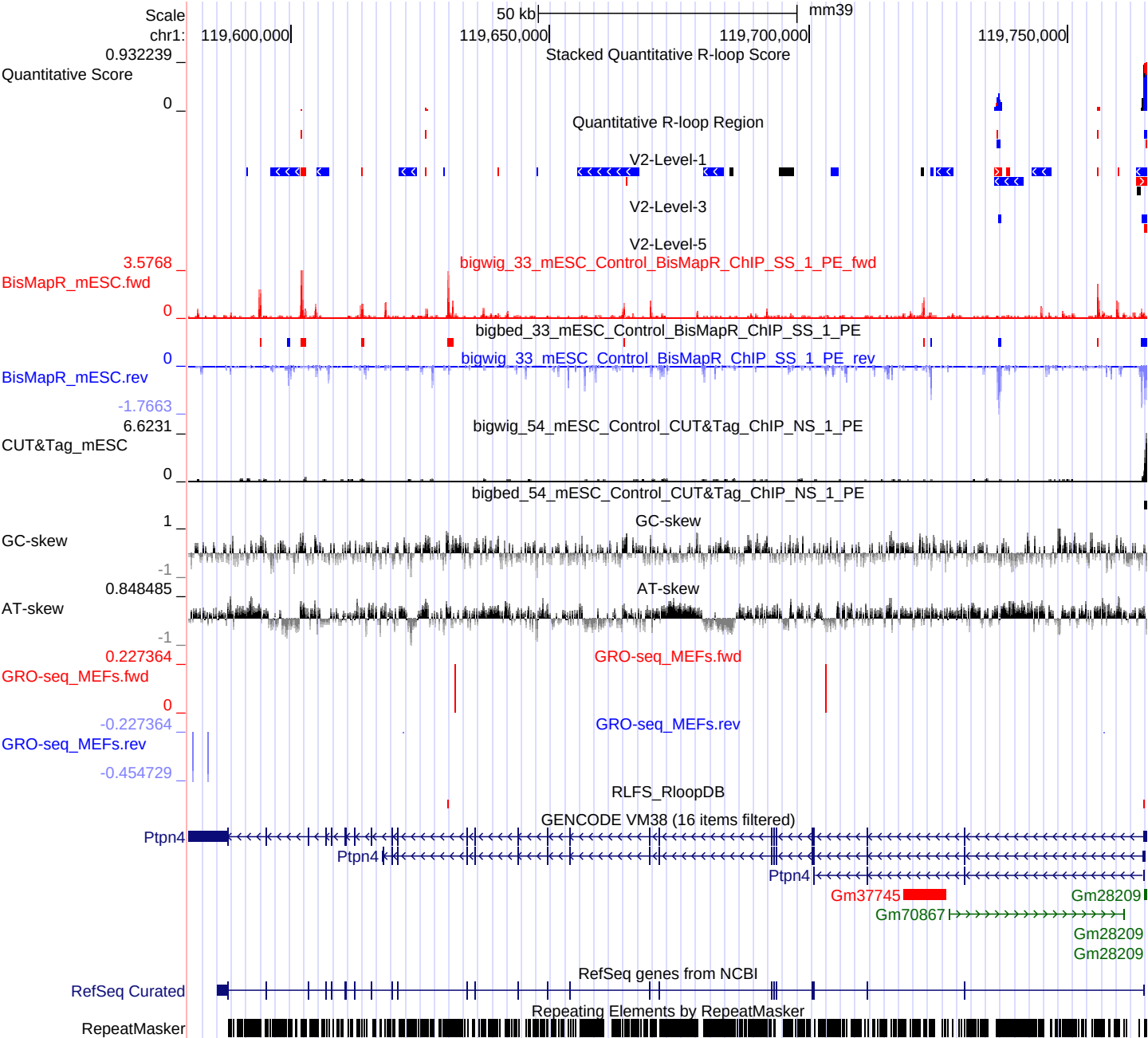

Scale
50 kb
mm39
chr1:
119,600,000
119,650,000
119,700,000
119,750,000
0.932239
Quantitative Score
0
Stacked Quantitative R-loop Score
Quantitative R-loop Region
V2-Level-1
V2-Level-3
V2-Level-5
3.5768
BisMapR_mESC.fwd
0
bigwig_33_mESC_Control_BisMapR_ChIP_SS_1_PE_fwd
bigbed_33_mESC_Control_BisMapR_ChIP_SS_1_PE
0
BisMapR_mESC.rev
bigwig_33_mESC_Control_BisMapR_ChIP_SS_1_PE_rev
-1.7663
6.6231
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
0.848485
AT-skew
-1
0.227364
GRO-seq_MEFs.fwd
0
GRO-seq_MEFs.fwd
-0.227364
GRO-seq_MEFs.rev
GRO-seq_MEFs.rev
-0.454729
RLFS_RloopDB
GENCODE VM38 (16 items filtered)
Ptpn4
Ptpn4
Ptpn4
Gm37745
Gm28209
Gm70867
Gm28209
Gm28209
RefSeq genes from NCBI
RefSeq Curated
Repeating Elements by RepeatMasker
RepeatMasker