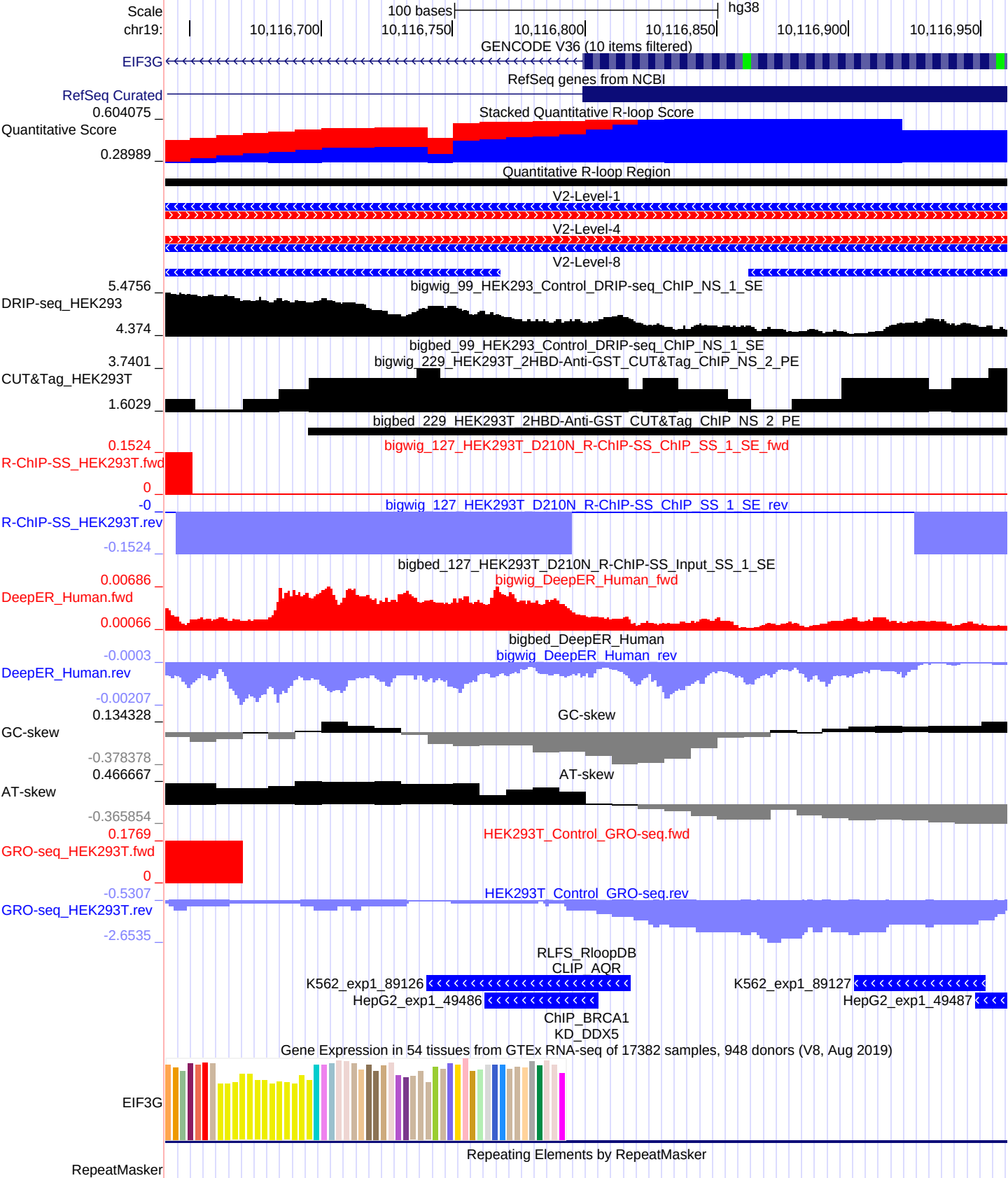
Scale
chr19:
100 bases
hg38
10,116,700
10,116,750
10,116,800
10,116,850
10,116,900
10,116,950
GENCODE V36 (10 items filtered)
EIF3G
RefSeq genes from NCBI
RefSeq Curated
0.604075
Stacked Quantitative R-loop Score
Quantitative Score
0.28989
Quantitative R-loop Region
V2-Level-1
V2-Level-4
V2-Level-8
5.4756
bigwig_99_HEK293_Control_DRIP-seq_ChIP_NS_1_SE
DRIP-seq_HEK293
4.374
bigbed_99_HEK293_Control_DRIP-seq_ChIP_NS_1_SE
3.7401
bigwig_229_HEK293T_2HBD-Anti-GST_CUT&Tag_ChIP_NS_2_PE
CUT&Tag_HEK293T
1.6029
bigbed_229_HEK293T_2HBD-Anti-GST_CUT&Tag_ChIP_NS_2_PE
0.1524
bigwig_127_HEK293T_D210N_R-ChIP-SS_ChIP_SS_1_SE_fwd
R-ChIP-SS_HEK293T.fwd
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-0
bigwig_127_HEK293T_D210N_R-ChIP-SS_ChIP_SS_1_SE_rev
R-ChIP-SS_HEK293T.rev
-0.1524
bigbed_127_HEK293T_D210N_R-ChIP-SS_Input_SS_1_SE
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bigwig_DeepER_Human_fwd
DeepER_Human.fwd
0.00066
bigbed_DeepER_Human
-0.0003
bigwig_DeepER_Human_rev
DeepER_Human.rev
-0.00207
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GC-skew
GC-skew
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AT-skew
AT-skew
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HEK293T_Control_GRO-seq.fwd
GRO-seq_HEK293T.fwd
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HEK293T_Control_GRO-seq.rev
GRO-seq_HEK293T.rev
-2.6535
RLFS_RloopDB
CLIP_AQR
K562_exp1_89126
K562_exp1_89127
HepG2_exp1_49486
HepG2_exp1_49487
ChIP_BRCA1
KD_DDX5
Gene Expression in 54 tissues from GTEx RNA-seq of 17382 samples, 948 donors (V8, Aug 2019)
EIF3G
Repeating Elements by RepeatMasker
RepeatMasker