Scale
chr19:
RPS11
5 kb
hg19
50,000,000
50,005,000
U56
hsa-mir-150
Placental Mammal Basewise Conservation by PhyloP
Mammal Cons
4
0
-4
Digital DNaseI Hypersensitivity Clusters from ENCODE
DNase Clusters
Open Chromatin by DNaseI HS from ENCODE/OpenChrom(Duke University)
GM12878 Pk
GM12878 DS
GM12878 OS
H1hESC Pk
H1hESC DS
H1hESC OS
K562 Pk
K562 DS
K562 OS
ENCODE Histone Modification Peaks and Signal based on Uniform processing pipeline
GM12878 H2A.Z
GM12878 H3K27ac
GM12878 H3K27me3
GM12878 H3K36me3
GM12878 H3K4me1
GM12878 H3K4me2
GM12878 H3K4me3
GM12878 H3K79me2
GM12878 H3K9ac
GM12878 H3K9me3
ENCODE Transcription Factor ChIP-seq Peaks and Signal based on Uniform processing pipeline
GM12878 CTCF Pk
GM12878 RAD21 Pk
GM12878 SMC3 Pk
GM12878 POL2 Pk
GM12878 POL2 Pk
GM12878 POL2 Pk
GM12878 BCL3 Pk
GM12878 CHD2 Pk
GM12878 EBF1 Pk
GM12878 ELF1 Pk
GM12878 FOXA3 Pk
GM12878 POU2F2 Pk
GM12878 PU1 Pk
GM12878 TCF12 Pk
GM12878 USF1 Pk
GM12878 GABP Pk
GM12878 BRCA1 Pk
GM12878 EBF1 Pk
GM12878 EGR1 Pk
GM12878 ETS1 Pk
GM12878 NRF1 Pk
GM12878 PAX5C20
GM12878 PAX5N19
GM12878 RFX5 Pk
GM12878 RXRA Pk
GM12878 SIN3AK20
GM12878 SIX5 Pk
GM12878 SP1 Pk
GM12878 SRF Pk
GM12878 TAF1 Pk
GM12878 TBP Pk
GM12878 USF2 Pk
GM12878 WHIP Pk
GM12878 YY1 Pk
GM12878 ZBTB33 Pk
GM12878 ZEB1 Pk
GM12878 ZNF143 Pk