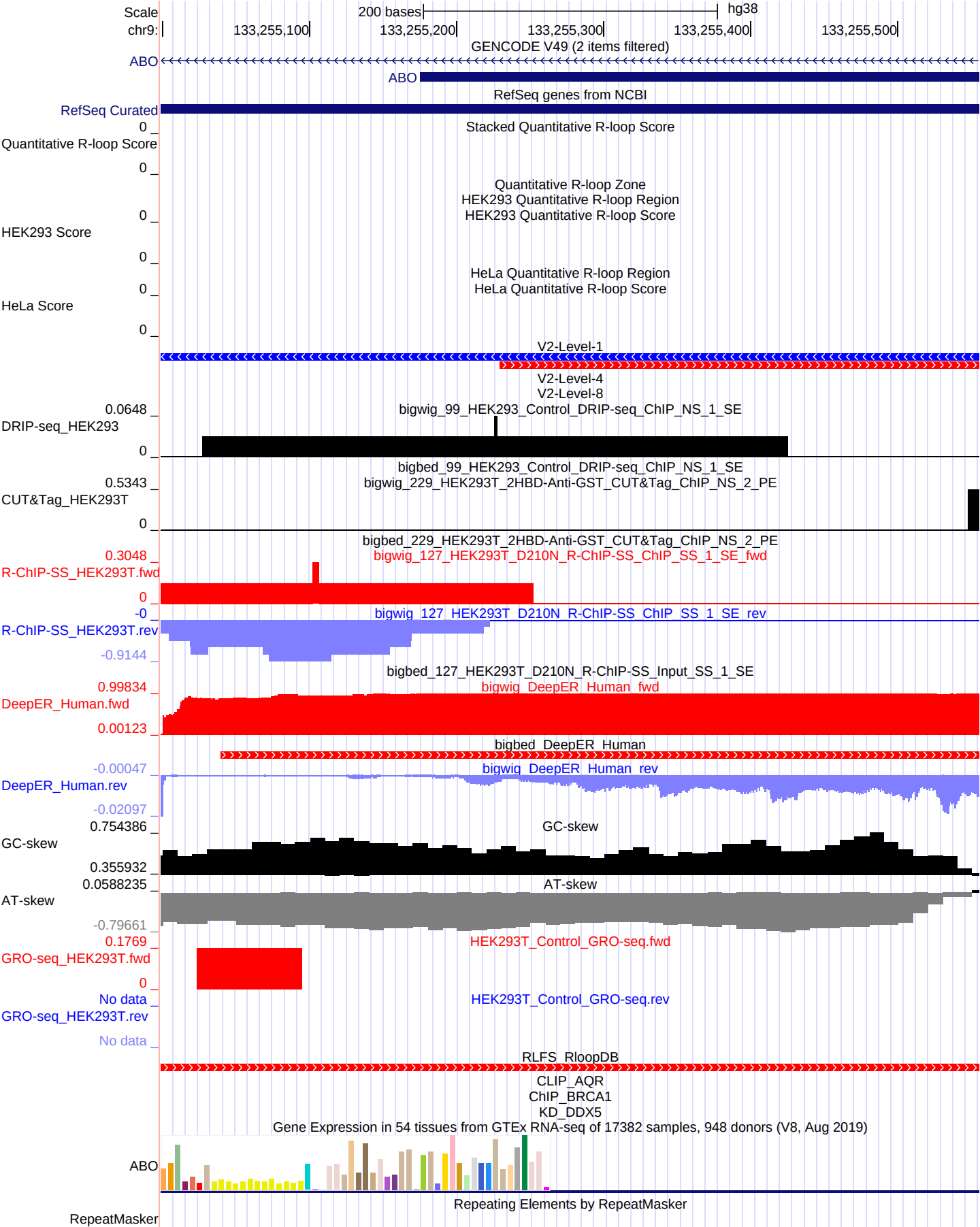

Scale
200 bases
hg38
chr9:
133,255,100
133,255,200
133,255,300
133,255,400
133,255,500
GENCODE V49 (2 items filtered)
ABO
ABO
RefSeq genes from NCBI
RefSeq Curated
0
Stacked Quantitative R-loop Score
Quantitative R-loop Score
0
Quantitative R-loop Zone
HEK293 Quantitative R-loop Region
HEK293 Quantitative R-loop Score
0
HEK293 Score
0
HeLa Quantitative R-loop Region
HeLa Quantitative R-loop Score
0
HeLa Score
0
V2-Level-1
V2-Level-4
V2-Level-8
0.0648
bigwig_99_HEK293_Control_DRIP-seq_ChIP_NS_1_SE
DRIP-seq_HEK293
0
bigbed_99_HEK293_Control_DRIP-seq_ChIP_NS_1_SE
0.5343
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
0.3048
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
-0.9144
bigbed_127_HEK293T_D210N_R-ChIP-SS_Input_SS_1_SE
0.99834
bigwig_DeepER_Human_fwd
DeepER_Human.fwd
0.00123
bigbed_DeepER_Human
-0.00047
bigwig_DeepER_Human_rev
DeepER_Human.rev
-0.02097
0.754386
GC-skew
GC-skew
0.355932
0.0588235
AT-skew
AT-skew
-0.79661
0.1769
HEK293T_Control_GRO-seq.fwd
GRO-seq_HEK293T.fwd
0
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)
ABO
Repeating Elements by RepeatMasker
RepeatMasker