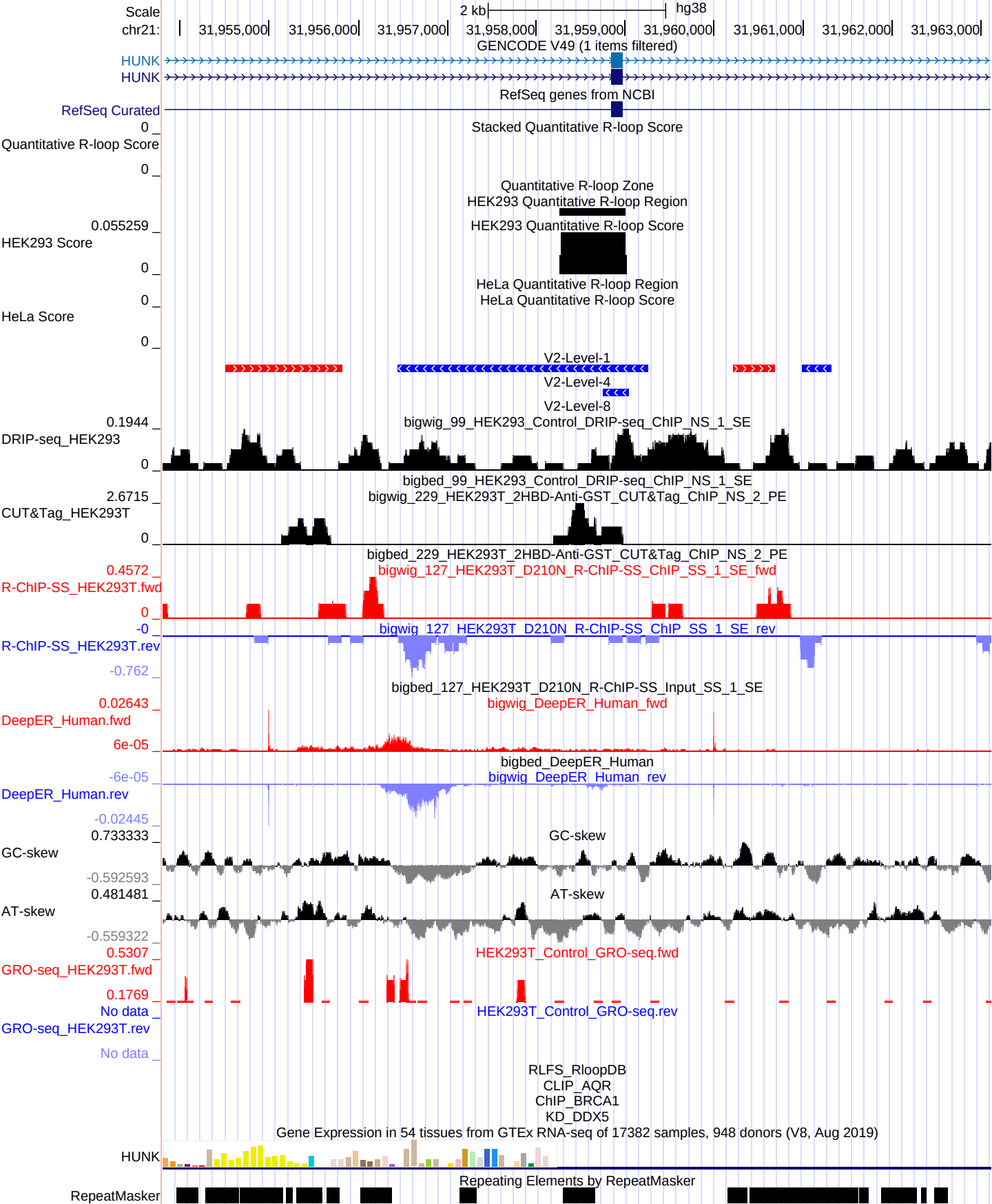

Scale
2 kb
hg38
chr21:
31,955,000
31,956,000
31,957,000
31,958,000
31,959,000
31,960,000
31,961,000
31,962,000
31,963,000
GENCODE V49 (1 items filtered)
HUNK
HUNK
RefSeq genes from NCBI
RefSeq Curated
0
Stacked Quantitative R-loop Score
Quantitative R-loop Score
0
Quantitative R-loop Zone
HEK293 Quantitative R-loop Region
0.055259
HEK293 Quantitative R-loop Score
HEK293 Score
0
HeLa Quantitative R-loop Region
0
HeLa Quantitative R-loop Score
HeLa Score
0
V2-Level-1
V2-Level-4
V2-Level-8
0.1944
bigwig_99_HEK293_Control_DRIP-seq_ChIP_NS_1_SE
DRIP-seq_HEK293
0
bigbed_99_HEK293_Control_DRIP-seq_ChIP_NS_1_SE
2.6715
bigwig_229_HEK293T_2HBD-Anti-GST_CUT&Tag_ChIP_NS_2_PE
CUT&Tag_HEK293T
0
bigbed_229_HEK293T_2HBD-Anti-GST_CUT&Tag_ChIP_NS_2_PE
0.4572
bigwig_127_HEK293T_D210N_R-ChIP-SS_ChIP_SS_1_SE_fwd
R-ChIP-SS_HEK293T.fwd
0
-0
bigwig_127_HEK293T_D210N_R-ChIP-SS_ChIP_SS_1_SE_rev
R-ChIP-SS_HEK293T.rev
-0.762
bigbed_127_HEK293T_D210N_R-ChIP-SS_Input_SS_1_SE
0.02643
bigwig_DeepER_Human_fwd
DeepER_Human.fwd
6e-05
bigbed_DeepER_Human
-6e-05
bigwig_DeepER_Human_rev
DeepER_Human.rev
-0.02445
0.733333
GC-skew
GC-skew
-0.592593
0.481481
AT-skew
AT-skew
-0.559322
0.5307
HEK293T_Control_GRO-seq.fwd
GRO-seq_HEK293T.fwd
0.1769
No data
HEK293T_Control_GRO-seq.rev
GRO-seq_HEK293T.rev
No data
RLFS_RloopDB
CLIP_AQR
ChIP_BRCA1
KD_DDX5
Gene Expression in 54 tissues from GTEx RNA-seq of 17382 samples, 948 donors (V8, Aug 2019)
HUNK
Repeating Elements by RepeatMasker
RepeatMasker