Scale
5 kb
hg38
chr17: 7,670,000 7,675,000 7,680,000 7,685,000
GENCODE V49 (9 items filtered)
TP53
WRAP53
RefSeq genes from NCBI
RefSeq Curated
0.338633
Quantitative R-loop Score
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Stacked Quantitative R-loop Score
Quantitative R-loop Zone
HEK293 Quantitative R-loop Region
2.49149
HEK293 Score
HEK293 Quantitative R-loop Score
HeLa Quantitative R-loop Region
2.82068
HeLa Score
HeLa Quantitative R-loop Score
V2-Level-1
V2-Level-4
V2-Level-8
1.134
DRIP-seq_HEK293
bigwig_99_HEK293_Control_DRIP-seq_ChIP_NS_1_SE
bigbed_99_HEK293_Control_DRIP-seq_ChIP_NS_1_SE
4.8087
CUT&Tag_HEK293T
bigwig_229_HEK293T_2HBD-Anti-GST_CUT&Tag_ChIP_NS_2_PE
bigbed_229_HEK293T_2HBD-Anti-GST_CUT&Tag_ChIP_NS_2_PE
0.4572
R-ChIP-SS_HEK293T.fwd
bigwig_127_HEK293T_D210N_R-ChIP-SS_ChIP_SS_1_SE_fwd
-0
R-ChIP-SS_HEK293T.rev
bigwig_127_HEK293T_D210N_R-ChIP-SS_ChIP_SS_1_SE_rev
-1.2192
bigbed_127_HEK293T_D210N_R-ChIP-SS_Input_SS_1_SE
0.99864
DeepER_Human.fwd
bigwig_DeepER_Human_fwd
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bigbed_DeepER_Human
-7e-05
DeepER_Human.rev
bigwig_DeepER_Human_rev
-0.99705
0.8
GC-skew
-0.702128
0.737705
AT-skew
-0.688889
0.3538
GRO-seq_HEK293T.fwd
HEK293T_Control_GRO-seq.fwd
0.1769
-0.1769
GRO-seq_HEK293T.rev
HEK293T_Control_GRO-seq.rev
-5.4839
RLFS_RloopDB
CLIP_AQR
HepG2_exp1_39750
HepG2_exp1_39751
K562_exp1_72852
K562_exp1_72853
HepG2_exp1_39752
K562_exp1_72854
K562_exp1_72855
HepG2_exp1_39753
K562_exp1_72856
K562_exp1_72857
HepG2_exp1_39754
K562_exp1_72858
HepG2_exp1_39756
K562_exp1_72859
HepG2_exp1_39757
K562_exp1_72860
HepG2_exp1_39755
K562_exp1_72861
HepG2_exp1_39758
K562_exp1_72862
HepG2_exp1_39759
K562_exp1_72863
HepG2_exp1_39760
K562_exp1_72864
ChIP_BRCA1
HeLa-S3_exp1_4490
GM12878_exp1_609
HepG2_exp2_3330
H1_exp1_771
HepG2_exp2_3331
KD_DDX5
Gene Expression in 54 tissues from GTEx RNA-seq of 17382 samples, 948 donors (V8, Aug 2019)
TP53
RP11-199F11.2
WRAP53
Repeating Elements by RepeatMasker
RepeatMasker