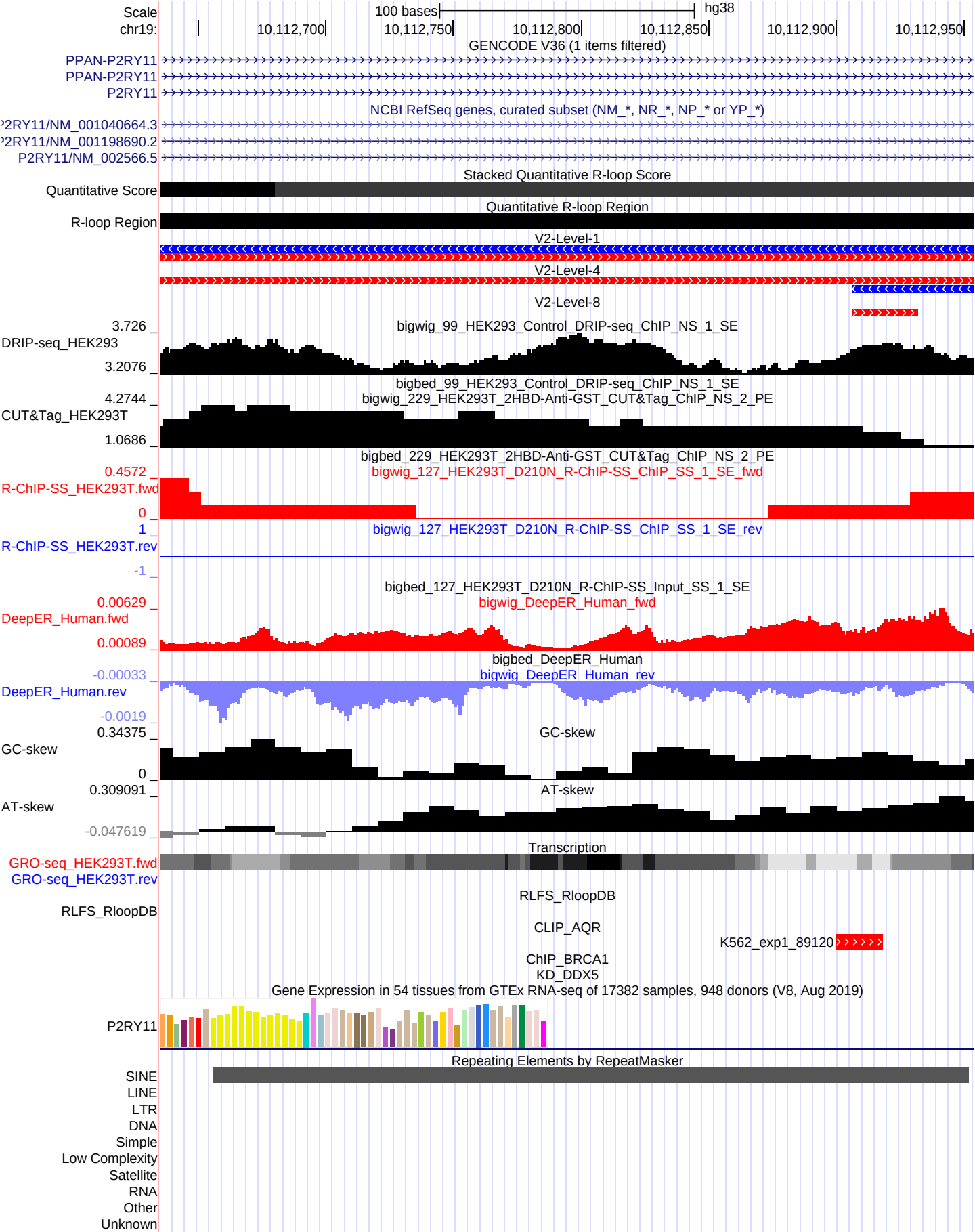
Scale
chr19:
100 bases
hg38
10,112,700
10,112,750
10,112,800
10,112,850
10,112,900
10,112,950
GENCODE V36 (1 items filtered)
PPAN-P2RY11
PPAN-P2RY11
P2RY11
NCBI RefSeq genes, curated subset (NM_*, NR_*, NP_* or YP_*)
P2RY11/NM_001040664.3
P2RY11/NM_001198690.2
P2RY11/NM_002566.5
Stacked Quantitative R-loop Score
Quantitative Score
Quantitative R-loop Region
R-loop Region
V2-Level-1
V2-Level-4
V2-Level-8
3.726
bigwig_99_HEK293_Control_DRIP-seq_ChIP_NS_1_SE
DRIP-seq_HEK293
3.2076
bigbed_99_HEK293_Control_DRIP-seq_ChIP_NS_1_SE
4.2744
bigwig_229_HEK293T_2HBD-Anti-GST_CUT&Tag_ChIP_NS_2_PE
CUT&Tag_HEK293T
1.0686
bigbed_229_HEK293T_2HBD-Anti-GST_CUT&Tag_ChIP_NS_2_PE
0.4572
bigwig_127_HEK293T_D210N_R-ChIP-SS_ChIP_SS_1_SE_fwd
R-ChIP-SS_HEK293T.fwd
0
1
bigwig_127_HEK293T_D210N_R-ChIP-SS_ChIP_SS_1_SE_rev
R-ChIP-SS_HEK293T.rev
-1
bigbed_127_HEK293T_D210N_R-ChIP-SS_Input_SS_1_SE
0.00629
bigwig_DeepER_Human_fwd
DeepER_Human.fwd
0.00089
bigbed_DeepER_Human
-0.00033
bigwig_DeepER_Human_rev
DeepER_Human.rev
-0.0019
0.34375
GC-skew
GC-skew
0
0.309091
AT-skew
AT-skew
-0.047619
Transcription
GRO-seq_HEK293T.fwd
GRO-seq_HEK293T.rev
RLFS_RloopDB
RLFS_RloopDB
CLIP_AQR
K562_exp1_89120
ChIP_BRCA1
KD_DDX5
Gene Expression in 54 tissues from GTEx RNA-seq of 17382 samples, 948 donors (V8, Aug 2019)
P2RY11
Repeating Elements by RepeatMasker
SINE
LINE
LTR
DNA
Simple
Low Complexity
Satellite
RNA
Other
Unknown