Scale
chr1:
50,000,000
100,000,000
150,000,000
200,000,000
100 Mb
hg38
RefSeq Curated
RefSeq genes from NCBI
Stacked Quantitative R-loop Score
Quantitative R-loop Score
Quantitative R-loop Zone
HEK293 Region
HEK293 Quantitative R-loop Region
HEK293 Score
HEK293 Quantitative R-loop Score
HeLa Region
HeLa Quantitative R-loop Region
HeLa Score
HeLa Quantitative R-loop Score
R-loop Zone Level
V2-Level-1
V2-Level-4
V2-Level-8
DRIP-seq_HEK293
R-loop Signal
DRIP-seq_HEK293
bigwig_99_HEK293_Control_DRIP-seq_ChIP_NS_1_SE
CUT&Tag_HEK293
R-loop Signal
CUT&Tag_HEK293
bigwig_229_HEK293T_2HBD-Anti-GST_CUT&Tag_ChIP_NS_2_PE
R-ChIP-SS_HEK293T.fwd
bigwig_127_HEK293T_D210N_R-ChIP-SS_ChIP_SS_1_SE_fwd
R-ChIP-SS_HEK293T.rev
bigwig_127_HEK293T_D210N_R-ChIP-SS_ChIP_SS_1_SE_rev
bigbed_127_HEK293T_D210N_R-ChIP-SS_Input_SS_1_SE
bigwig_DeepER_Human_fwd
DeepER_Human.fwd
DeepER
bigwig_DeepER_Human_rev
DeepER_Human
DeepER_Human.rev
GC-skew
AT-skew
GRO-seq_HEK293T.fwd
HEK293T_Control_GRO-seq.fwd
GRO-seq_HEK293T.rev
HEK293T_Control_GRO-seq.rev
RLFS_RloopDB
Regulator
ChIP_BRCA1
CLIP_AQR
KD_DDX5
GTEx Gene V8
Gene Expression in 54 tissues from GTEx RNA-seq of 17382 samples, 948 donors (V8, Aug 2019)
RepeatMasker
Repeating Elements by RepeatMasker
zoom in to <= 10,000,000 bases to view items