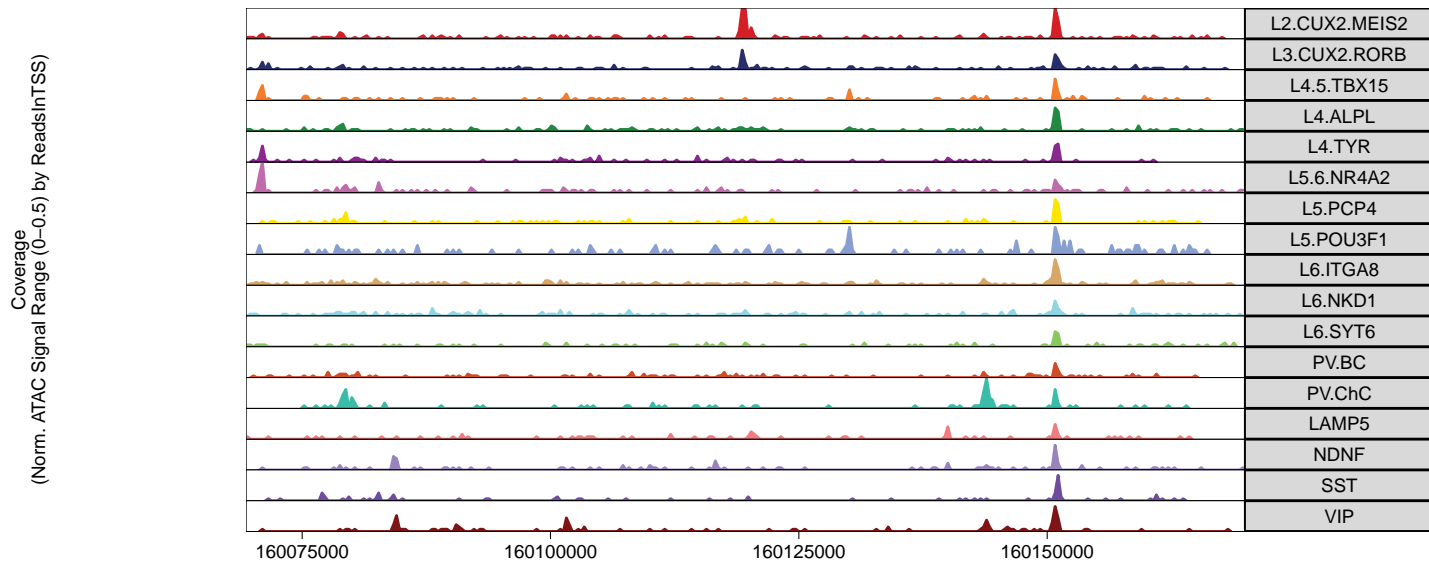


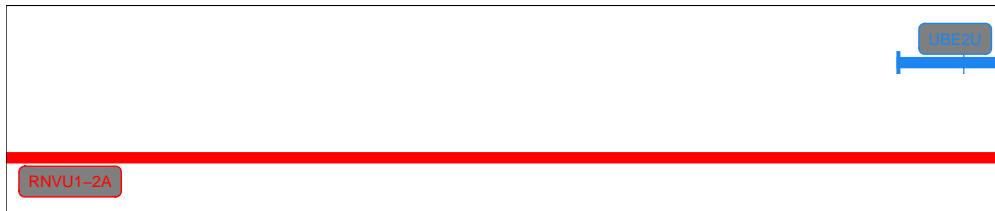
## chr1:160069359–160169860 Marker Peak Tracks for: L2.CUX2.MEIS2



L2.CUX2.MEIS2

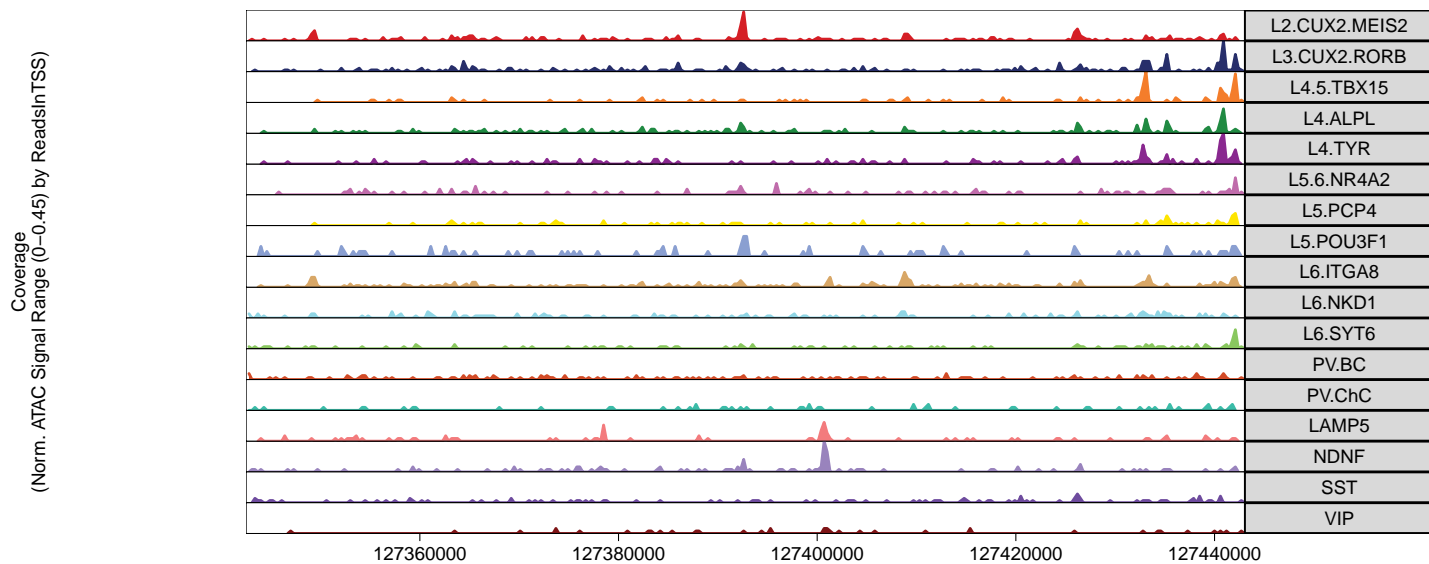


Peaks



Genes

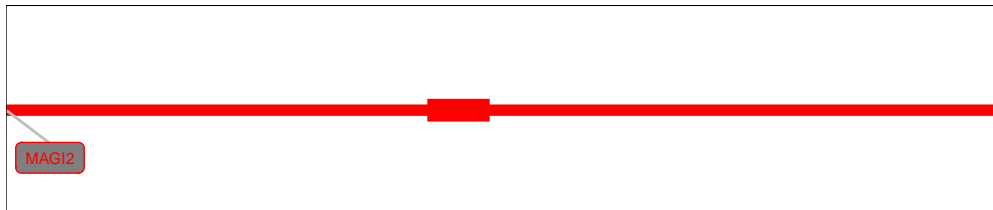
## chr3:127342511–127443012 Marker Peak Tracks for: L2.CUX2.MEIS2



L2.CUX2.MEIS2



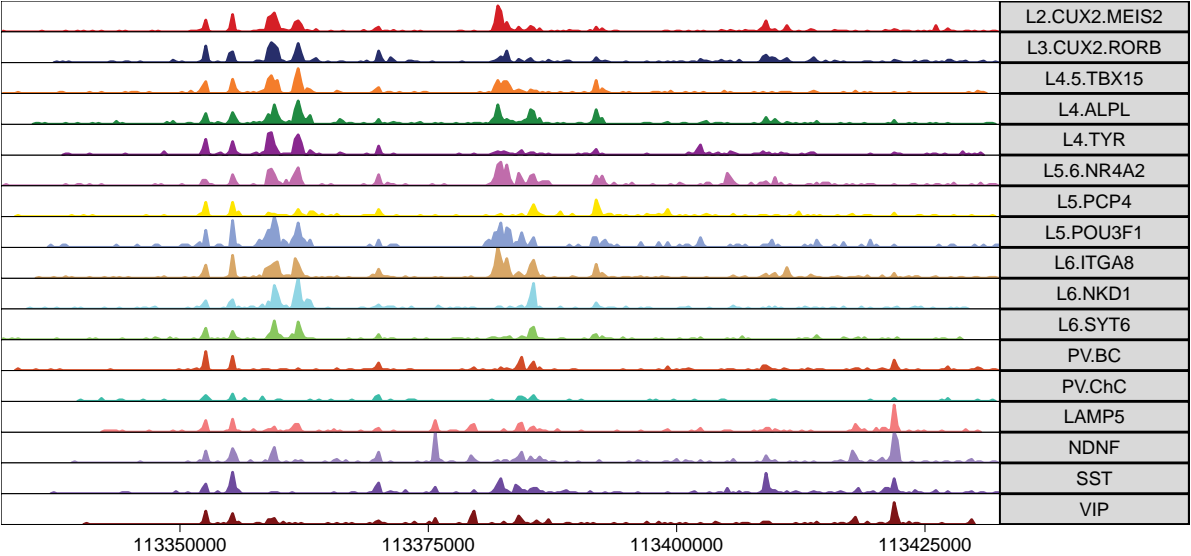
Peaks



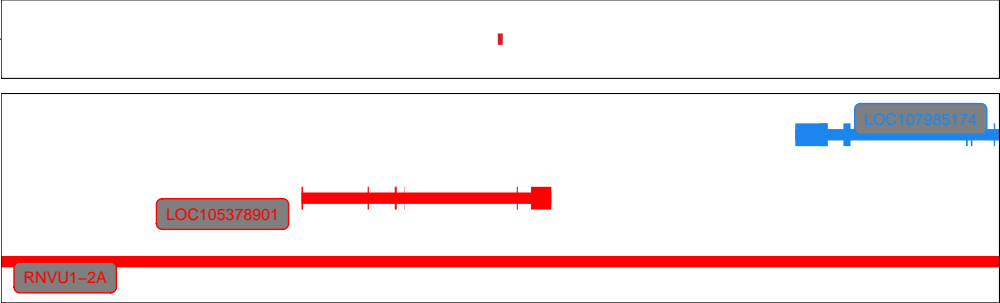
Genes

chr1:113331996–113432497 Marker Peak Tracks for: L2.CUX2.MEIS2

Coverage  
(Norm. ATAC Signal Range (0–0.92) by ReadsInTSS)



L2.CUX2.MEIS2

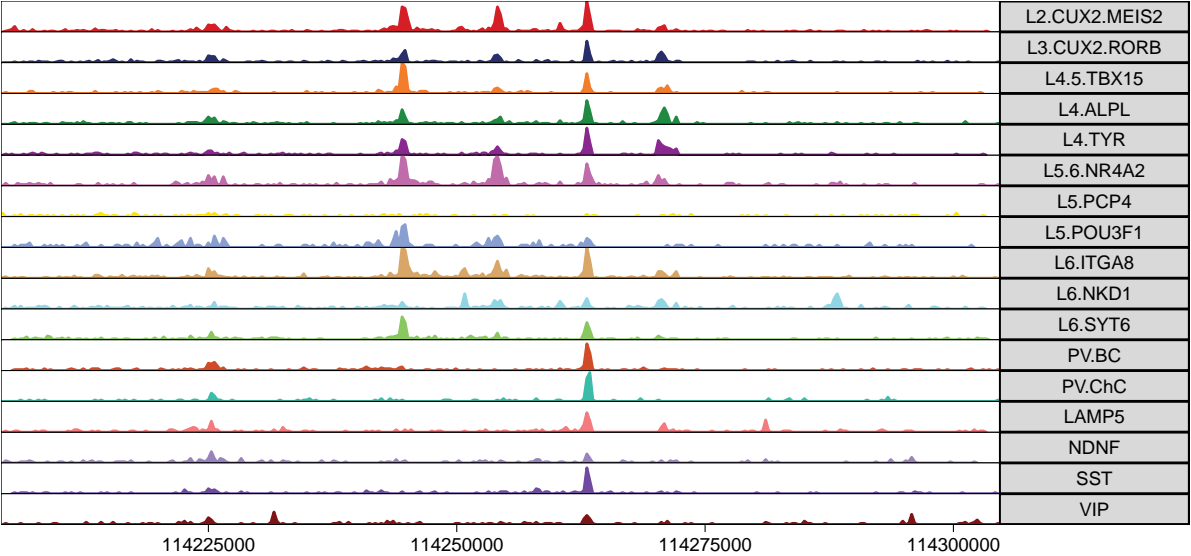


Peaks

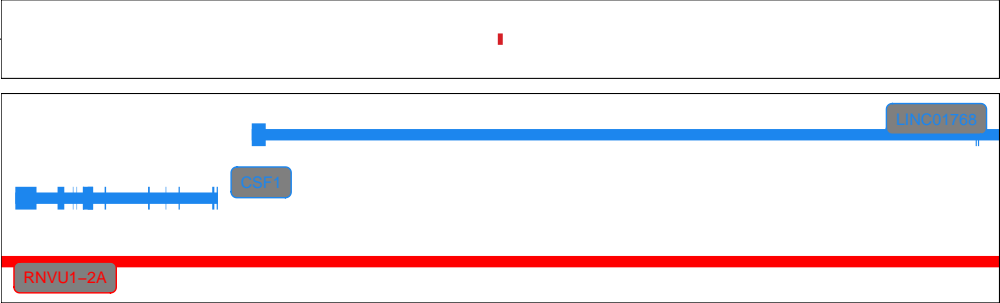
Genes

chr1:114204133–114304634 Marker Peak Tracks for: L2.CUX2.MEIS2

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

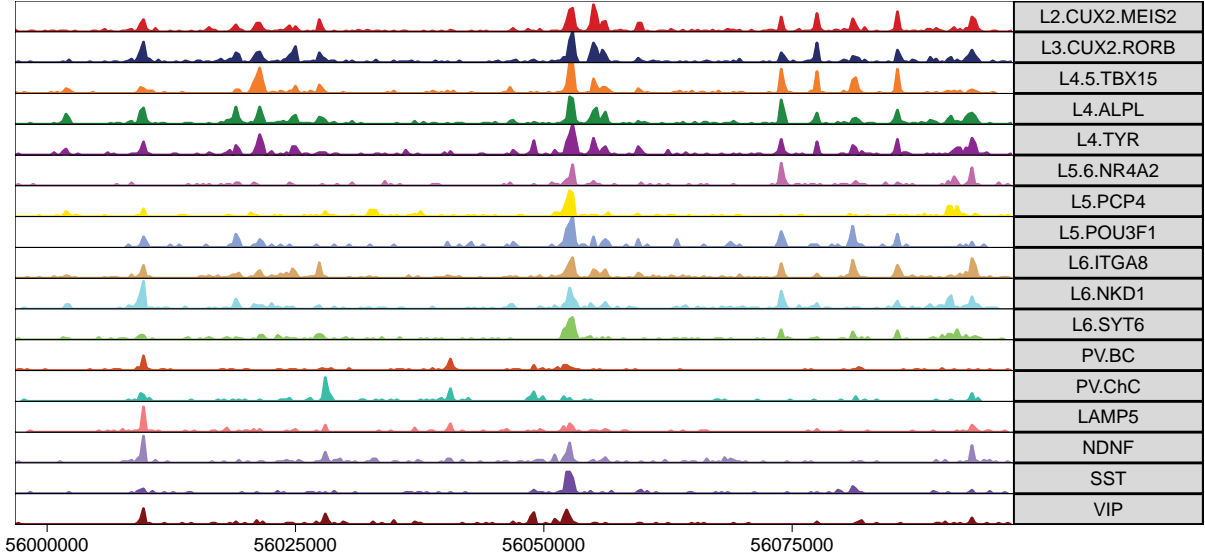


L2.CUX2.MEIS2

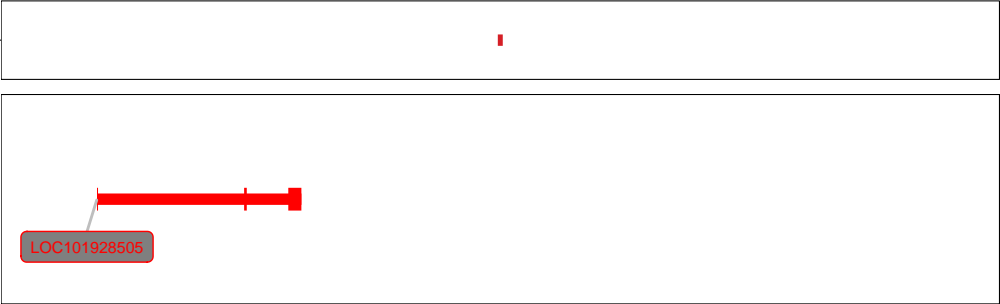


chr6:55996776–56097277 Marker Peak Tracks for: L2.CUX2.MEIS2

Coverage  
(Norm. ATAC Signal Range (0–0.85) by ReadsInTSS)



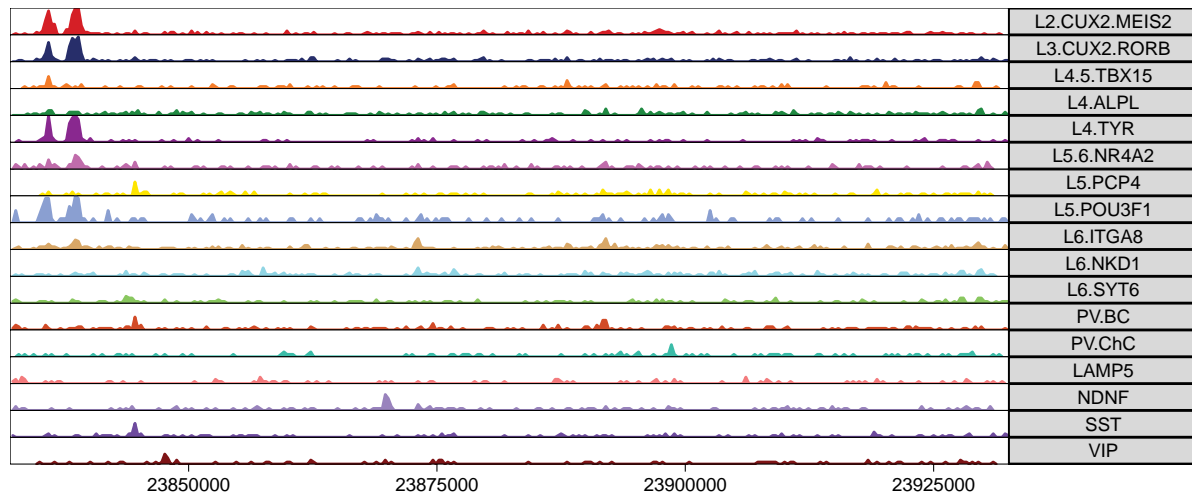
L2.CUX2.MEIS2



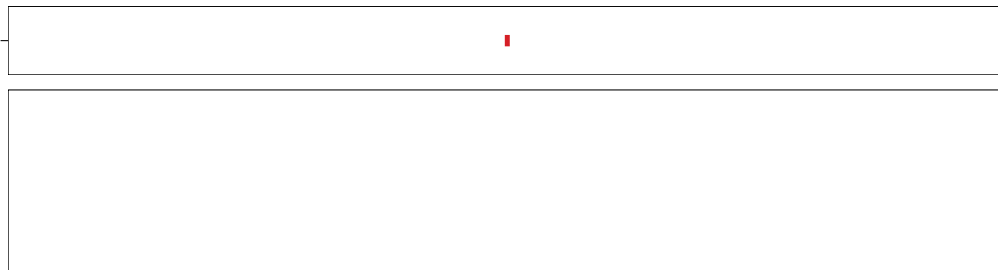
Peaks

Genes

## chr10:23832031–23932532 Marker Peak Tracks for: L2.CUX2.MEIS2

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

L2.CUX2.MEIS2



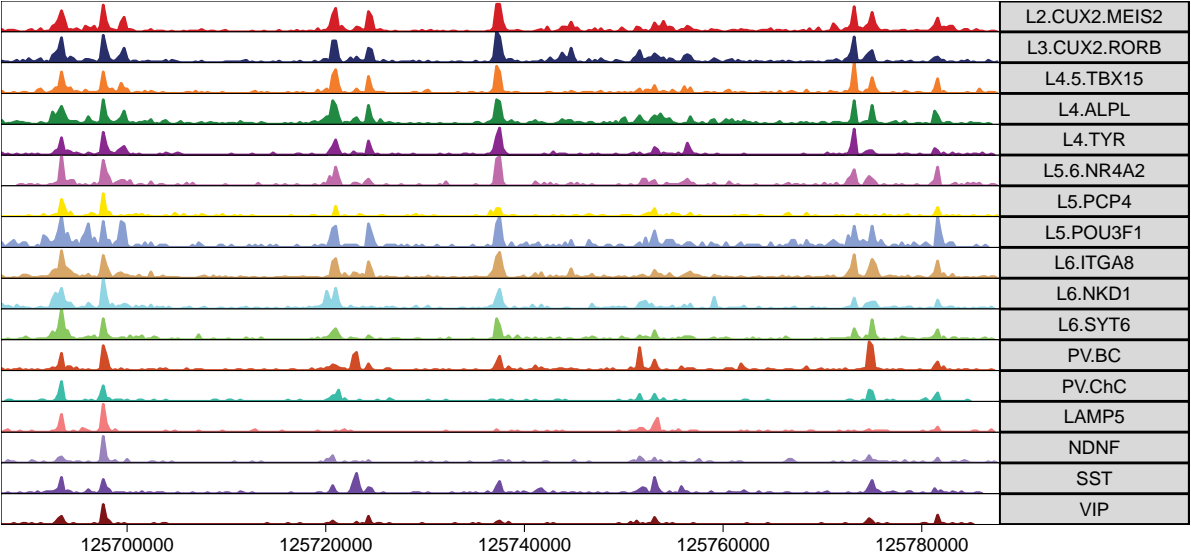
Peaks

GeneTrack

strand • •

chr14:125687320-125787821 Marker Peak Tracks for: L2.CUX2.MEIS2

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



L2.CUX2.MEIS2

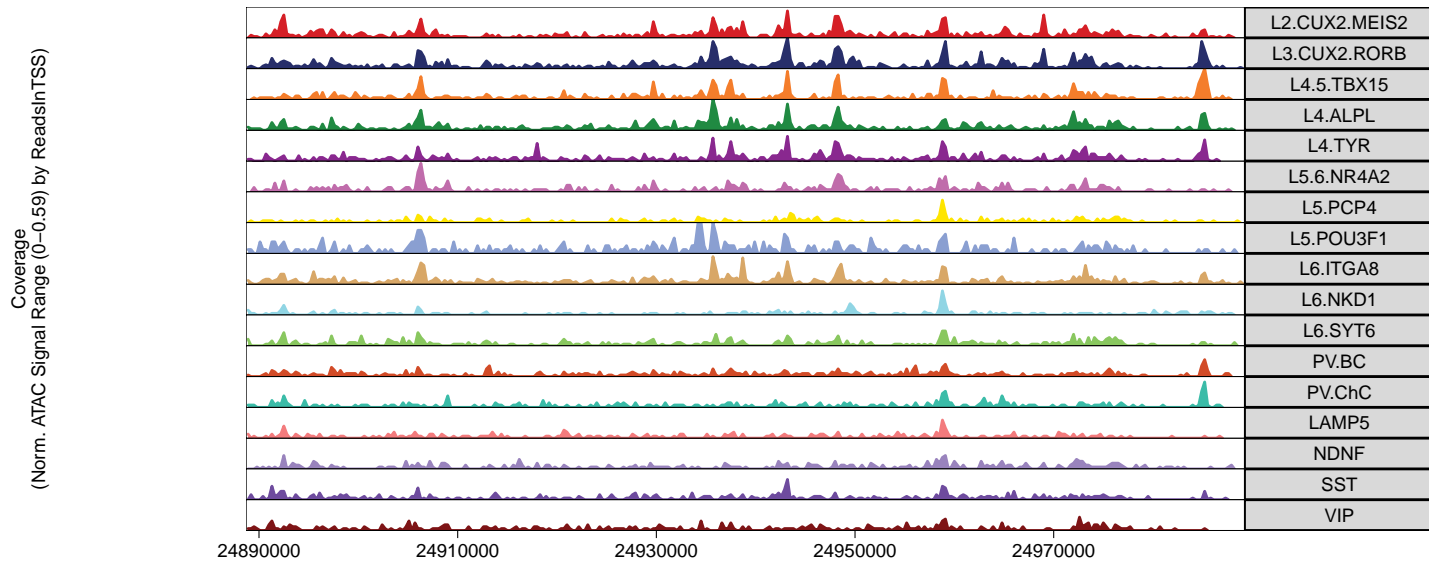


Peaks



Genes

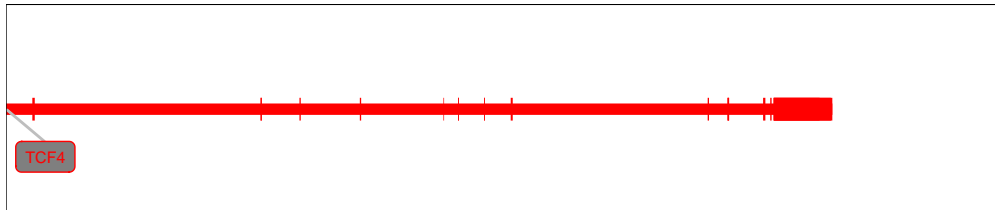
## chr18:24888703–24989204 Marker Peak Tracks for: L2.CUX2.MEIS2



L2.CUX2.MEIS2



Peaks

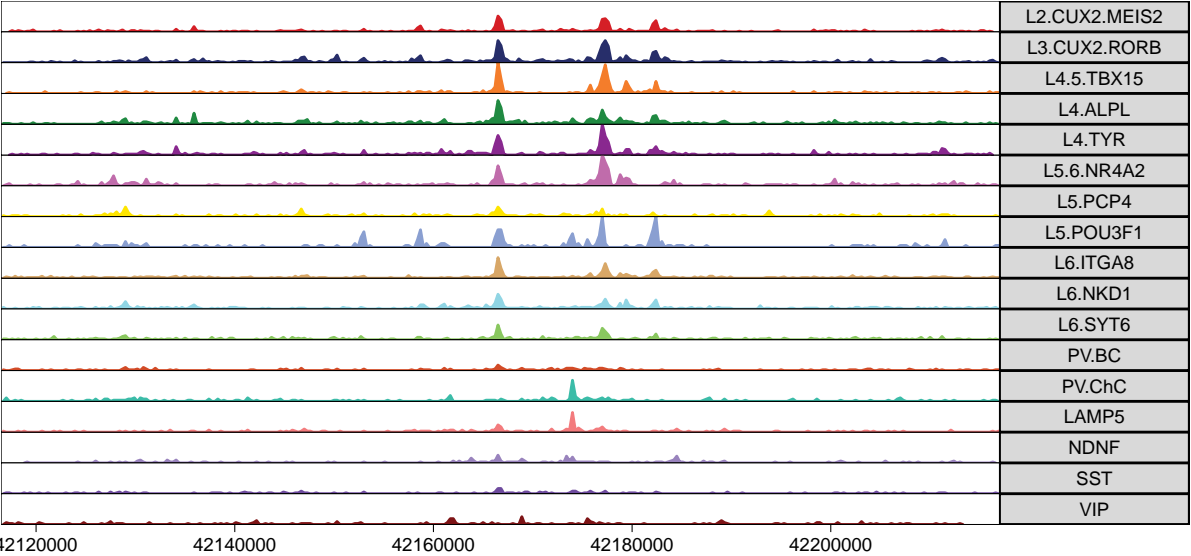


Genes

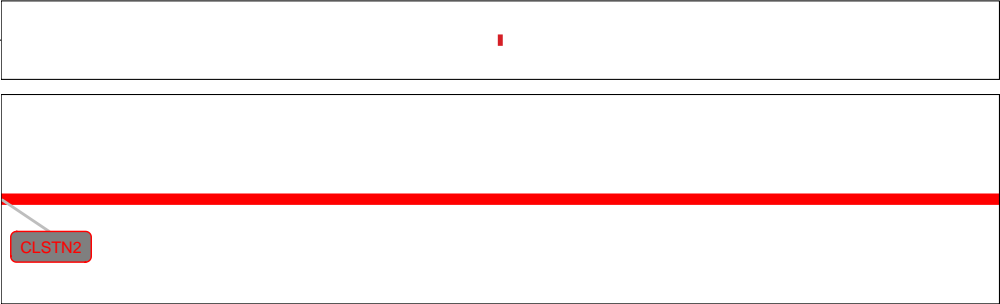


chr2:42116480–42216981 Marker Peak Tracks for: L2.CUX2.MEIS2

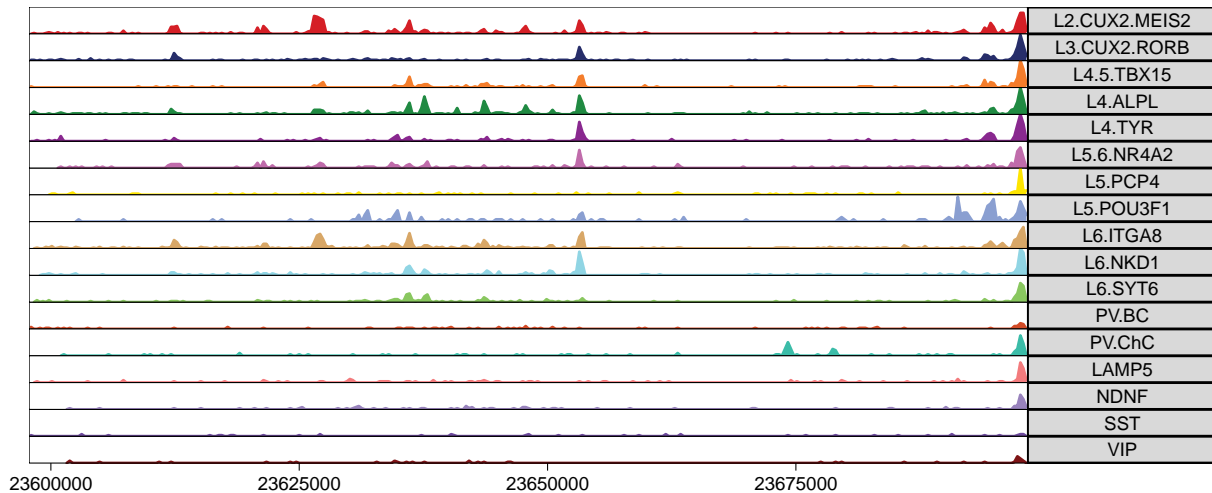
Coverage  
(Norm. ATAC Signal Range (0–1.15) by ReadsInTSS)



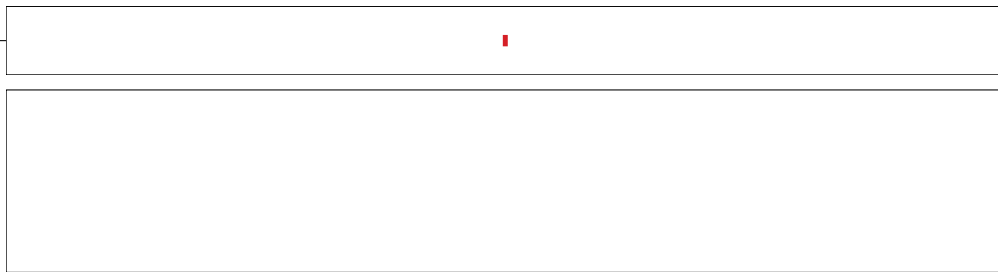
L2.CUX2.MEIS2



## chr15:23597809–23698310 Marker Peak Tracks for: L2.CUX2.MEIS2

Coverage  
(Norm. ATAC Signal Range (0–0.87) by ReadsInTSS)

L2.CUX2.MEIS2

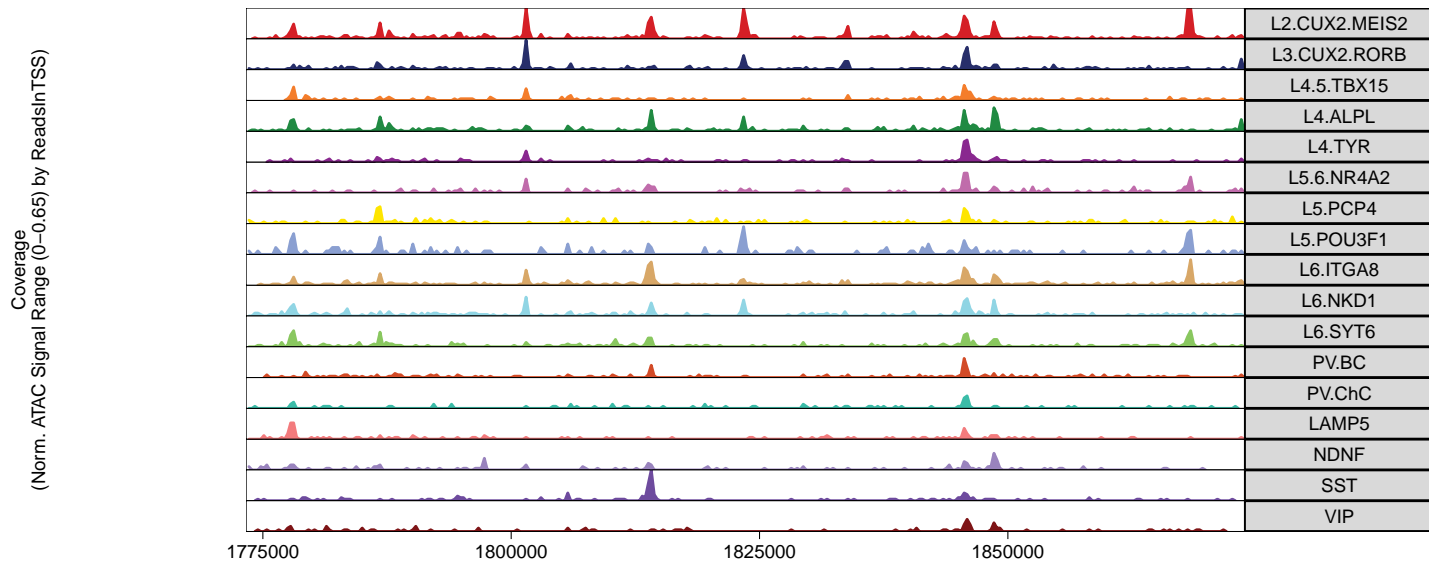


Peaks

GeneTrack

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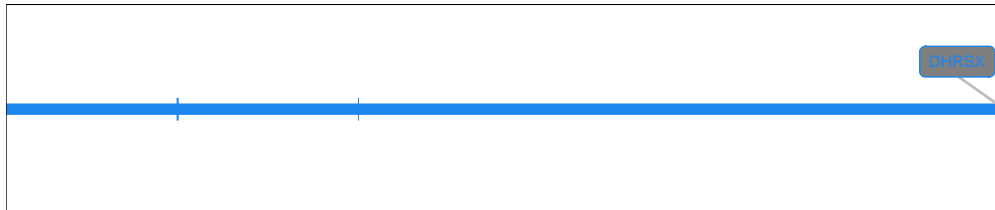
## chrX:1773316–1873817 Marker Peak Tracks for: L2.CUX2.MEIS2



L2.CUX2.MEIS2

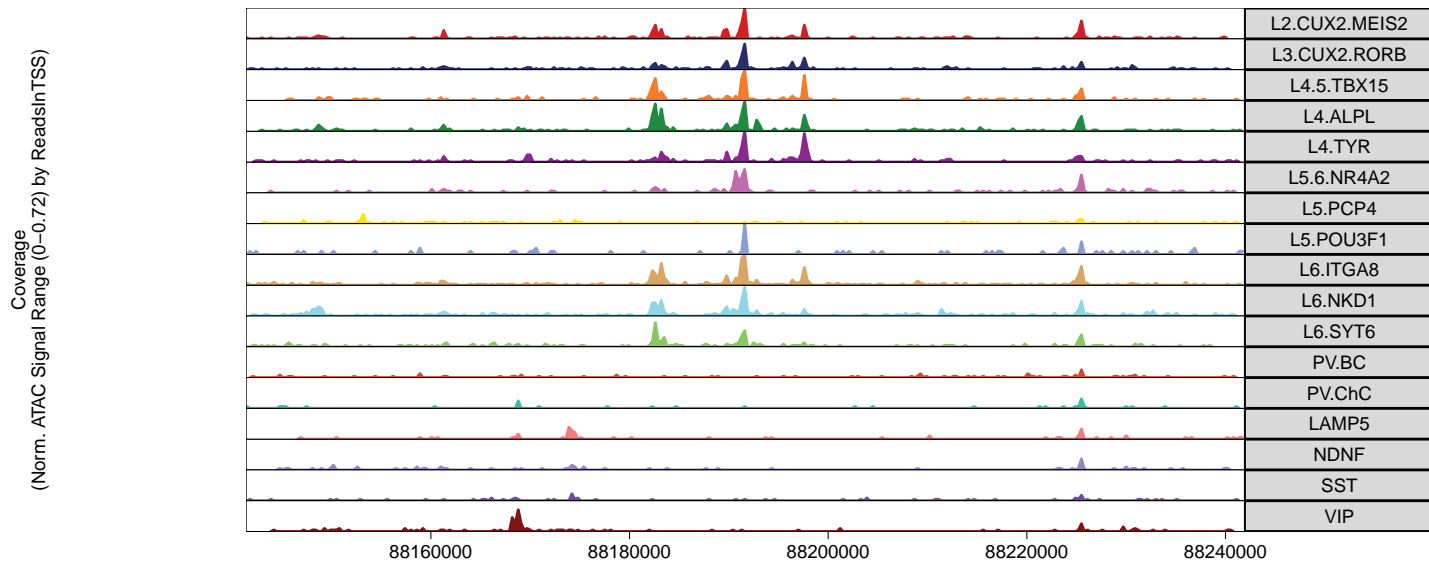


Peaks



Genes

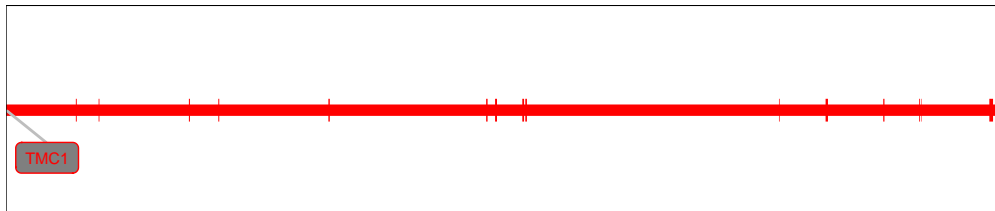
## chr15:88141406–88241907 Marker Peak Tracks for: L2.CUX2.MEIS2



L2.CUX2.MEIS2

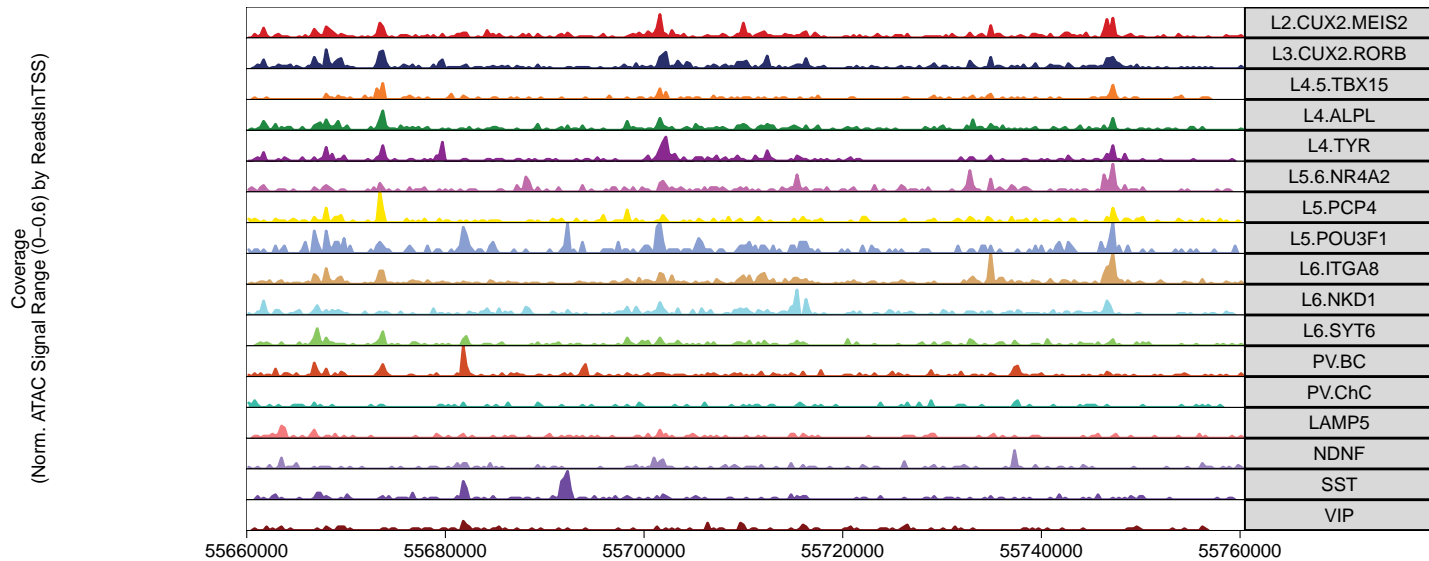


Peaks



Genes

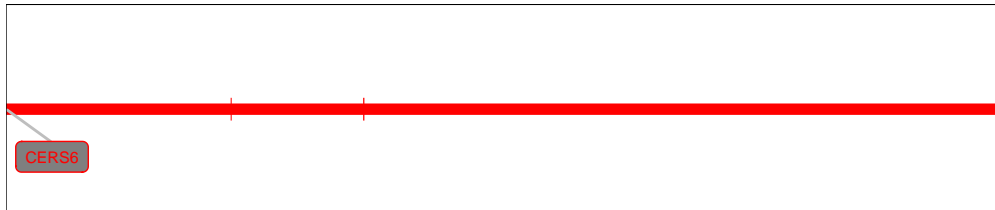
## chr12:55659940–55760441 Marker Peak Tracks for: L2.CUX2.MEIS2



L2.CUX2.MEIS2

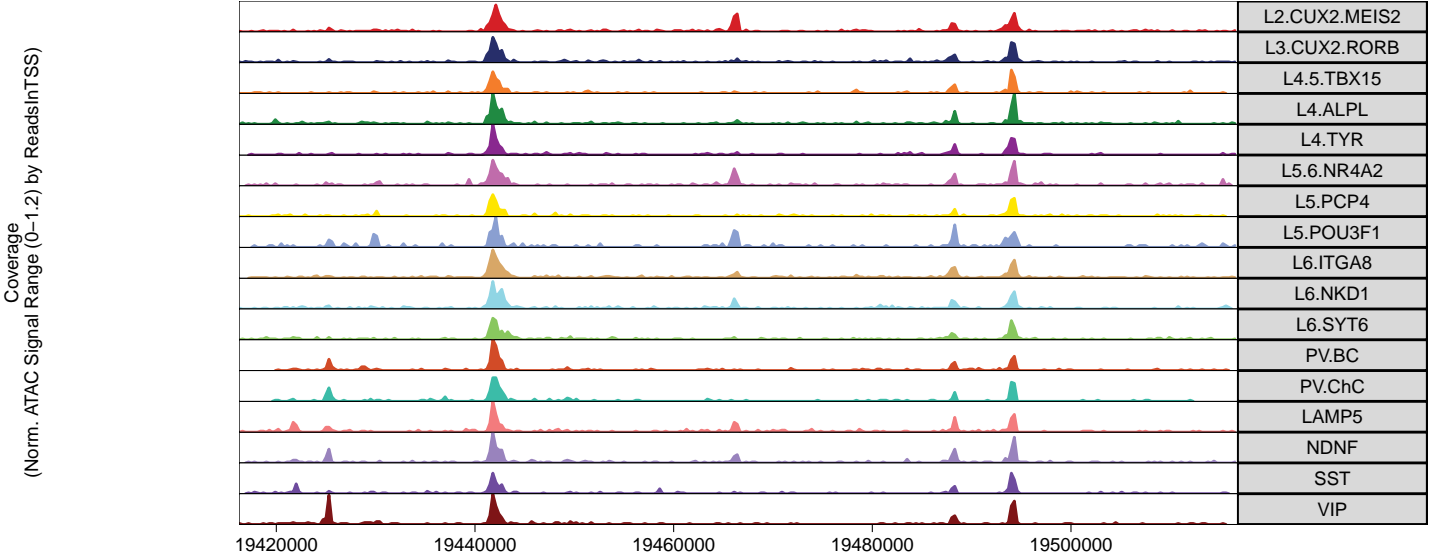


Peaks

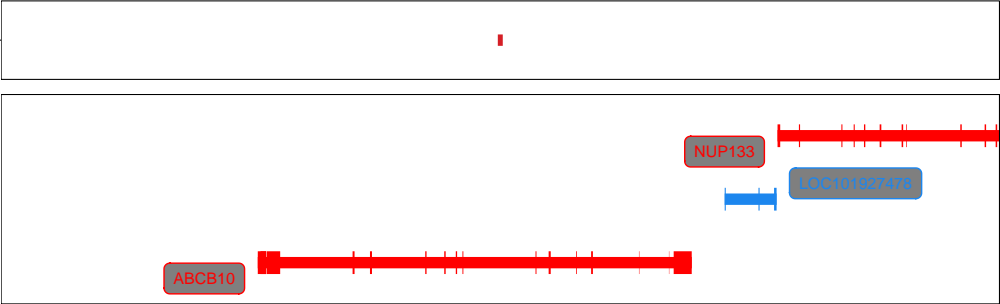


Genes

chr1:19416257–19516758 Marker Peak Tracks for: L2.CUX2.MEIS2



L2.CUX2.MEIS2

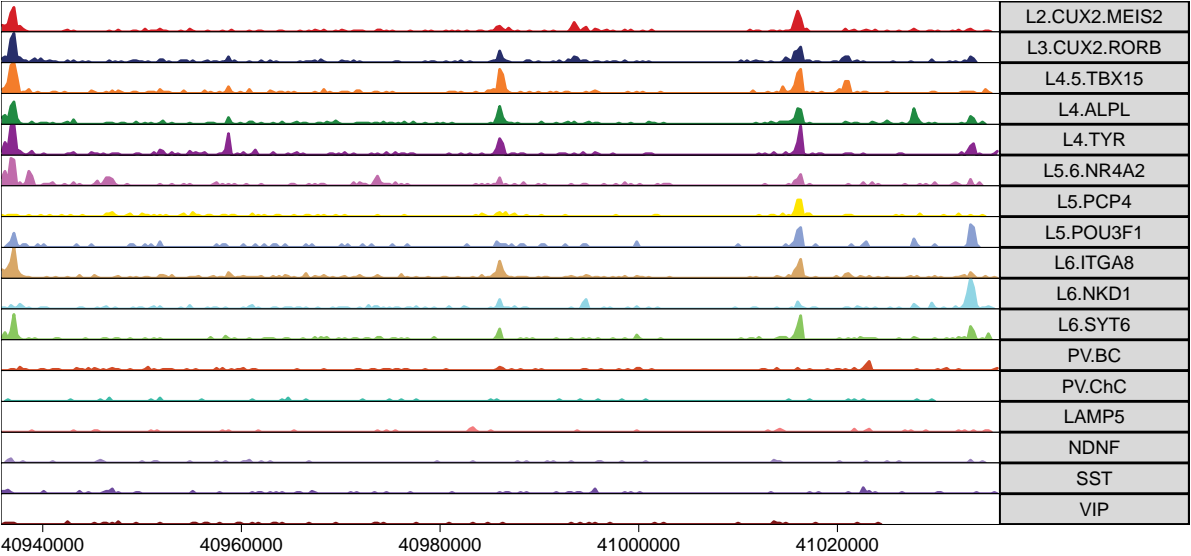


Peaks

Genes

chr11:40935810–41036311 Marker Peak Tracks for: L2.CUX2.MEIS2

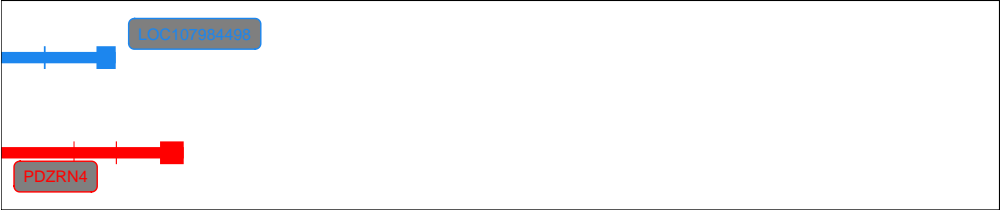
Coverage  
(Norm. ATAC Signal Range (0–0.8) by ReadsInTSS)



L2.CUX2.MEIS2



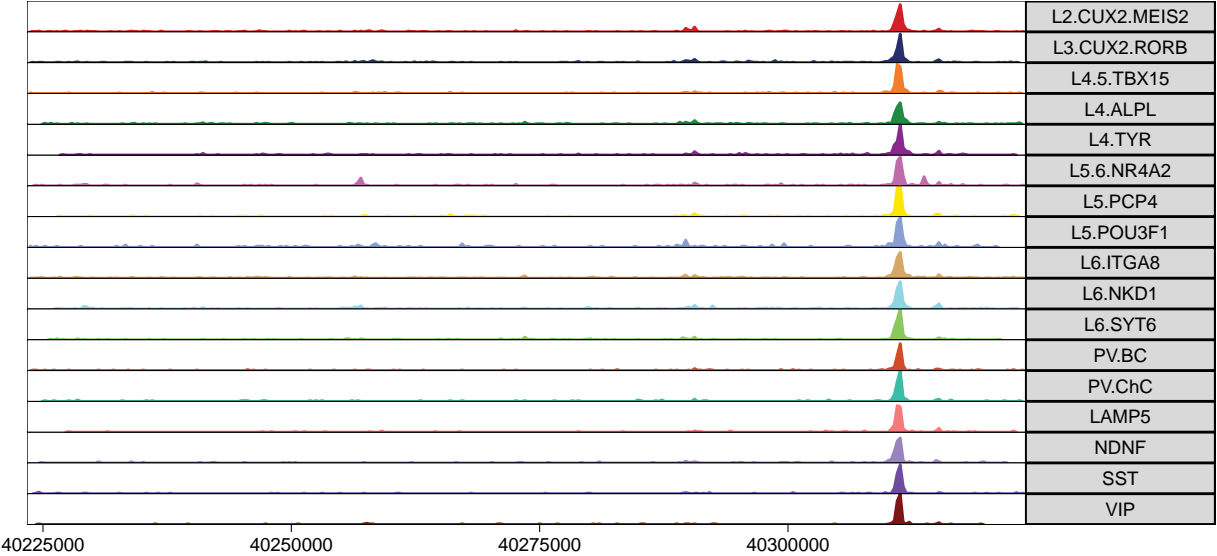
Peaks



Genes

chr13:40223404–40323905 Marker Peak Tracks for: L2.CUX2.MEIS2

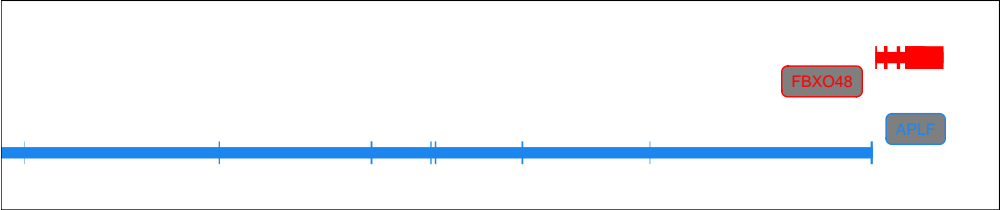
Coverage  
(Norm. ATAC Signal Range (0–1.83) by ReadsInTSS)



L2.CUX2.MEIS2



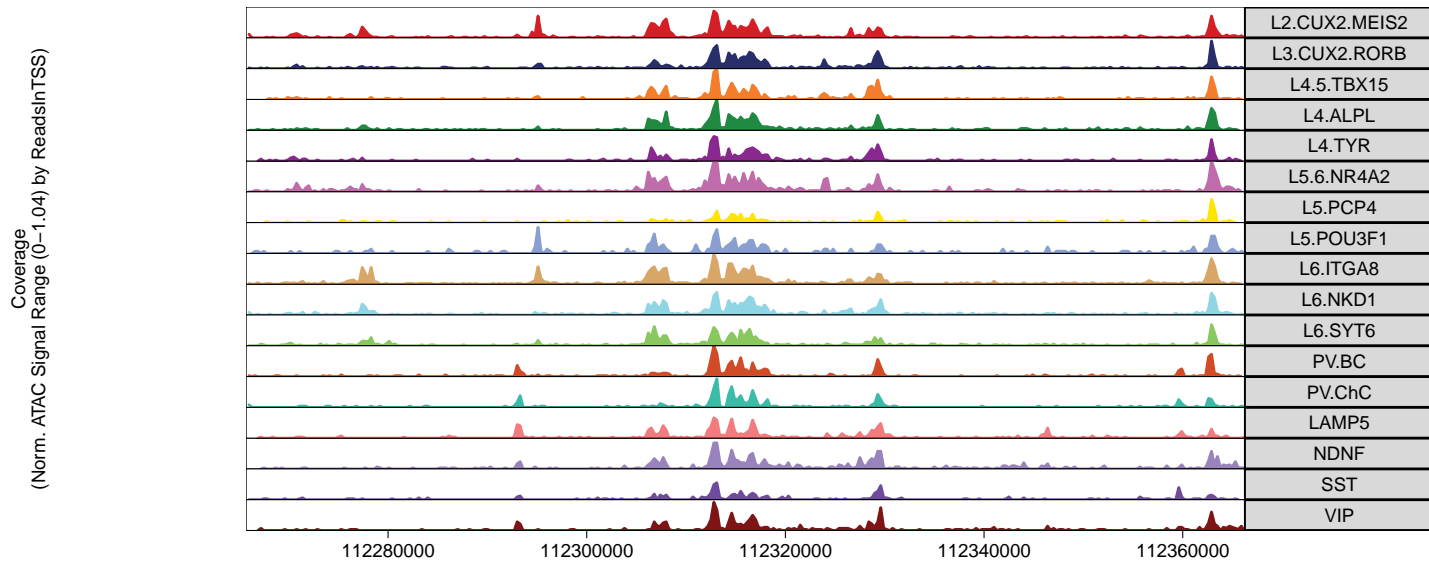
Peaks



Genes



## chr15:112265712–112366213 Marker Peak Tracks for: L2.CUX2.MEIS2



L2.CUX2.MEIS2



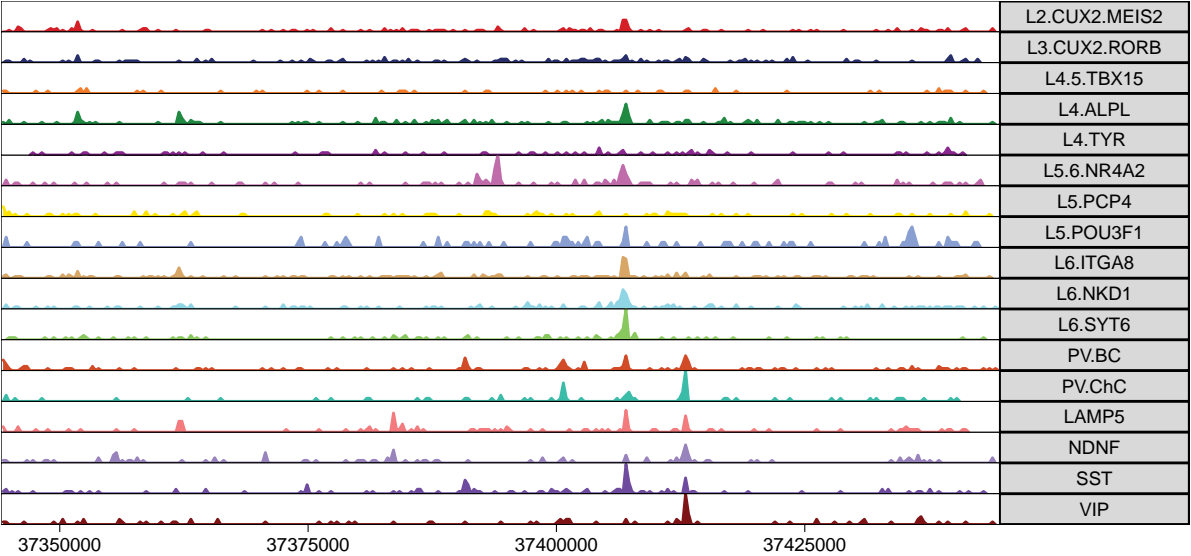
Peaks



Genes

chr12:37344097–37444598 Marker Peak Tracks for: L2.CUX2.MEIS2

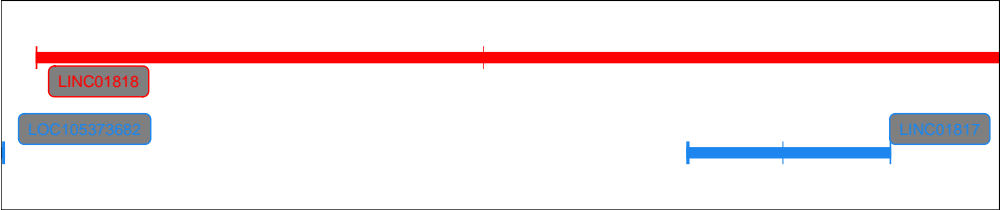
Coverage  
(Norm. ATAC Signal Range (0–0.44) by ReadsInTSS)



L2.CUX2.MEIS2

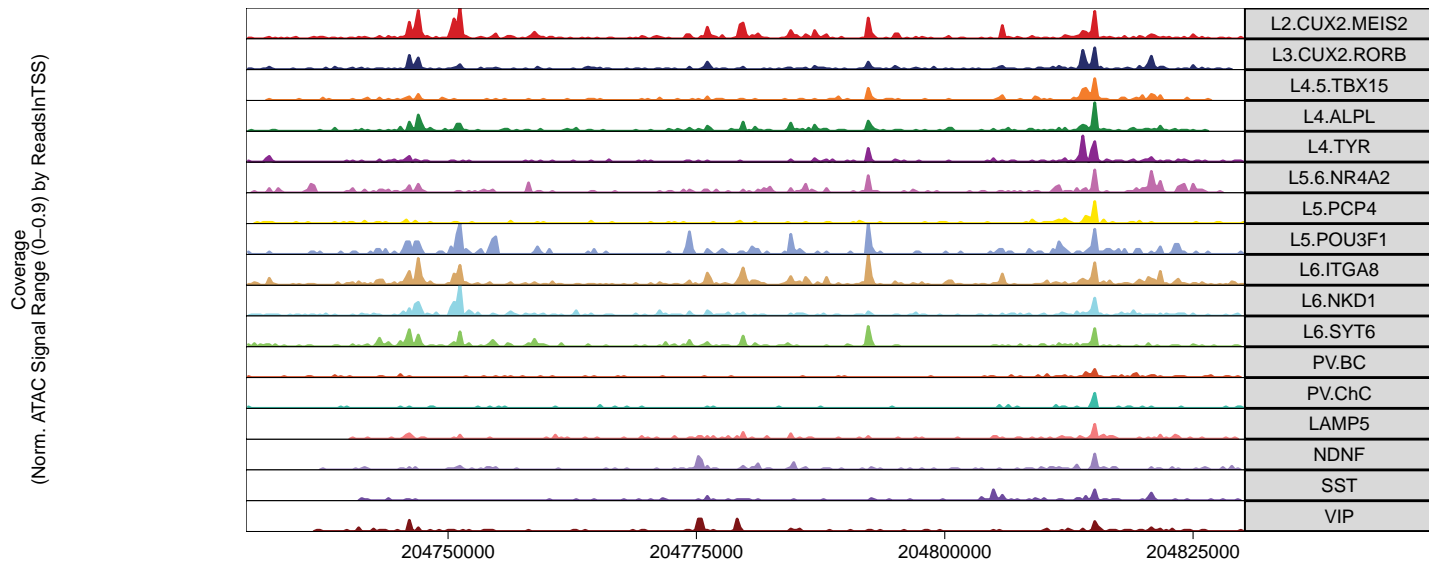


Peaks



Genes

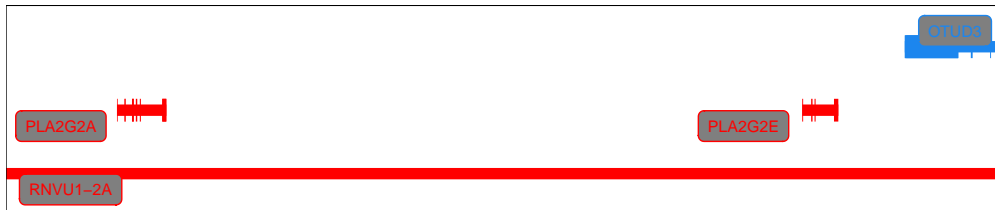
## chr1:204729670–204830171 Marker Peak Tracks for: L2.CUX2.MEIS2



L2.CUX2.MEIS2



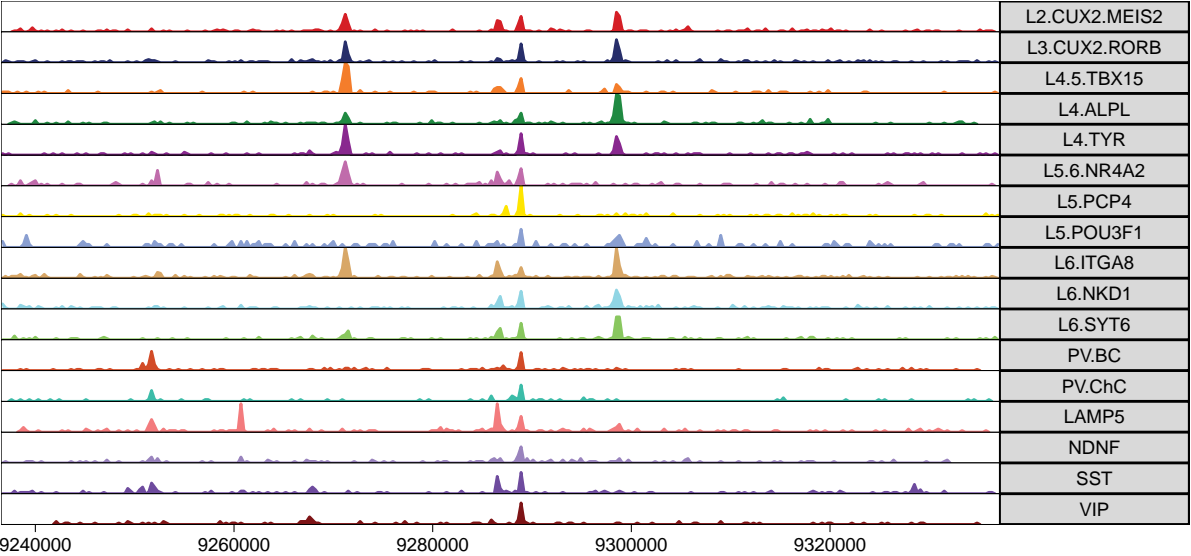
Peaks



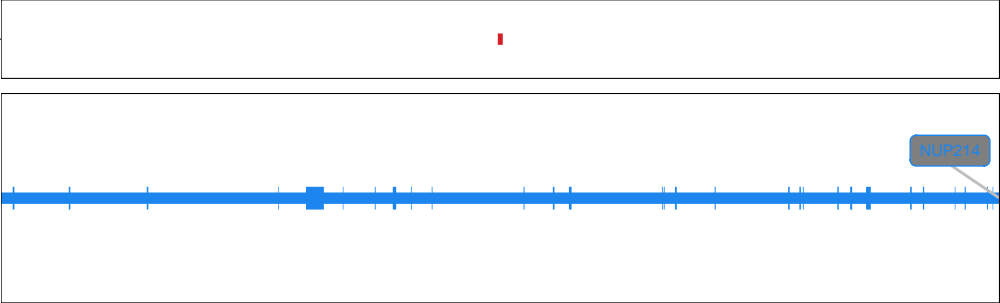
Genes

chr15:9236555-9337056 Marker Peak Tracks for: L2.CUX2.MEIS2

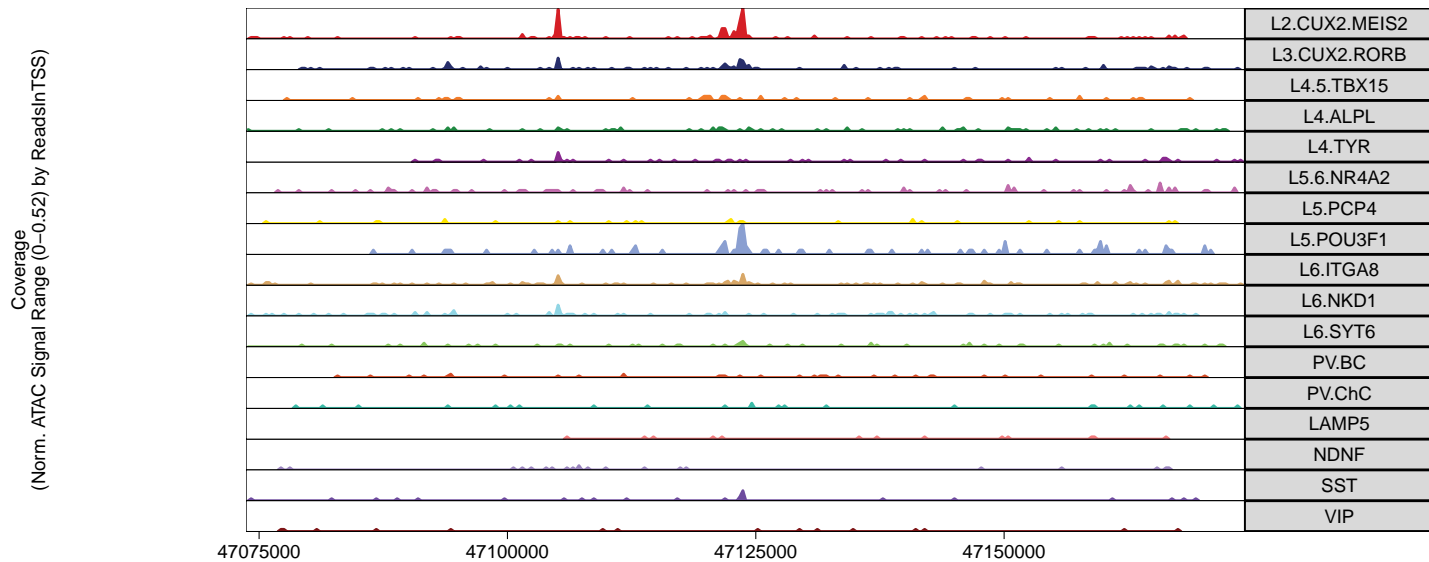
Coverage  
(Norm. ATAC Signal Range (0-0.75) by ReadsInTSS)



L2.CUX2.MEIS2



## chr1:47073692–47174193 Marker Peak Tracks for: L2.CUX2.MEIS2



L2.CUX2.MEIS2



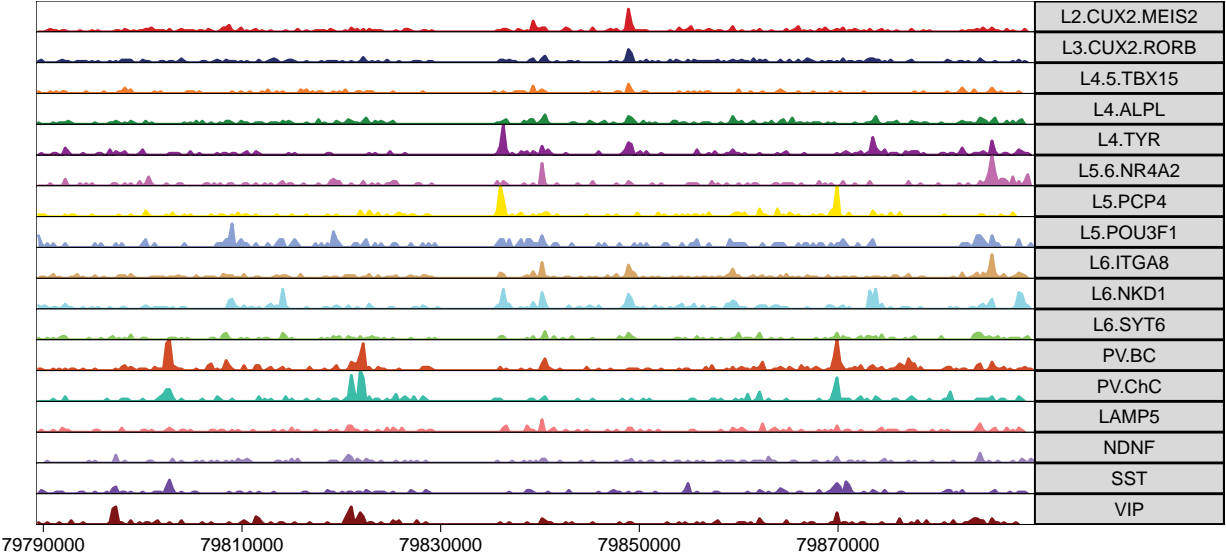
Peaks



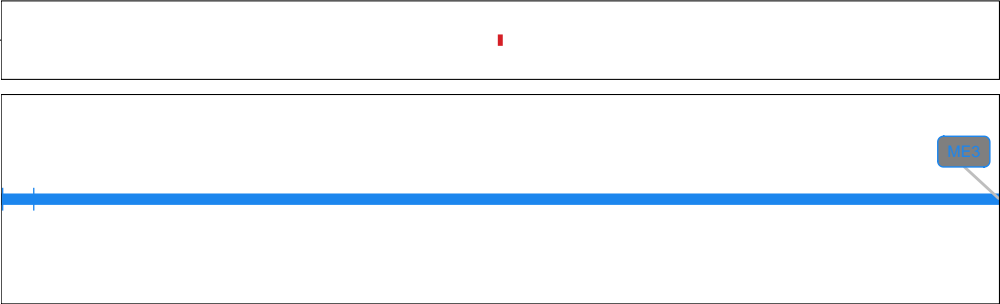
Genes

chr14:79789273–79889774 Marker Peak Tracks for: L2.CUX2.MEIS2

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



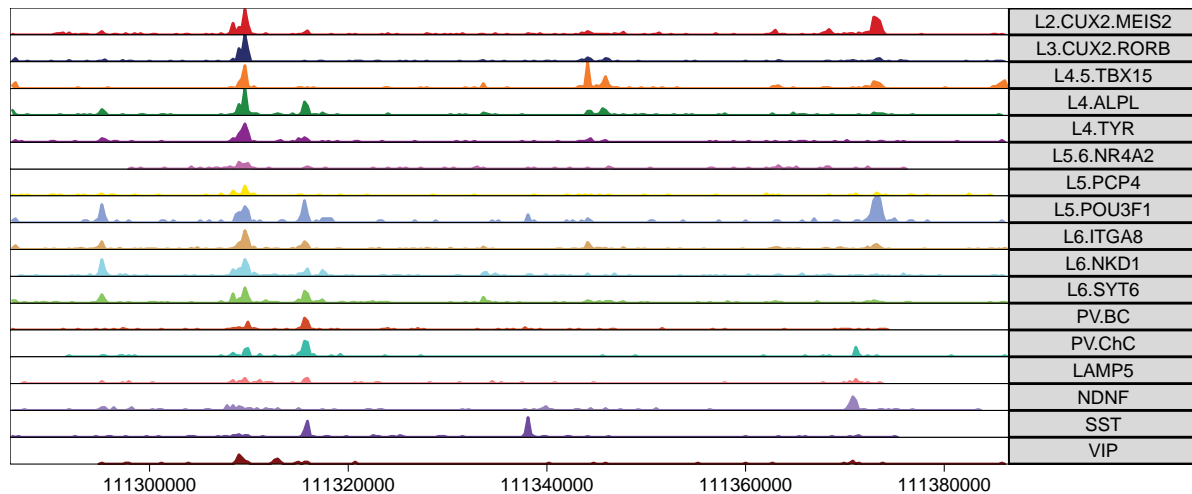
L2.CUX2.MEIS2



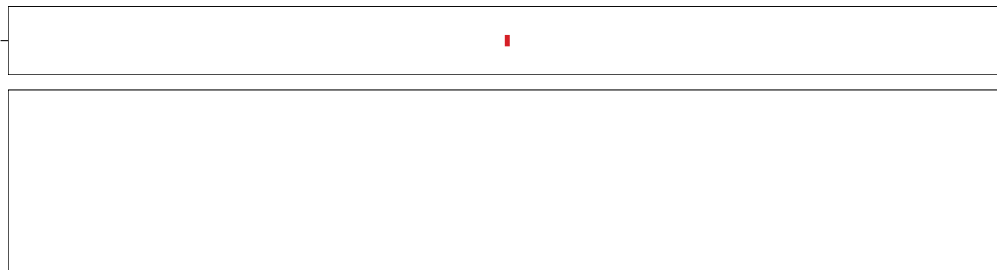
Peaks

Genes

## chr6:111285962–111386463 Marker Peak Tracks for: L2.CUX2.MEIS2

Coverage  
(Norm. ATAC Signal Range (0–0.98) by ReadsInTSS)

L2.CUX2.MEIS2

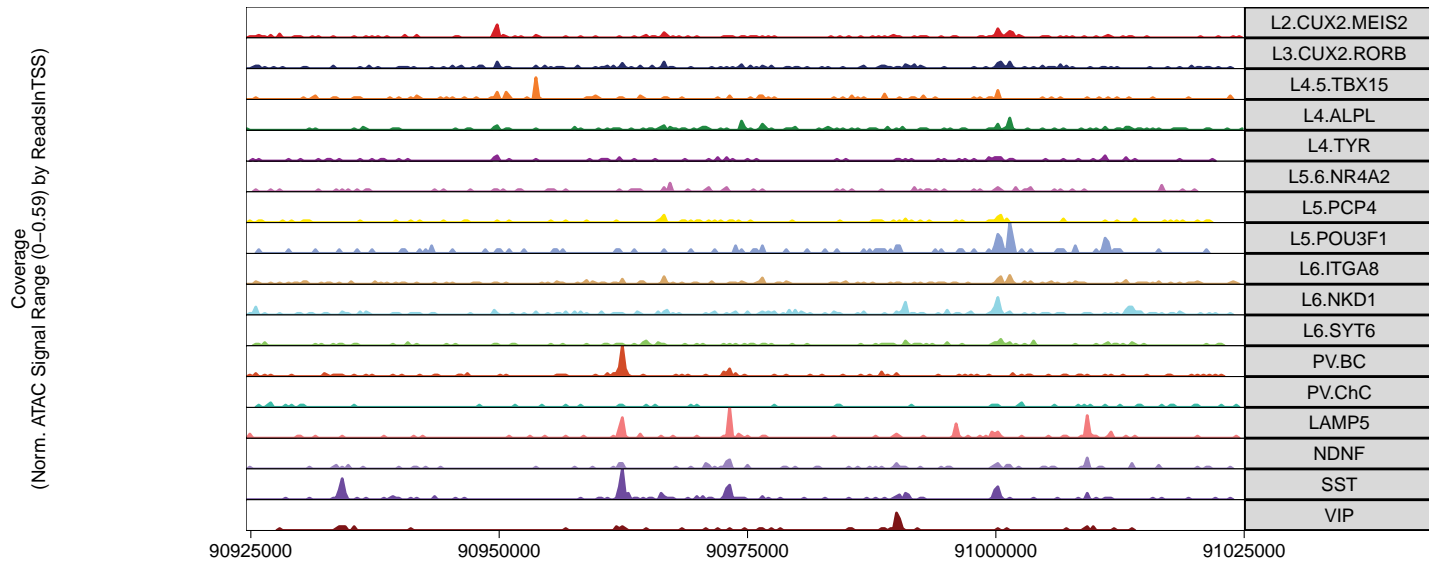


Peaks

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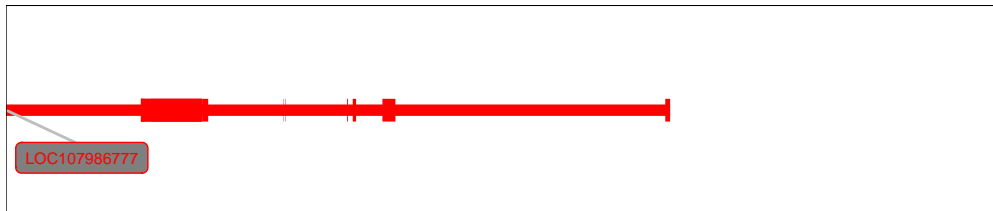
## chr3:90924525–91025026 Marker Peak Tracks for: L2.CUX2.MEIS2



L2.CUX2.MEIS2



Peaks

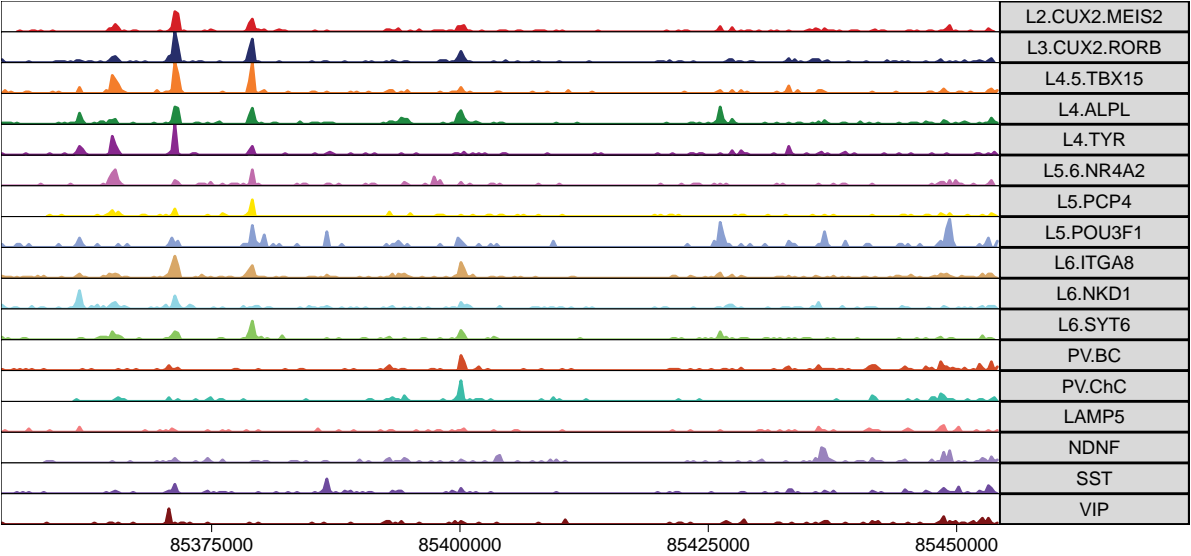


Genes

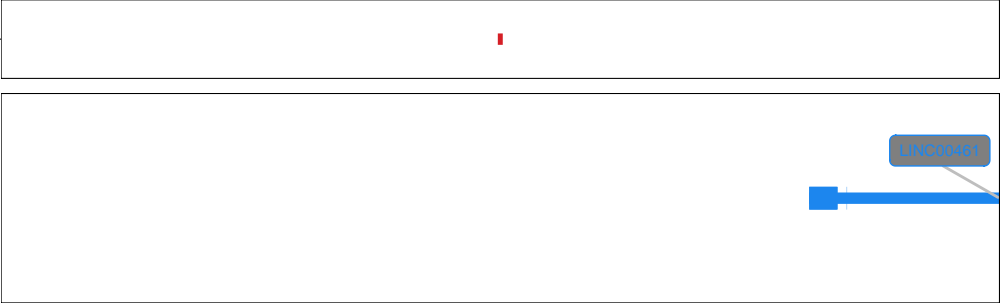


chr6:85353809–85454310 Marker Peak Tracks for: L2.CUX2.MEIS2

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



L2.CUX2.MEIS2

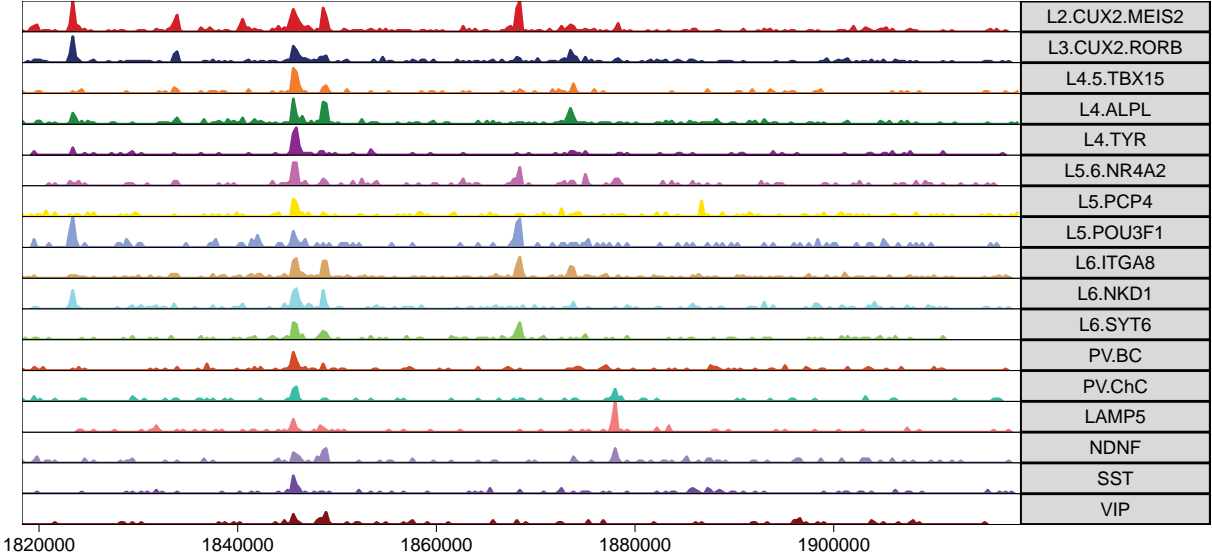


Peaks

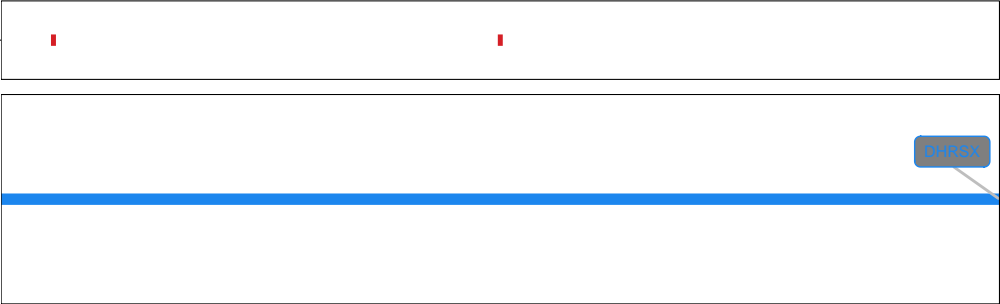
Genes

chrX:1818298–1918799 Marker Peak Tracks for: L2.CUX2.MEIS2

Coverage  
(Norm. ATAC Signal Range (0–0.56) by ReadsInTSS)



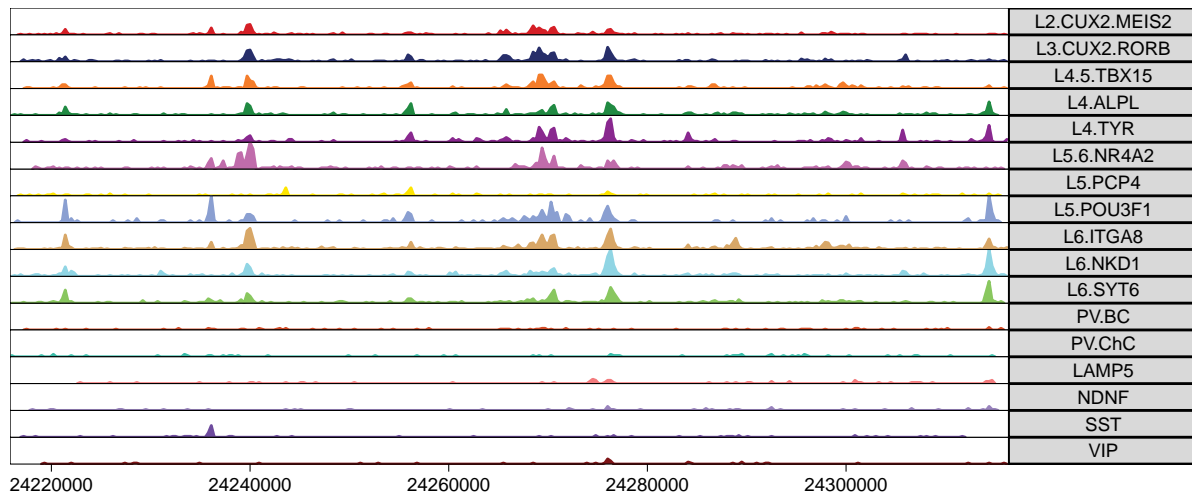
L2.CUX2.MEIS2



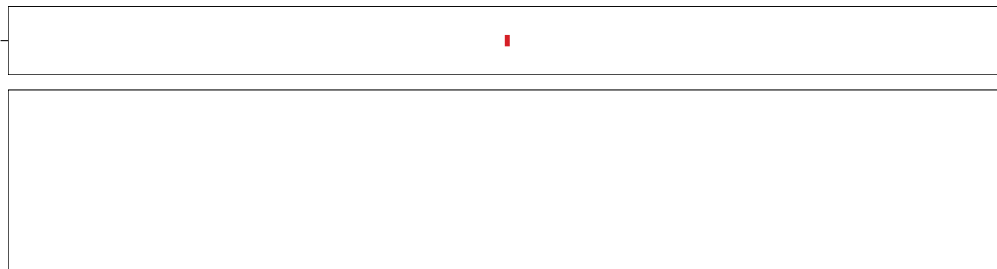
Peaks

Genes

## chr11:24215846–24316347 Marker Peak Tracks for: L2.CUX2.MEIS2

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

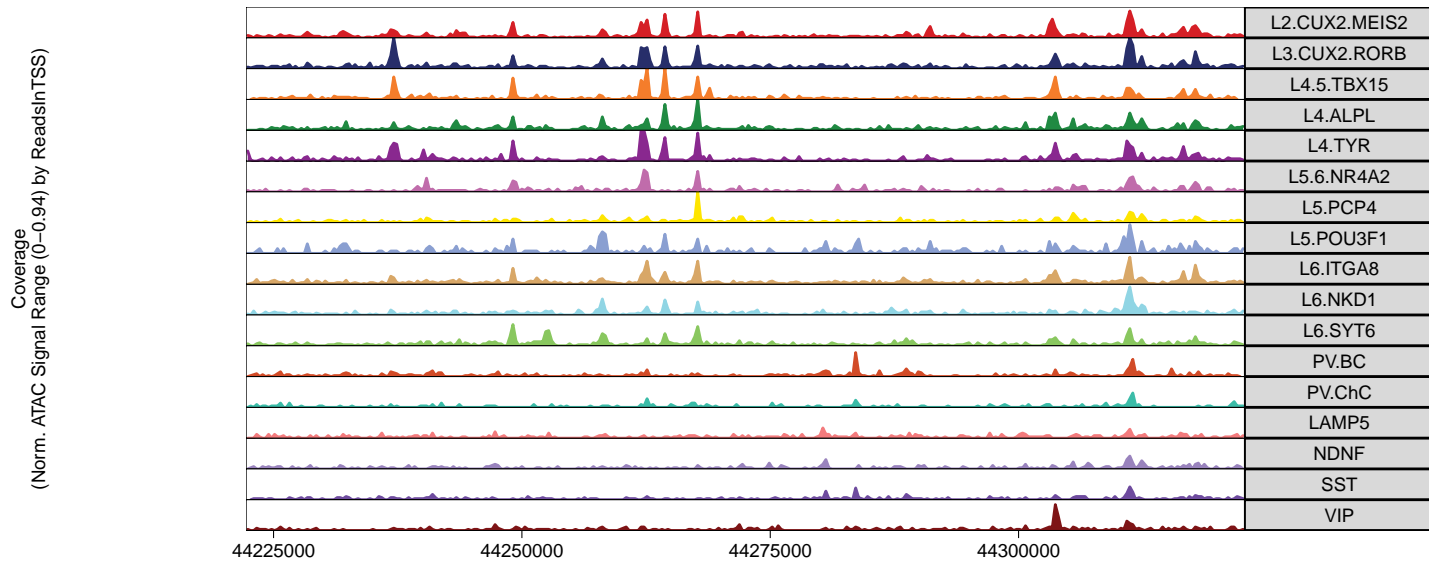
L2.CUX2.MEIS2



Peaks

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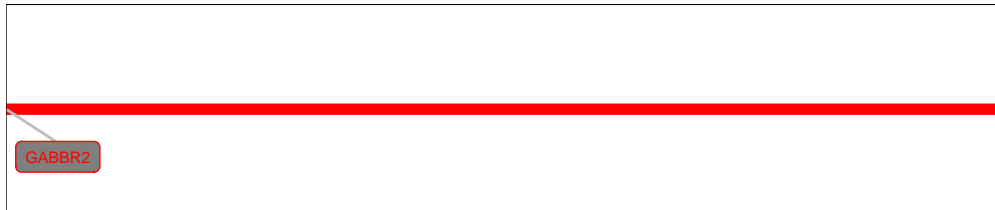
## chr15:44222237–44322738 Marker Peak Tracks for: L2.CUX2.MEIS2



L2.CUX2.MEIS2



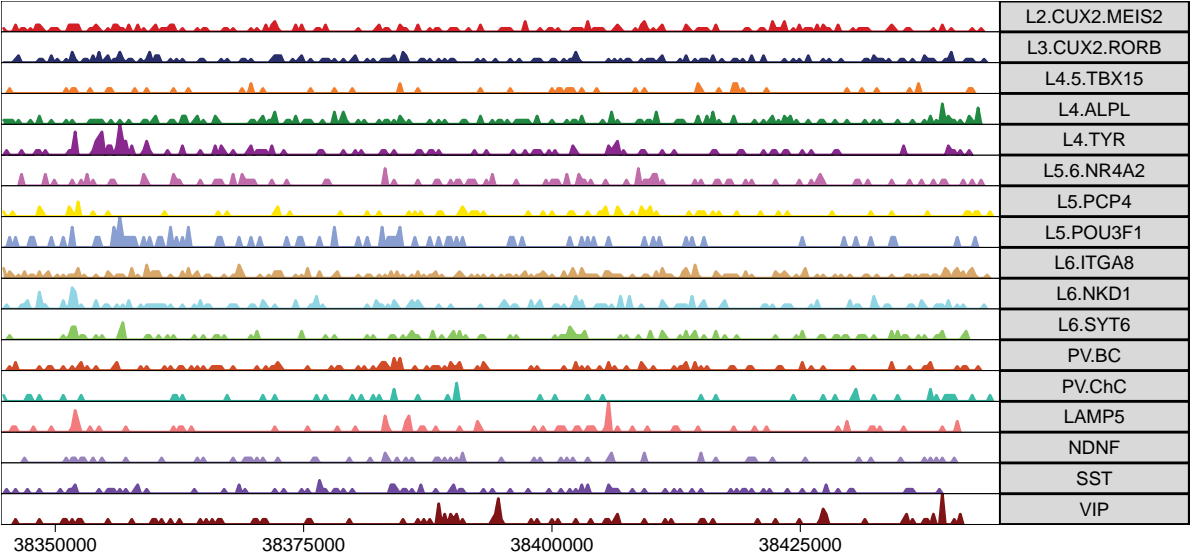
Peaks



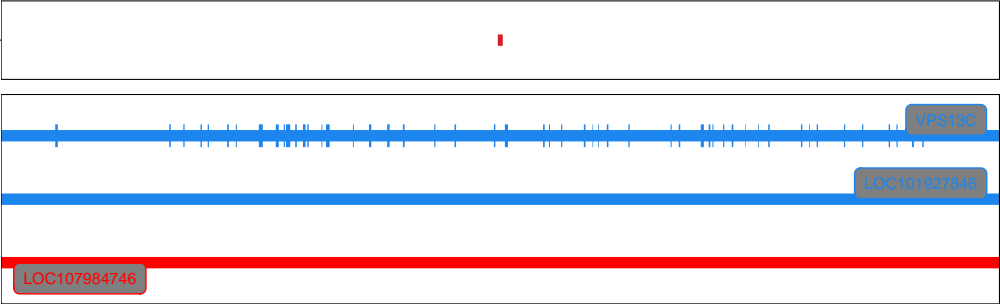
Genes

chr7:38344547-38445048 Marker Peak Tracks for: L2.CUX2.MEIS2

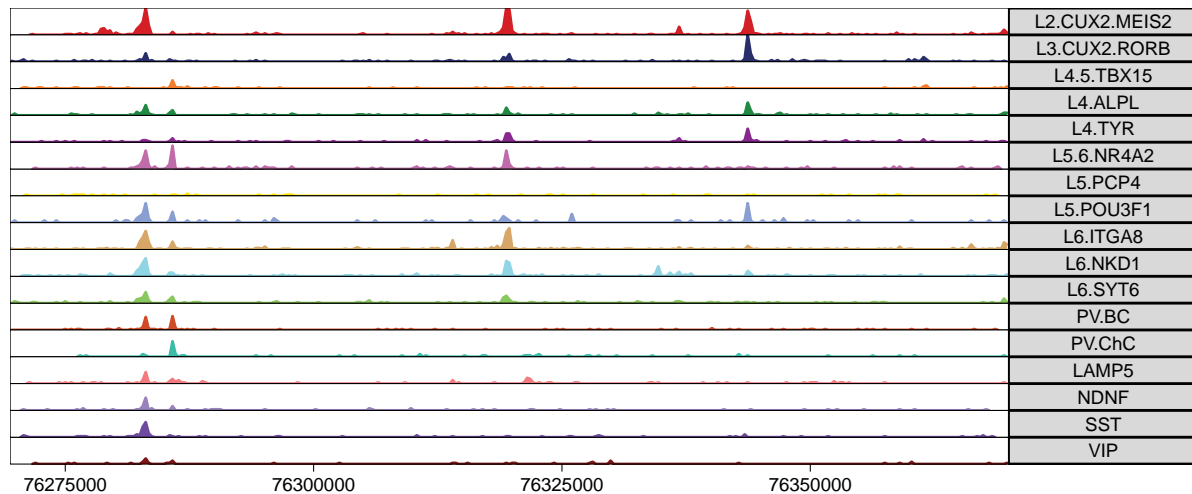
Coverage  
(Norm. ATAC Signal Range (0-0.22) by ReadsInTSS)



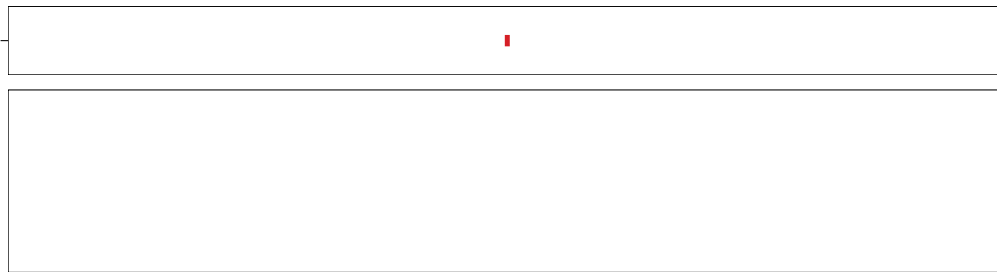
L2.CUX2.MEIS2



## chr1:76269448–76369949 Marker Peak Tracks for: L2.CUX2.MEIS2

Coverage  
(Norm. ATAC Signal Range (0–0.92) by ReadsInTSS)

L2.CUX2.MEIS2

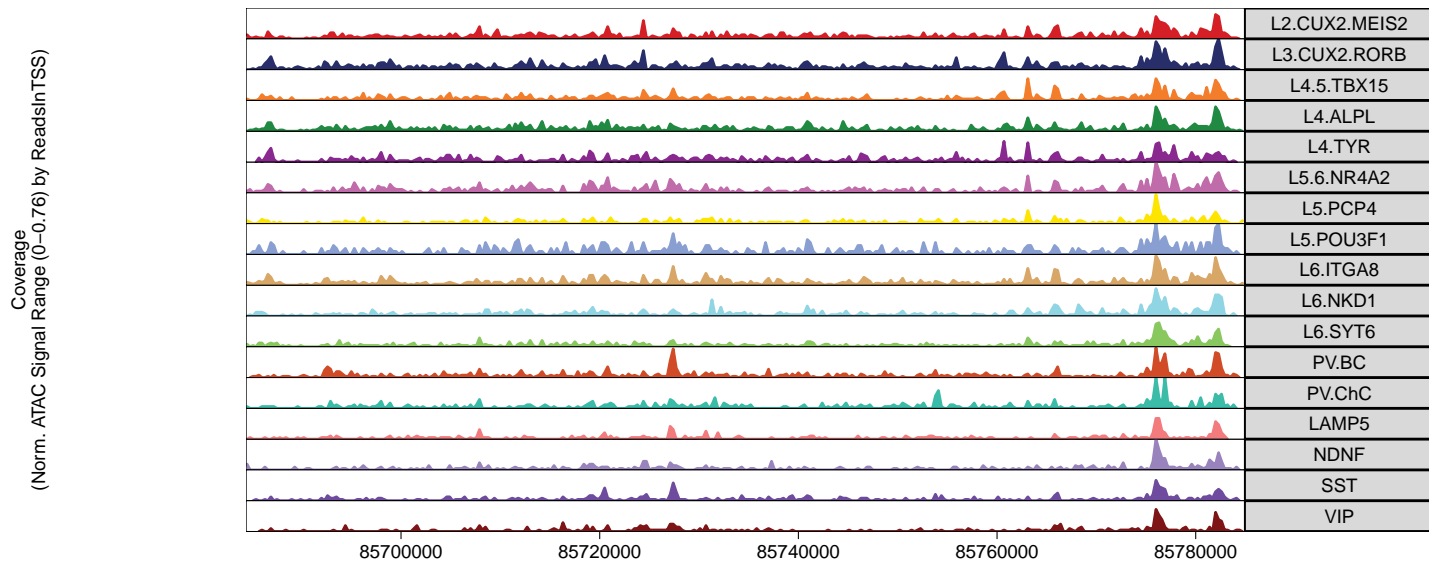


Peaks

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## chr6:85684405–85784906 Marker Peak Tracks for: L2.CUX2.MEIS2



L2.CUX2.MEIS2



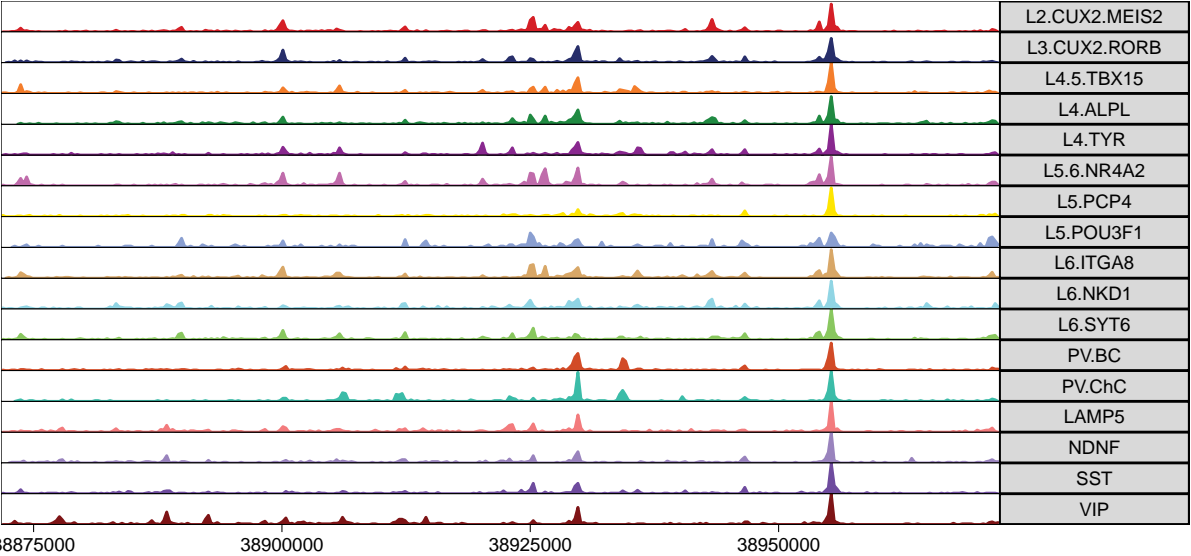
Peaks



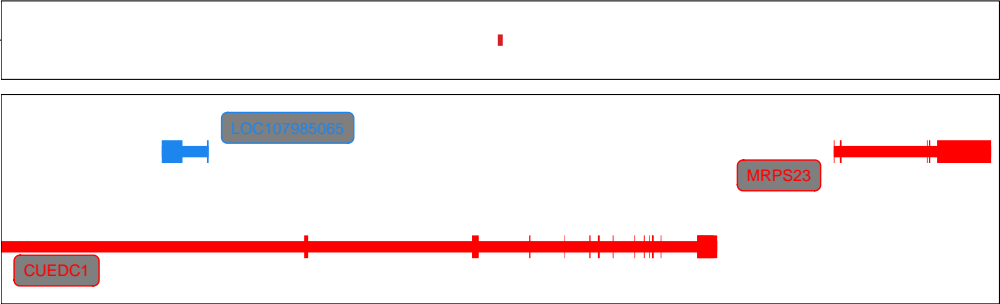
Genes

chr16:38871731–38972232 Marker Peak Tracks for: L2.CUX2.MEIS2

Coverage  
(Norm. ATAC Signal Range (0–1.72) by ReadsInTSS)

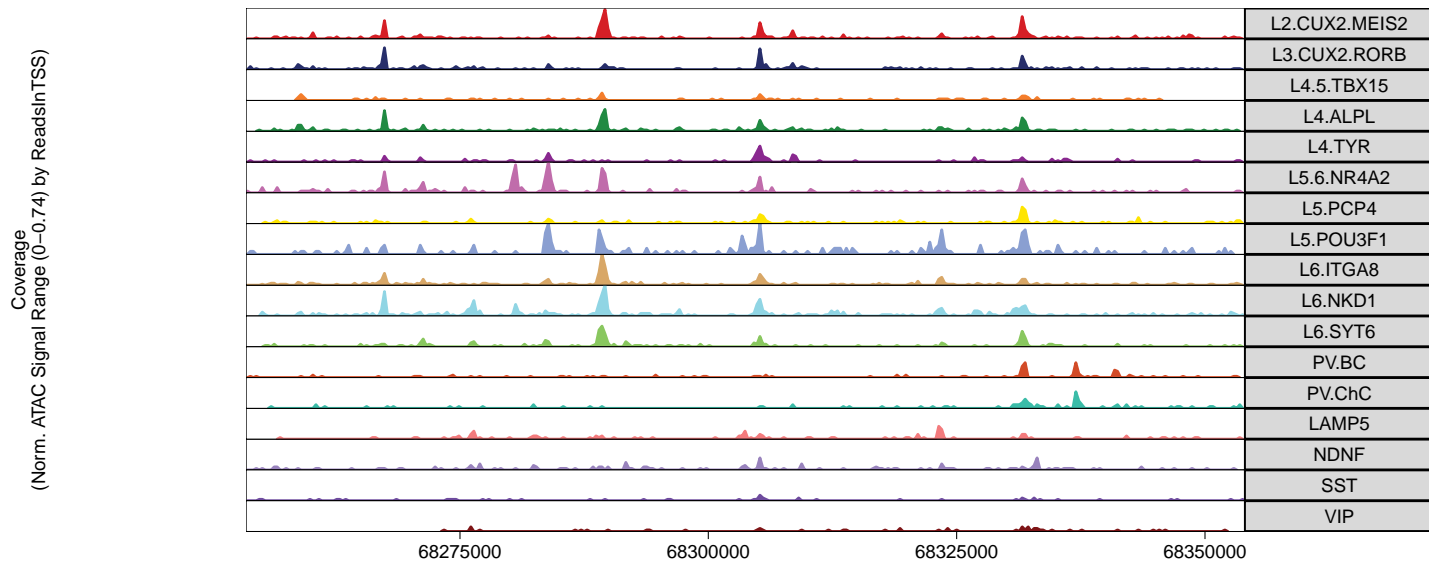


L2.CUX2.MEIS2

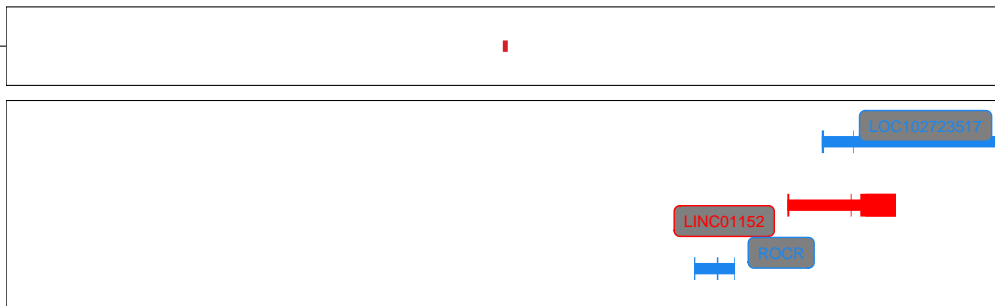




## chr16:68253472–68353973 Marker Peak Tracks for: L2.CUX2.MEIS2

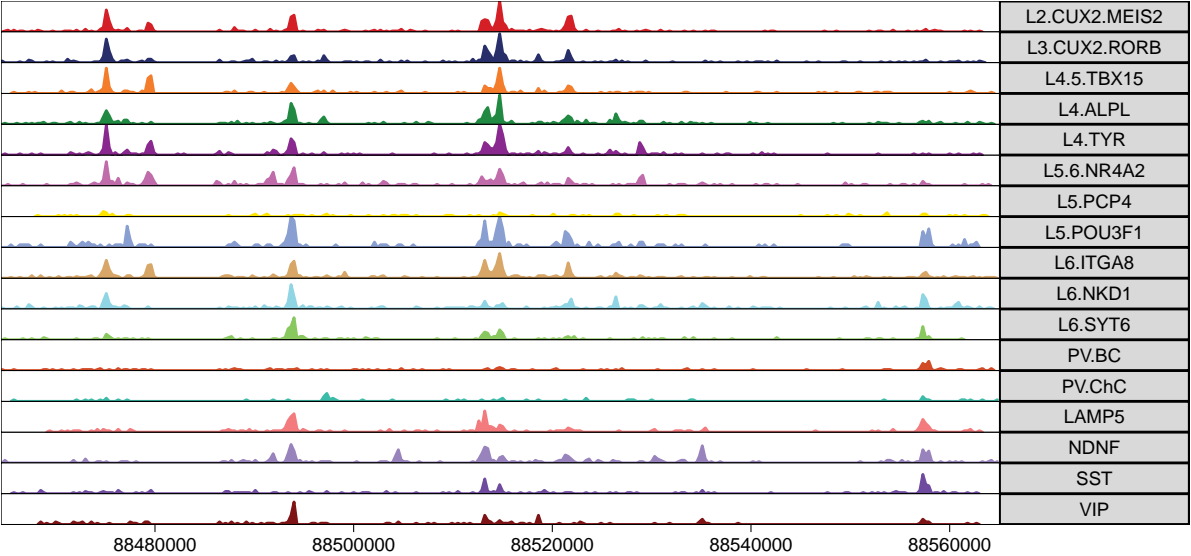


L2.CUX2.MEIS2



chr3:88464511–88565012 Marker Peak Tracks for: L2.CUX2.MEIS2

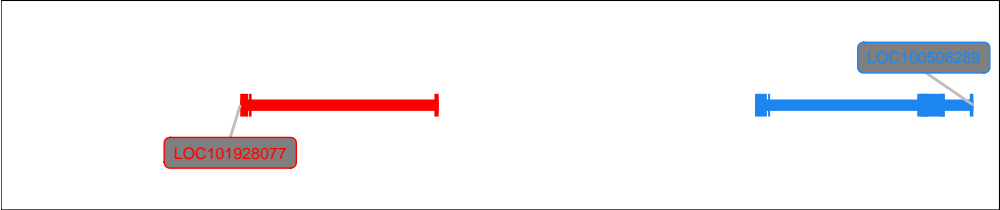
Coverage  
(Norm. ATAC Signal Range (0–0.85) by ReadsInTSS)



L2.CUX2.MEIS2

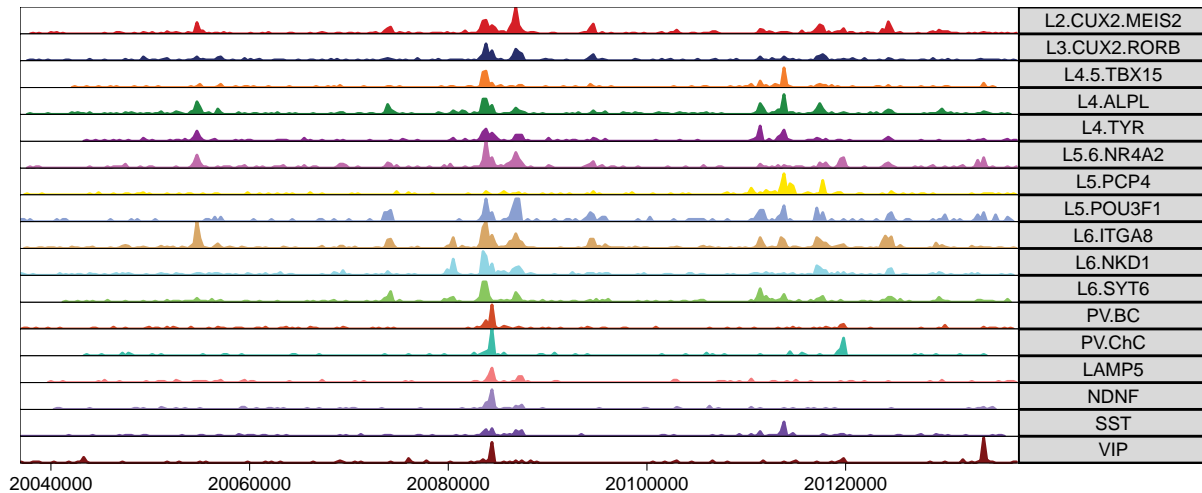


Peaks

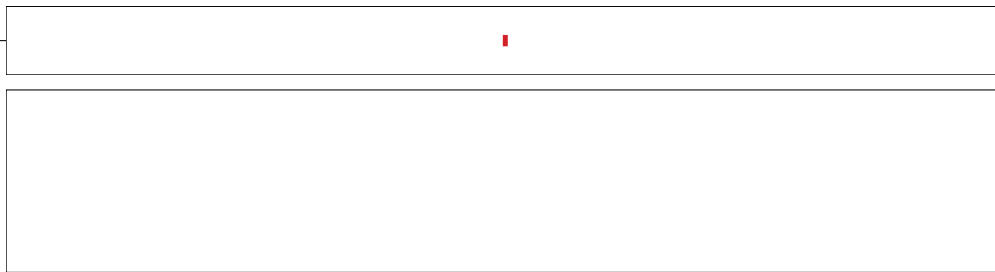


Genes

## chr1:20036899–20137400 Marker Peak Tracks for: L2.CUX2.MEIS2

Coverage  
(Norm. ATAC Signal Range (0–0.86) by ReadsInTSS)

L2.CUX2.MEIS2

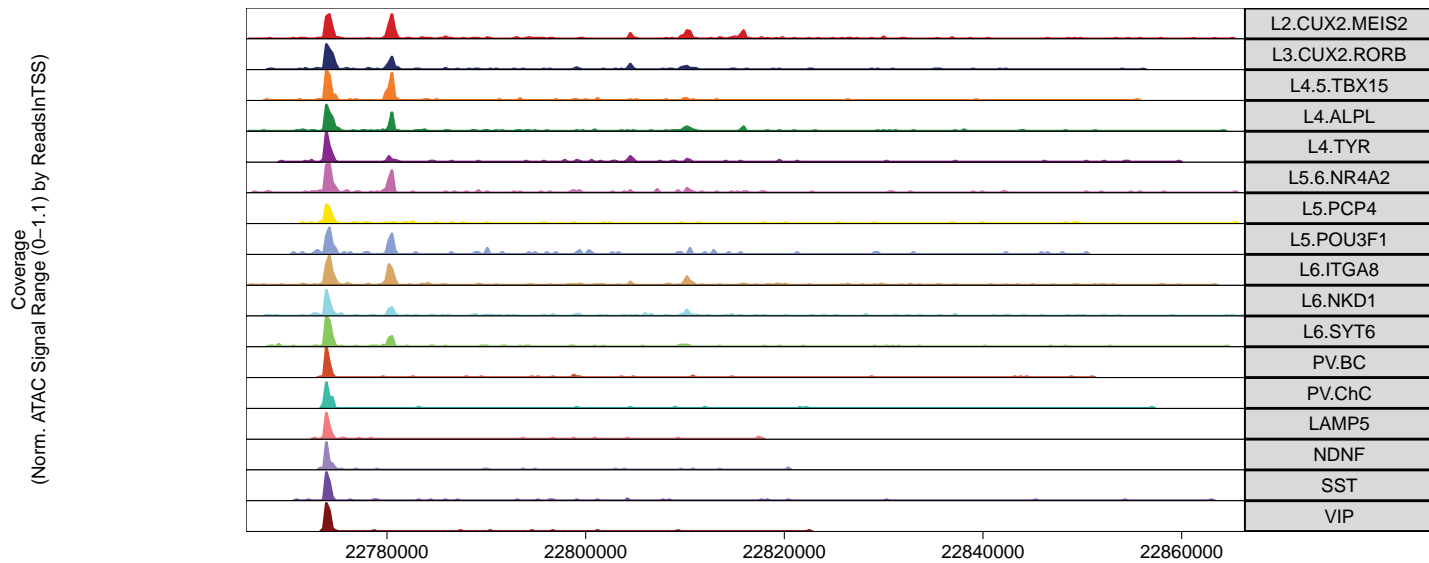


Peaks

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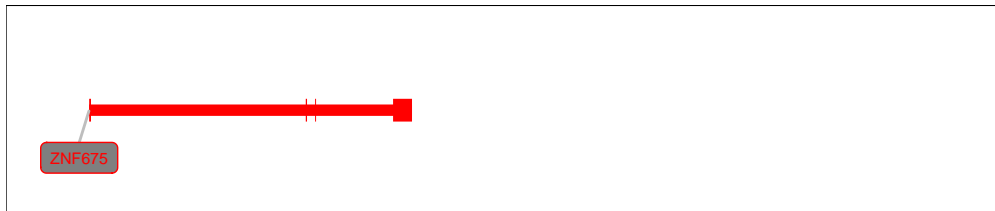
## chr19:22765819–22866320 Marker Peak Tracks for: L2.CUX2.MEIS2



L2.CUX2.MEIS2

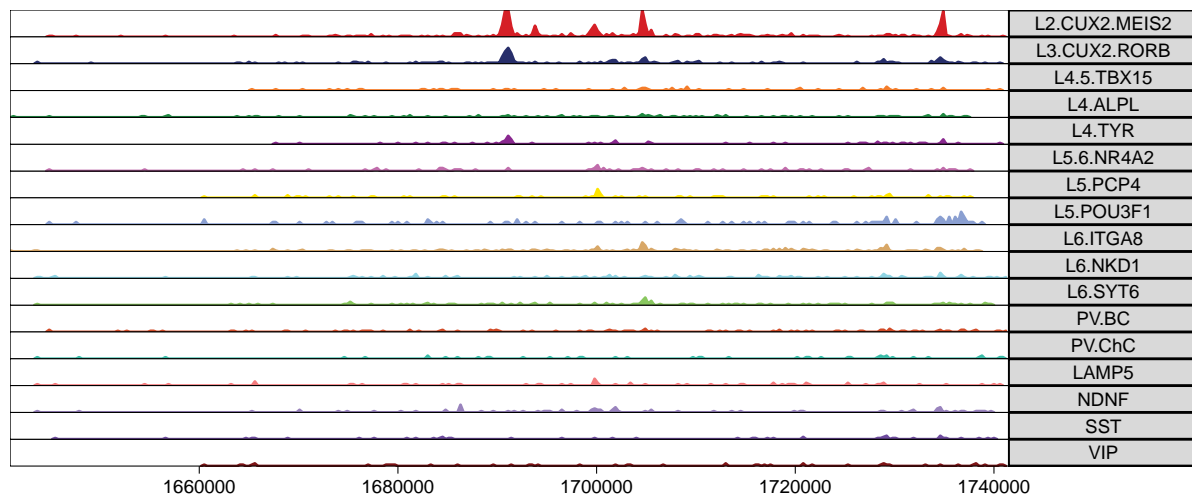


Peaks

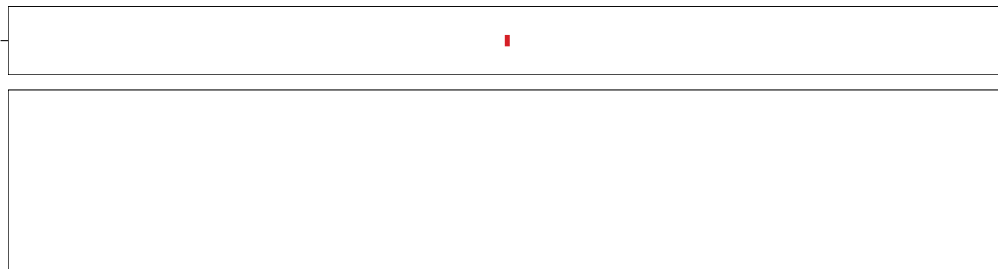


Genes

## chrX:1640959–1741460 Marker Peak Tracks for: L2.CUX2.MEIS2

Coverage  
(Norm. ATAC Signal Range (0–0.77) by ReadsInTSS)

L2.CUX2.MEIS2

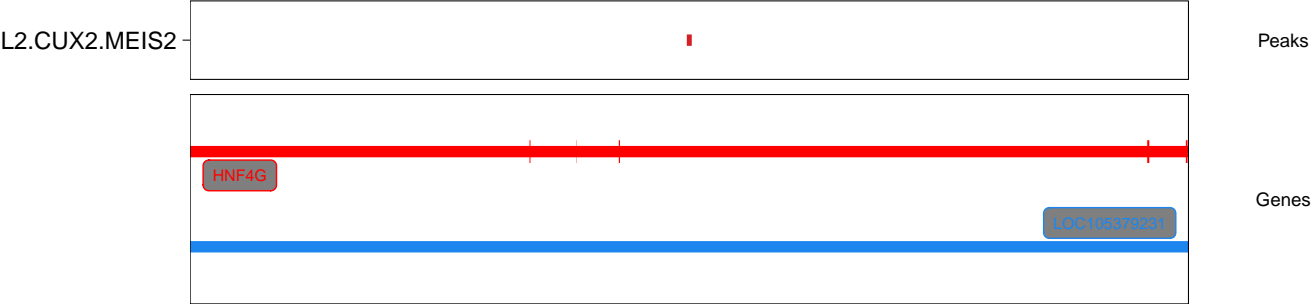
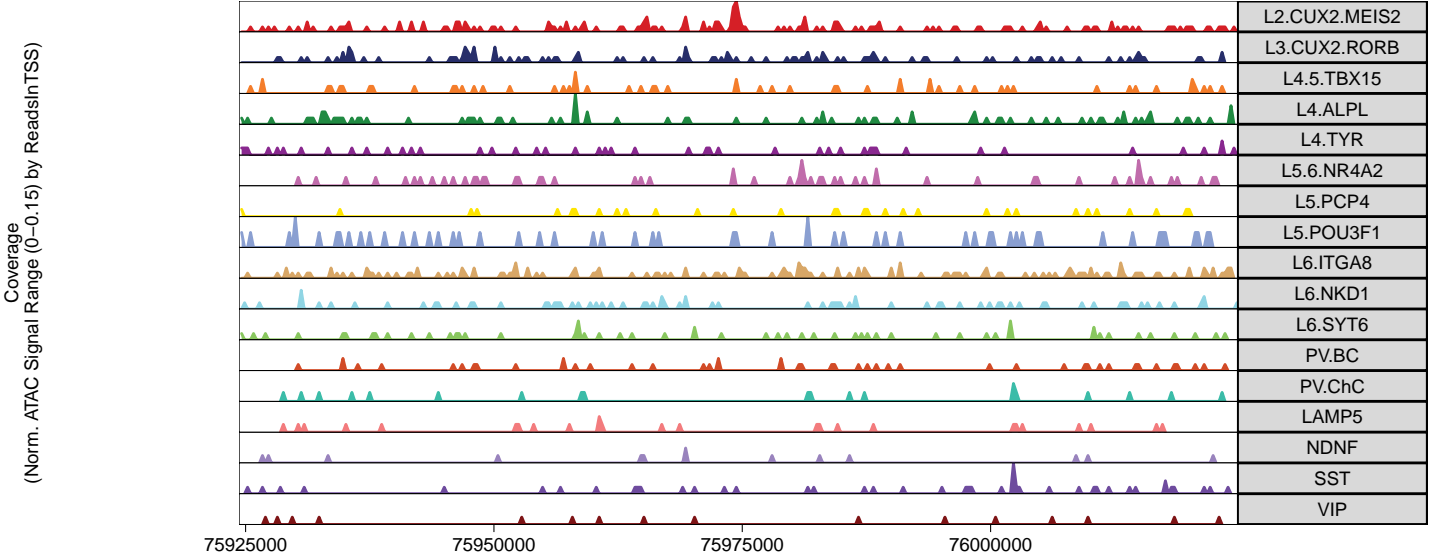


Peaks

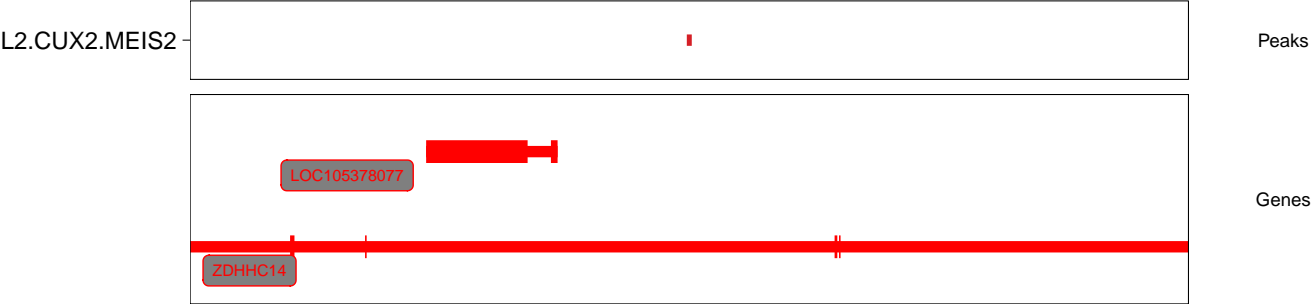
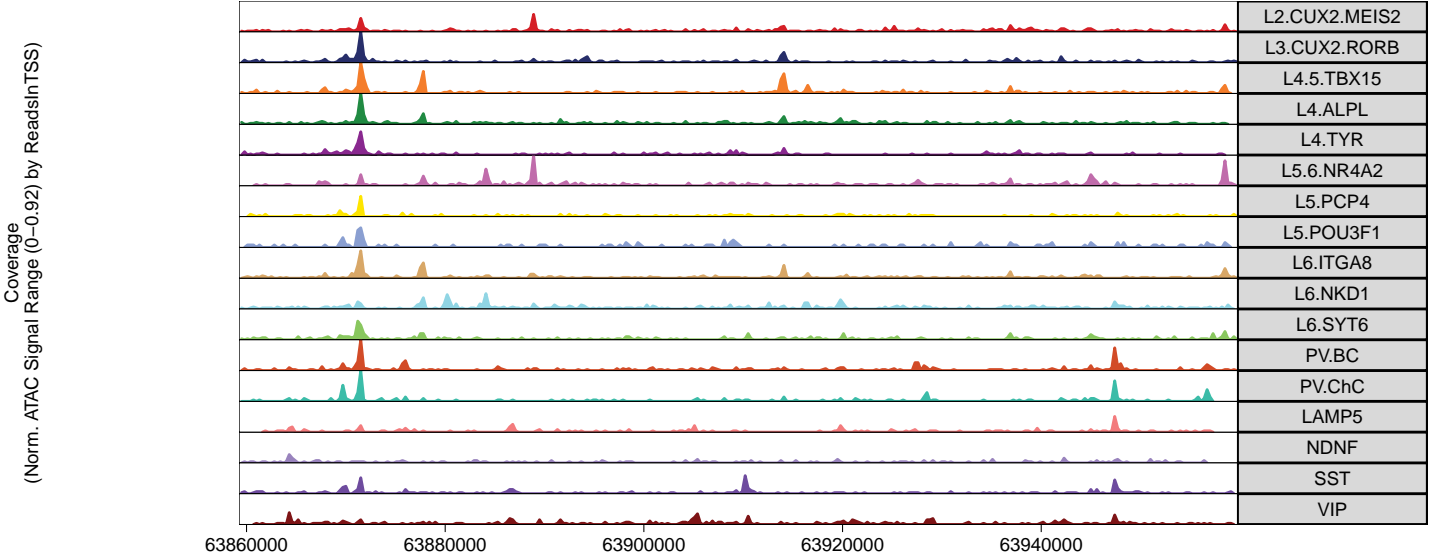
GeneTrack

strand • •

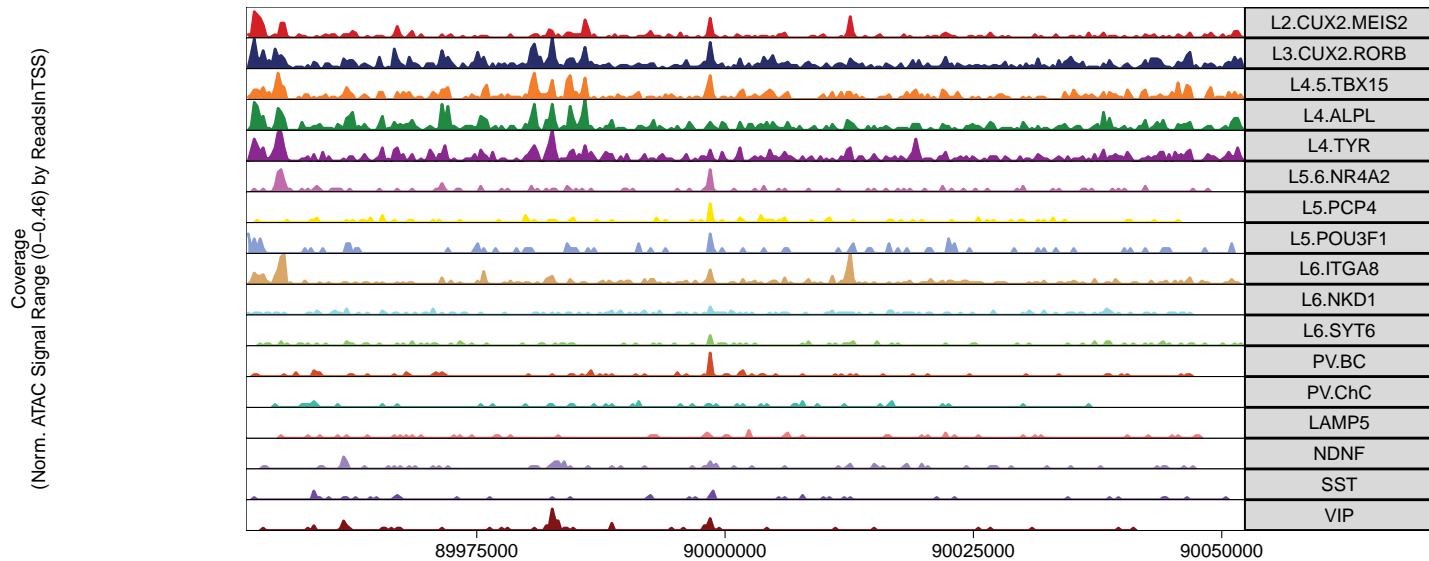
chr8:75924348–76024849 Marker Peak Tracks for: L2.CUX2.MEIS2



chr4:63859254–63959755 Marker Peak Tracks for: L2.CUX2.MEIS2



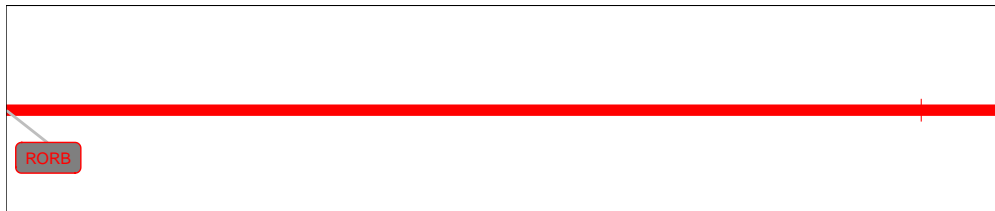
## chr15:89951783–90052284 Marker Peak Tracks for: L2.CUX2.MEIS2



L2.CUX2.MEIS2



Peaks

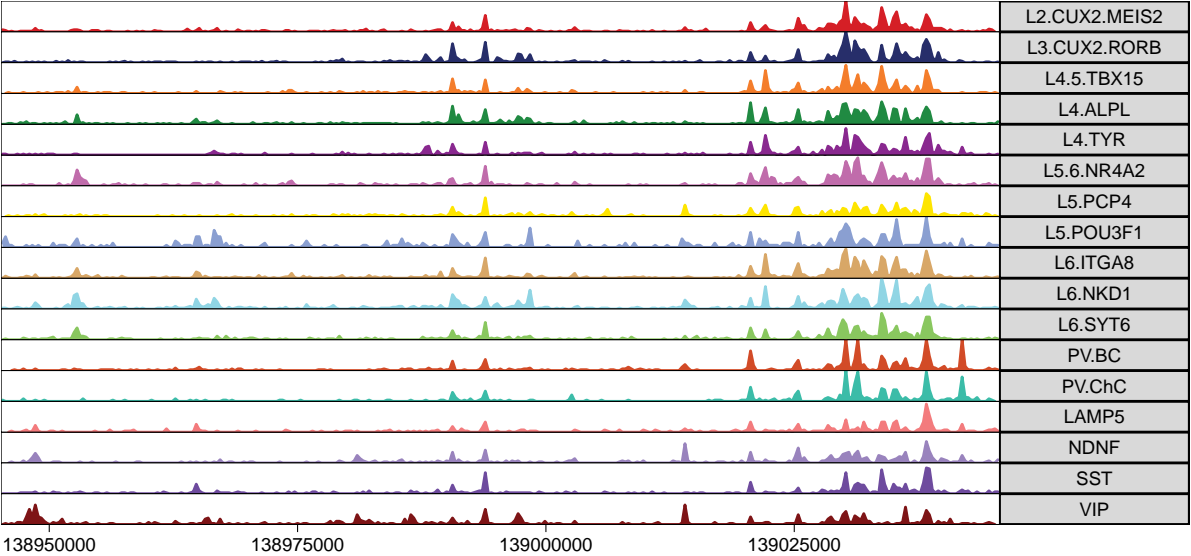


Genes



chr7:138945156–139045657 Marker Peak Tracks for: L2.CUX2.MEIS2

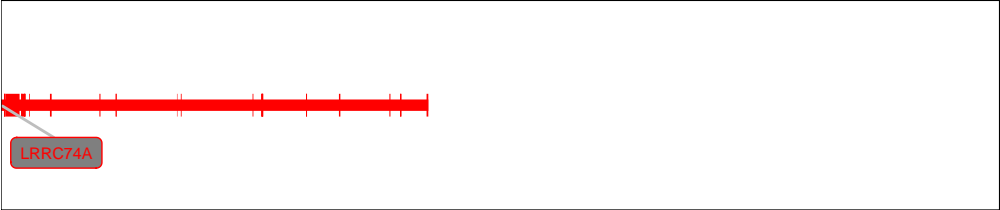
Coverage  
(Norm. ATAC Signal Range (0–1.05) by ReadsInTSS)



L2.CUX2.MEIS2

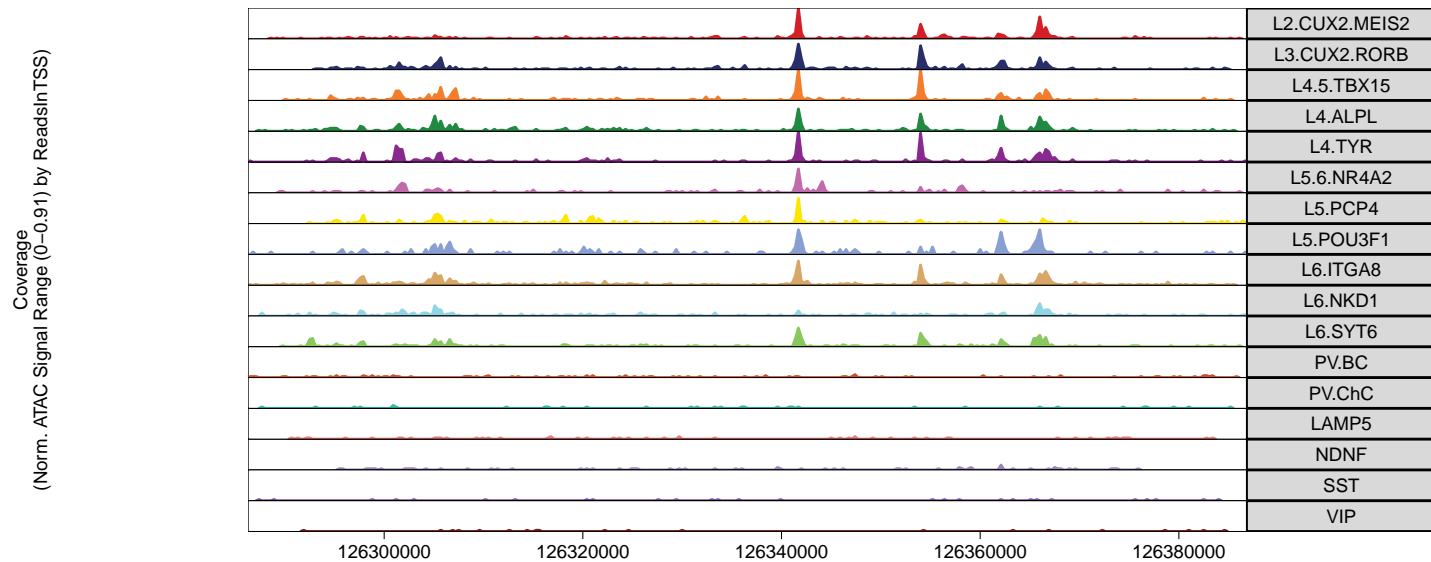


Peaks

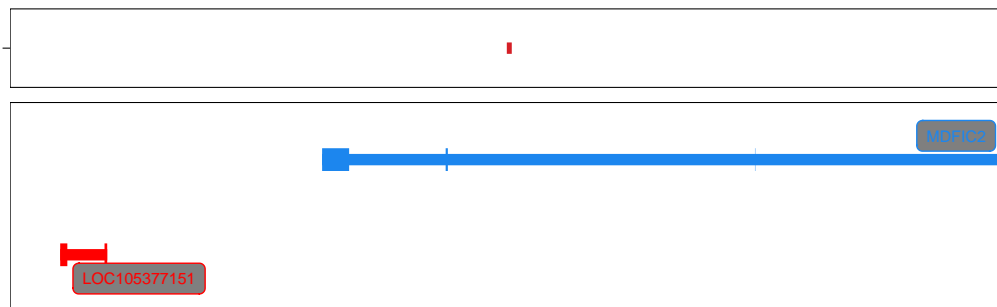


Genes

## chr2:126286304–126386805 Marker Peak Tracks for: L2.CUX2.MEIS2



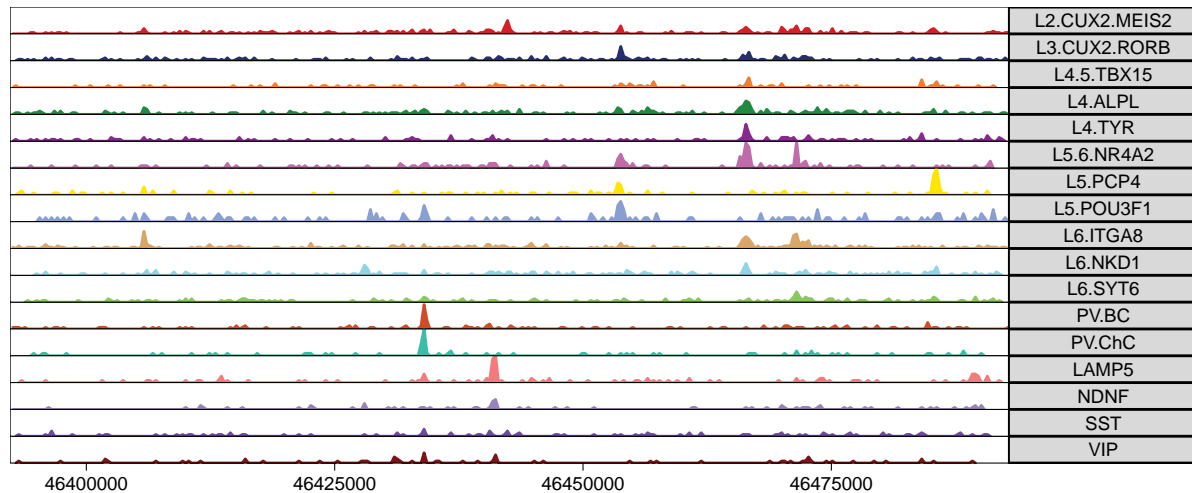
L2.CUX2.MEIS2



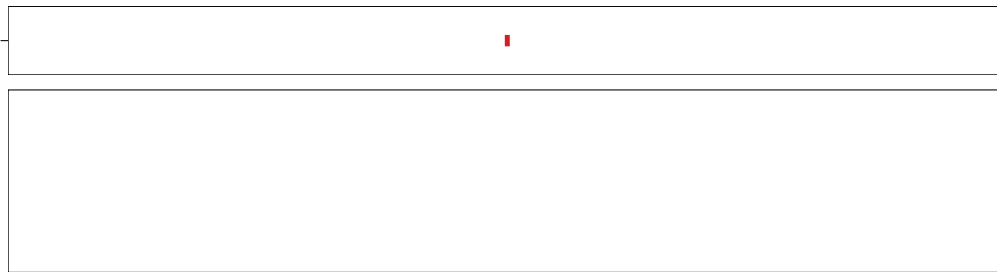
Peaks

Genes

## chr13:46392327–46492828 Marker Peak Tracks for: L2.CUX2.MEIS2

Coverage  
(Norm. ATAC Signal Range (0–0.49) by ReadsInTSS)

L2.CUX2.MEIS2

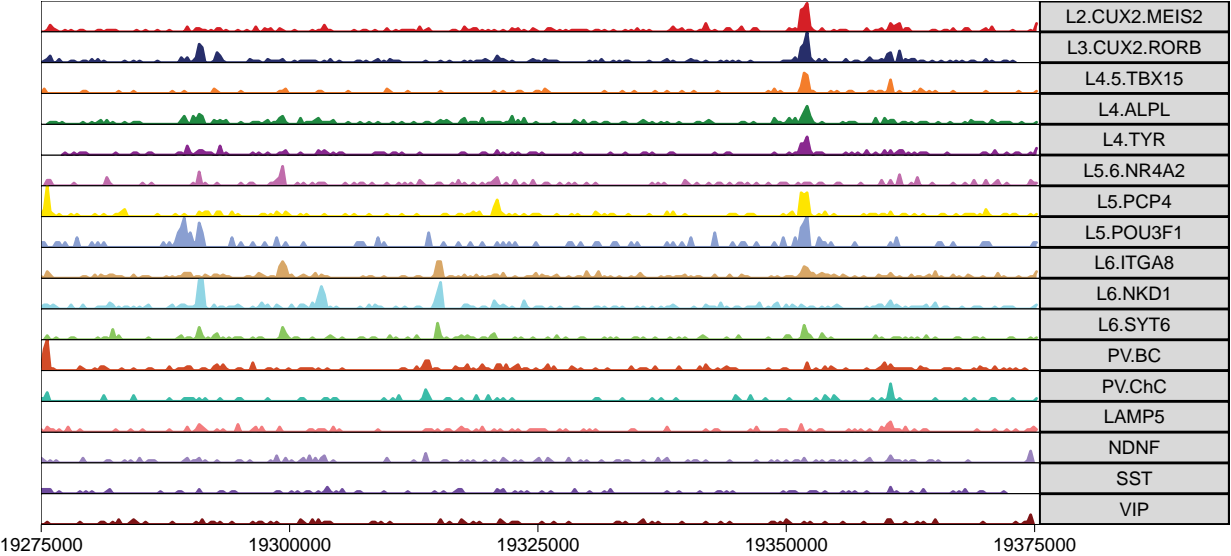


Peaks

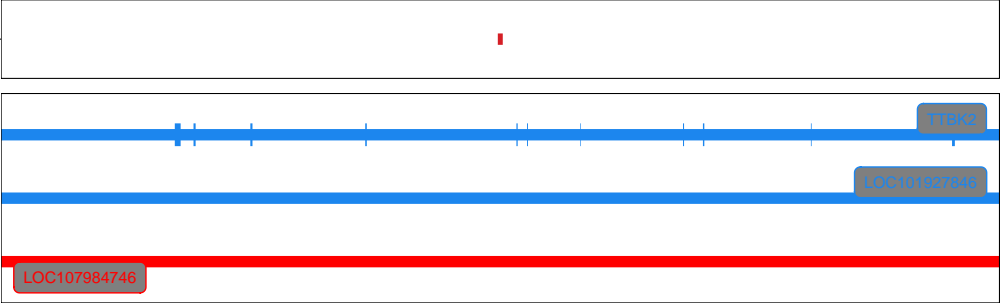
strand ● ·

chr7:19274987–19375488 Marker Peak Tracks for: L2.CUX2.MEIS2

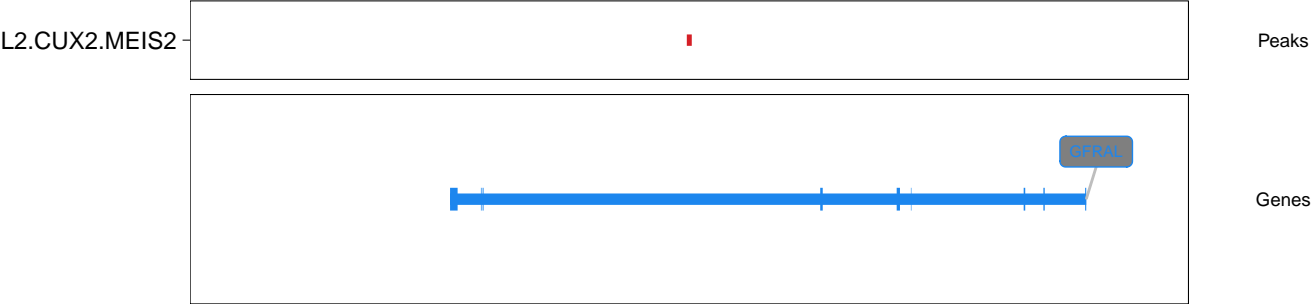
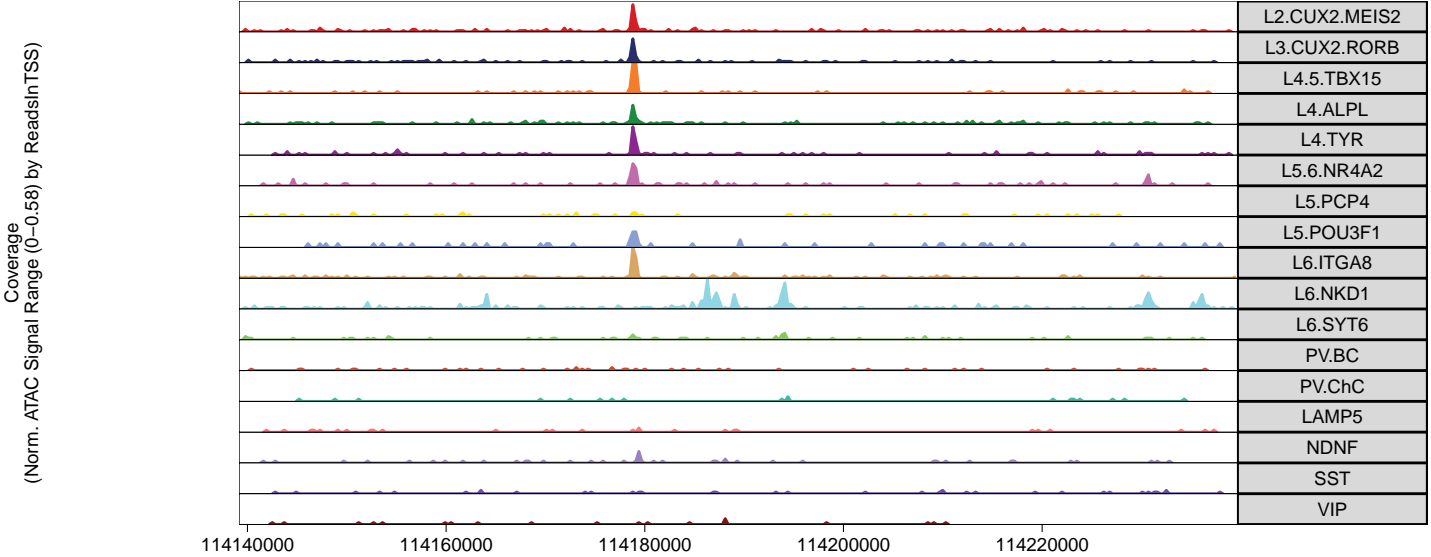
Coverage  
(Norm. ATAC Signal Range (0–0.46) by ReadsInTSS)



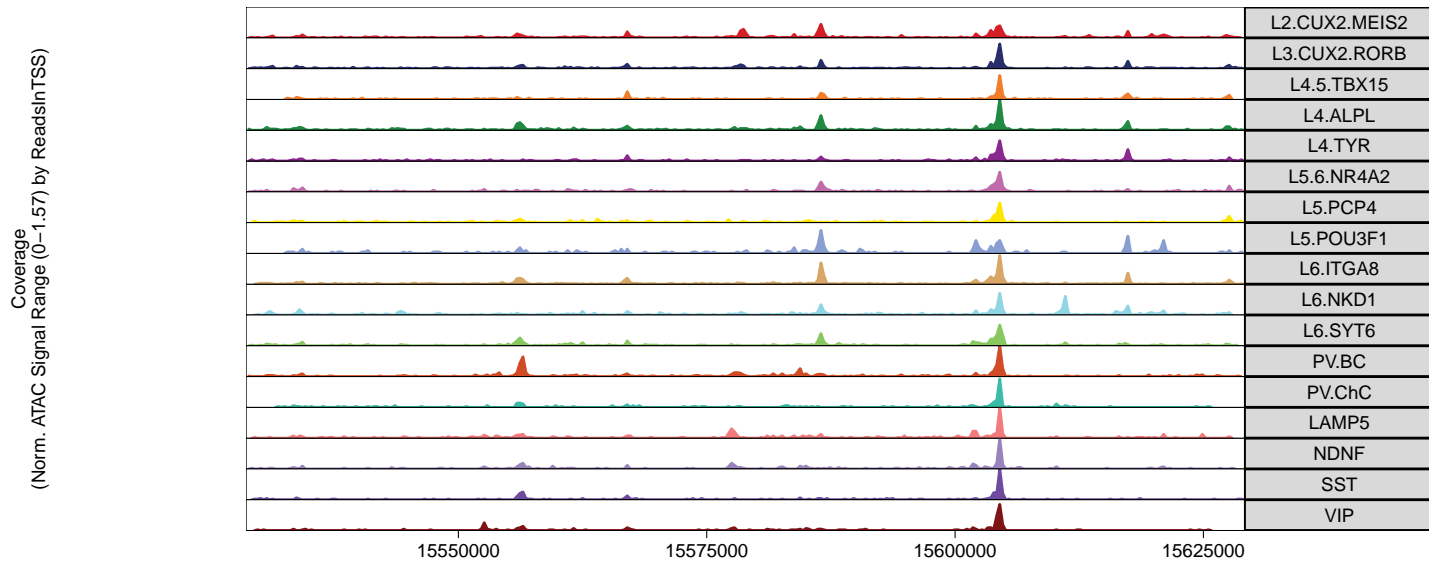
L2.CUX2.MEIS2



chr4:114139161–114239662 Marker Peak Tracks for: L2.CUX2.MEIS2



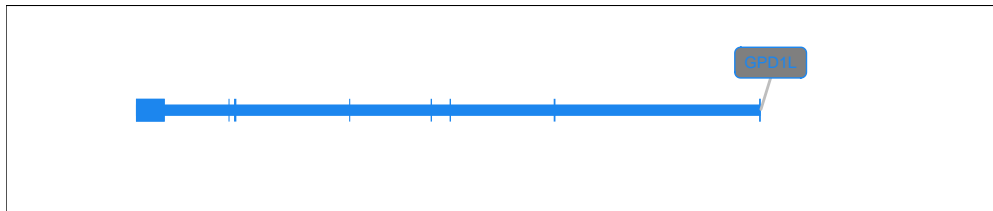
## chr2:15528631–15629132 Marker Peak Tracks for: L2.CUX2.MEIS2



L2.CUX2.MEIS2

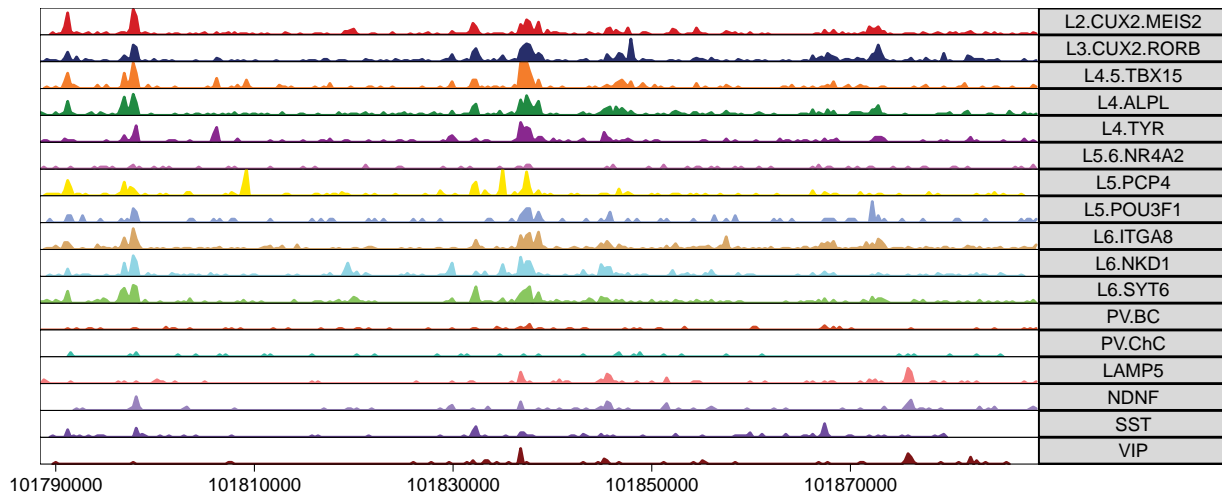


Peaks

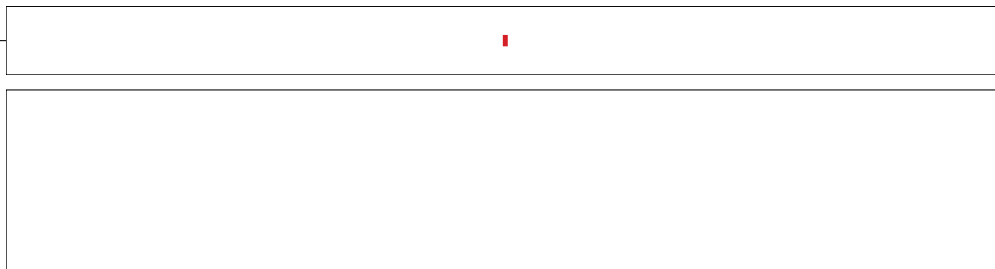


Genes

## chr3:101788427–101888928 Marker Peak Tracks for: L2.CUX2.MEIS2

Coverage  
(Norm. ATAC Signal Range (0–0.56) by ReadsInTSS)

L2.CUX2.MEIS2

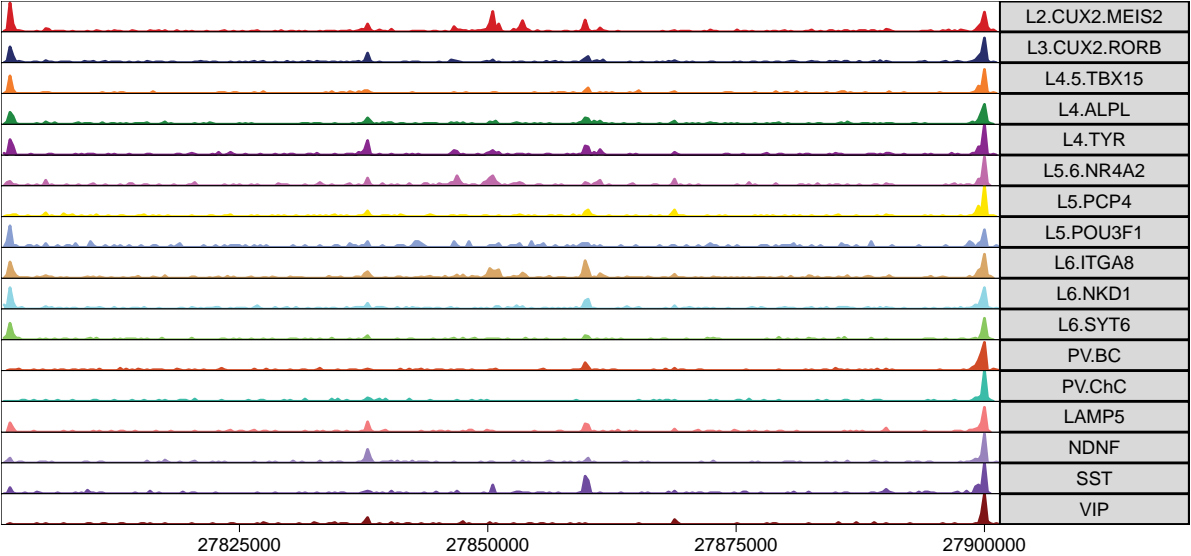


Peaks

GeneTrack

chr10:27801007–27901508 Marker Peak Tracks for: L2.CUX2.MEIS2

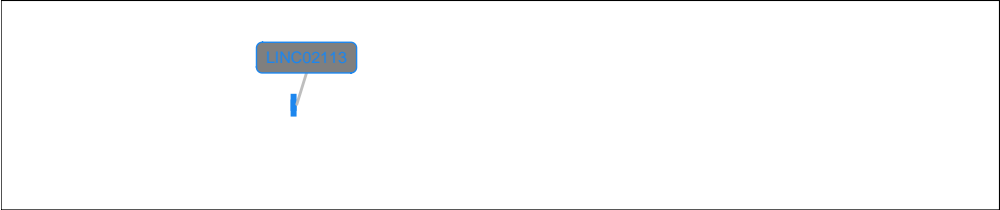
Coverage  
(Norm. ATAC Signal Range (0–1.15) by ReadsInTSS)



L2.CUX2.MEIS2



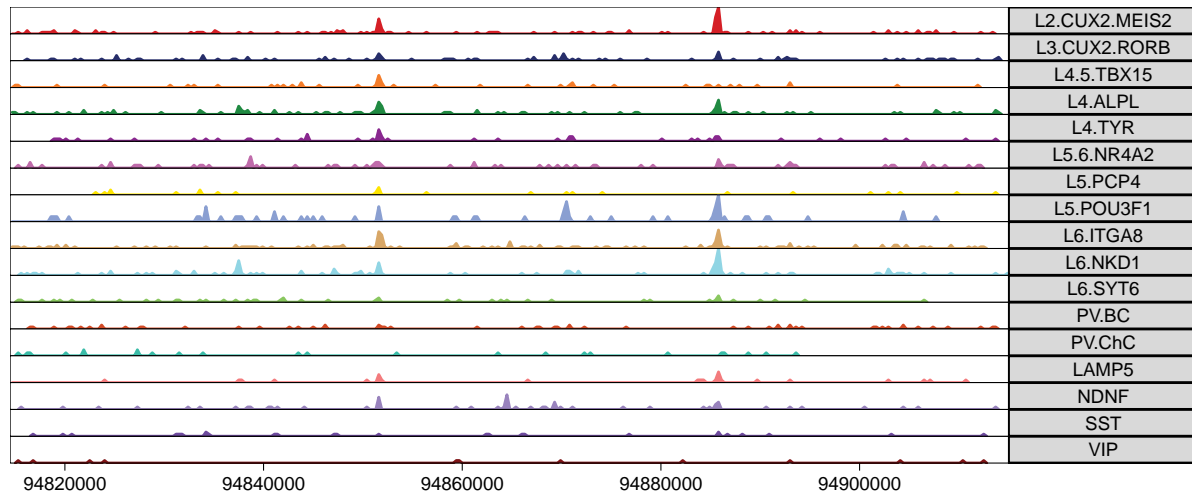
Peaks



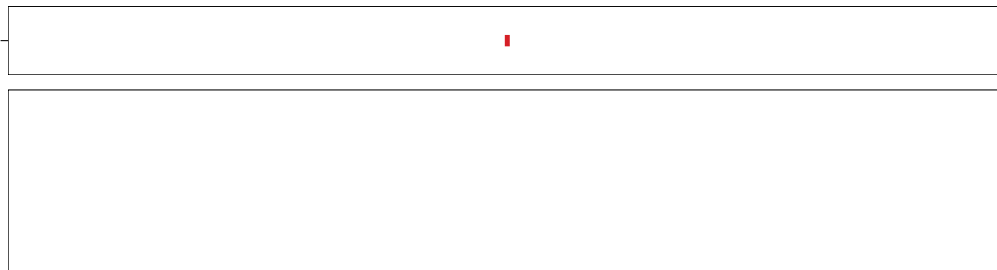
Genes



## chr6:94814486–94914987 Marker Peak Tracks for: L2.CUX2.MEIS2

Coverage  
(Norm. ATAC Signal Range (0–0.39) by ReadsInTSS)

L2.CUX2.MEIS2

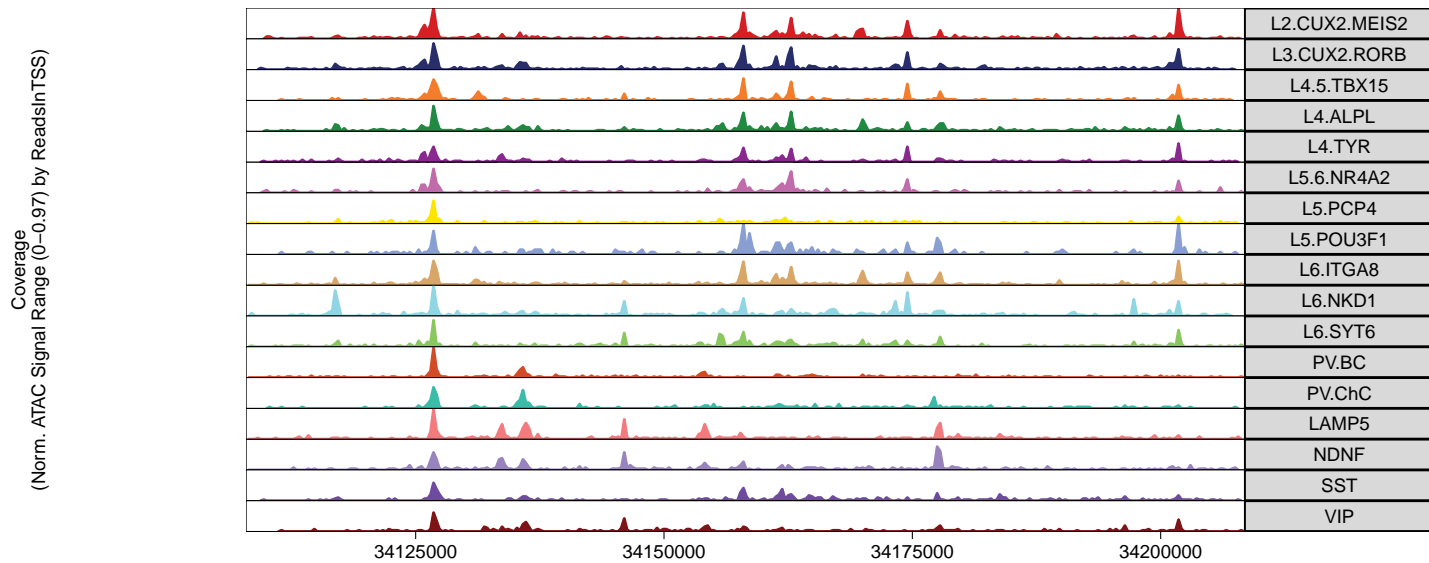


Peaks

GeneTrack

strand ● ·

## chr20:34107932–34208433 Marker Peak Tracks for: L2.CUX2.MEIS2



L2.CUX2.MEIS2

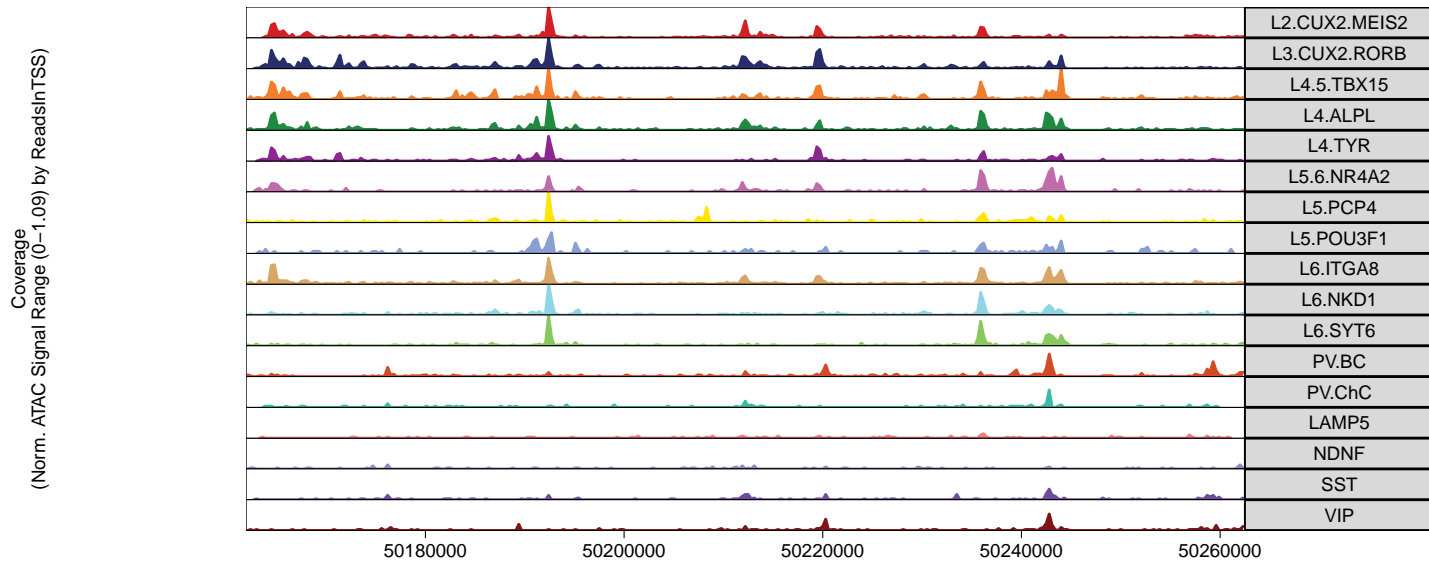


Peaks



Genes

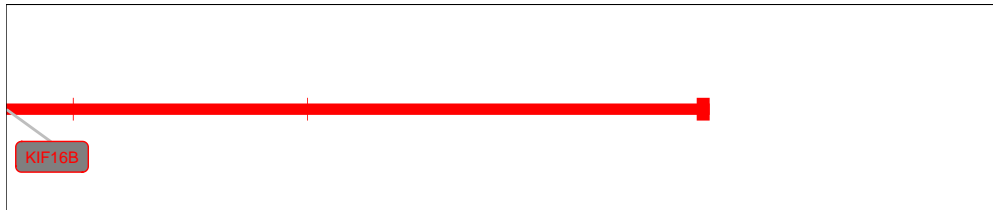
## chr10:50161950–50262451 Marker Peak Tracks for: L2.CUX2.MEIS2



L2.CUX2.MEIS2

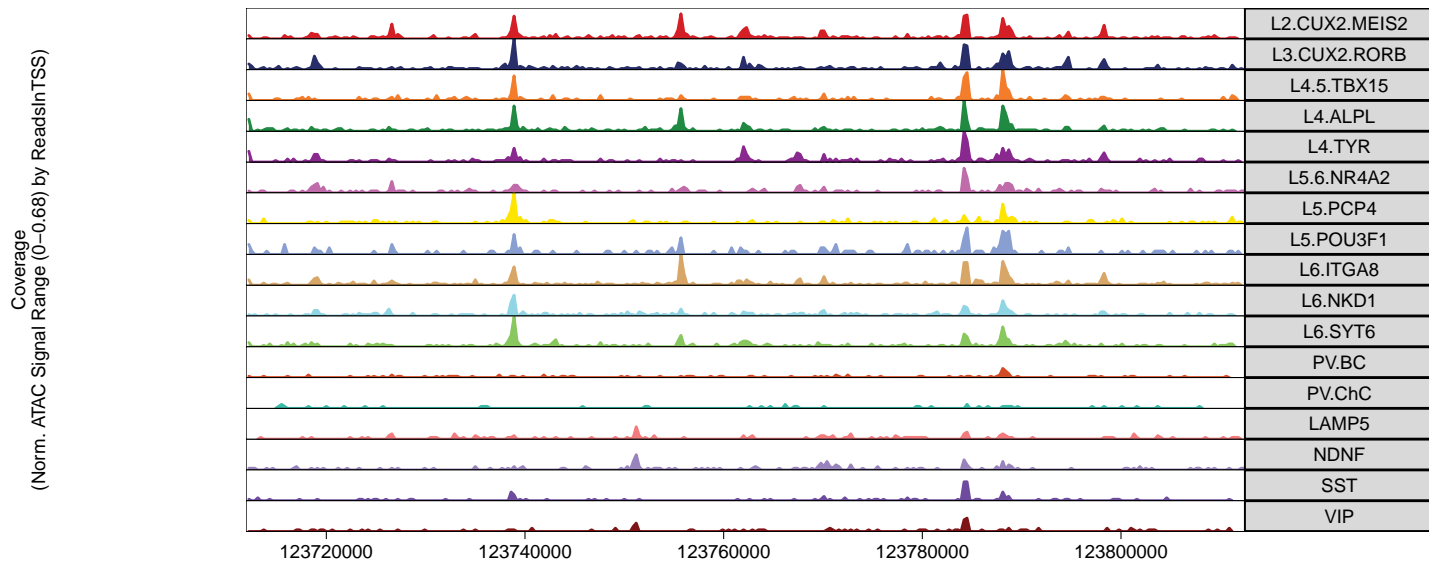


Peaks



Genes

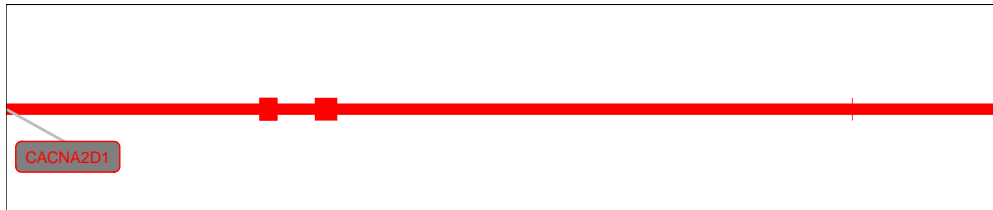
## chr3:123711926–123812427 Marker Peak Tracks for: L2.CUX2.MEIS2



L2.CUX2.MEIS2



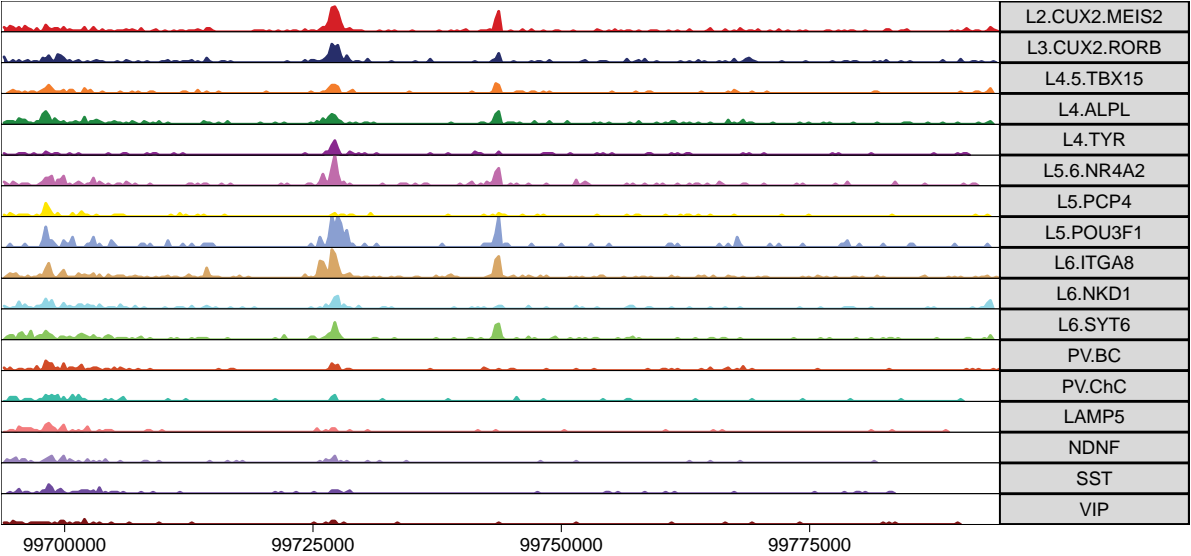
Peaks



Genes

chr7:99693605–99794106 Marker Peak Tracks for: L2.CUX2.MEIS2

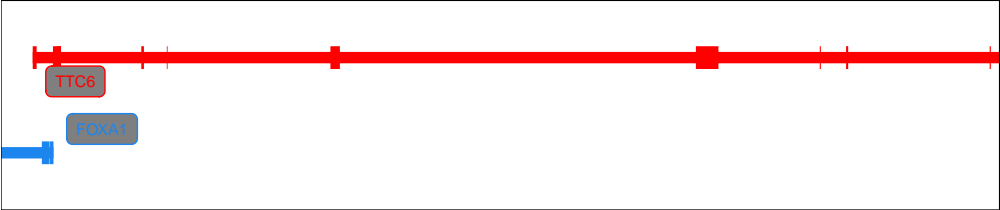
Coverage  
(Norm. ATAC Signal Range (0–0.66) by ReadsInTSS)



L2.CUX2.MEIS2

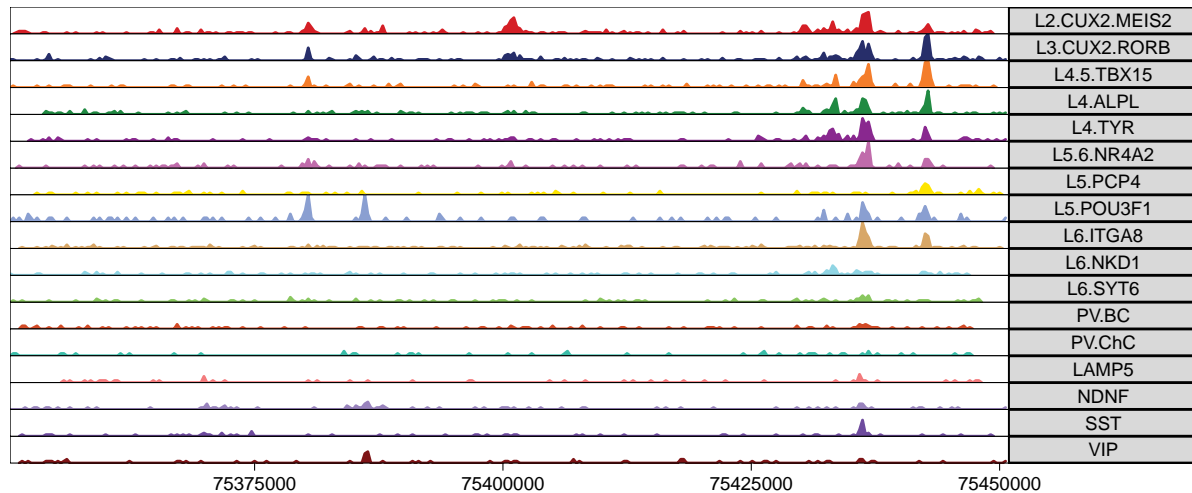


Peaks

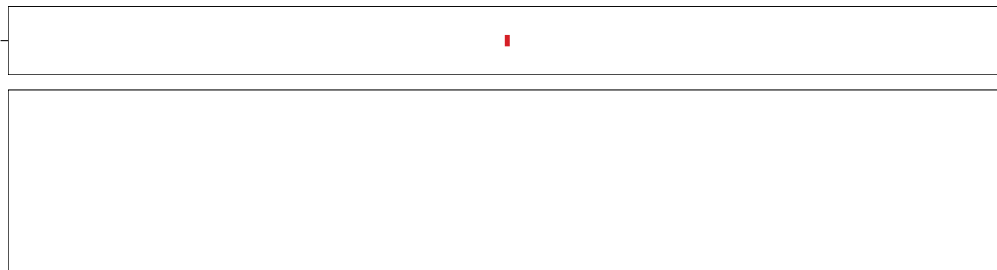


Genes

## chr4:75350388–75450889 Marker Peak Tracks for: L2.CUX2.MEIS2

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

L2.CUX2.MEIS2

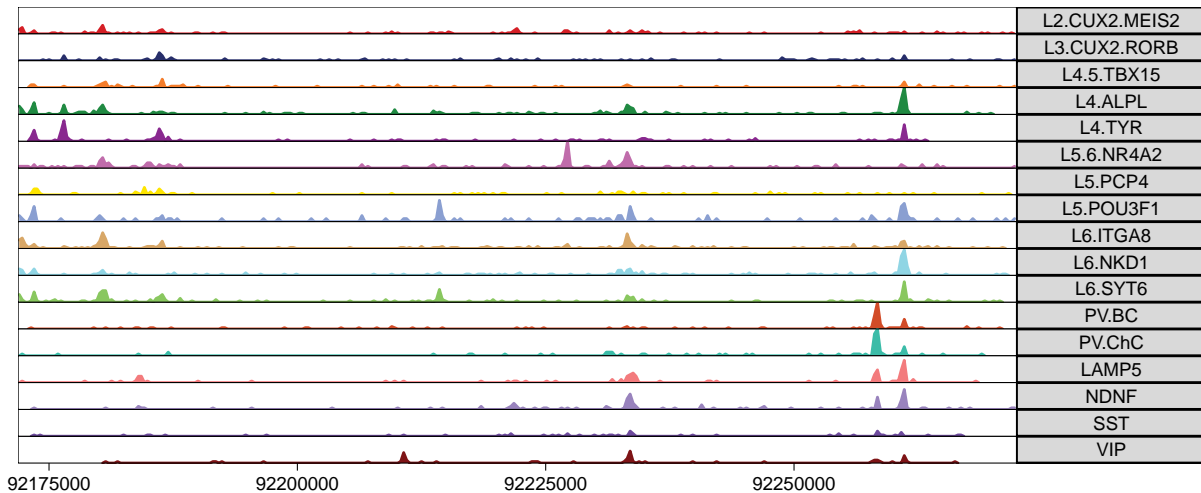


Peaks

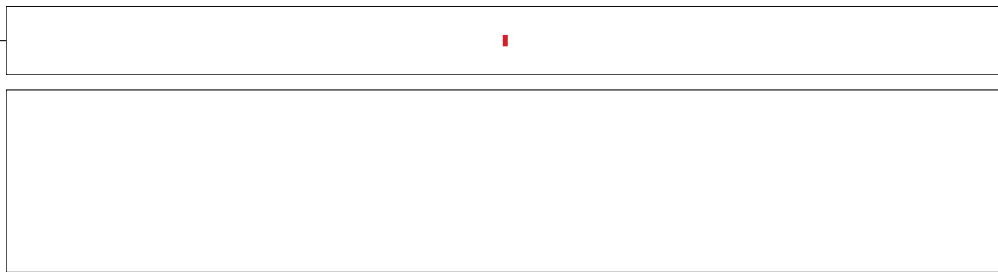
GeneTrack

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## chr7:92171889–92272390 Marker Peak Tracks for: L2.CUX2.MEIS2

Coverage  
(Norm. ATAC Signal Range (0–0.65) by ReadsInTSS)

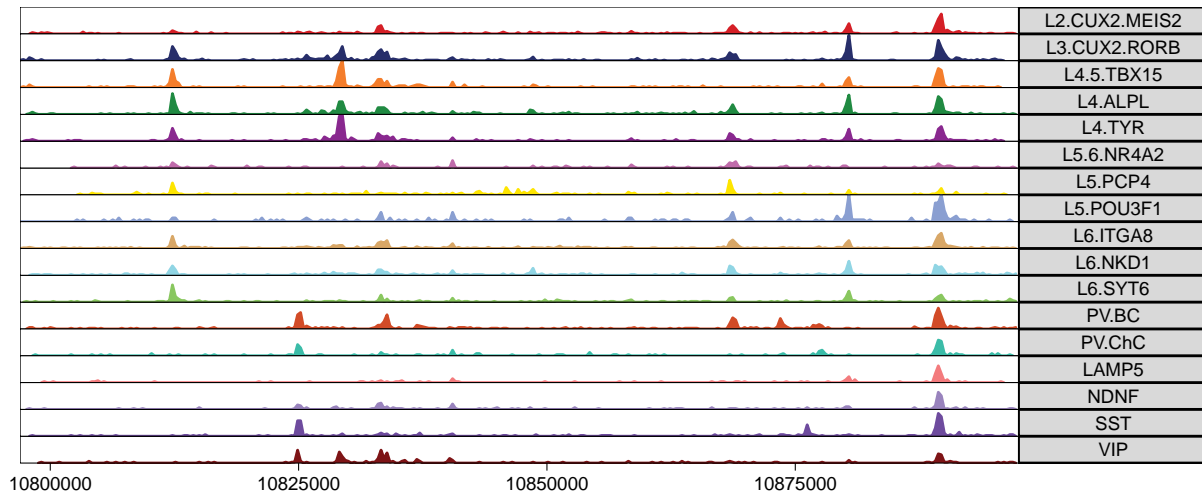
L2.CUX2.MEIS2



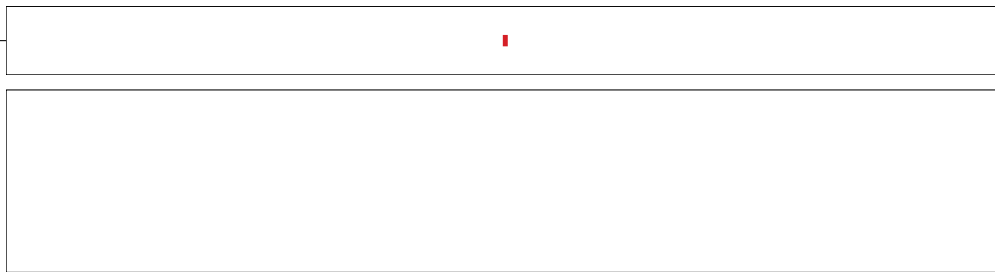
Peaks

strand ● ·

## chr6:10796965–10897466 Marker Peak Tracks for: L2.CUX2.MEIS2

Coverage  
(Norm. ATAC Signal Range (0–1.05) by ReadsInTSS)

L2.CUX2.MEIS2

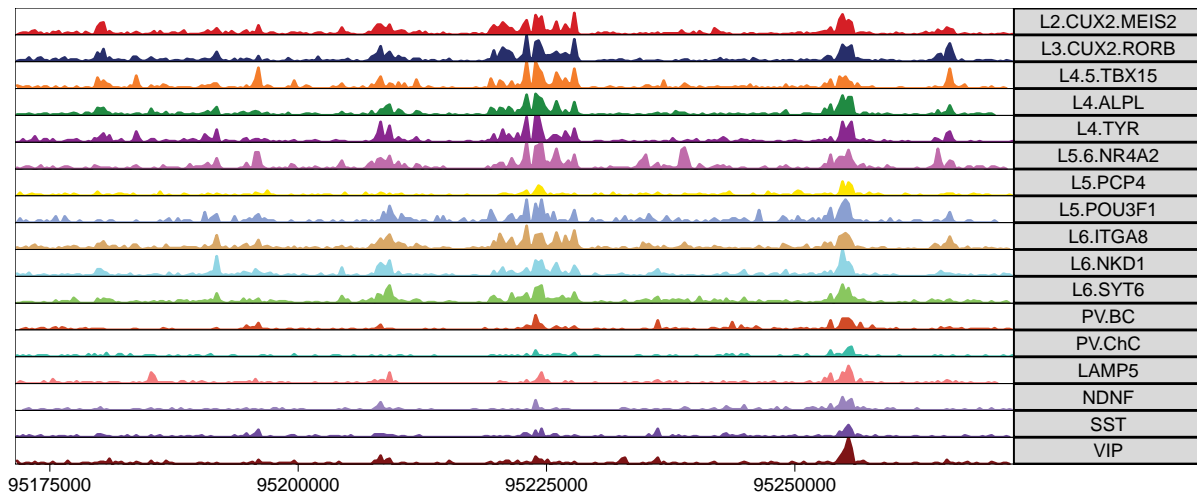


Peaks

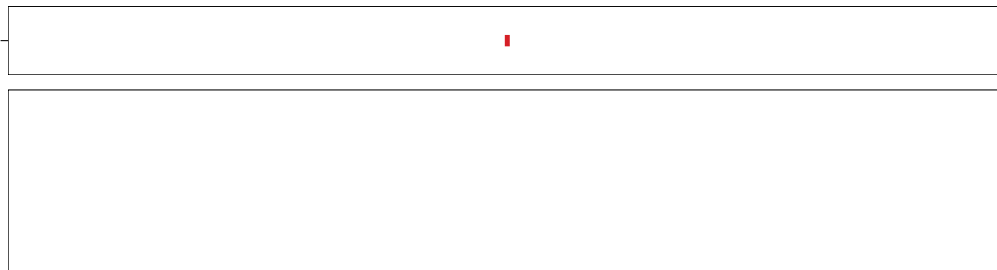
strand • •



## chr17:95171505–95272006 Marker Peak Tracks for: L2.CUX2.MEIS2

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

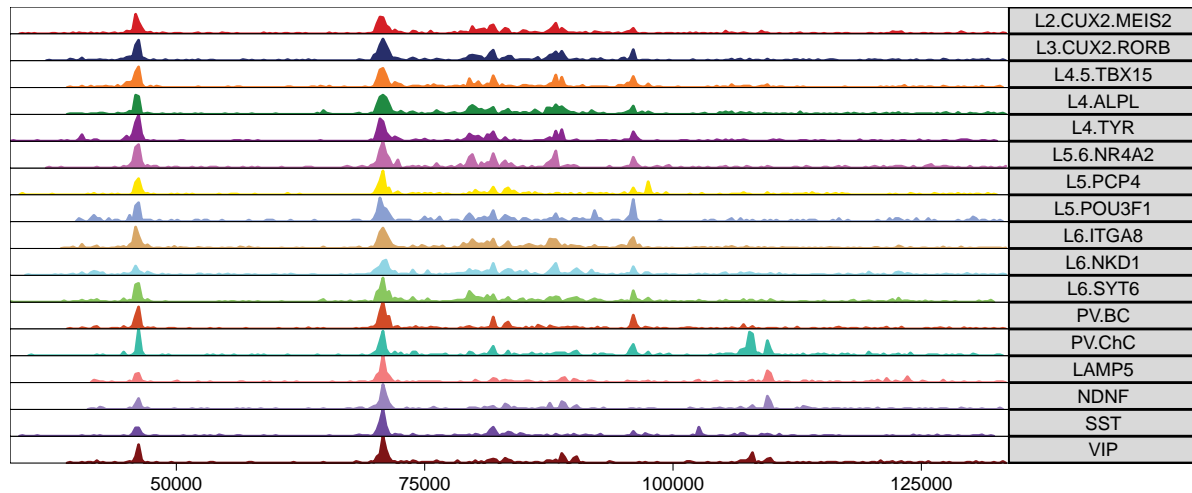
L2.CUX2.MEIS2



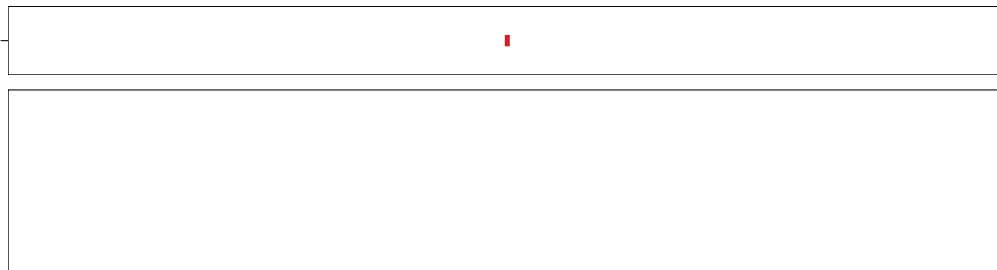
Peaks

strand • ·

## chr11:33266–133767 Marker Peak Tracks for: L2.CUX2.MEIS2

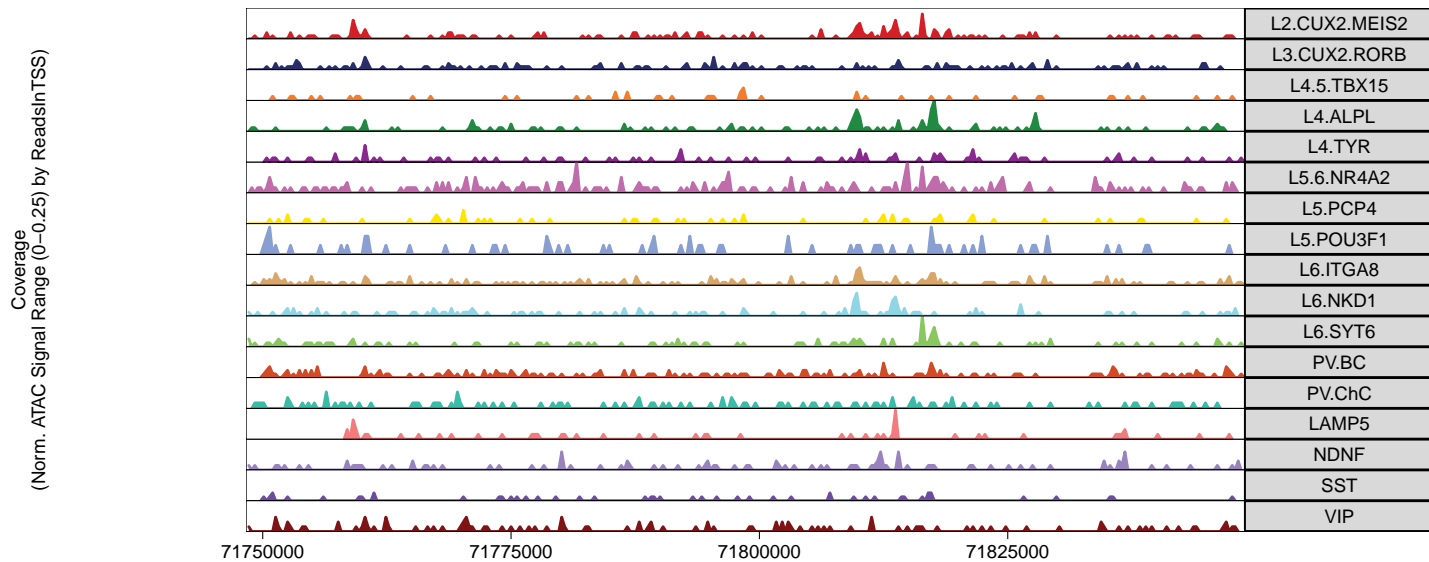
Coverage  
(Norm. ATAC Signal Range (0–1.25) by ReadsInTSS)

L2.CUX2.MEIS2



strand ● ·

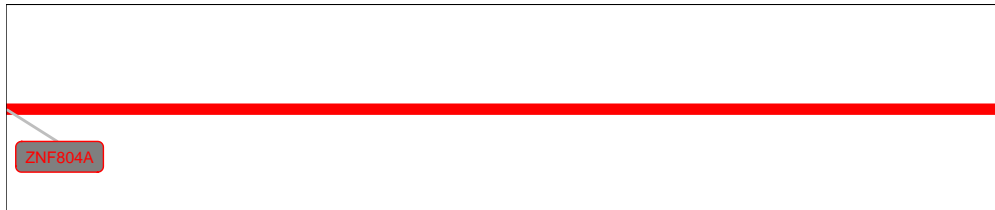
## chr12:71748329–71848830 Marker Peak Tracks for: L2.CUX2.MEIS2



L2.CUX2.MEIS2

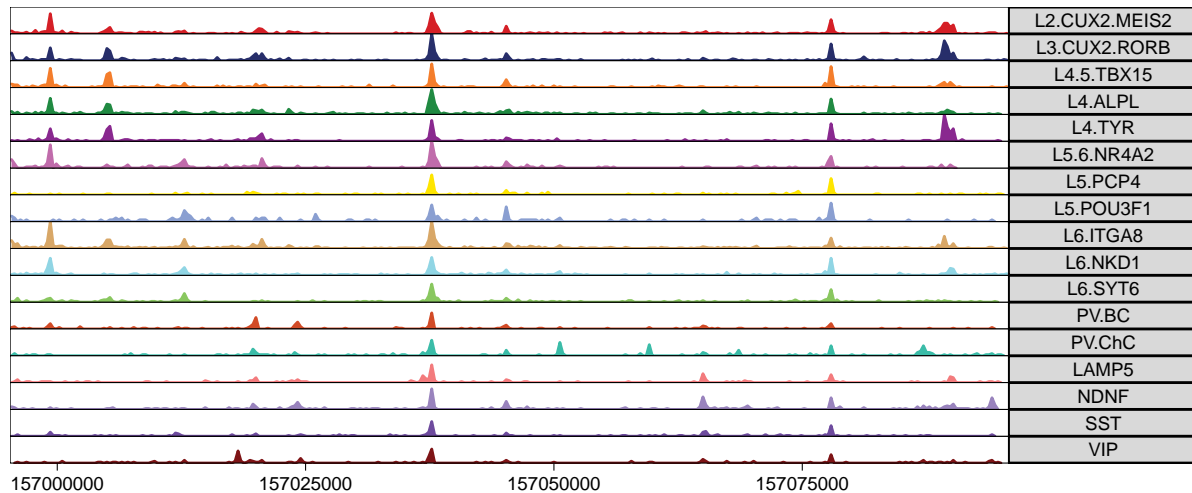


Peaks

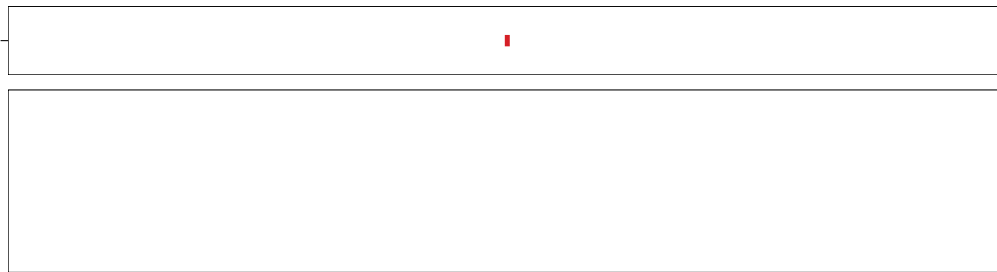


Genes

## chr5:156995255–157095756 Marker Peak Tracks for: L2.CUX2.MEIS2

Coverage  
(Norm. ATAC Signal Range (0–1.06) by ReadsInTSS)

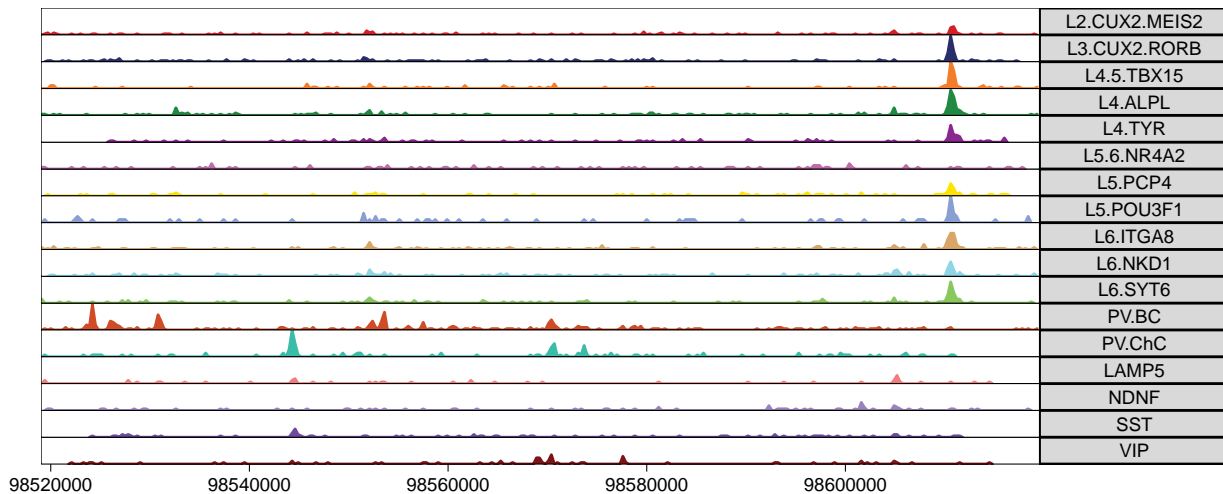
L2.CUX2.MEIS2



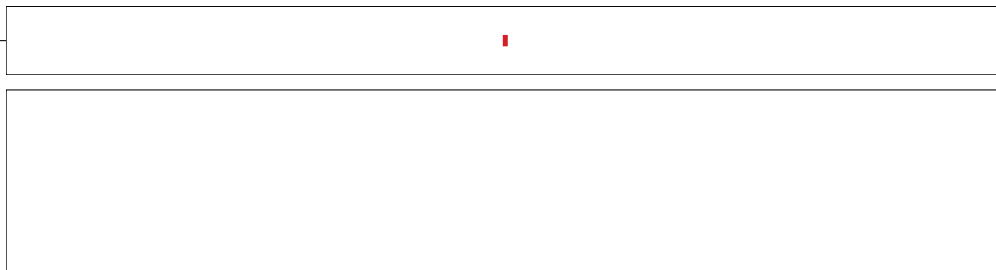
Peaks

GeneTrack

## chr12:98519032–98619533 Marker Peak Tracks for: L2.CUX2.MEIS2

Coverage  
(Norm. ATAC Signal Range (0–0.63) by ReadsInTSS)

L2.CUX2.MEIS2

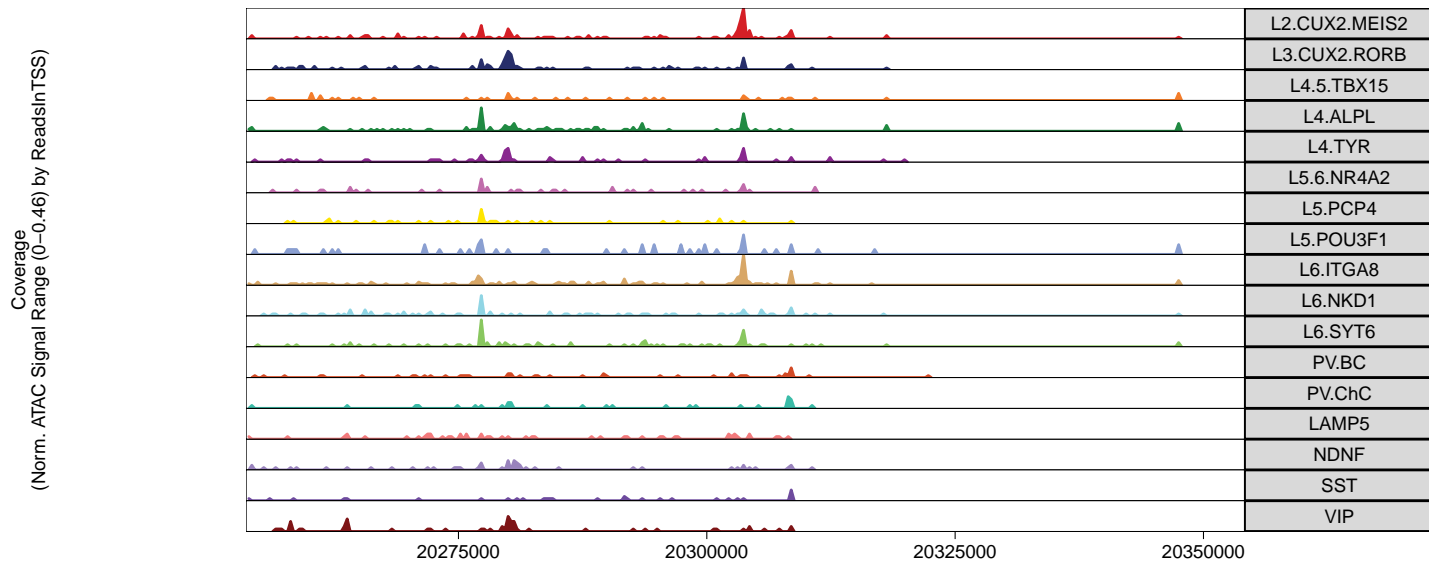


Peaks

GeneTrack

strand ● ·

## chr1:20253624–20354125 Marker Peak Tracks for: L2.CUX2.MEIS2



L2.CUX2.MEIS2

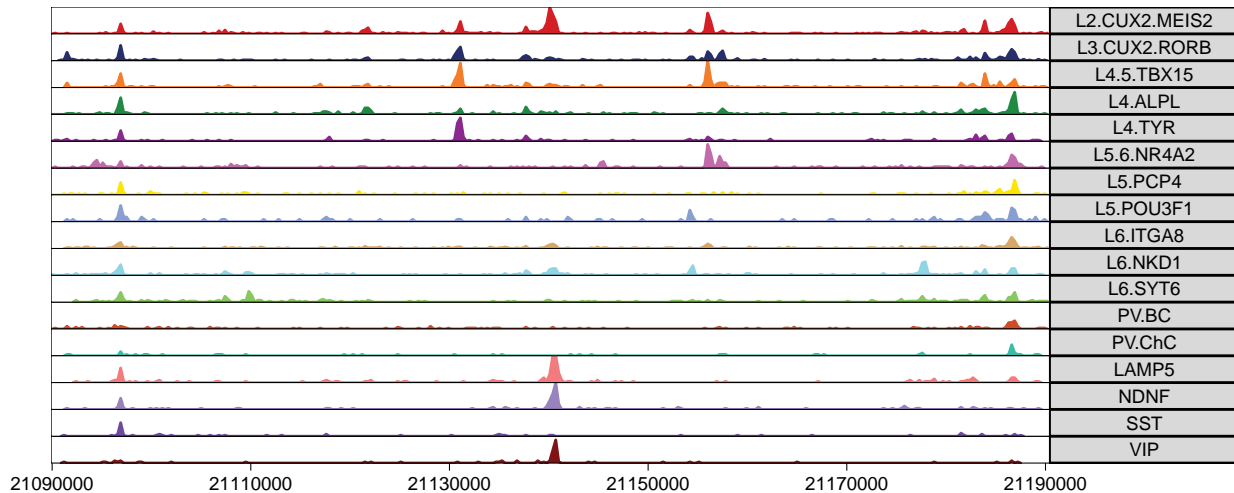


Peaks

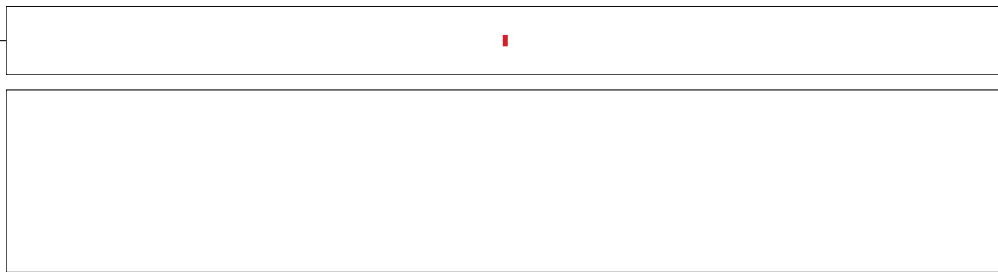


Genes

## chr1:21089899–21190400 Marker Peak Tracks for: L2.CUX2.MEIS2

Coverage  
(Norm. ATAC Signal Range (0–0.85) by ReadsInTSS)

L2.CUX2.MEIS2

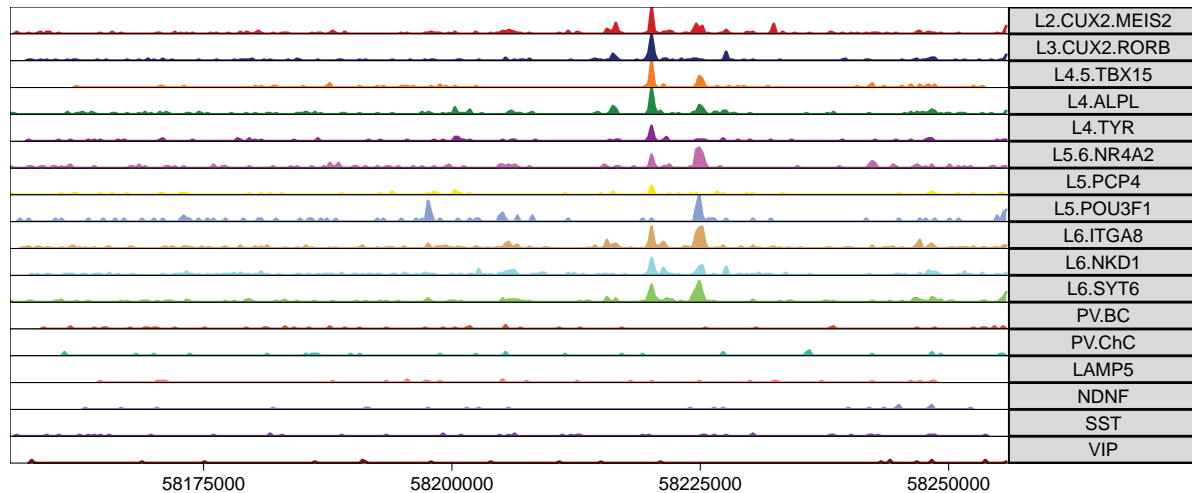


Peaks

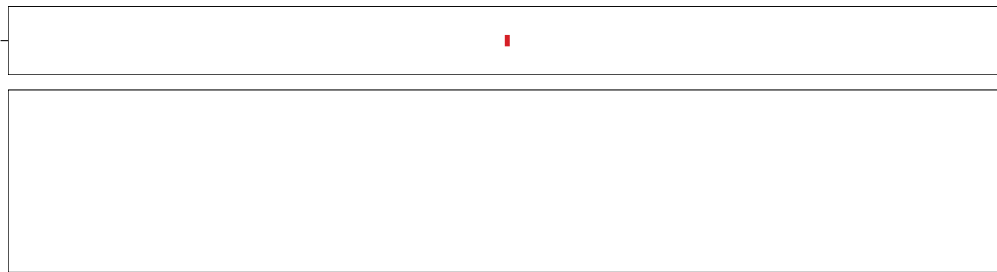
GeneTrack

strand • •

## chr3:58155524–58256025 Marker Peak Tracks for: L2.CUX2.MEIS2

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

L2.CUX2.MEIS2



Peaks

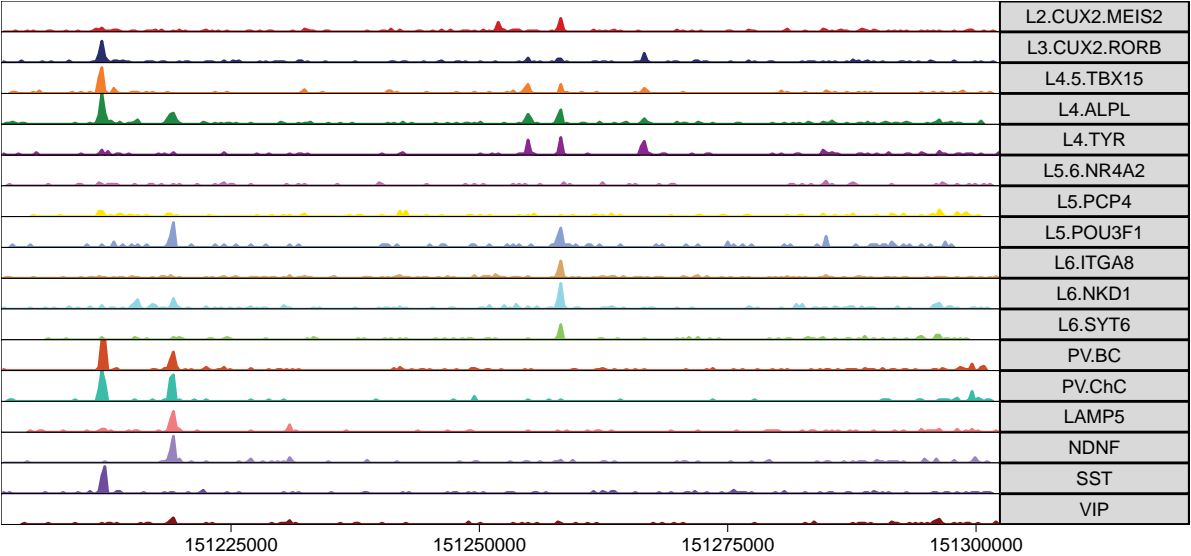
GeneTrack

strand ● ·

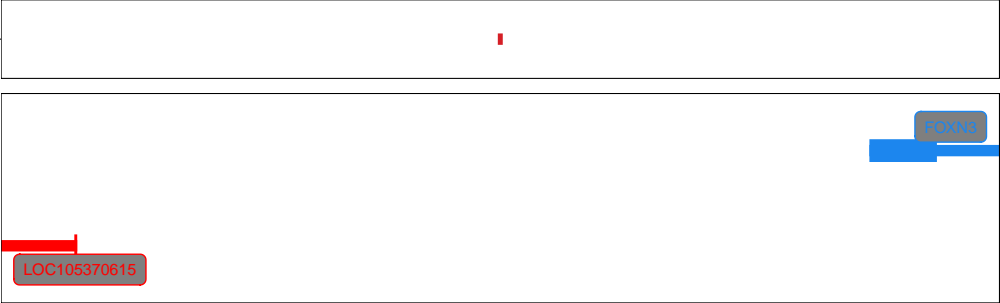


chr7:151201855–151302356 Marker Peak Tracks for: L2.CUX2.MEIS2

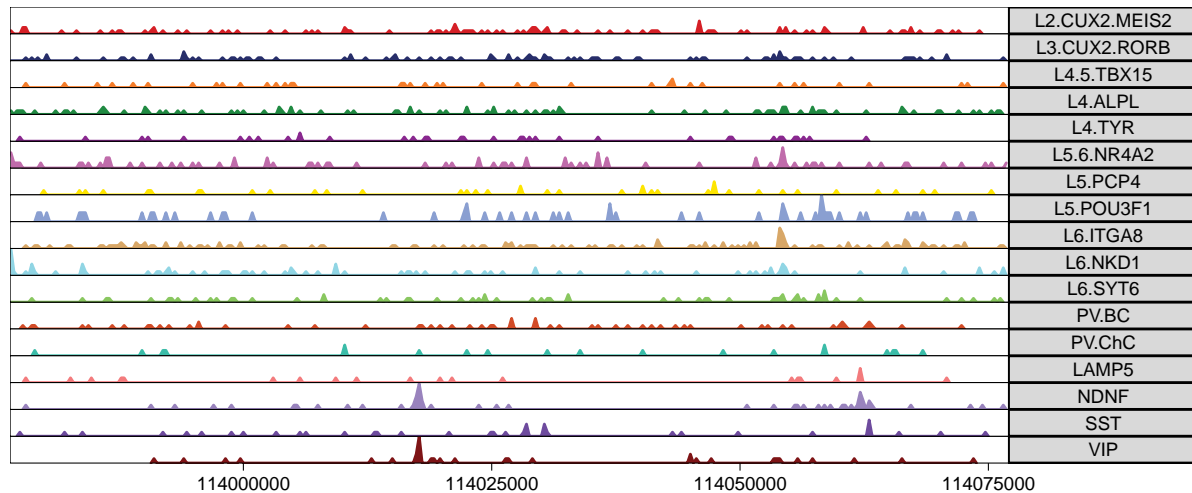
Coverage  
(Norm. ATAC Signal Range (0–0.82) by ReadsInTSS)



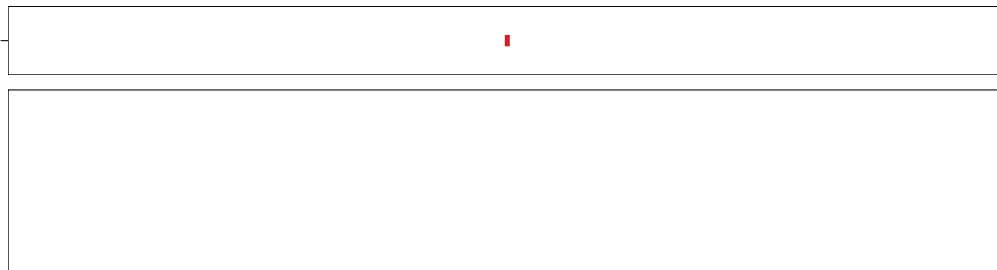
L2.CUX2.MEIS2



## chr5:113976519–114077020 Marker Peak Tracks for: L2.CUX2.MEIS2

Coverage  
(Norm. ATAC Signal Range (0–0.22) by ReadsInTSS)

L2.CUX2.MEIS2

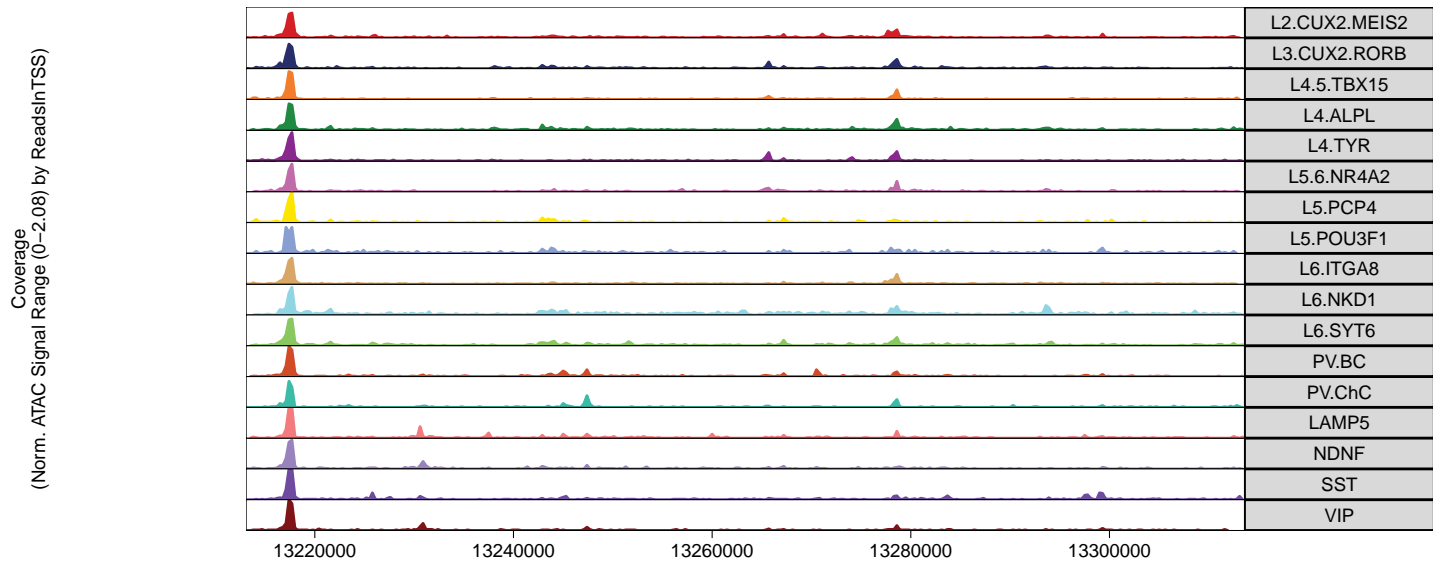


Peaks

GeneTrack

strand ● ·

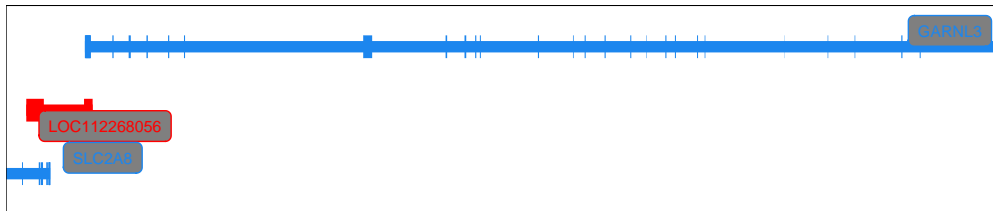
## chr15:13213086–13313587 Marker Peak Tracks for: L2.CUX2.MEIS2



L2.CUX2.MEIS2

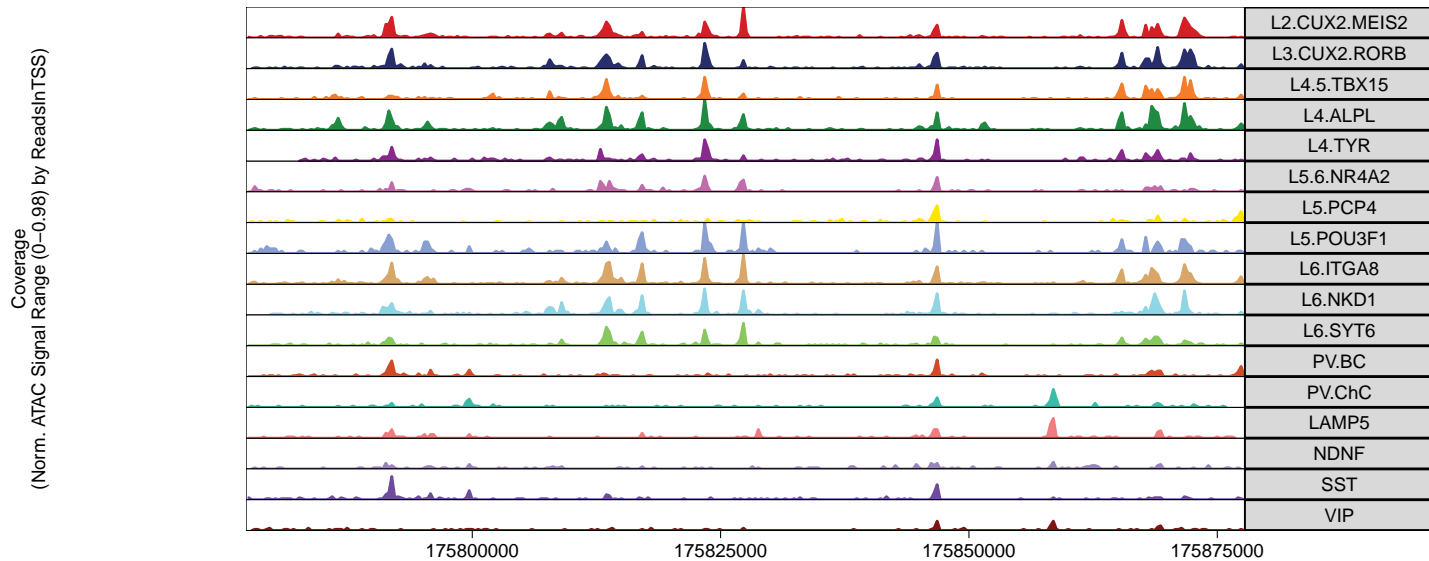


Peaks



Genes

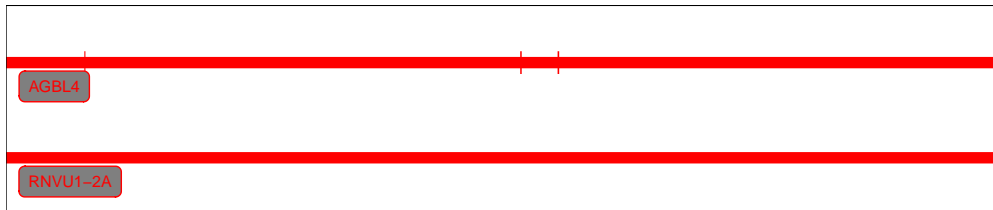
## chr1:175777234–175877735 Marker Peak Tracks for: L2.CUX2.MEIS2



L2.CUX2.MEIS2

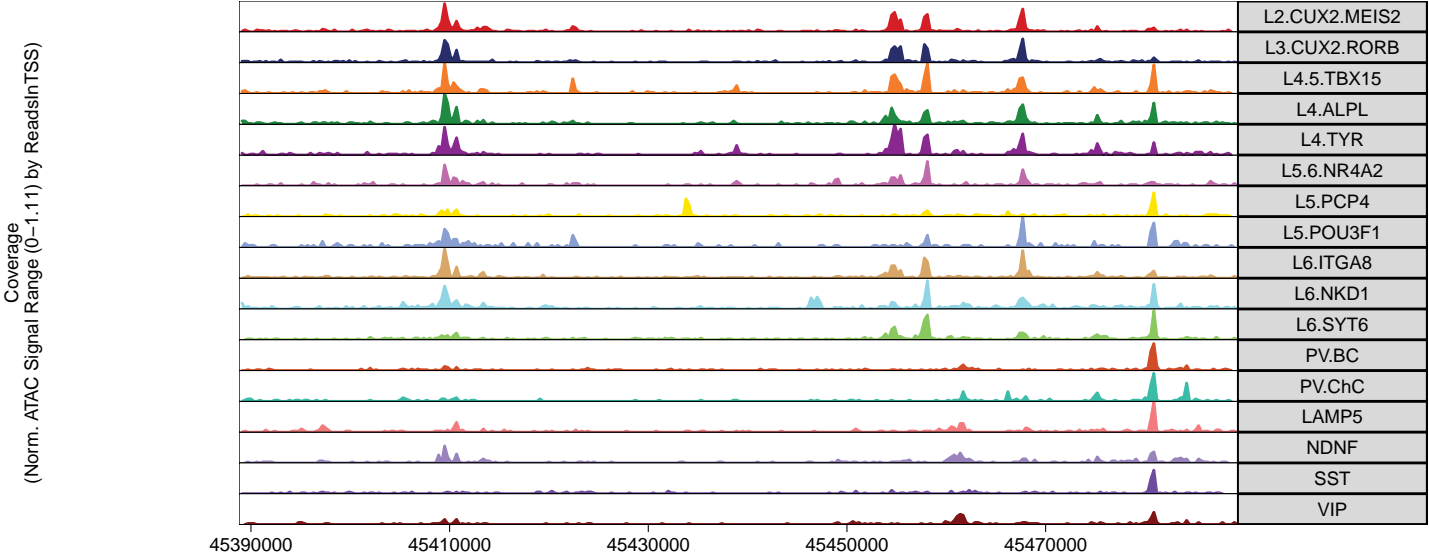


Peaks

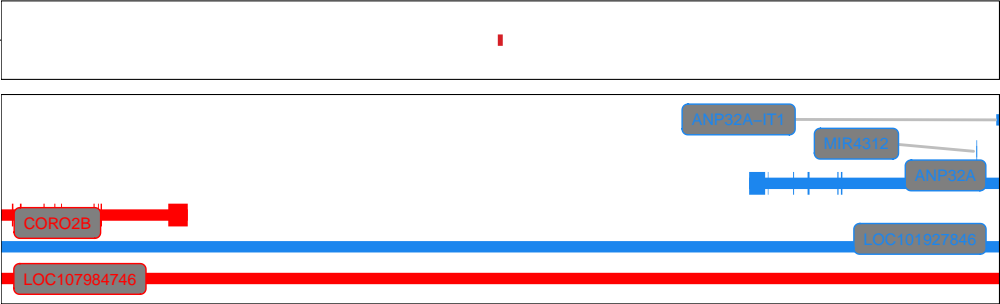


Genes

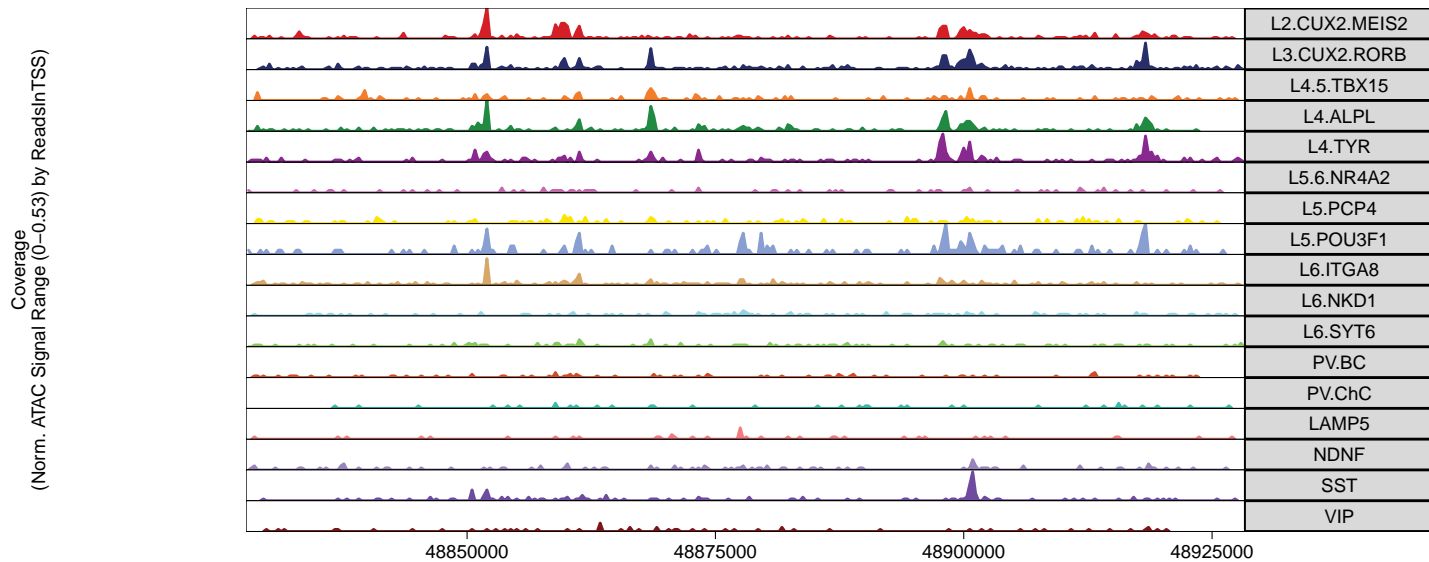
chr7:45388810–45489311 Marker Peak Tracks for: L2.CUX2.MEIS2



L2.CUX2.MEIS2



## chr20:48827757–48928258 Marker Peak Tracks for: L2.CUX2.MEIS2



L2.CUX2.MEIS2



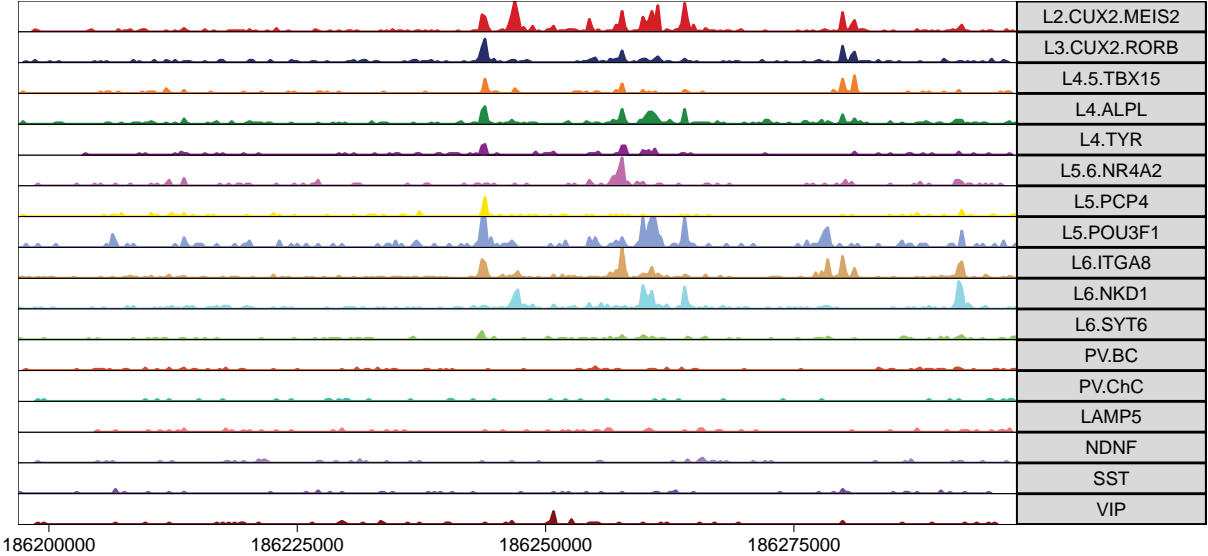
Peaks



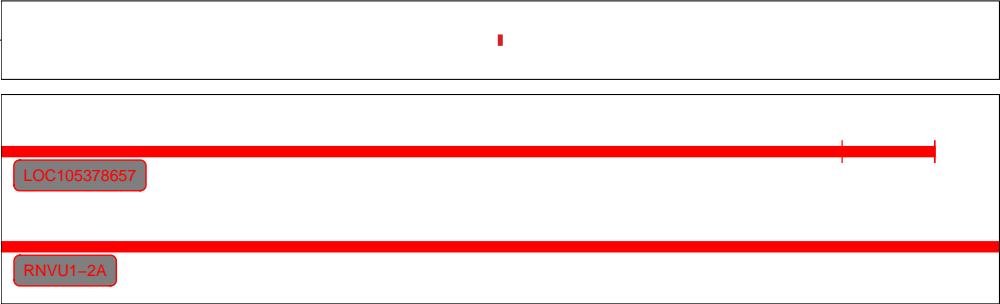
Genes

chr1:186196899–186297400 Marker Peak Tracks for: L2.CUX2.MEIS2

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



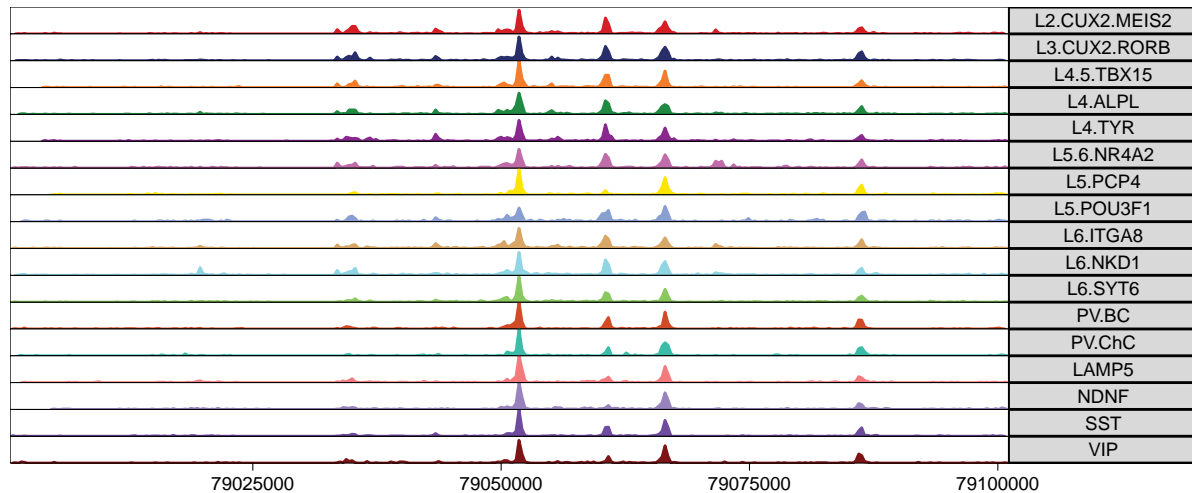
L2.CUX2.MEIS2



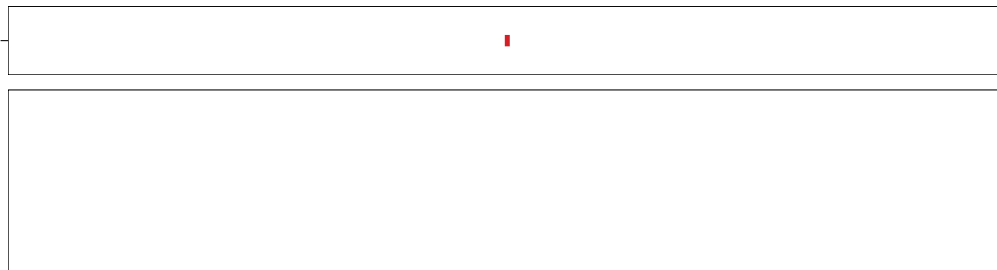
Peaks

Genes

## chr16:79000569–79101070 Marker Peak Tracks for: L2.CUX2.MEIS2

Coverage  
(Norm. ATAC Signal Range (0–2.44) by ReadsInTSS)

L2.CUX2.MEIS2

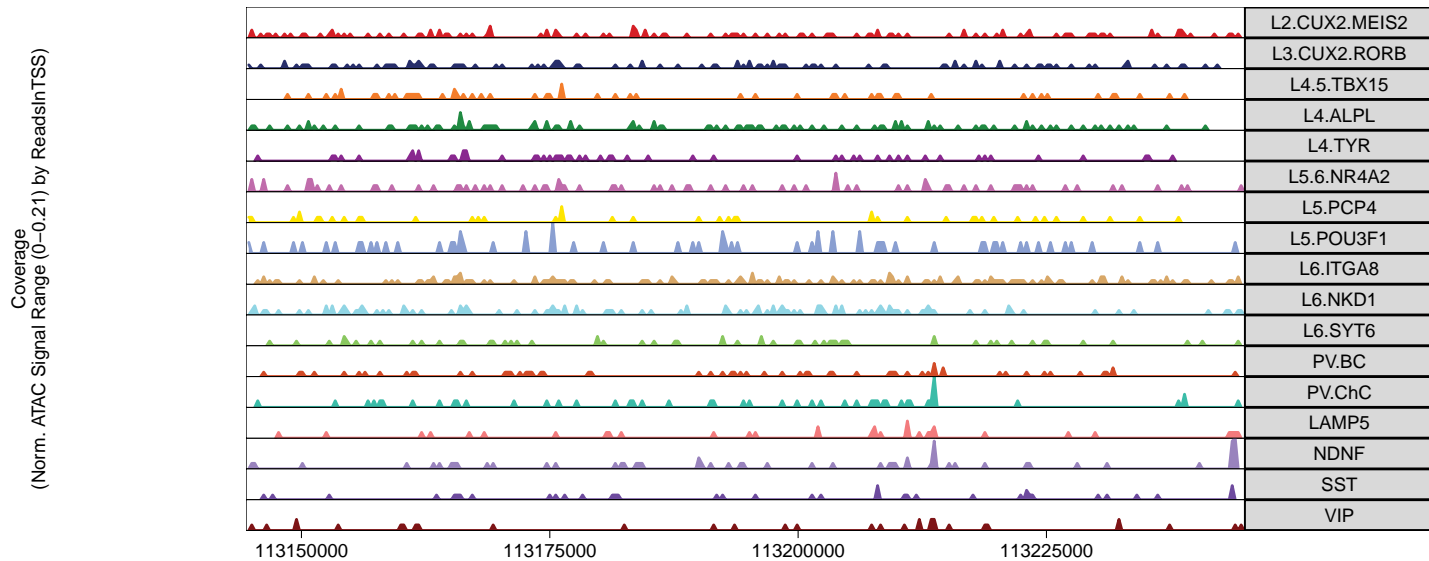


Peaks

strand ● ·



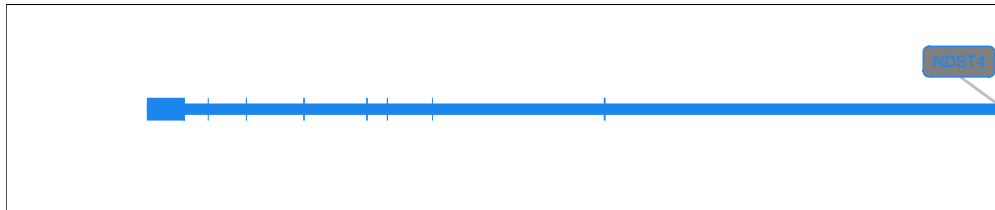
## chr5:113144430–113244931 Marker Peak Tracks for: L2.CUX2.MEIS2



L2.CUX2.MEIS2



Peaks



Genes