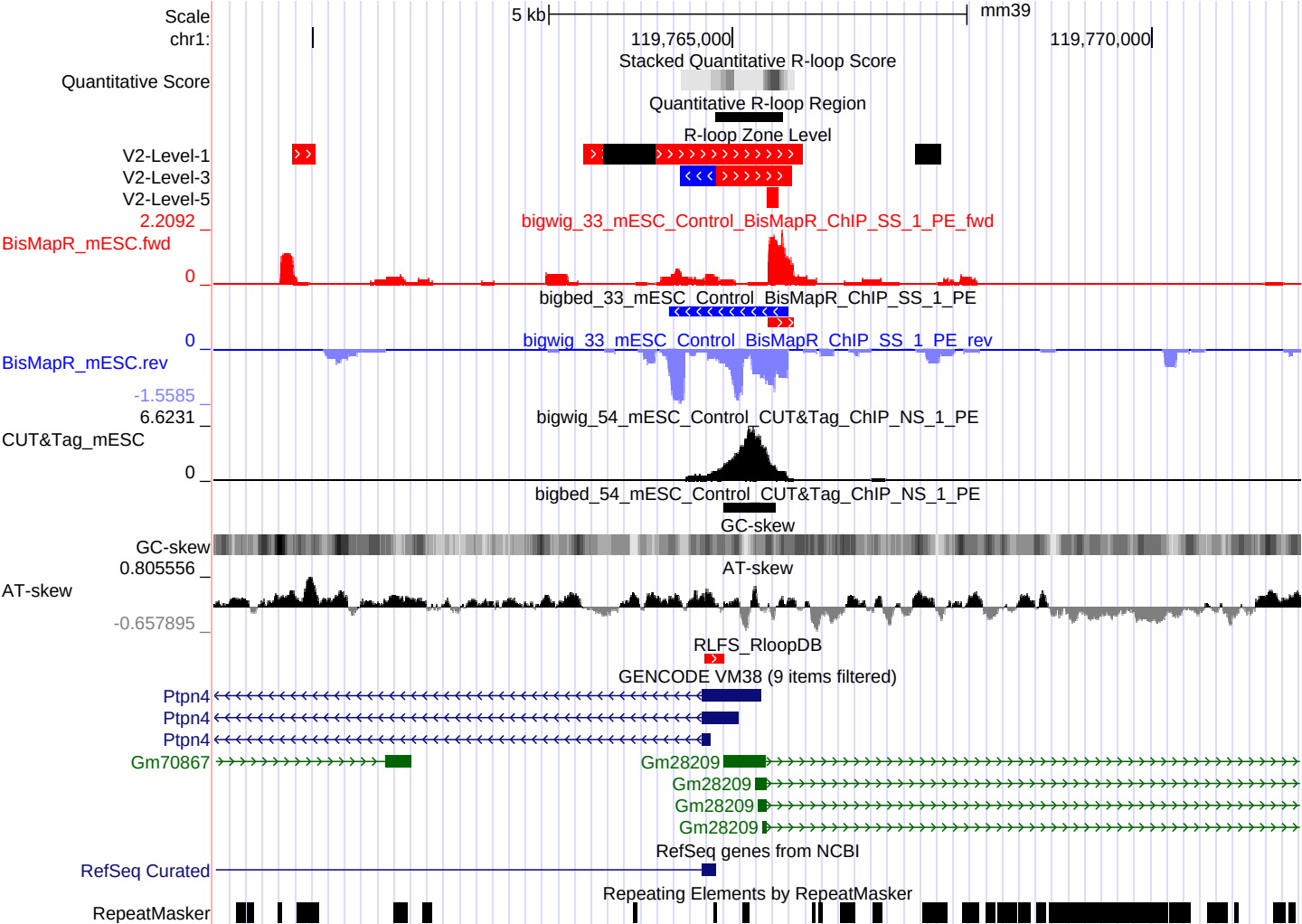
Scale
chr1:
5 kb
mm39
119,765,000
119,770,000
Stacked Quantitative R-loop Score
Quantitative Score
Quantitative R-loop Region
R-loop Zone Level
V2-Level-1
V2-Level-3
V2-Level-5
2.2092
bigwig_33_mESC_Control_BisMapR_ChIP_SS_1_PE_fwd
BisMapR_mESC.fwd
0
bigbed_33_mESC_Control_BisMapR_ChIP_SS_1_PE
0
bigwig_33_mESC_Control_BisMapR_ChIP_SS_1_PE_rev
BisMapR_mESC.rev
-1.5585
6.6231
bigwig_54_mESC_Control_CUT&Tag_ChIP_NS_1_PE
CUT&Tag_mESC
0
bigbed_54_mESC_Control_CUT&Tag_ChIP_NS_1_PE
GC-skew
GC-skew
0.805556
AT-skew
AT-skew
-0.657895
RLFS_RloopDB
GENCODE VM38 (9 items filtered)
Ptpn4
Ptpn4
Ptpn4
Gm70867
Gm28209
Gm28209
Gm28209
Gm28209
RefSeq genes from NCBI
RefSeq Curated
Repeating Elements by RepeatMasker
RepeatMasker