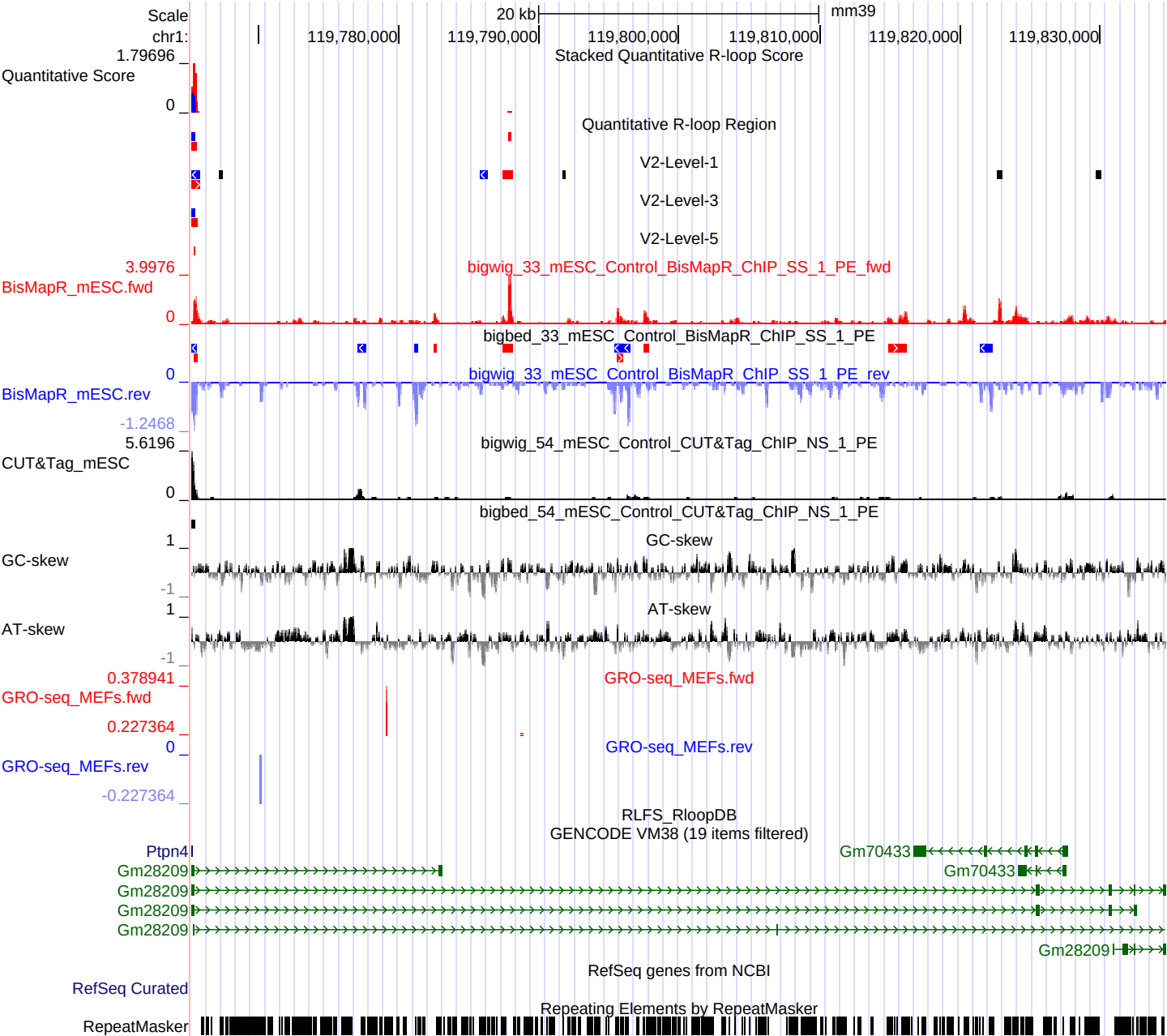

Scale
20 kb
mm39
chr1:
119,780,000
119,790,000
119,800,000
119,810,000
119,820,000
119,830,000
1.79696
Quantitative Score
0
Stacked Quantitative R-loop Score
Quantitative R-loop Region
V2-Level-1
V2-Level-3
V2-Level-5
3.9976
BisMapR_mESC.fwd
0
bigwig_33_mESC_Control_BisMapR_ChIP_SS_1_PE_fwd
bigbed_33_mESC_Control_BisMapR_ChIP_SS_1_PE
0
BisMapR_mESC.rev
-1.2468
bigwig_33_mESC_Control_BisMapR_ChIP_SS_1_PE_rev
5.6196
CUT&Tag_mESC
0
bigwig_54_mESC_Control_CUT&Tag_ChIP_NS_1_PE
bigbed_54_mESC_Control_CUT&Tag_ChIP_NS_1_PE
1
GC-skew
-1
GC-skew
1
AT-skew
-1
AT-skew
0.378941
GRO-seq_MEFs.fwd
GRO-seq_MEFs.fwd
0.227364
0
GRO-seq_MEFs.rev
GRO-seq_MEFs.rev
-0.227364
RLFS_RloopDB
GENCODE VM38 (19 items filtered)
Ptpn4
Gm28209
Gm70433
RefSeq genes from NCBI
RefSeq Curated
Repeating Elements by RepeatMasker
RepeatMasker