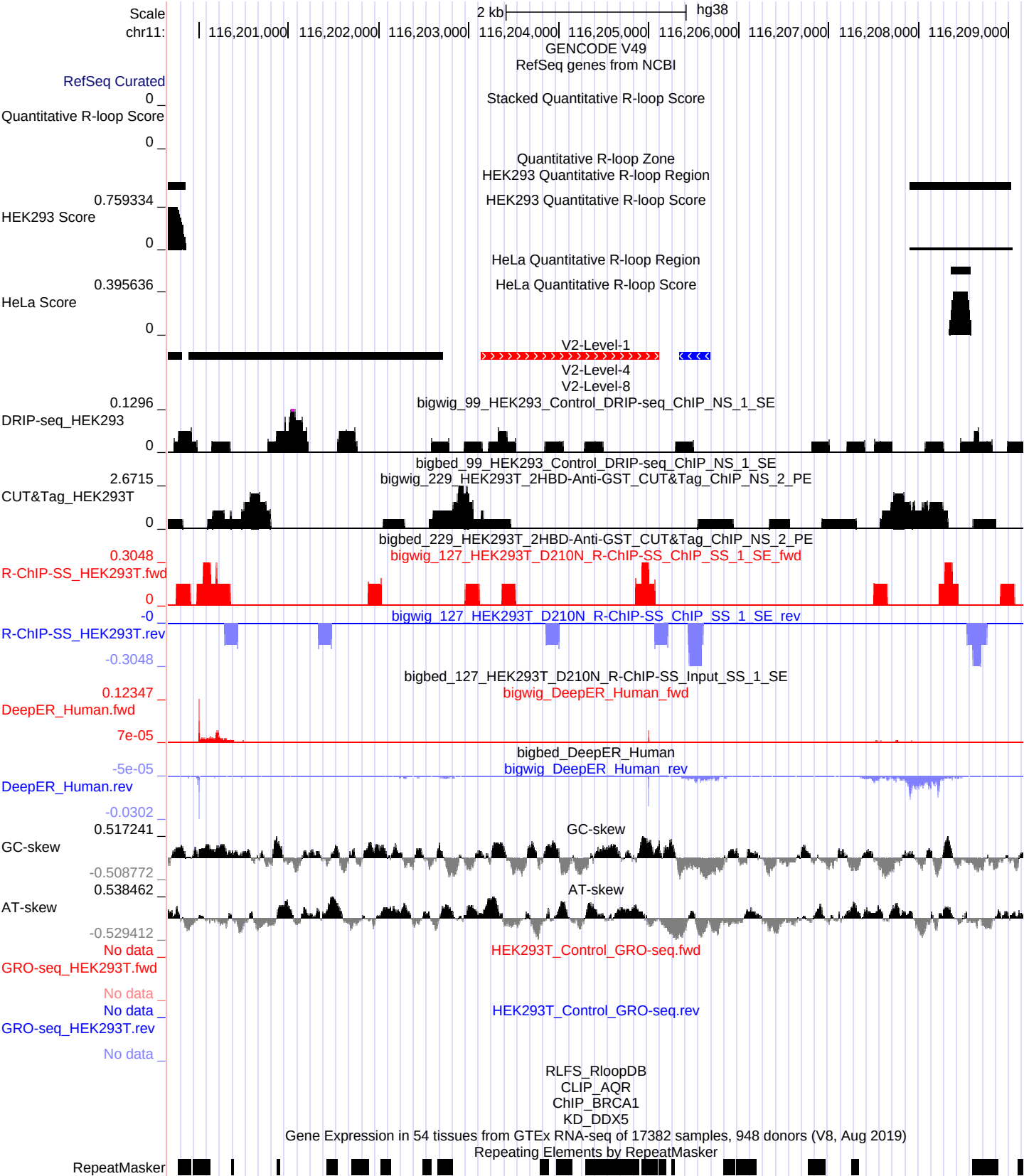

Scale
2 kb
hg38
chr11:
116,201,000
116,202,000
116,203,000
116,204,000
116,205,000
116,206,000
116,207,000
116,208,000
116,209,000
GENCODE V49
RefSeq genes from NCBI
RefSeq Curated
0
Stacked Quantitative R-loop Score
Quantitative R-loop Score
0
Quantitative R-loop Zone
HEK293 Quantitative R-loop Region
0.759334
HEK293 Quantitative R-loop Score
HEK293 Score
0
HeLa Quantitative R-loop Region
0.395636
HeLa Quantitative R-loop Score
HeLa Score
0
V2-Level-1
V2-Level-4
V2-Level-8
0.1296
bigwig_99_HEK293_Control_DRIP-seq_ChIP_NS_1_SE
DRIP-seq_HEK293
0
bigbed_99_HEK293_Control_DRIP-seq_ChIP_NS_1_SE
2.6715
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
-0.3048
bigbed_127_HEK293T_D210N_R-ChIP-SS_Input_SS_1_SE
0.12347
bigwig_DeepER_Human_fwd
DeepER_Human.fwd
7e-05
bigbed_DeepER_Human
-5e-05
bigwig_DeepER_Human_rev
DeepER_Human.rev
-0.0302
0.517241
GC-skew
GC-skew
-0.508772
0.538462
AT-skew
AT-skew
-0.529412
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)
Repeating Elements by RepeatMasker
RepeatMasker