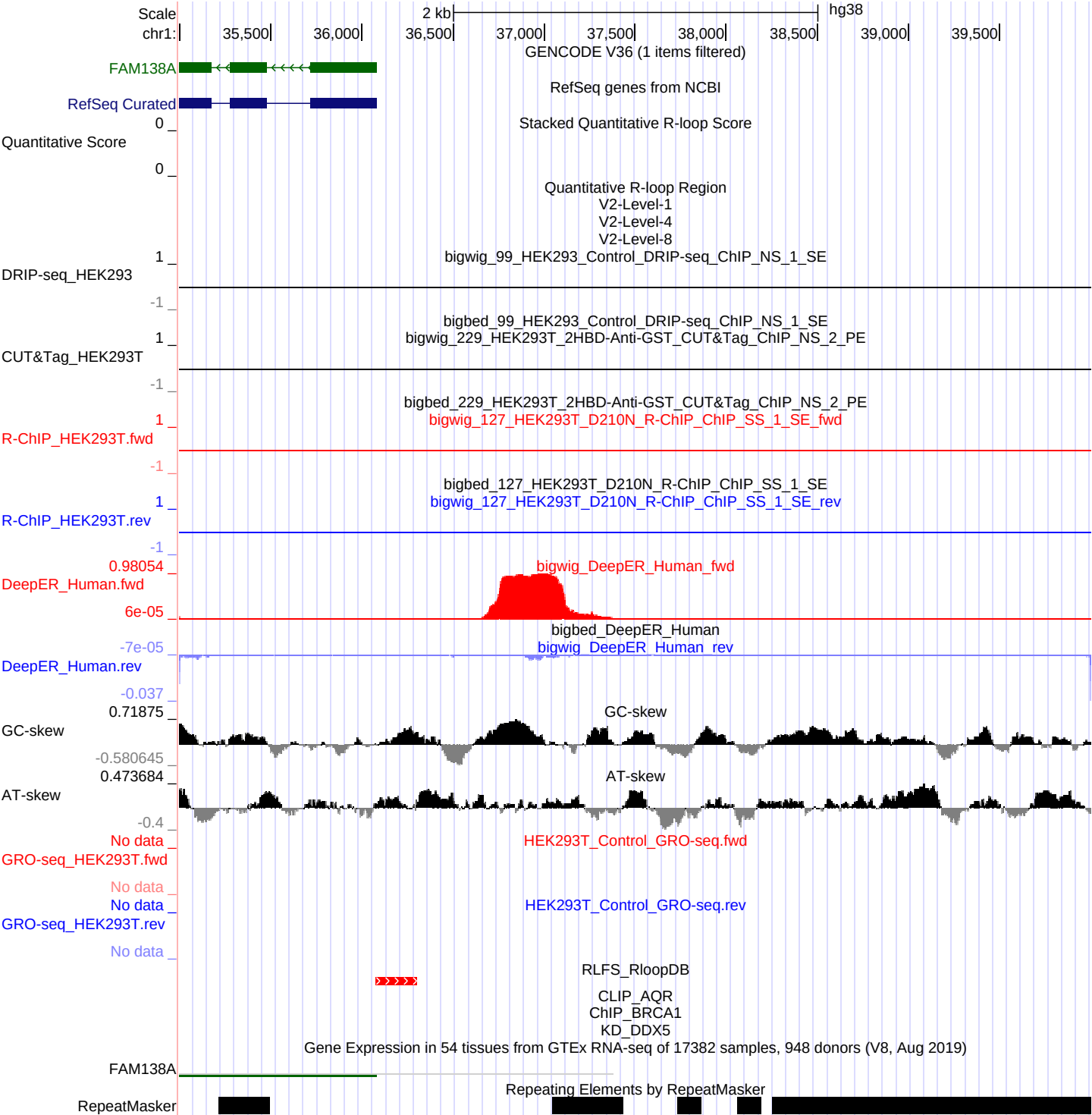
Scale
chr1:
2 kb
hg38
35,500
36,000
36,500
37,000
37,500
38,000
38,500
39,000
39,500
GENCODE V36 (1 items filtered)
FAM138A
RefSeq genes from NCBI
RefSeq Curated
Stacked Quantitative R-loop Score
0
Quantitative Score
0
Quantitative R-loop Region
V2-Level-1
V2-Level-4
V2-Level-8
1
bigwig_99_HEK293_Control_DRIP-seq_ChIP_NS_1_SE
DRIP-seq_HEK293
-1
bigbed_99_HEK293_Control_DRIP-seq_ChIP_NS_1_SE
1
bigwig_229_HEK293T_2HBD-Anti-GST_CUT&Tag_ChIP_NS_2_PE
CUT&Tag_HEK293T
-1
bigbed_229_HEK293T_2HBD-Anti-GST_CUT&Tag_ChIP_NS_2_PE
1
bigwig_127_HEK293T_D210N_R-ChIP_ChIP_SS_1_SE_fwd
R-ChIP_HEK293T.fwd
-1
bigbed_127_HEK293T_D210N_R-ChIP_ChIP_SS_1_SE
1
bigwig_127_HEK293T_D210N_R-ChIP_ChIP_SS_1_SE_rev
R-ChIP_HEK293T.rev
-1
0.98054
bigwig_DeepER_Human_fwd
DeepER_Human.fwd
6e-05
bigbed_DeepER_Human
-7e-05
bigwig_DeepER_Human_rev
DeepER_Human.rev
-0.037
0.71875
GC-skew
GC-skew
-0.580645
0.473684
AT-skew
AT-skew
-0.4
No data
HEK293T_Control_GRO-seq.fwd
GRO-seq_HEK293T.fwd
No data
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)
FAM138A
Repeating Elements by RepeatMasker
RepeatMasker