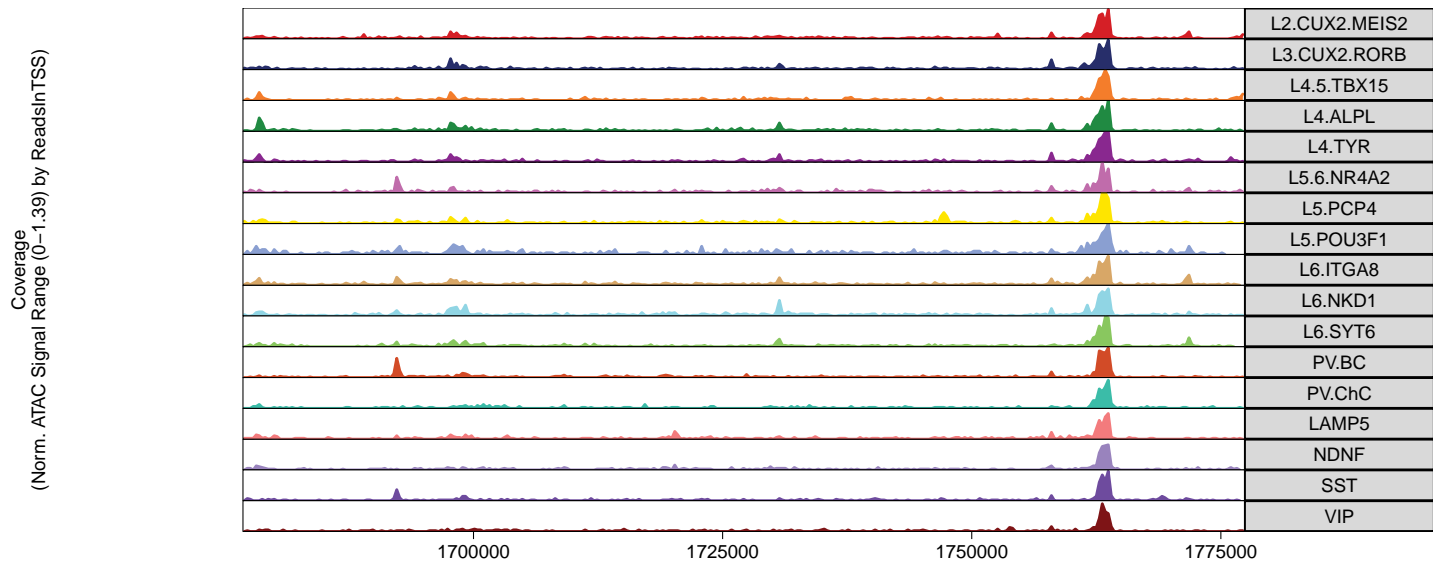


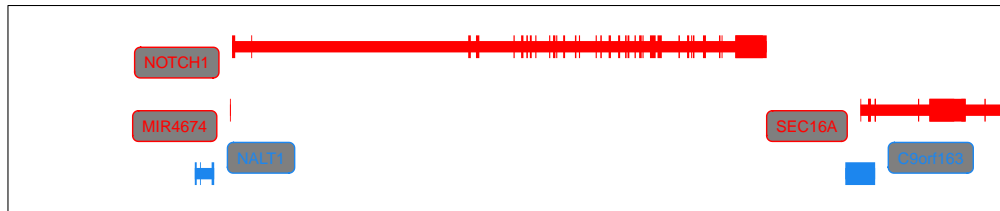
chr15:1676865–1777366 Marker Peak Tracks for: L3.CUX2.RORB



L3.CUX2.RORB

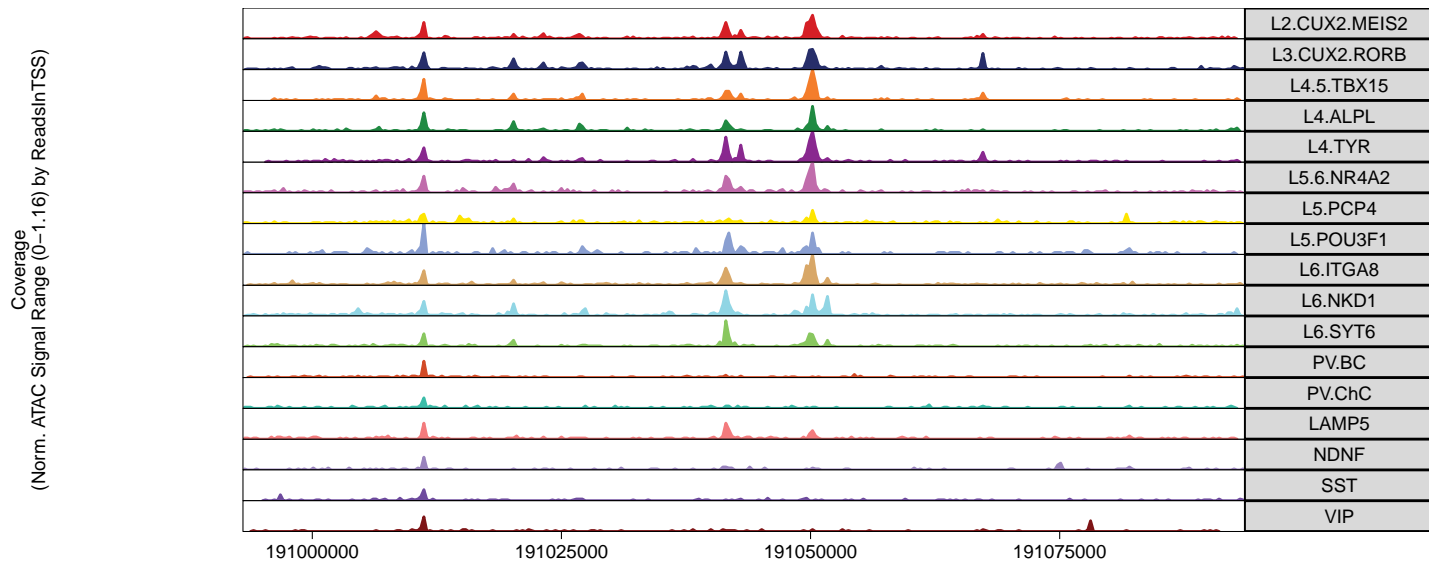


Peaks



Genes

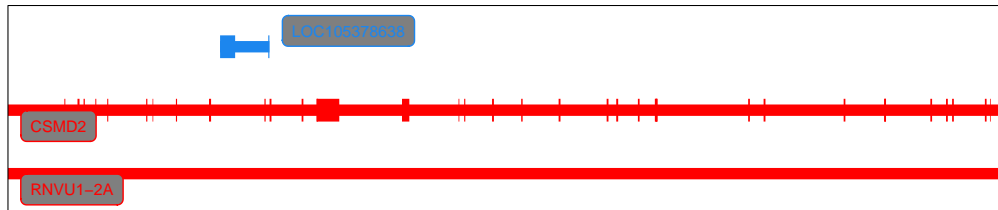
chr1:190993041–191093542 Marker Peak Tracks for: L3.CUX2.RORB



L3.CUX2.RORB

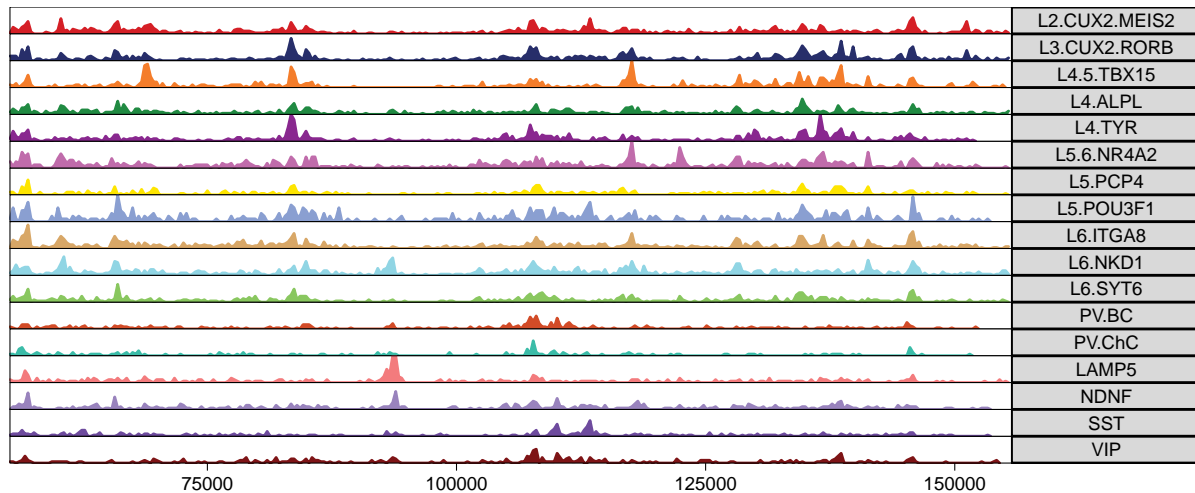


Peaks

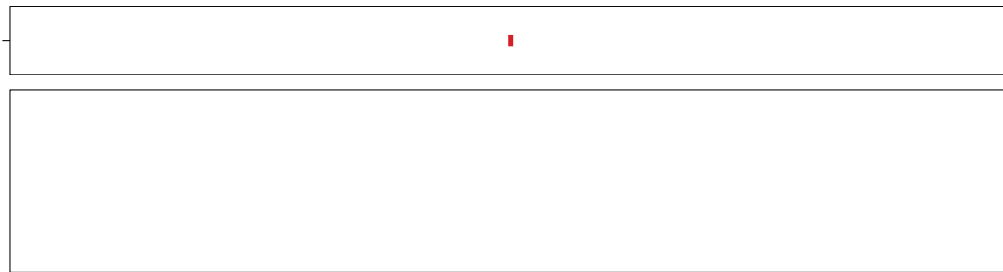


Genes

chr15:55188–155689 Marker Peak Tracks for: L3.CUX2.RORB

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

L3.CUX2.RORB

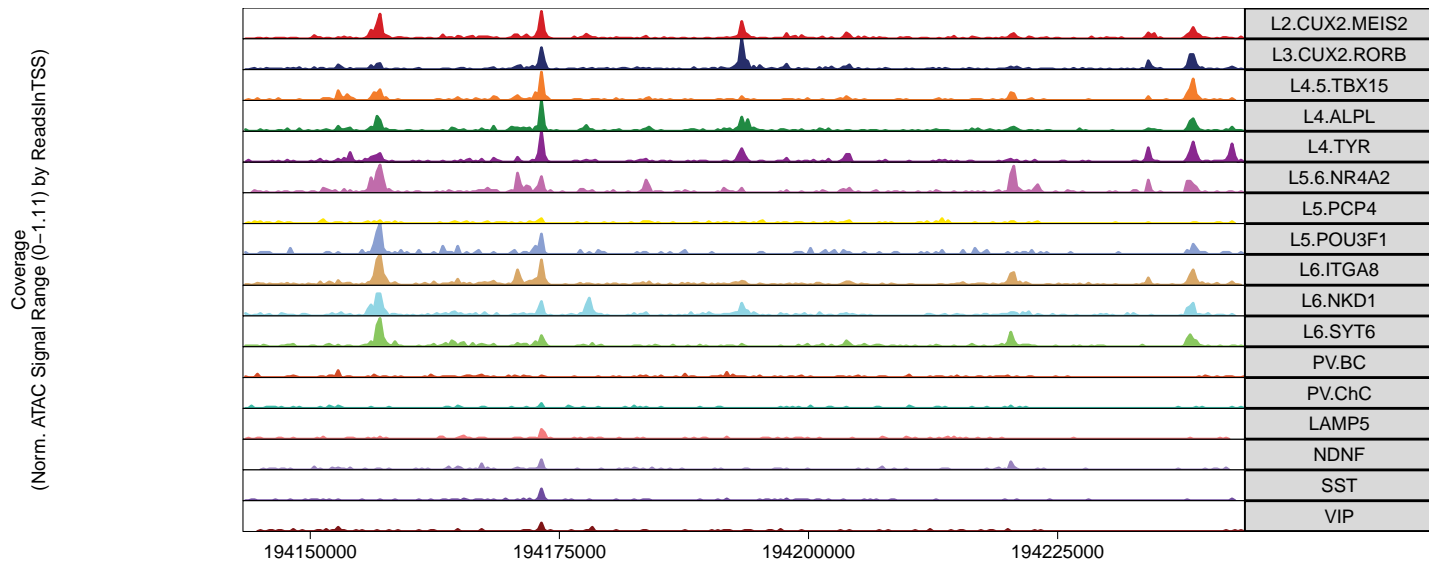


Peaks

GeneTrack

strand • •

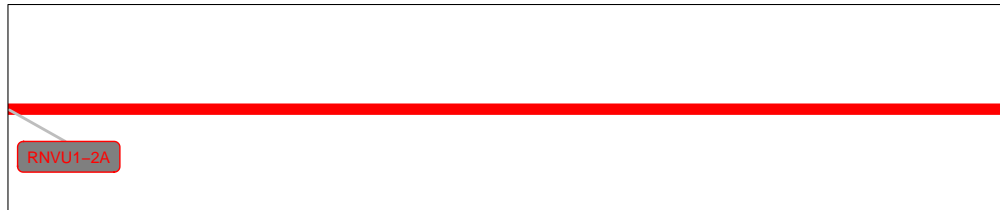
chr1:194143245–194243746 Marker Peak Tracks for: L3.CUX2.RORB



L3.CUX2.RORB

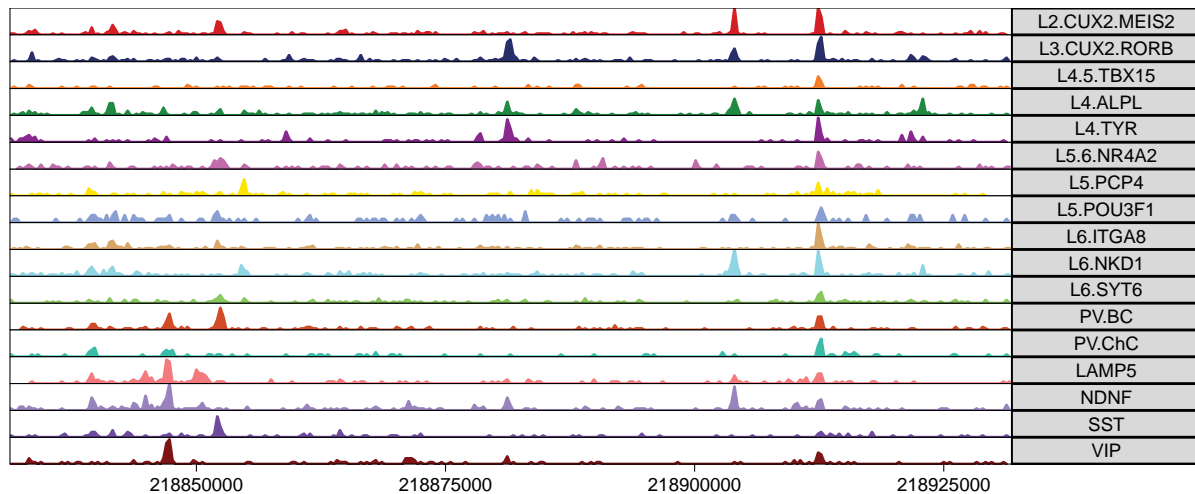


Peaks

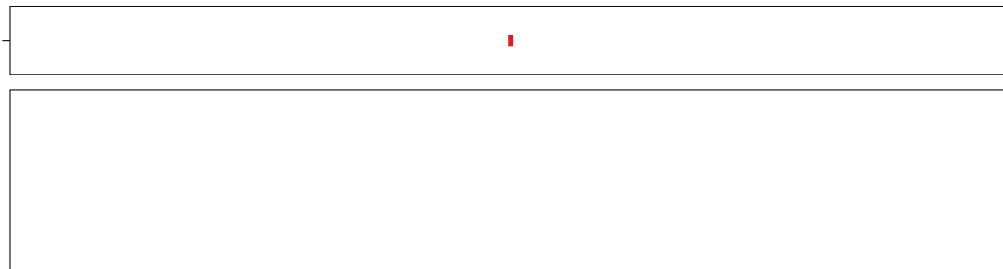


Genes

chr1:218831291–218931792 Marker Peak Tracks for: L3.CUX2.RORB

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

L3.CUX2.RORB

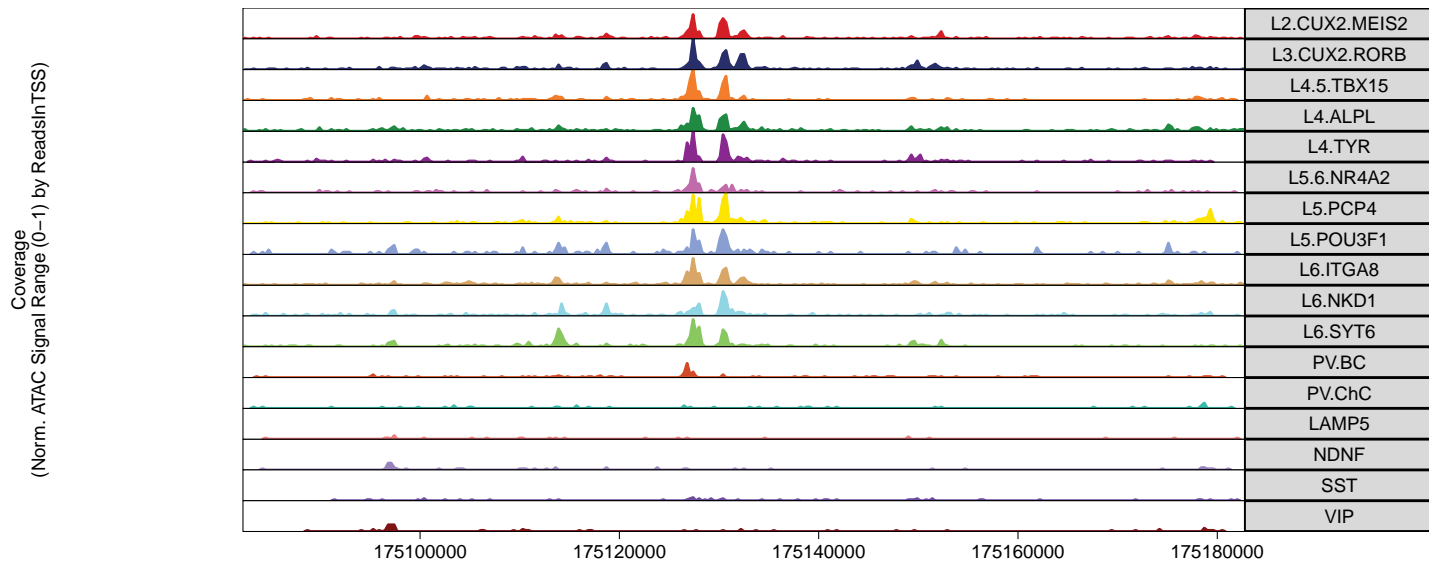


Peaks

GeneTrack

strand • •

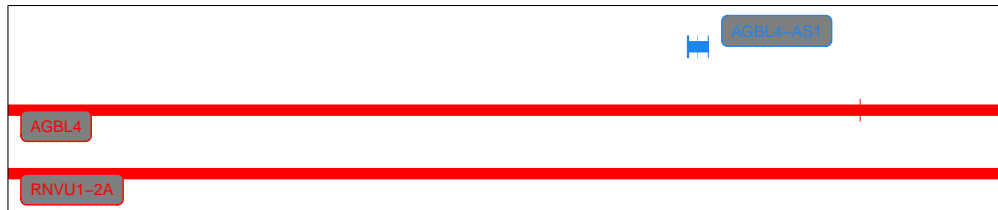
chr1:175082226–175182727 Marker Peak Tracks for: L3.CUX2.RORB



L3.CUX2.RORB

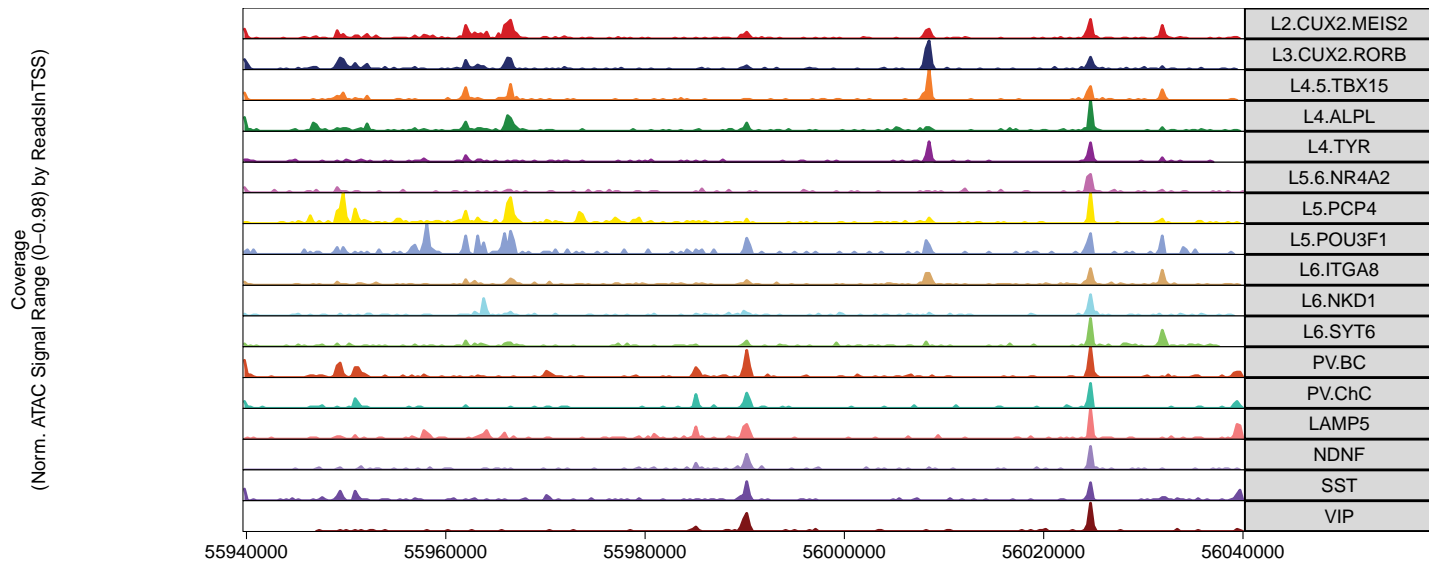


Peaks



Genes

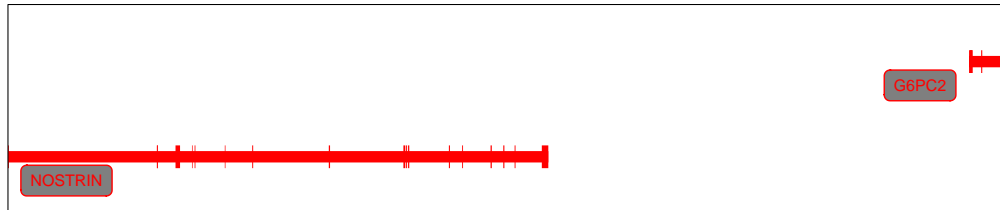
chr12:55939645–56040146 Marker Peak Tracks for: L3.CUX2.RORB



L3.CUX2.RORB



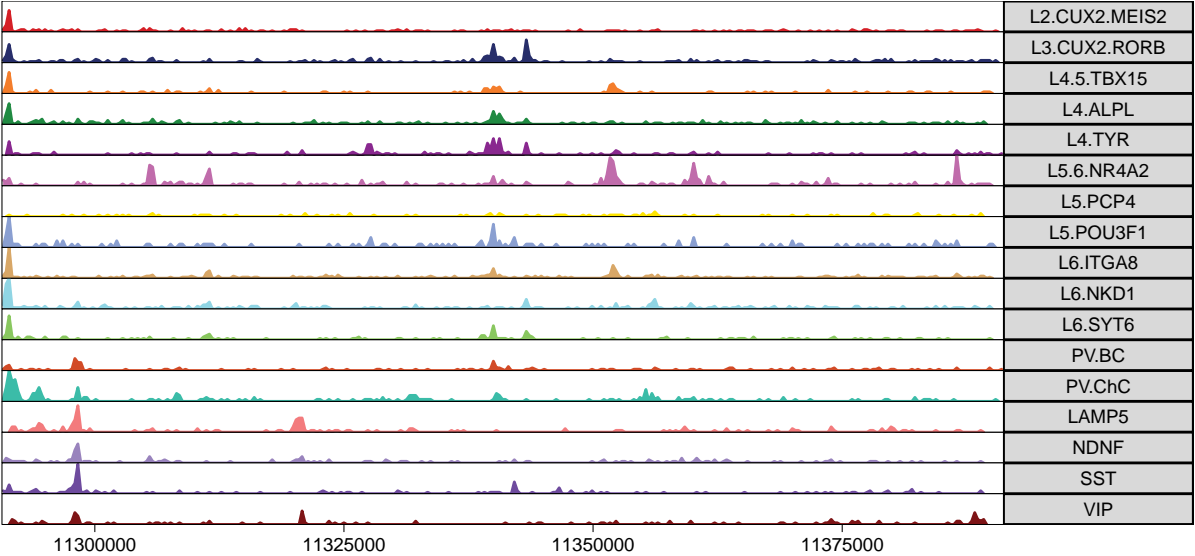
Peaks



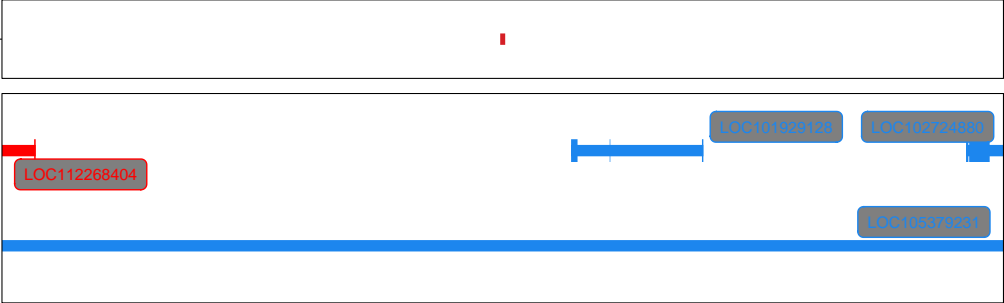
Genes

chr8:11290681–11391182 Marker Peak Tracks for: L3.CUX2.RORB

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



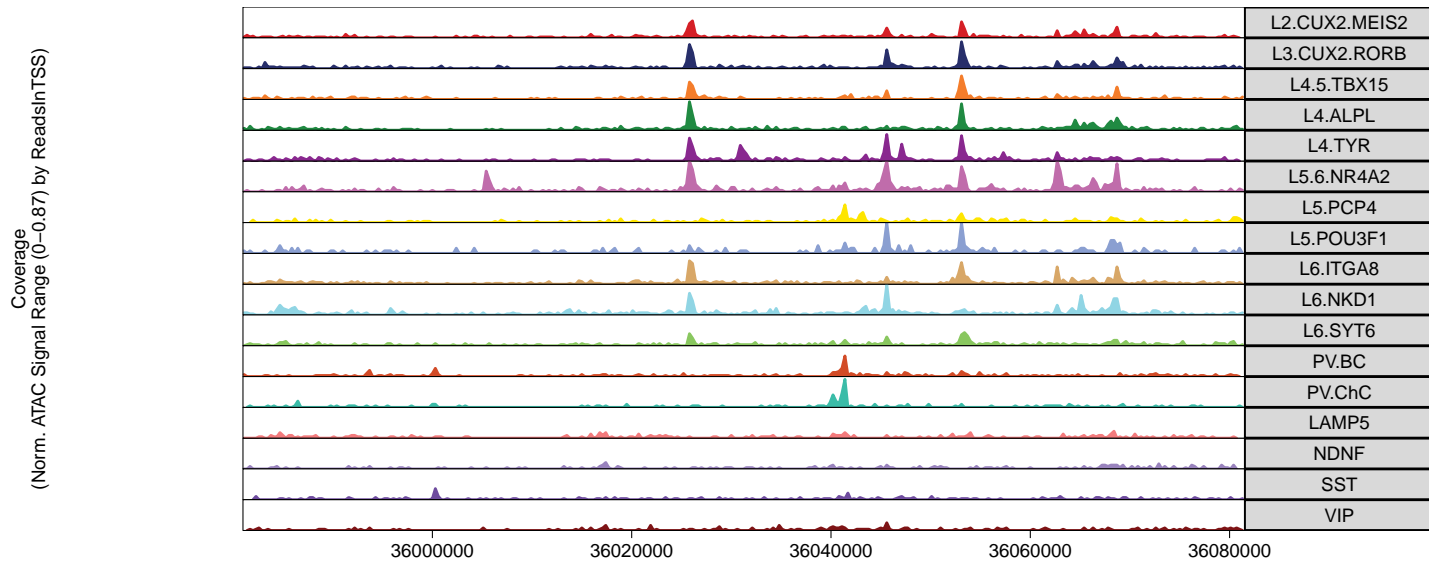
L3.CUX2.RORB



Peaks

Genes

chr18:35980990–36081491 Marker Peak Tracks for: L3.CUX2.RORB



L3.CUX2.RORB

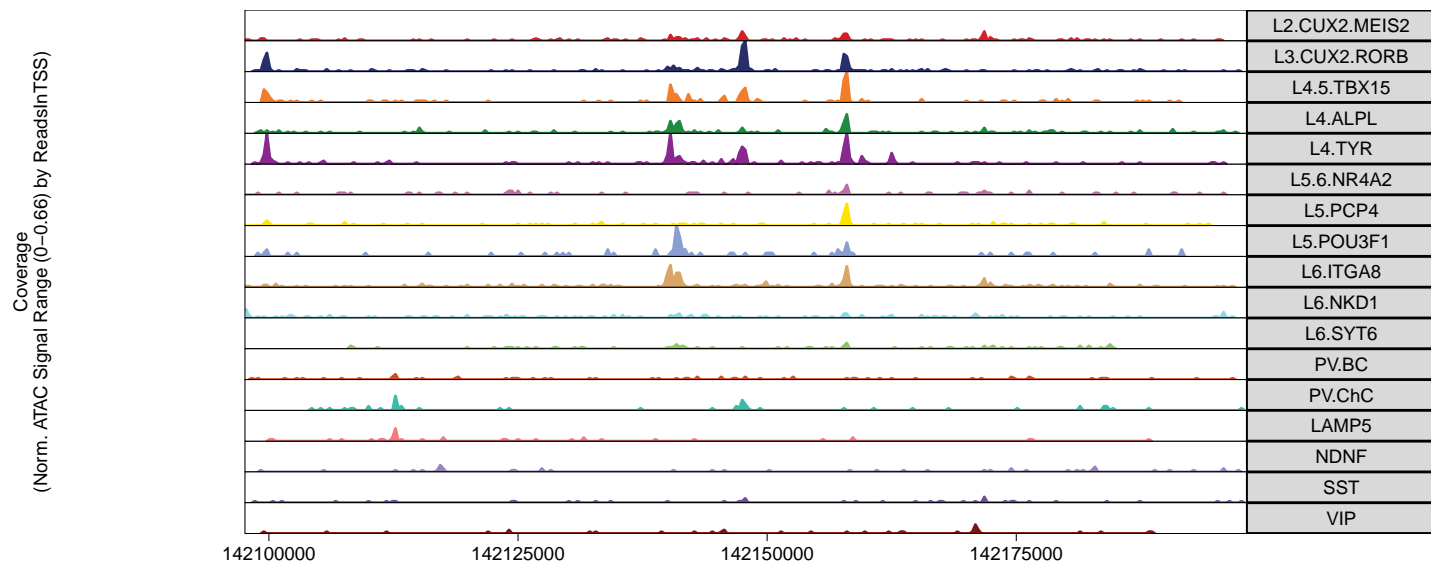


Peaks



Genes

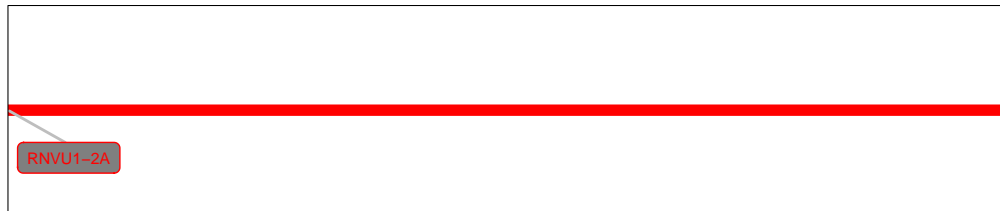
chr1:142097597–142198098 Marker Peak Tracks for: L3.CUX2.RORB



L3.CUX2.RORB

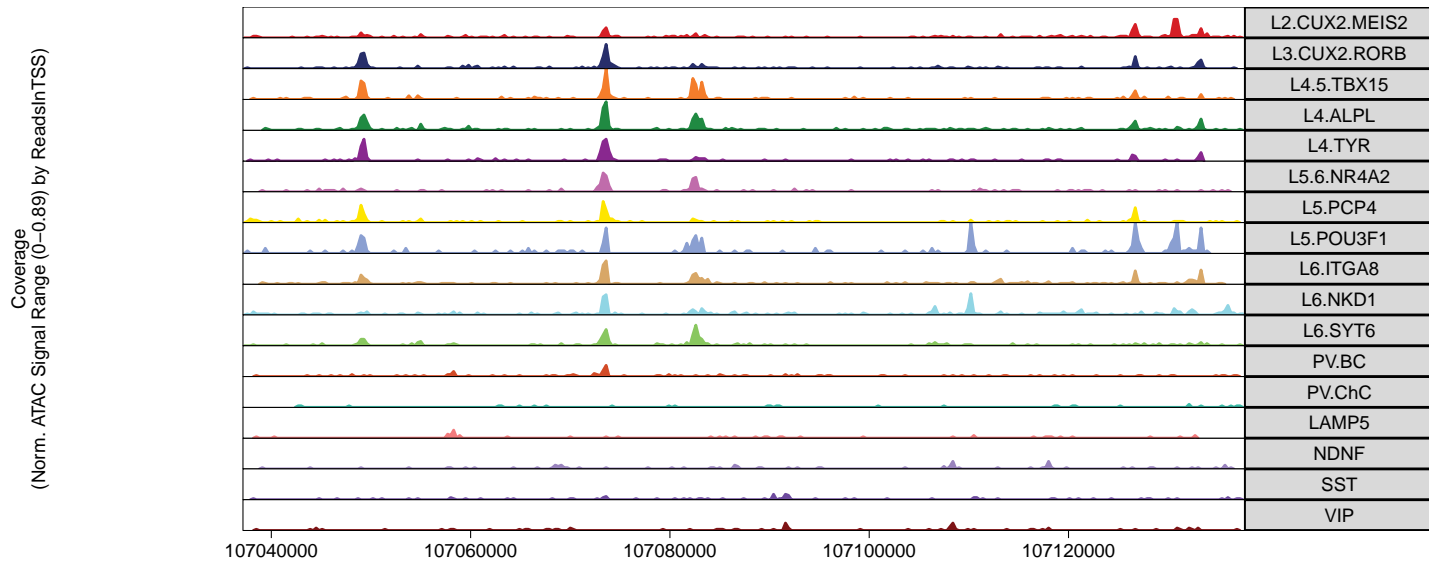


Peaks



Genes

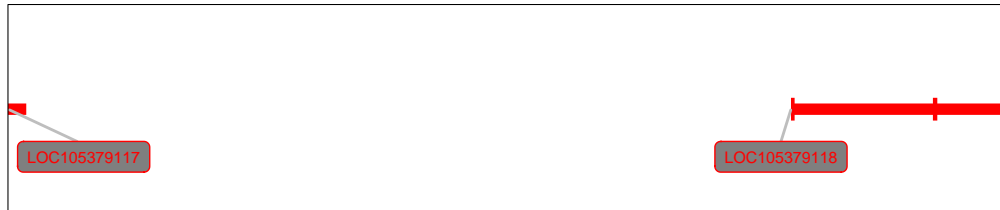
chr6:107037154–107137655 Marker Peak Tracks for: L3.CUX2.RORB



L3.CUX2.RORB

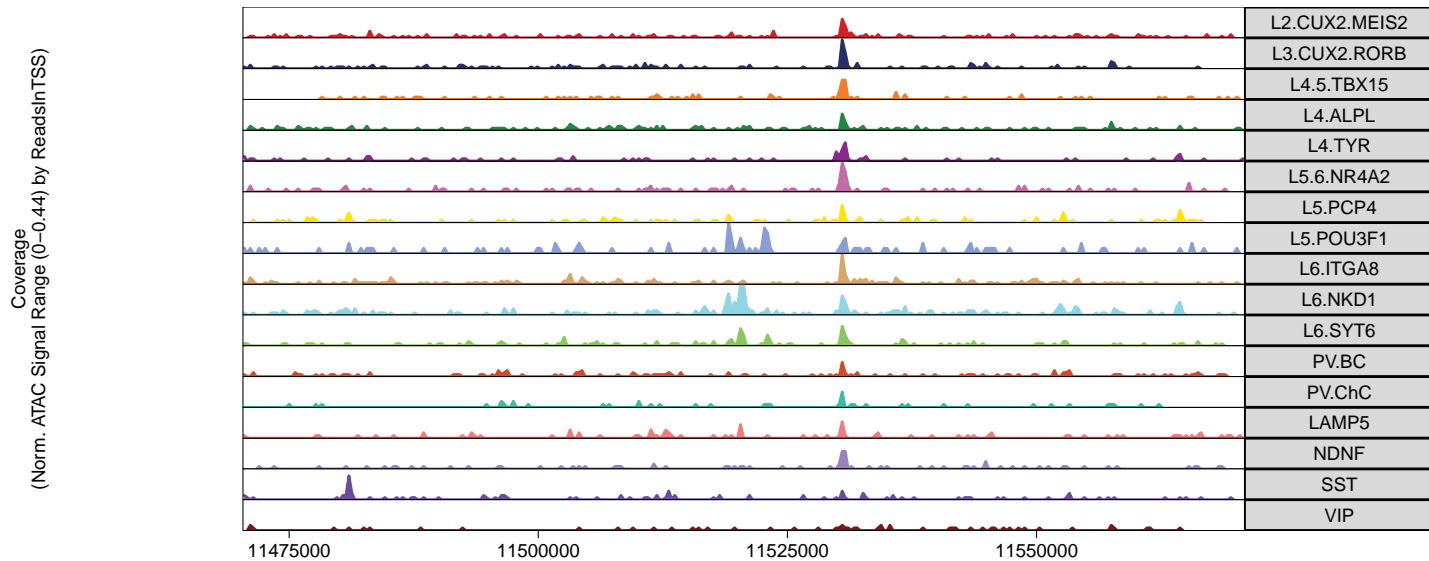


Peaks

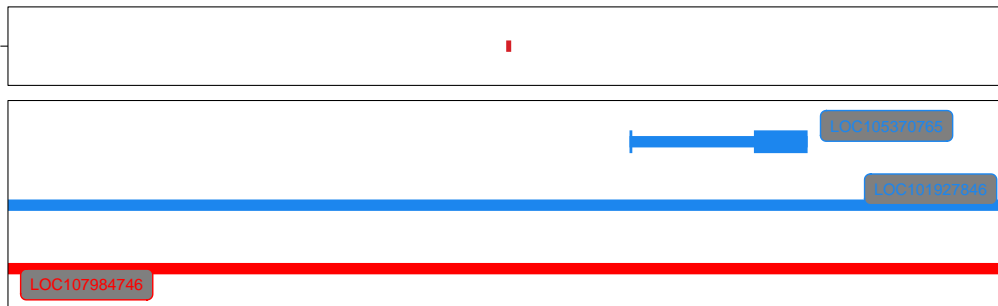


Genes

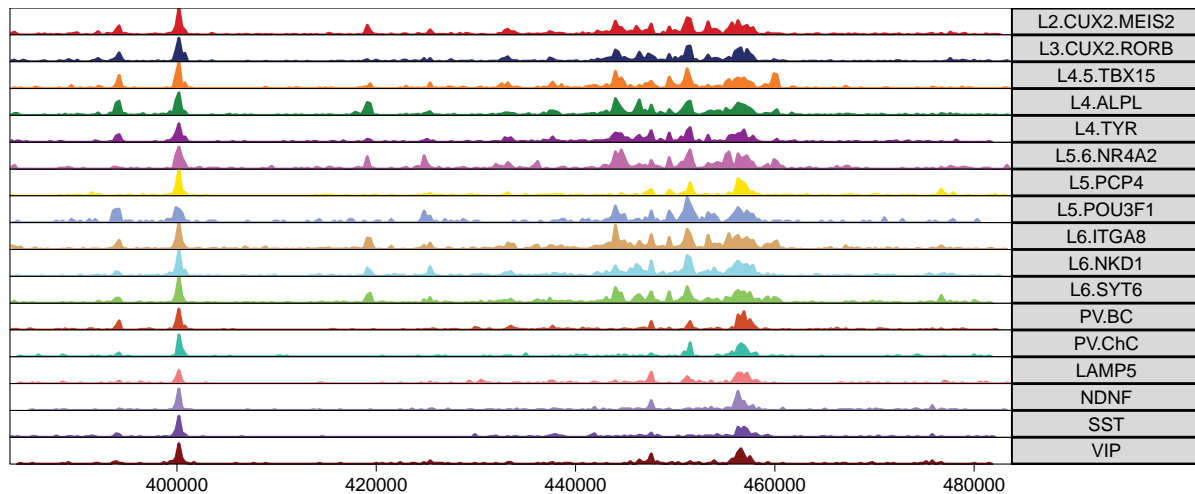
chr7:11470364–11570865 Marker Peak Tracks for: L3.CUX2.RORB



L3.CUX2.RORB



chr17:383242–483743 Marker Peak Tracks for: L3.CUX2.RORB

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

L3.CUX2.RORB

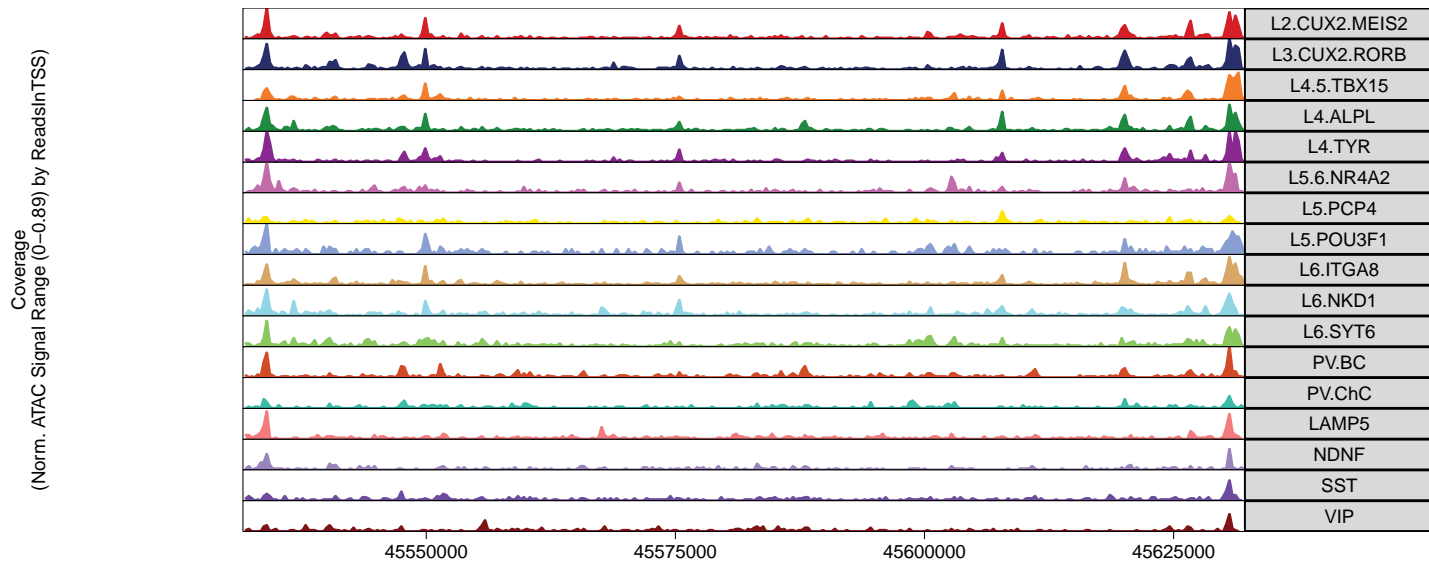


Peaks

GeneTrack

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chr14:45531605–45632106 Marker Peak Tracks for: L3.CUX2.RORB



L3.CUX2.RORB

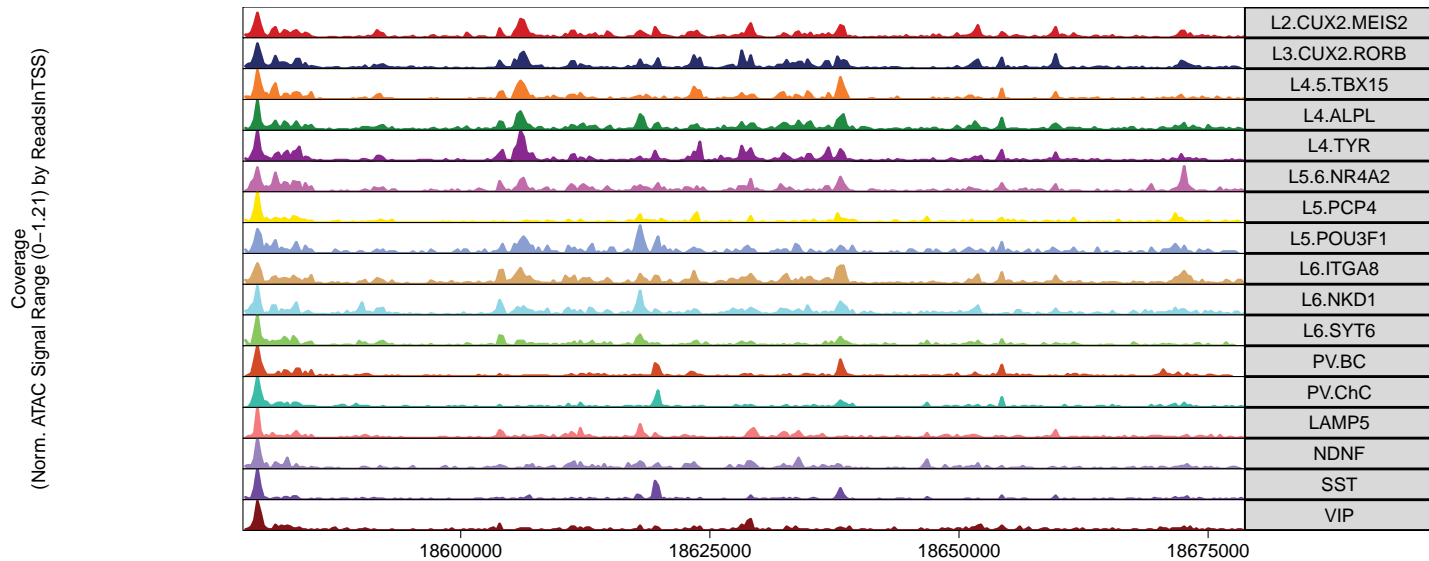


Peaks



Genes

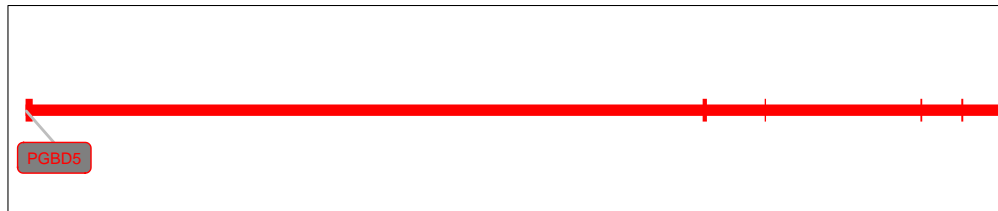
chr1:18578145–18678646 Marker Peak Tracks for: L3.CUX2.RORB



L3.CUX2.RORB

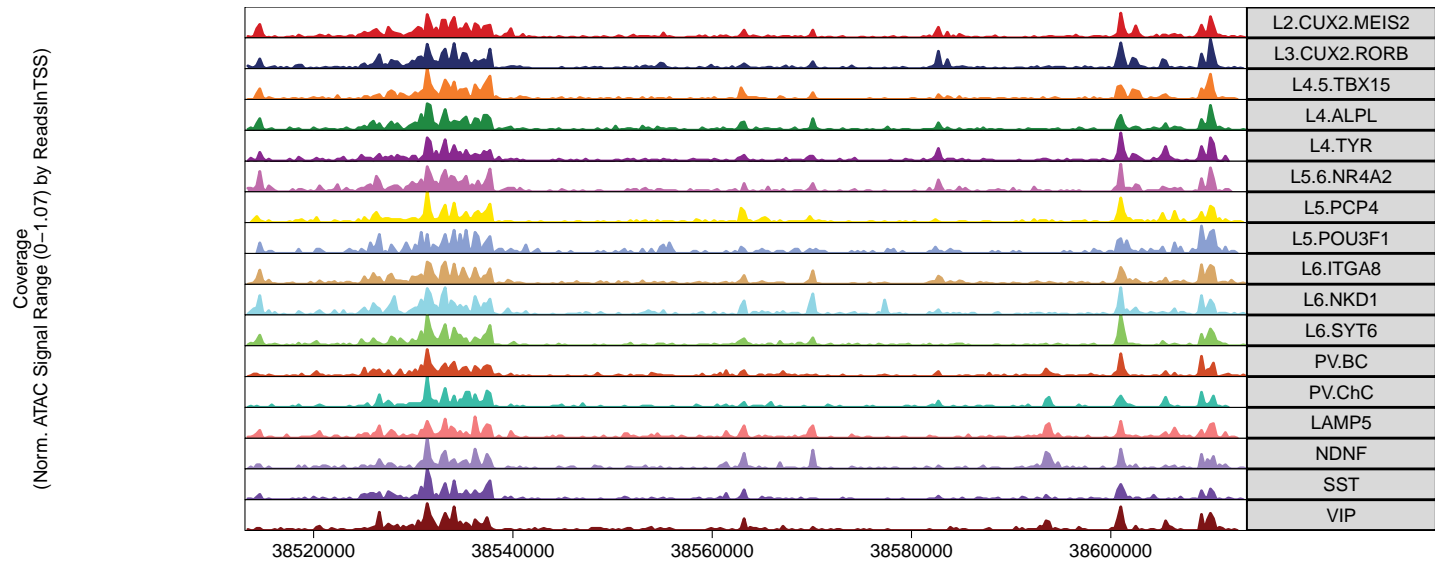


Peaks



Genes

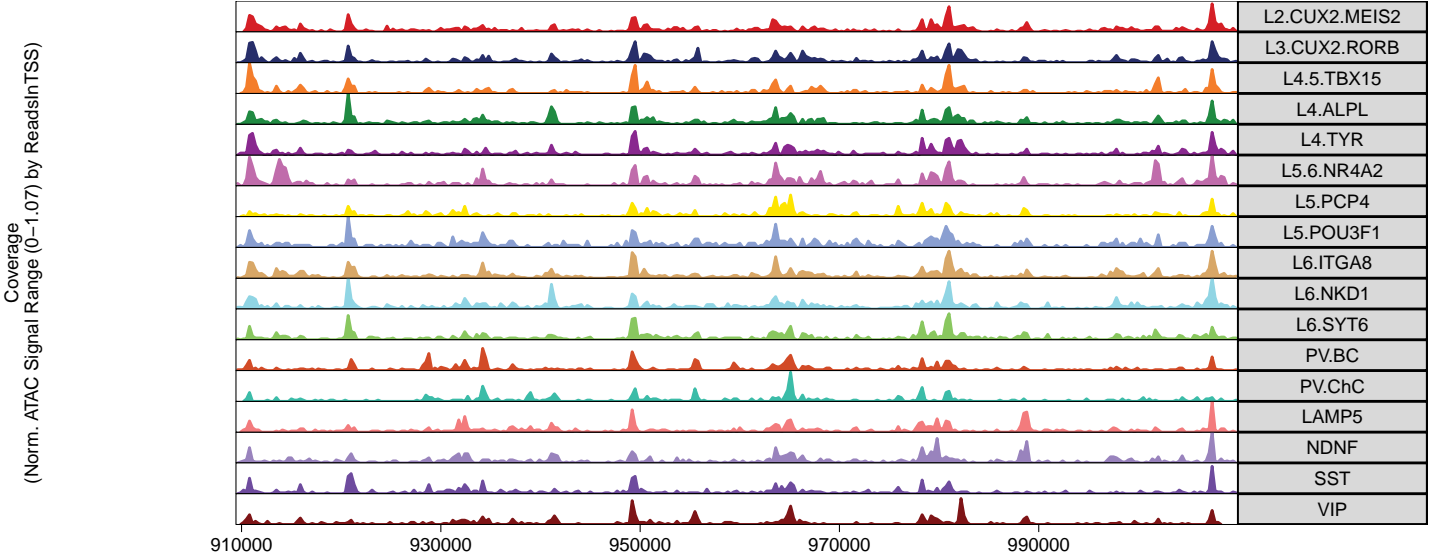
chr3:38513110–38613611 Marker Peak Tracks for: L3.CUX2.RORB



L3.CUX2.RORB



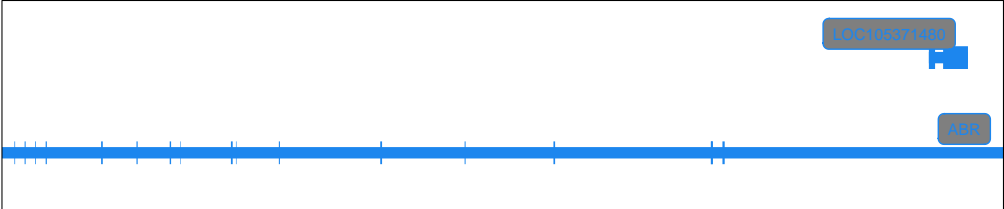
chr16:909439–1009940 Marker Peak Tracks for: L3.CUX2.RORB



L3.CUX2.RORB

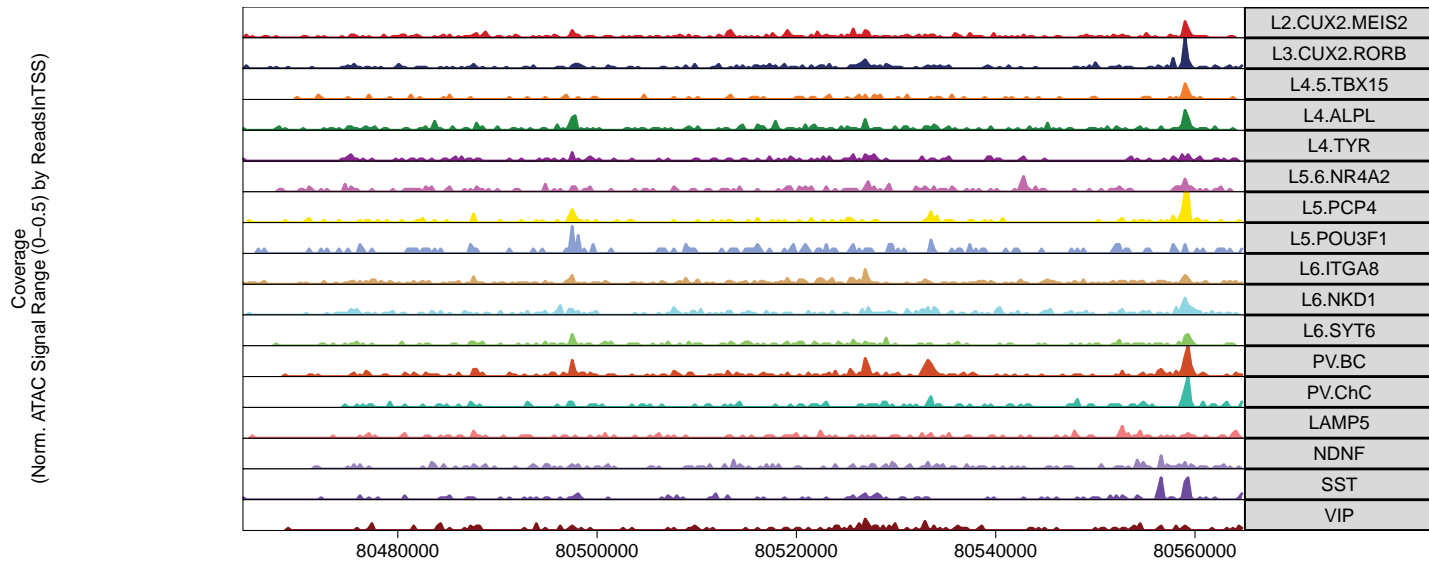


Peaks



Genes

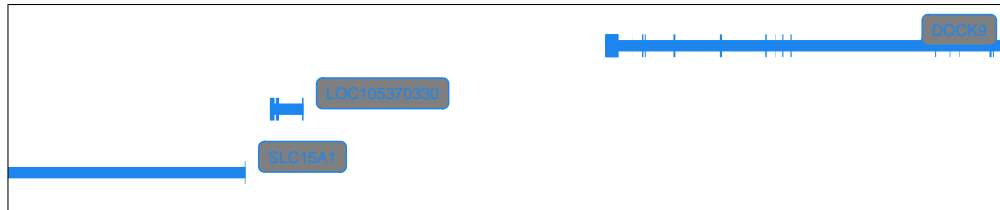
chr17:80464460–80564961 Marker Peak Tracks for: L3.CUX2.RORB



L3.CUX2.RORB

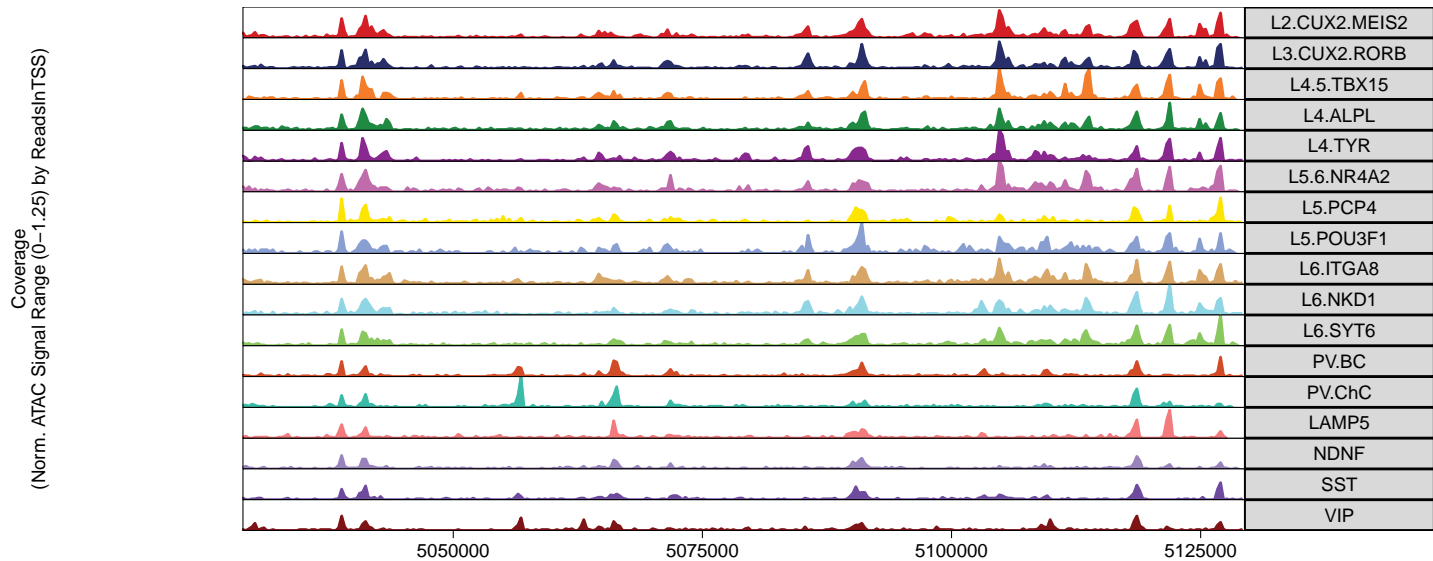


Peaks

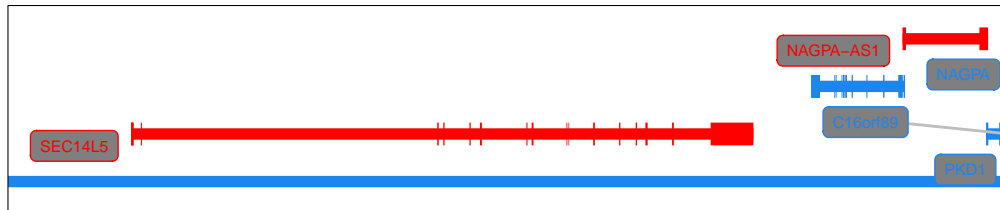


Genes

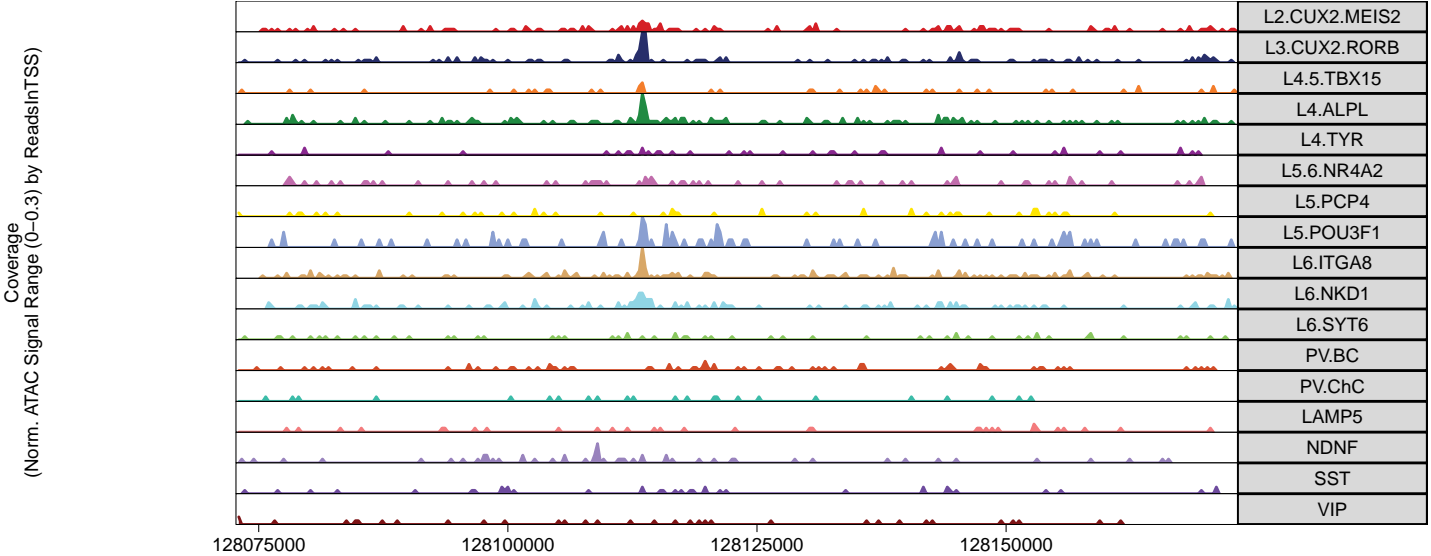
chr20:5028895–5129396 Marker Peak Tracks for: L3.CUX2.RORB



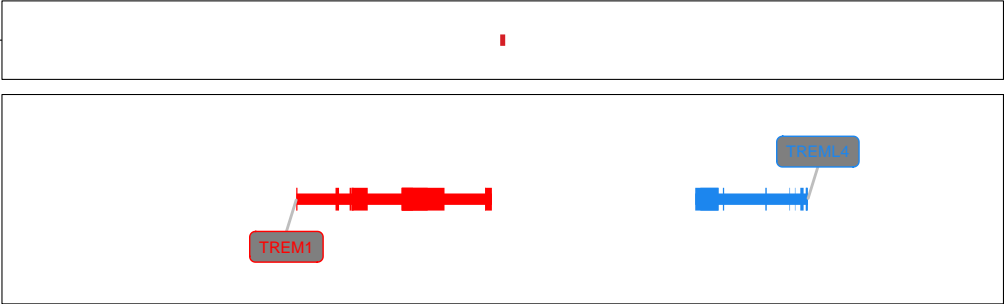
L3.CUX2.RORB



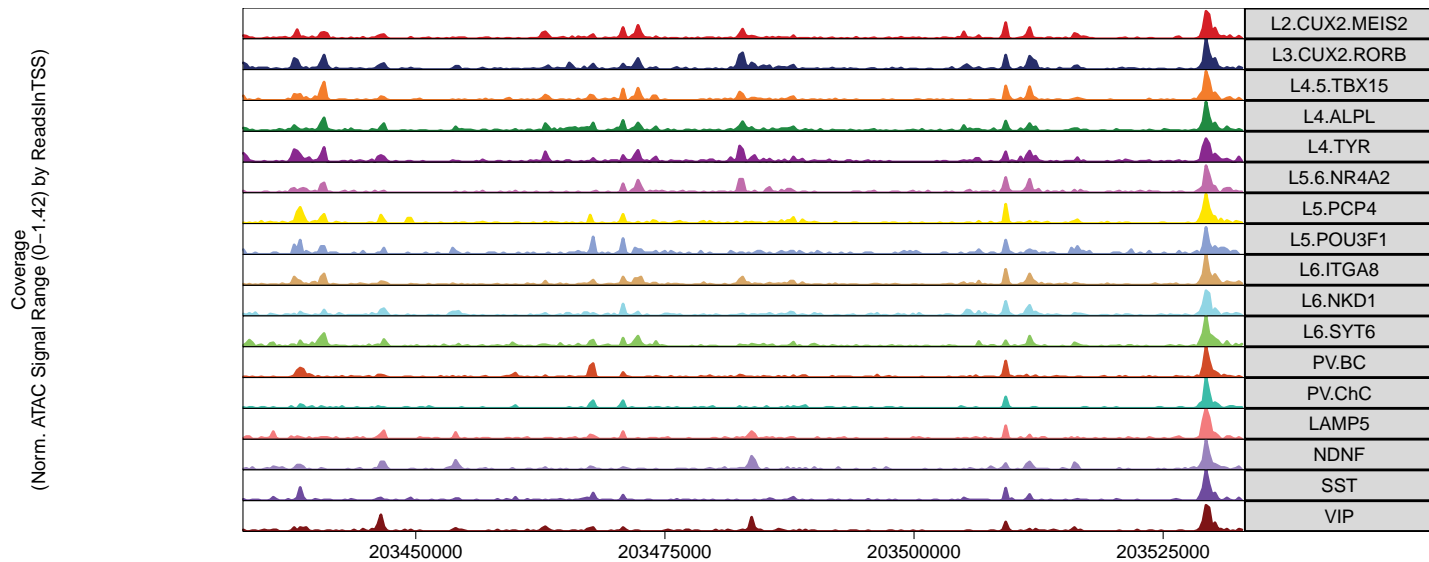
chr4:128072715–128173216 Marker Peak Tracks for: L3.CUX2.RORB



L3.CUX2.RORB



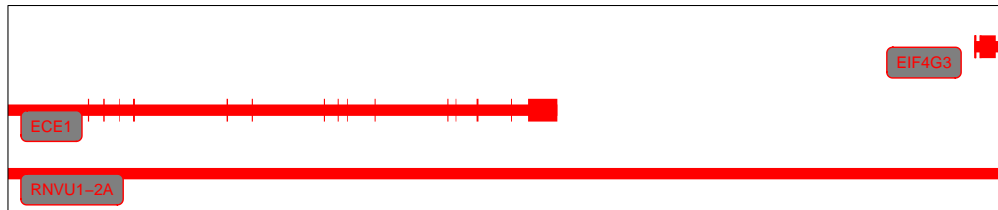
chr1:203432657–203533158 Marker Peak Tracks for: L3.CUX2.RORB



L3.CUX2.RORB



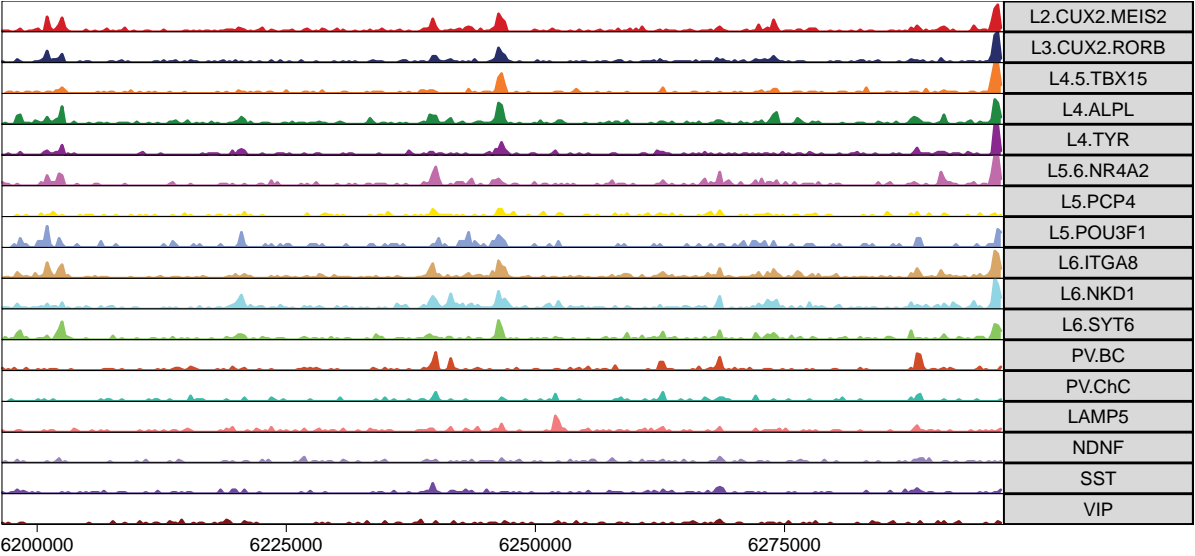
Peaks



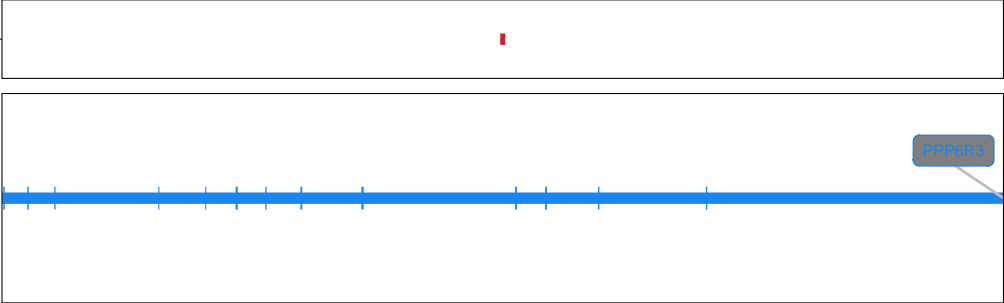
Genes

chr14:6196467–6296968 Marker Peak Tracks for: L3.CUX2.RORB

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



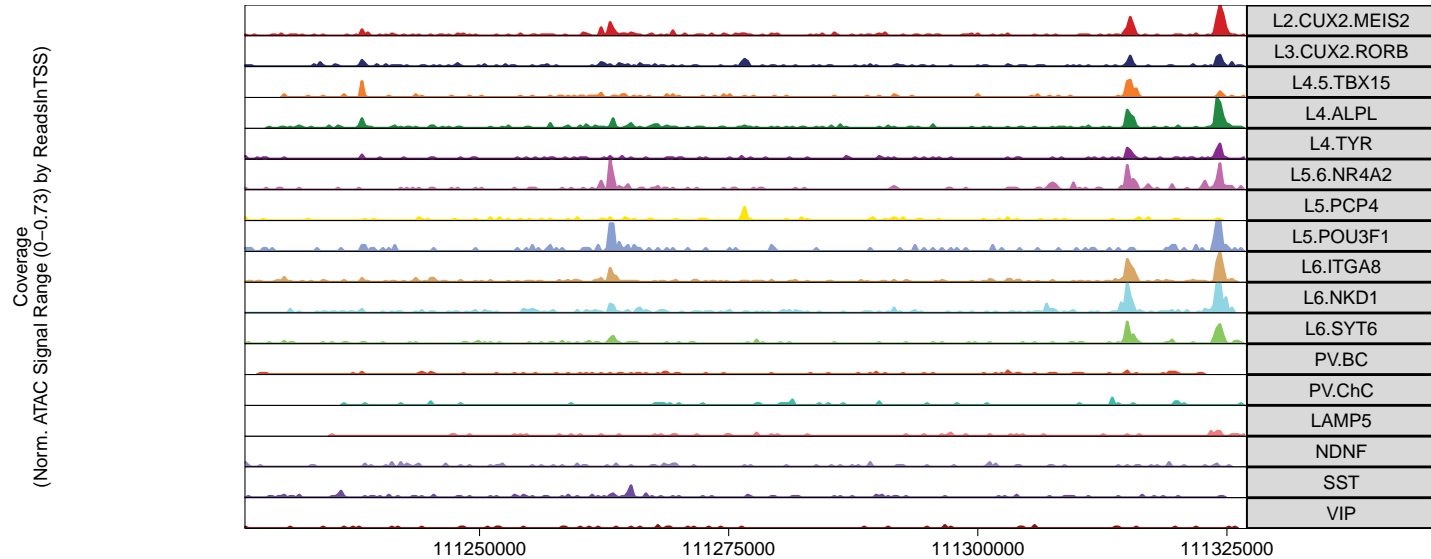
L3.CUX2.RORB



Peaks

Genes

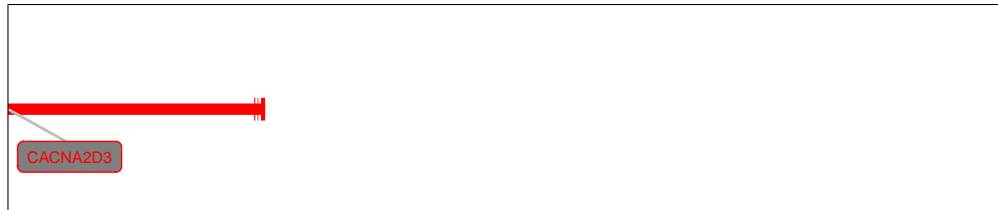
chr2:111226459–111326960 Marker Peak Tracks for: L3.CUX2.RORB



L3.CUX2.RORB

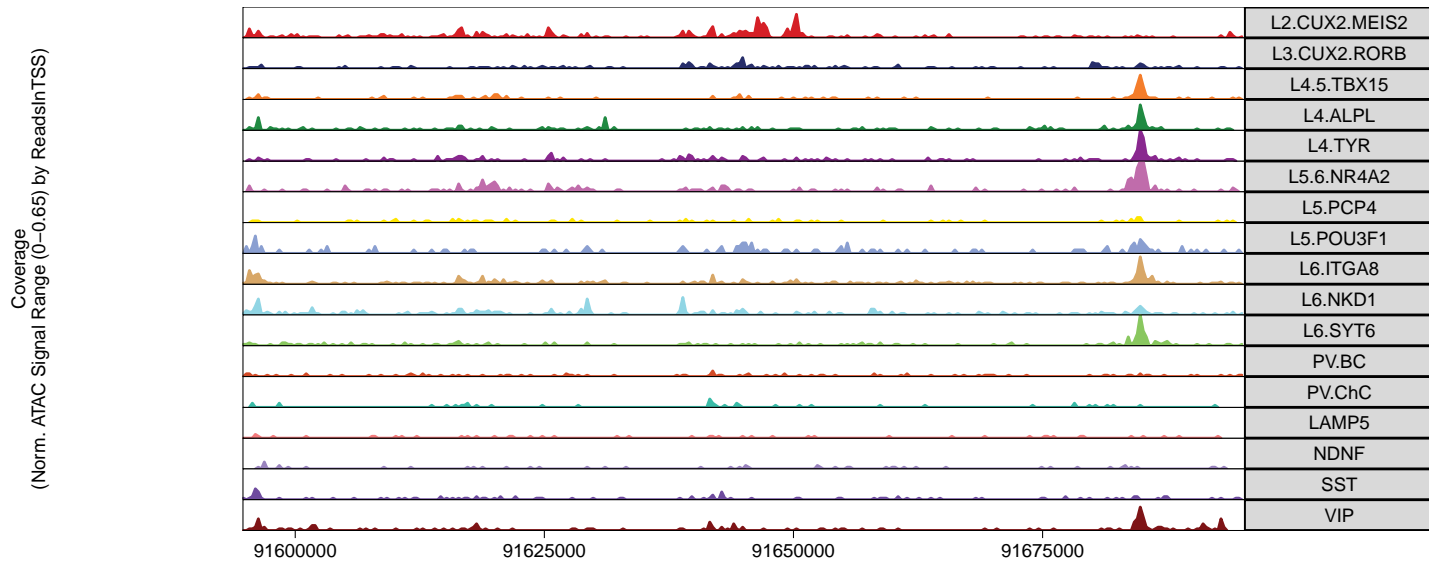


Peaks



Genes

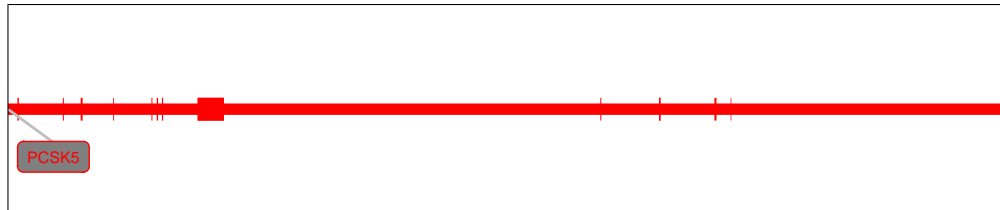
chr15:91594754–91695255 Marker Peak Tracks for: L3.CUX2.RORB



L3.CUX2.RORB



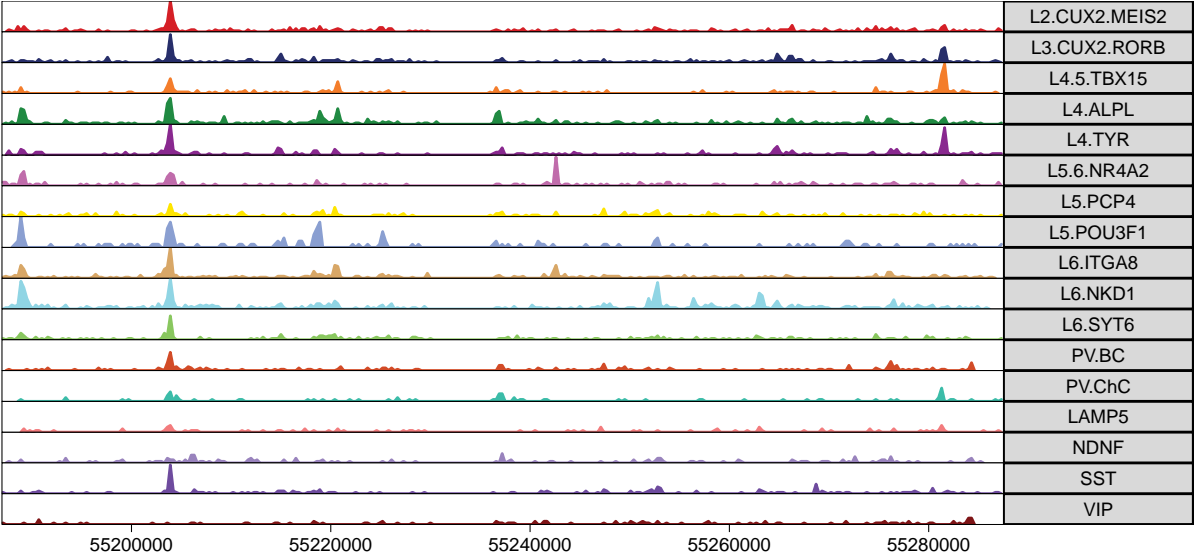
Peaks



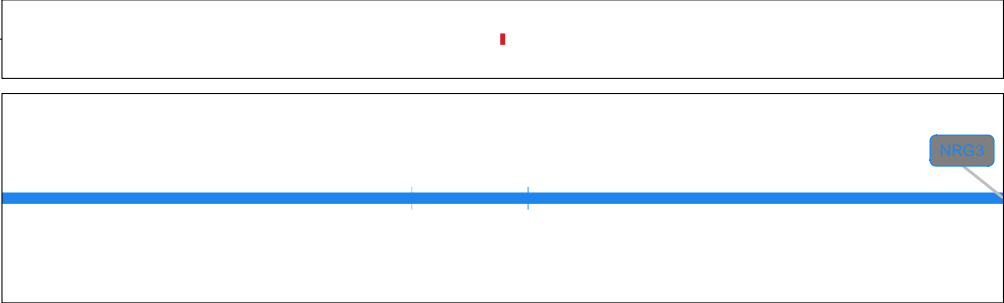
Genes

chr9:55187002–55287503 Marker Peak Tracks for: L3.CUX2.RORB

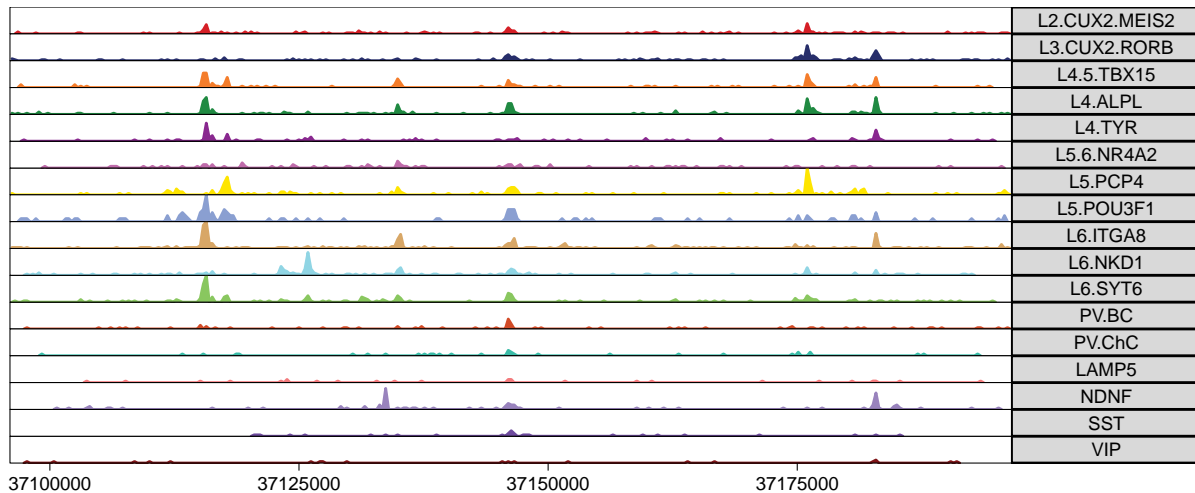
Coverage
(Norm. ATAC Signal Range (0–0.71) by ReadsInTSS)



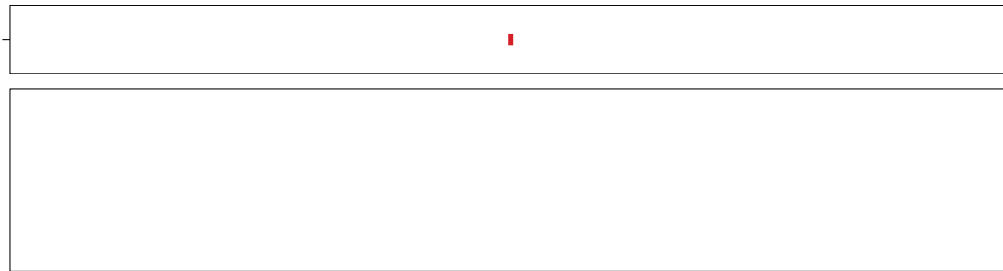
L3.CUX2.RORB



chr17:37095995–37196496 Marker Peak Tracks for: L3.CUX2.RORB

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

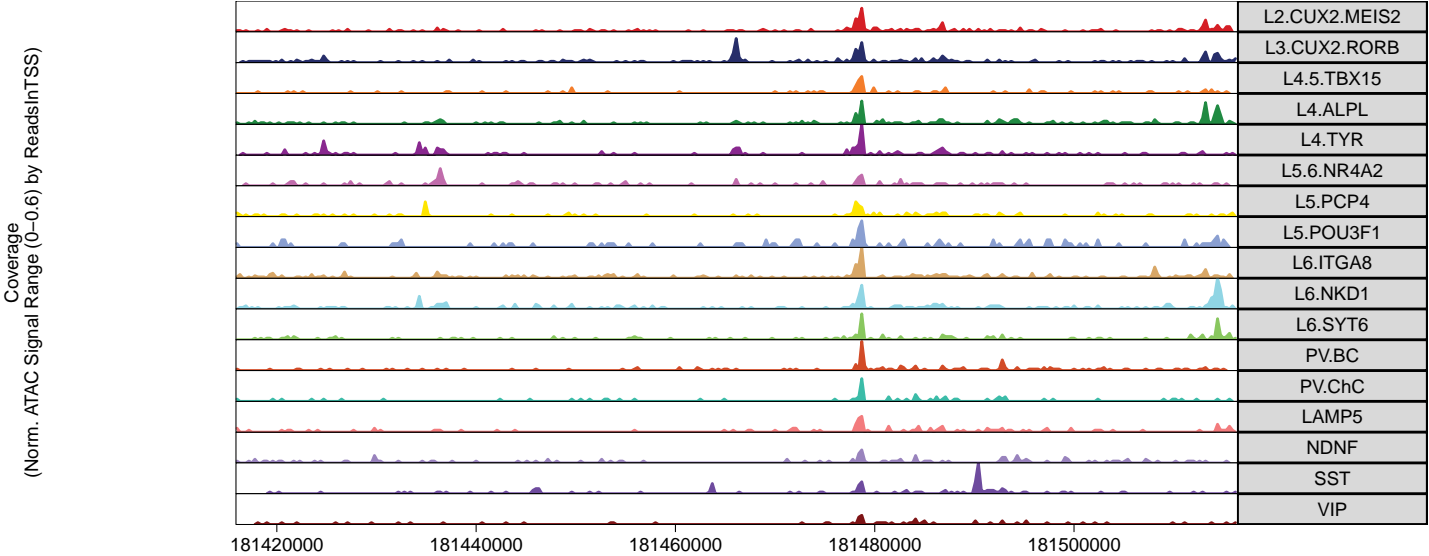
L3.CUX2.RORB



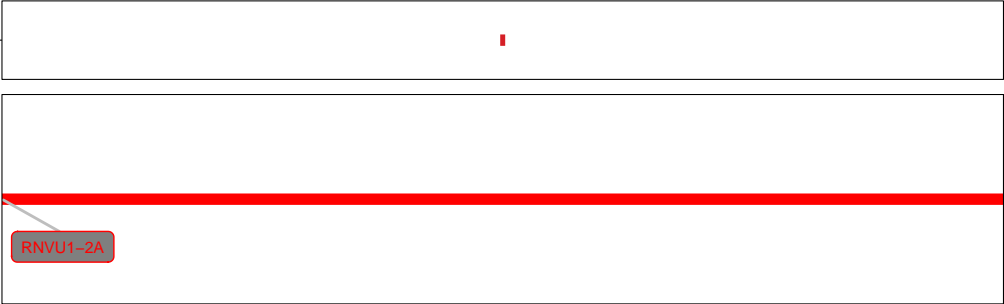
Peaks

strand • •

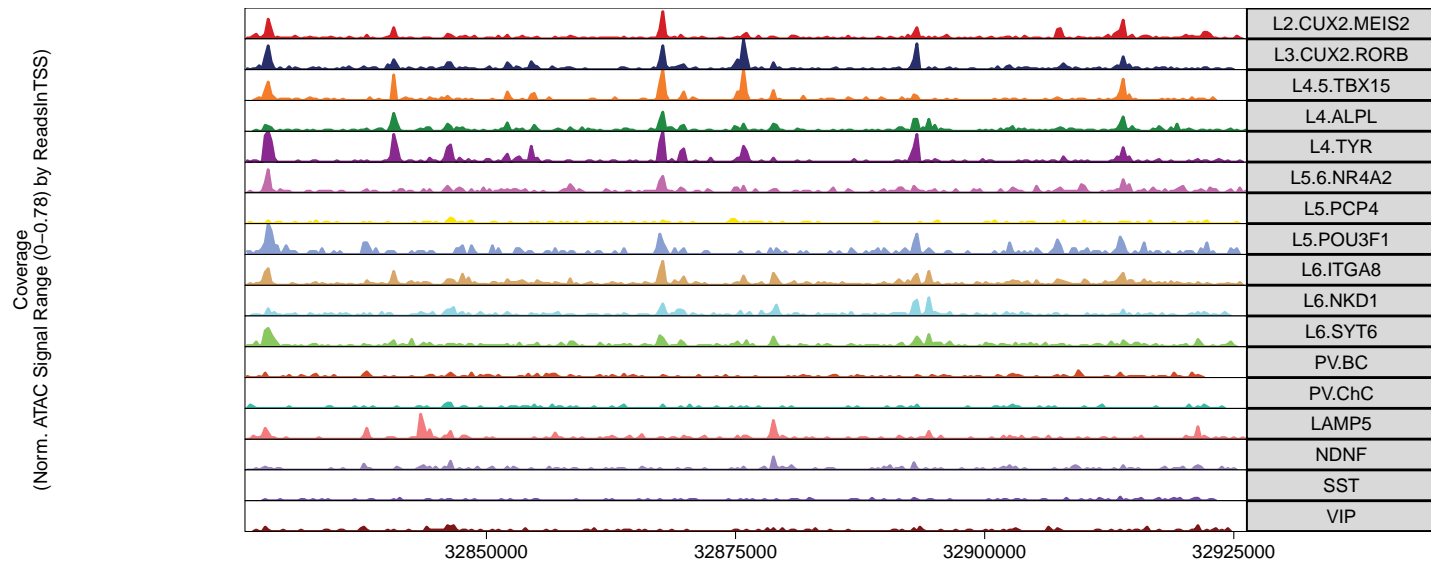
chr1:181415898–181516399 Marker Peak Tracks for: L3.CUX2.RORB



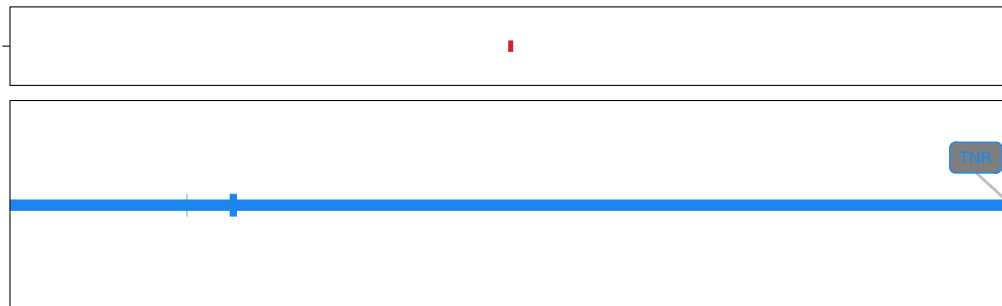
L3.CUX2.RORB



chr1:32825768–32926269 Marker Peak Tracks for: L3.CUX2.RORB

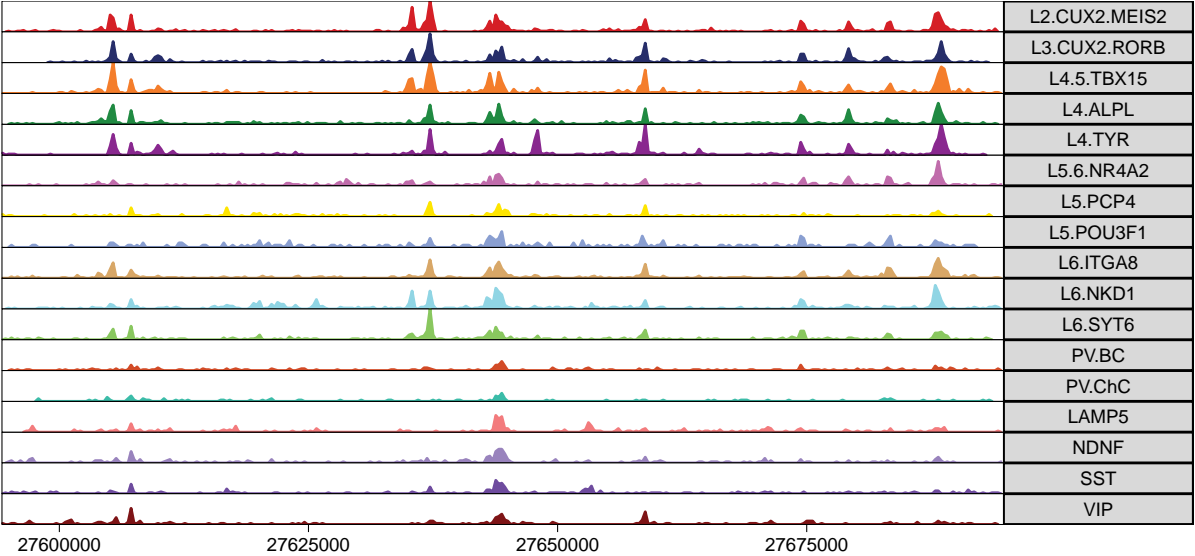


L3.CUX2.RORB

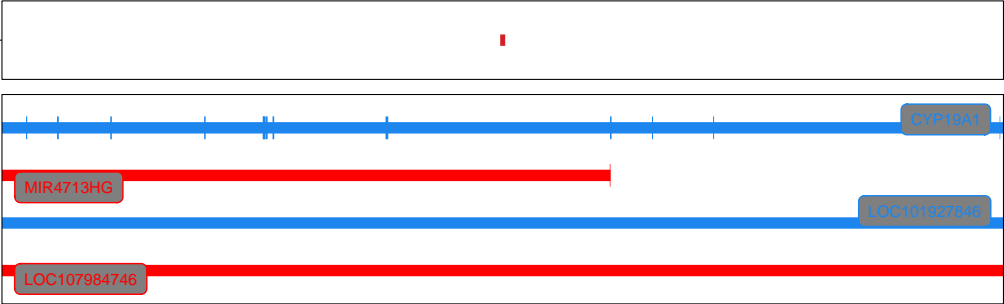


chr7:27594241–27694742 Marker Peak Tracks for: L3.CUX2.RORB

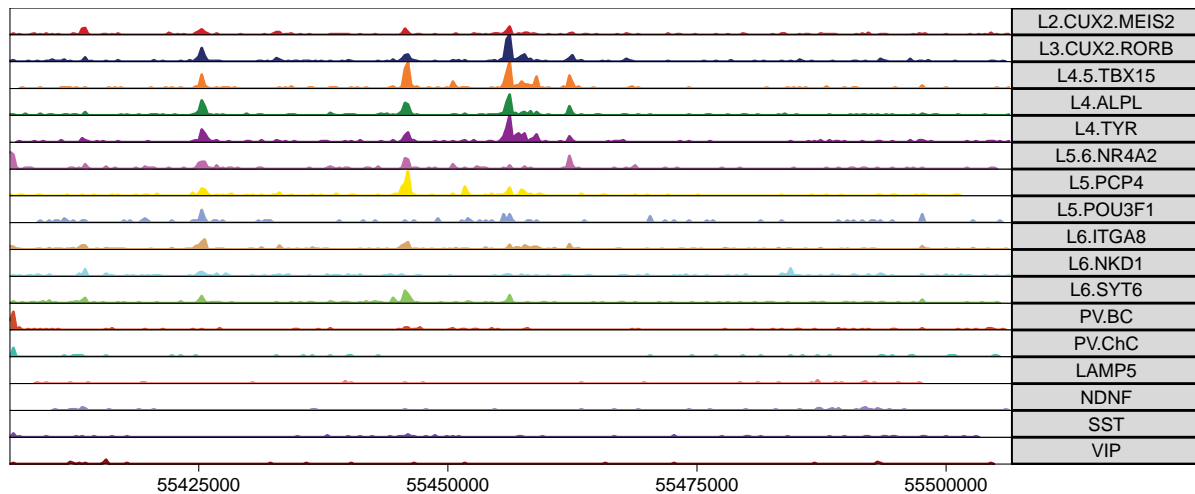
Coverage
(Norm. ATAC Signal Range (0–1) by ReadsInTSS)



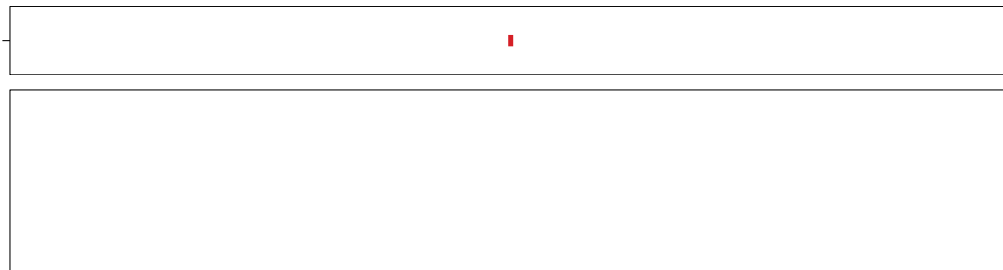
L3.CUX2.RORB



chr1:55406056–55506557 Marker Peak Tracks for: L3.CUX2.RORB

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

L3.CUX2.RORB



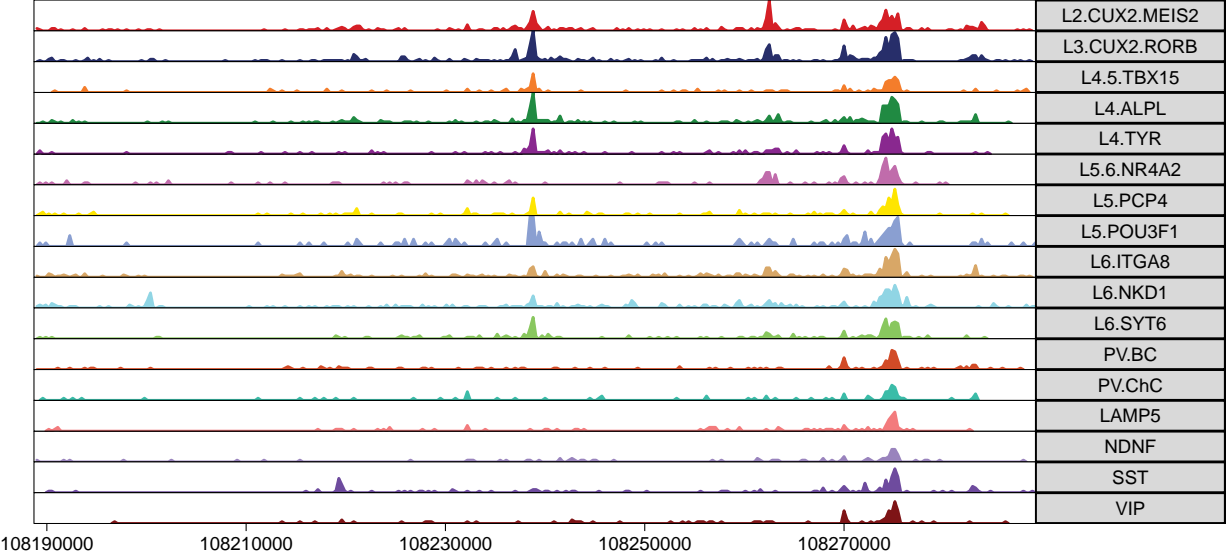
Peaks

GeneTrack

strand • •

chr1:108188707-108289208 Marker Peak Tracks for: L3.CUX2.RORB

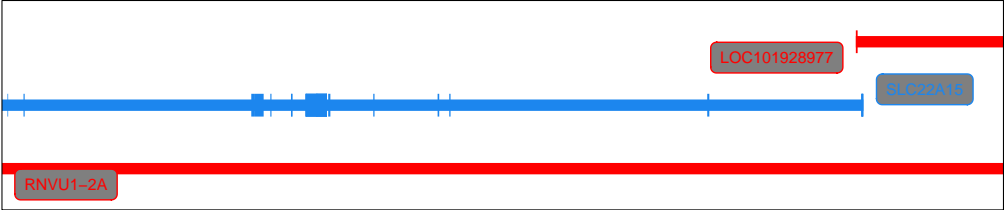
Coverage
(Norm. ATAC Signal Range (0-0.62) by ReadsInTSS)



L3.CUX2.RORB



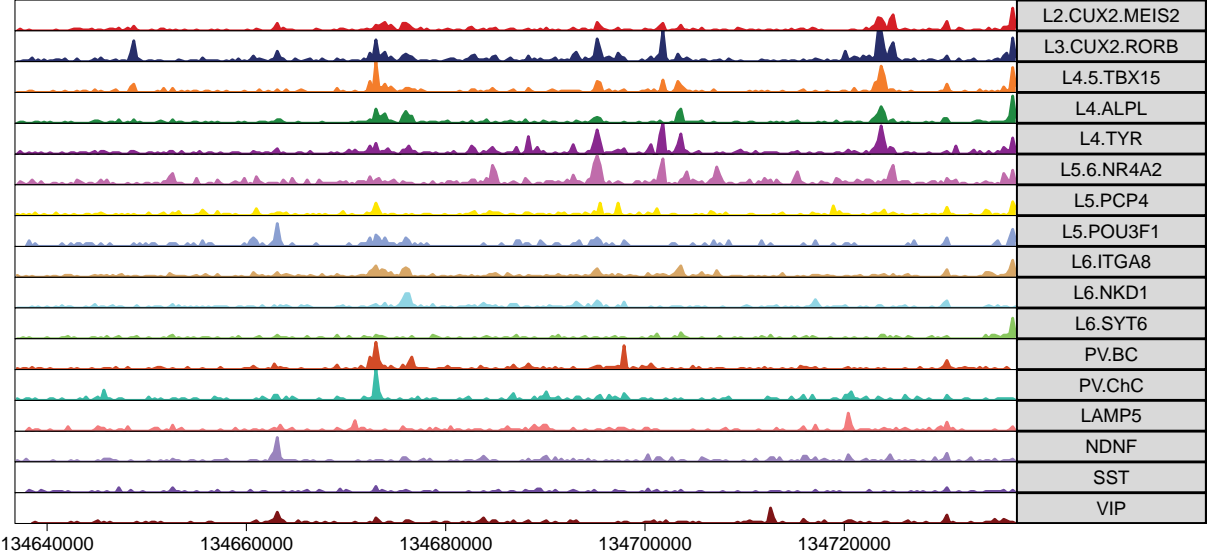
Peaks



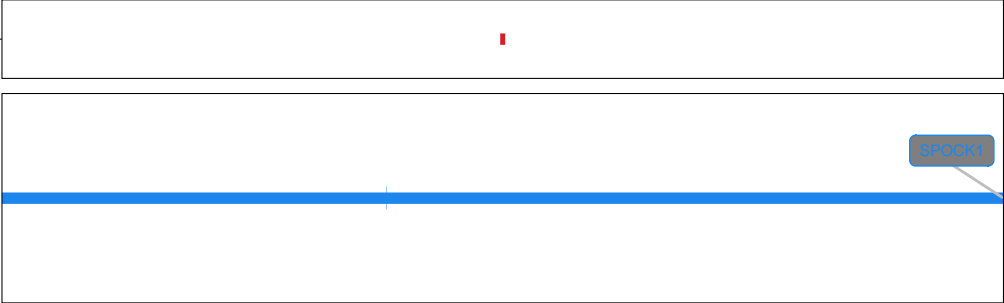
Genes

chr6:134636775–134737276 Marker Peak Tracks for: L3.CUX2.RORB

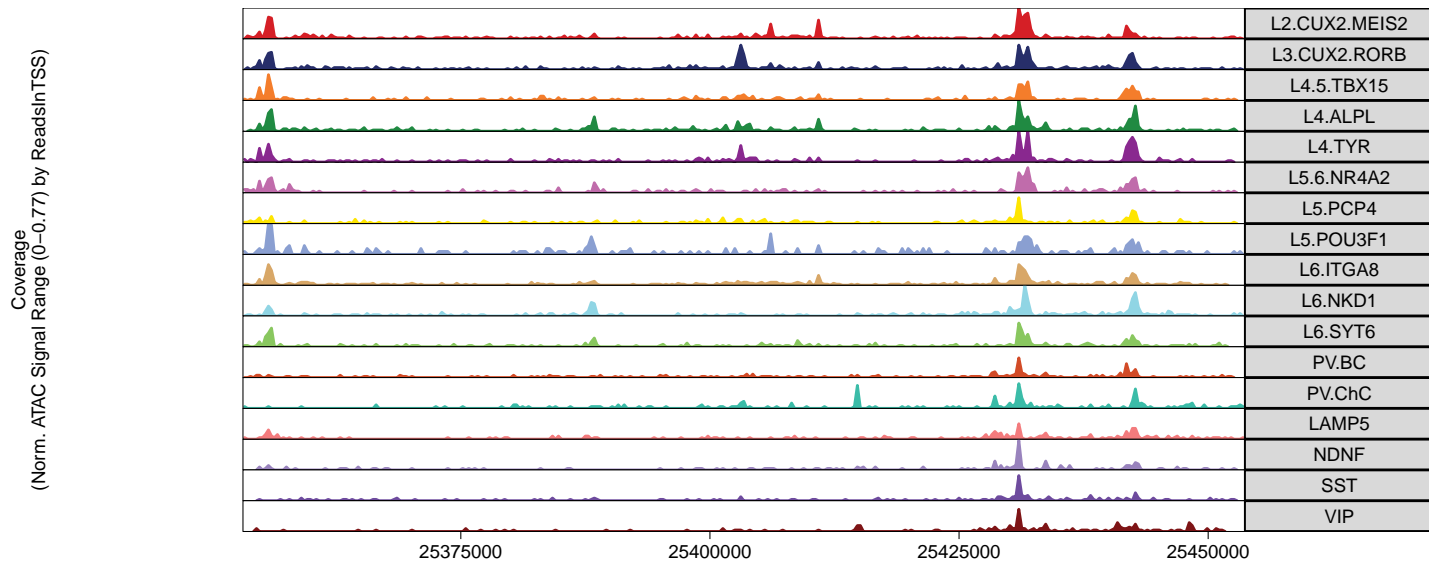
Coverage
(Norm. ATAC Signal Range (0–0.79) by ReadsInTSS)



L3.CUX2.RORB



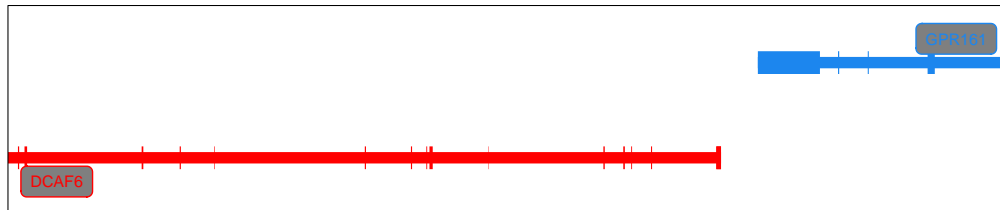
chr1:25353139–25453640 Marker Peak Tracks for: L3.CUX2.RORB



L3.CUX2.RORB

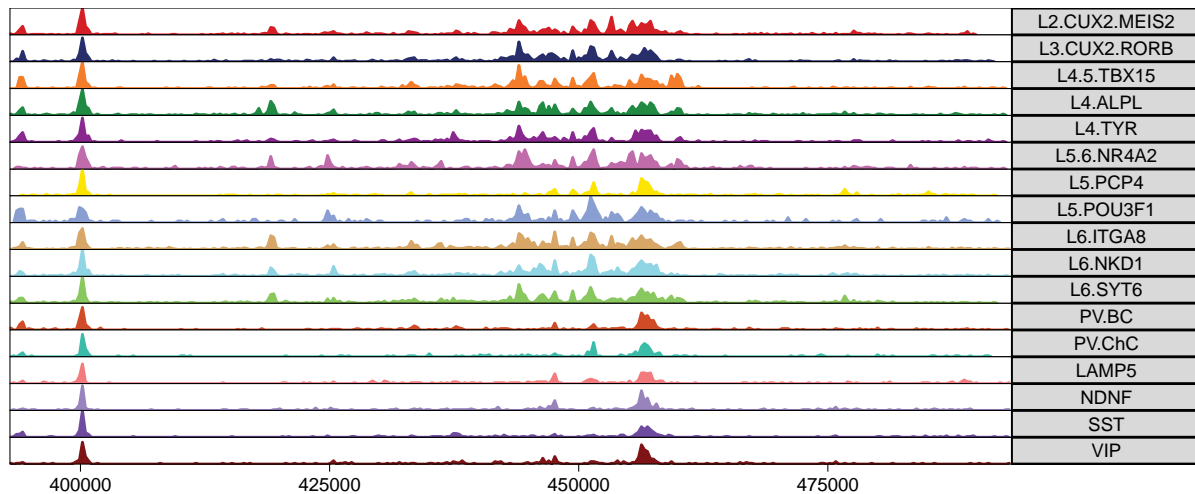


Peaks



Genes

chr17:392924–493425 Marker Peak Tracks for: L3.CUX2.RORB

Coverage
(Norm. ATAC Signal Range (0–1.16) by ReadsInTSS)

L3.CUX2.RORB



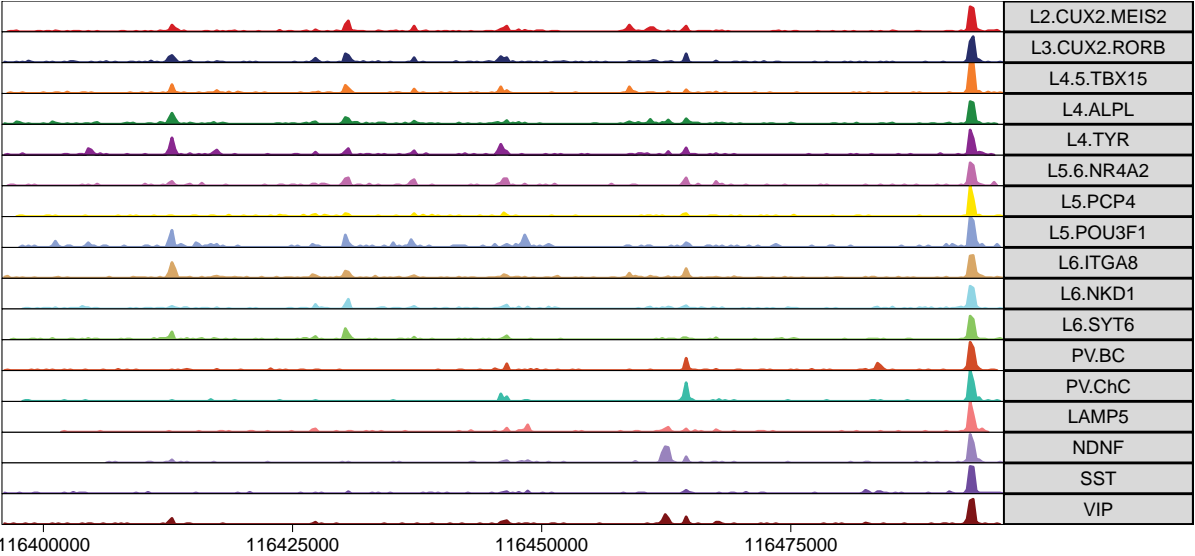
Peaks

GeneTrack

strand • ·

chr7:116395839–116496340 Marker Peak Tracks for: L3.CUX2.RORB

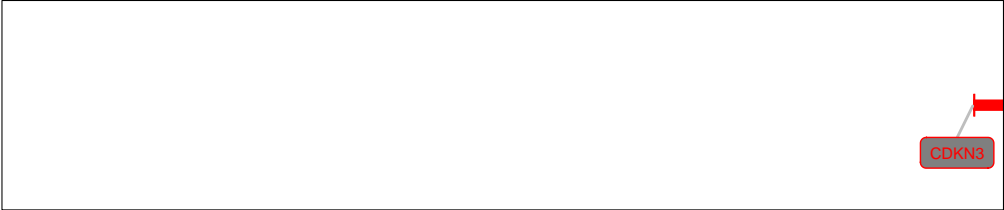
Coverage
(Norm. ATAC Signal Range (0–1.45) by ReadsInTSS)



L3.CUX2.RORB

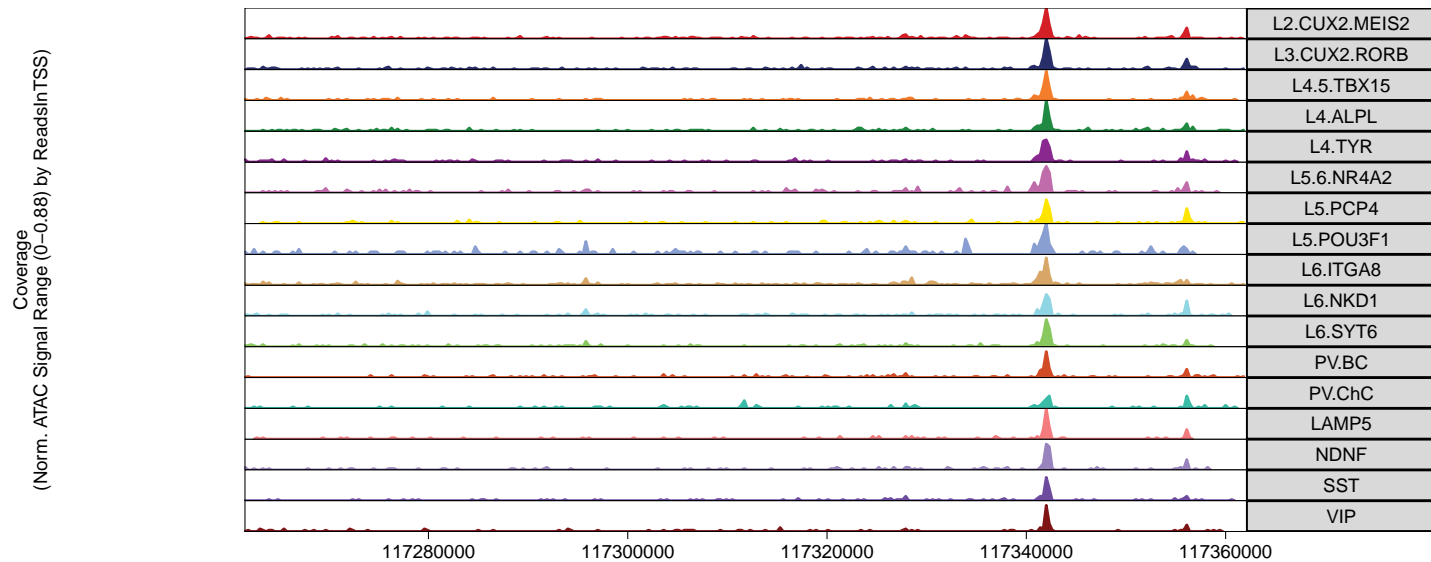


Peaks



Genes

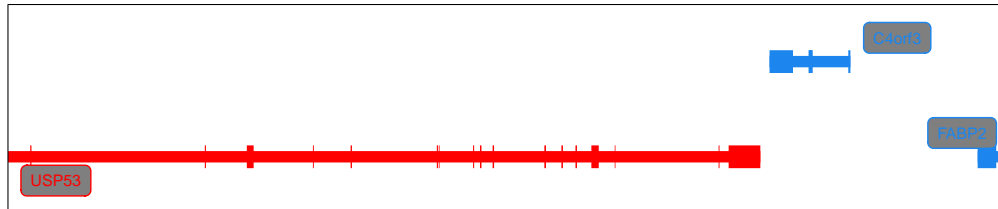
chr5:117261588–117362089 Marker Peak Tracks for: L3.CUX2.RORB



L3.CUX2.RORB

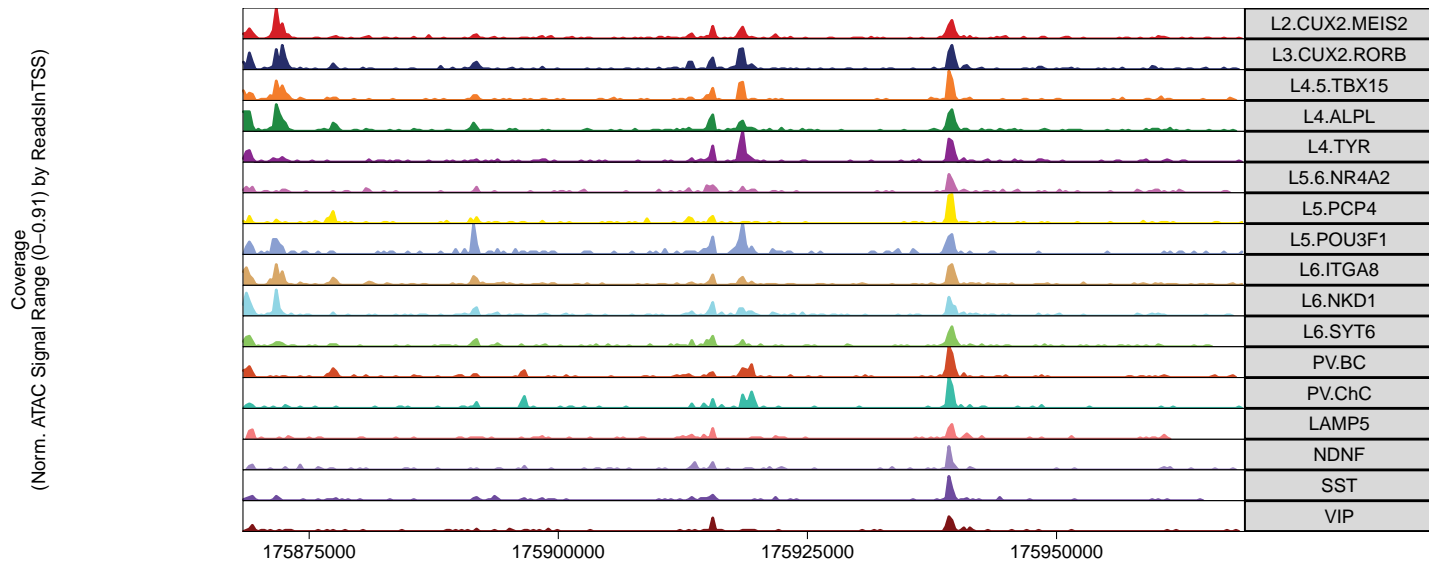


Peaks



Genes

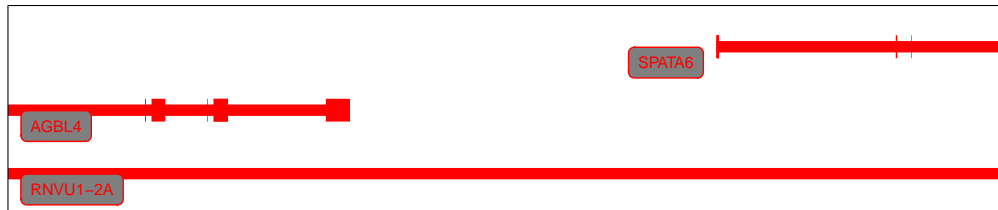
chr1:175868355–175968856 Marker Peak Tracks for: L3.CUX2.RORB



L3.CUX2.RORB



Peaks



Genes