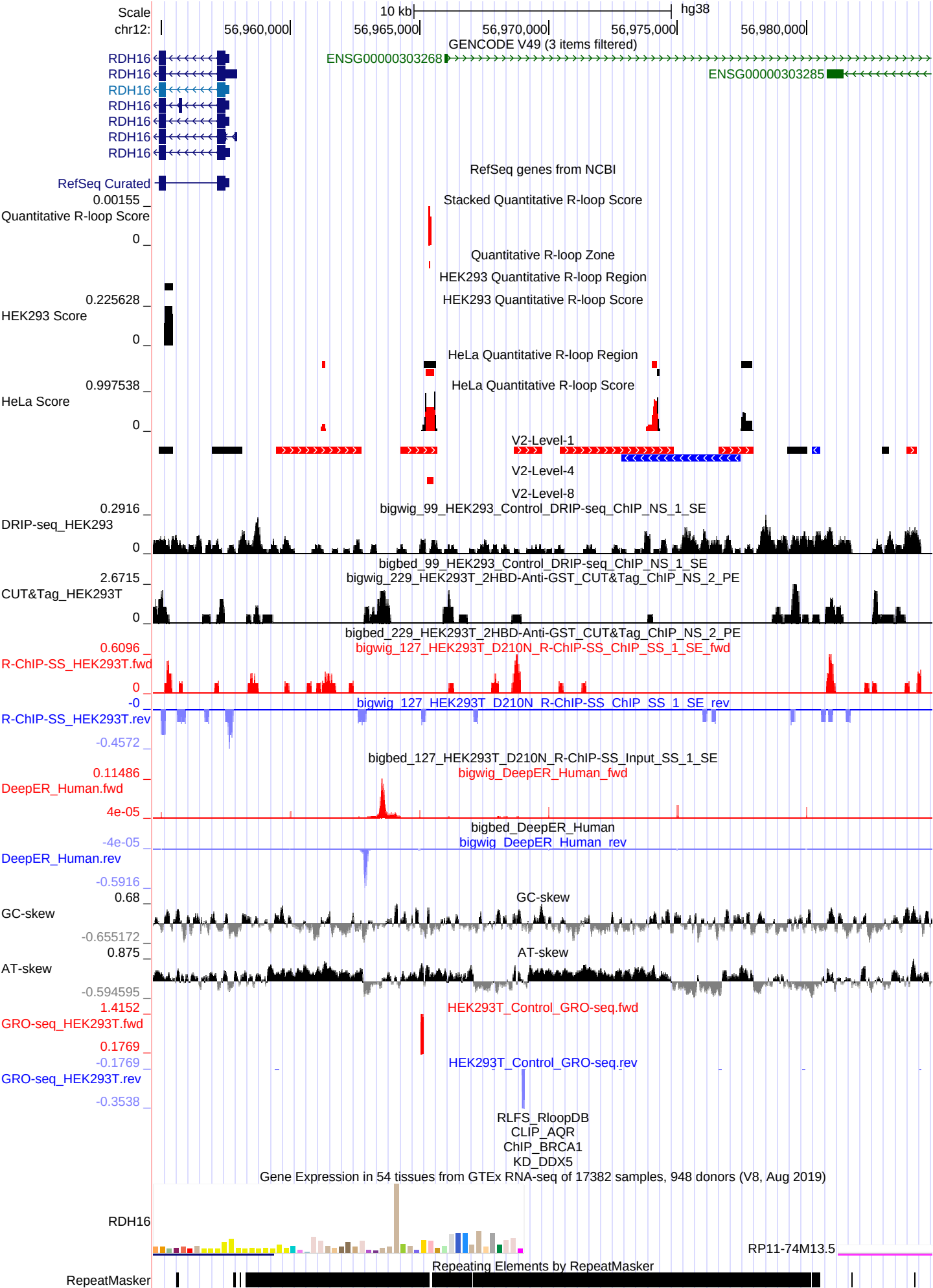

Scale
10 kb
hg38
chr12:
56,960,000
56,965,000
56,970,000
56,975,000
56,980,000
GENCODE V49 (3 items filtered)
RDH16
ENSG00000303268
ENSG00000303285
RefSeq genes from NCBI
RefSeq Curated
0.00155
Quantitative R-loop Score
Stacked Quantitative R-loop Score
0
Quantitative R-loop Zone
HEK293 Quantitative R-loop Region
0.225628
HEK293 Score
HEK293 Quantitative R-loop Score
HeLa Quantitative R-loop Region
0.997538
HeLa Score
HeLa Quantitative R-loop Score
V2-Level-1
V2-Level-4
V2-Level-8
0.2916
DRIP-seq_HEK293
bigwig_99_HEK293_Control_DRIP-seq_ChIP_NS_1_SE
bigbed_99_HEK293_Control_DRIP-seq_ChIP_NS_1_SE
2.6715
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
0.6096
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
-0.4572
0.11486
DeepER_Human.fwd
bigbed_127_HEK293T_D210N_R-ChIP-SS_Input_SS_1_SE
bigwig_DeepER_Human_fwd
4e-05
-4e-05
DeepER_Human.rev
bigbed_DeepER_Human
bigwig_DeepER_Human_rev
-0.5916
0.68
GC-skew
-0.655172
0.875
AT-skew
-0.594595
1.4152
GRO-seq_HEK293T.fwd
HEK293T_Control_GRO-seq.fwd
0.1769
-0.1769
GRO-seq_HEK293T.rev
HEK293T_Control_GRO-seq.rev
-0.3538
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)
RDH16
RP11-74M13.5
RepeatMasker
Repeating Elements by RepeatMasker