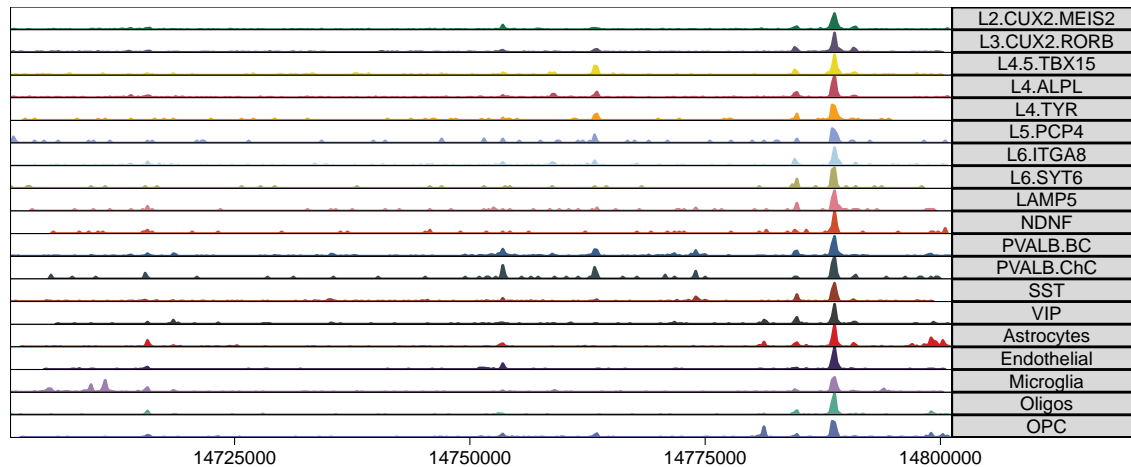


Coverage
(Norm. ATAC Signal Range (0-2.08) by Reads/nTSS)

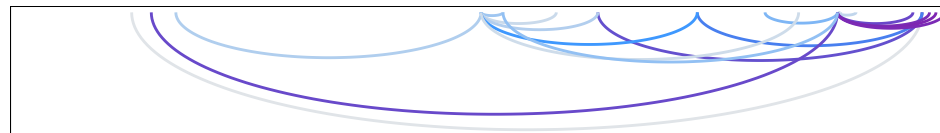
chr10:14701195-14801196



PVALB.BC



Peaks



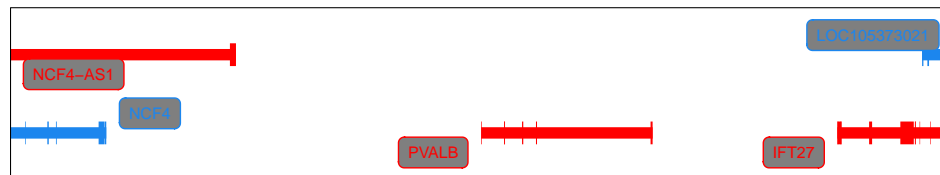
Peak2GeneLinks

value

0.8

0.6

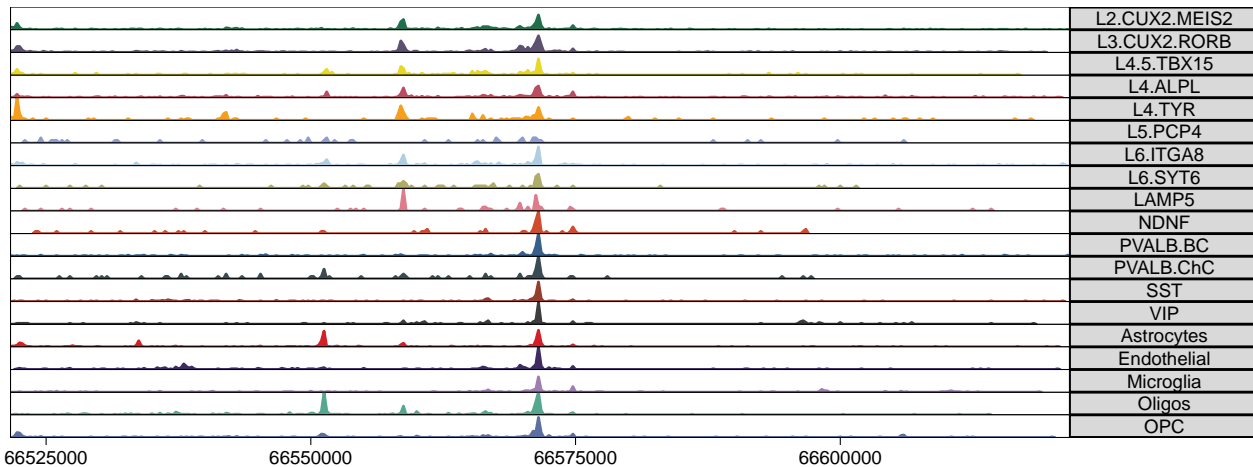
0.4



Genes

Coverage
(Norm. ATAC Signal Range (0–1.66) by Reads/nTSS)

chr9:66521641–66621642



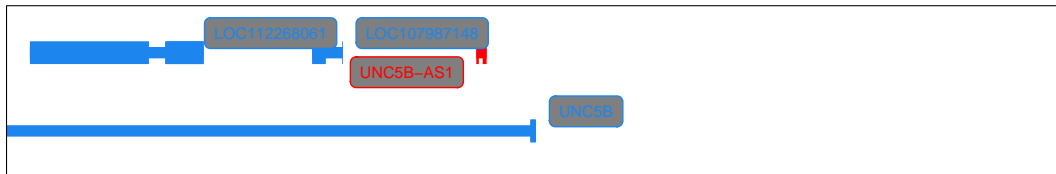
PVALB.BC



Peaks



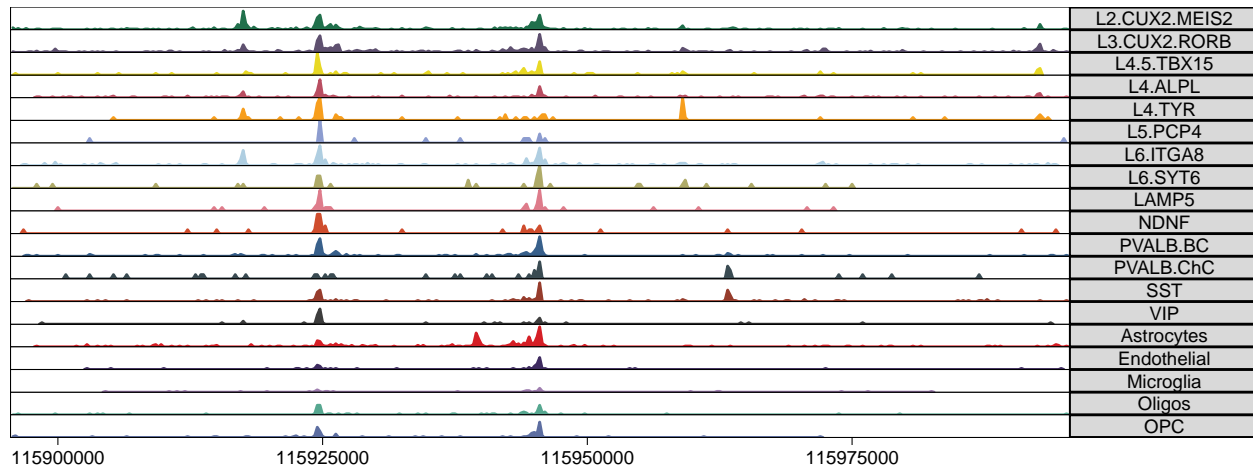
LoopTrack



Genes

Coverage
(Norm. ATAC Signal Range (0-0.94) by Reads/nTSS)

chr8:115895528-115995529



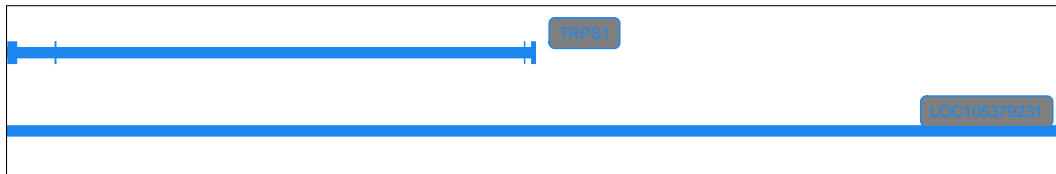
PVALB.BC



Peaks

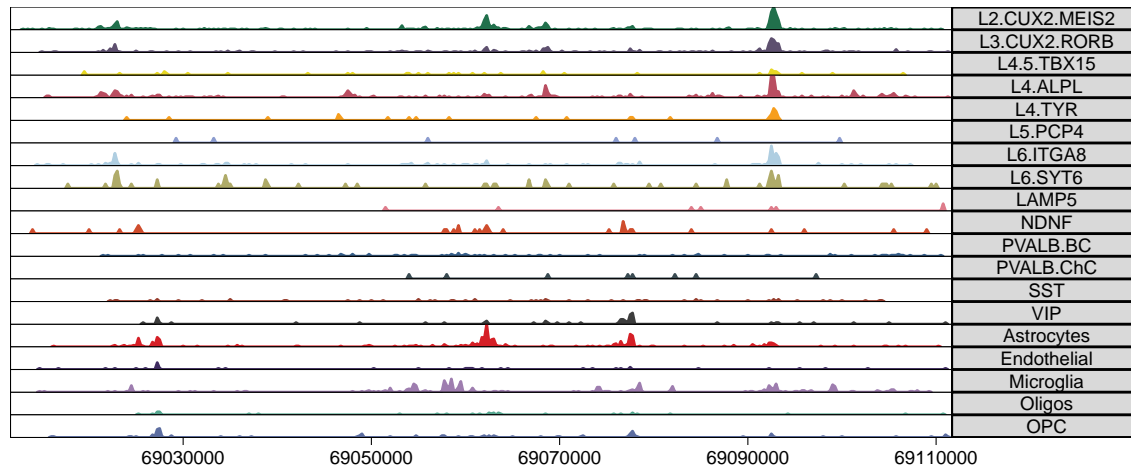


LoopTrack



Genes

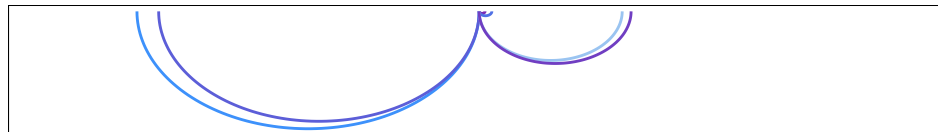
chr15:69011659-69111660

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

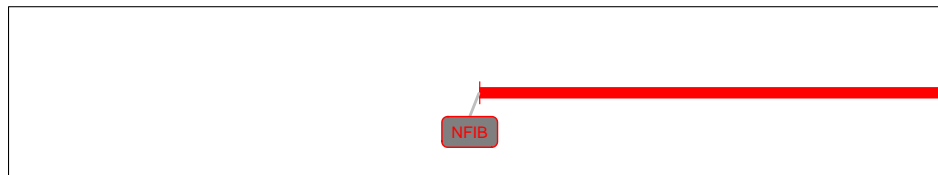
PVALB.BC



Peaks



Peak2GeneLinks



Genes

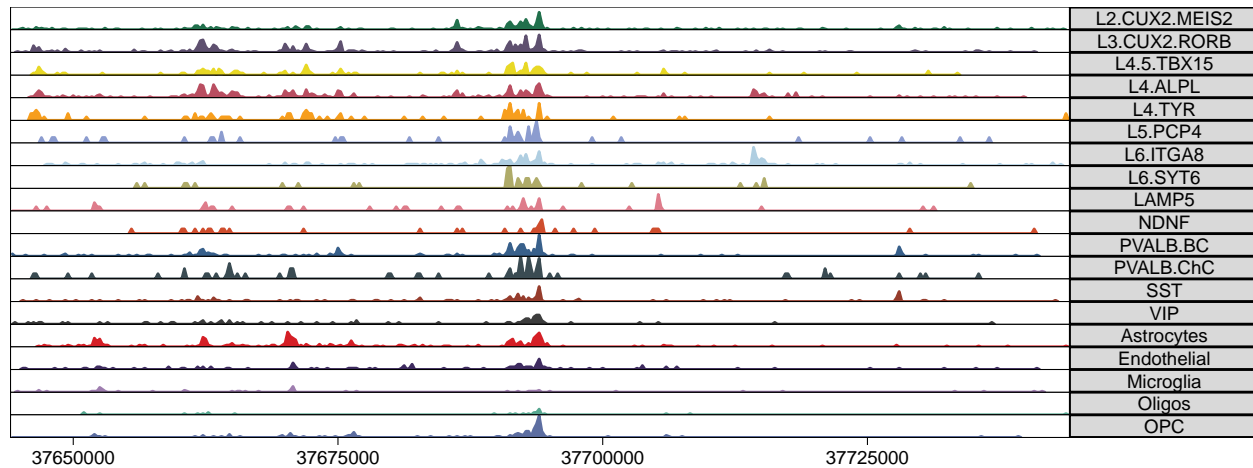
value

0.8

0.6

0.4

chr7:37644076-37744077

Coverage
(Norm. ATAC Signal Range (0-0.79) by Reads/nTSS)

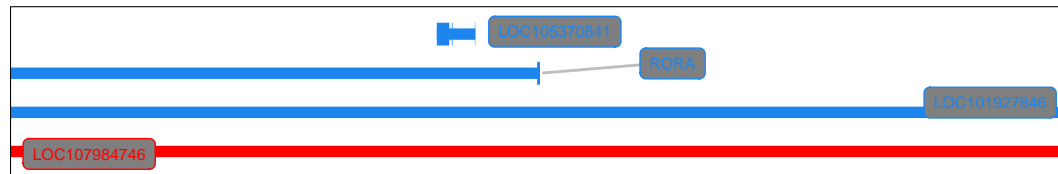
PVALB.BC



Peaks

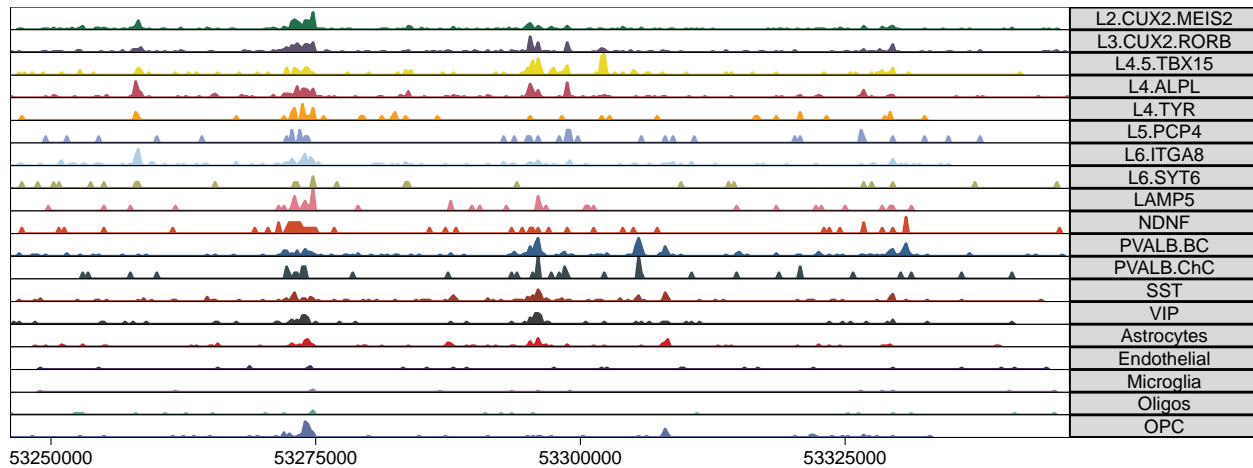


LoopTrack



Genes

chr12:53246181-53346182

Coverage
(Norm. ATAC Signal Range (0-0.66) by Reads/nTSS)

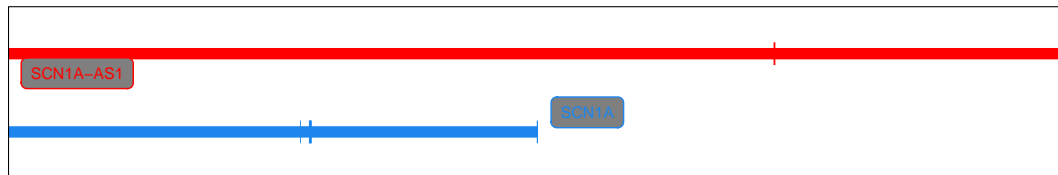
PVALB.BC



Peaks



LoopTrack



Genes

