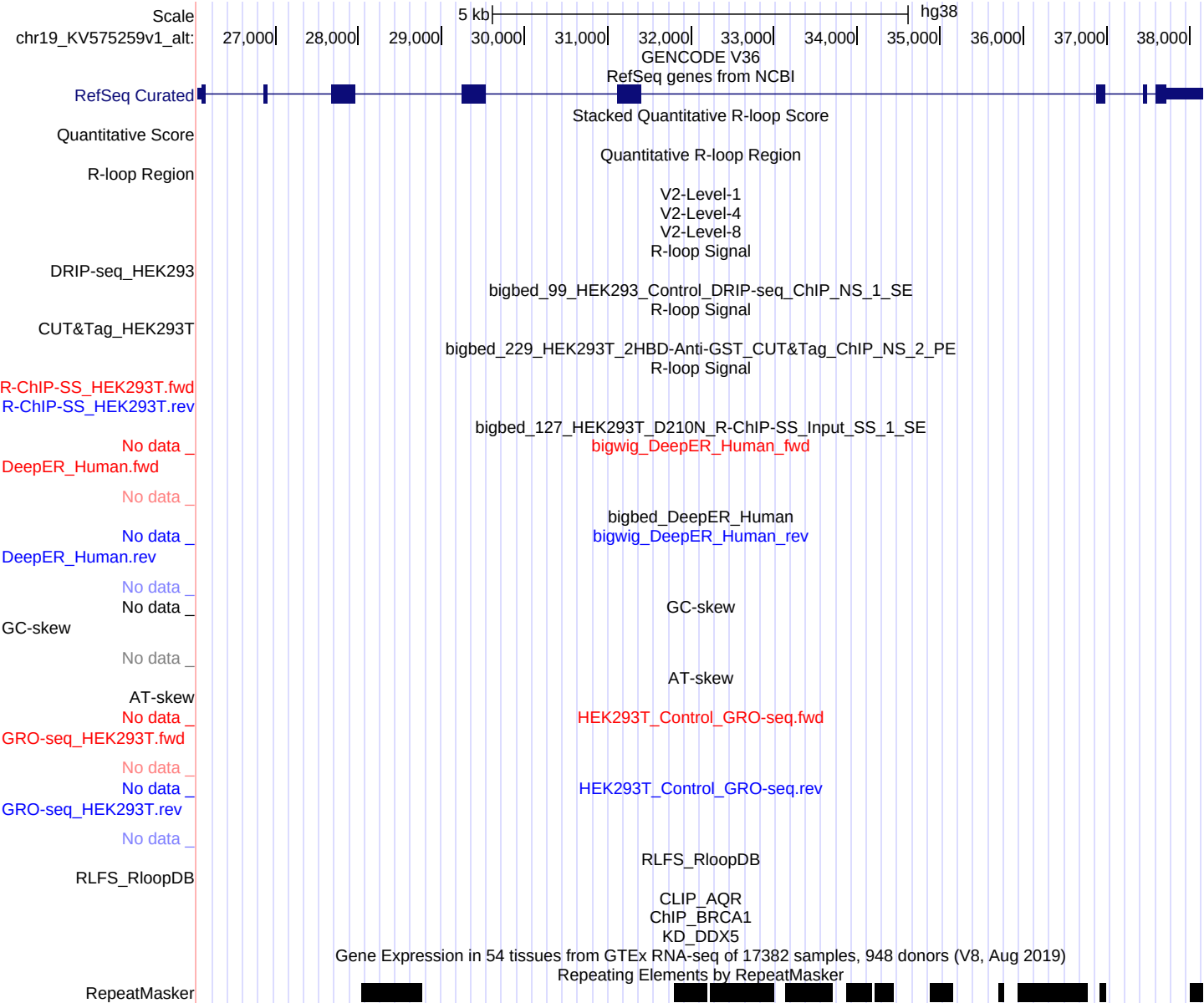
Scale
chr19_KV575259v1_alt:
5 kb
hg38
27,000
28,000
29,000
30,000
31,000
32,000
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34,000
35,000
36,000
37,000
38,000
GENCODE V36
RefSeq genes from NCBI
RefSeq Curated
Stacked Quantitative R-loop Score
Quantitative Score
Quantitative R-loop Region
R-loop Region
V2-Level-1
V2-Level-4
V2-Level-8
R-loop Signal
DRIP-seq_HEK293
bigbed_99_HEK293_Control_DRIP-seq_ChIP_NS_1_SE
R-loop Signal
CUT&Tag_HEK293T
bigbed_229_HEK293T_2HBD-Anti-GST_CUT&Tag_ChIP_NS_2_PE
R-loop Signal
R-ChIP-SS_HEK293T.fwd
R-ChIP-SS_HEK293T.rev
bigbed_127_HEK293T_D210N_R-ChIP-SS_Input_SS_1_SE
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bigwig_DeepER_Human_fwd
DeepER_Human.fwd
No data
bigbed_DeepER_Human
No data
bigwig_DeepER_Human_rev
DeepER_Human.rev
No data
No data
GC-skew
GC-skew
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AT-skew
AT-skew
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HEK293T_Control_GRO-seq.fwd
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HEK293T_Control_GRO-seq.rev
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RLFS_RloopDB
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)
Repeating Elements by RepeatMasker
RepeatMasker