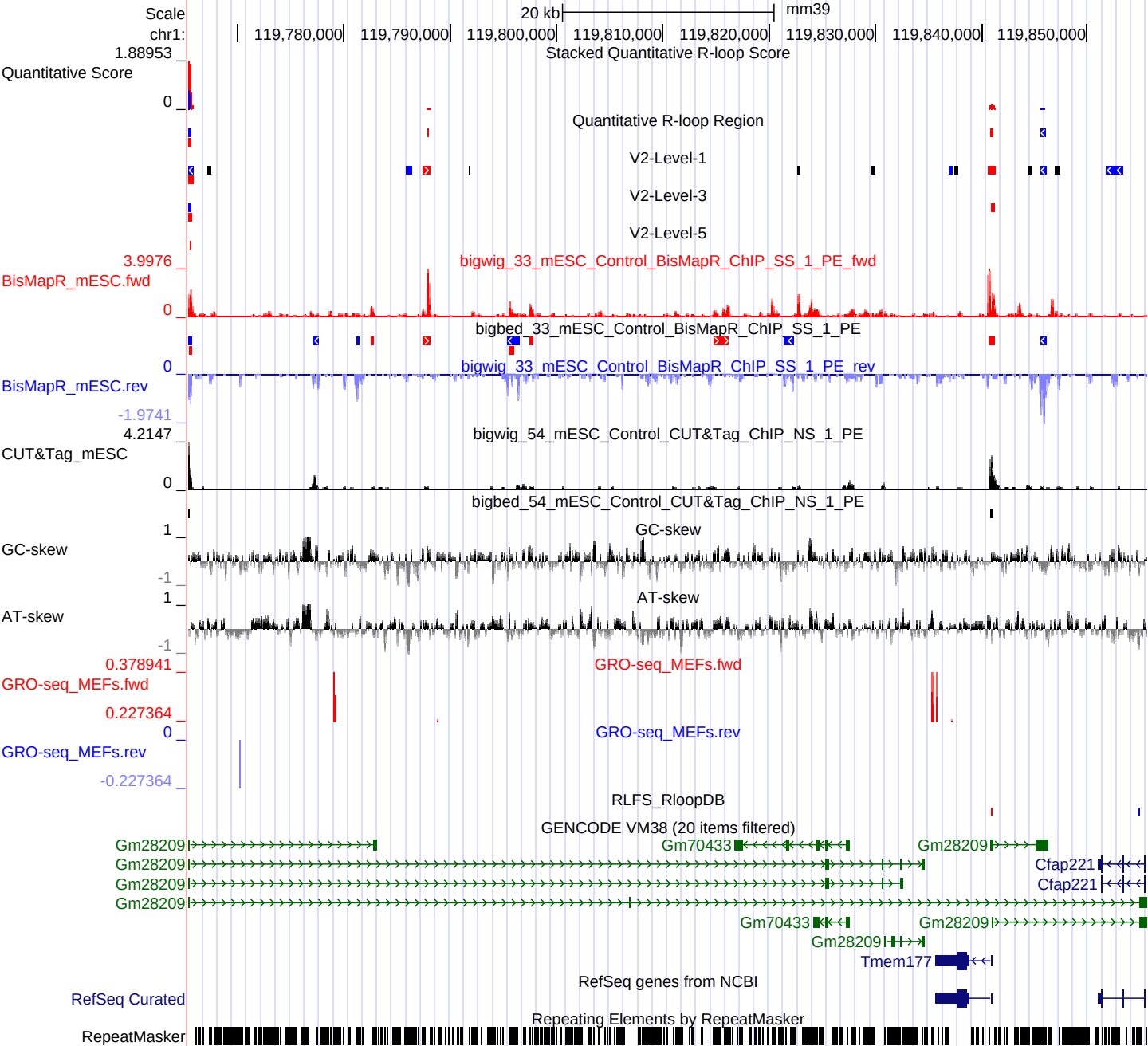
Scale
20 kb
mm39
chr1:
119,780,000
119,790,000
119,800,000
119,810,000
119,820,000
119,830,000
119,840,000
119,850,000
1.88953
Stacked Quantitative R-loop Score
Quantitative Score
0
Quantitative R-loop Region
V2-Level-1
V2-Level-3
V2-Level-5
3.9976
bigwig_33_mESC_Control_BisMapR_ChIP_SS_1_PE_fwd
BisMapR_mESC.fwd
0
bigbed_33_mESC_Control_BisMapR_ChIP_SS_1_PE
0
bigwig_33_mESC_Control_BisMapR_ChIP_SS_1_PE_rev
BisMapR_mESC.rev
-1.9741
4.2147
bigwig_54_mESC_Control_CUT&Tag_ChIP_NS_1_PE
CUT&Tag_mESC
0
bigbed_54_mESC_Control_CUT&Tag_ChIP_NS_1_PE
1
GC-skew
GC-skew
-1
1
AT-skew
AT-skew
-1
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 (20 items filtered)
Gm28209
Gm70433
Gm28209
Gm28209
Cfap221
Gm28209
Cfap221
Gm28209
Gm70433
Gm28209
Gm28209
Tmem177
RefSeq genes from NCBI
RefSeq Curated
Repeating Elements by RepeatMasker
RepeatMasker