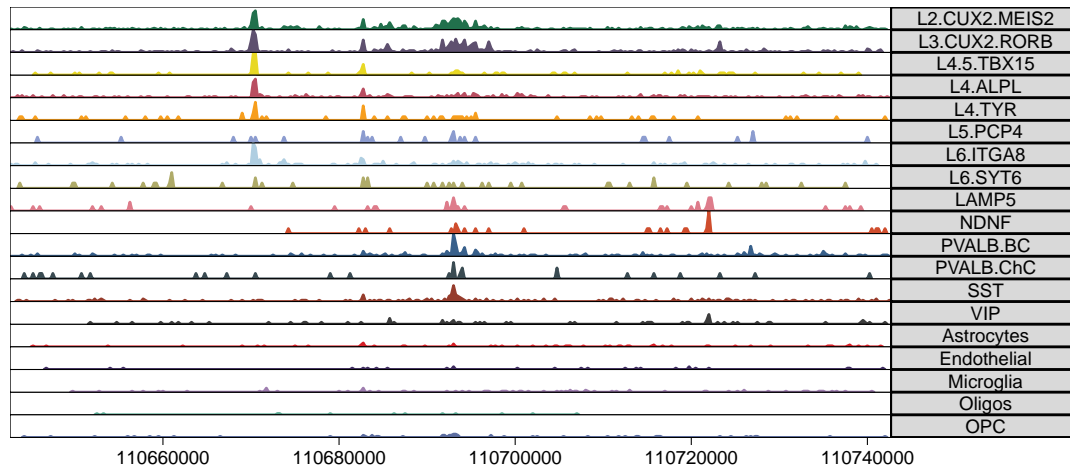


Coverage
(Norm. ATAC Signal Range (0-0.74) by Reads/nTSS)

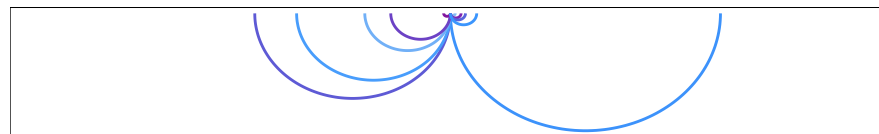
chr11:110642646-110742647



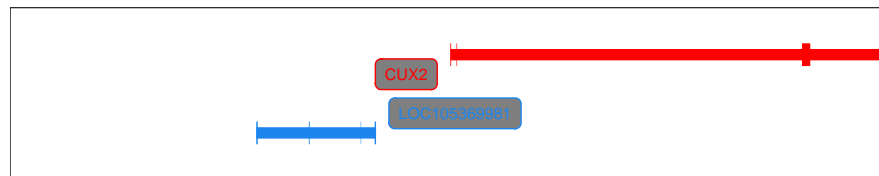
L2.CUX2.MEIS2
L3.CUX2.RORB



Peaks

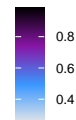


Peak2GeneLinks



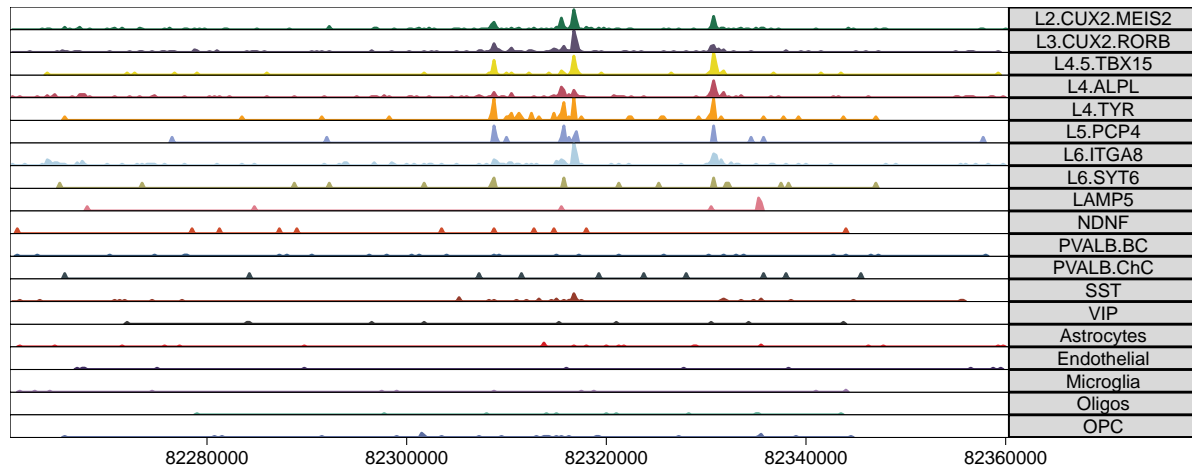
Genes

value



Coverage
(Norm. ATAC Signal Range (0-0.75) by Reads/nTSS)

chr14:82260279-82360280



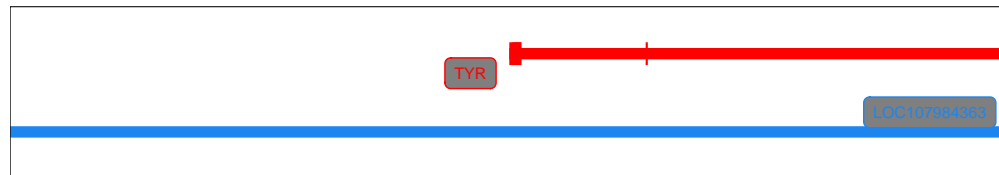
L2.CUX2.MEIS2
L3.CUX2.RORB



Peaks



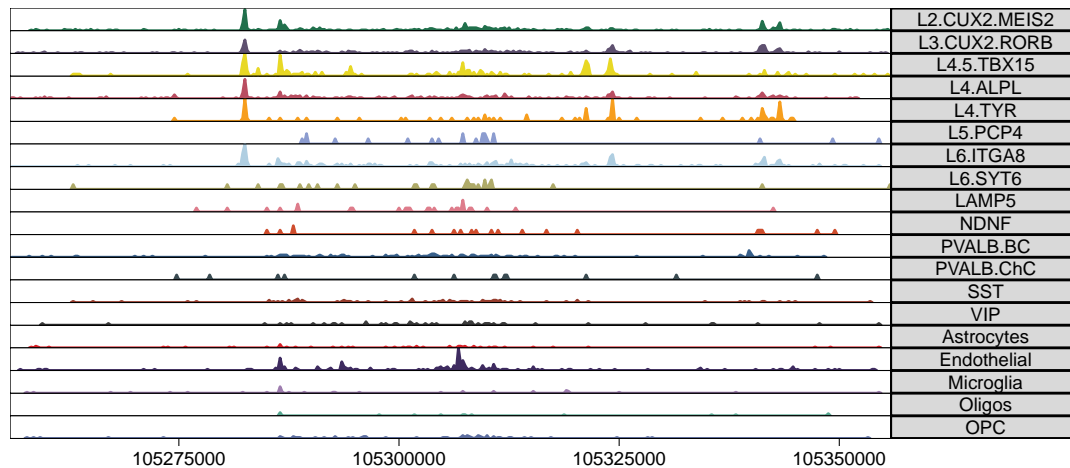
LoopTrack



Genes

Coverage
(Norm. ATAC Signal Range (0-0.82) by Reads/nTSS)

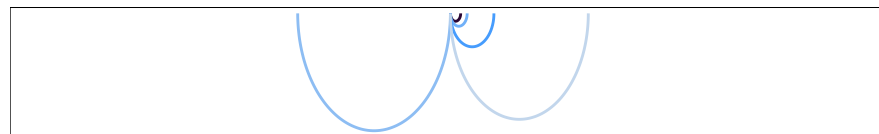
chr1:105255838-105355839



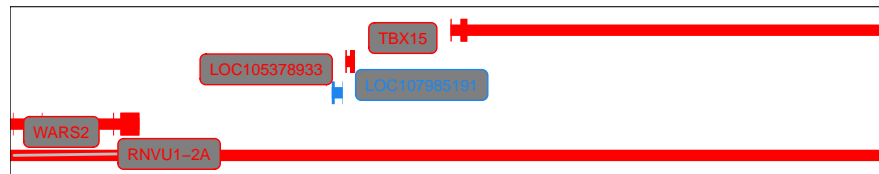
L2.CUX2.MEIS2
L3.CUX2.RORB



Peaks

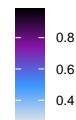


Peak2GeneLinks



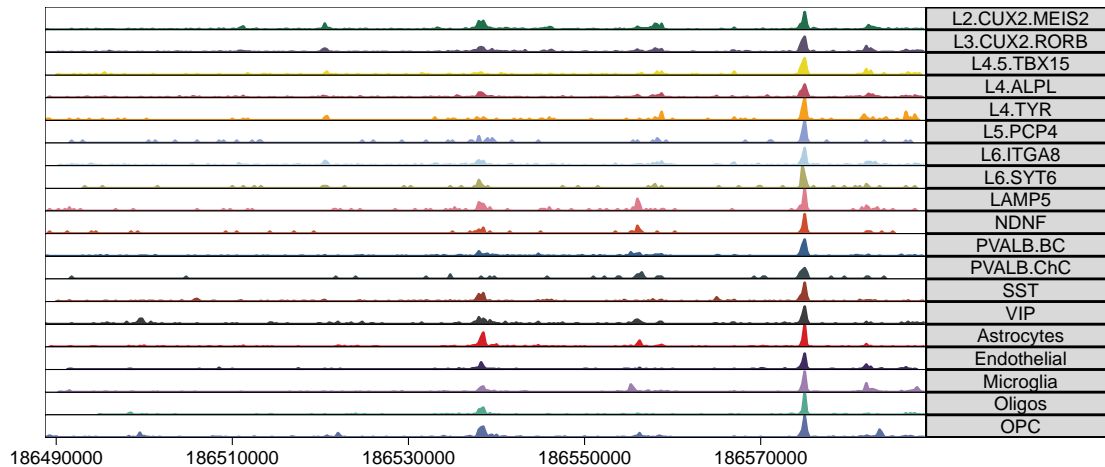
Genes

value



Coverage
(Norm. ATAC Signal Range (0-1.88) by Reads/nTSS)

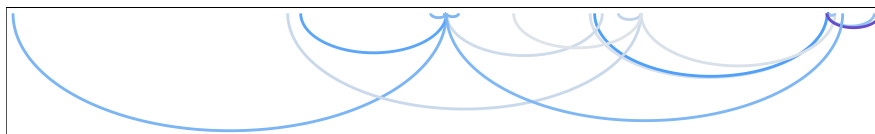
chr1:186488741-186588742



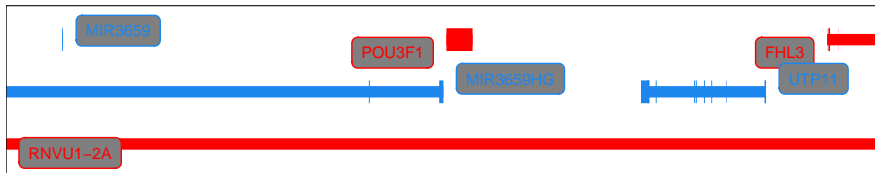
L2.CUX2.MEIS2
L3.CUX2.RORB



Peaks



Peak2GeneLinks

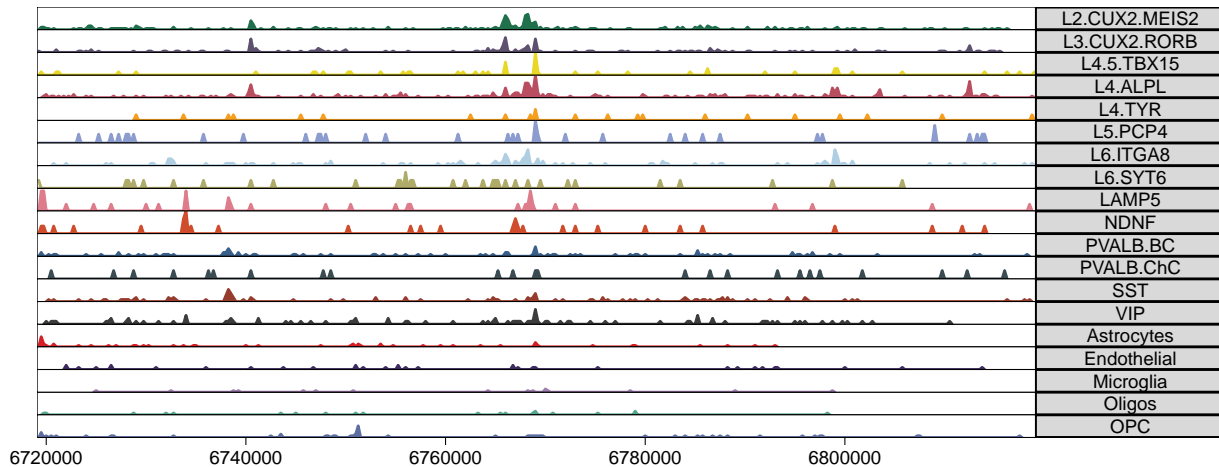


Genes

value



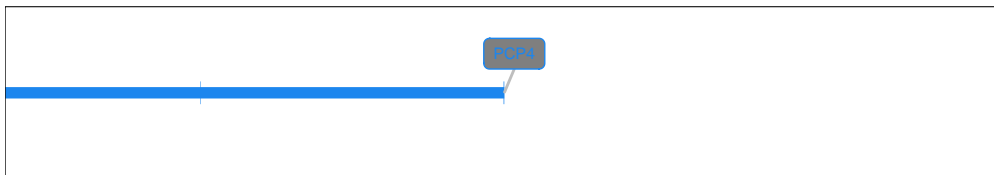
chr3:6719088–6819089

Coverage
(Norm. ATAC Signal Range (0–0.49) by Reads/nTSS)L2.CUX2.MEIS2
L3.CUX2.RORB

Peaks

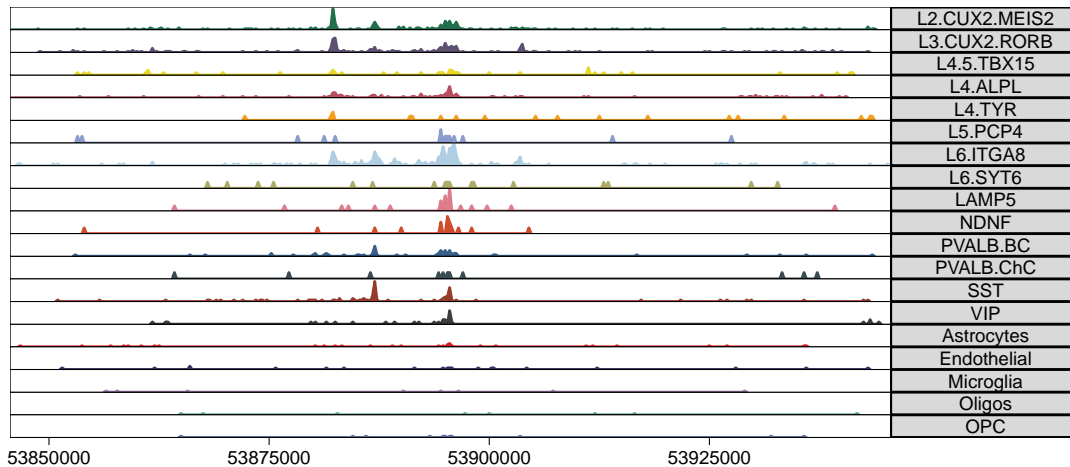


LoopTrack

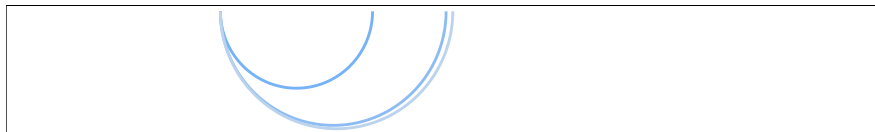


Genes

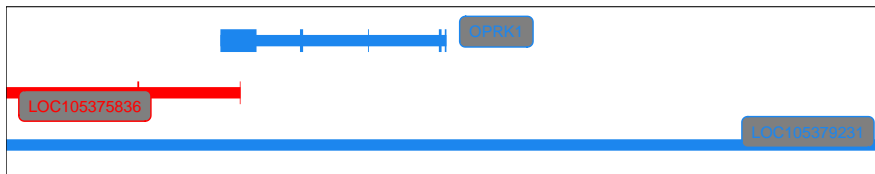
chr8:53845588-53945589

Coverage
(Norm. ATAC Signal Range (0-0.64) by Reads/nTSS)L2.CUX2.MEIS2
L3.CUX2.RORB

Peaks

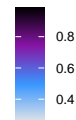


Peak2GeneLinks

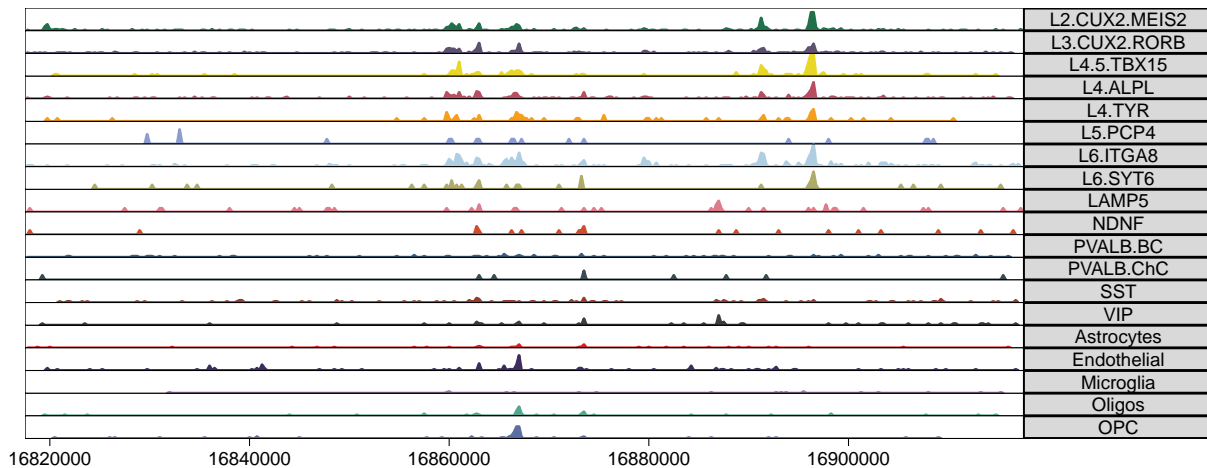


Genes

value



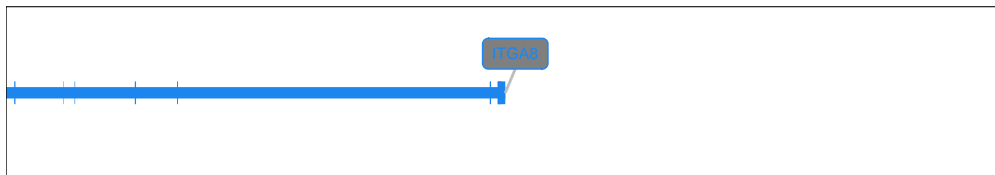
chr9:16817524–16917525

Coverage
(Norm. ATAC Signal Range (0–0.88) by Reads/nTSS)L2.CUX2.MEIS2
L3.CUX2.RORB

Peaks

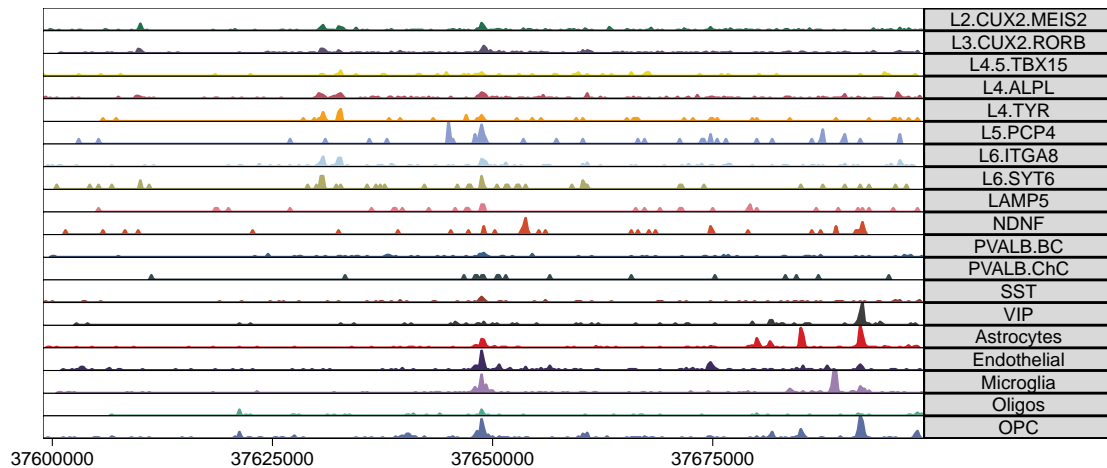


LoopTrack

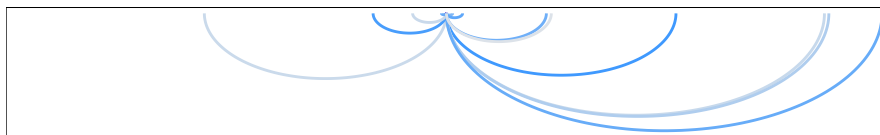


Genes

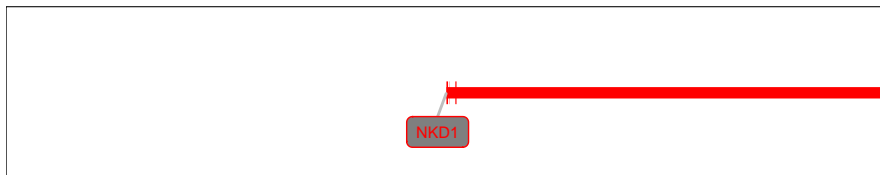
chr20:37598957-37698958

Coverage
(Norm. ATAC Signal Range (0-0.89) by Reads/nTSS)L2.CUX2.MEIS2
L3.CUX2.RORB

Peaks

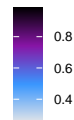


Peak2GeneLinks



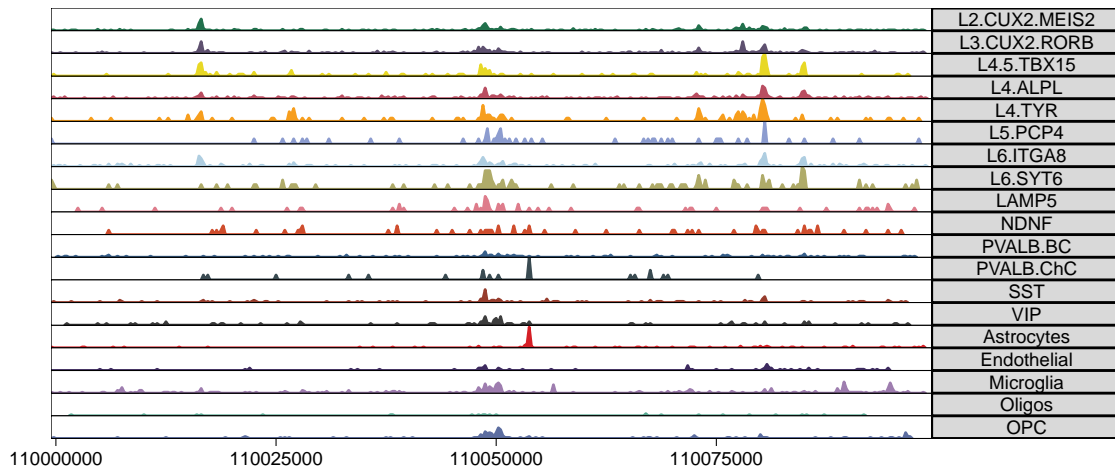
Genes

value



Coverage
(Norm. ATAC Signal Range (0-0.83) by Reads/nTSS)

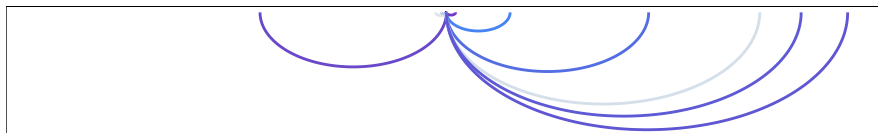
chr1:109999458-110099459



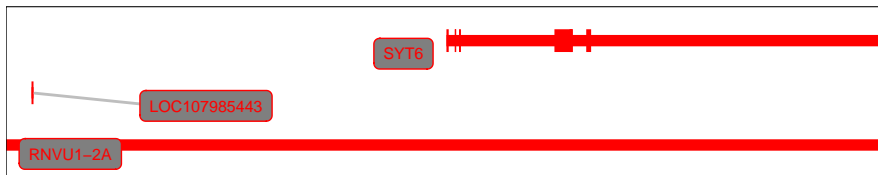
L2.CUX2.MEIS2
L3.CUX2.RORB



Peaks

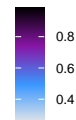


Peak2GeneLinks

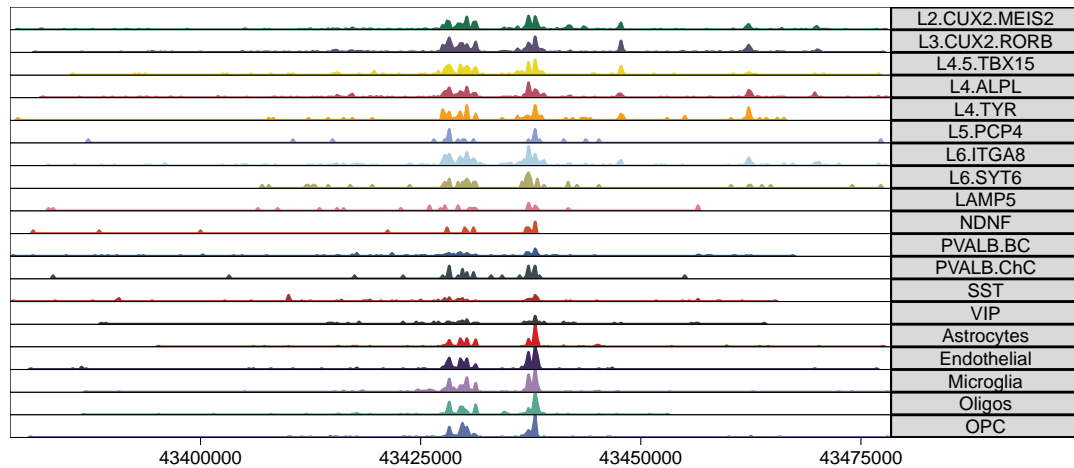


Genes

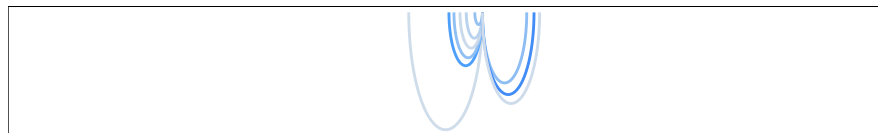
value



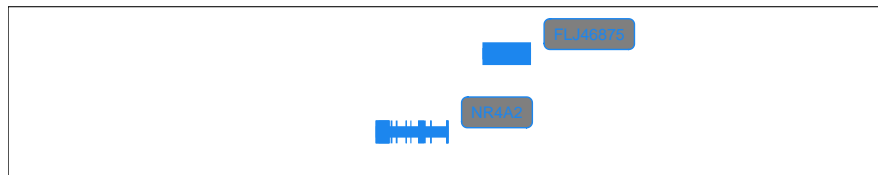
chr12:43378380-43478381

Coverage
(Norm. ATAC Signal Range (0-1.25) by Reads/nTSS)L2.CUX2.MEIS2
L3.CUX2.RORB

Peaks



Peak2GeneLinks



Genes

value



