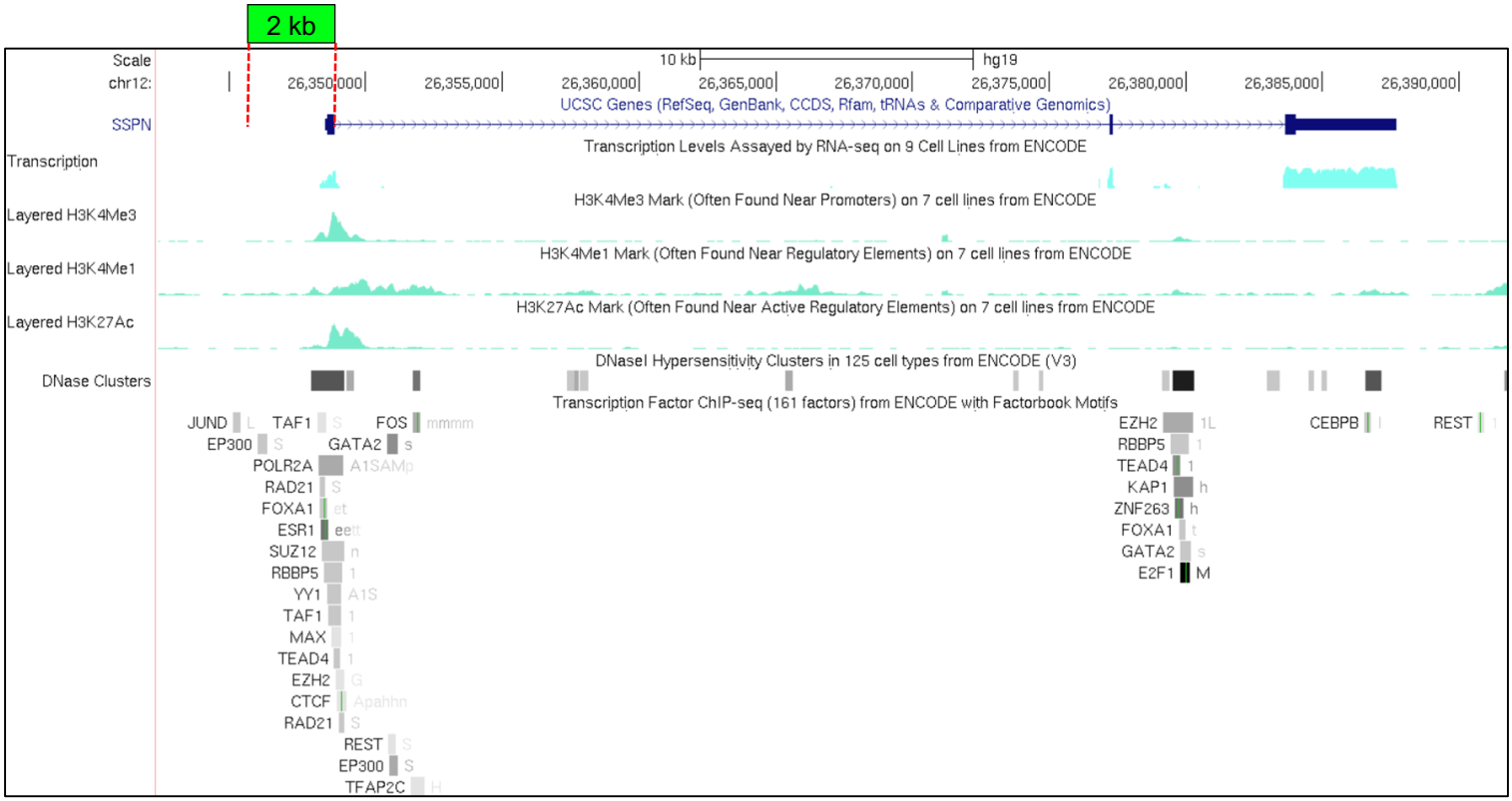
2 kb
Scale
chr12:
10 kb
hg19
26,350,000
26,355,000
26,360,000
26,365,000
26,370,000
26,375,000
26,380,000
26,385,000
26,390,000
UCSC Genes (RefSeq, GenBank, CCDS, Rfam, tRNAs & Comparative Genomics)
SSPN
Transcription
Transcription Levels Assayed by RNA-seq on 9 Cell Lines from ENCODE
Layered H3K4Me3
H3K4Me3 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE
Layered H3K4Me1
H3K4Me1 Mark (Often Found Near Regulatory Elements) on 7 cell lines from ENCODE
Layered H3K27Ac
H3K27Ac Mark (Often Found Near Active Regulatory Elements) on 7 cell lines from ENCODE
DNase Clusters
DNaseI Hypersensitivity Clusters in 125 cell types from ENCODE (V3)
Transcription Factor ChIP-seq (161 factors) from ENCODE with Factorbook Motifs
JUND
TAF1
FOS
EP300
GATA2
POLR2A
RAD21
FOXA1
ESR1
SUZ12
RBBP5
YY1
TAF1
MAX
TEAD4
EZH2
CTCF
RAD21
REST
EP300
TFAP2C
EZH2
RBBP5
TEAD4
KAP1
ZNF263
FOXA1
GATA2
E2F1
CEBPB
REST