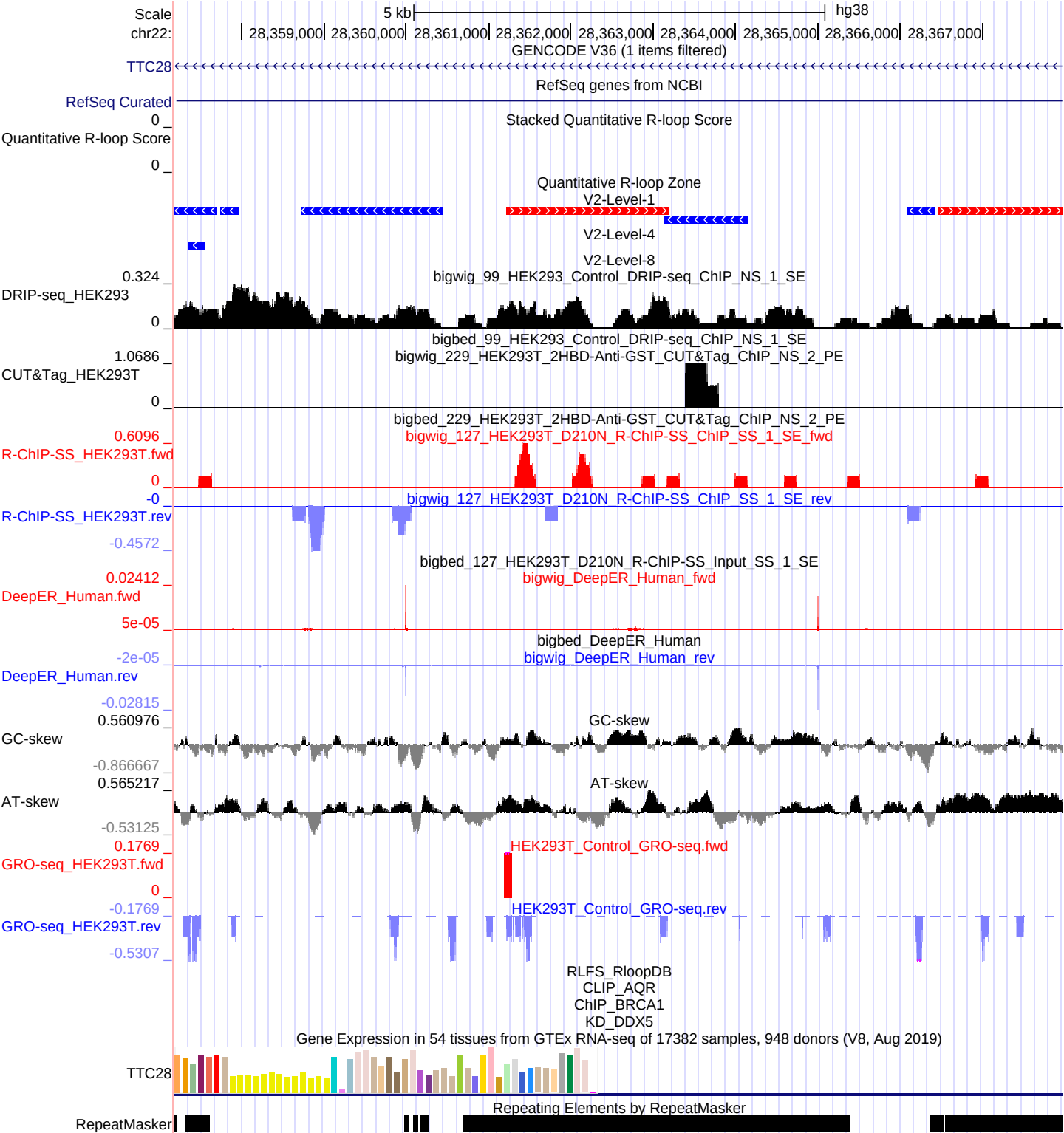
Scale
5 kb
hg38
chr22:
28,359,000
28,360,000
28,361,000
28,362,000
28,363,000
28,364,000
28,365,000
28,366,000
28,367,000
GENCODE V36 (1 items filtered)
TTC28
RefSeq genes from NCBI
RefSeq Curated
0
Stacked Quantitative R-loop Score
Quantitative R-loop Score
0
Quantitative R-loop Zone
V2-Level-1
V2-Level-4
V2-Level-8
0.324
bigwig_99_HEK293_Control_DRIP-seq_ChIP_NS_1_SE
DRIP-seq_HEK293
0
bigbed_99_HEK293_Control_DRIP-seq_ChIP_NS_1_SE
1.0686
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
0.6096
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
-0.4572
bigbed_127_HEK293T_D210N_R-ChIP-SS_Input_SS_1_SE
0.02412
bigwig_DeepER_Human_fwd
DeepER_Human.fwd
5e-05
bigbed_DeepER_Human
-2e-05
bigwig_DeepER_Human_rev
DeepER_Human.rev
-0.02815
0.560976
GC-skew
GC-skew
-0.866667
0.565217
AT-skew
AT-skew
-0.53125
0.1769
HEK293T_Control_GRO-seq.fwd
GRO-seq_HEK293T.fwd
0
-0.1769
HEK293T_Control_GRO-seq.rev
GRO-seq_HEK293T.rev
-0.5307
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)
TTC28
Repeating Elements by RepeatMasker
RepeatMasker