Scale
chr7: 113,750,000 113,800,000 113,850,000 113,900,000 113,950,000 114,000,000 114,050,000 114,100,000 114,150,000 114,200,000 114,250,000 114,300,000 114,350,000
200 kb hg19

User Supplied Track

FOXP2

GENCODE V47lift37 (18 items filtered out)

ENSG00000299157 ENSG00000299157 ENSG00000299139 RPL36P13 ENSG00000224595 FOXP2 ENSG00000282859 RNA5SP238 MIR3666 ENSG00000303050 AC020606.1

100 vertebrates conservation by PhastCons

Layered H3K27Ac — H3K27Ac Mark (Often Found Near Active Regulatory Elements) on 7 cell lines from ENCODE

GH Reg Elems (DE) — GeneHancer Regulatory Elements and Gene Interactions

PPP1R3A FOXP2 — GeneCards genes TSS (Double Elite)

GH07J113918 GH07J114047 GH07J114051 PPP1R3A GH07J114081 — Interactions between GeneHancer regulatory elements and genes (Double Elite)

PPP1R3A FOXP2 — Clustered interactions of GeneHancer regulatory elements and genes (Double Elite)

Transcription — ln(x+1) — Transcription Levels Assayed by RNA-seq on 9 Cell Lines from ENCODE

Layered H3K4Me1 — H3K4Me1 Mark (Often Found Near Regulatory Elements) on 7 cell lines from ENCODE

Layered H3K4Me3 — H3K4Me3 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE

Txn Factor ChIP — Transcription Factor ChIP-seq Clusters (161 factors) from ENCODE with Factorbook Motifs

100 vertebrates Basewise Conservation by PhyloP

DNaseI Hypersensitivity Clusters in 125 cell types from ENCODE (V3)

Transcription Factor ChIP-seq Clusters (338 factors, 130 cell types) from ENCODE 3

Transcription Factor ChIP-seq from ENCODE (V2)

Repeating Elements by RepeatMasker

SINE LINE LTR DNA Simple Low Complexity Satellite RNA Other Unknown