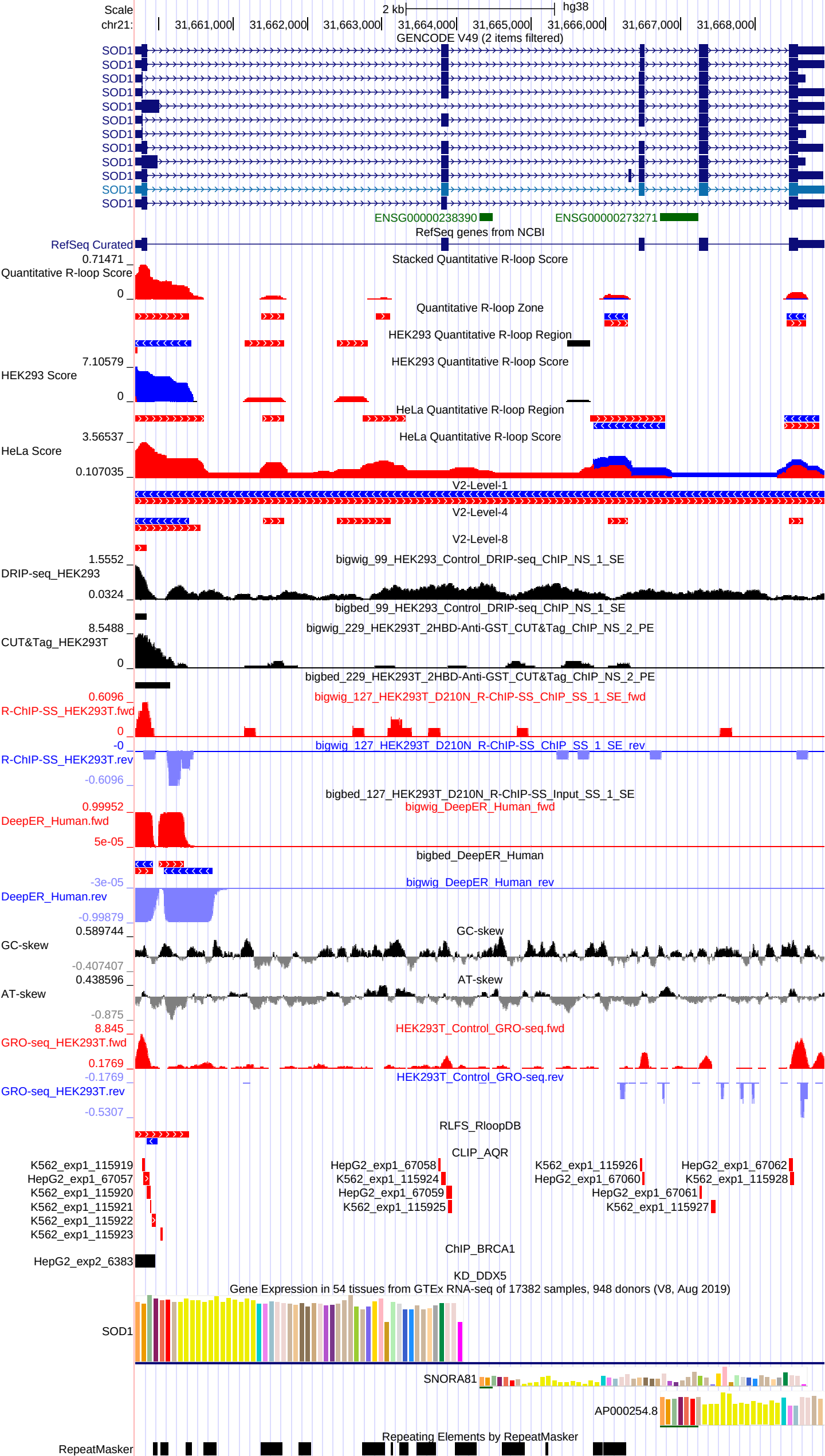

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GENCODE V49 (2 items filtered)
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ENSG00000273271
RefSeq genes from NCBI
RefSeq Curated
Stacked Quantitative R-loop Score
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Quantitative R-loop Score
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Quantitative R-loop Zone
HEK293 Quantitative R-loop Region
HEK293 Quantitative R-loop Score
7.10579
HEK293 Score
0
HeLa Quantitative R-loop Region
HeLa Quantitative R-loop Score
3.56537
HeLa Score
0.107035
V2-Level-1
V2-Level-4
V2-Level-8
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DRIP-seq_HEK293
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CUT&Tag_HEK293T
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bigbed_229_HEK293T_2HBD-Anti-GST_CUT&Tag_ChIP_NS_2_PE
bigwig_127_HEK293T_D210N_R-ChIP-SS_ChIP_SS_1_SE_fwd
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bigbed_DeepER_Human
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bigwig_DeepER_Human_rev
DeepER_Human.rev
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GC-skew
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GC-skew
-0.407407
AT-skew
0.438596
AT-skew
-0.875
HEK293T_Control_GRO-seq.fwd
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GRO-seq_HEK293T.rev
-0.5307
RLFS_RloopDB
CLIP_AQR
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ChIP_BRCA1
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KD_DDX5
Gene Expression in 54 tissues from GTEx RNA-seq of 17382 samples, 948 donors (V8, Aug 2019)
SOD1
SNORA81
AP000254.8
Repeating Elements by RepeatMasker
RepeatMasker