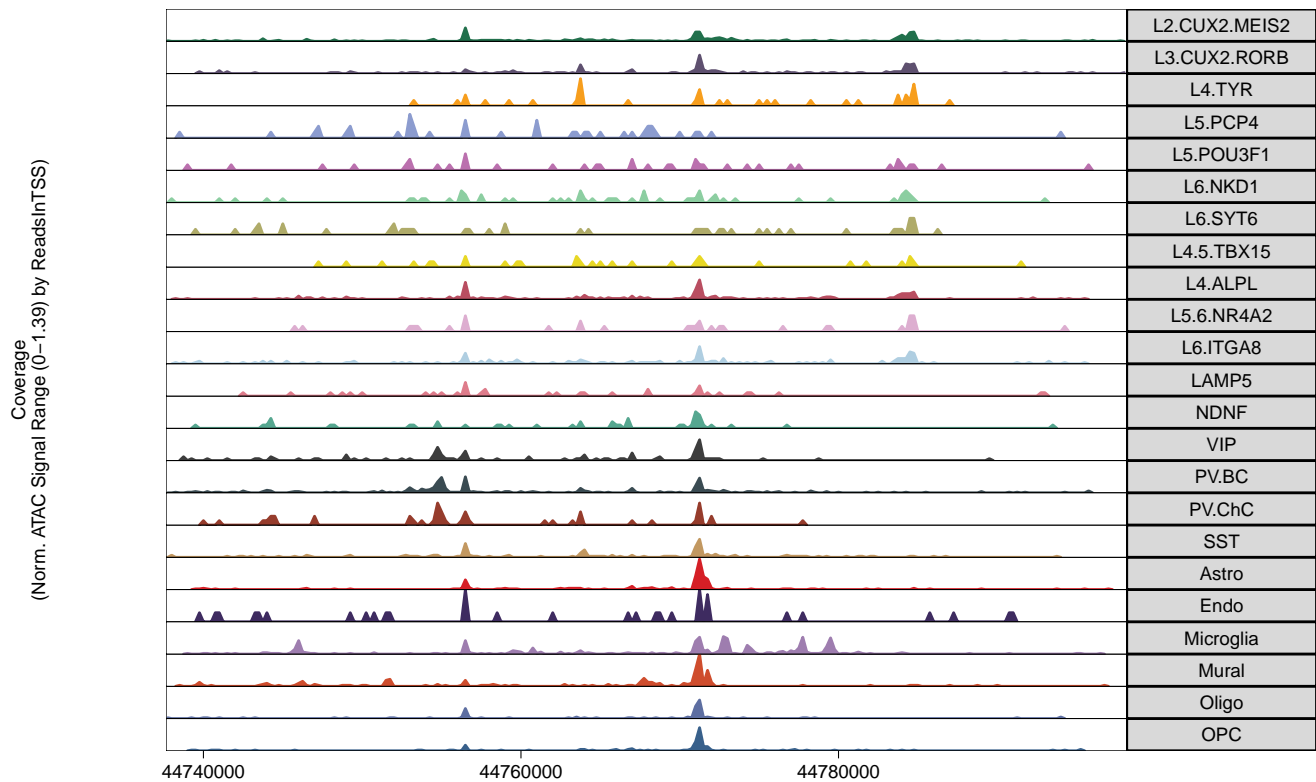
Peaks



LoopTrack



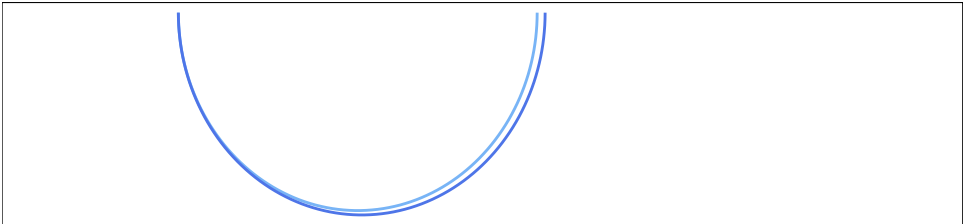
Genes



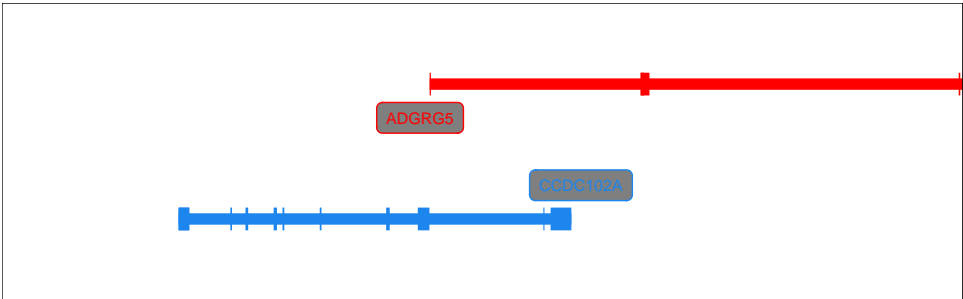
L5.PCP4



Peaks



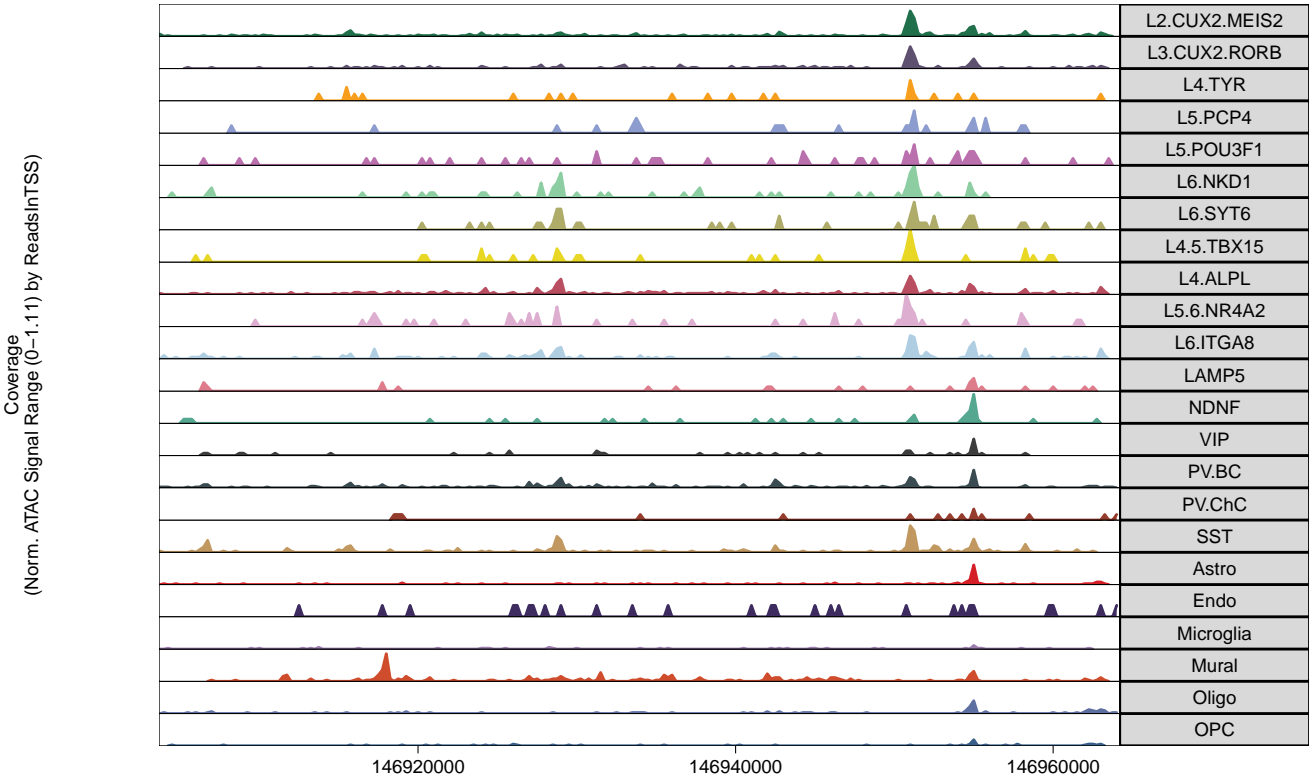
Peak2GeneLinks



Genes

value

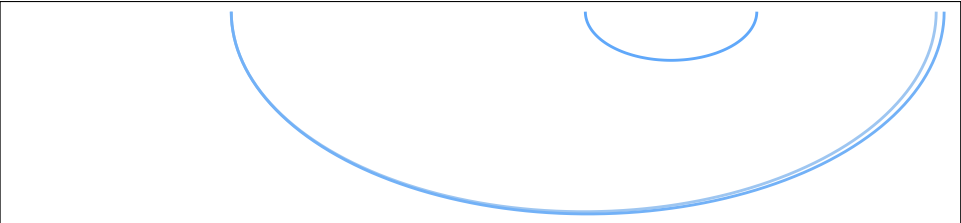




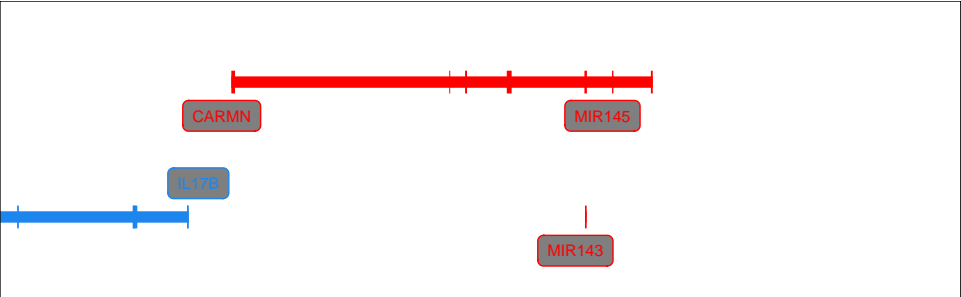
L5.PCP4



Peaks

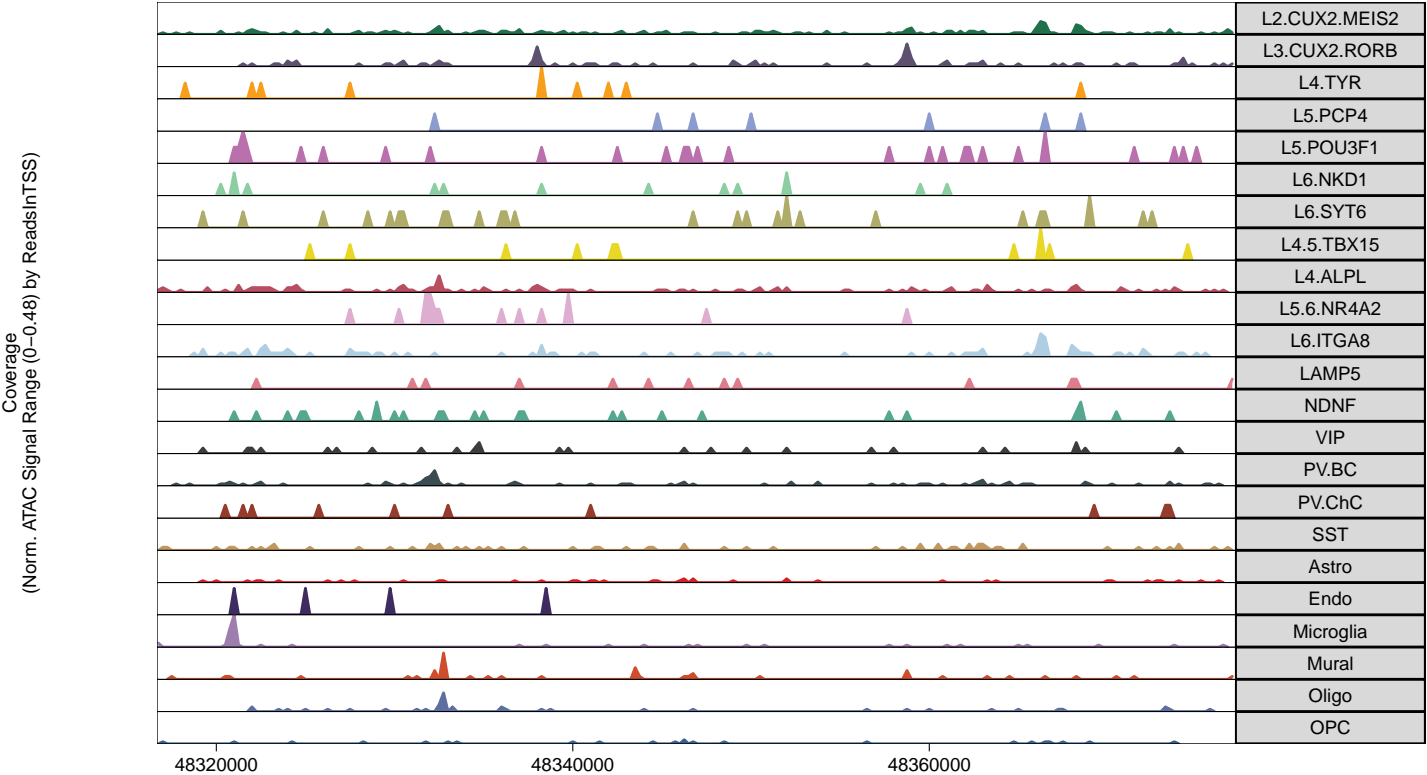


Peak2GeneLinks

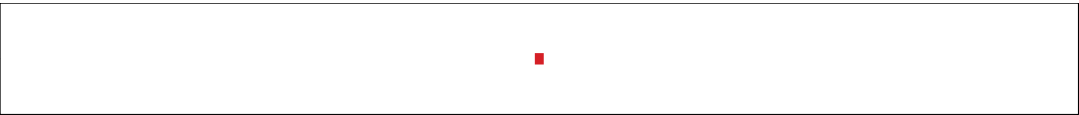


Genes

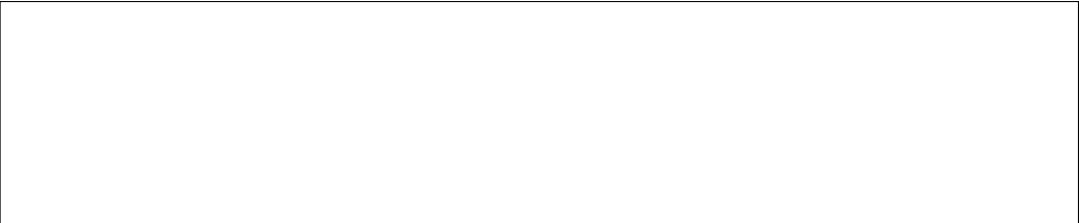
chr8:48316677-48377178 Marker Peak Tracks for: L5.PCP4



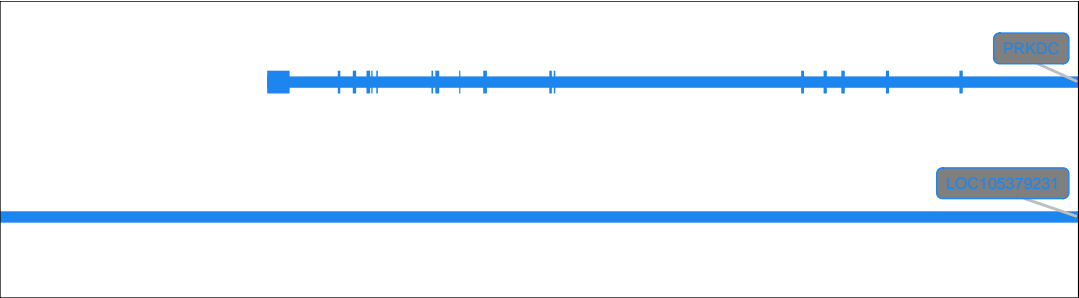
L5.PCP4



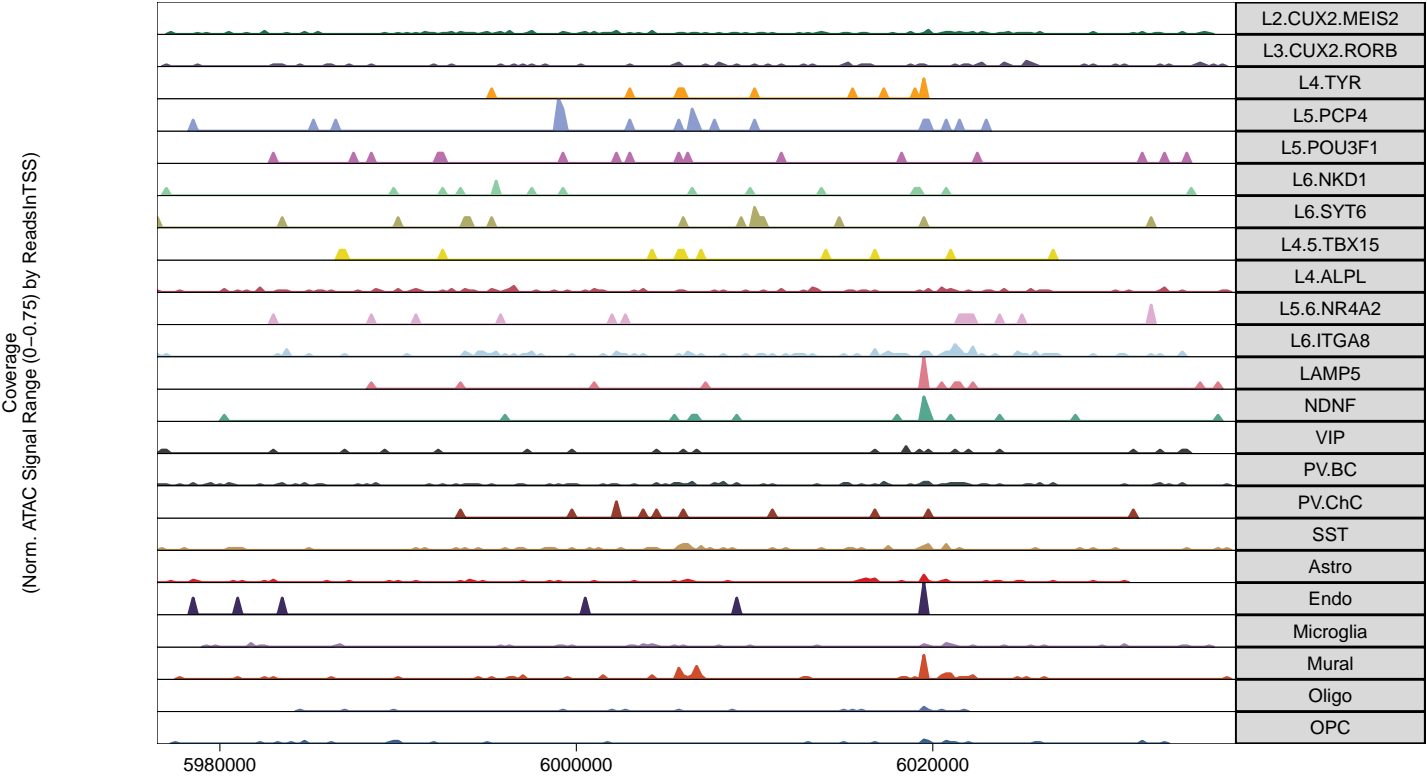
Peaks



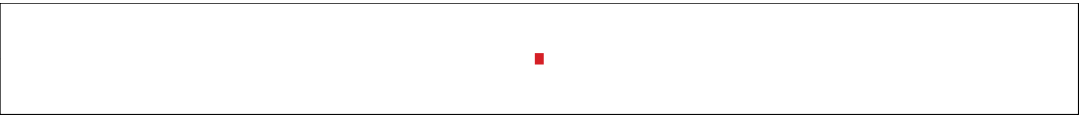
LoopTrack



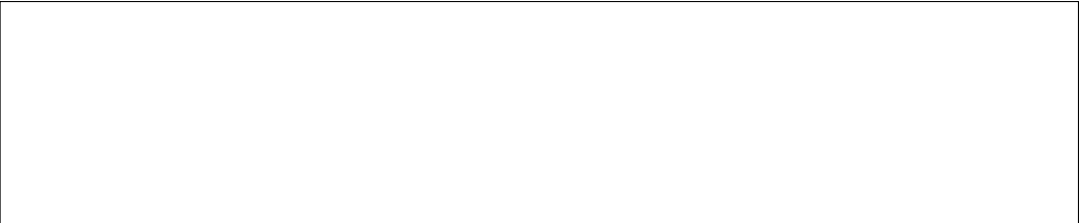
Genes



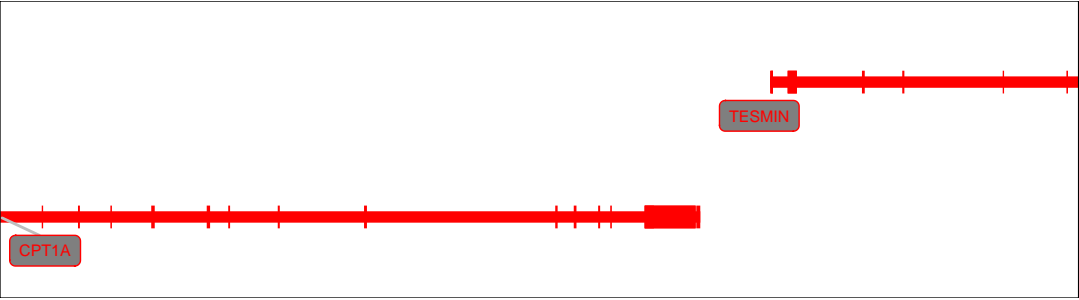
L5.PCP4



Peaks

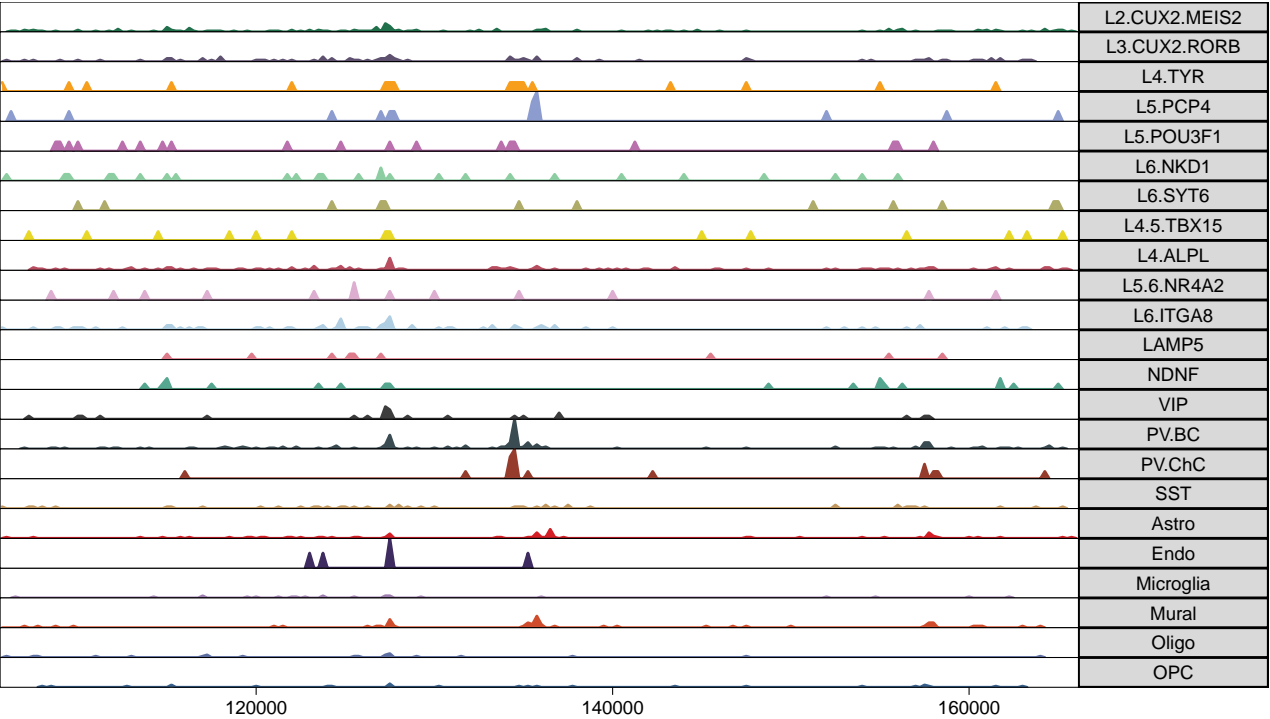


LoopTrack

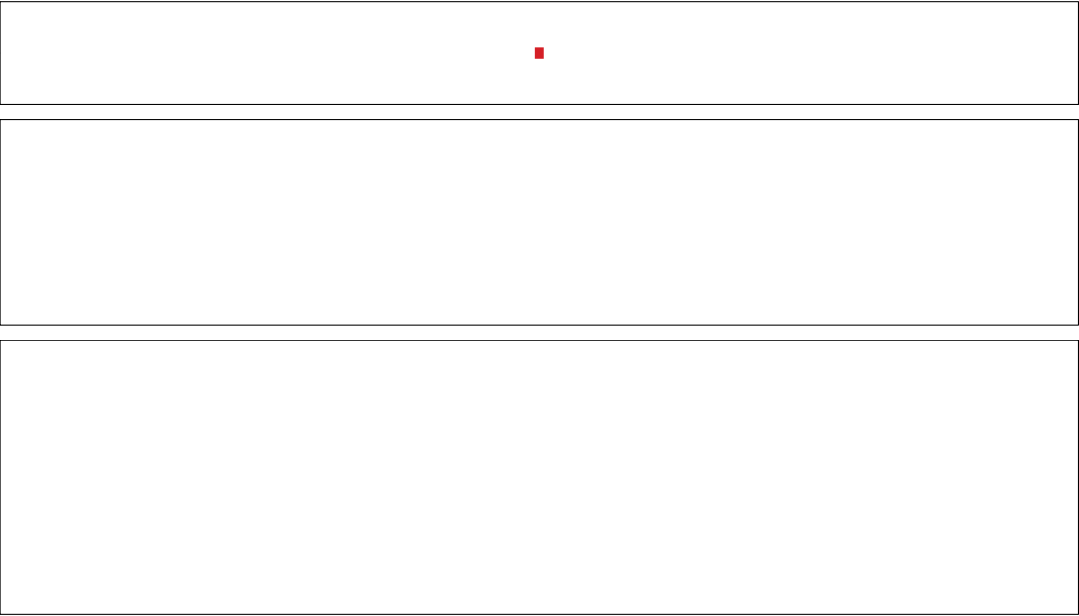


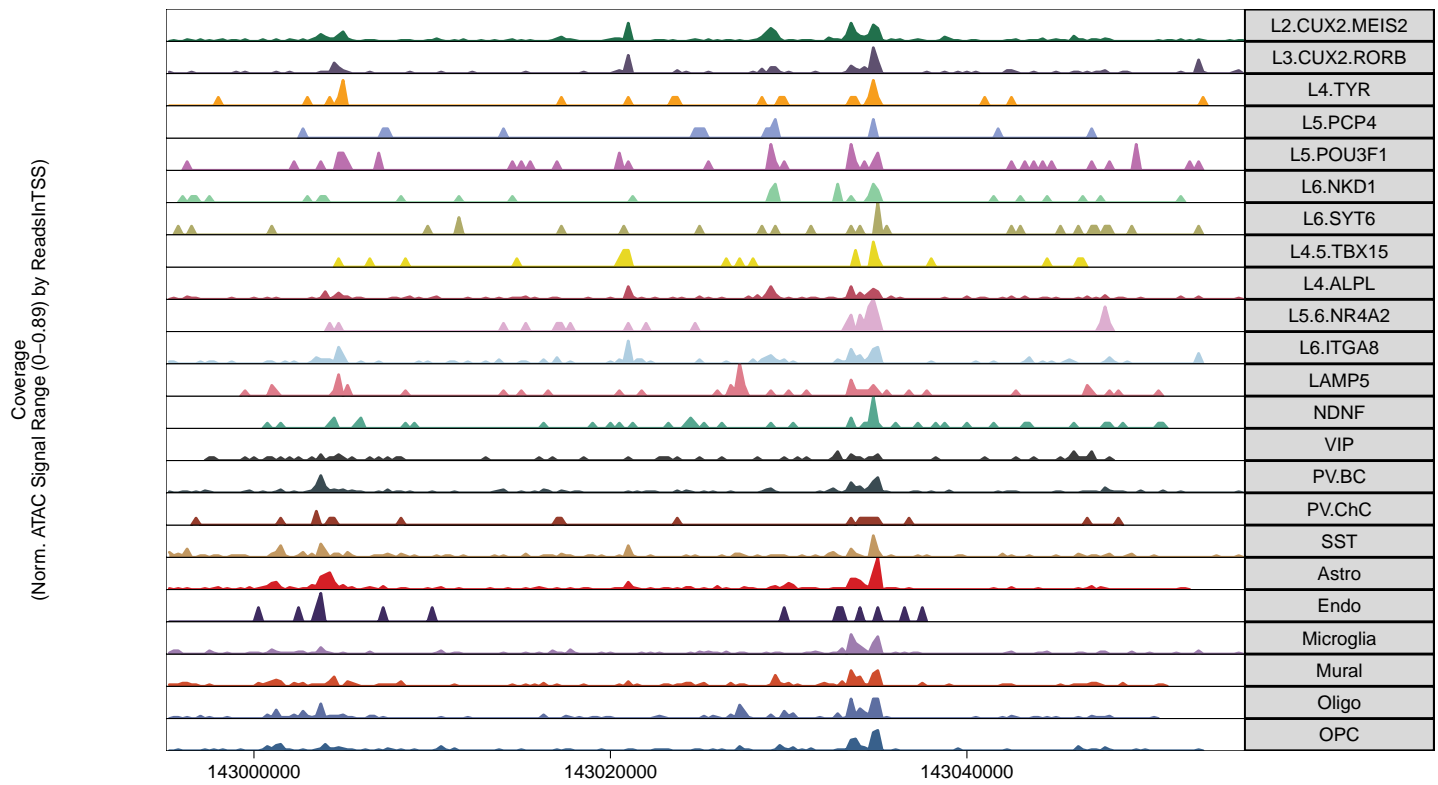
Genes

Coverage
(Norm. ATAC Signal Range (0–0.77) by Reads in TSS)

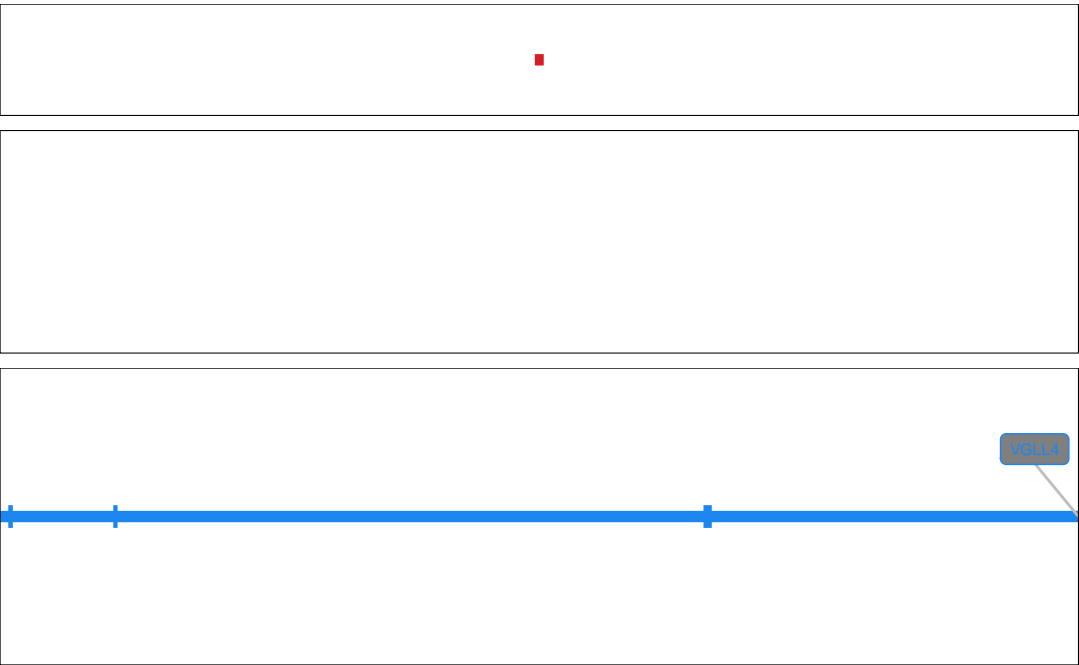


L5.PCP4

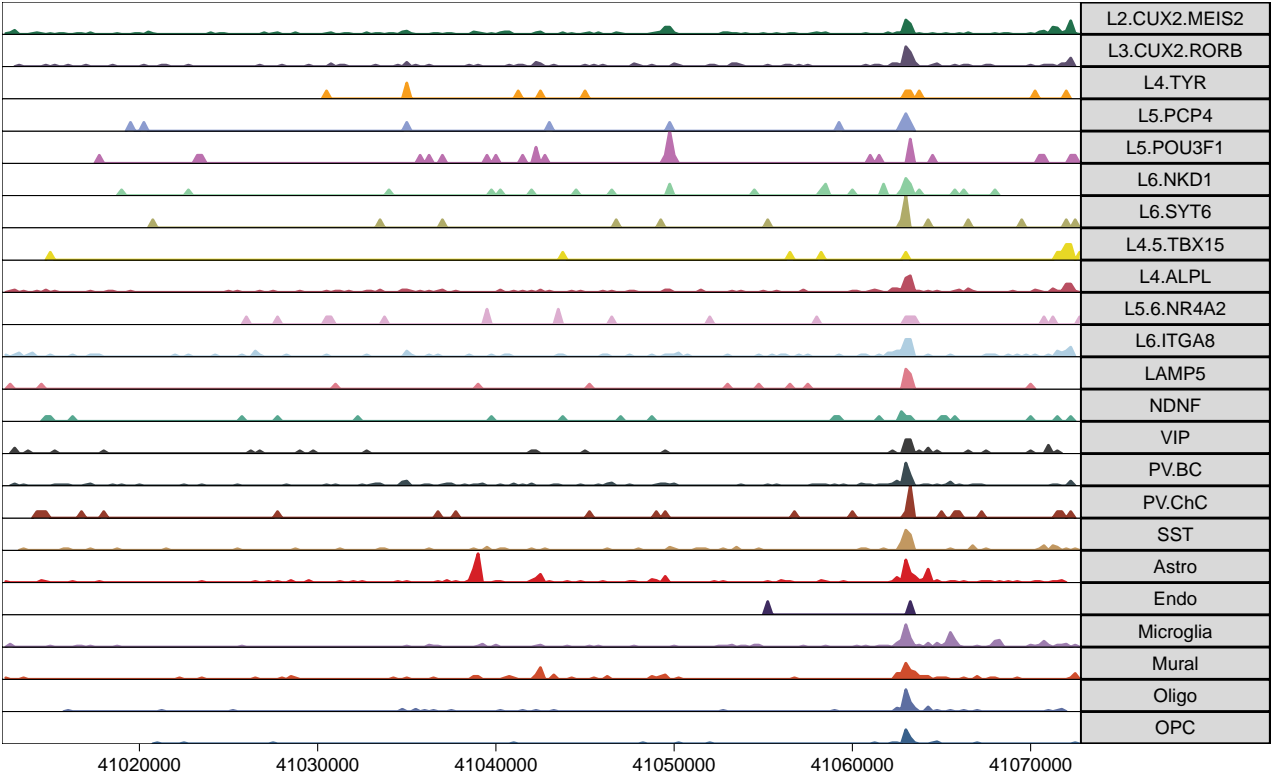




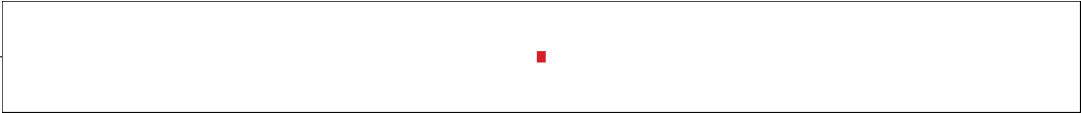
L5.PCP4



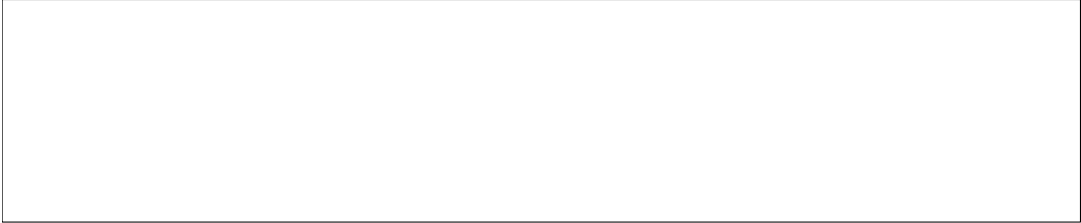
Coverage
(Norm. ATAC Signal Range (0–0.95) by ReadsInTSS)



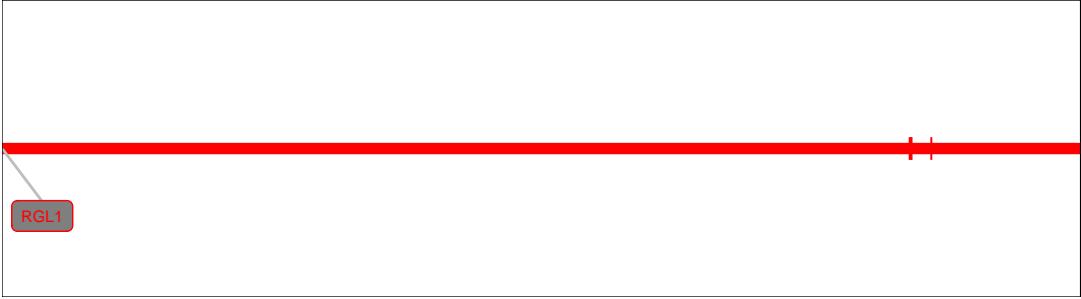
L5.PCP4



Peaks

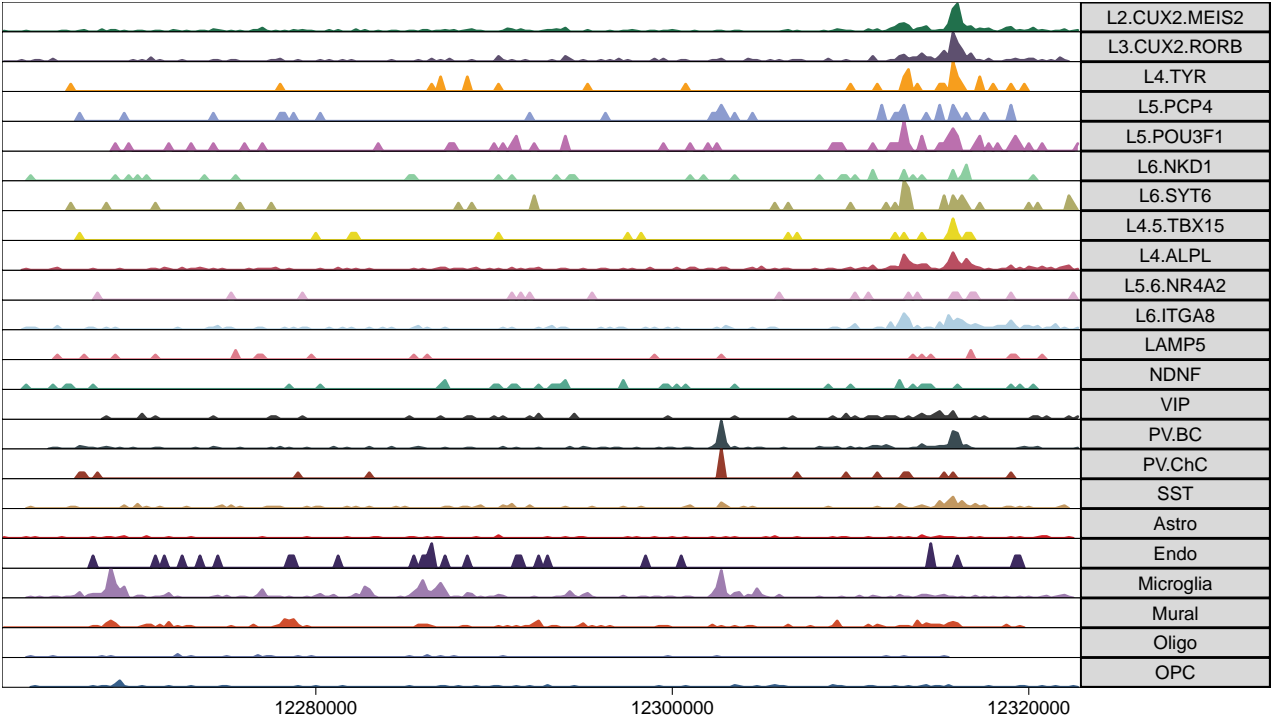


LoopTrack

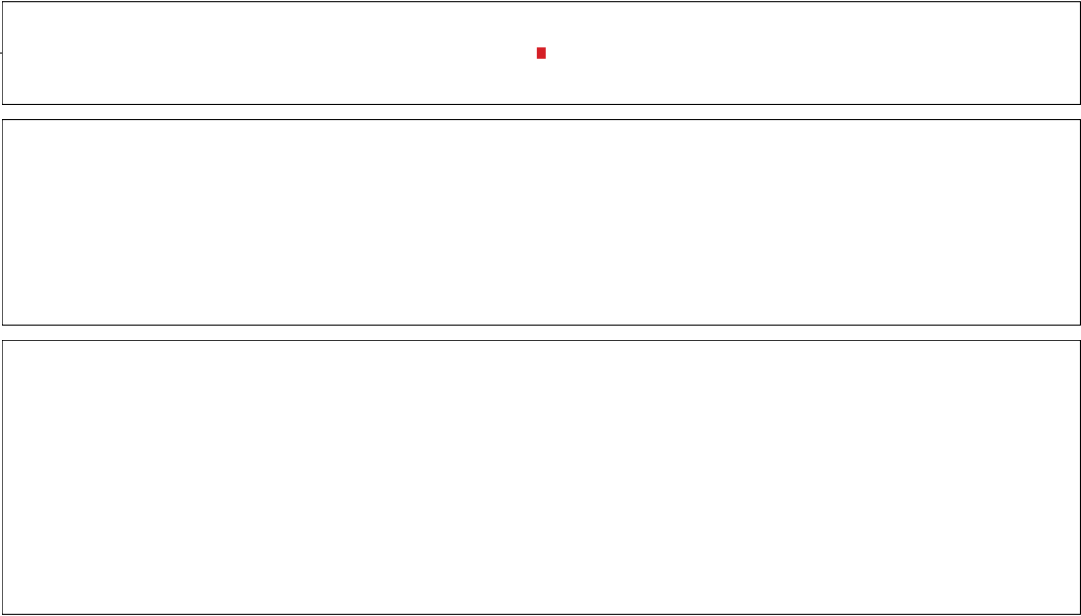


Genes

Coverage
(Norm. ATAC Signal Range (0–0.95) by Reads in TSS)



L5.PCP4

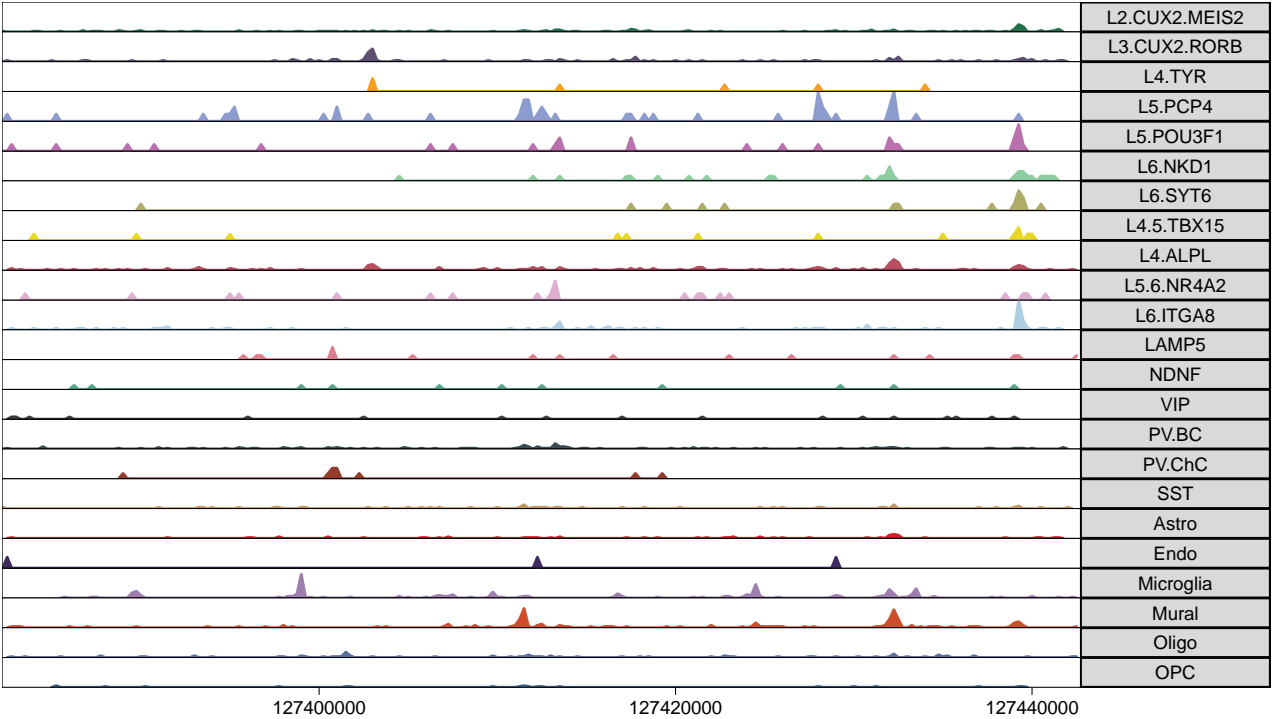


Peaks

LoopTrack

GeneTrack

Coverage
(Norm. ATAC Signal Range (0–1.07) by Reads in TSS)



L5.PCP4

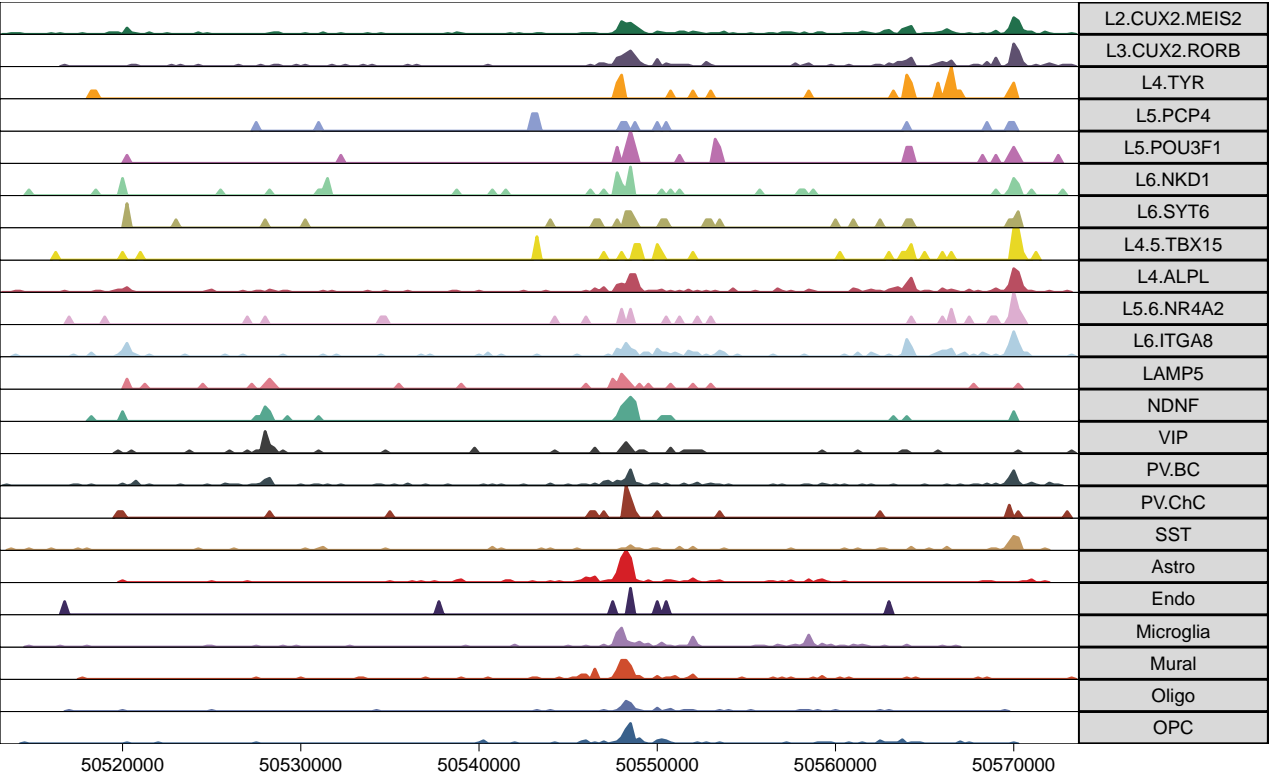


Peaks

LoopTrack

GeneTrack

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



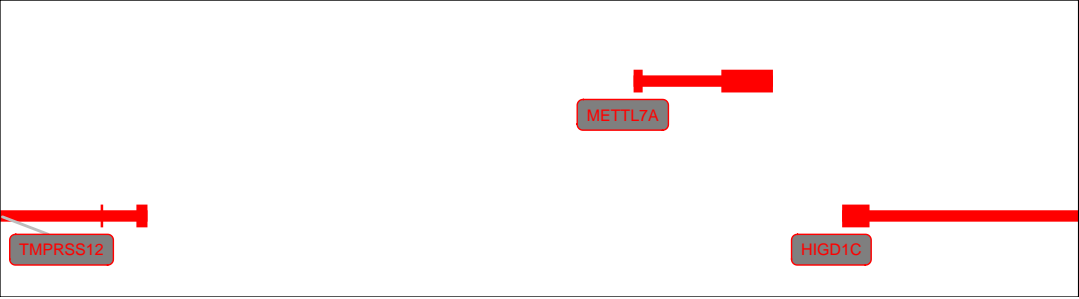
L5.PCP4



Peaks

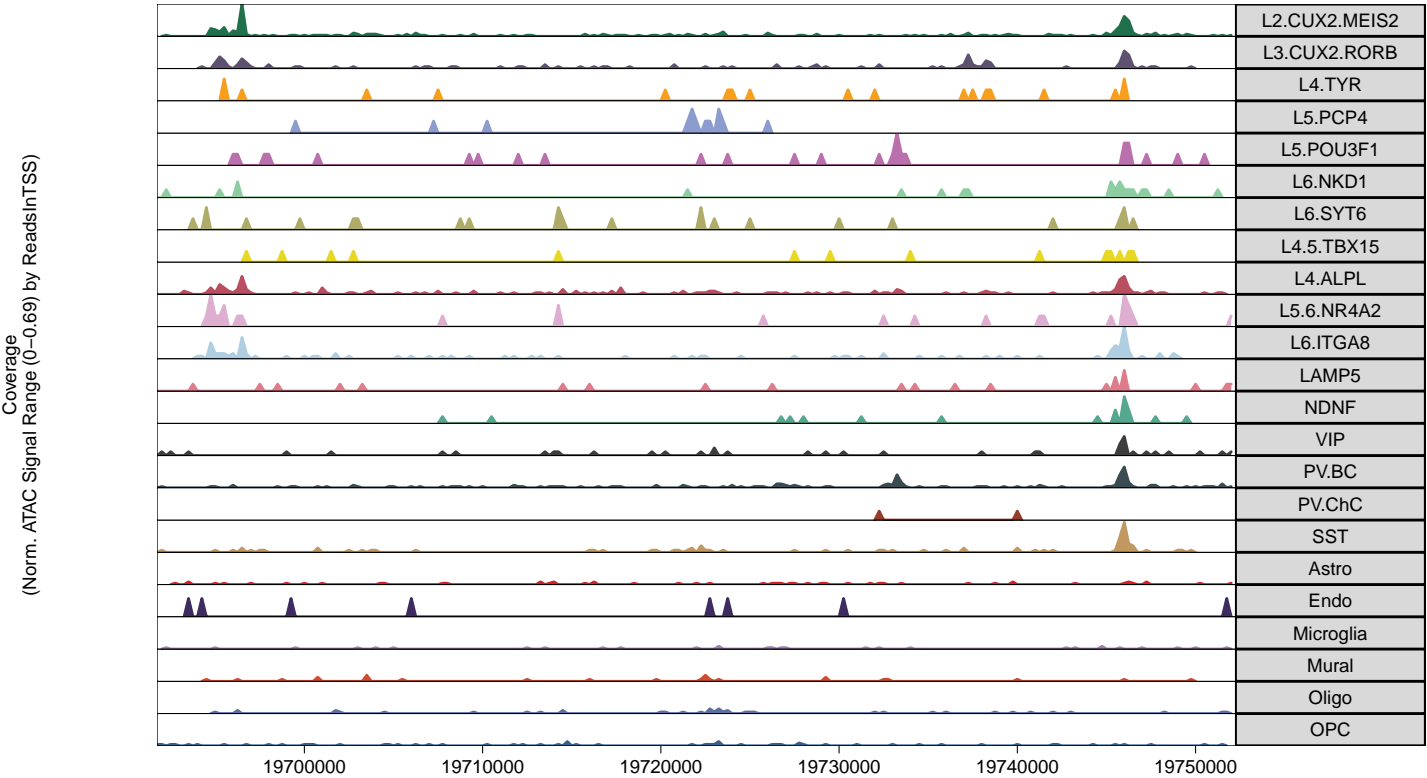


LoopTrack

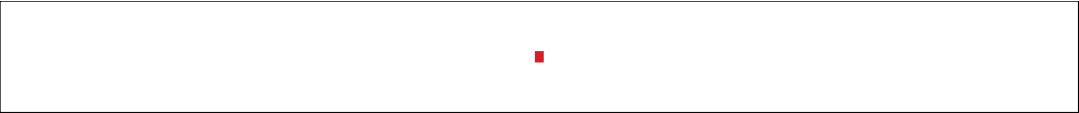


Genes

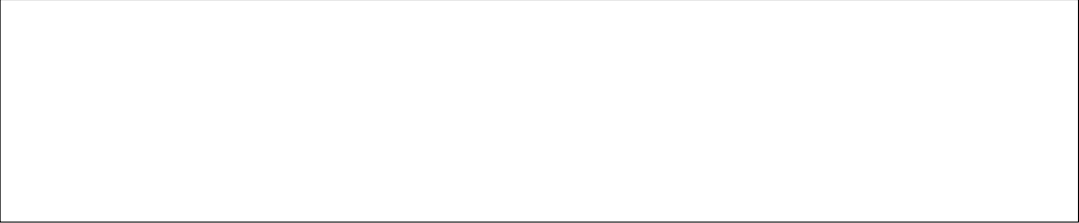
chr14:19691737-19752238 Marker Peak Tracks for: L5.PCP4



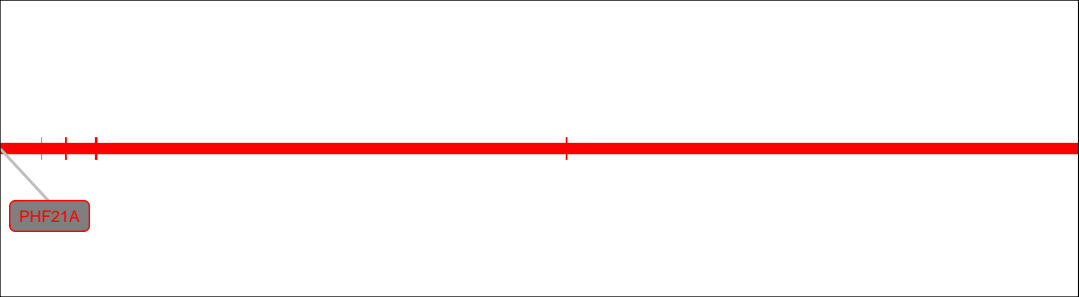
L5.PCP4



Peaks

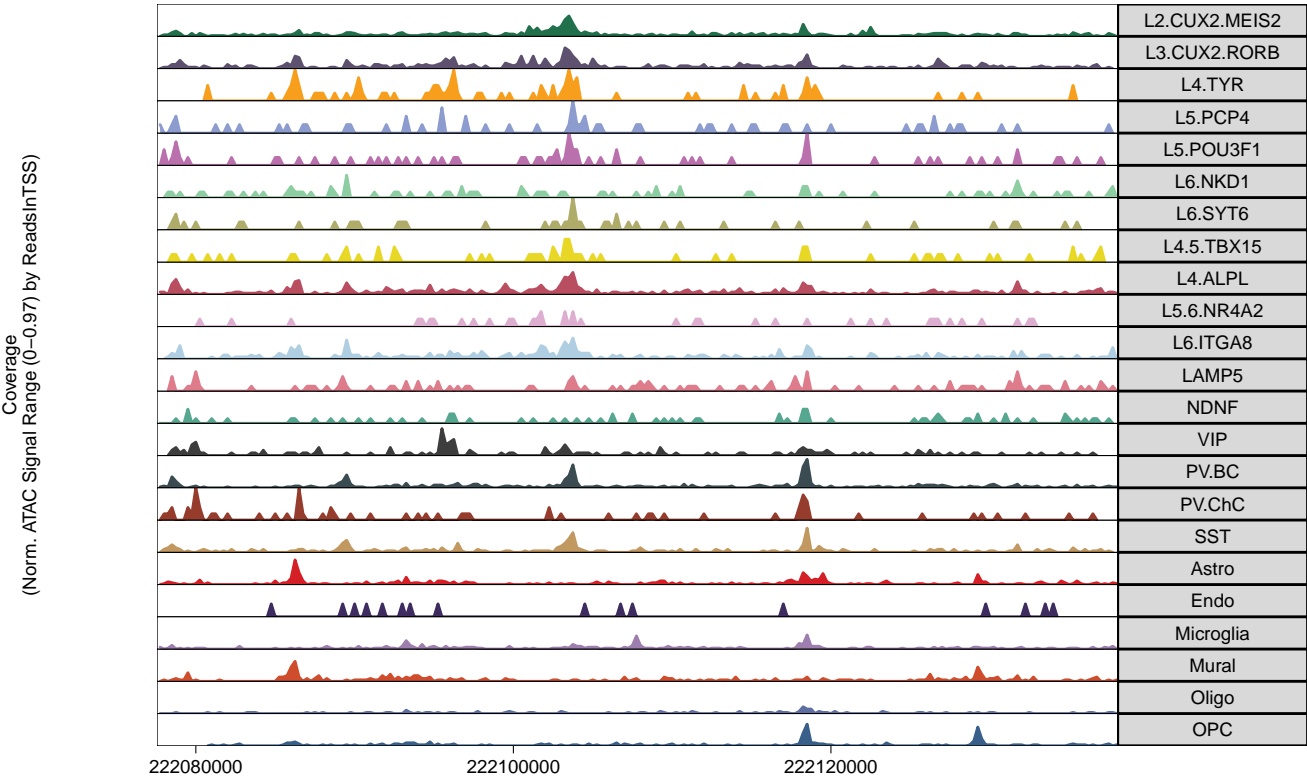


LoopTrack



Genes

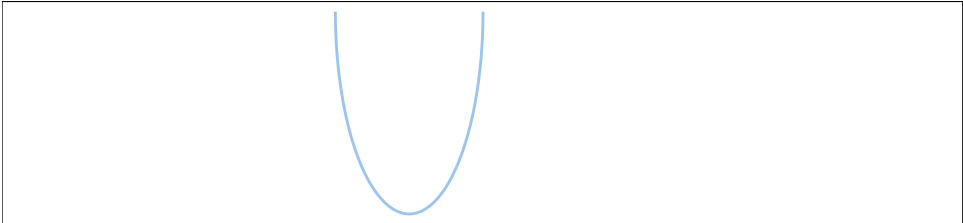
chr1:222077561–222138062 Marker Peak Tracks for: L5.PCP4



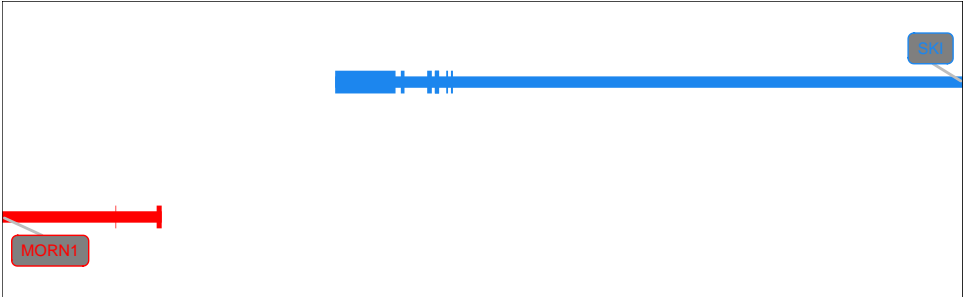
L5.PCP4



Peaks

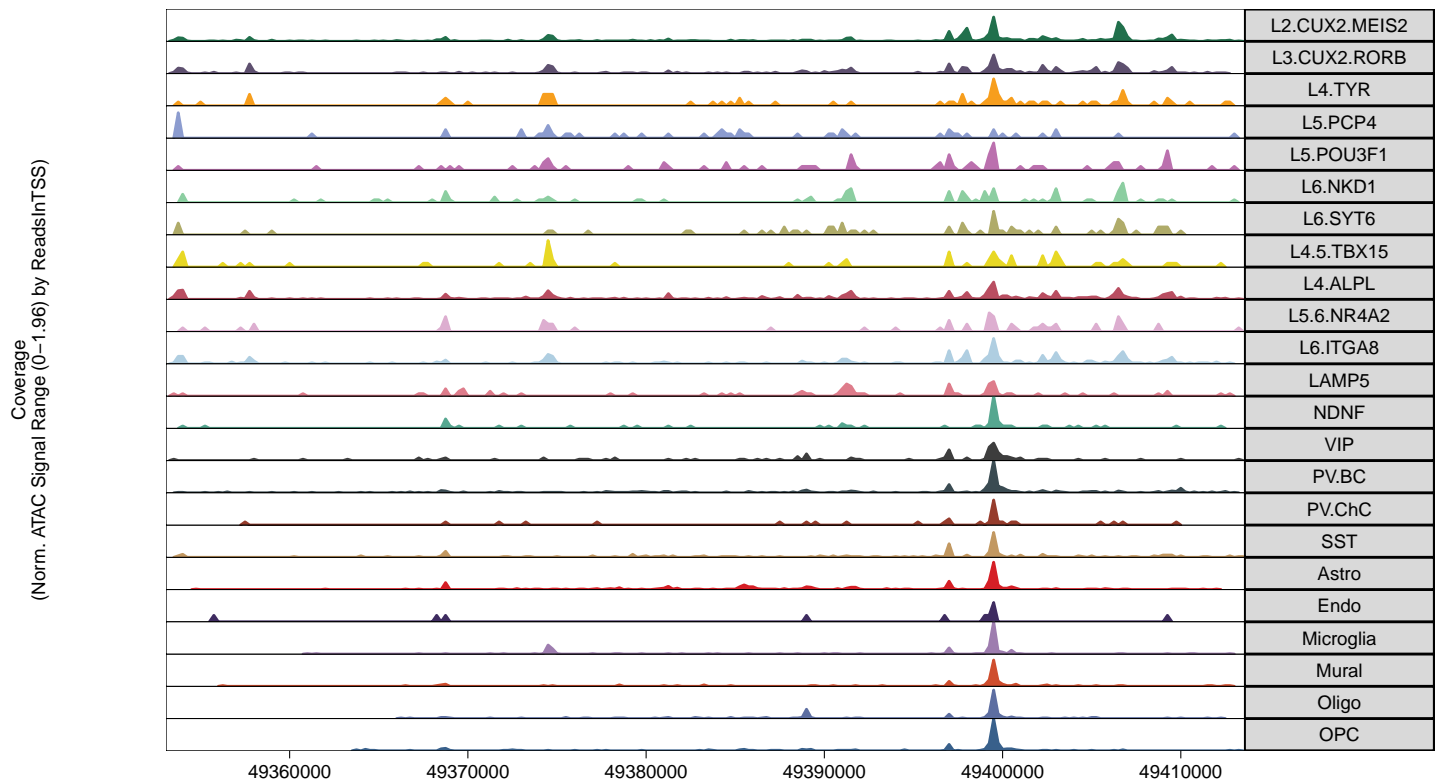


Peak2GeneLinks



Genes

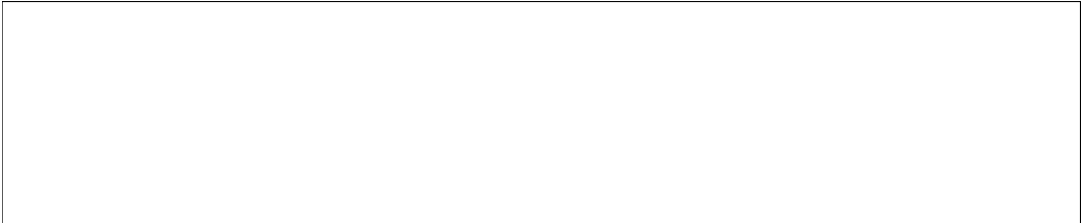




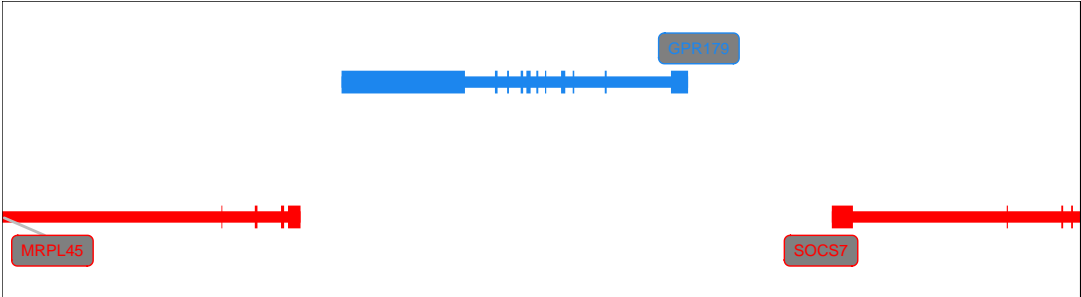
L5.PCP4



Peaks

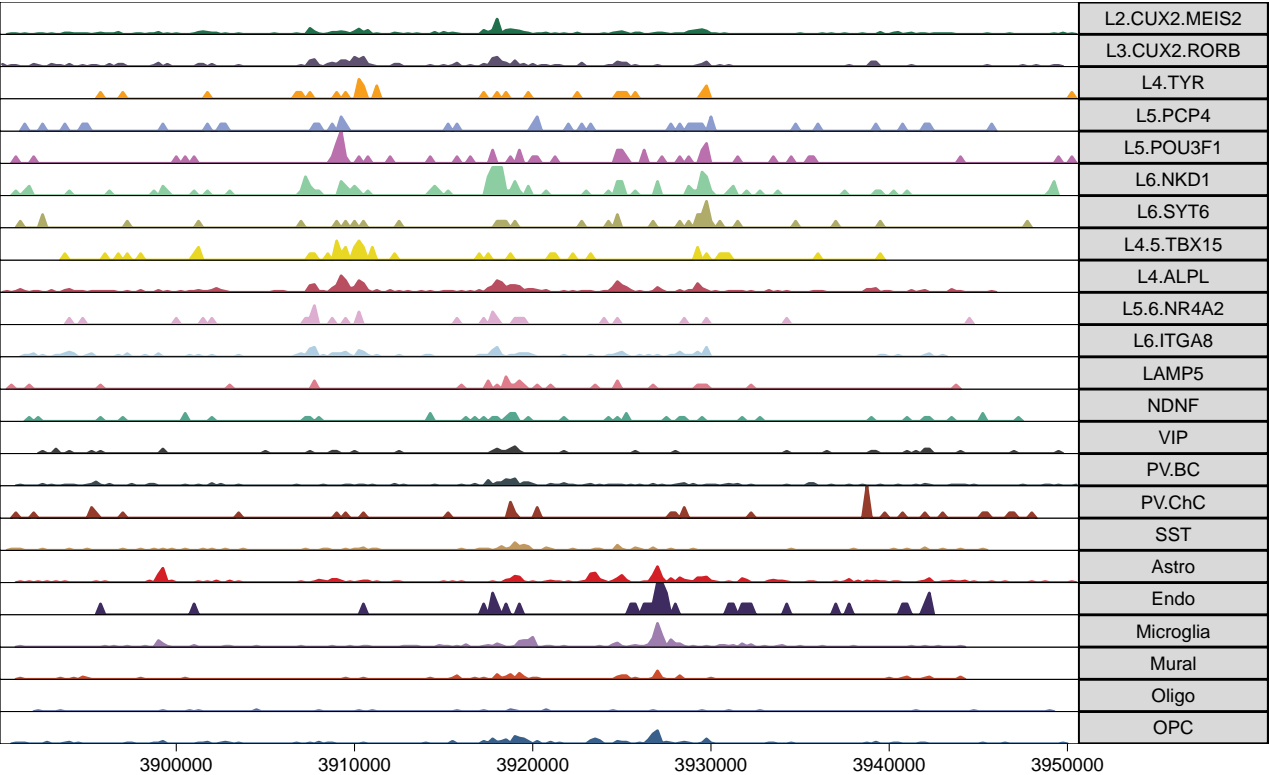


LoopTrack



Genes

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



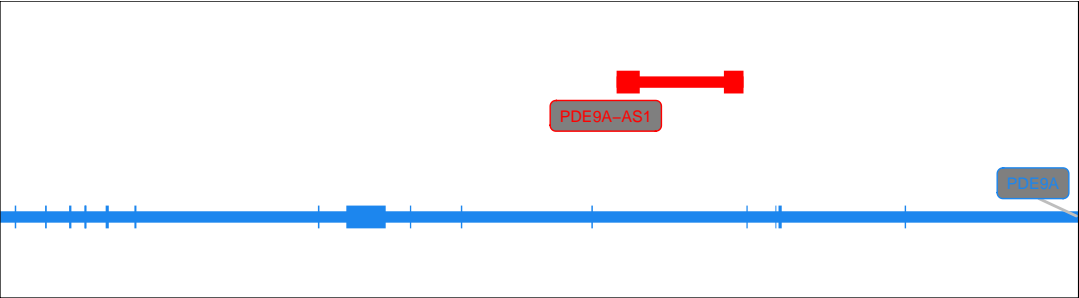
L5.PCP4



Peaks

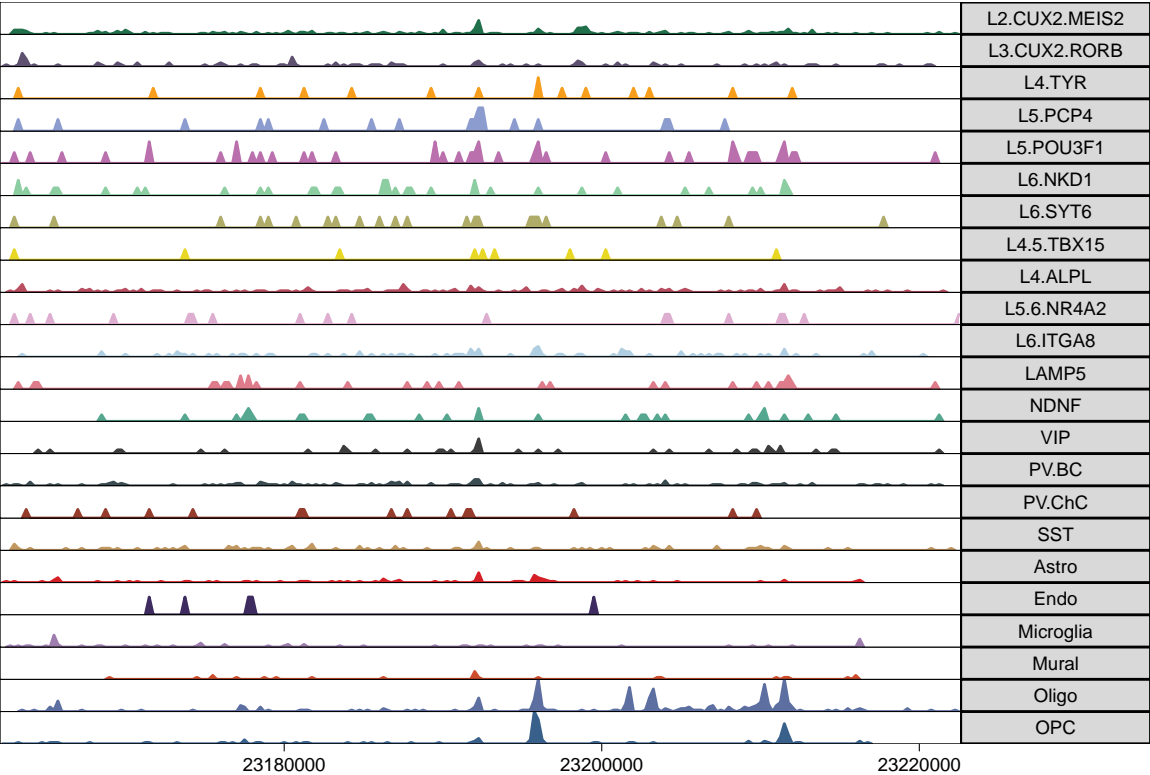


LoopTrack



Genes

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



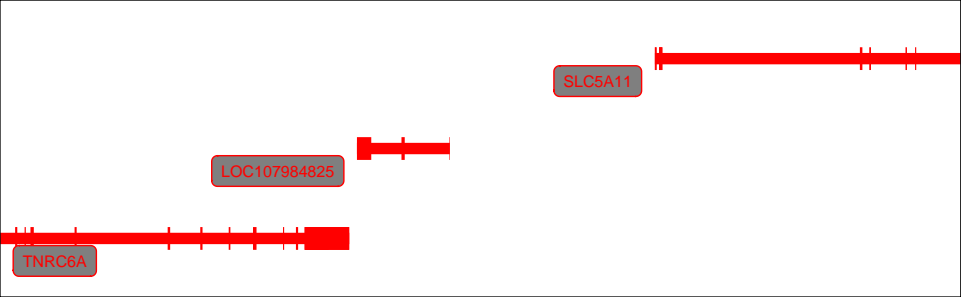
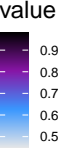
L5.PCP4



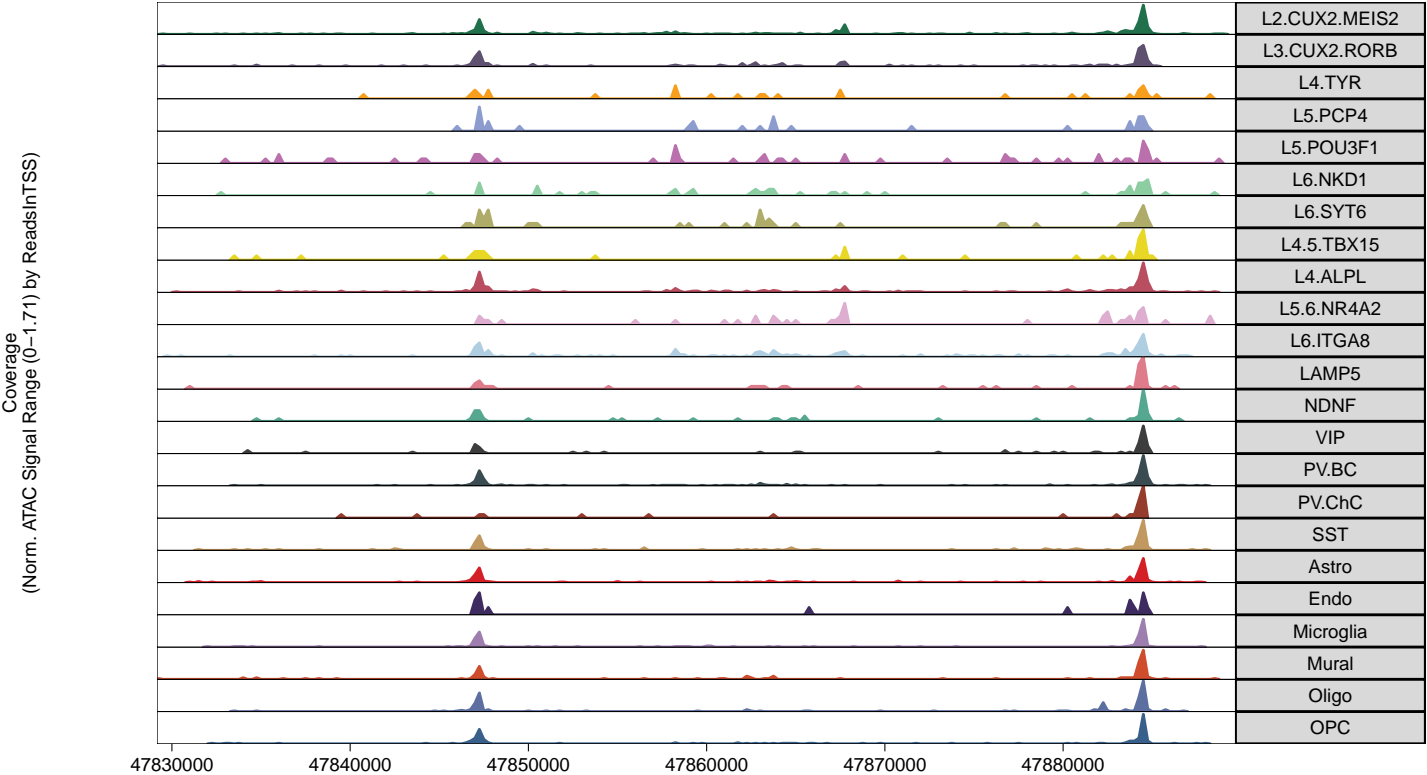
Peaks



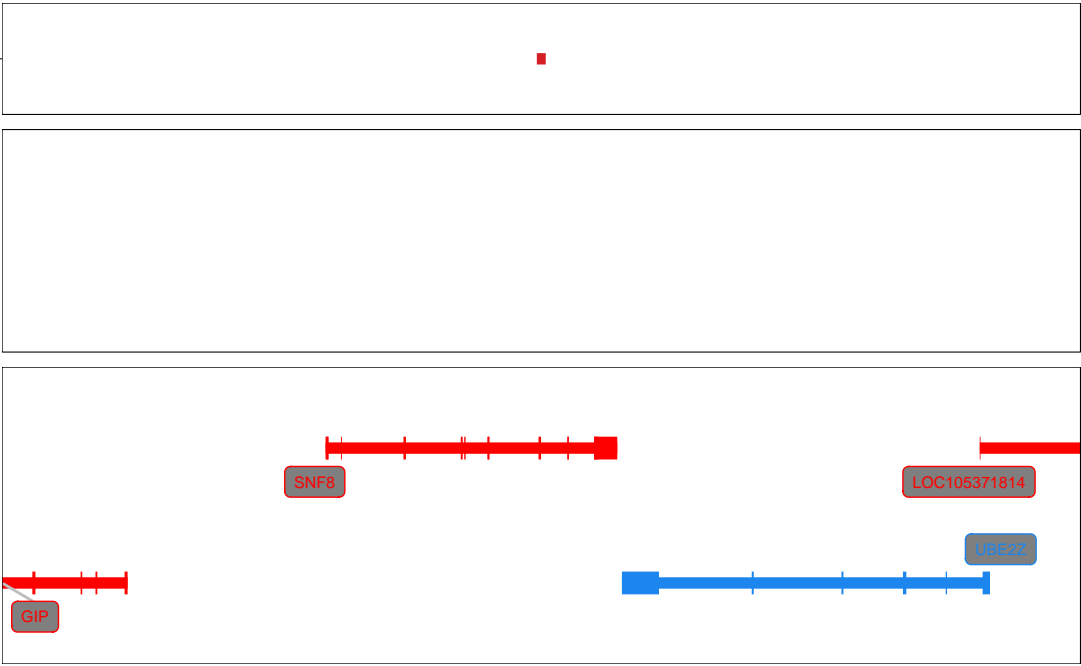
Peak2GeneLinks



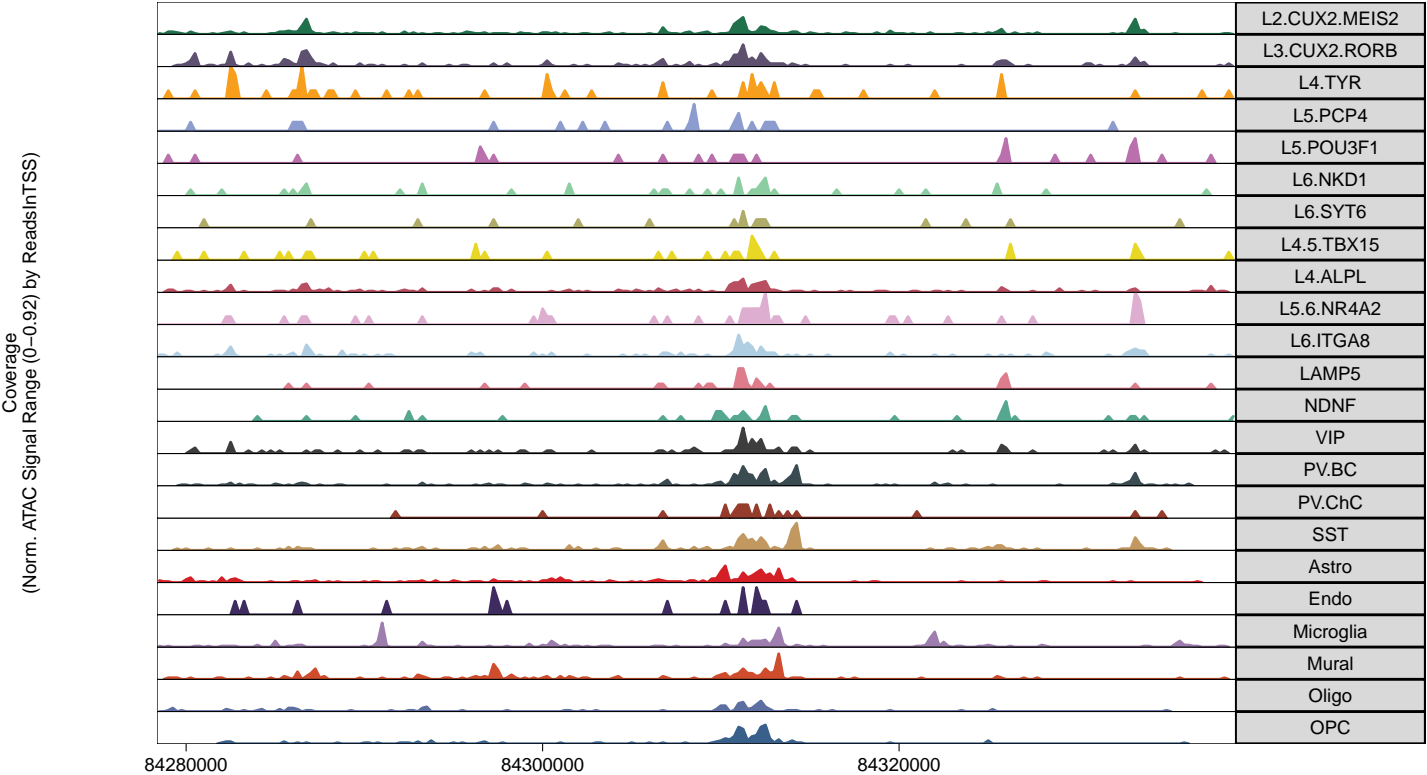
Genes



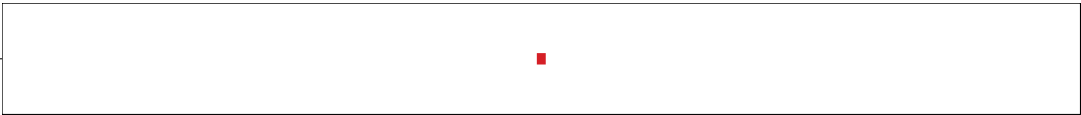
L5.PCP4



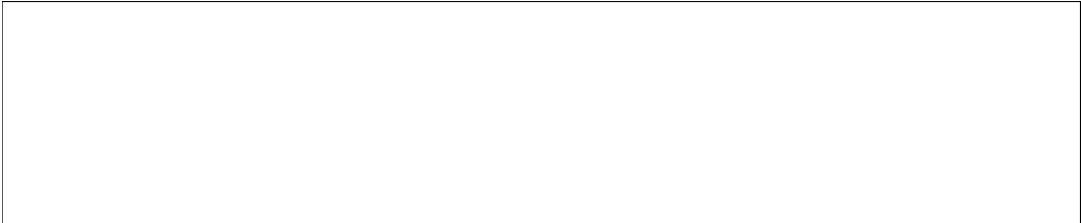
chr1:84278371-84338872 Marker Peak Tracks for: L5.PCP4



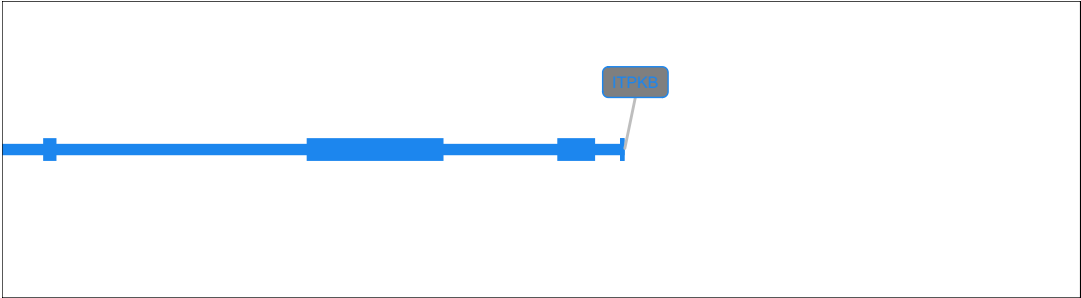
L5.PCP4



Peaks

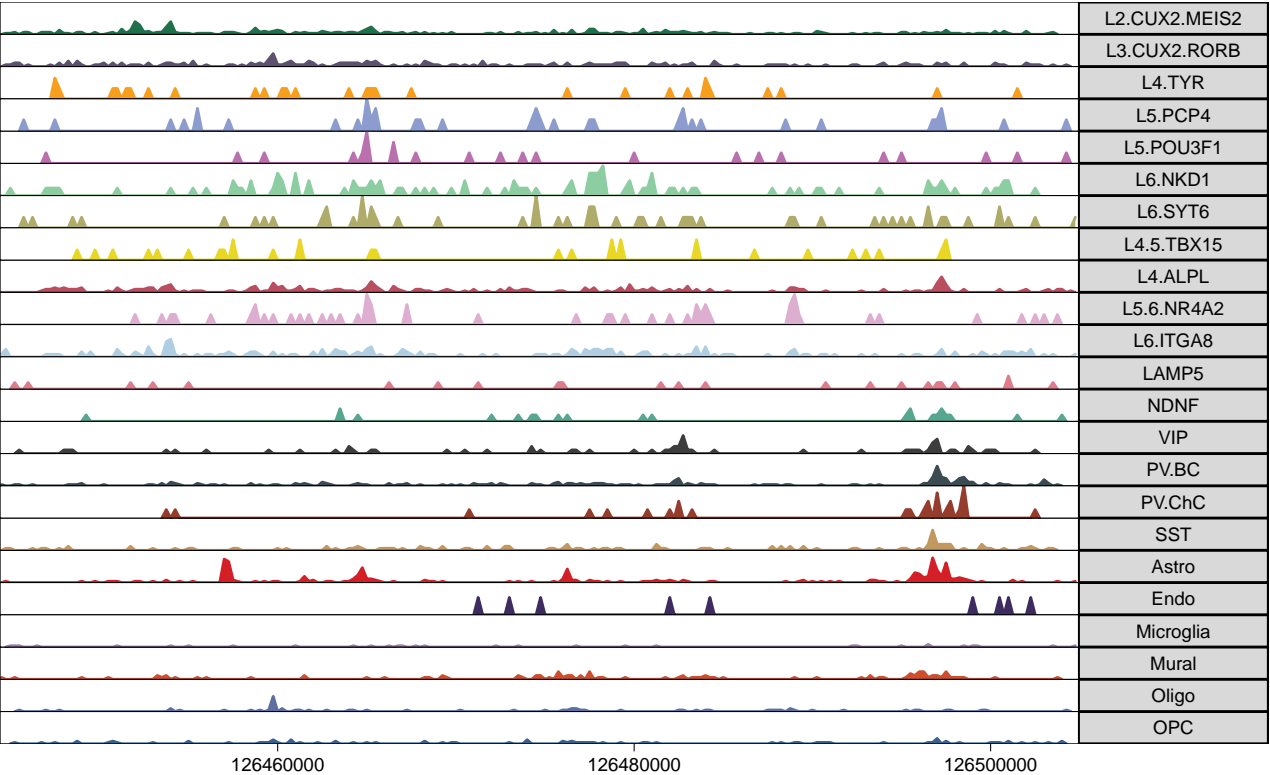


LoopTrack



Genes

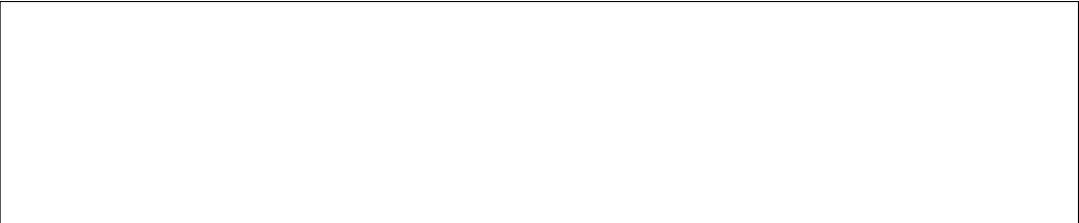
Coverage
(Norm. ATAC Signal Range (0-0.73) by ReadsInTSS)



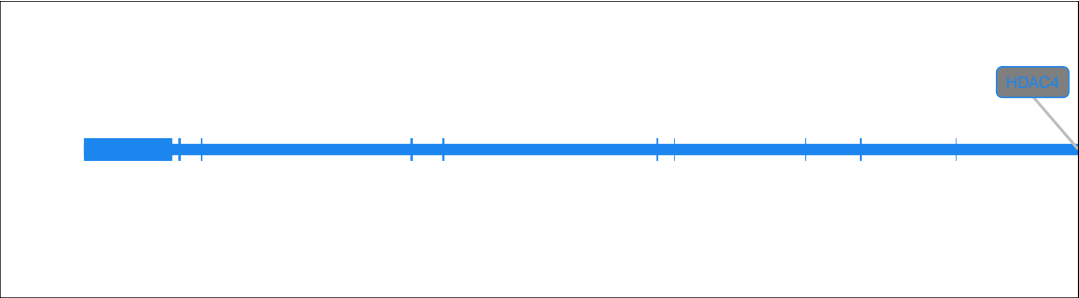
L5.PCP4



Peaks

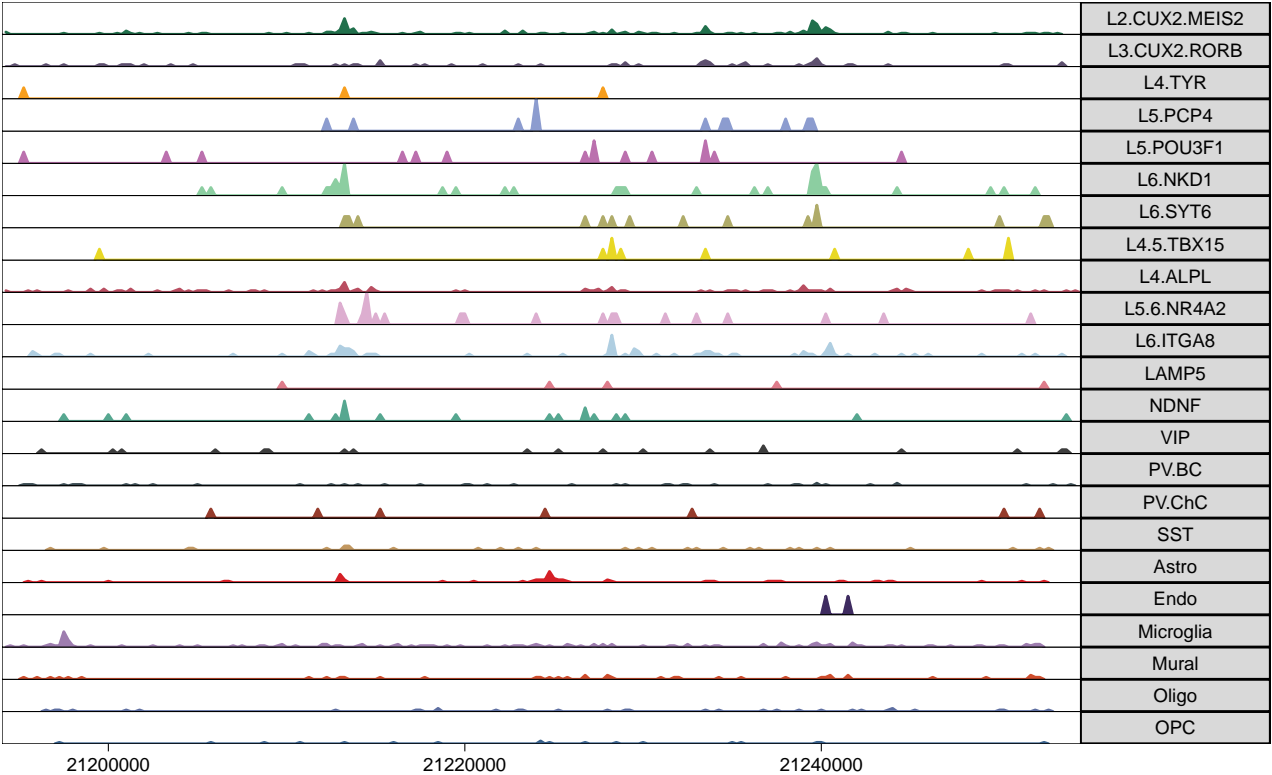


LoopTrack



Genes

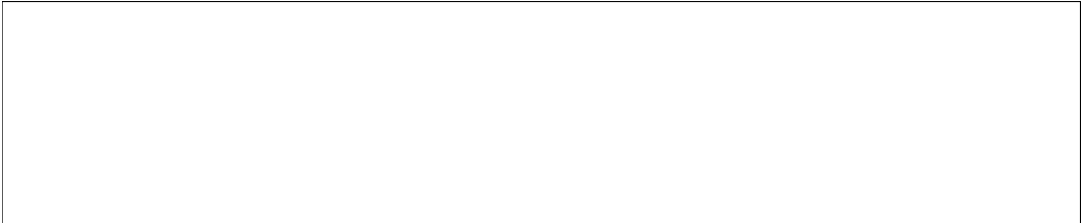
Coverage
(Norm. ATAC Signal Range 0–0.68) by ReadsInTSS)



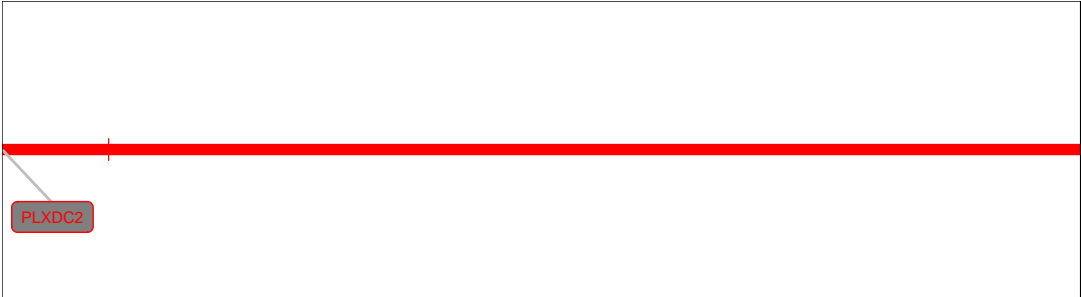
L5.PCP4



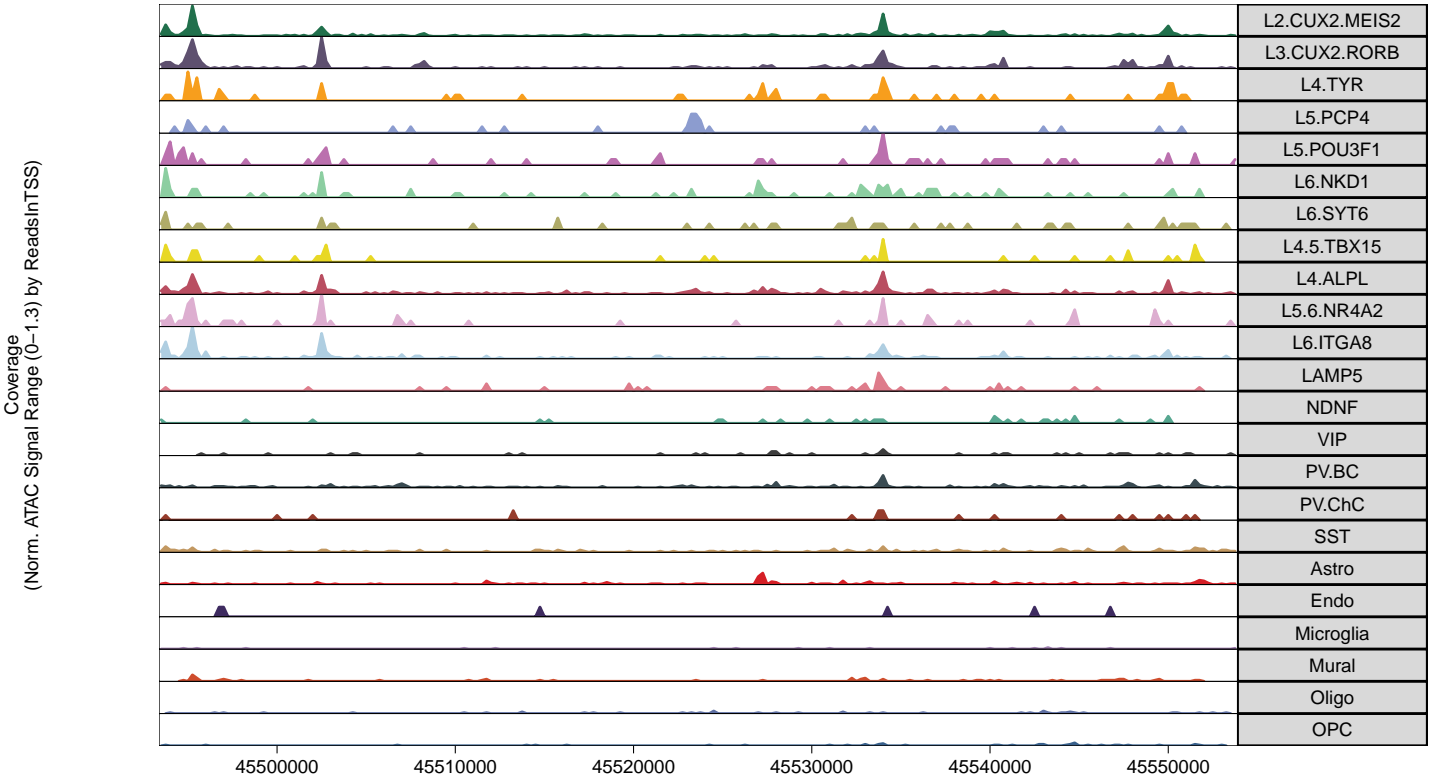
Peaks



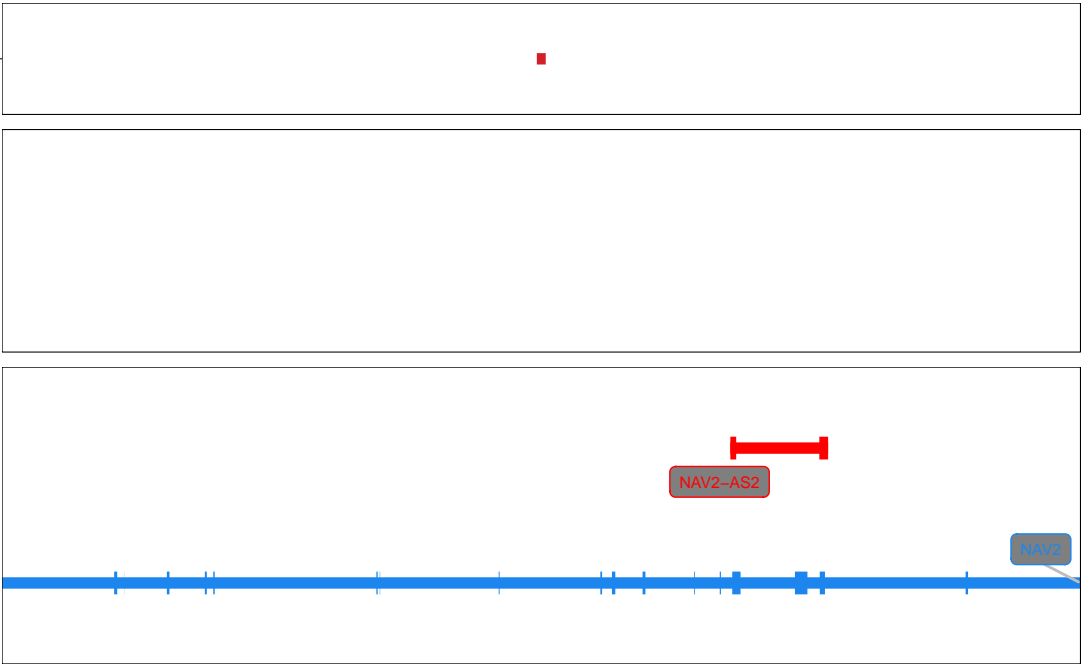
LoopTrack



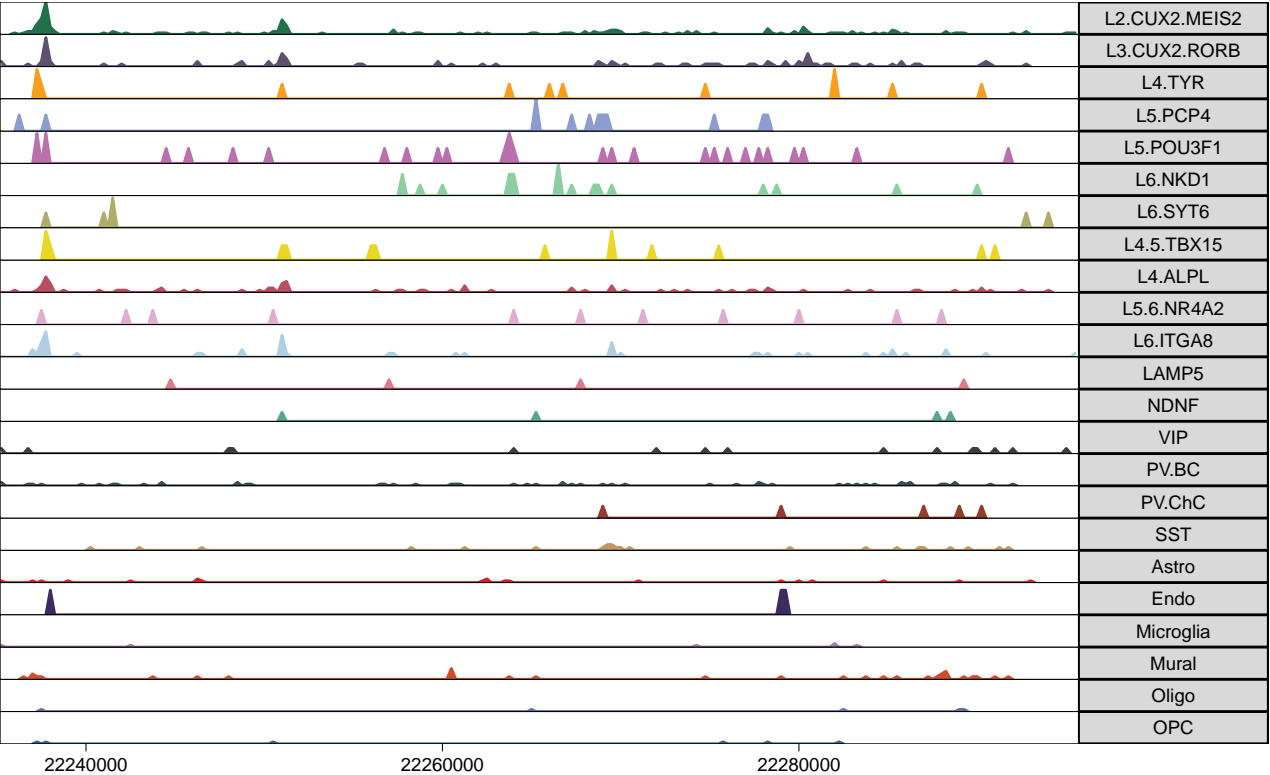
Genes



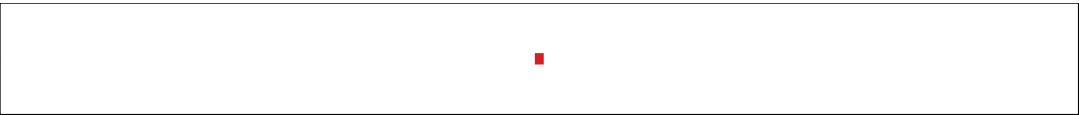
L5.PCP4



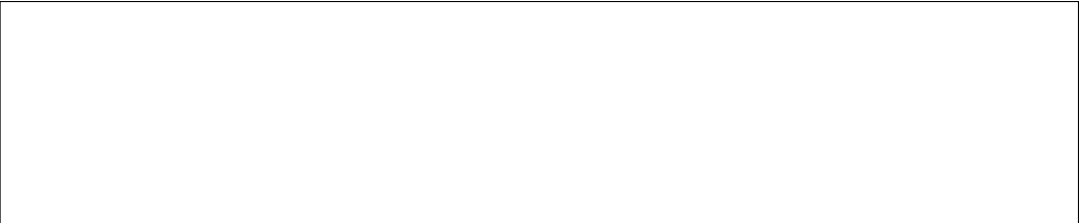
Coverage
(Norm. ATAC Signal Range (0-0.51) by ReadsInTSS)



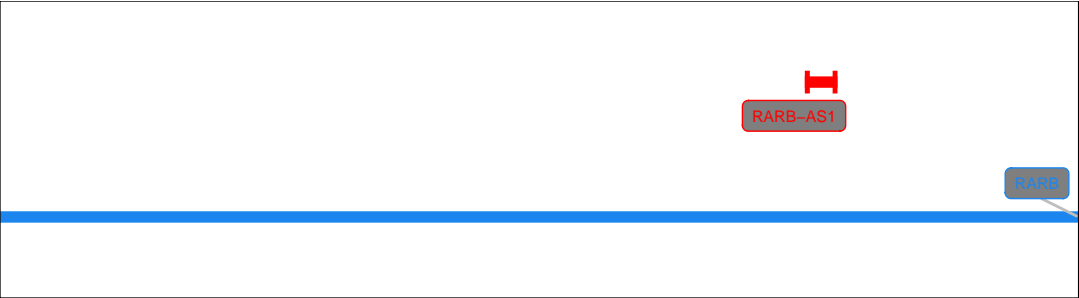
L5.PCP4



Peaks

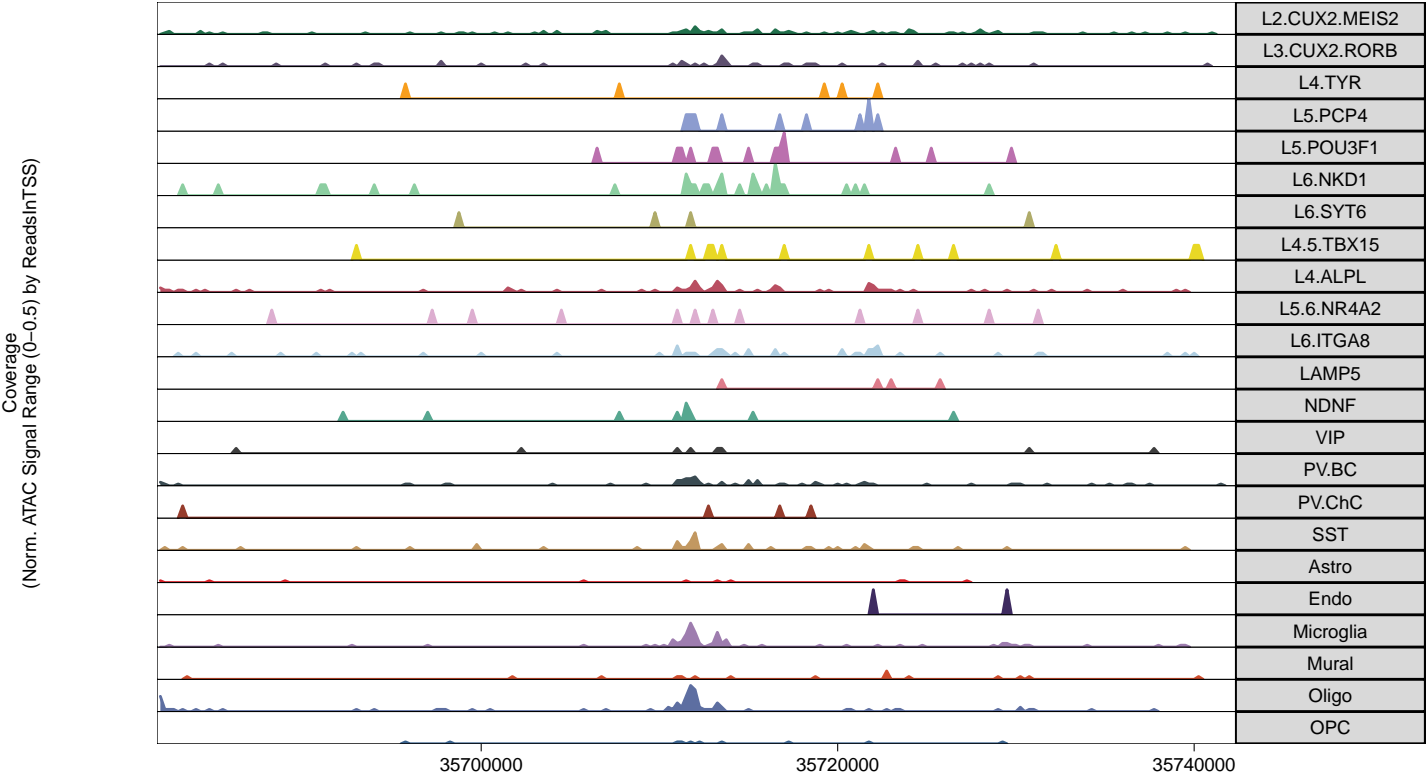


LoopTrack



Genes

chr10:35681816–35742317 Marker Peak Tracks for: L5.PCP4



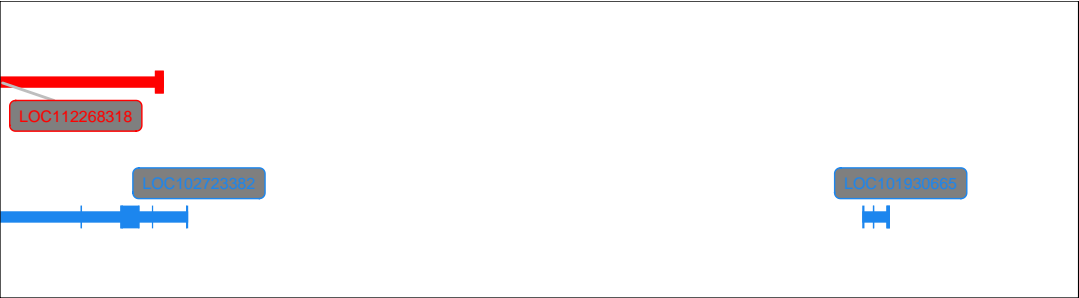
L5.PCP4



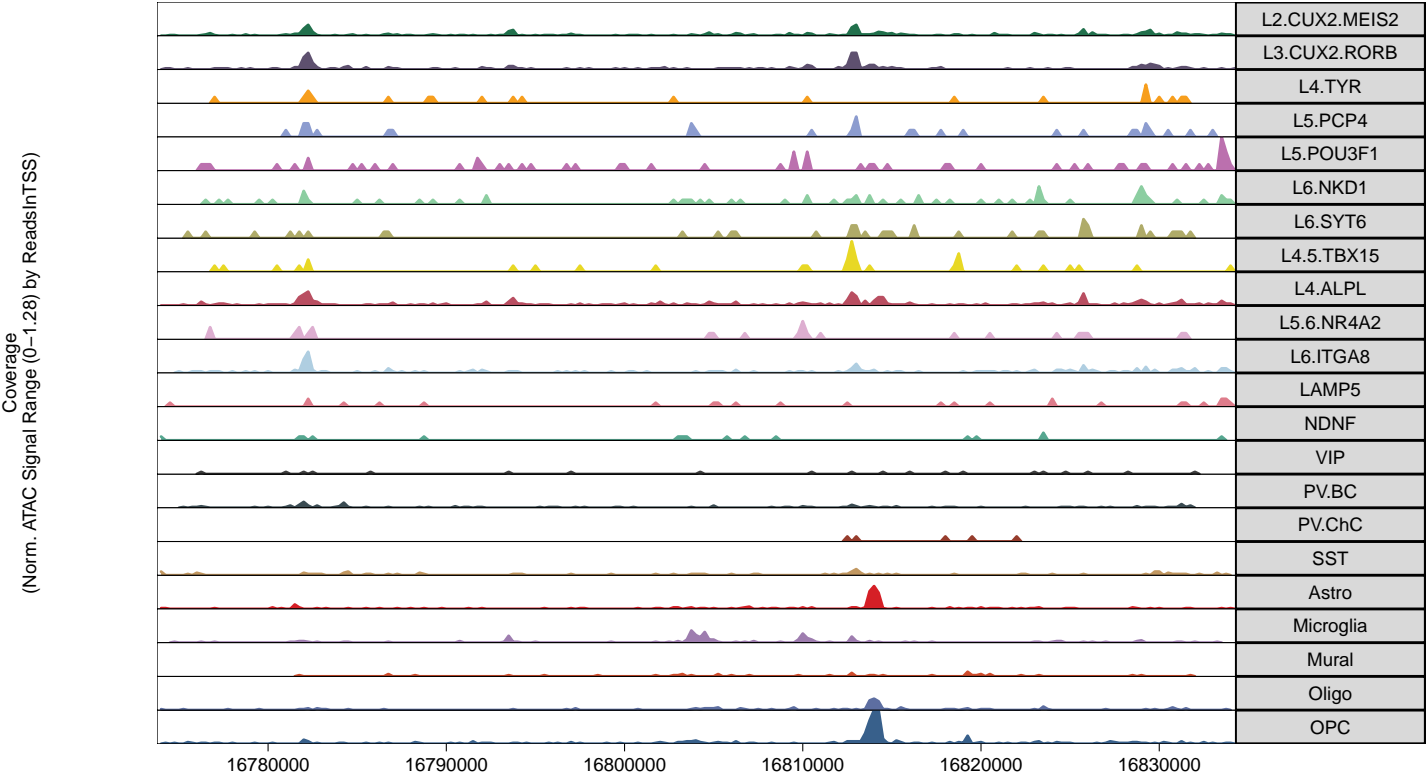
Peaks



LoopTrack



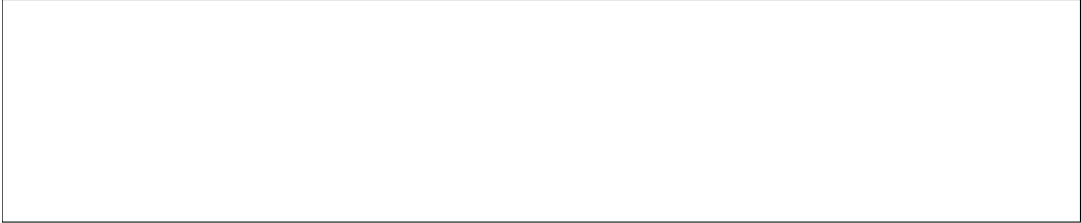
Genes



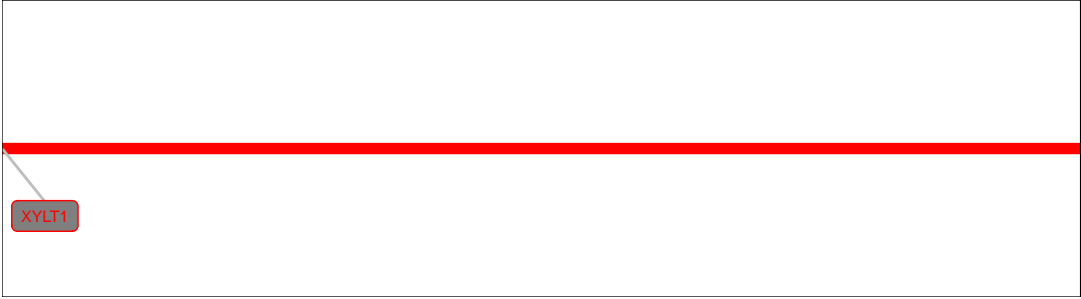
L5.PCP4



Peaks

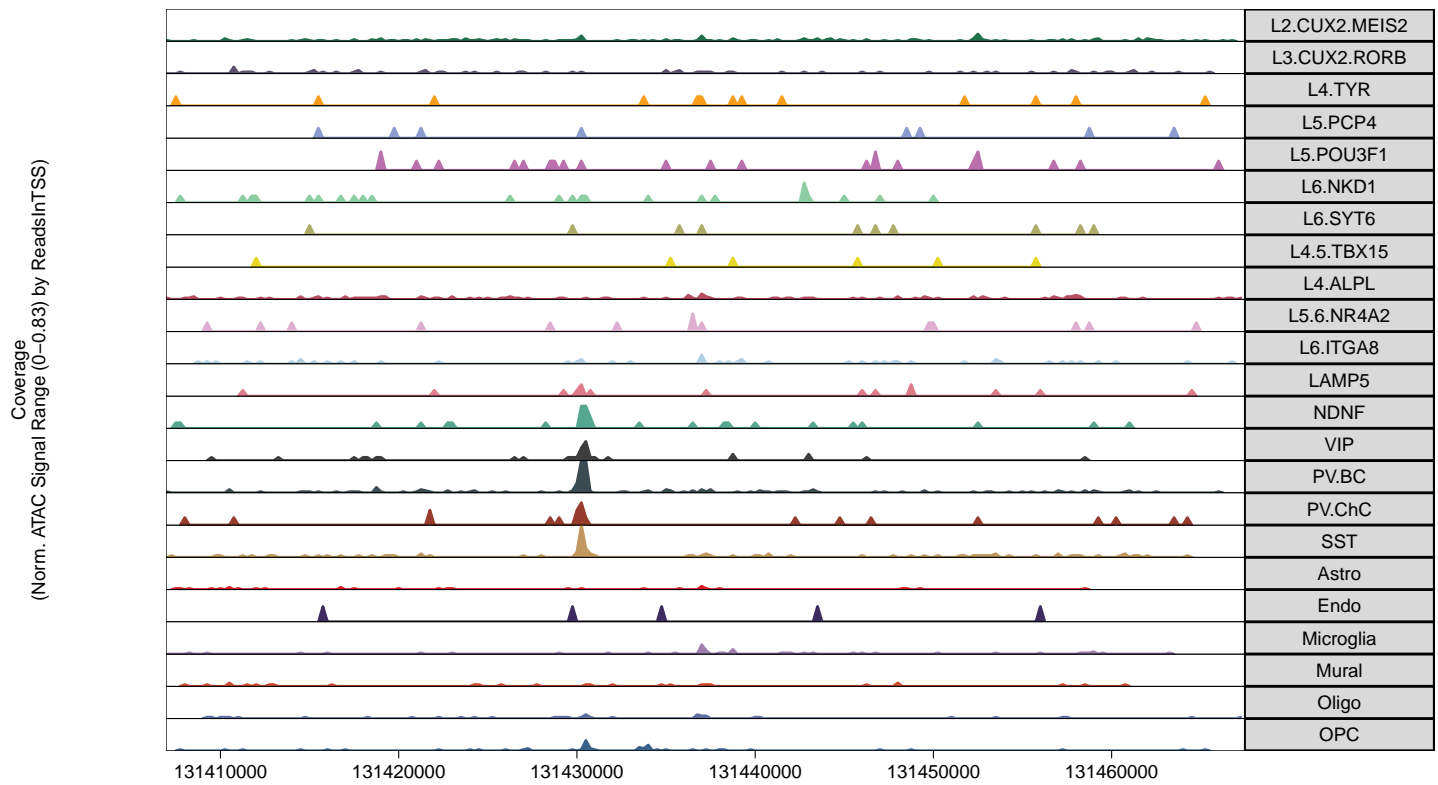


LoopTrack



Genes

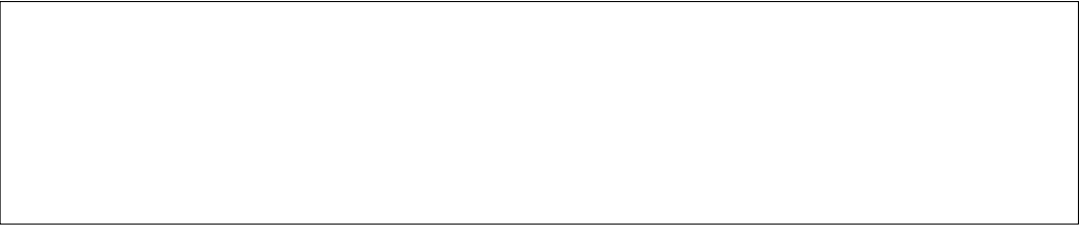
chr1:131406950–131467451 Marker Peak Tracks for: L5.PCP4



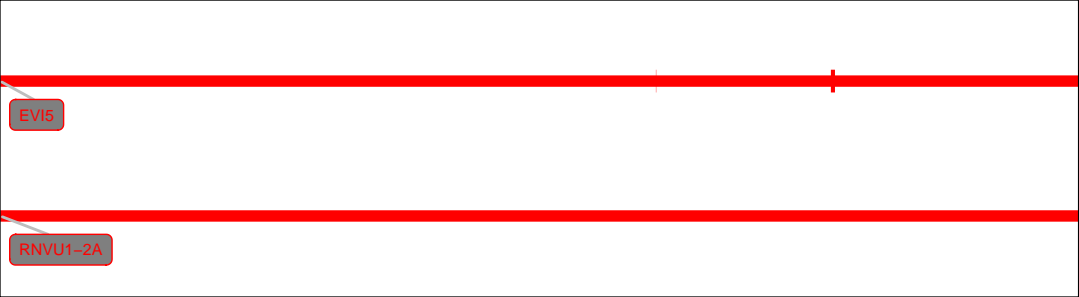
L5.PCP4



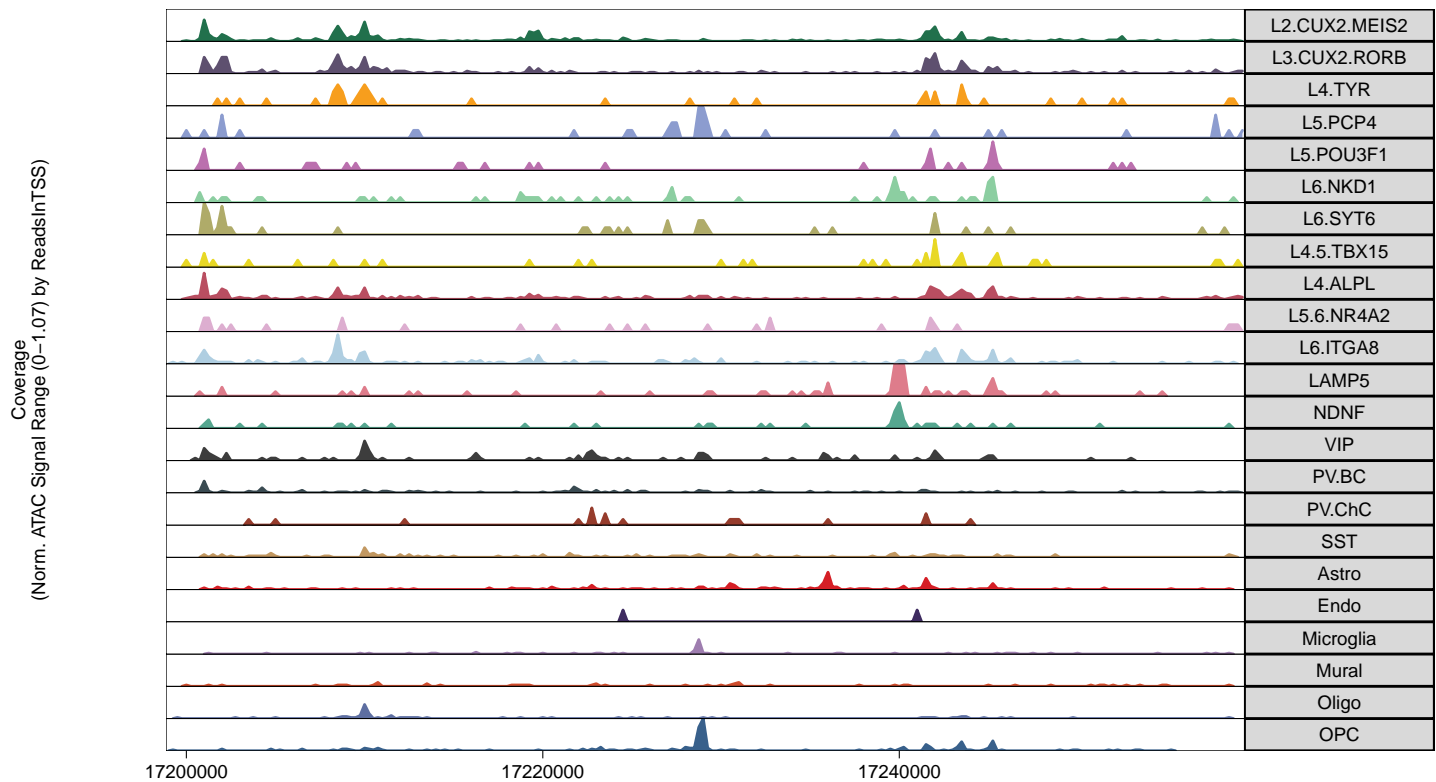
Peaks



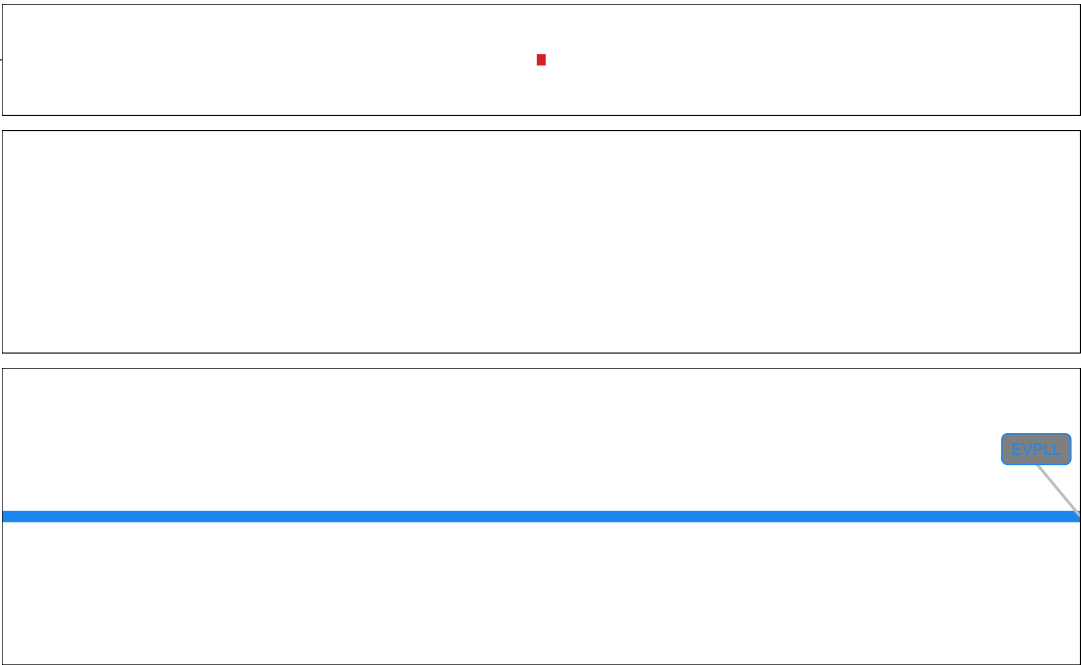
LoopTrack



Genes



L5.PCP4

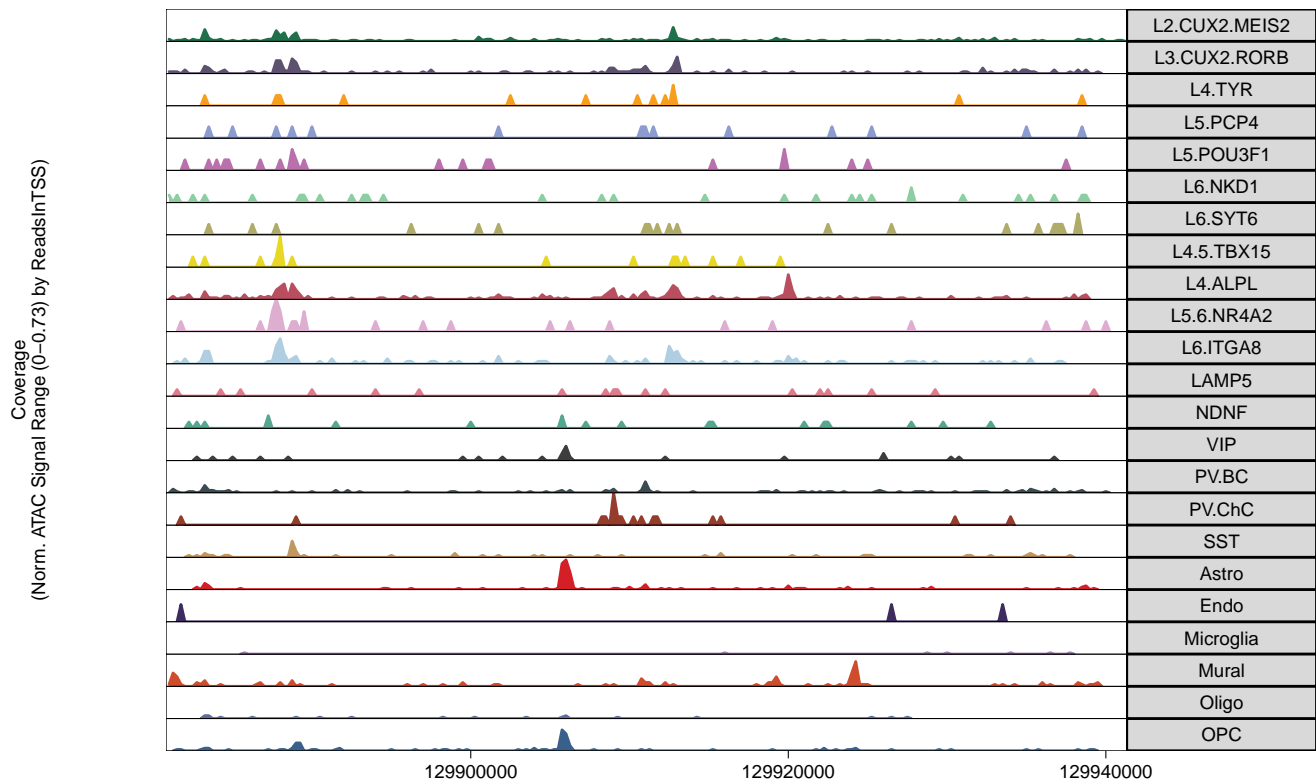


Peaks

LoopTrack

Genes

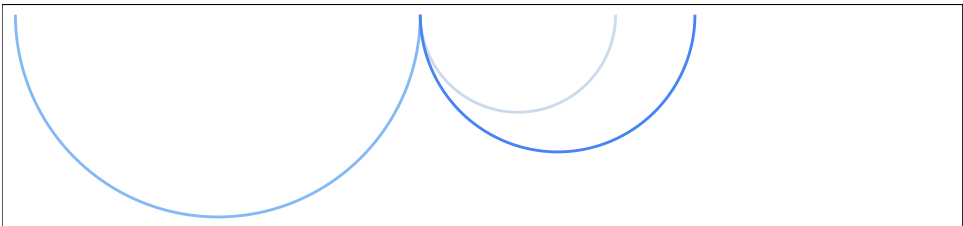
chr11:129880813–129941314 Marker Peak Tracks for: L5.PCP4



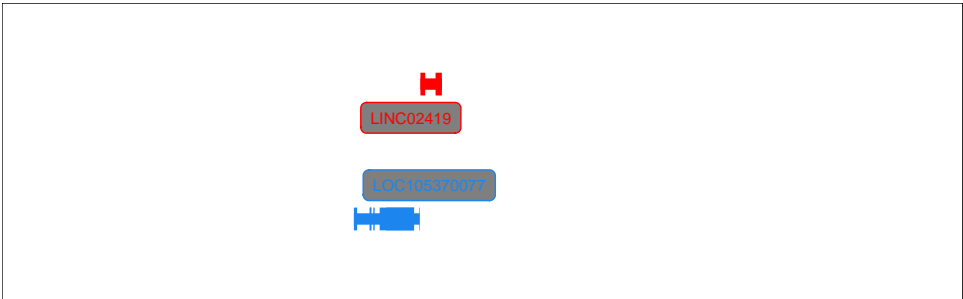
L5.PCP4



Peaks



Peak2GeneLinks

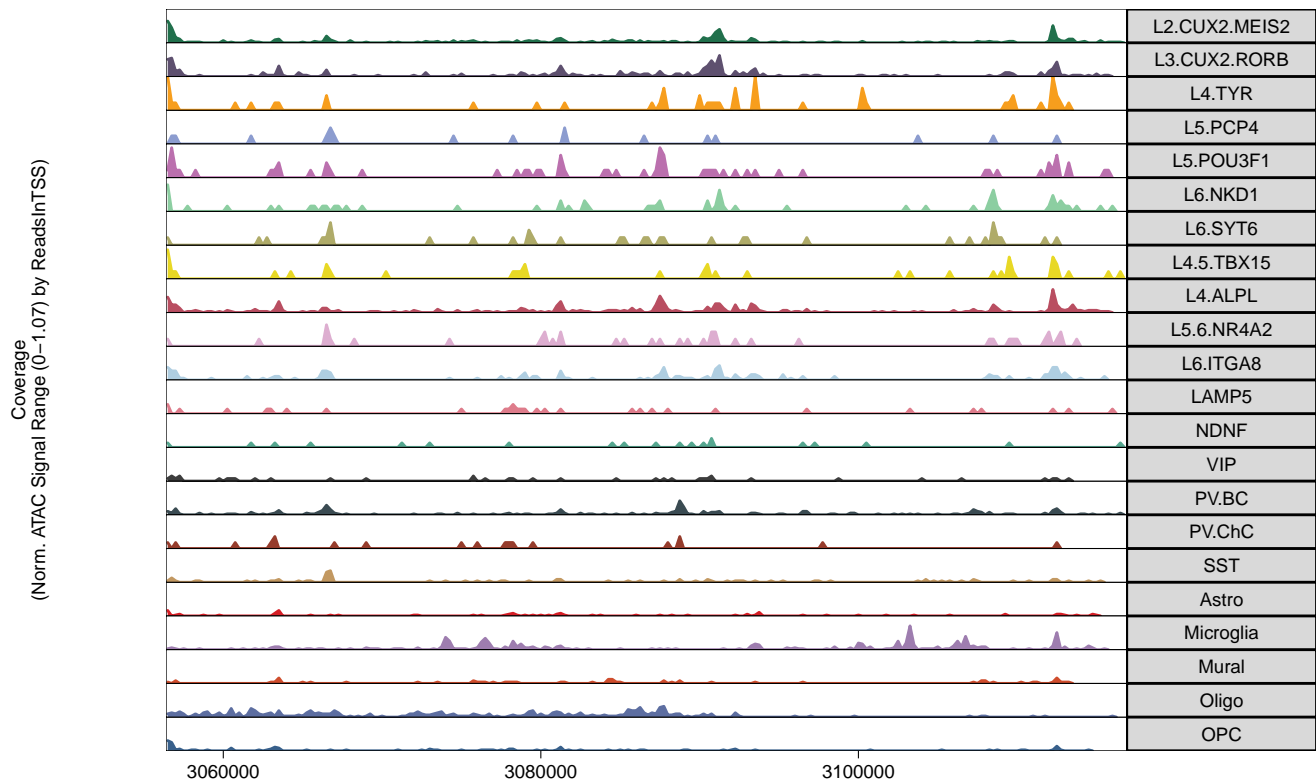


Genes

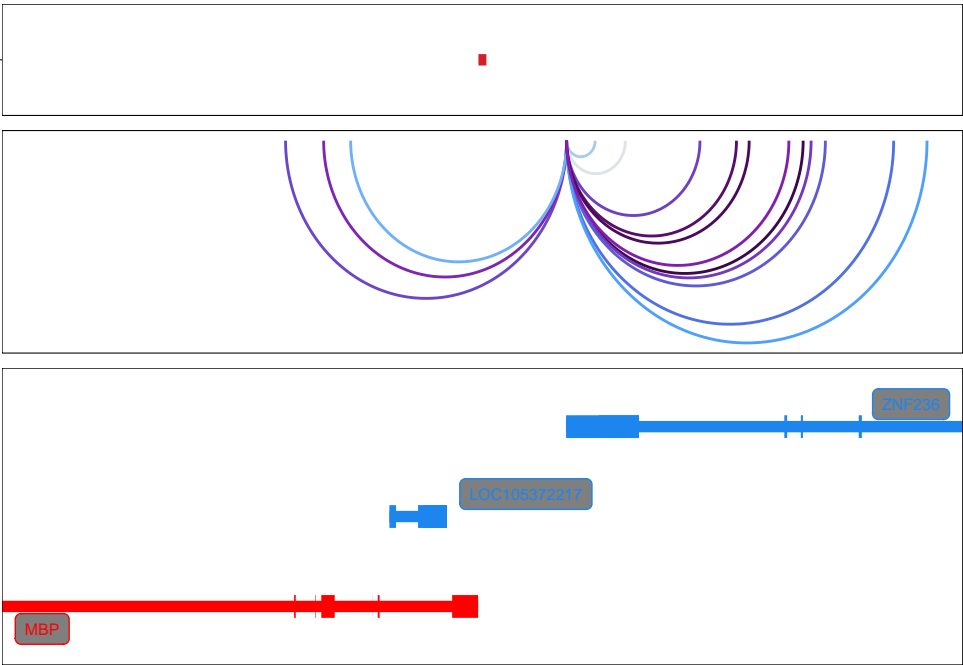
value



chr18:3056398–3116899 Marker Peak Tracks for: L5.PCP4



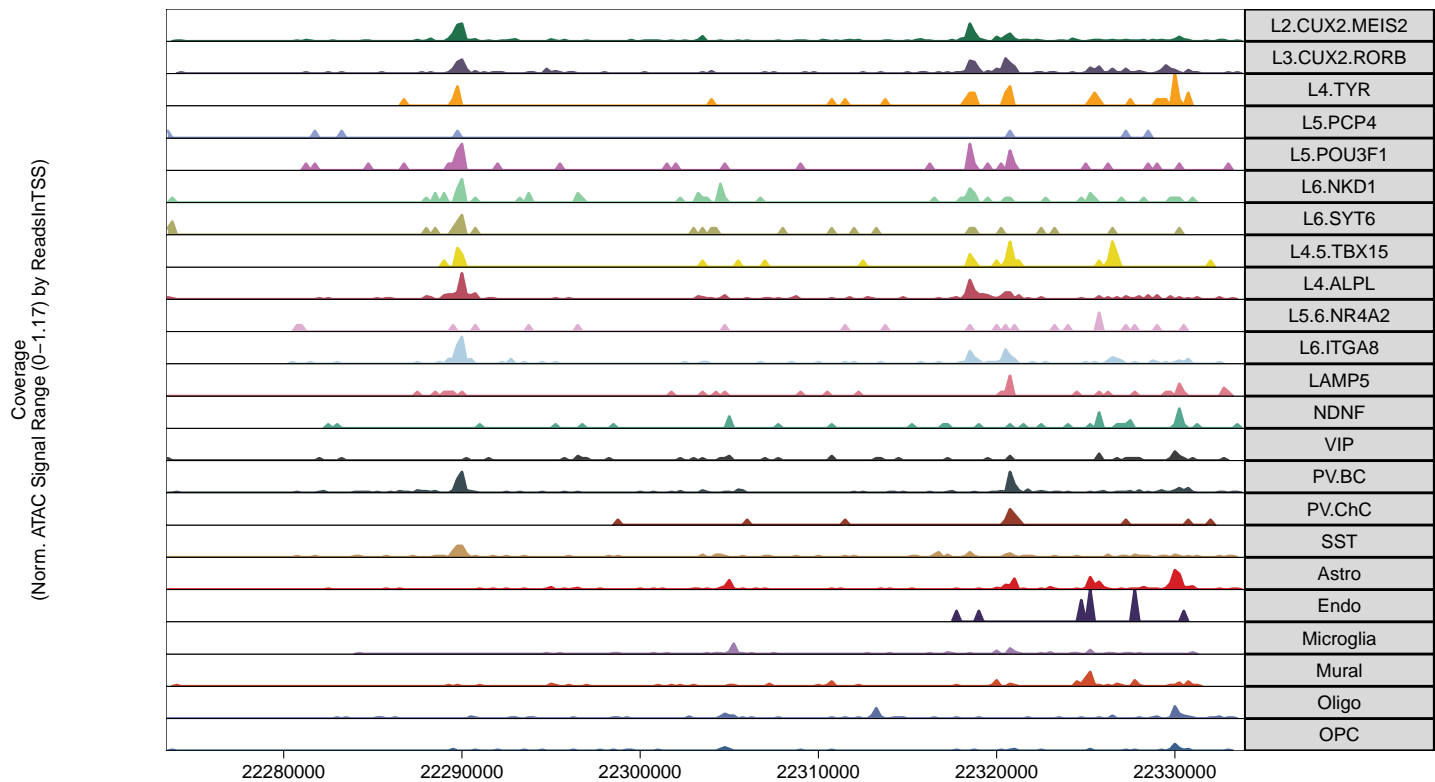
L5.PCP4



value



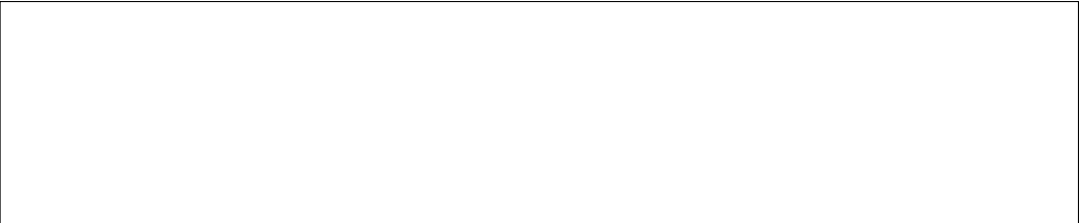
chr15:22273402–22333903 Marker Peak Tracks for: L5.PCP4



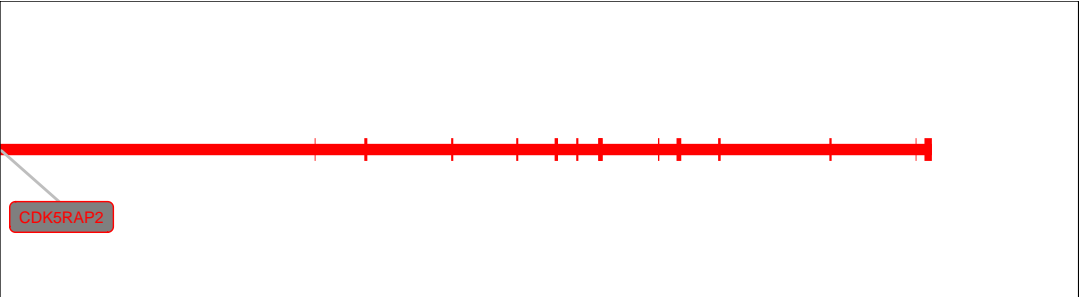
L5.PCP4



Peaks

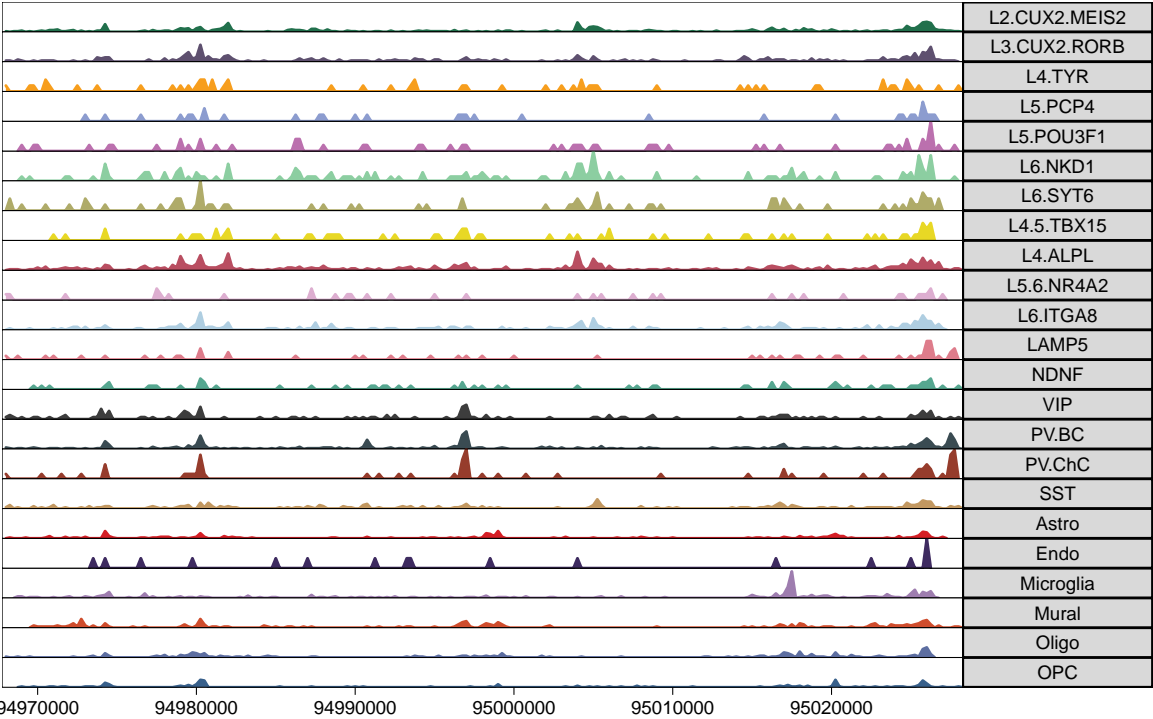


LoopTrack



Genes

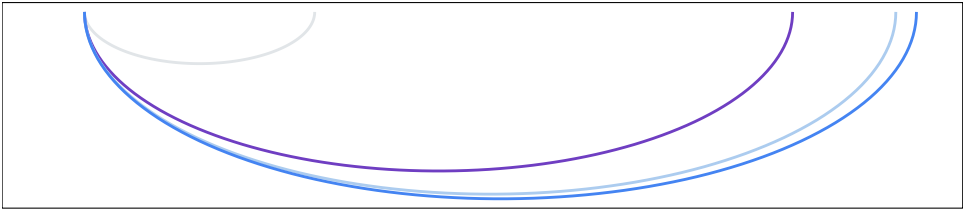
Coverage
(Norm. ATAC Signal Range (0–1.8) by Reads in TSS)



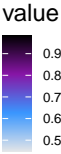
L5.PCP4



Peaks



Peak2GeneLinks

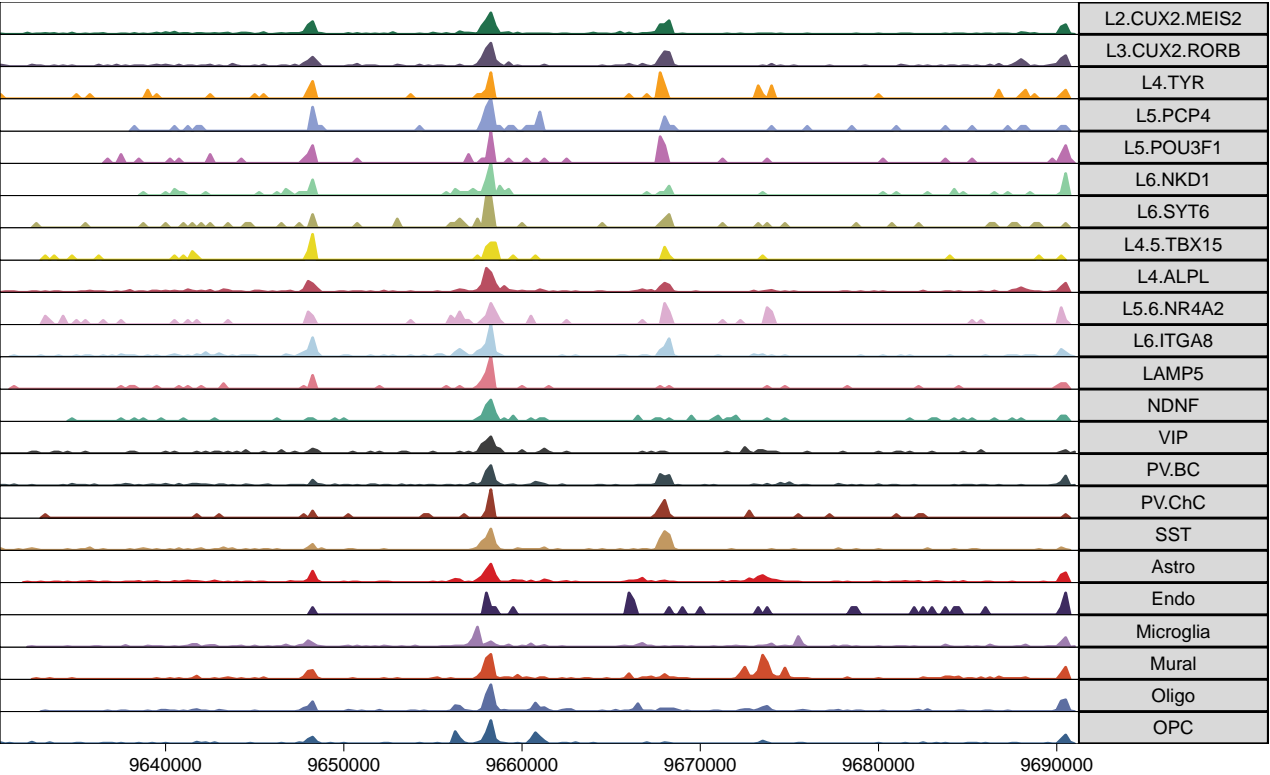


GeneTrack

strand • •

chr14:9630716-9691217 Marker Peak Tracks for: L5.PCP4

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



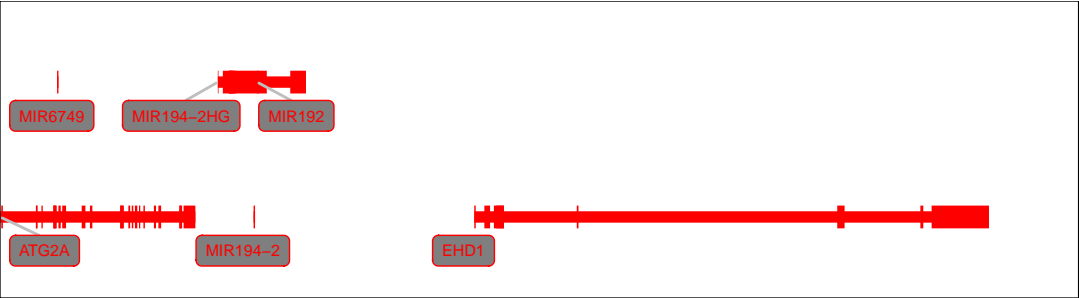
L5.PCP4



Peaks



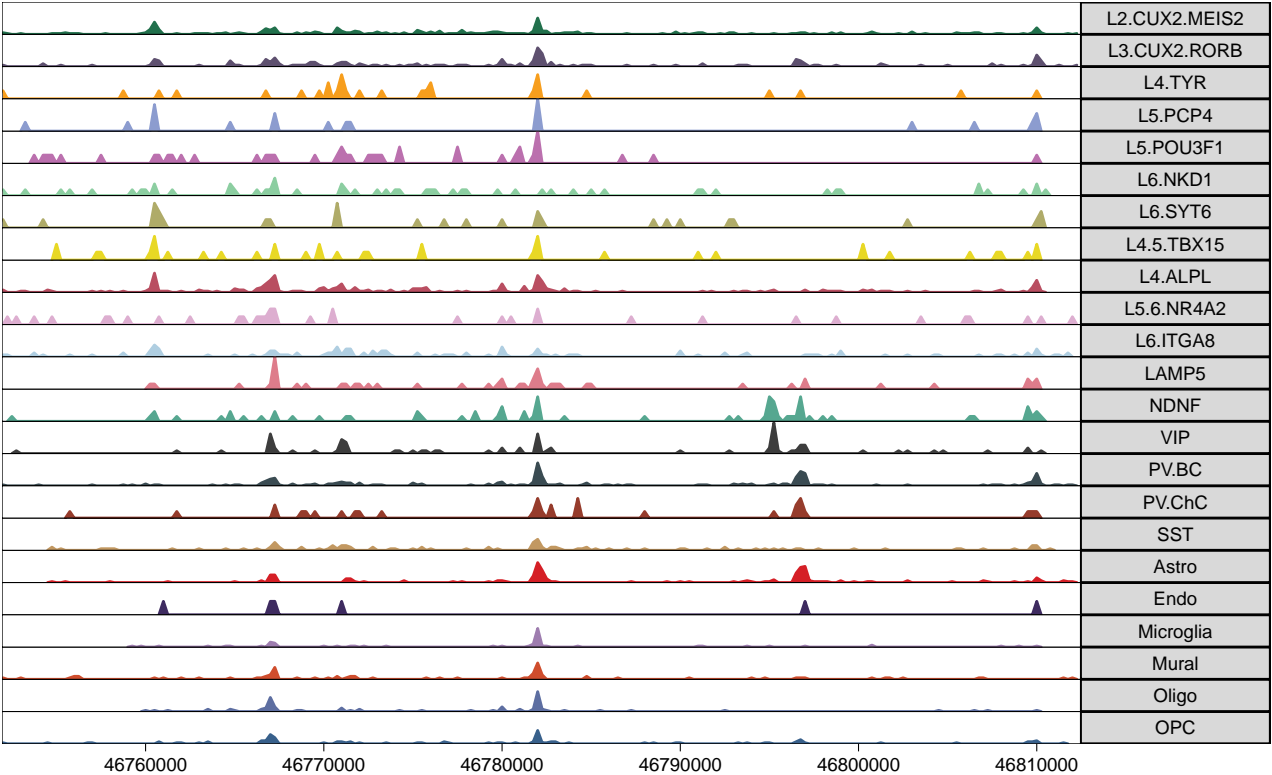
LoopTrack



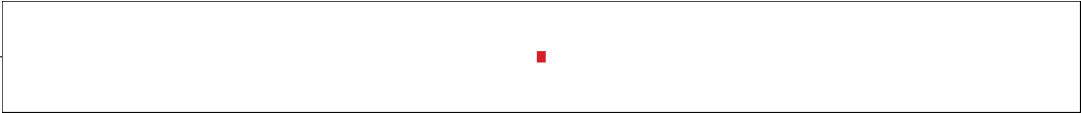
Genes

chr3:46751950–46812451 Marker Peak Tracks for: L5.PCP4

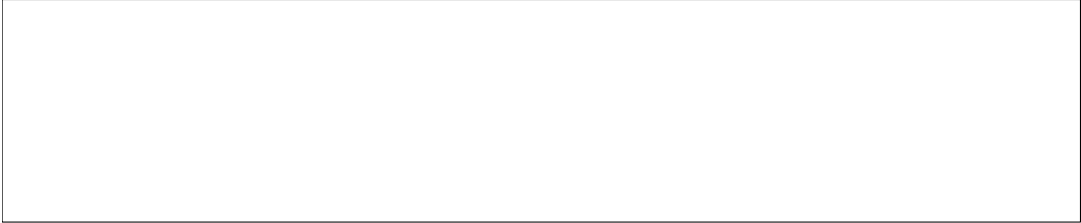
Coverage
(Norm. ATAC Signal Range (0–0.94) by ReadsInTSS)



L5.PCP4



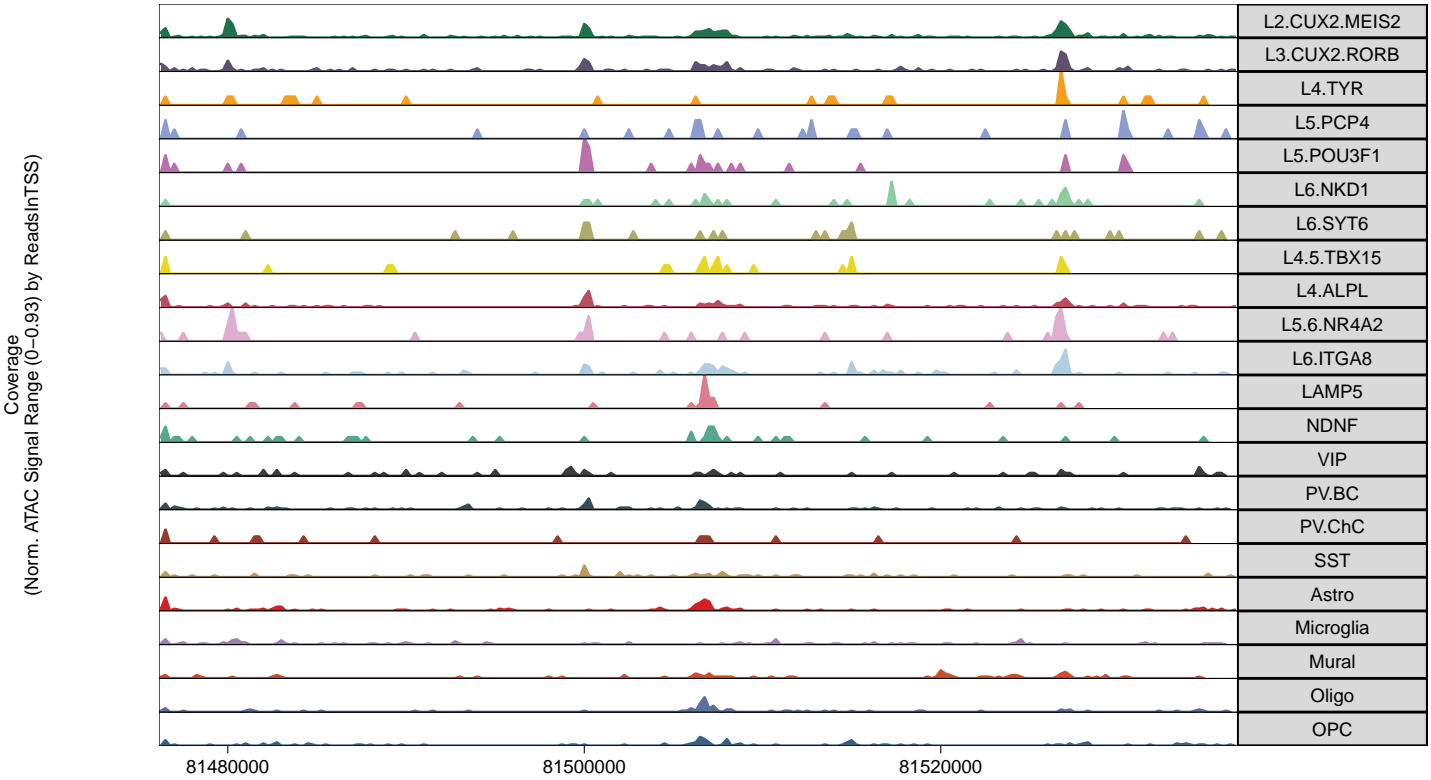
Peaks



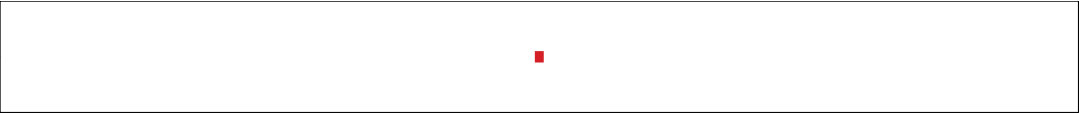
LoopTrack



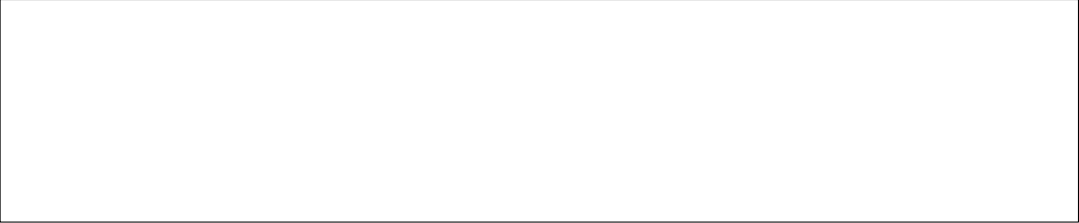
Genes



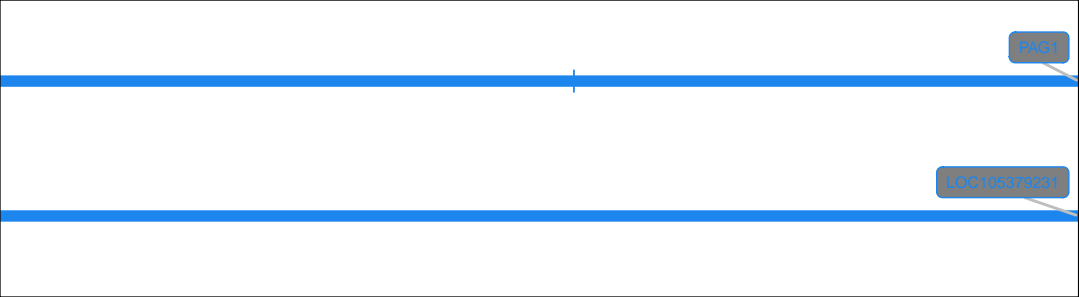
L5.PCP4



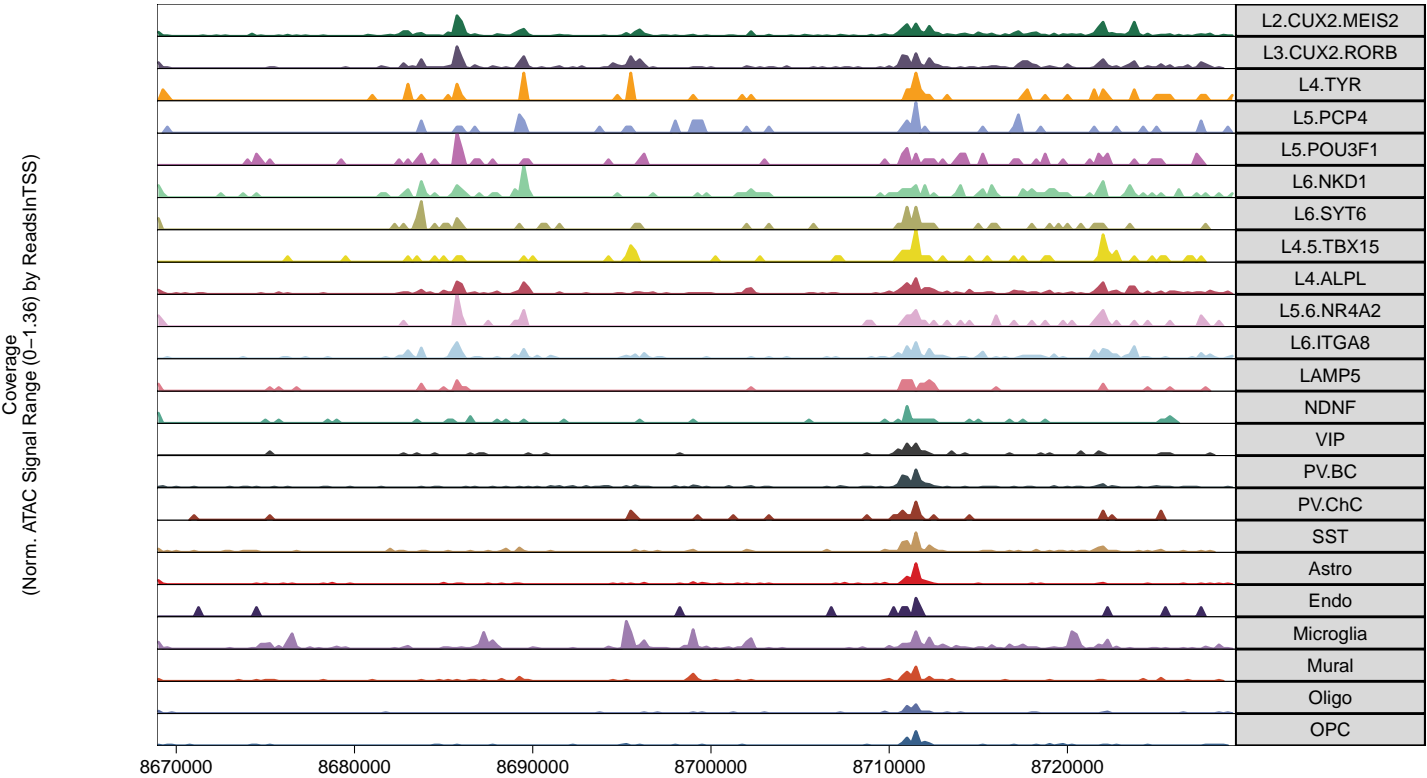
Peaks



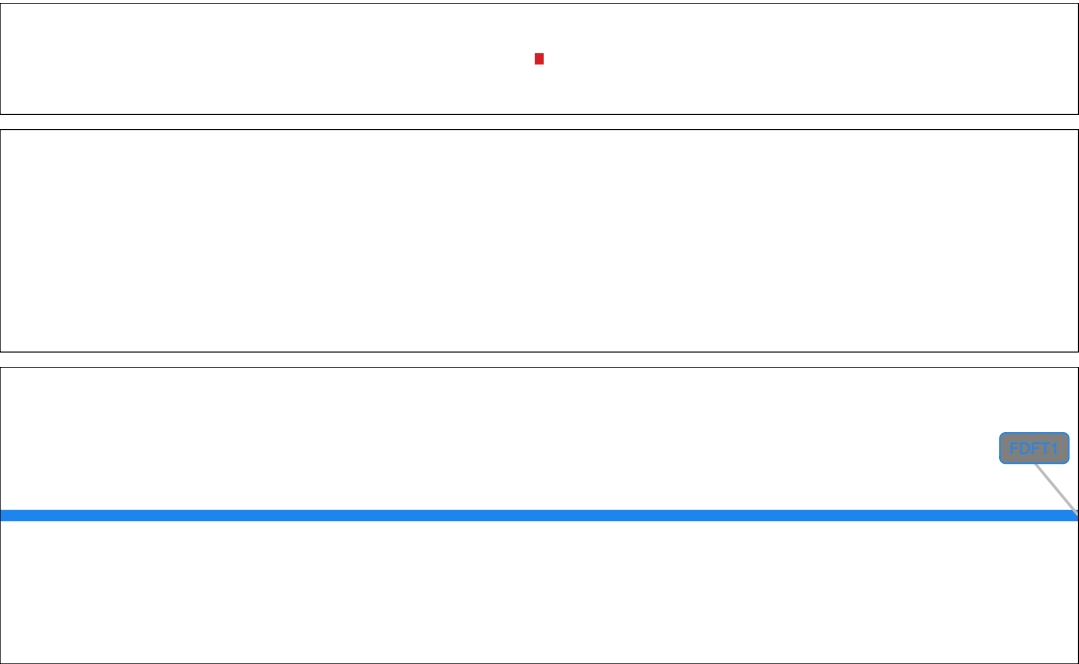
LoopTrack



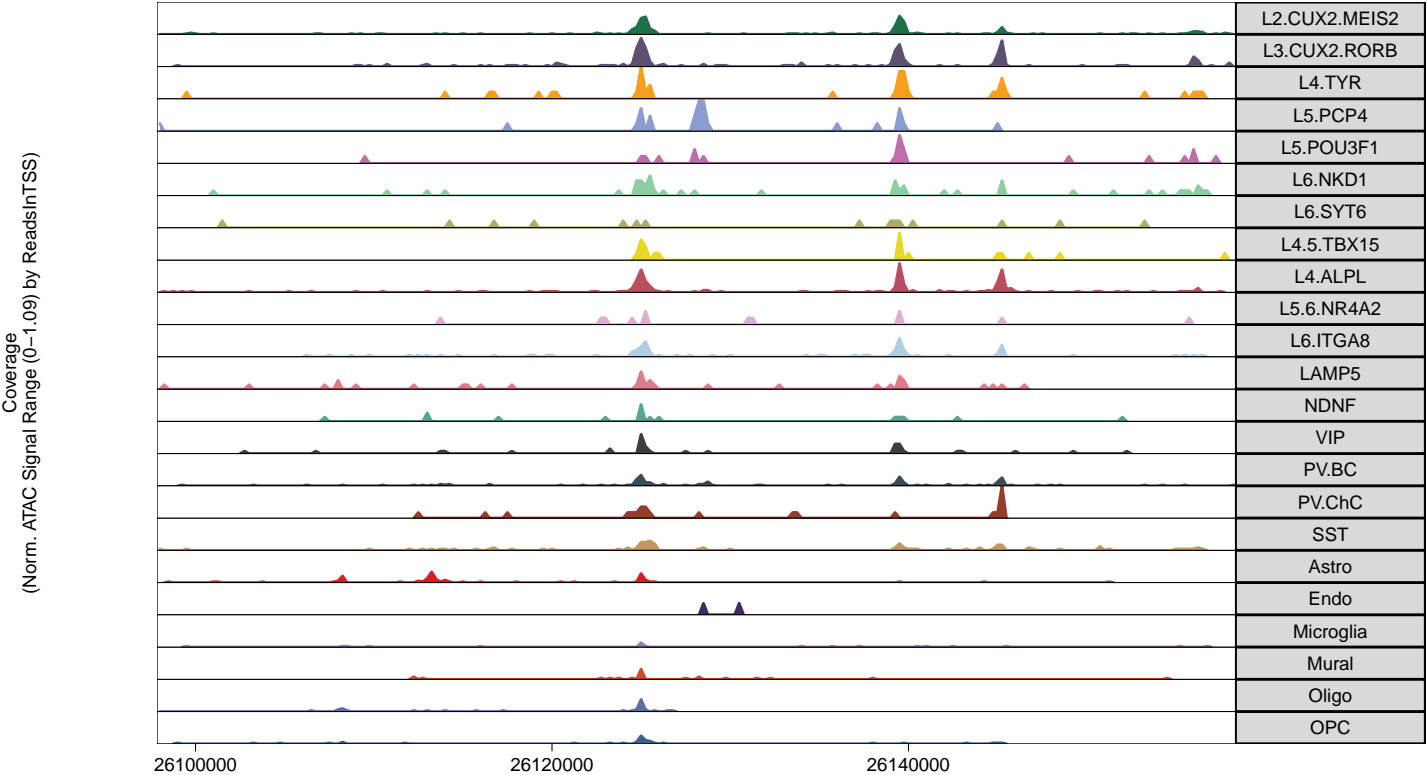
Genes



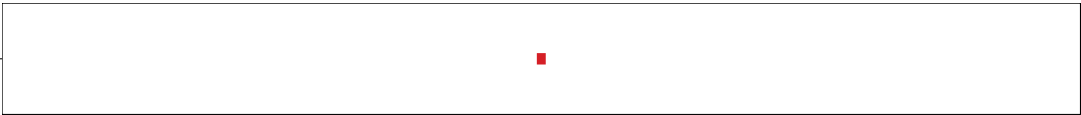
L5.PCP4



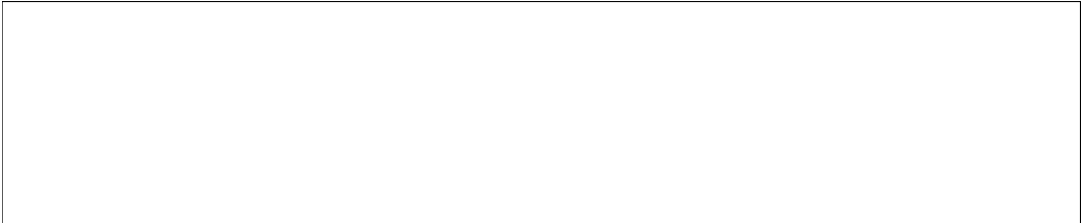
chr9:26097845-26158346 Marker Peak Tracks for: L5.PCP4



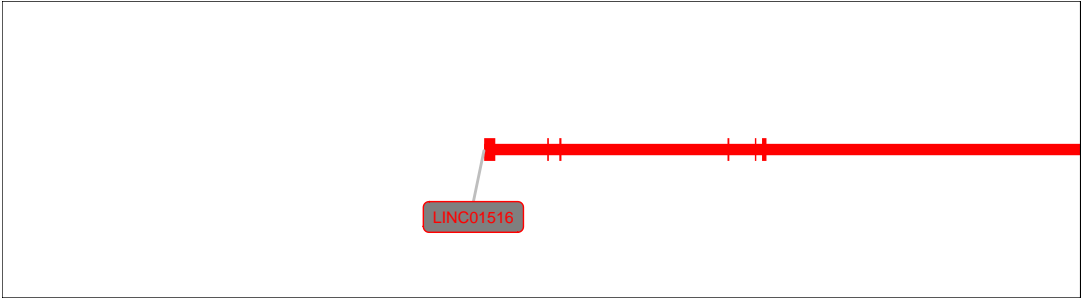
L5.PCP4



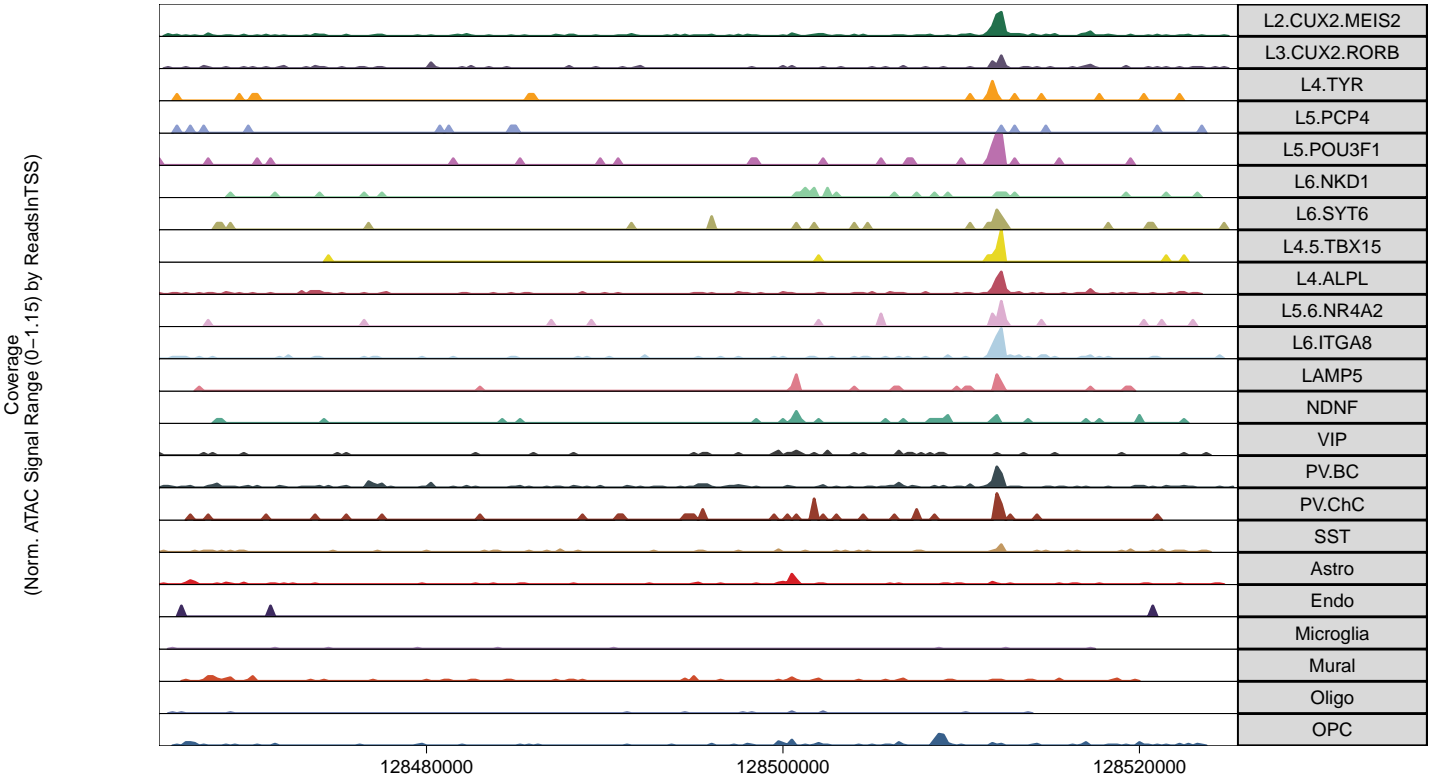
Peaks



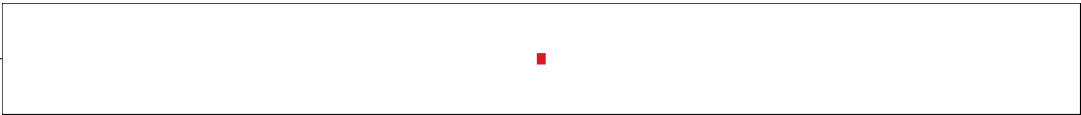
LoopTrack



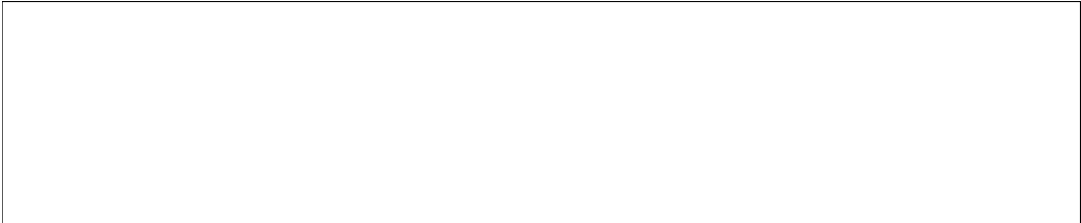
Genes



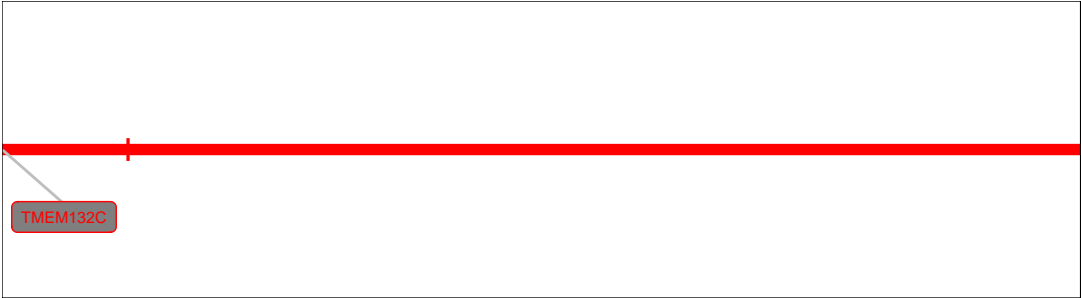
L5.PCP4



Peaks



LoopTrack



Genes