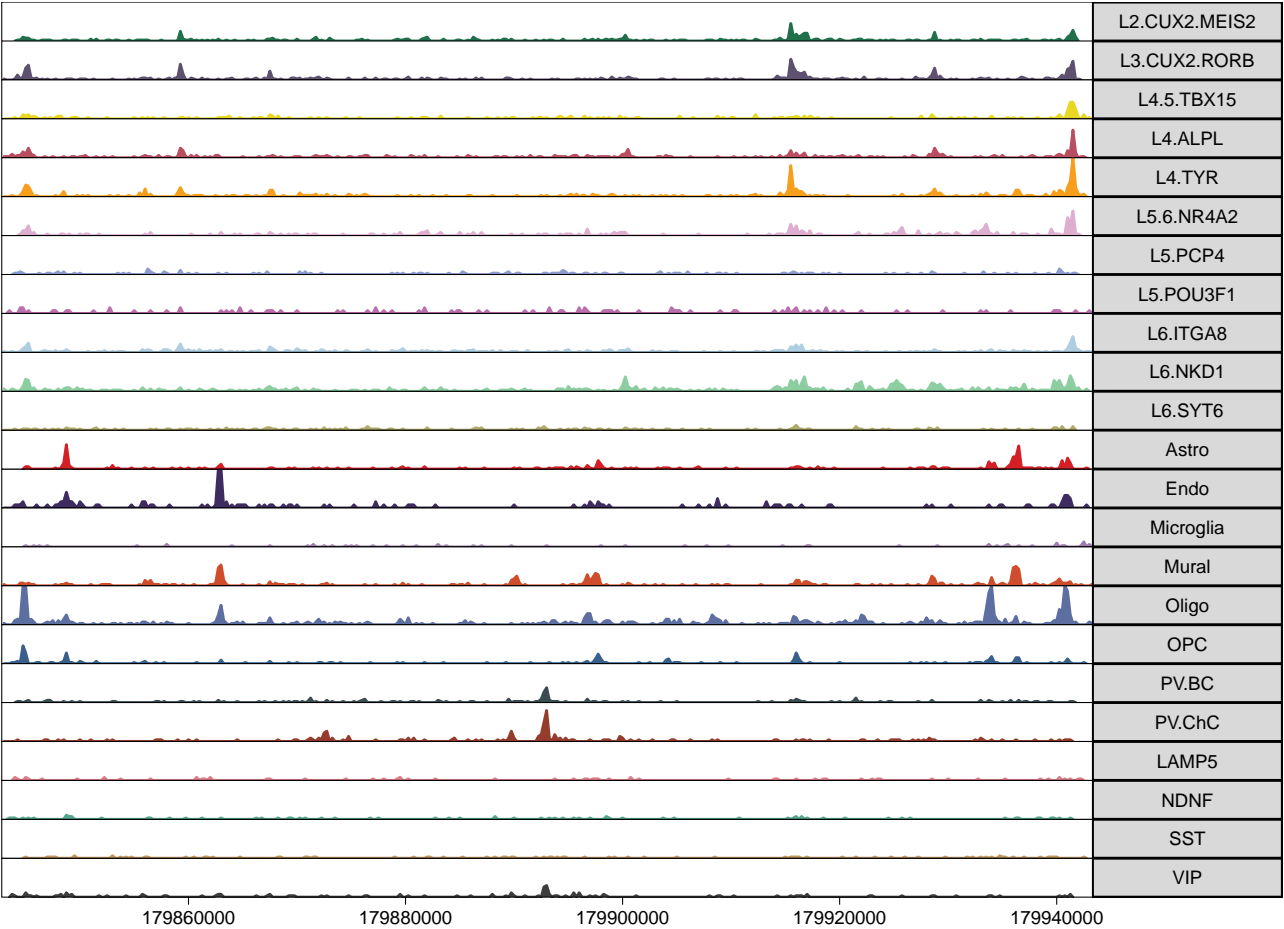
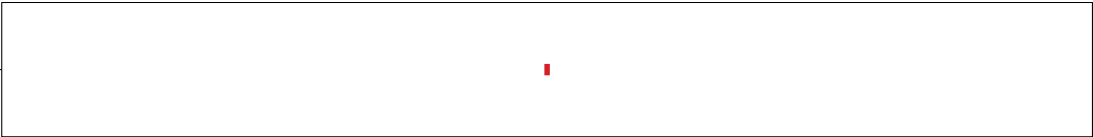


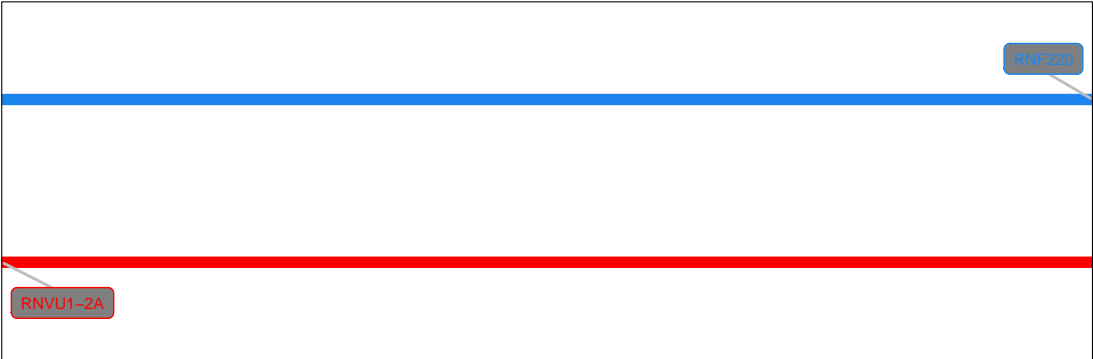
Coverage  
(Norm. ATAC Signal Range (0–0.99) by ReadsInTSS)



PV.ChC



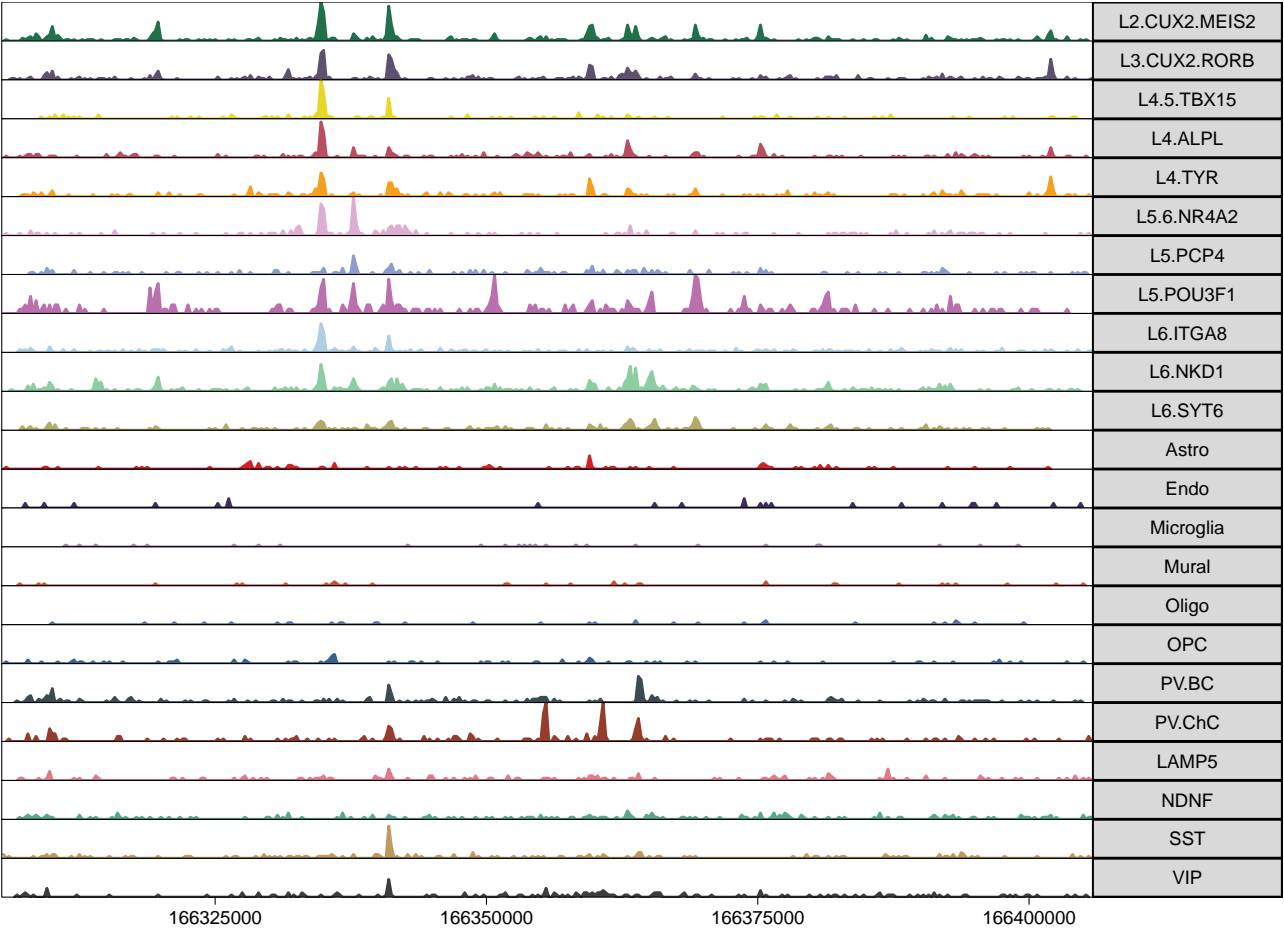
Peaks



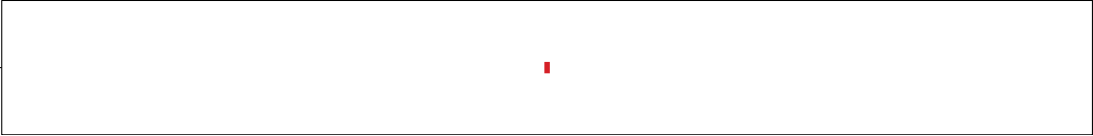
Genes

chr1:166305339–166405840 Marker Peak Tracks for: PV.ChC

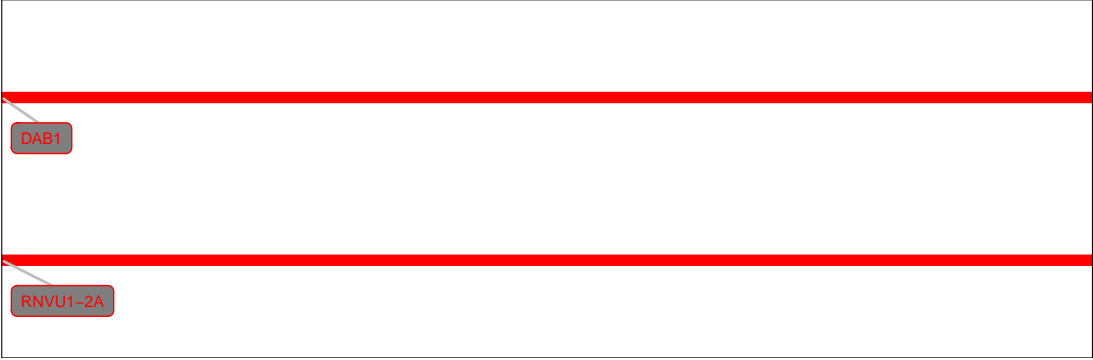
Coverage  
(Norm. ATAC Signal Range (0–0.67) by ReadsInTSS)



PV.ChC

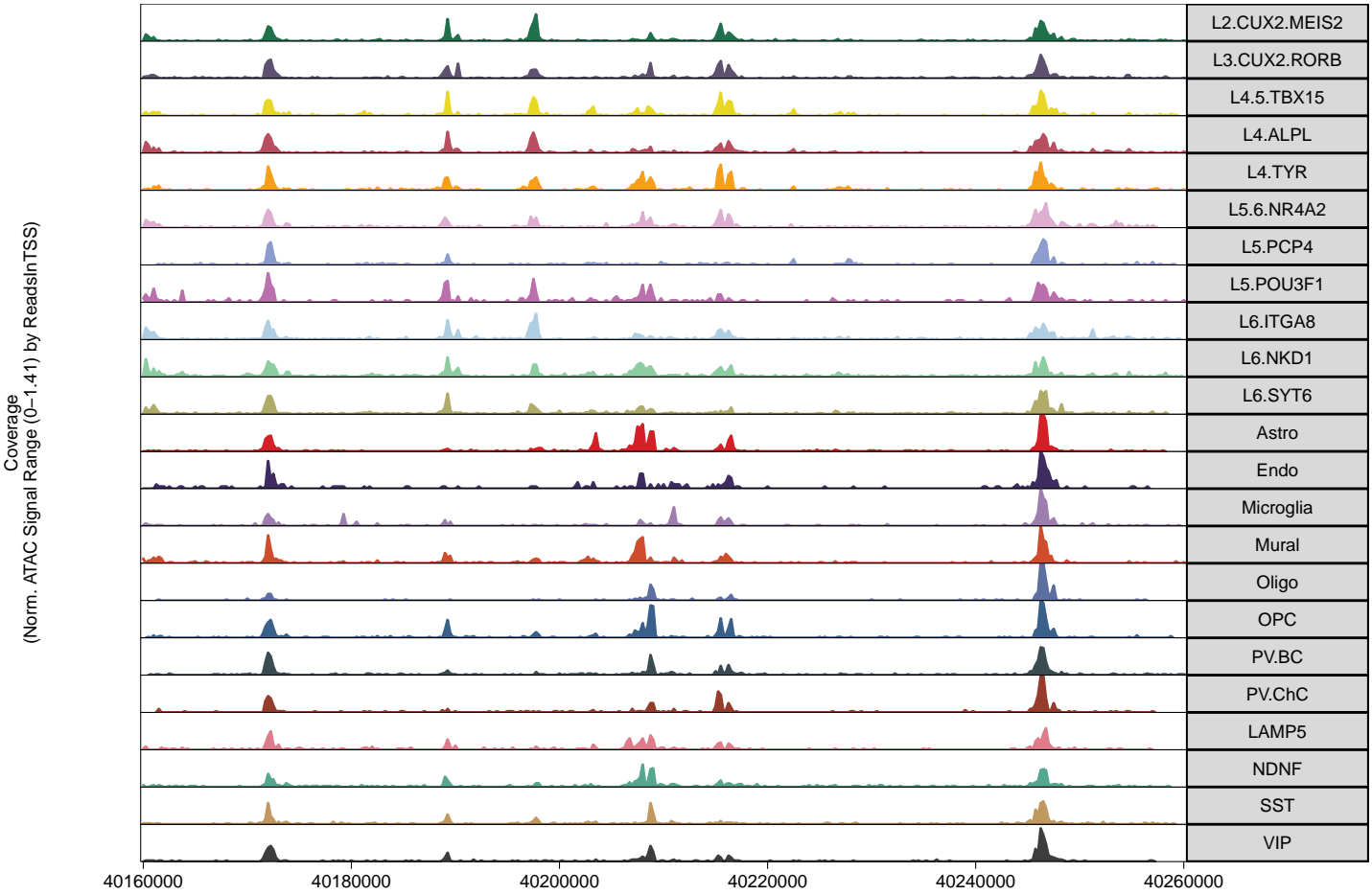


Peaks

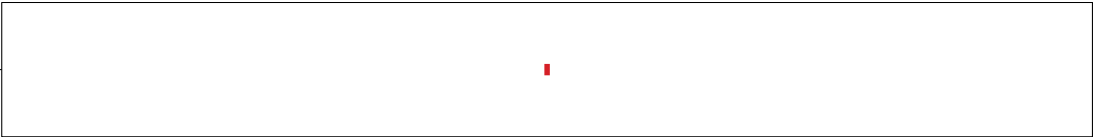


Genes

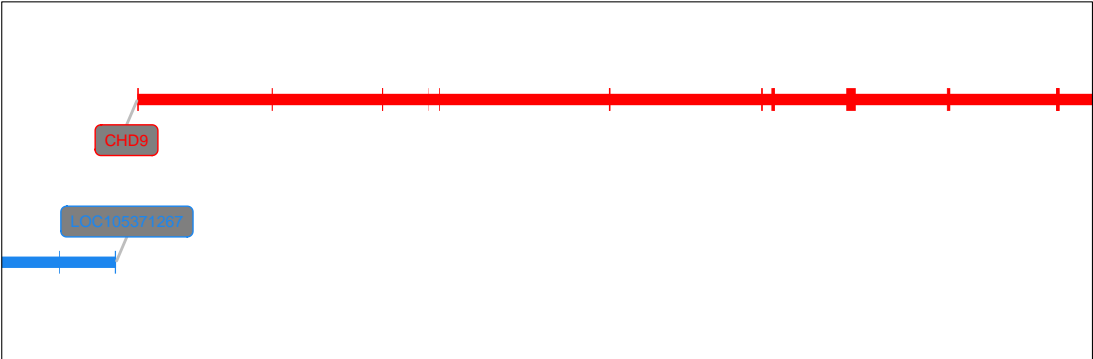
chr20:40159753–40260254 Marker Peak Tracks for: PV.ChC



PV.ChC



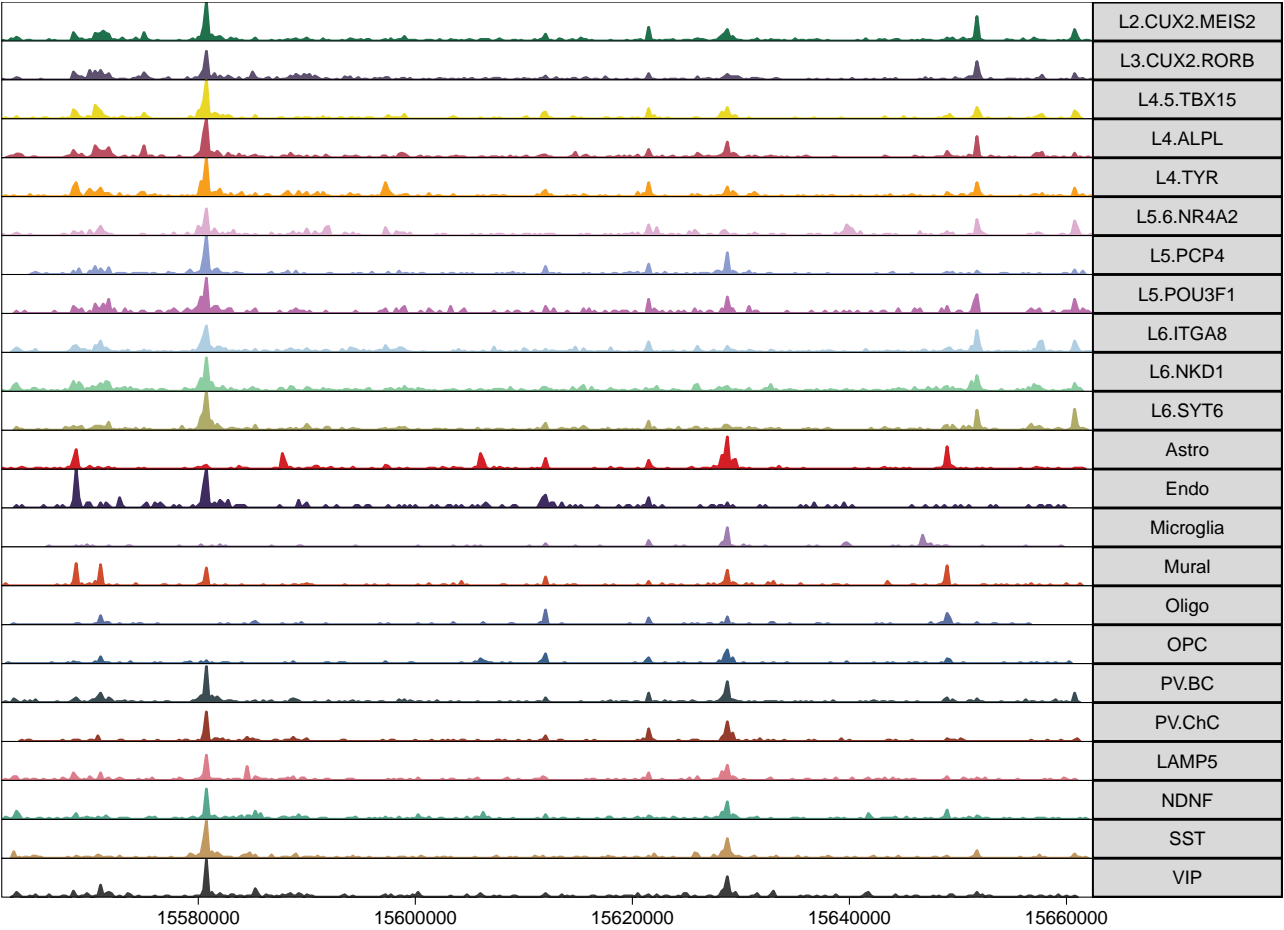
Peaks



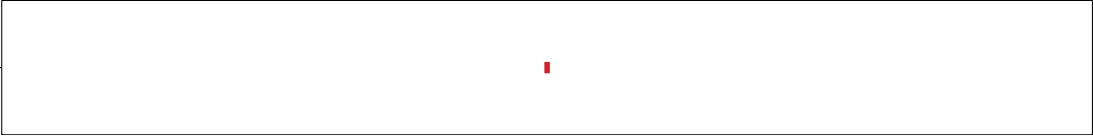
Genes

chr10:15561892-15662393 Marker Peak Tracks for: PV.ChC

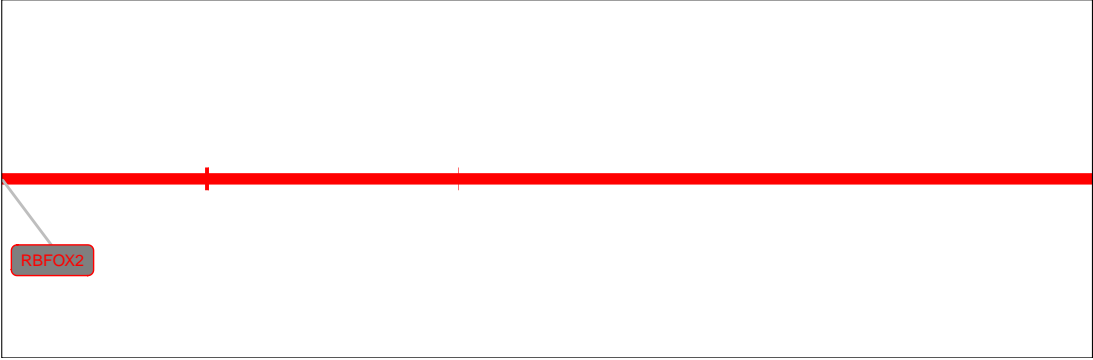
Coverage  
(Norm. ATAC Signal Range (0-1.22) by ReadsInTSS)



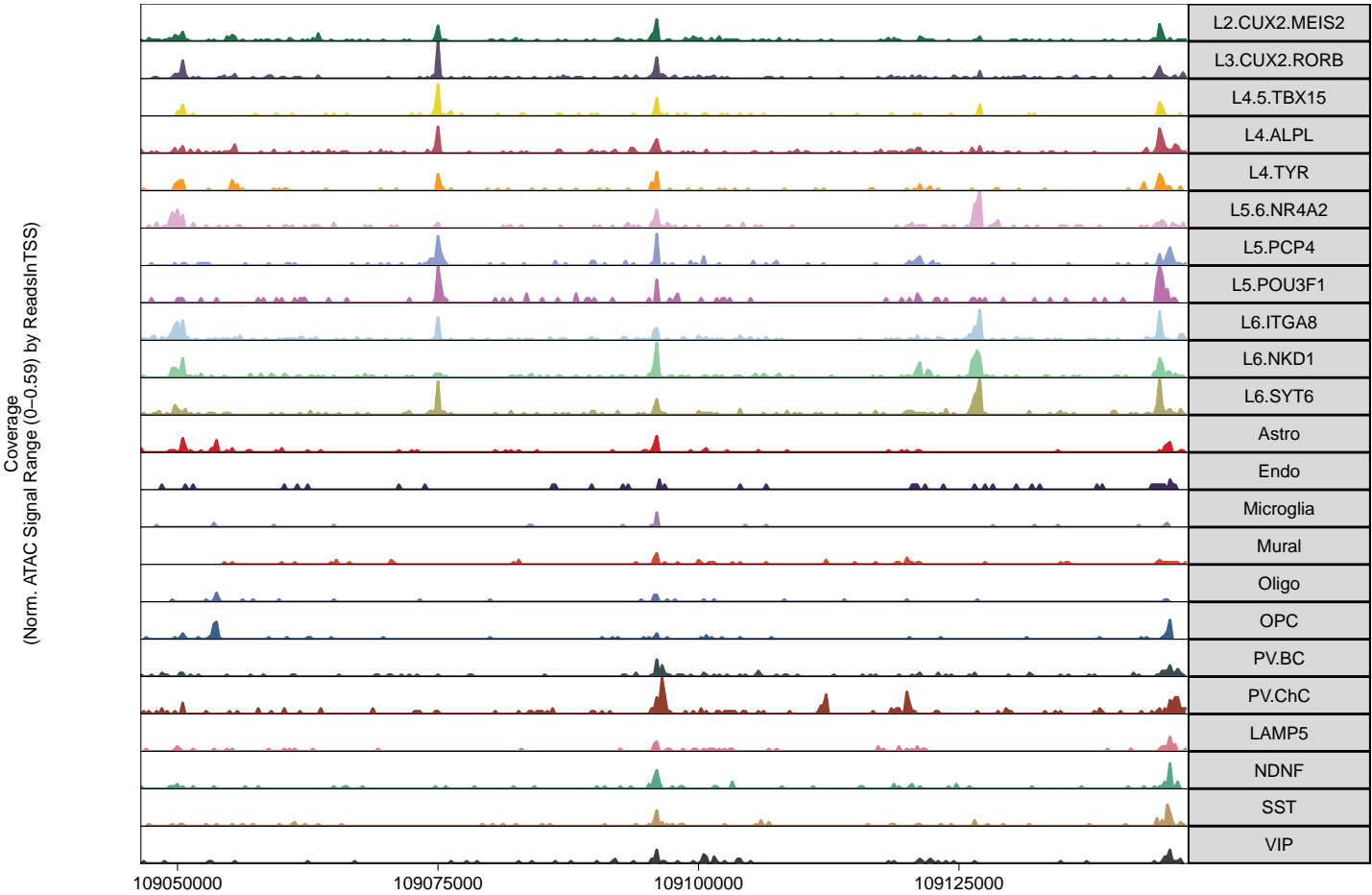
PV.ChC



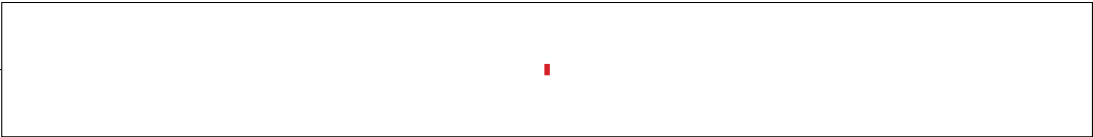
Peaks



Genes



PV.ChC

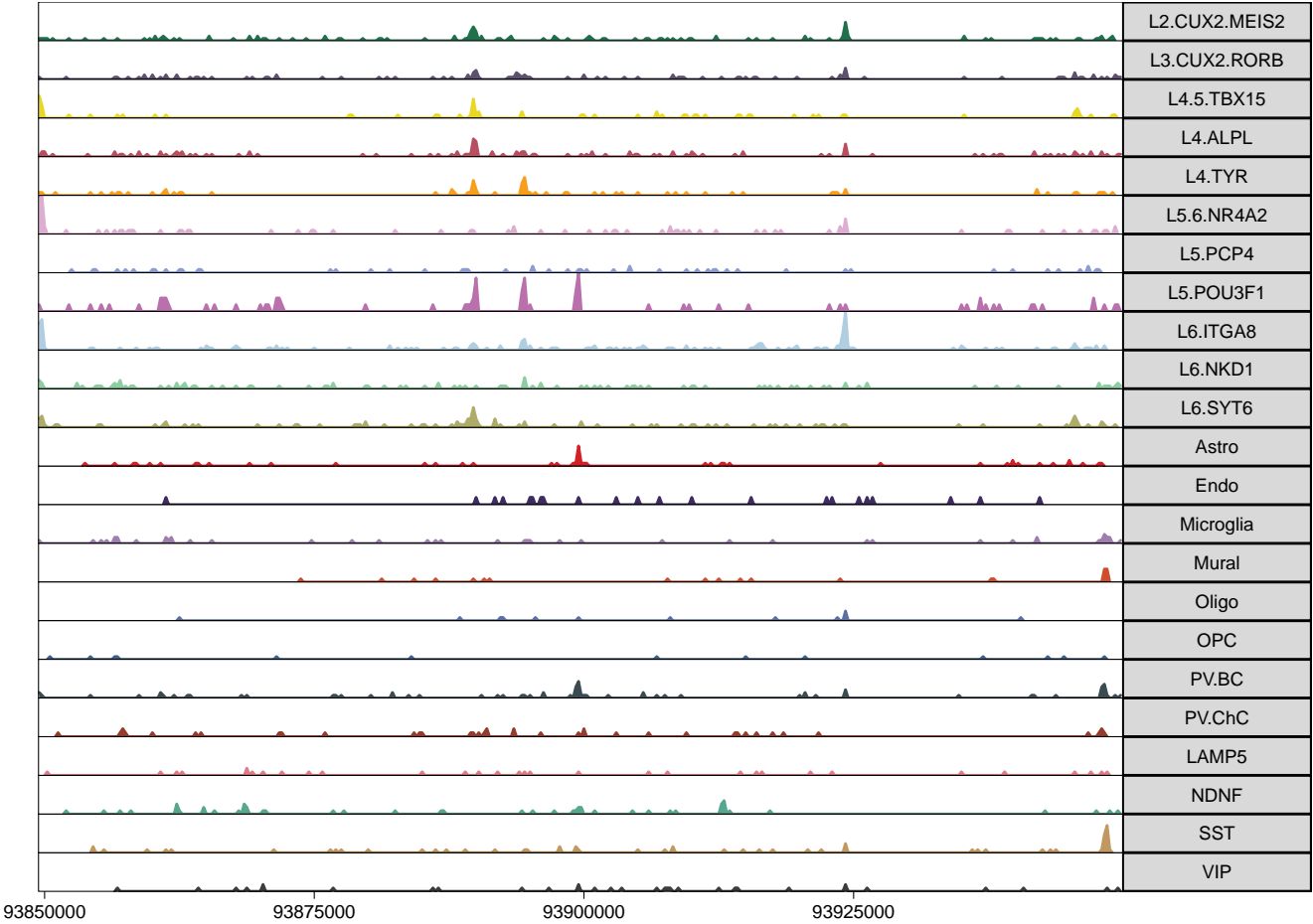


Peaks

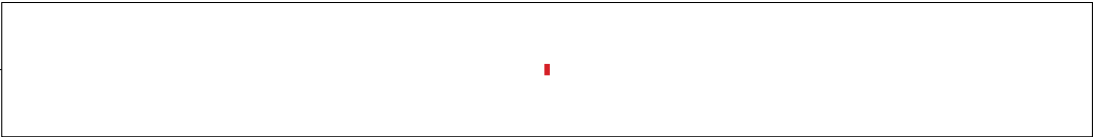


Genes

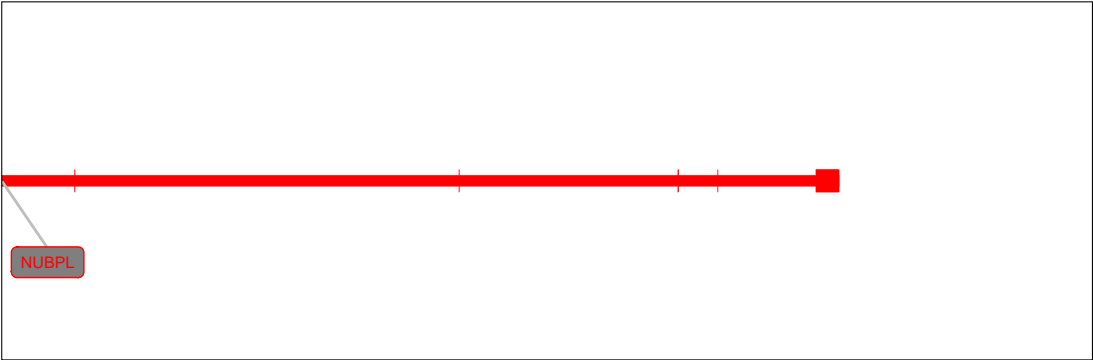
Coverage  
(Norm. ATAC Signal Range (0–0.42) by ReadsInTSS)



PV.ChC

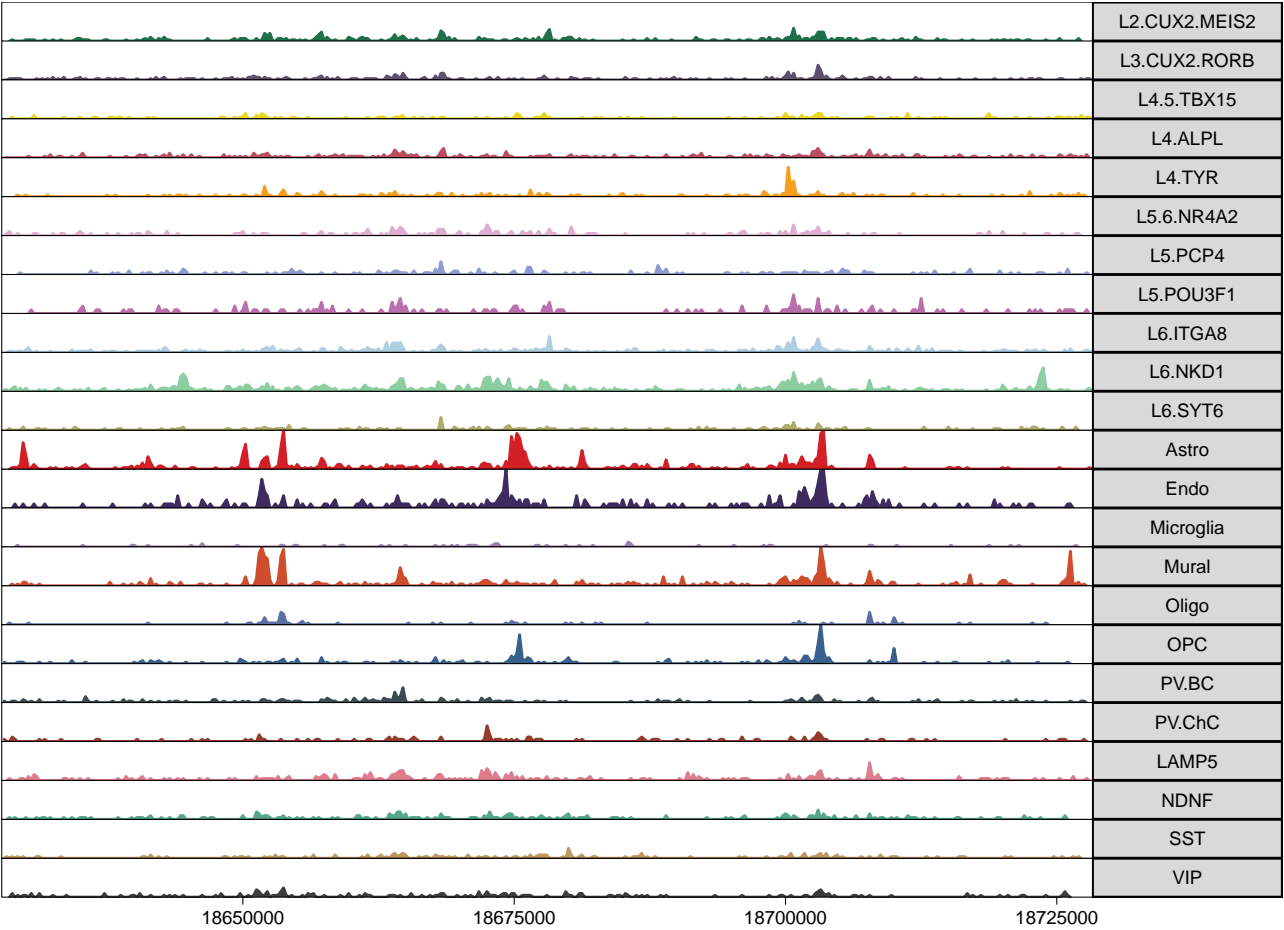


Peaks

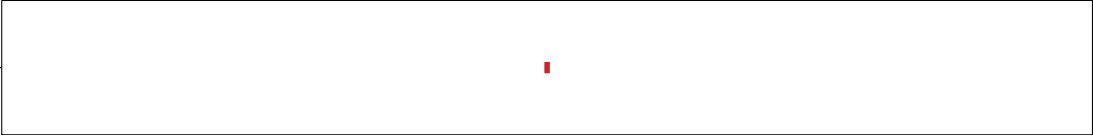


Genes

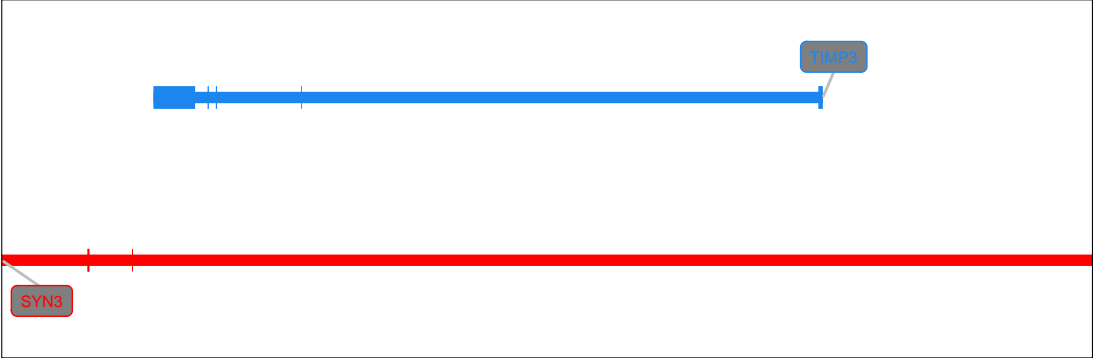
Coverage  
(Norm. ATAC Signal Range (0–0.76) by ReadsInTSS)



PV.ChC

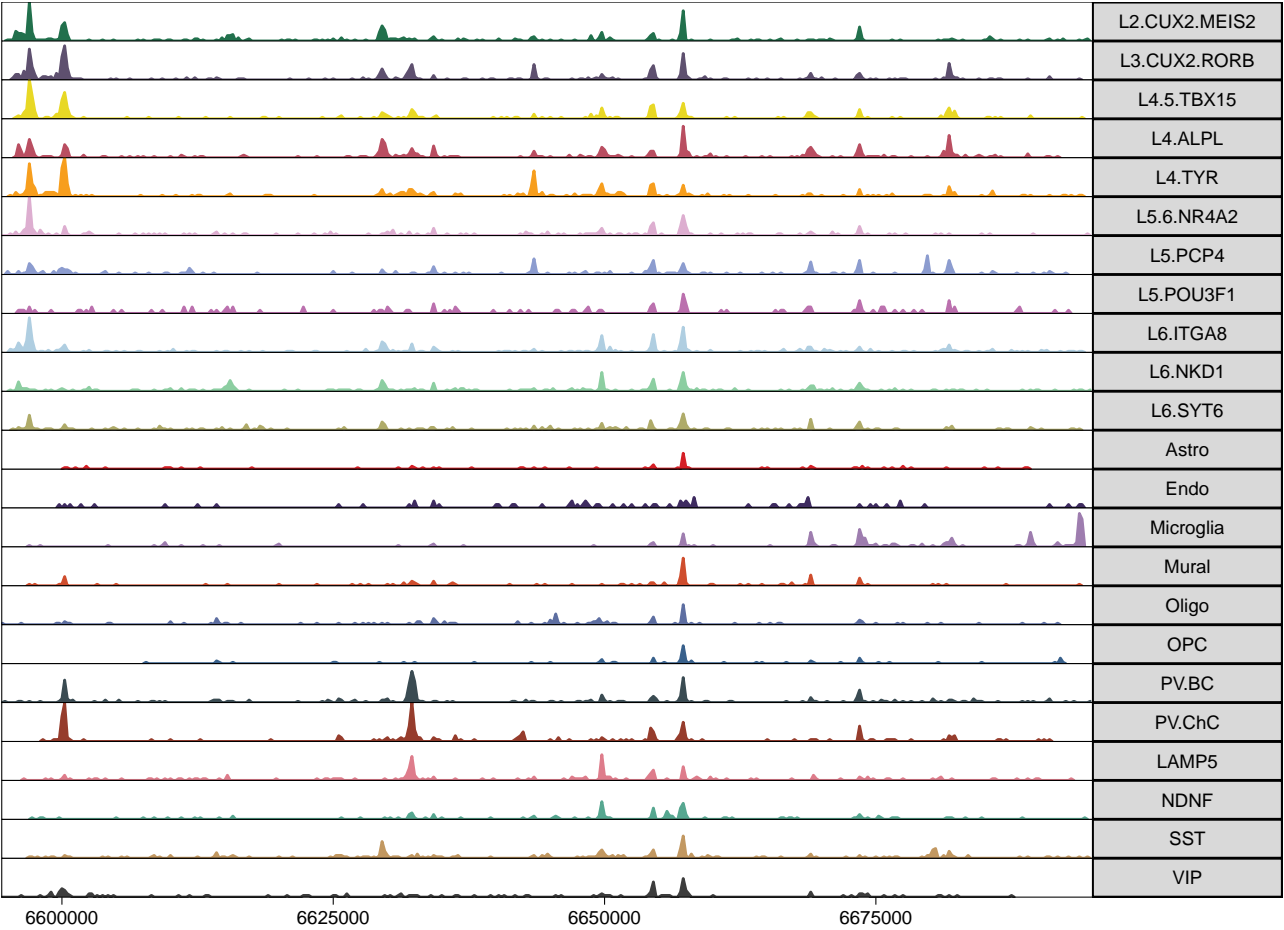


Peaks

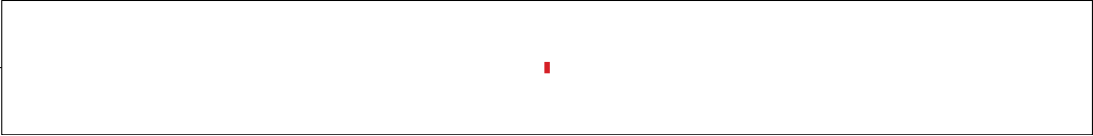


Genes

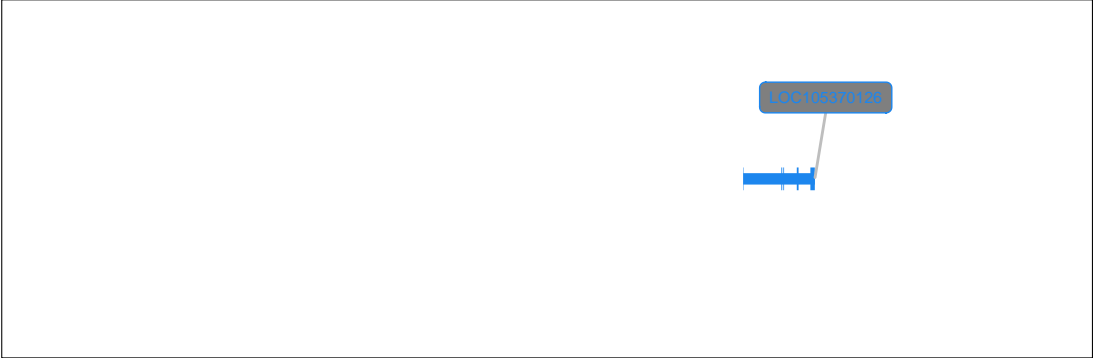
Coverage  
(Norm. ATAC Signal Range (0–0.89) by ReadsInTSS)



PV.ChC



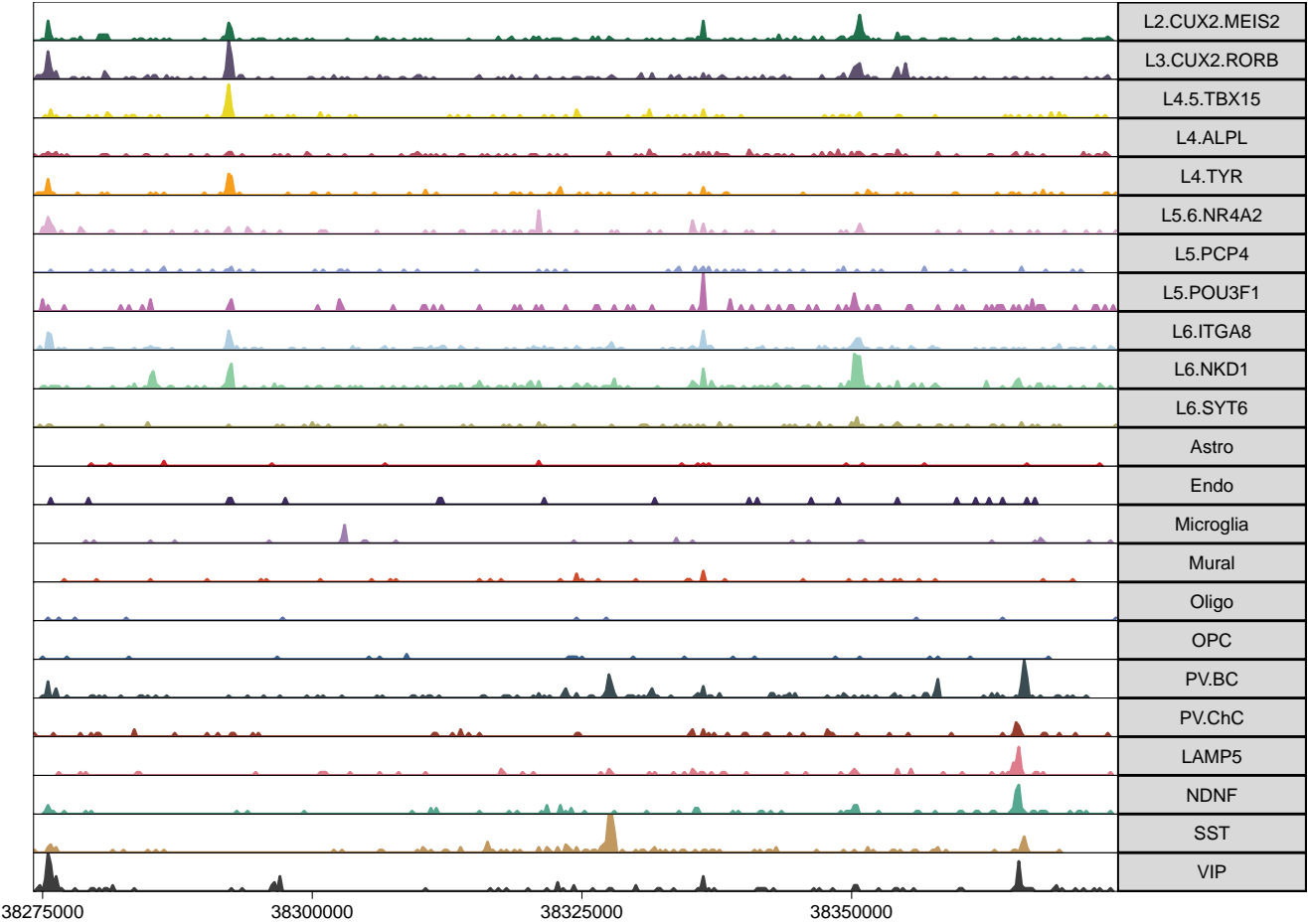
Peaks



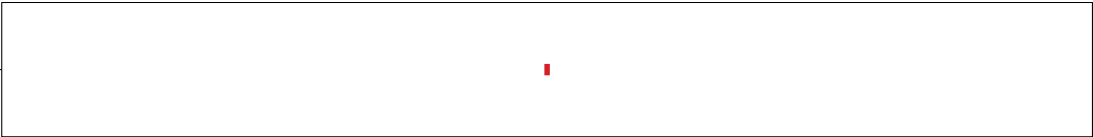
Genes



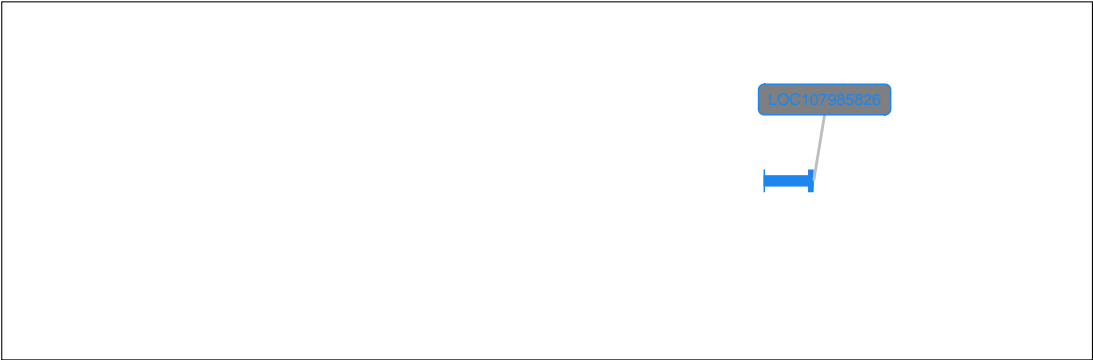
Coverage  
(Norm. ATAC Signal Range (0–0.48) by ReadsInTSS)



PV.ChC

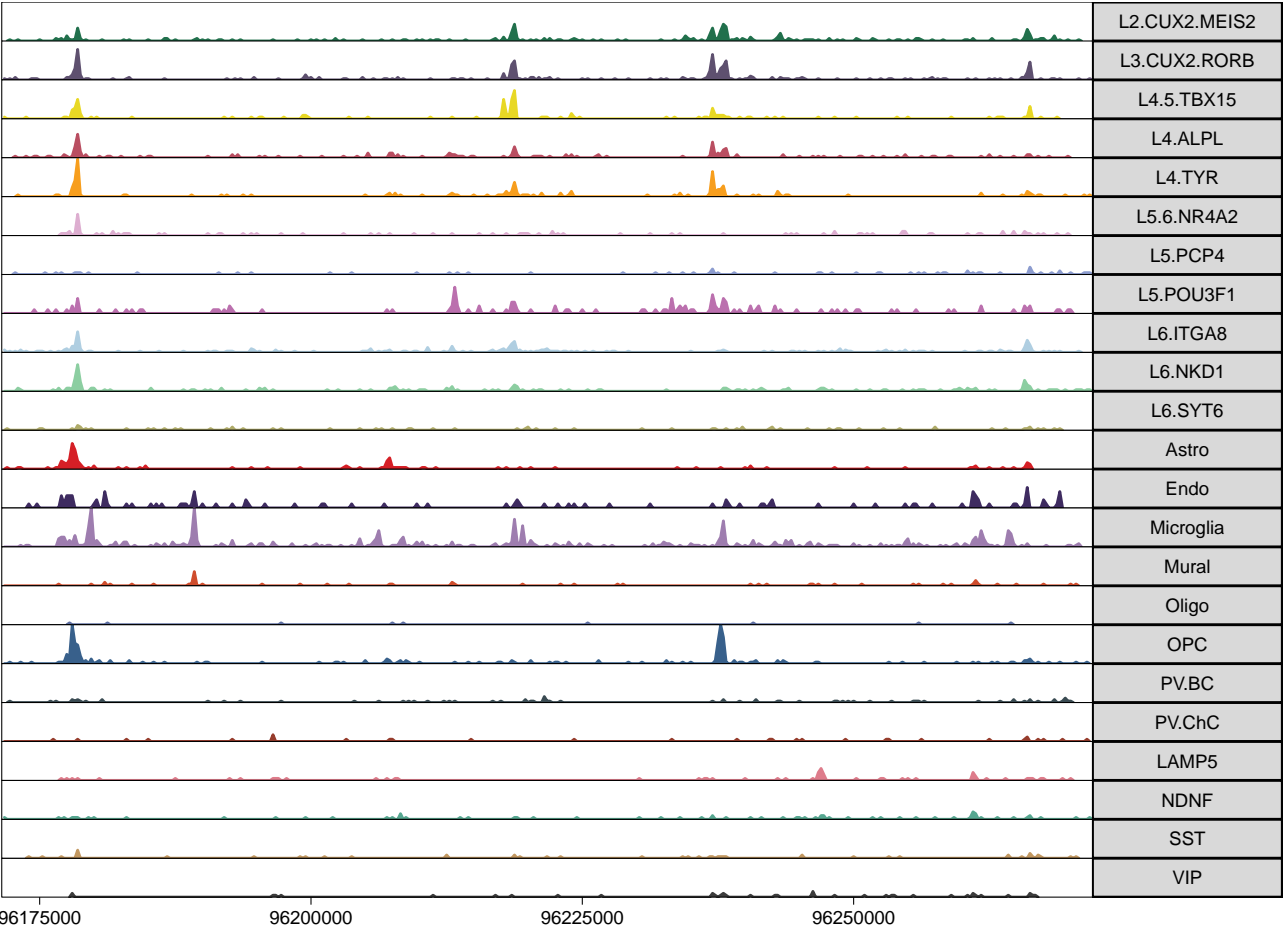


Peaks

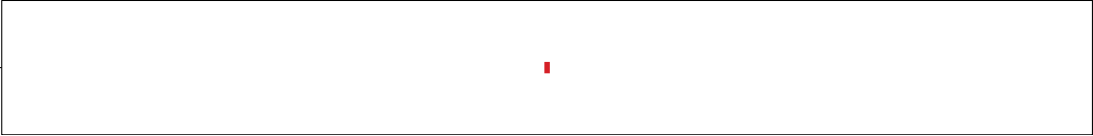


Genes

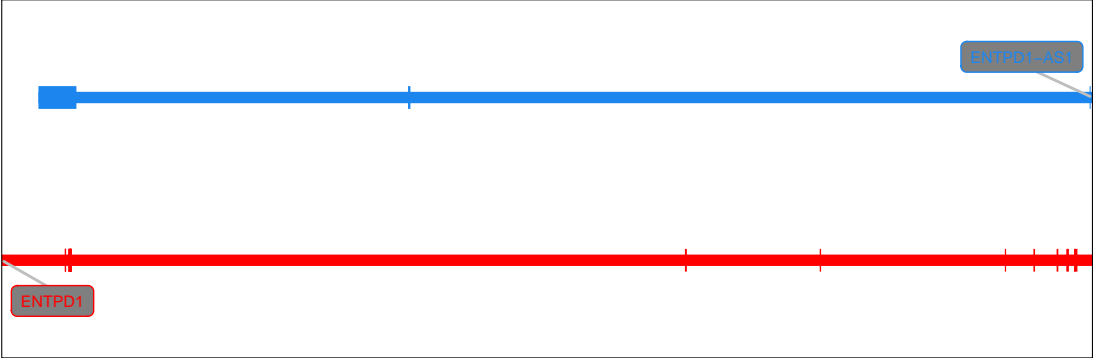
Coverage  
(Norm. ATAC Signal Range (0–0.76) by ReadsInTSS)



PV.ChC

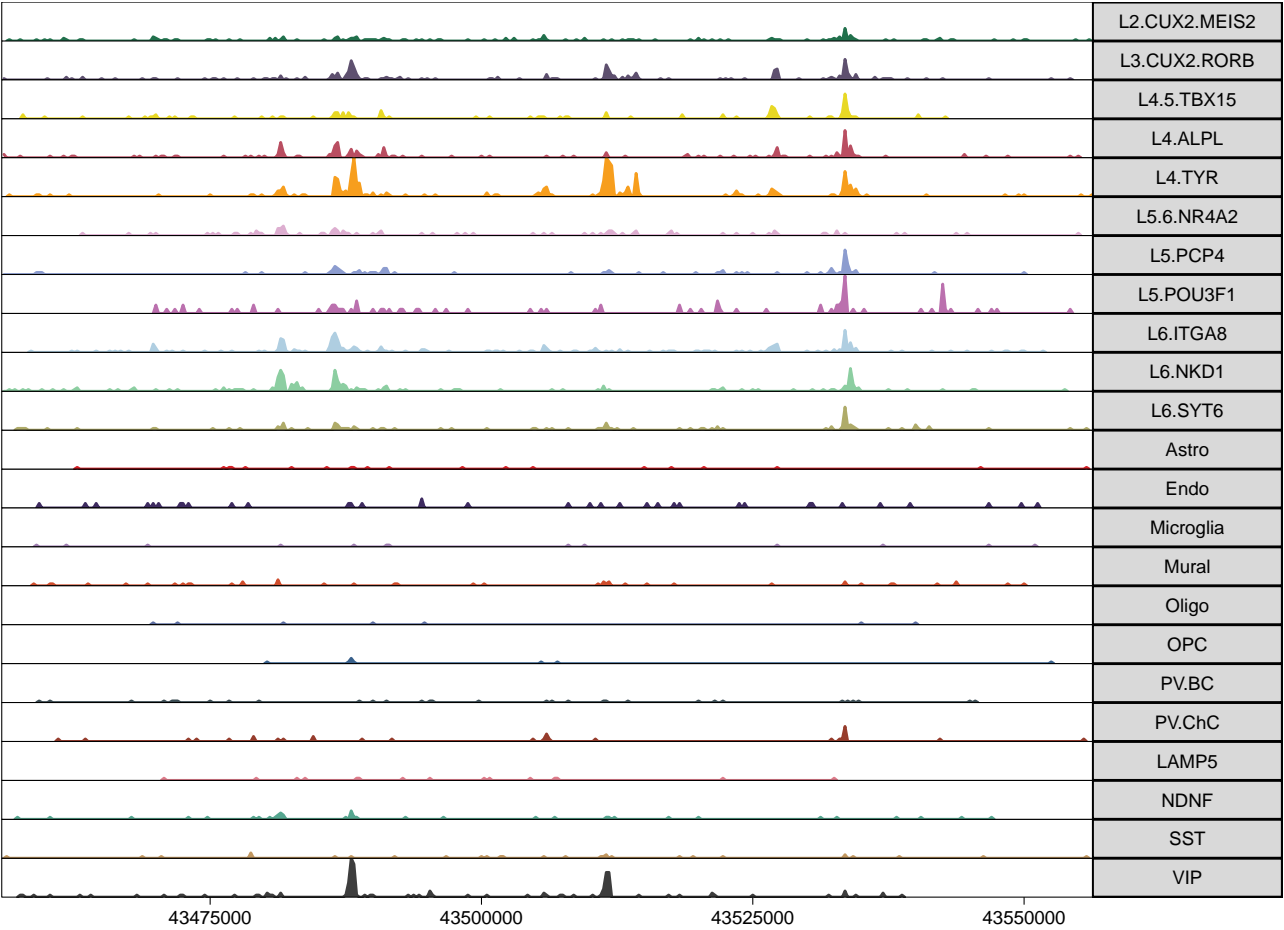


Peaks

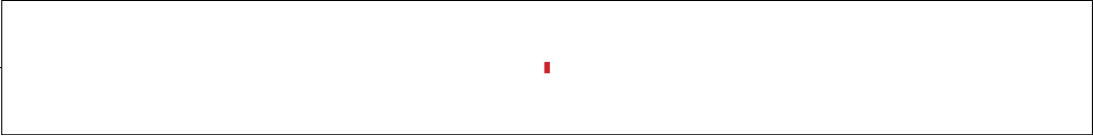


Genes

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



PV.ChC



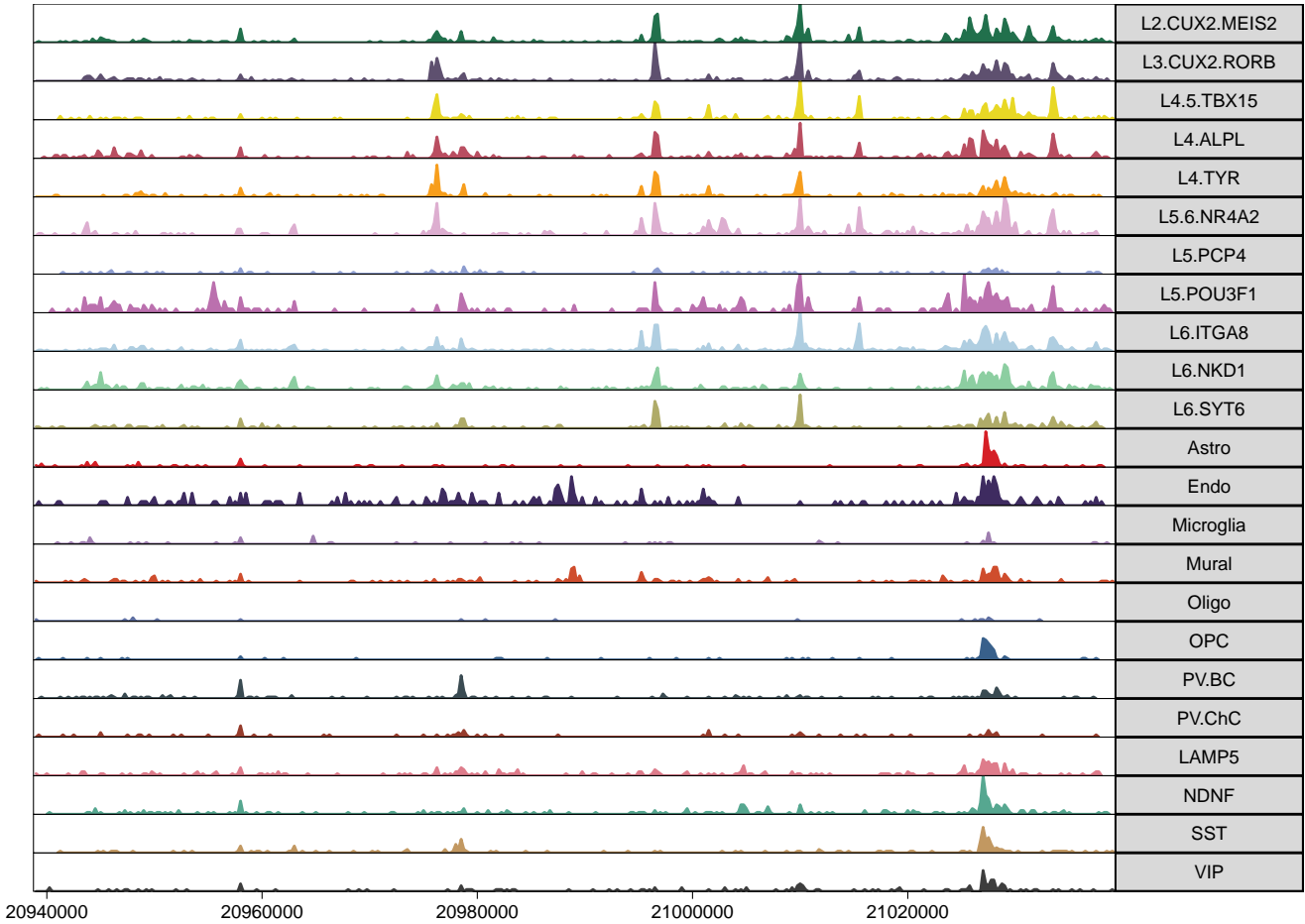
Peaks



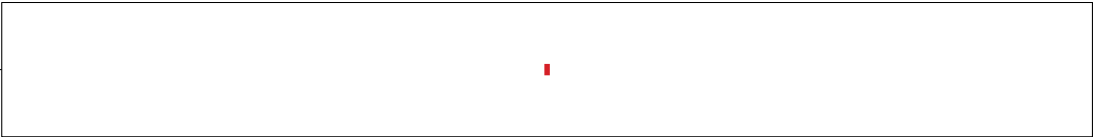
Genes

chr3:20938758–21039259 Marker Peak Tracks for: PV.ChC

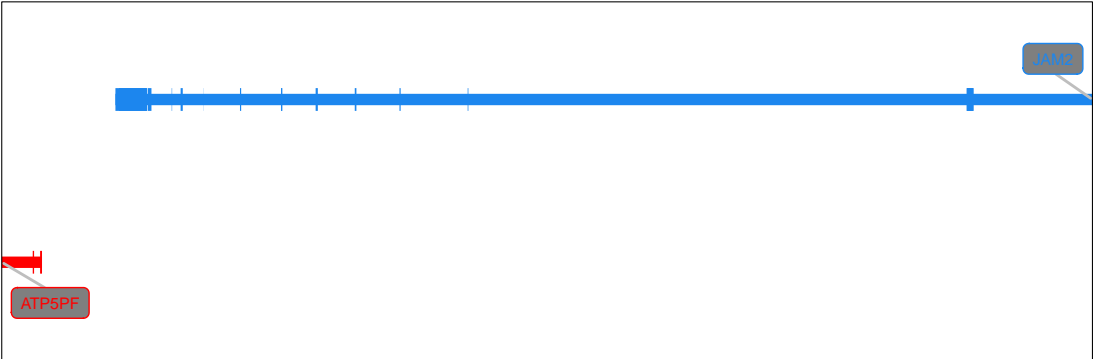
Coverage  
(Norm. ATAC Signal Range (0–0.75) by ReadsInTSS)



PV.ChC



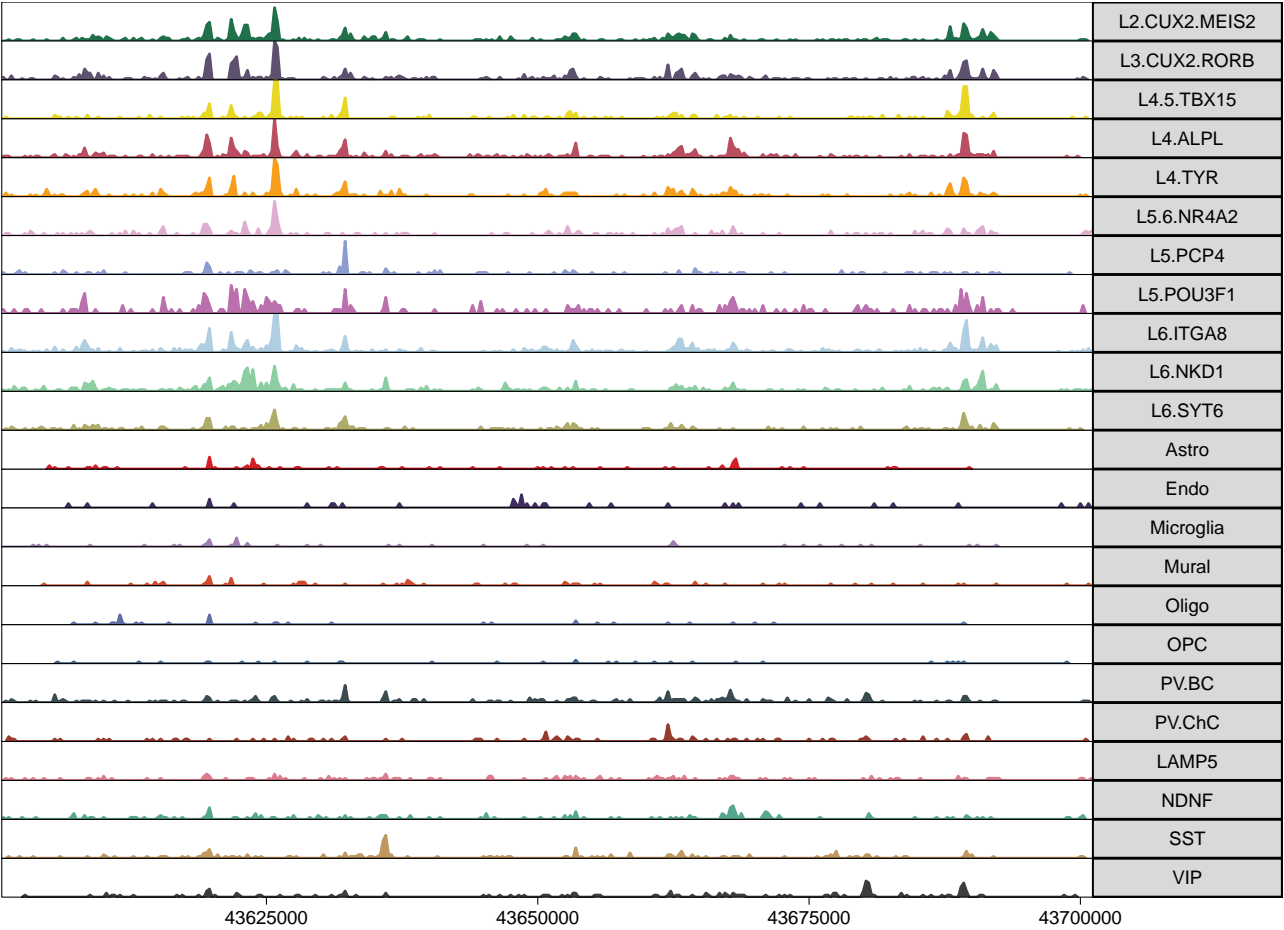
Peaks



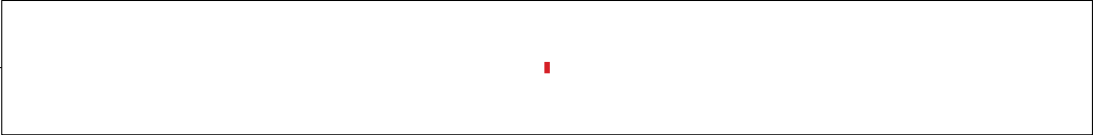
Genes

chr6:43600610–43701111 Marker Peak Tracks for: PV.ChC

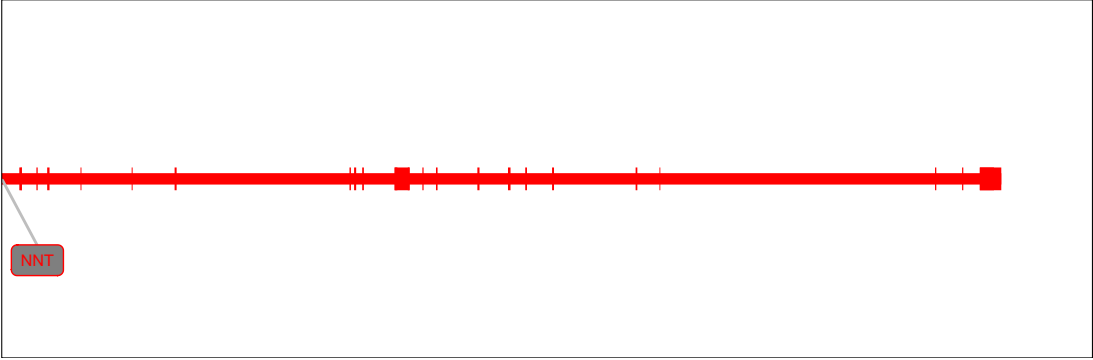
Coverage  
(Norm. ATAC Signal Range (0–0.71) by ReadsInTSS)



PV.ChC



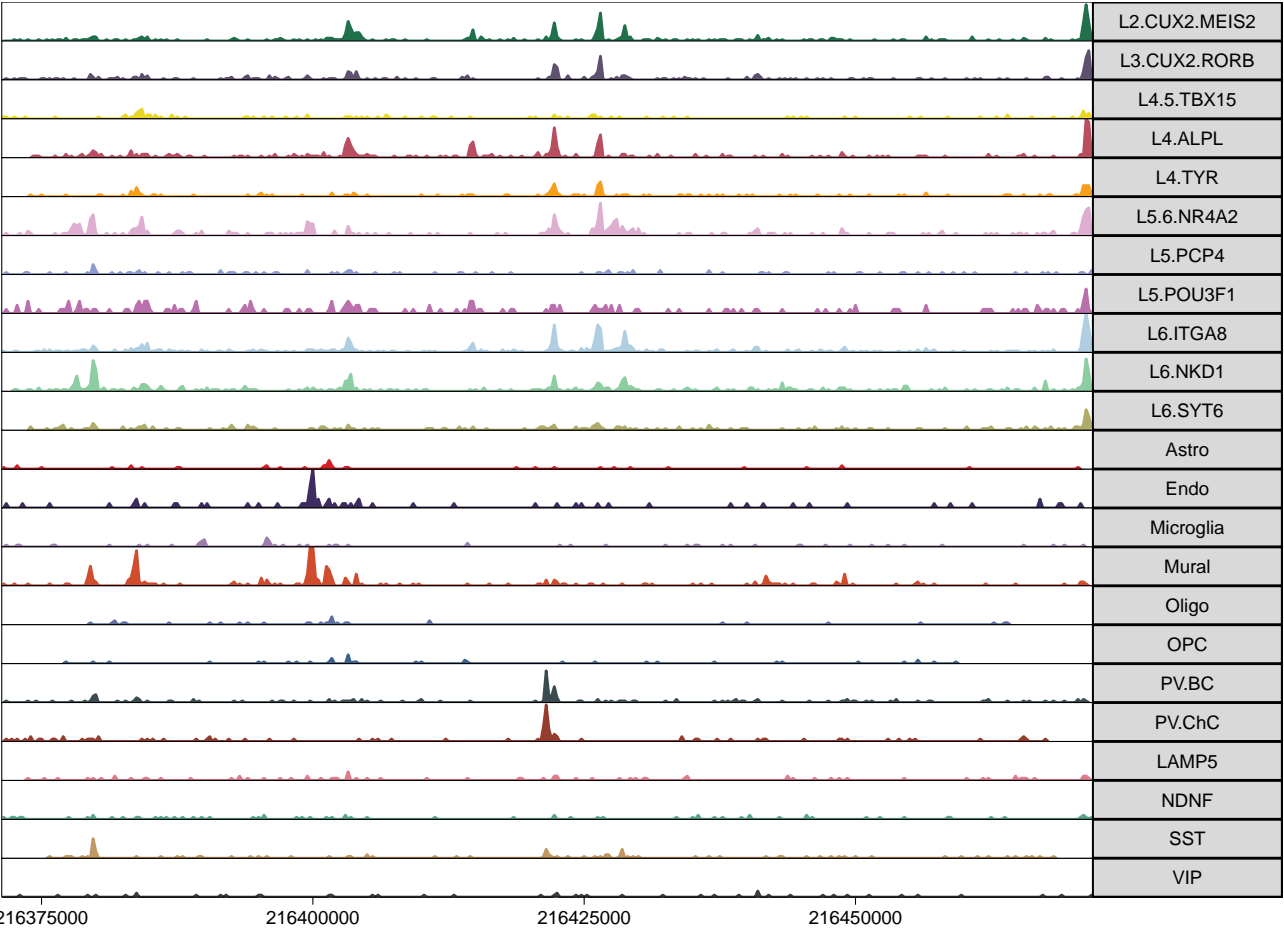
Peaks



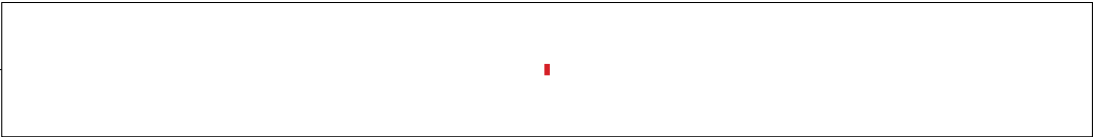
Genes

chr1:216371333-216471834 Marker Peak Tracks for: PV.ChC

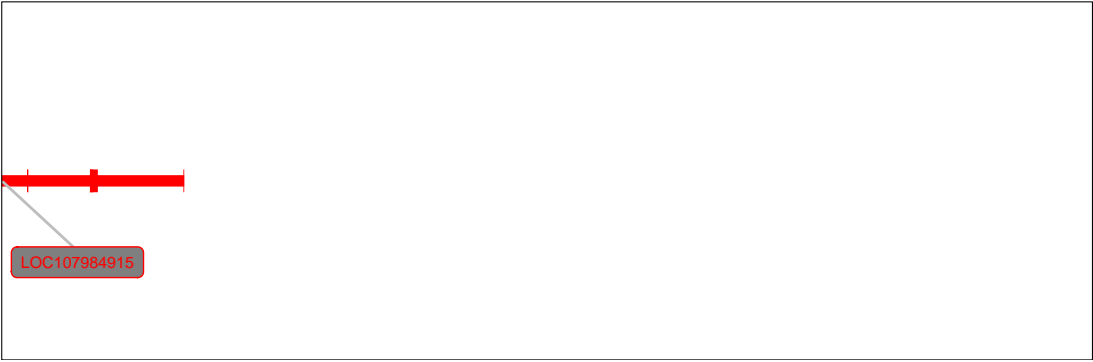
Coverage  
(Norm. ATAC Signal Range (0-0.7) by ReadsInTSS)



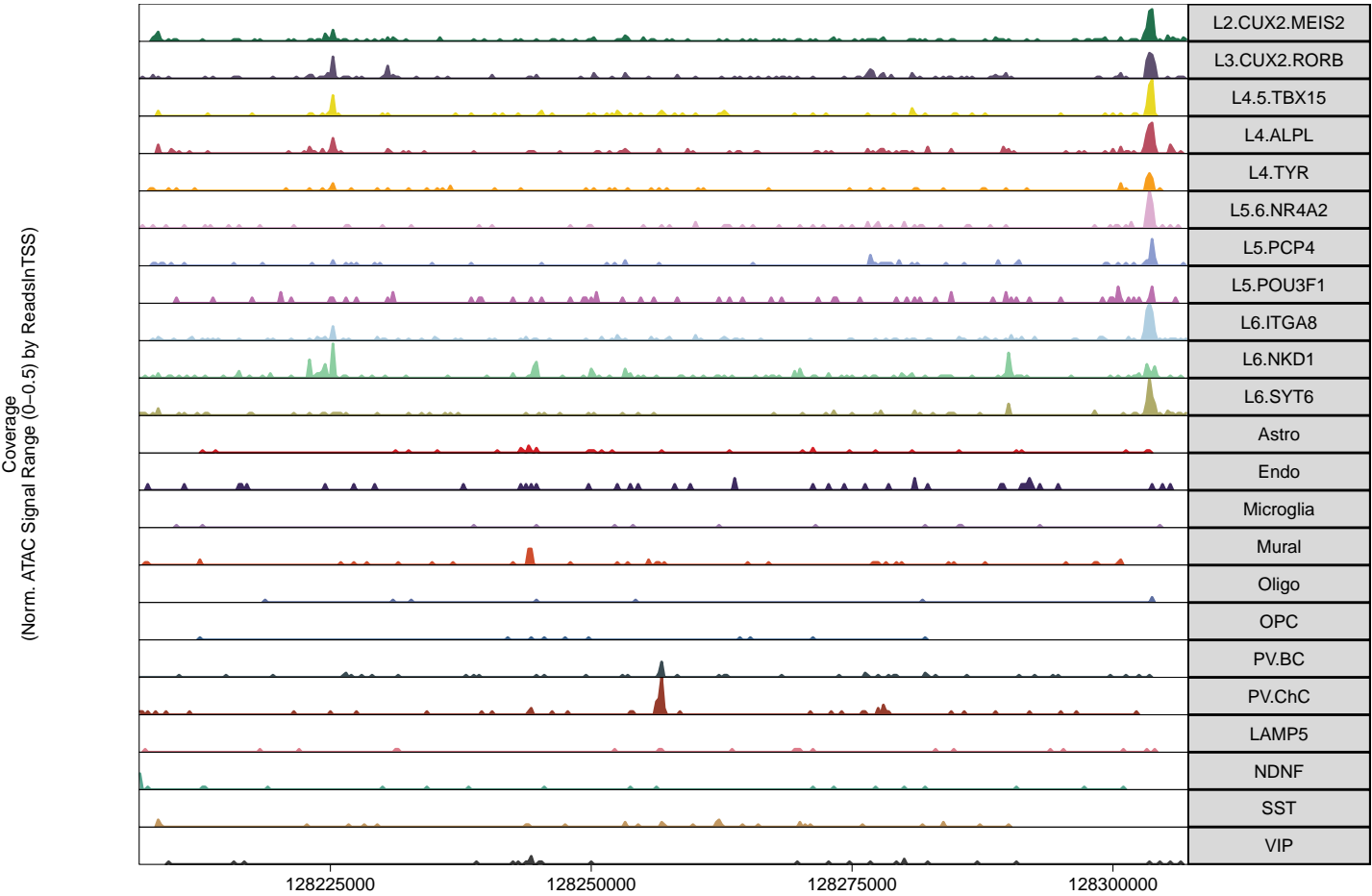
PV.ChC



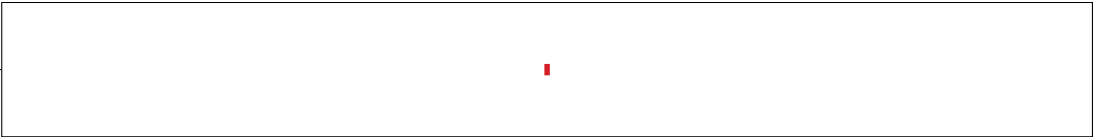
Peaks



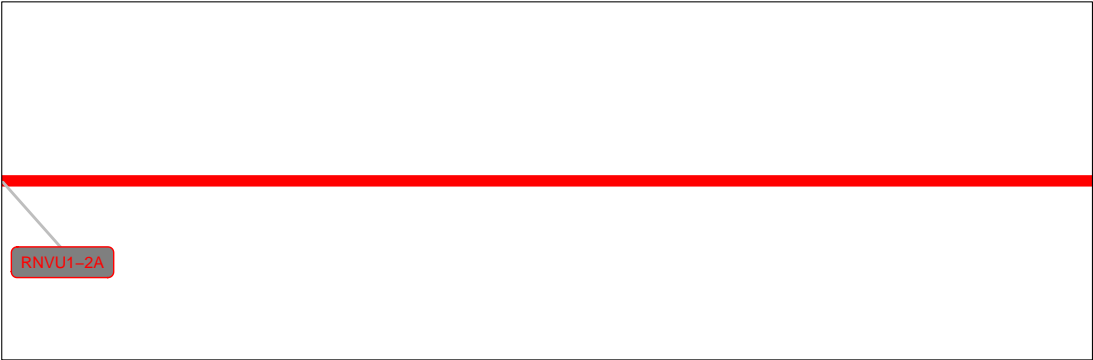
Genes



PV.ChC

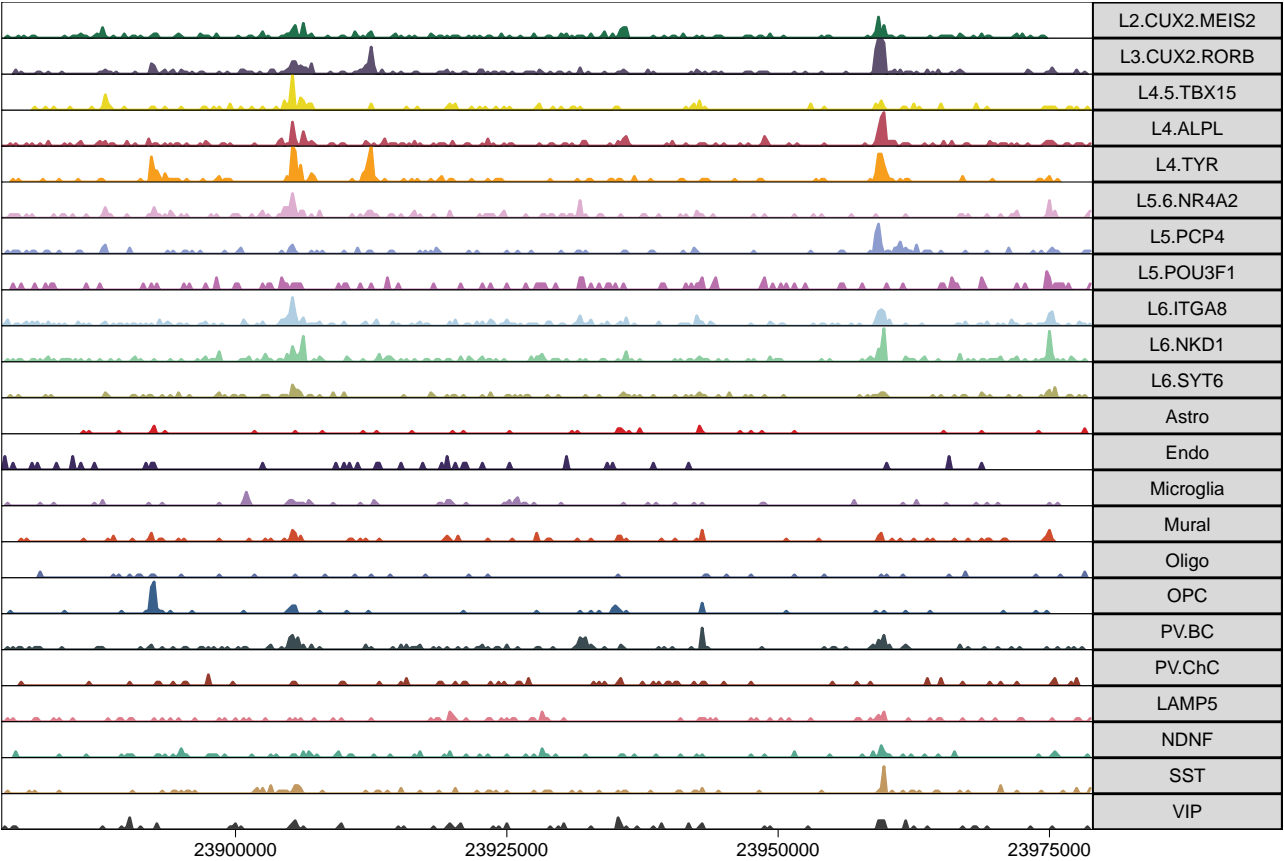


Peaks



Genes

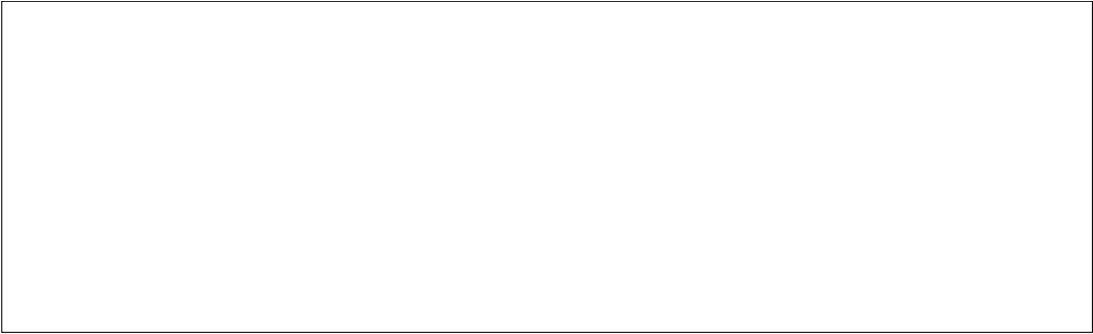
Coverage  
(Norm. ATAC Signal Range (0–0.43) by ReadsInTSS)



PV.ChC



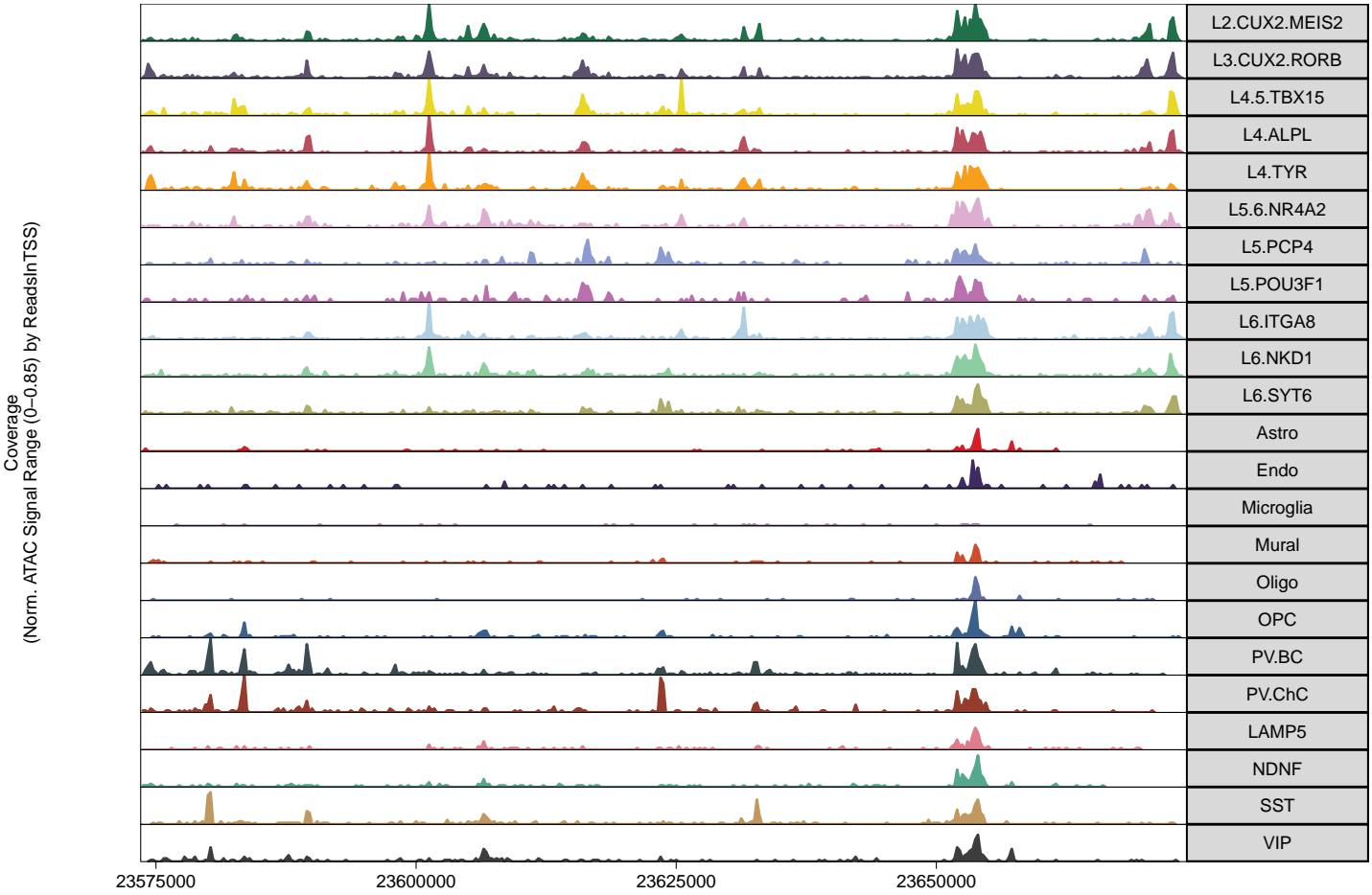
Peaks



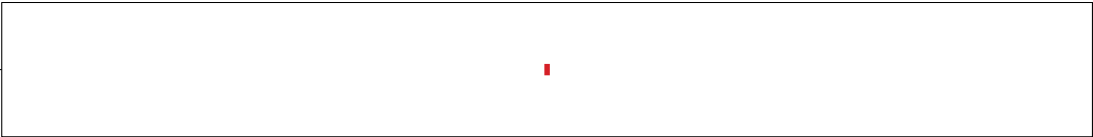
GeneTrack



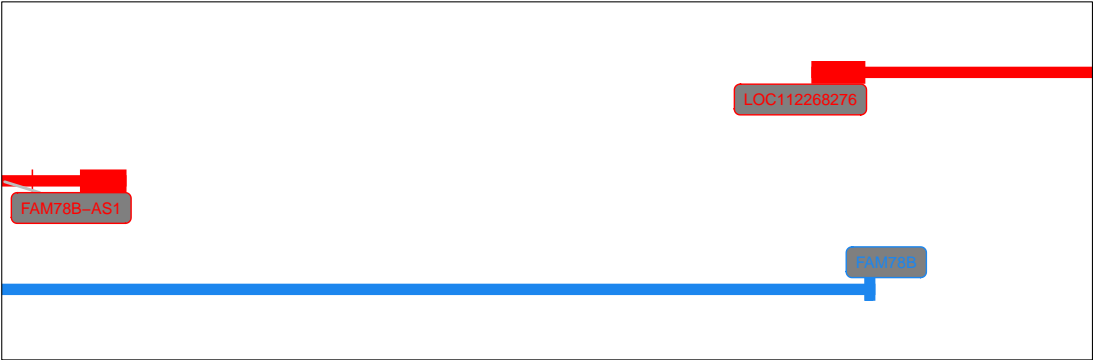
chr1:23573533–23674034 Marker Peak Tracks for: PV.ChC



PV.ChC

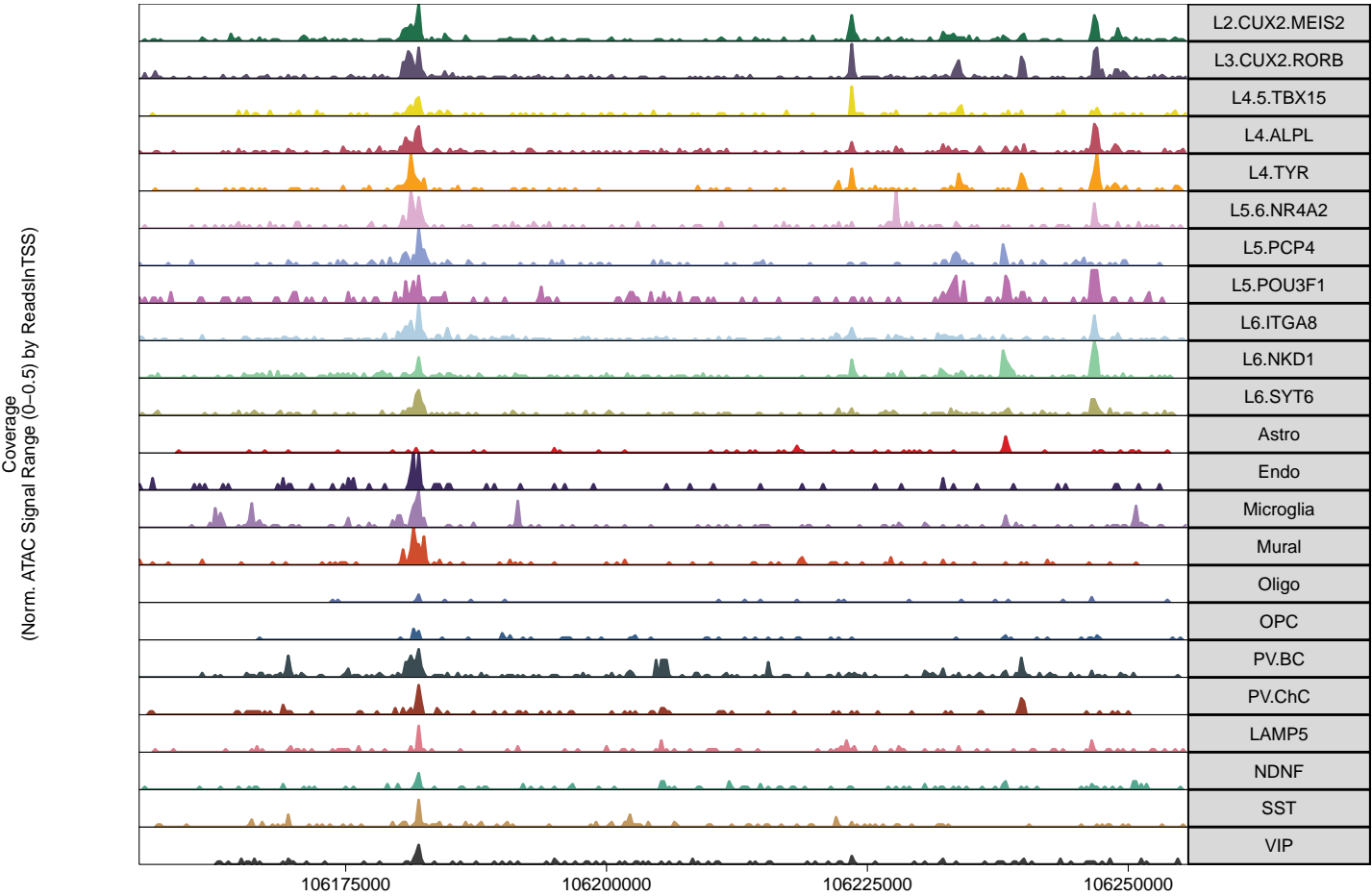


Peaks

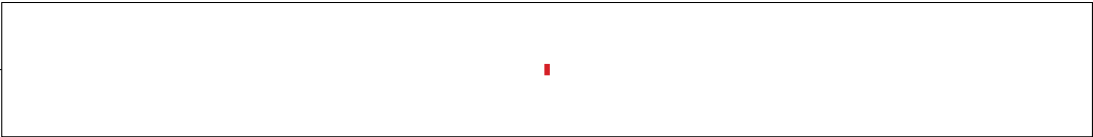


Genes

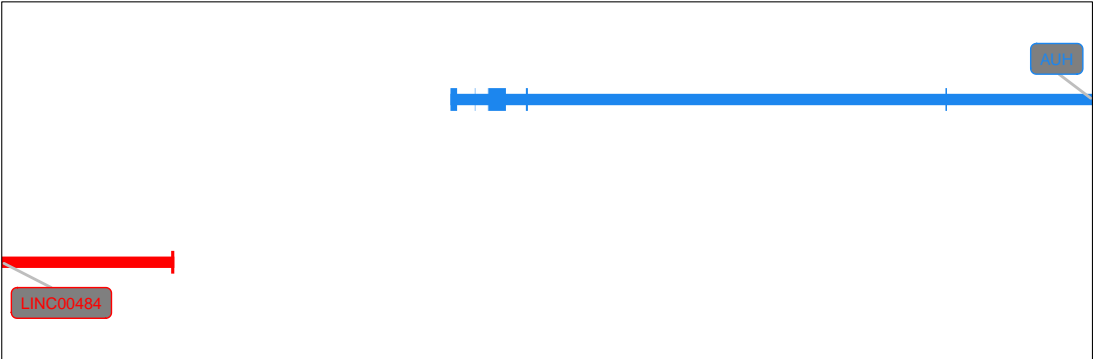
chr15:106155209–106255710 Marker Peak Tracks for: PV.ChC



PV.ChC

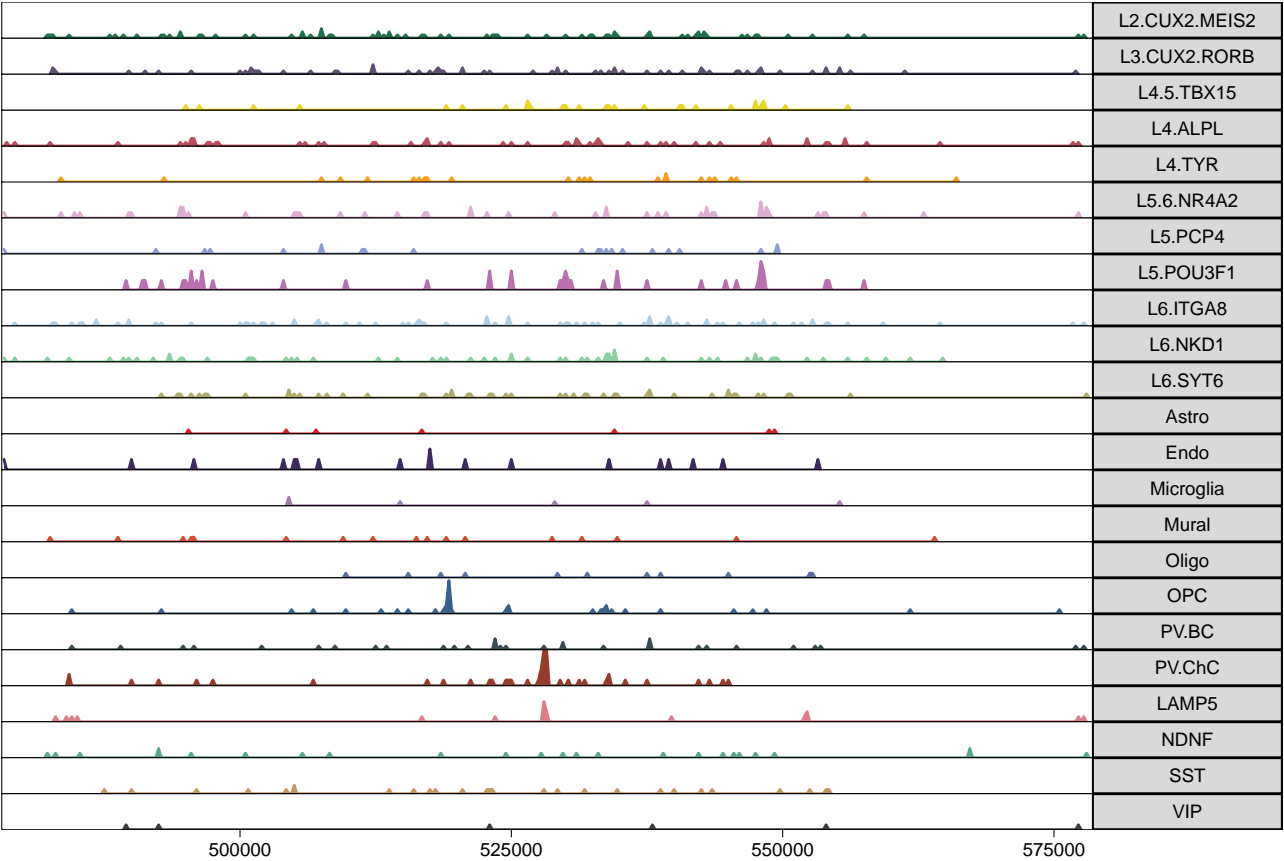


Peaks



Genes

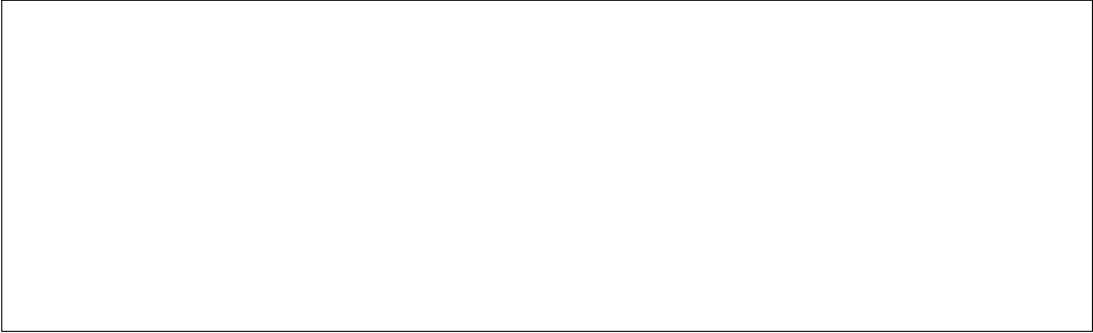
Coverage  
(Norm. ATAC Signal Range (0–0.28) by ReadsInTSS)



PV.ChC

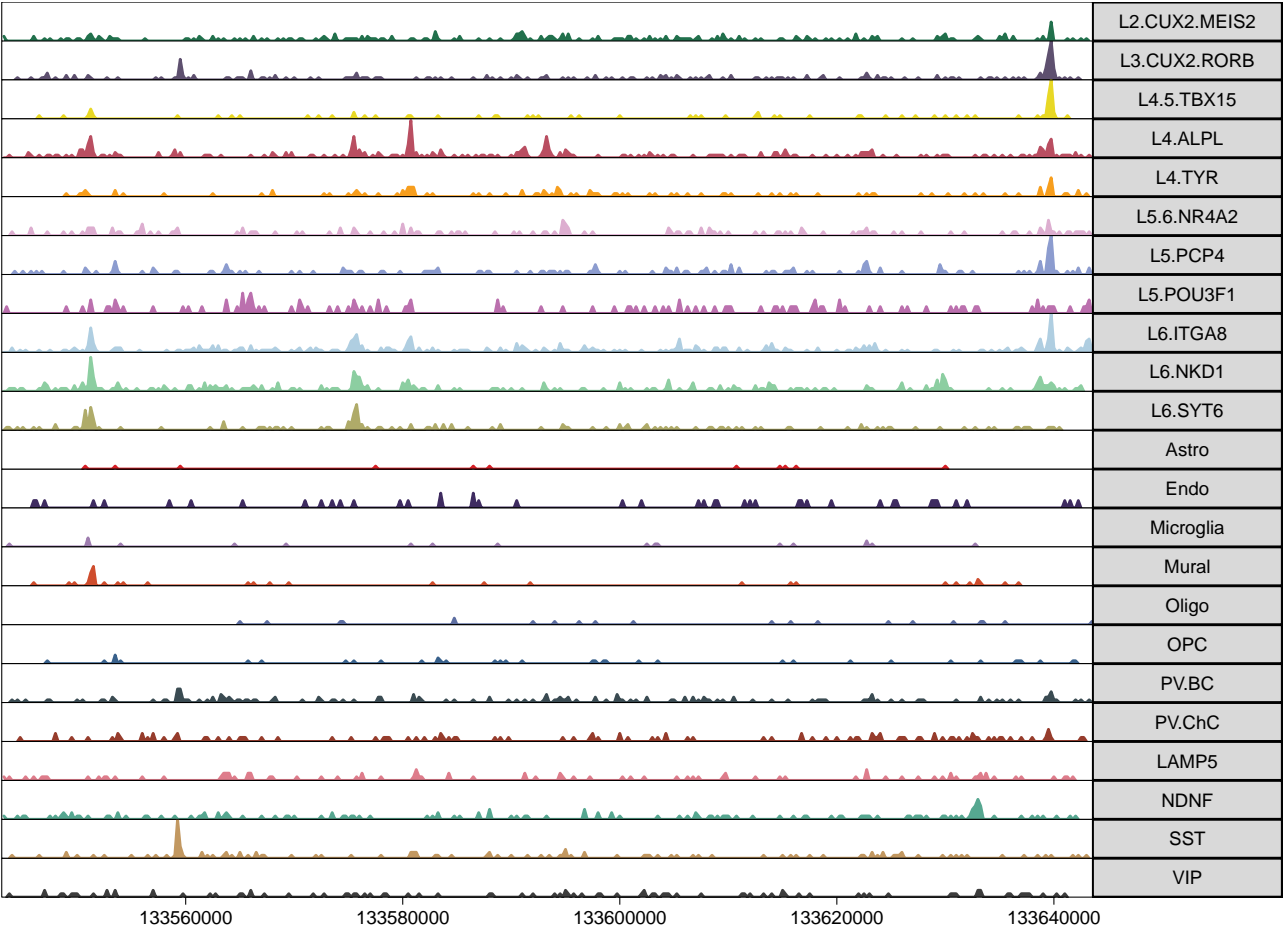


Peaks

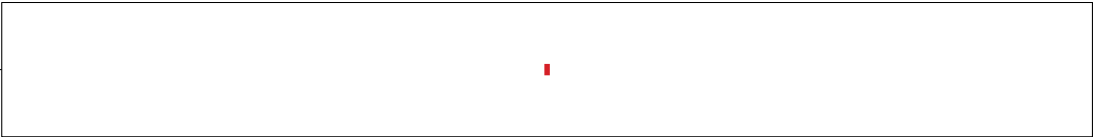


GeneTrack

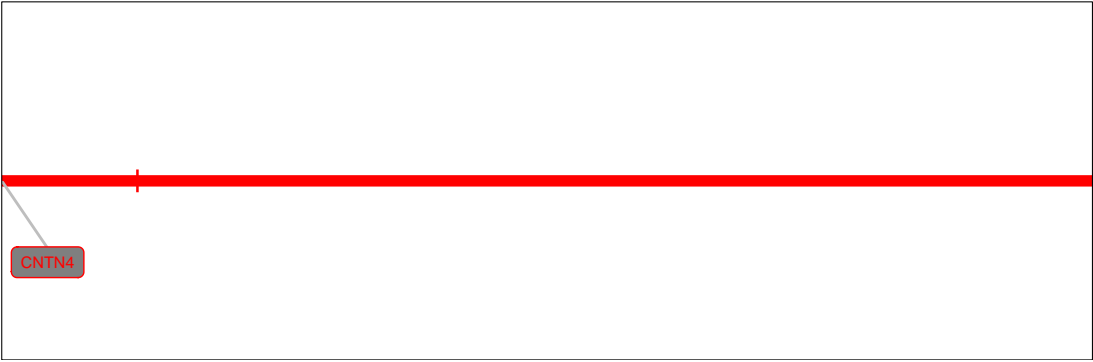
Coverage  
(Norm. ATAC Signal Range (0-0.42) by ReadsInTSS)



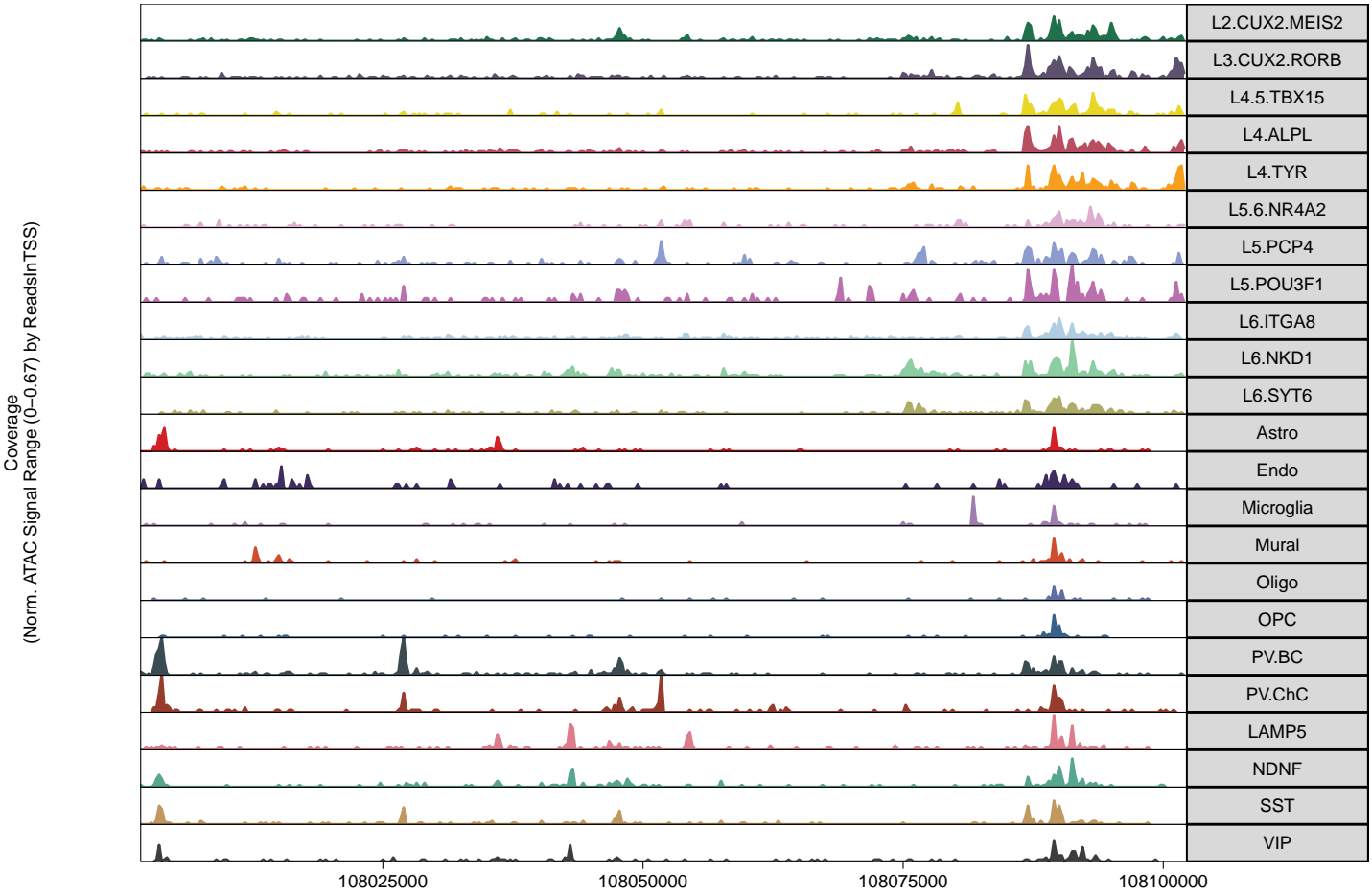
PV.ChC



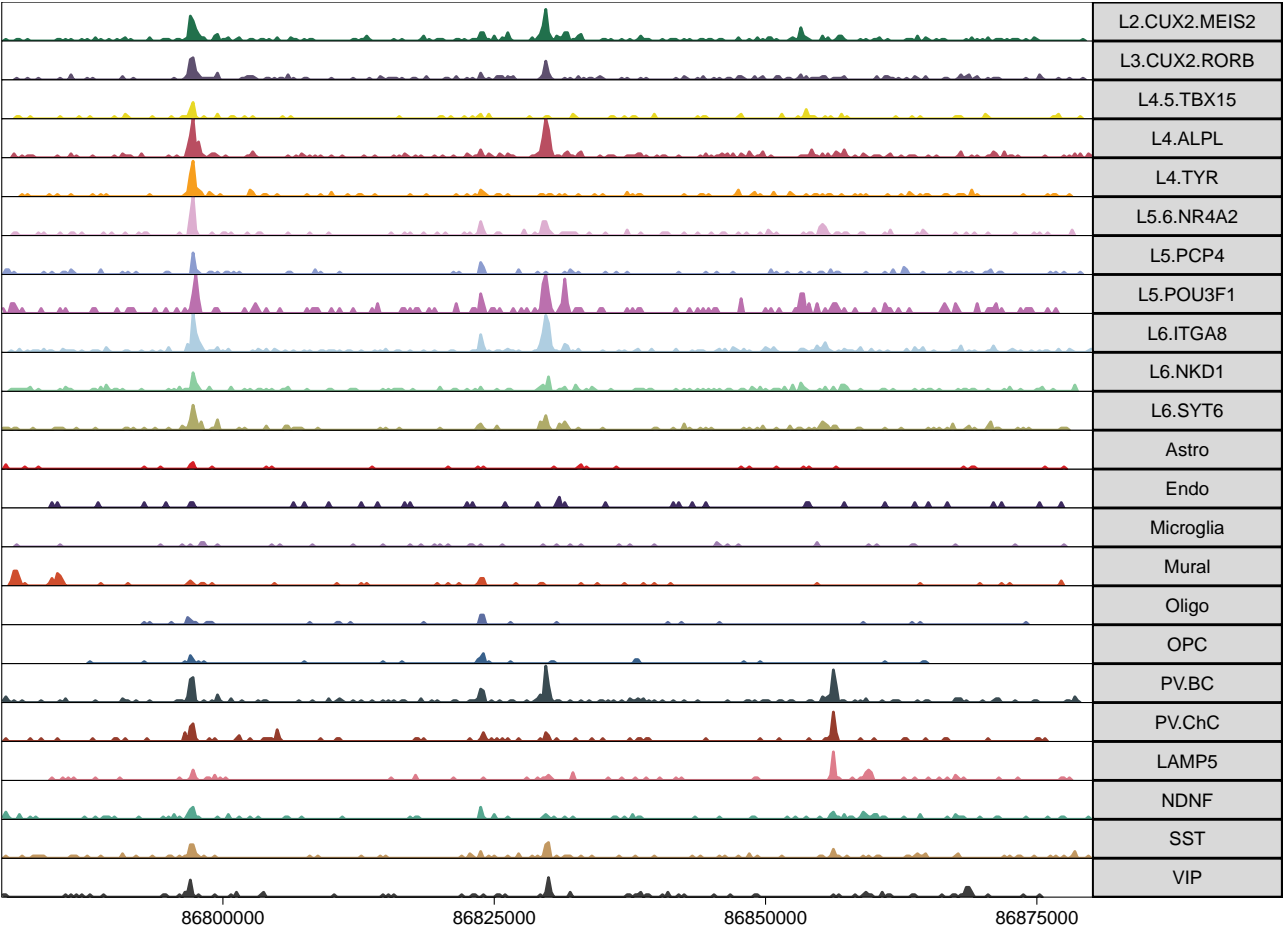
Peaks



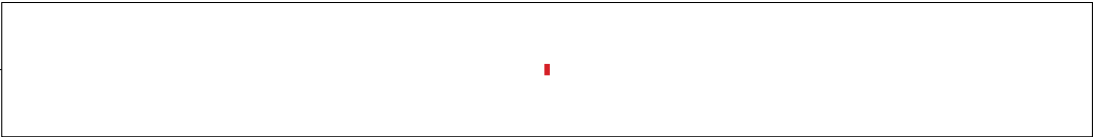
Genes



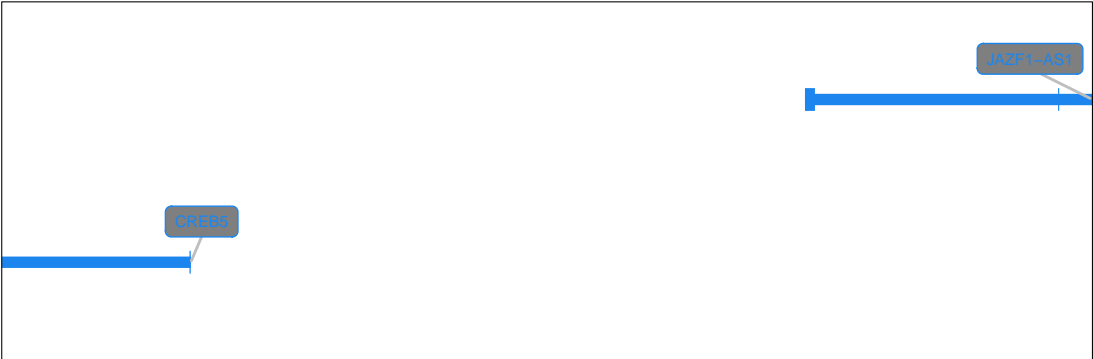
Coverage  
(Norm. ATAC Signal Range (0–0.58) by ReadsInTSS)



PV.ChC

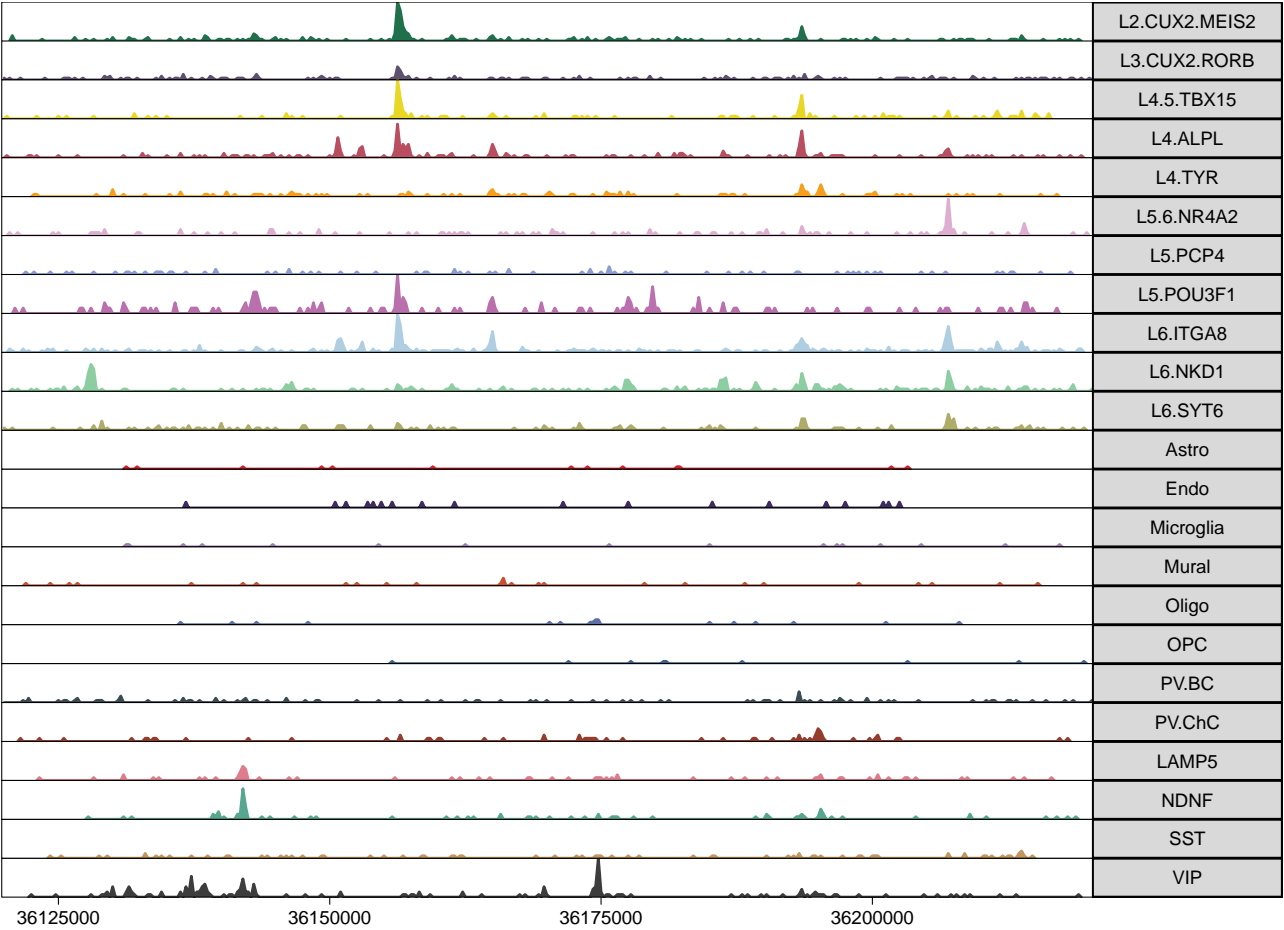


Peaks

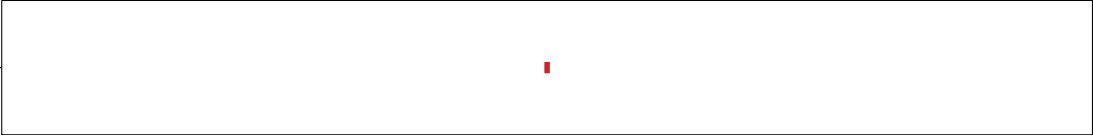


Genes

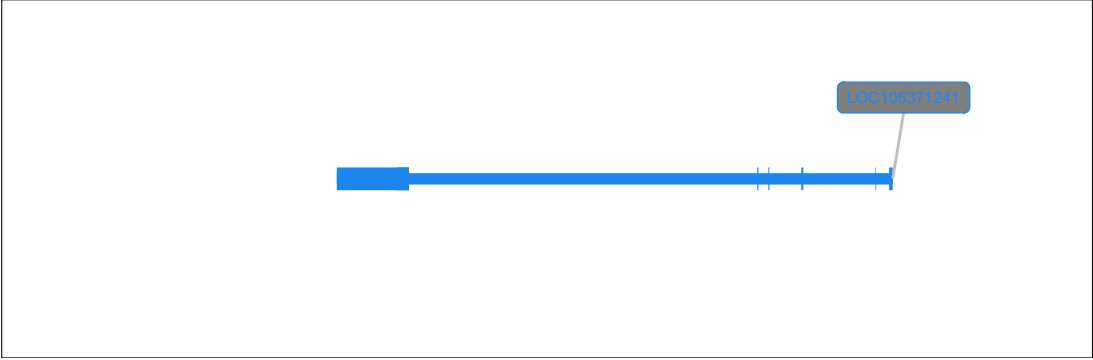
Coverage  
(Norm. ATAC Signal Range (0–0.53) by ReadsInTSS)



PV.ChC




Peaks



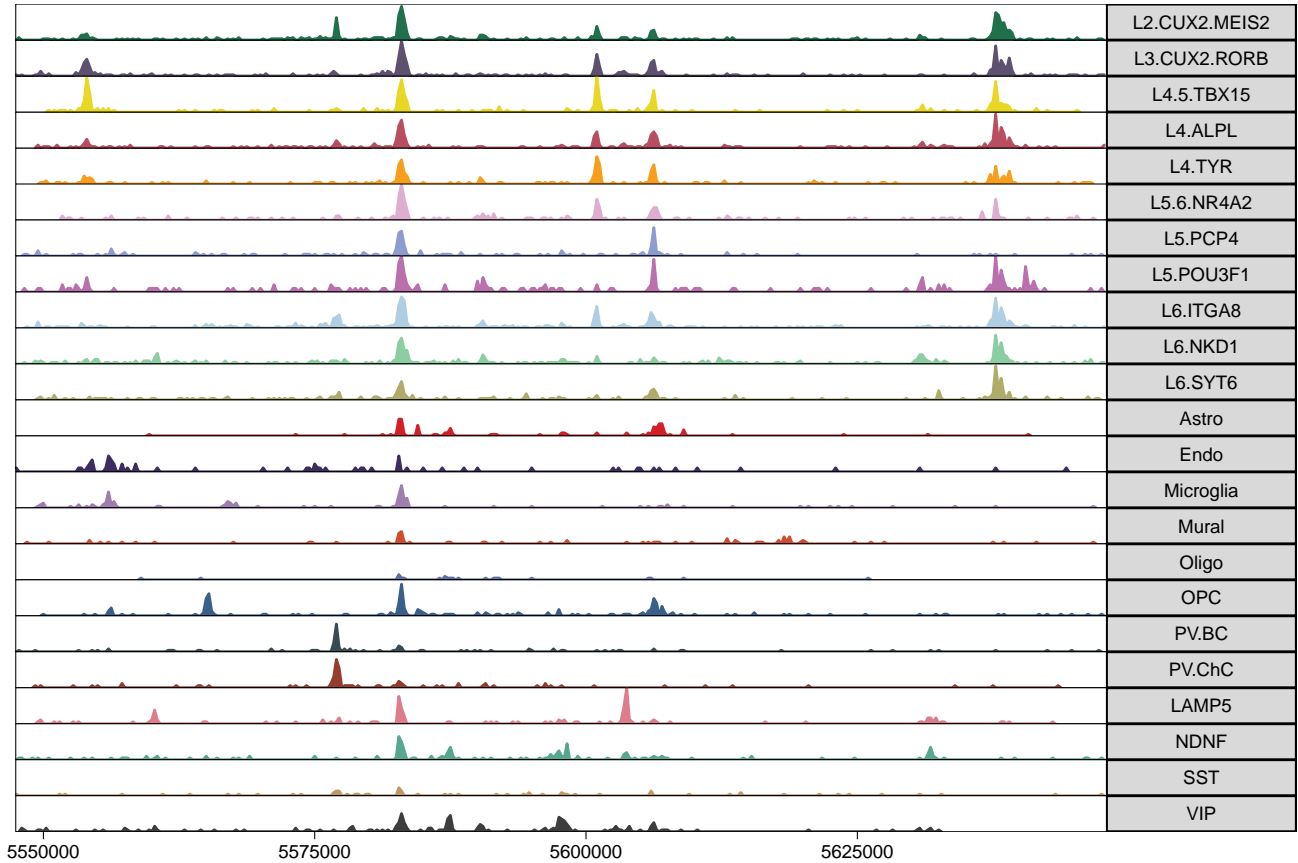
Genes

	L2.CUX2.MEIS2
	L3.CUX2.RORB
	L4.5.TBX15
	L4.ALPL
	L4.TYR
	L5.6.NR4A2
	L5.PCP4
	L5.POU3F1
	L6.ITGA8
	L6.NKD1
	L6.SYT6
	Astro
	Endo
	Microglia
	Mural
	Oligo
	OPC
	PV.BC
	PV.ChC
	LAMP5
	NDNF
	SST
	VIP





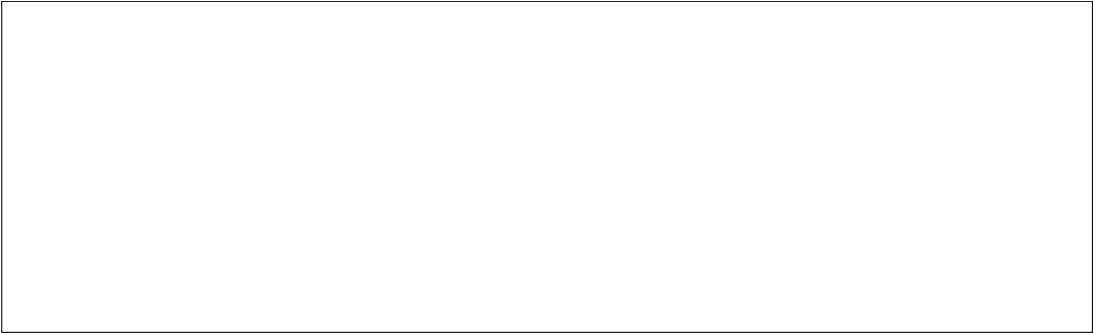
Coverage  
(Norm. ATAC Signal Range (0–0.73) by ReadsInTSS)



PV.ChC

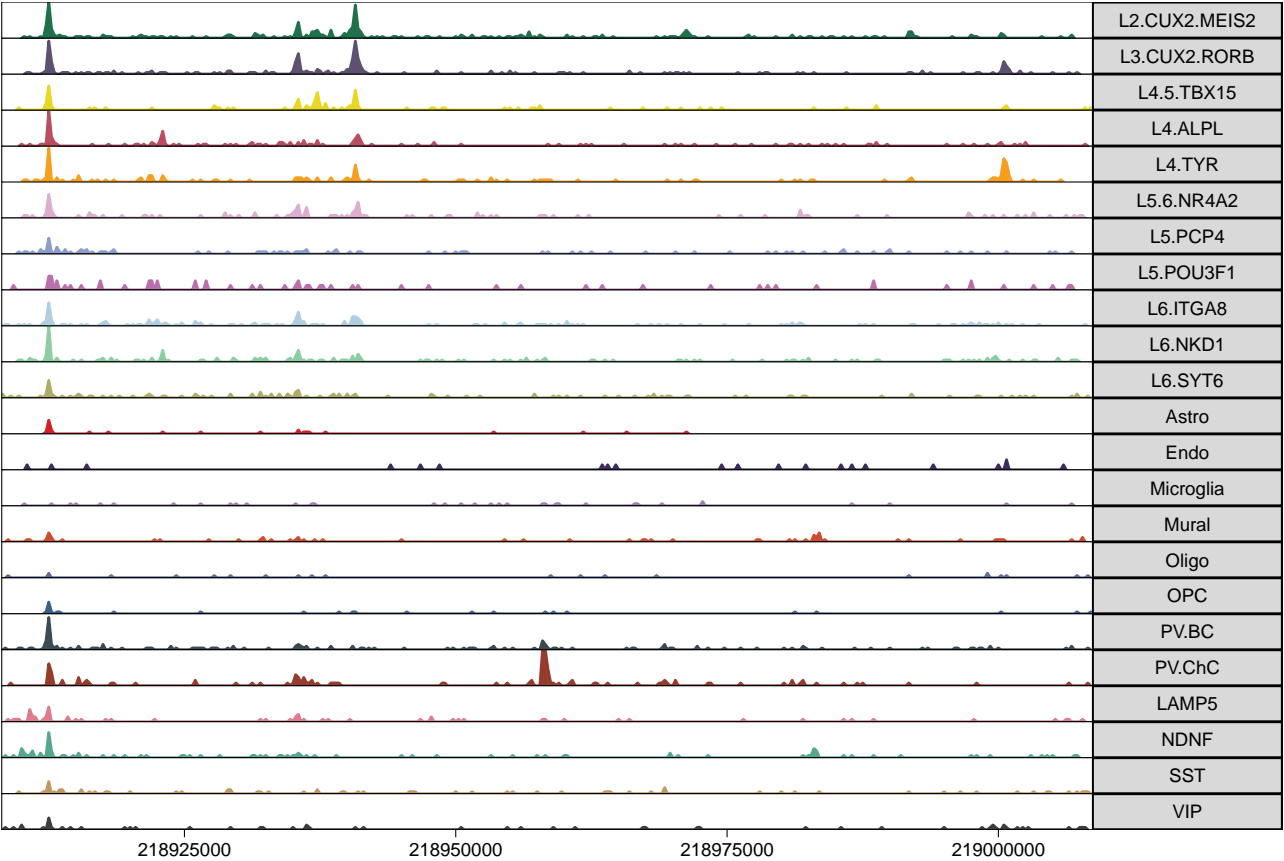


Peaks



GeneTrack

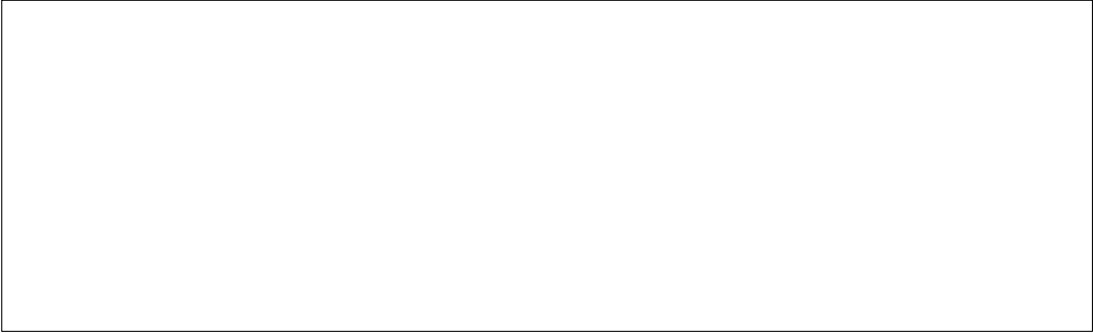
Coverage  
(Norm. ATAC Signal Range (0–0.57) by ReadsInTSS)



PV.ChC

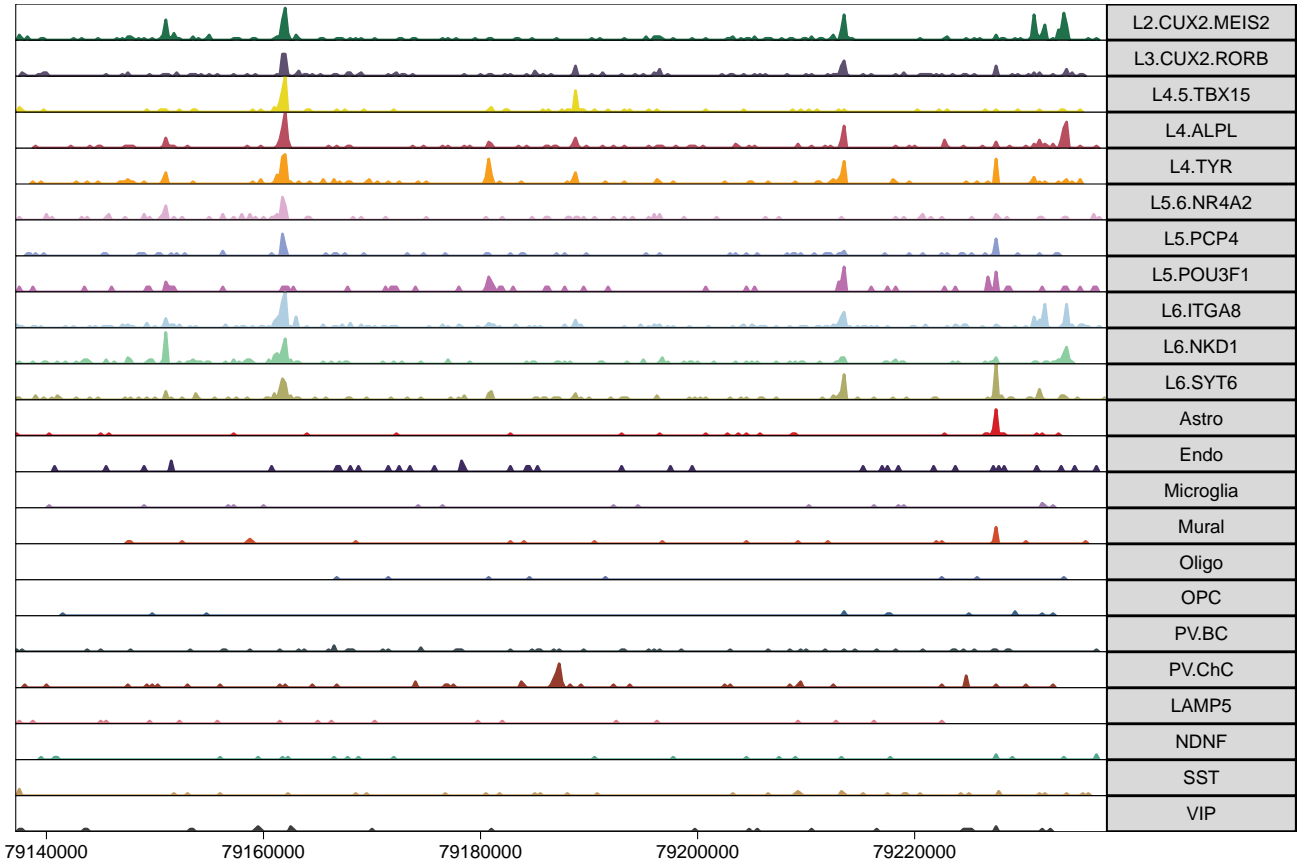


Peaks



GeneTrack

Coverage  
(Norm. ATAC Signal Range (0–0.53) by ReadsInTSS)



PV.ChC

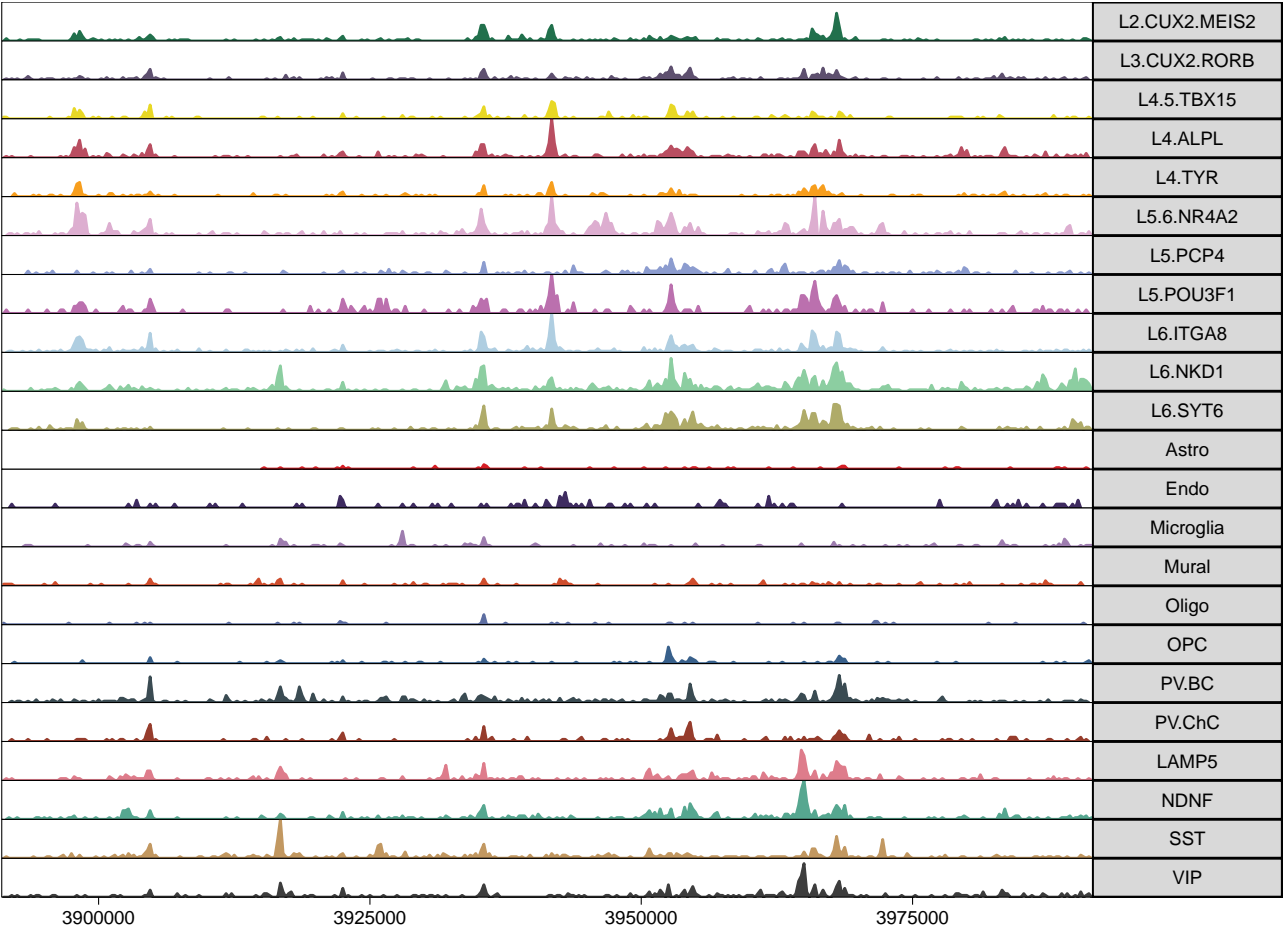


Peaks

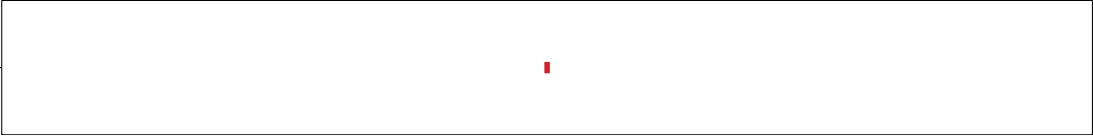
GeneTrack

chr4:3891075–3991576 Marker Peak Tracks for: PV.ChC

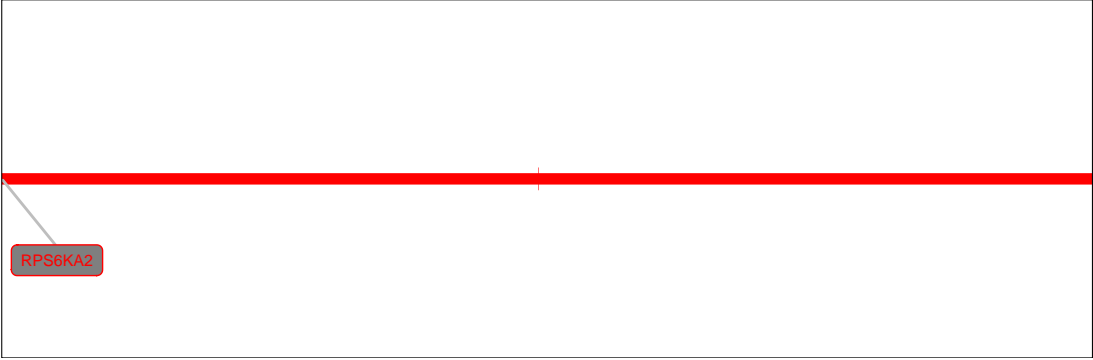
Coverage  
(Norm. ATAC Signal Range (0–0.8) by ReadsInTSS)



PV.ChC

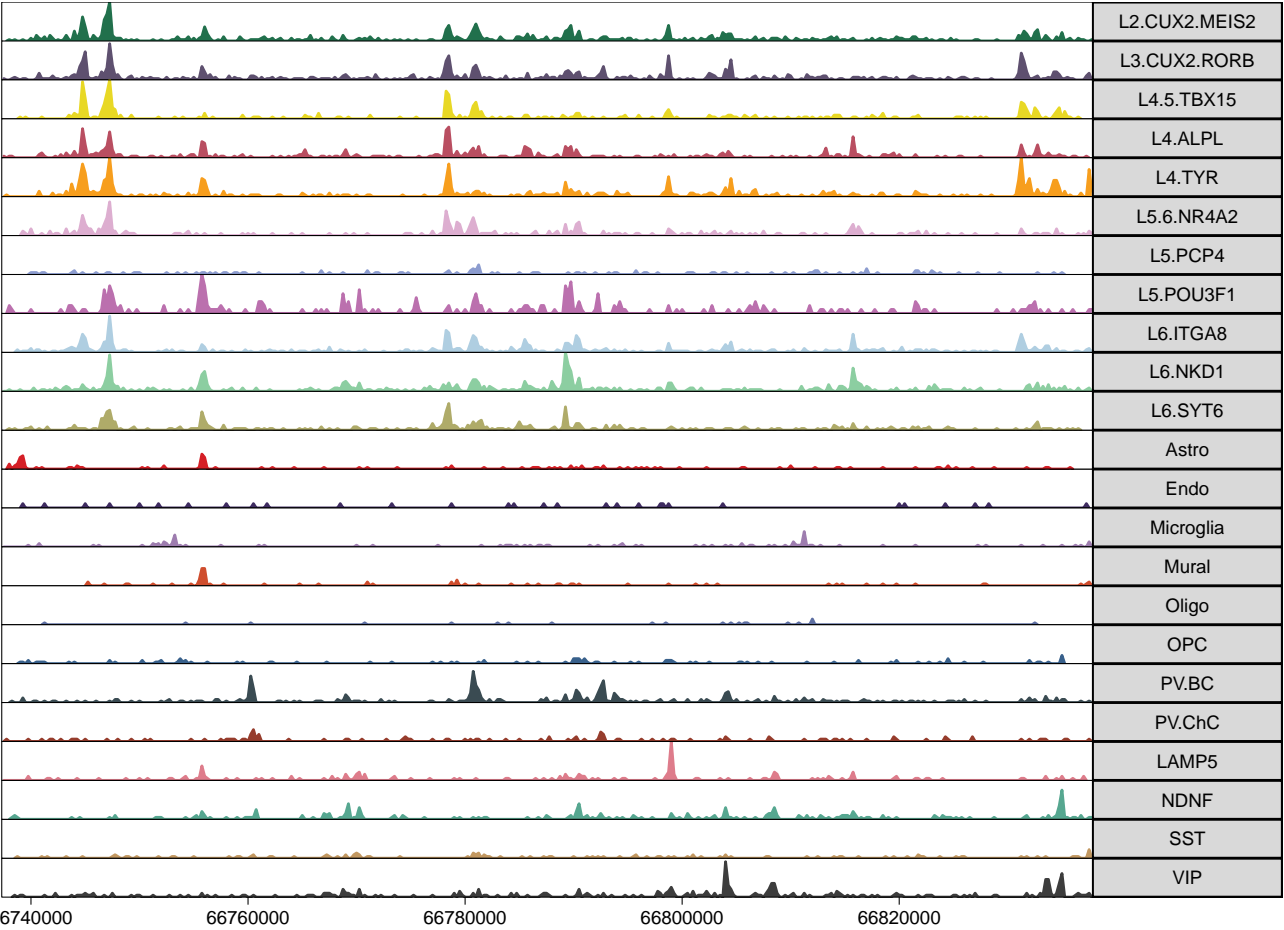


Peaks

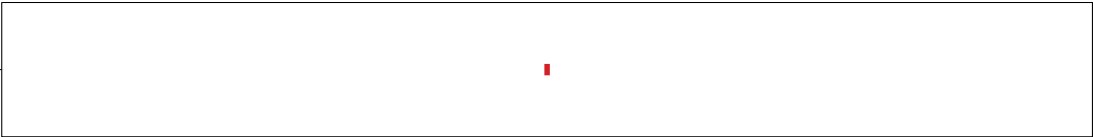


Genes

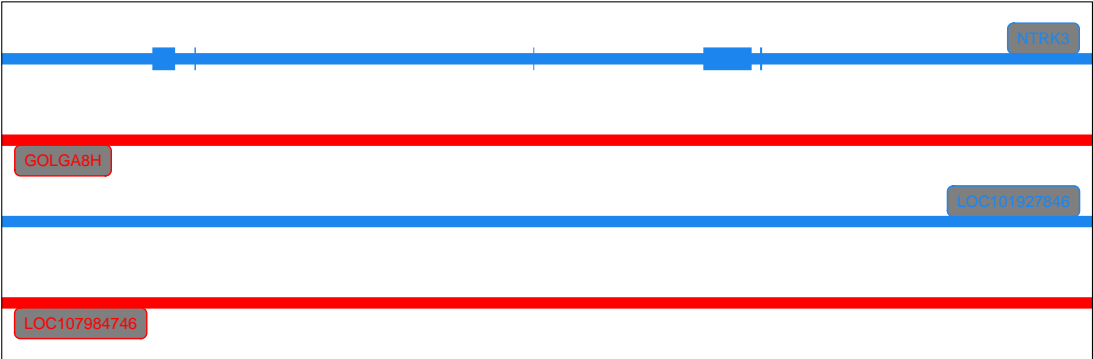
Coverage  
(Norm. ATAC Signal Range (0–0.72) by ReadsInTSS)



PV.ChC

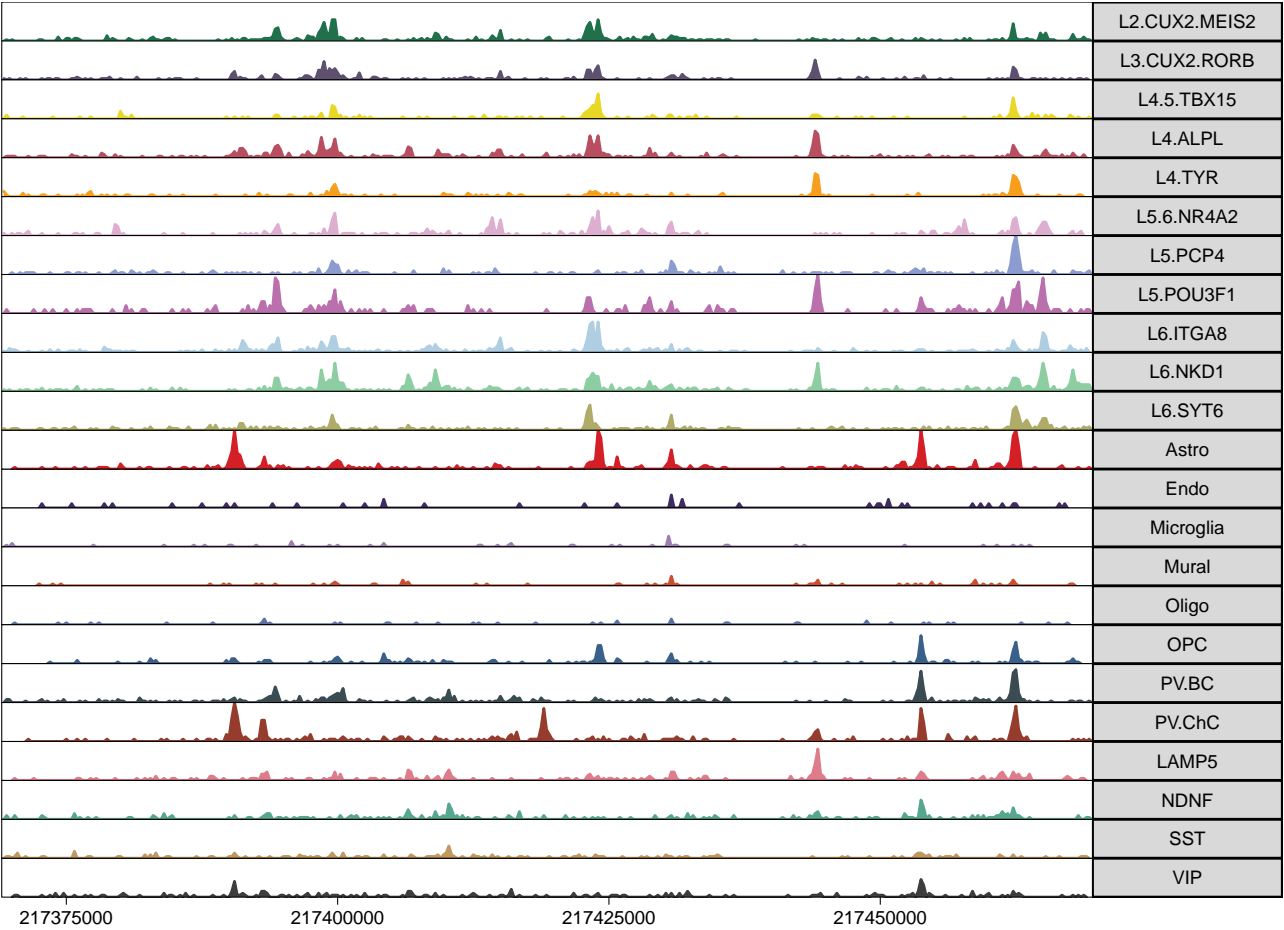


Peaks

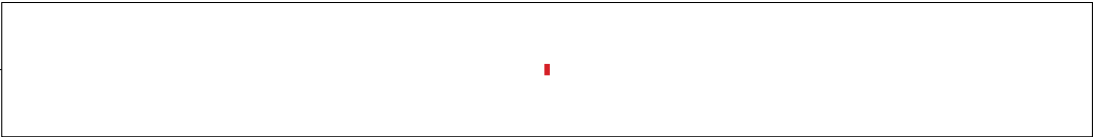


Genes

Coverage  
(Norm. ATAC Signal Range (0–0.72) by ReadsInTSS)



PV.ChC

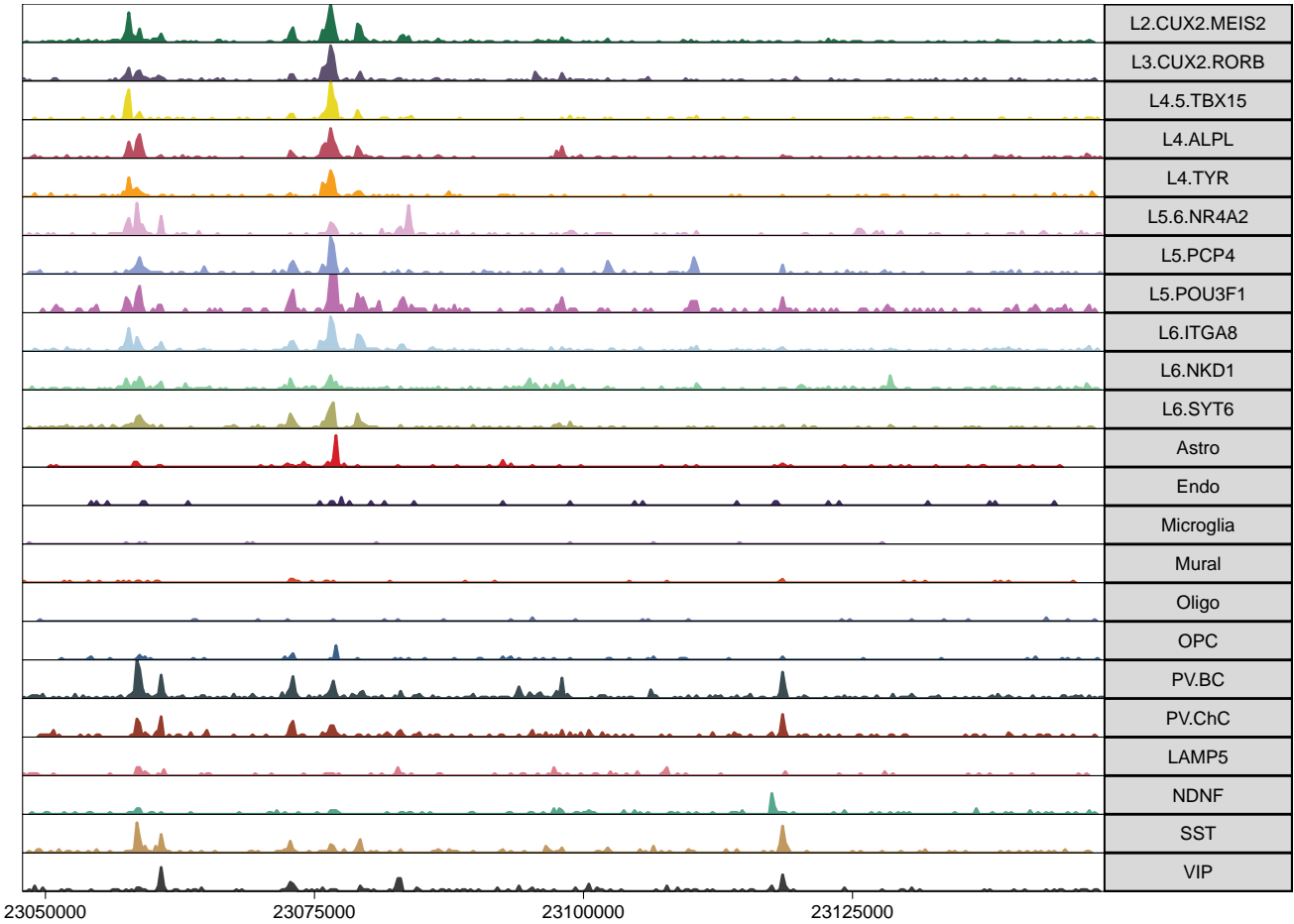


Peaks

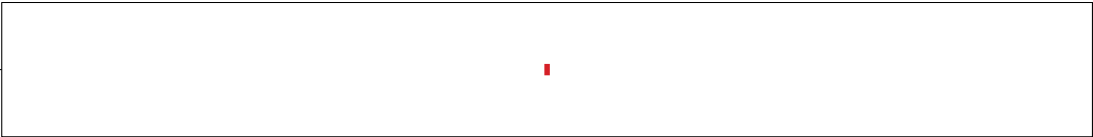


Genes

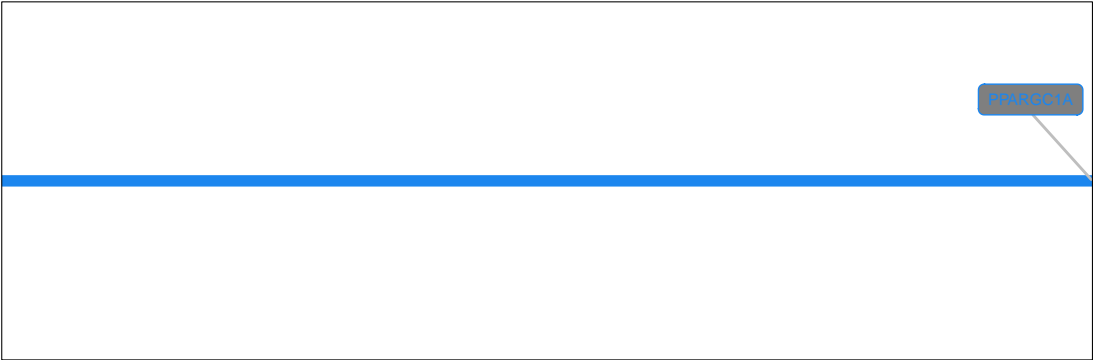
Coverage  
(Norm. ATAC Signal Range (0–0.75) by ReadsInTSS)



PV.ChC

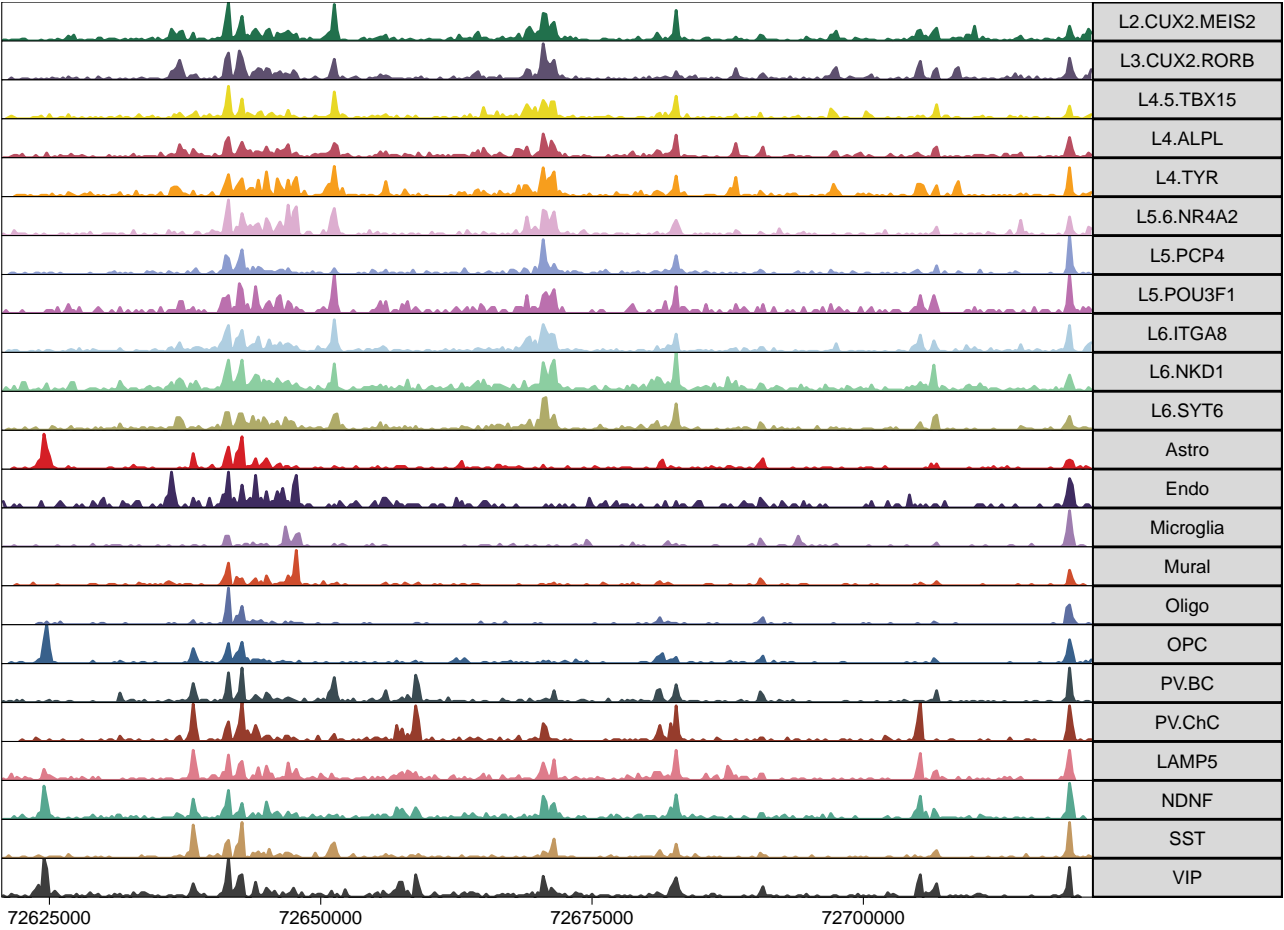


Peaks

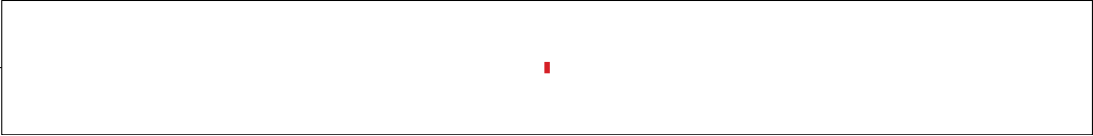


Genes

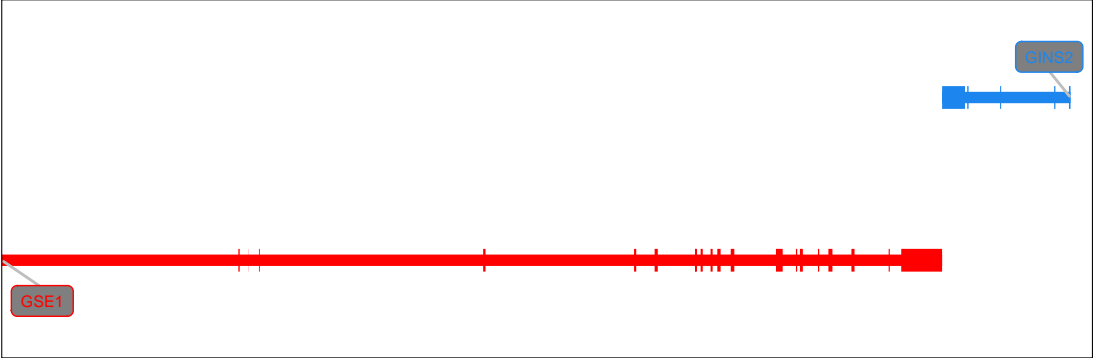
Coverage  
(Norm. ATAC Signal Range (0–0.96) by ReadsInTSS)



PV.ChC



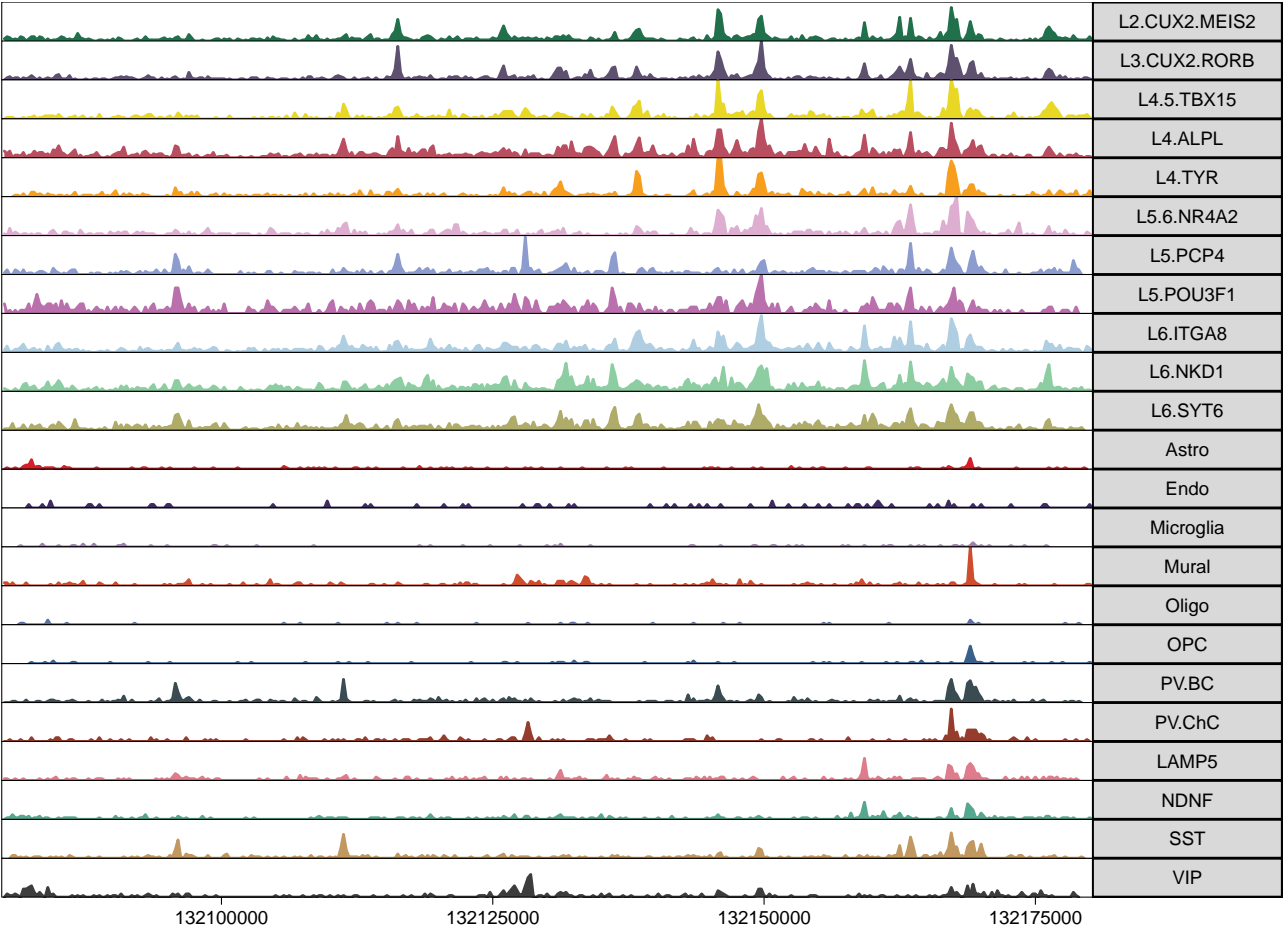
Peaks



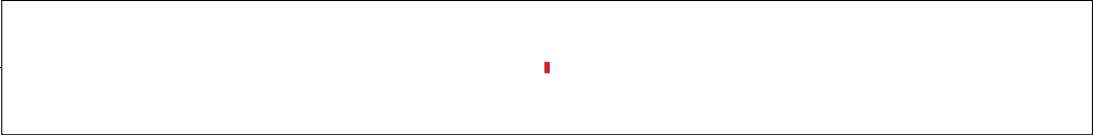
Genes



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



PV.ChC

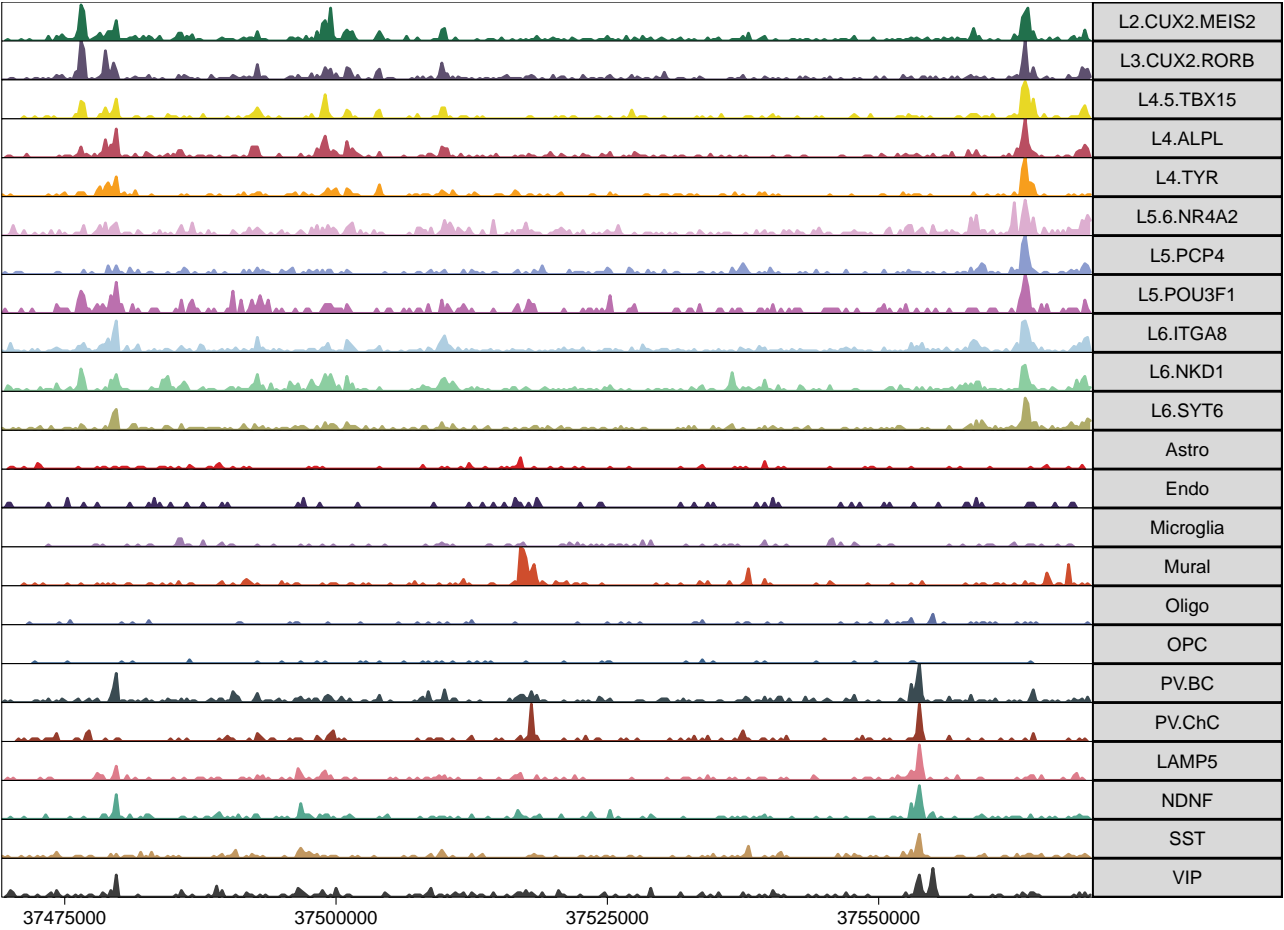


Peaks

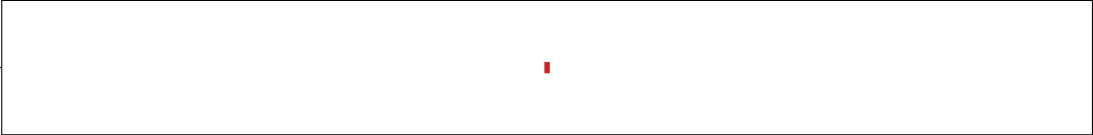


Genes

Coverage  
(Norm. ATAC Signal Range (0–0.64) by ReadsInTSS)



PV.ChC

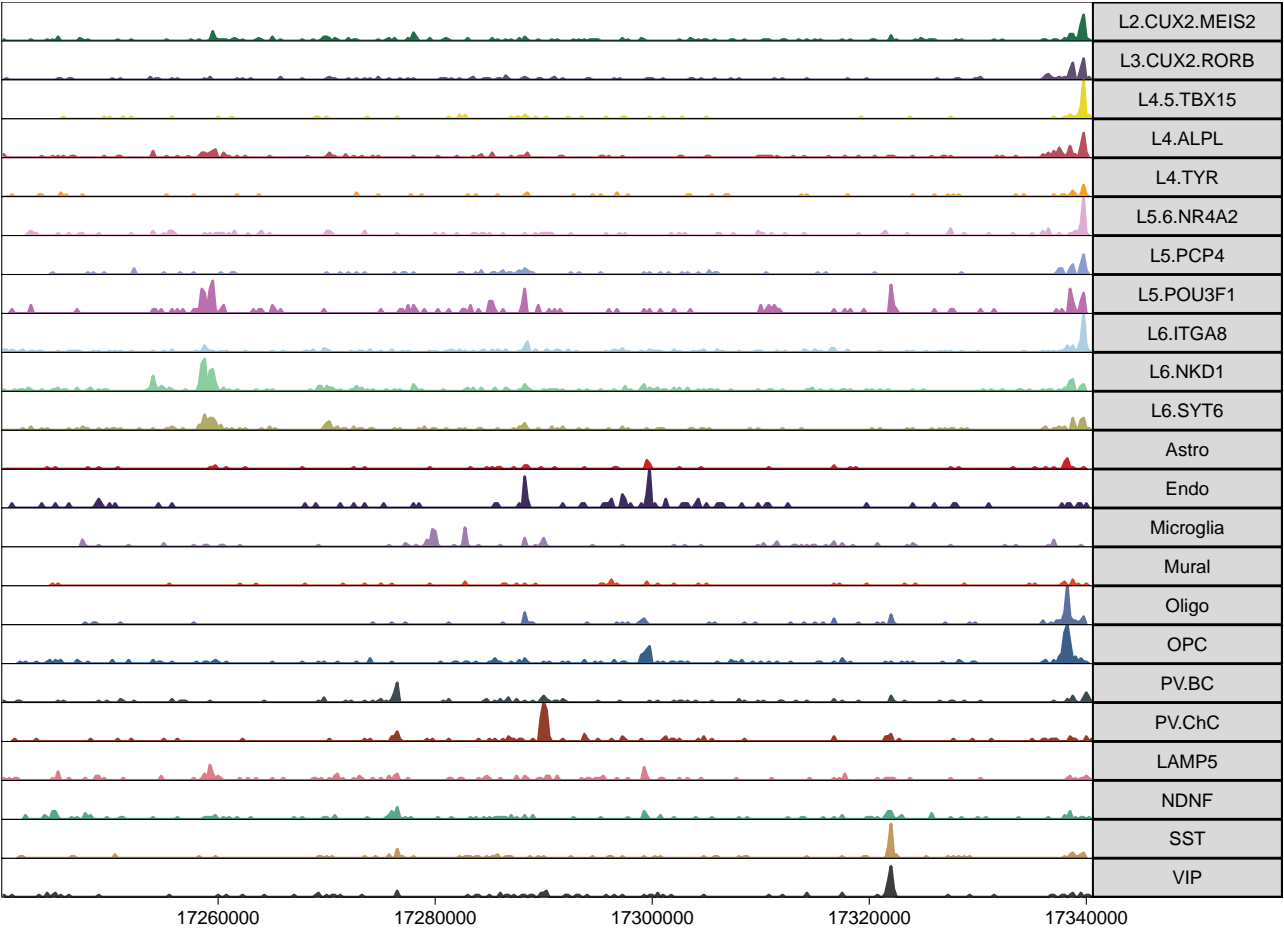


Peaks

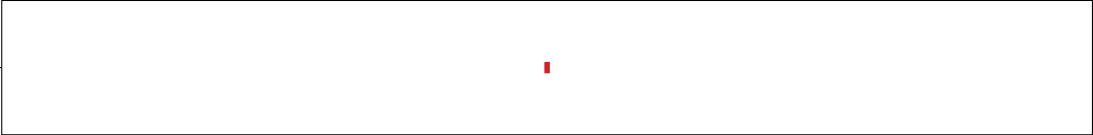


Genes

Coverage  
(Norm. ATAC Signal Range (0–0.7) by ReadsInTSS)



PV.ChC

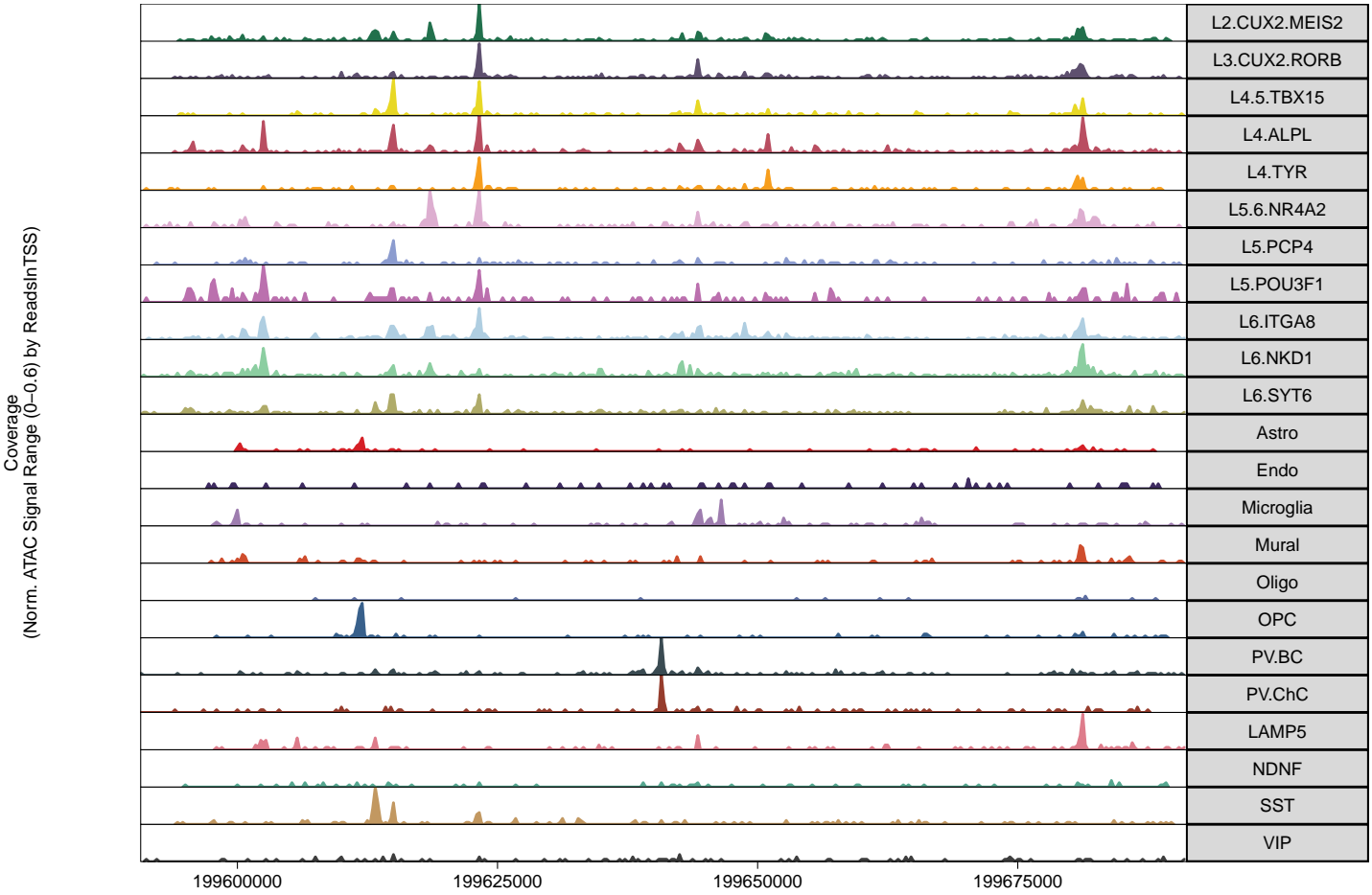


Peaks

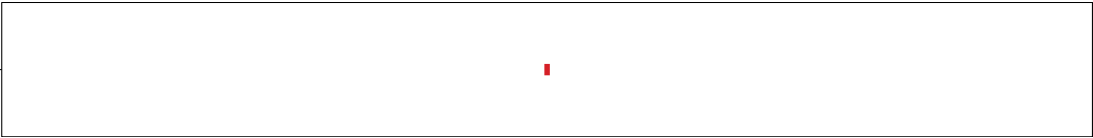


Genes

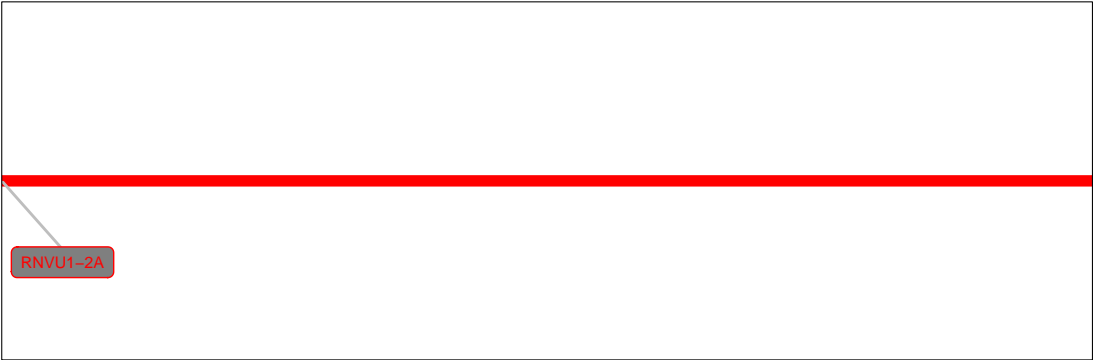
chr1:199590708–199691209 Marker Peak Tracks for: PV.ChC



PV.ChC

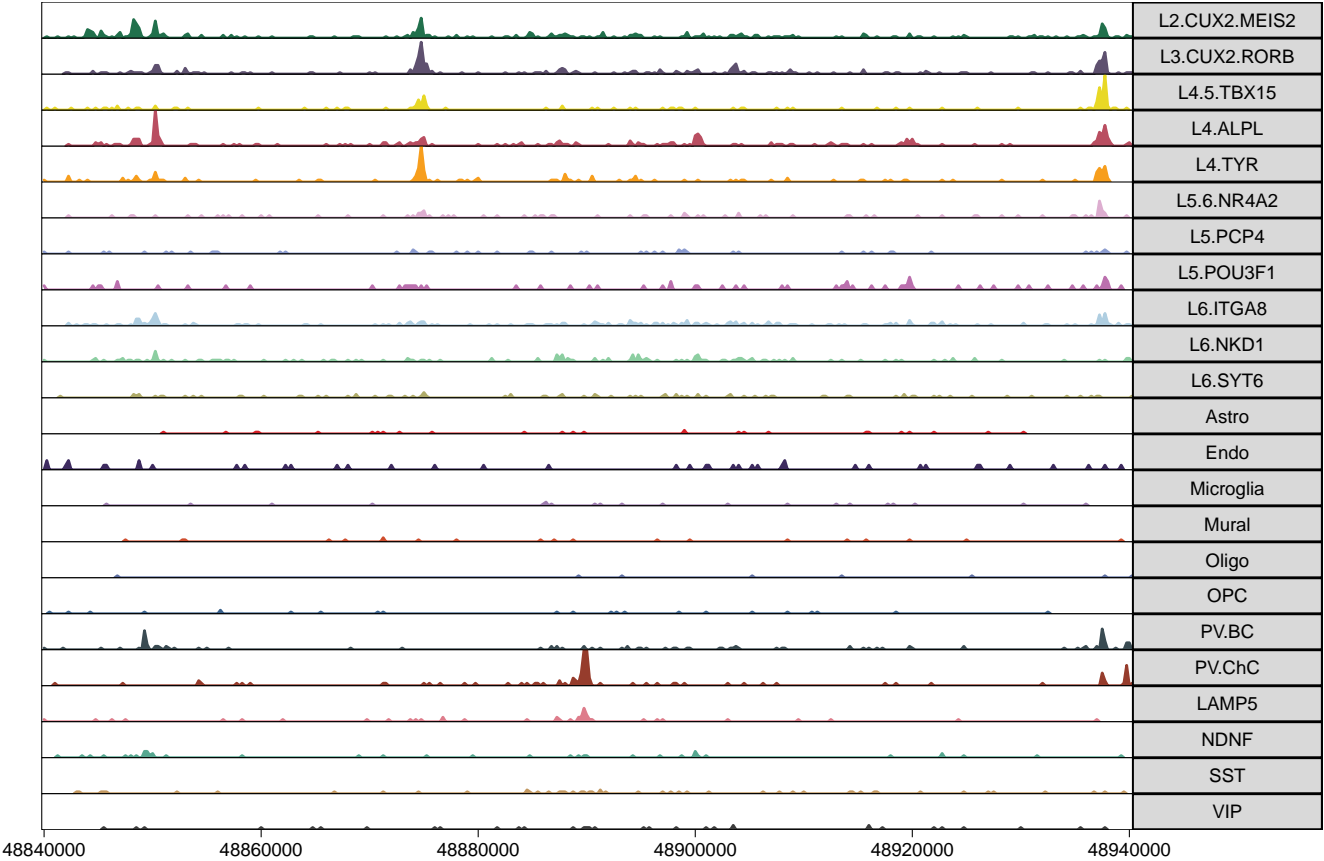


Peaks



Genes

Coverage  
(Norm. ATAC Signal Range (0–0.62) by ReadsInTSS)

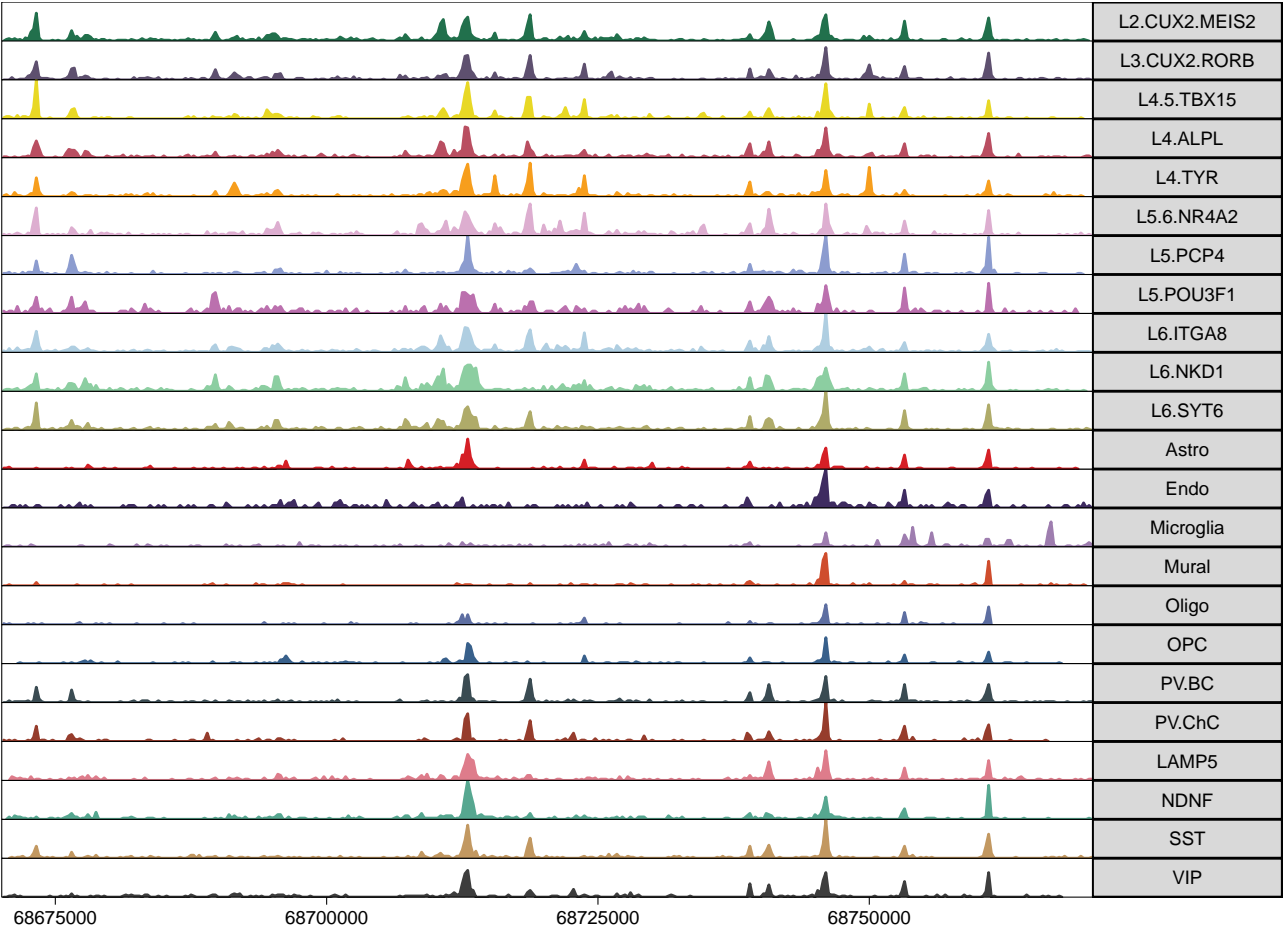


PV.ChC

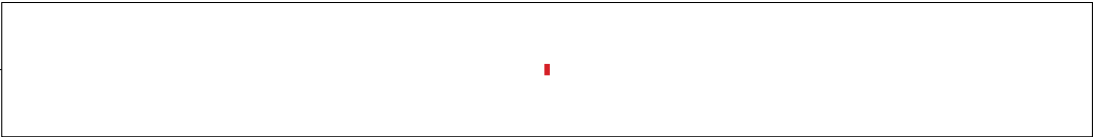
Peaks

GeneTrack

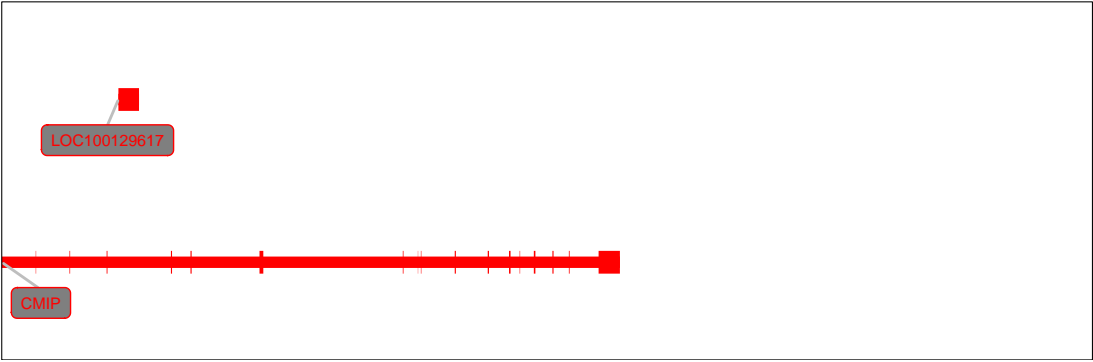
Coverage  
(Norm. ATAC Signal Range (0–1.23) by ReadsInTSS)



PV.ChC

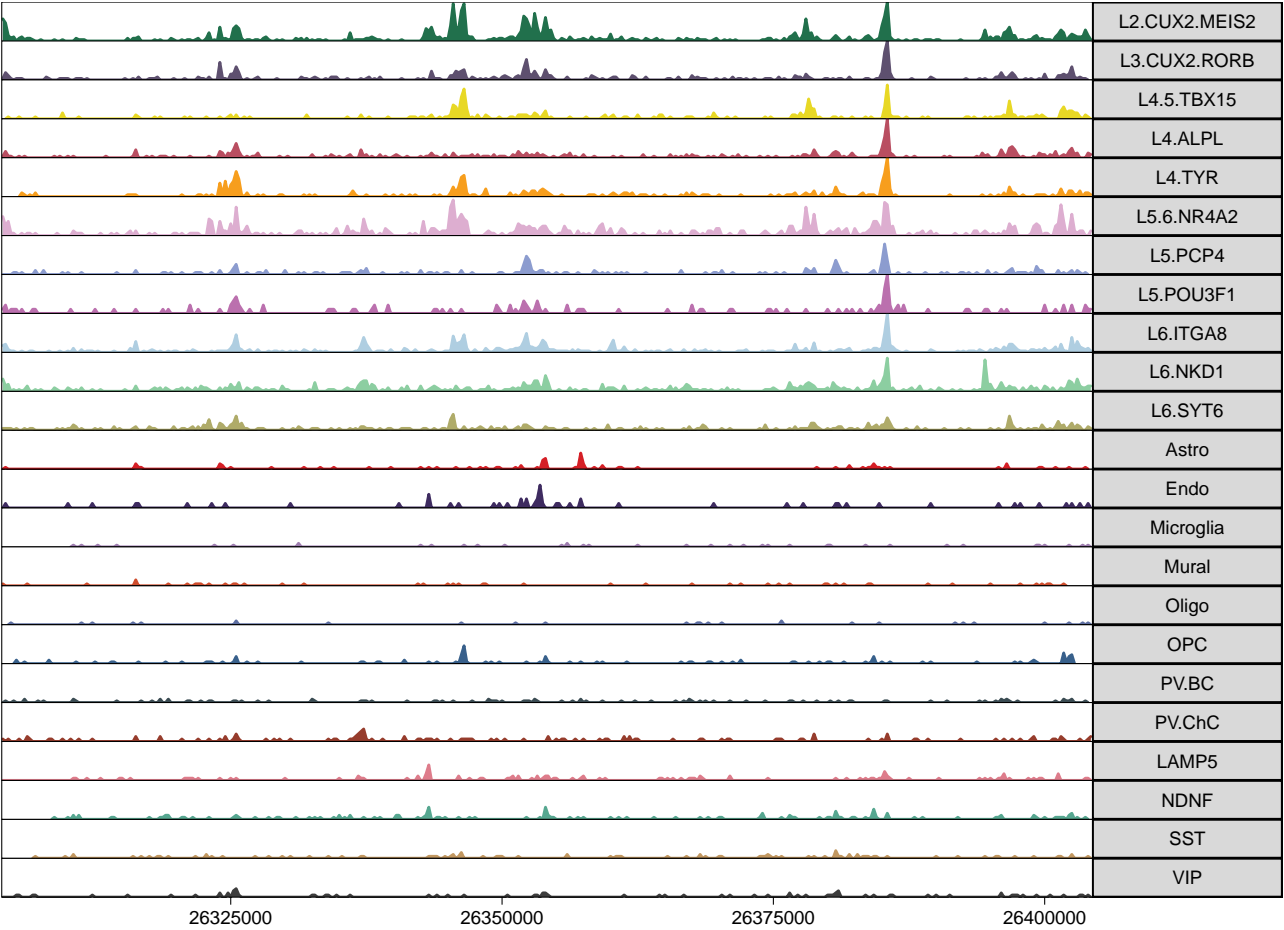


Peaks

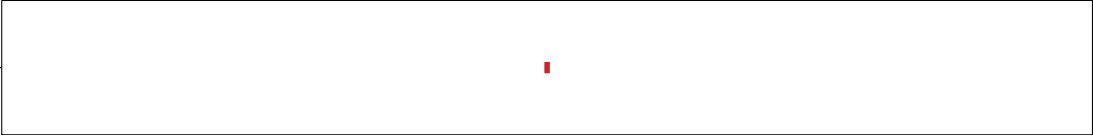


Genes

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



PV.ChC

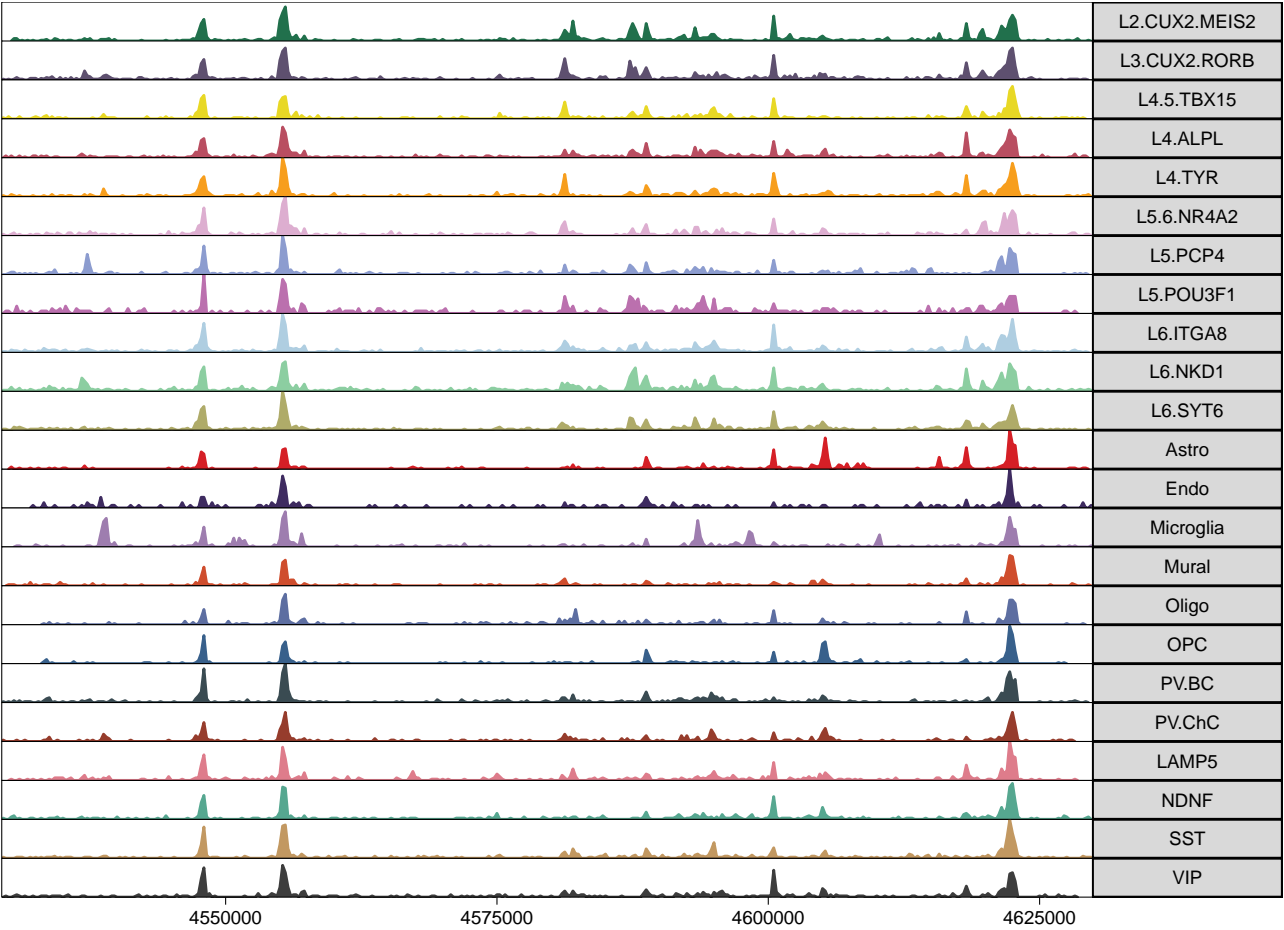


Peaks

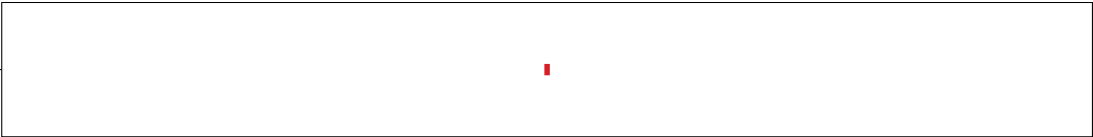


Genes

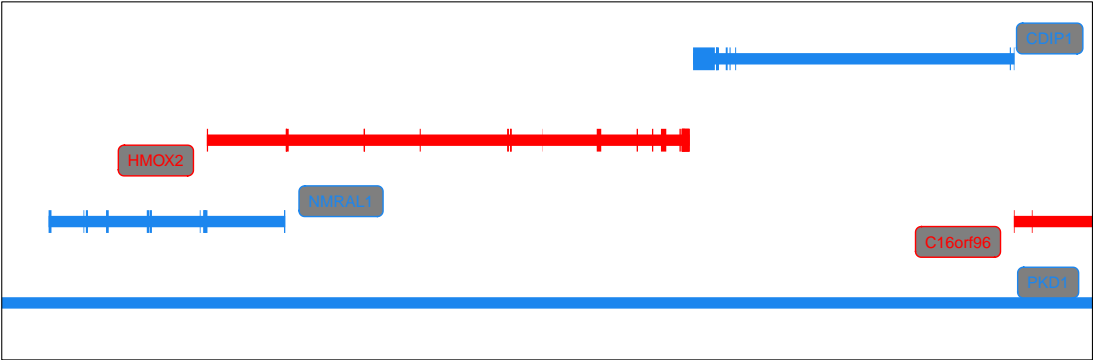
Coverage  
(Norm. ATAC Signal Range (0–1.17) by ReadsInTSS)



PV.ChC



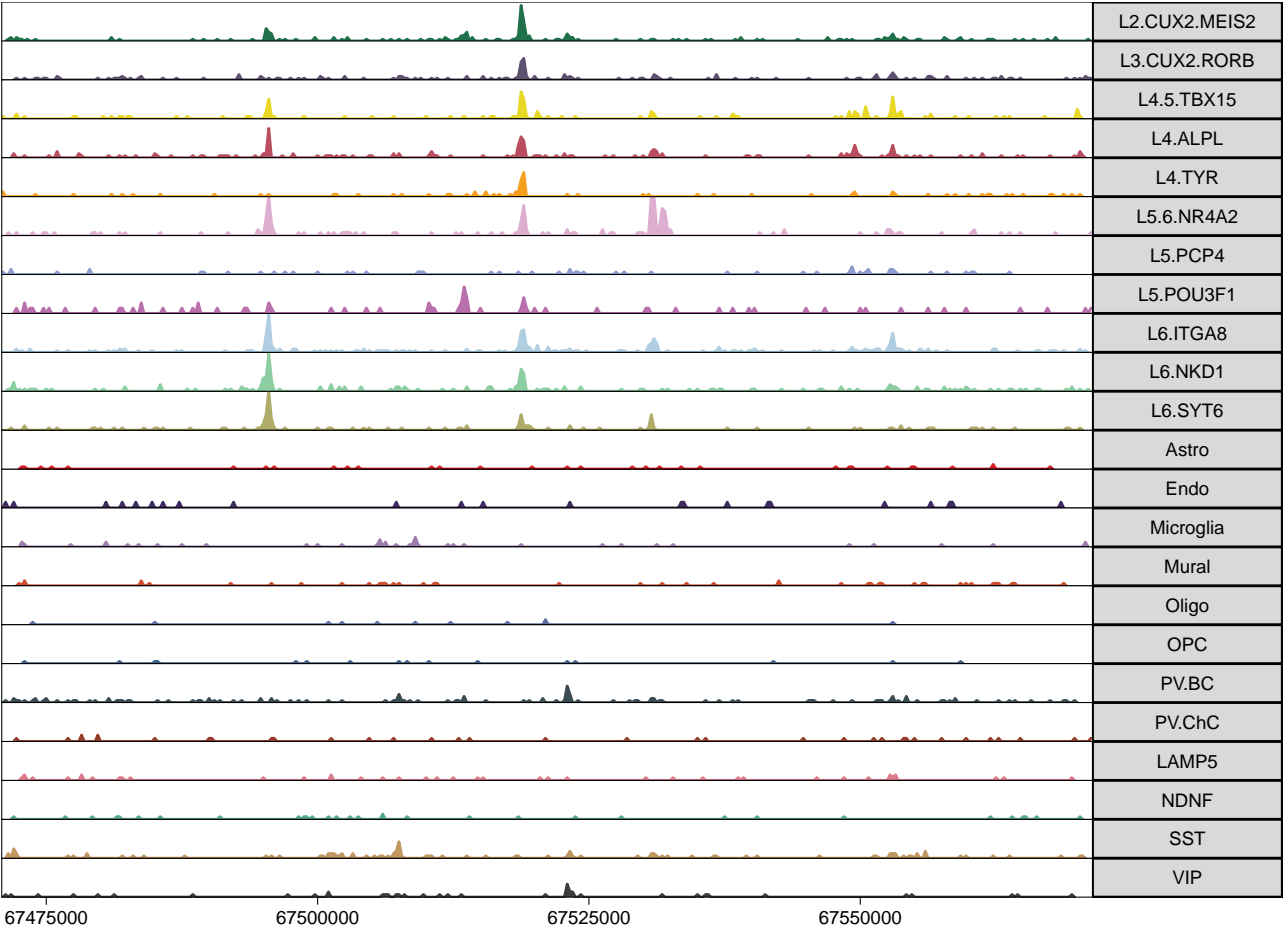
Peaks



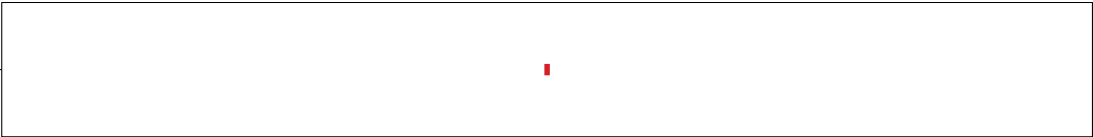
Genes



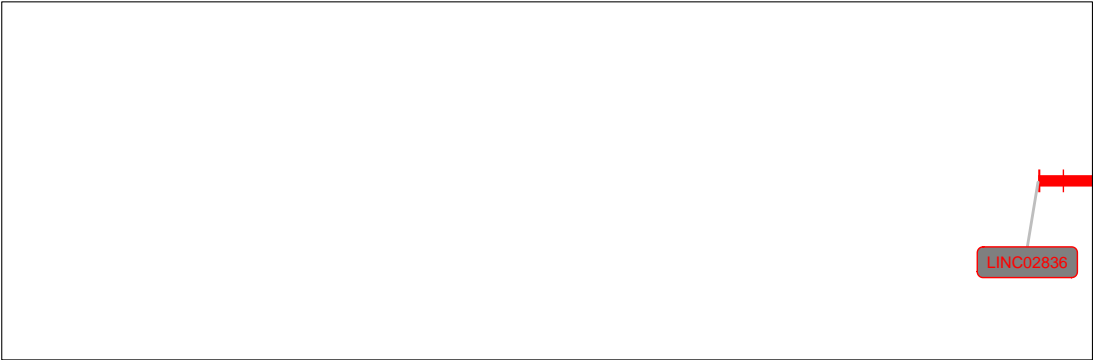
Coverage  
(Norm. ATAC Signal Range (0–0.54) by ReadsInTSS)



PV.ChC



Peaks



Genes