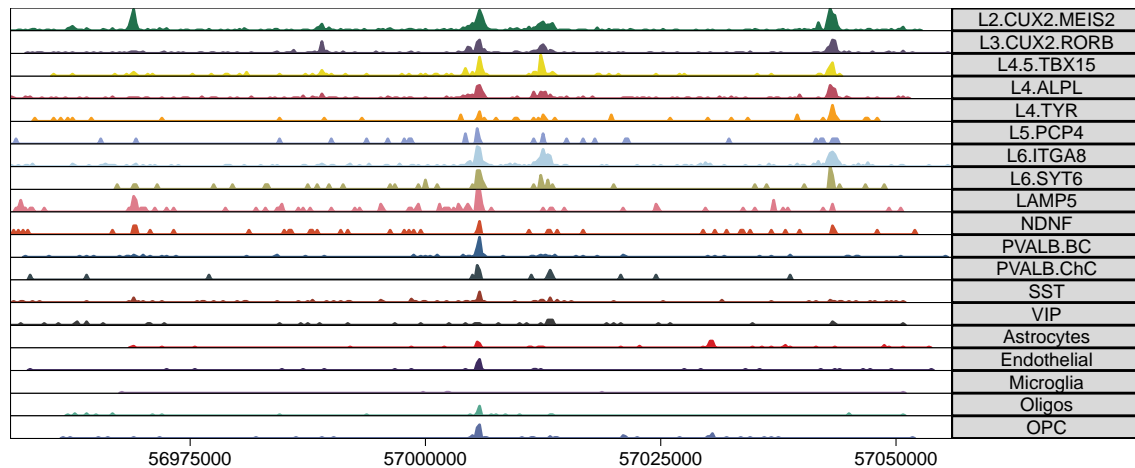
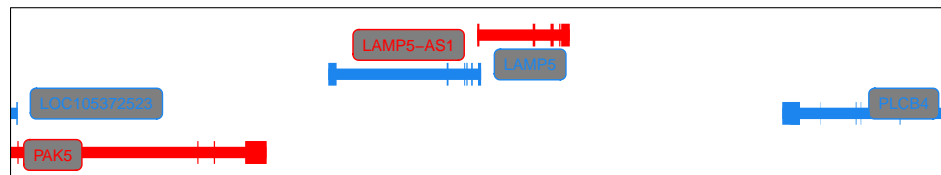
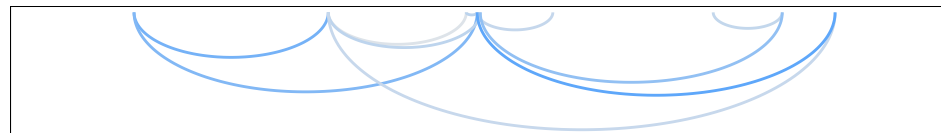
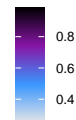


chr10:56955915-57055916

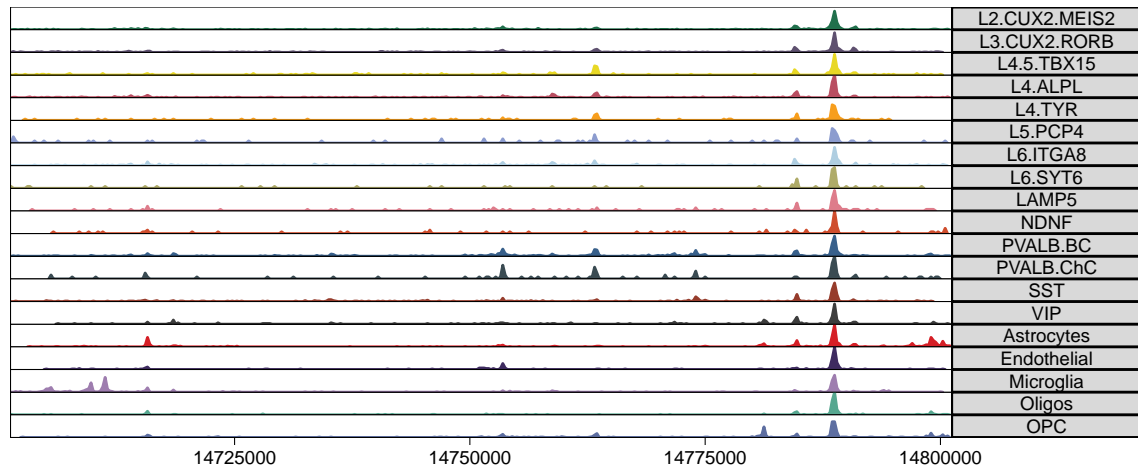
Coverage  
(Norm. ATAC Signal Range (0-0.83) by Reads/nTSS)LAMP5  
PVALB.BC  
SST  
VIP

value



chr10:14701195-14801196

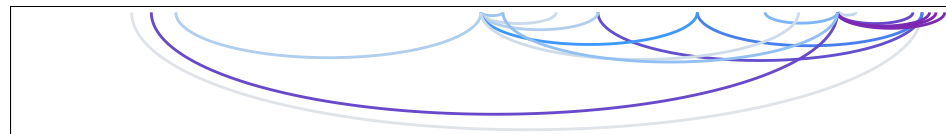
Coverage  
(Norm. ATAC Signal Range (0-2.06) by Reads/nTSS)



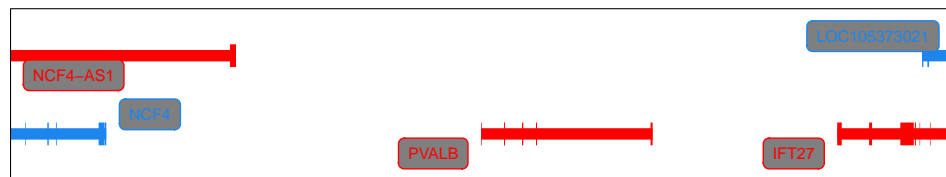
LAMP5  
PVALB.BC  
SST  
VIP



Peaks



Peak2GeneLinks

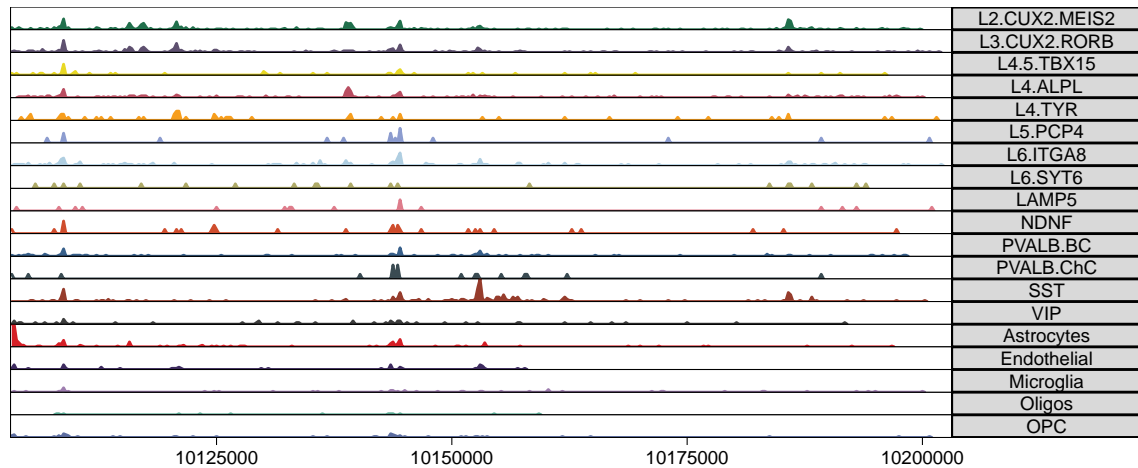


Genes

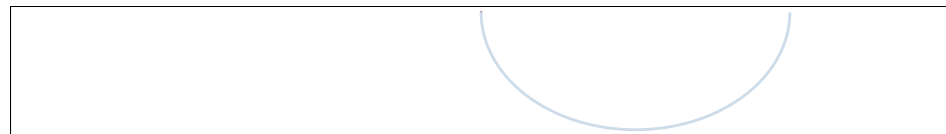
value



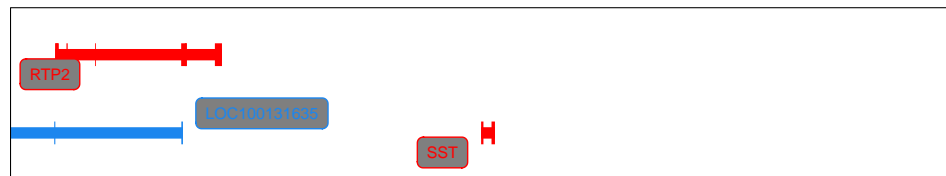
chr2:10103111-10203112

Coverage  
(Norm. ATAC Signal Range (0-0.87) by Reads/nTSS)LAMP5  
PVALB.BC  
SST  
VIP

Peaks

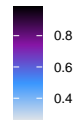


Peak2GeneLinks

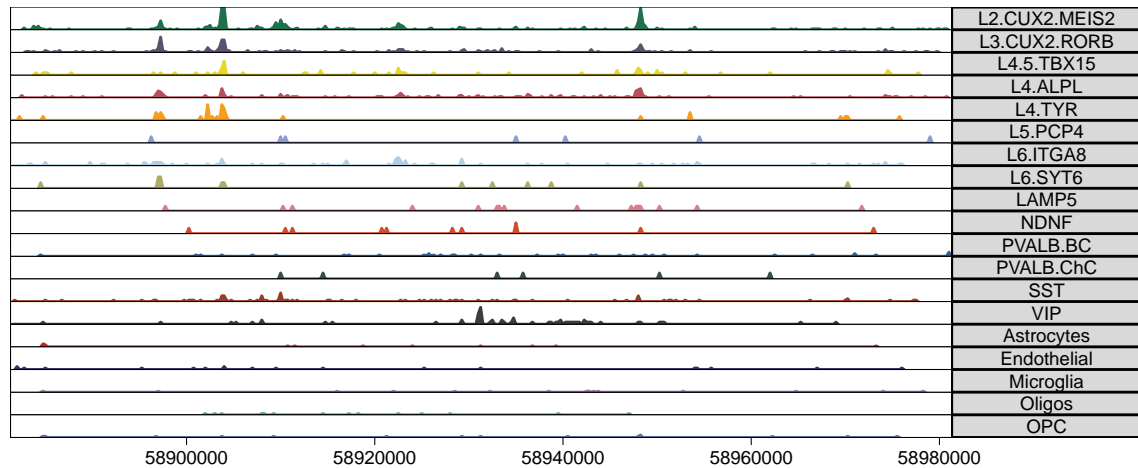


Genes

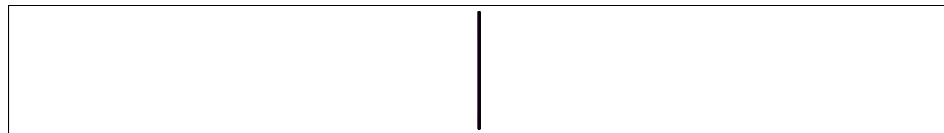
value



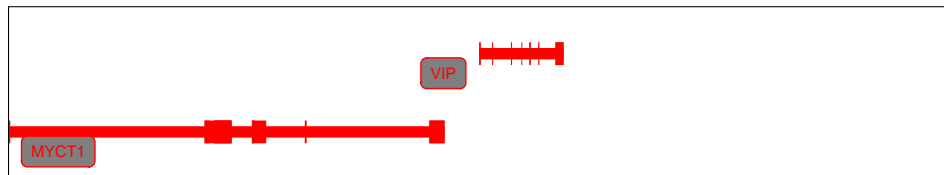
chr4:58881300-58981301

Coverage  
(Norm. ATAC Signal Range (0-0.68) by Reads/nTSS)LAMP5  
PVALB.BC  
SST  
VIP

Peaks

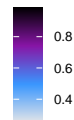


Peak2GeneLinks

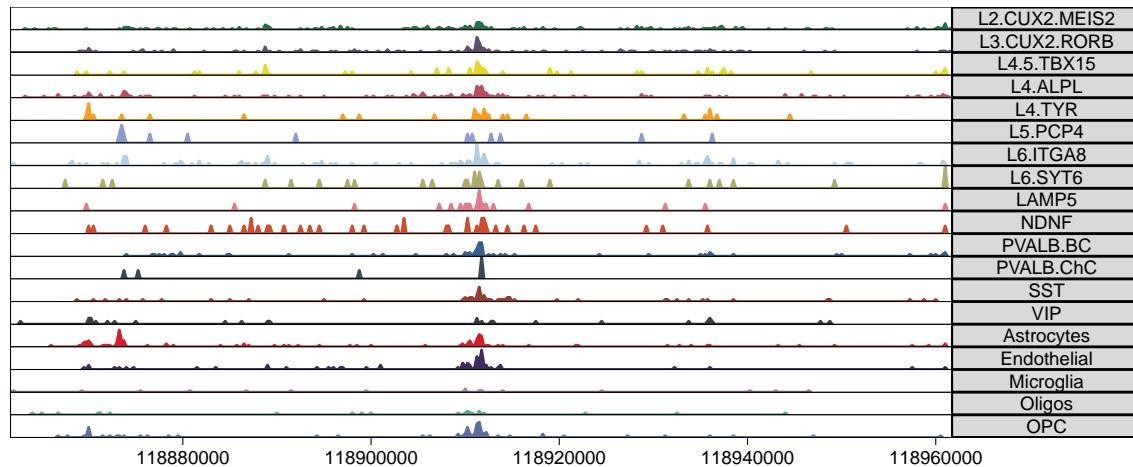


Genes

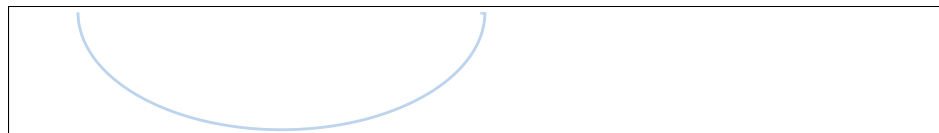
value



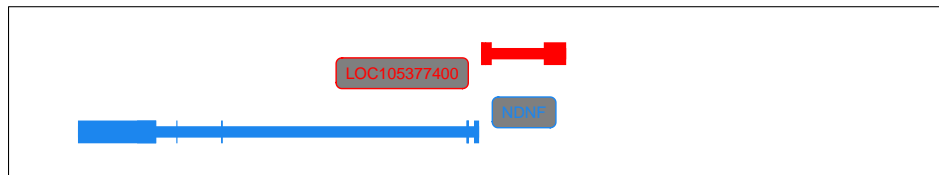
chr5:118861694-118961695

Coverage  
(Norm. ATAC Signal Range (0-0.47) by Reads/nTSS)LAMP5  
PVALB.BC  
SST  
VIP

Peaks



Peak2GeneLinks



Genes

value

