
Scale
chr1:
500 bases
hg38
109,548,500
109,549,000
GENCODE V49
GENCODE V49 (1 items filtered)
NCBI RefSeq genes, curated subset (NM_*, NR_*, NP_* or YP_*)
GNAI3/NM_006496.4
Stacked Quantitative R-loop Score
Quantitative R-loop Score
Quantitative R-loop Zone
Quantitative R-loop Zone
HEK293 Quantitative R-loop Region
HEK293 Region
HEK293 Quantitative R-loop Score
HEK293 Score
HeLa Quantitative R-loop Region
HeLa Region
HeLa Quantitative R-loop Score
HeLa Score
R-loop Zone Level
V2-Level-1
V2-Level-4
V2-Level-8
R-loop Signal
DRIP-seq_HEK293
DRIP-seq_HEK293
CUT&Tag_HEK293T
CUT&Tag_HEK293T
R-ChIP-SS_HEK293T.fwd
R-ChIP-SS_HEK293T.rev
R-ChIP-SS_HEK293T
DeepER
DeepER_Human.fwd
DeepER_Human
DeepER_Human.rev
GC-skew
GC-skew
AT-skew
AT-skew
Transcription
GRO-seq_HEK293T.fwd
GRO-seq_HEK293T.rev
RLFS_RloopDB
RLFS_RloopDB
Regulator
CLIP_AQR
ChIP_BRCA1
KD_DDX5
Gene Expression in 54 tissues from GTEx RNA-seq of 17382 samples, 948 donors (V8, Aug 2019)
GTEx Gene V8
Repeating Elements by RepeatMasker
SINE
LINE
LTR
DNA
Simple
Low Complexity
Satellite
RNA
Other
Unknown