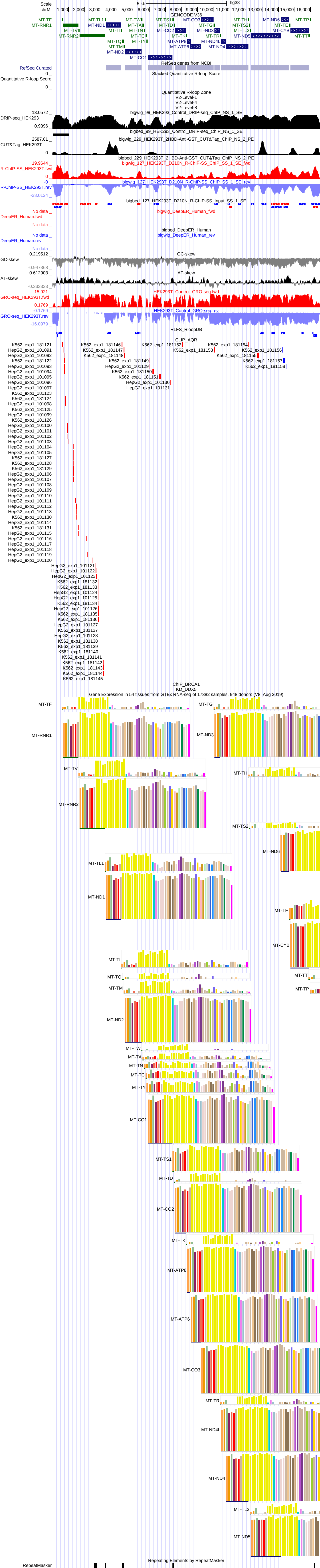
Scale
chrM:
5 kb
hg38
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GENCODE V36
MT-TF MT-RNR1 MT-TV MT-RNR2 MT-TL1 MT-ND1 MT-TI MT-TQ MT-TM MT-ND2 MT-TW MT-TA MT-TN MT-TC MT-TY MT-CO1 MT-TS1 MT-CO2 MT-TD MT-TK MT-ATP8 MT-ATP6 MT-CO3 MT-TG MT-ND3 MT-TR MT-ND4L MT-ND4 MT-TH MT-TS2 MT-TL2 MT-ND5 MT-ND6 MT-TE MT-CYB MT-TT MT-TP
RefSeq genes from NCBI
RefSeq Curated
Stacked Quantitative R-loop Score
Quantitative R-loop Score
Quantitative R-loop Zone
V2-Level-1
V2-Level-4
V2-Level-8
bigwig_99_HEK293_Control_DRIP-seq_ChIP_NS_1_SE
DRIP-seq_HEK293
bigbed_99_HEK293_Control_DRIP-seq_ChIP_NS_1_SE
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CUT&Tag_HEK293T
bigbed_229_HEK293T_2HBD-Anti-GST_CUT&Tag_ChIP_NS_2_PE
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R-ChIP-SS_HEK293T.fwd
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R-ChIP-SS_HEK293T.rev
bigbed_127_HEK293T_D210N_R-ChIP-SS_Input_SS_1_SE
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DeepER_Human.rev
GC-skew
AT-skew
HEK293T_Control_GRO-seq.fwd
GRO-seq_HEK293T.fwd
HEK293T_Control_GRO-seq.rev
GRO-seq_HEK293T.rev
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