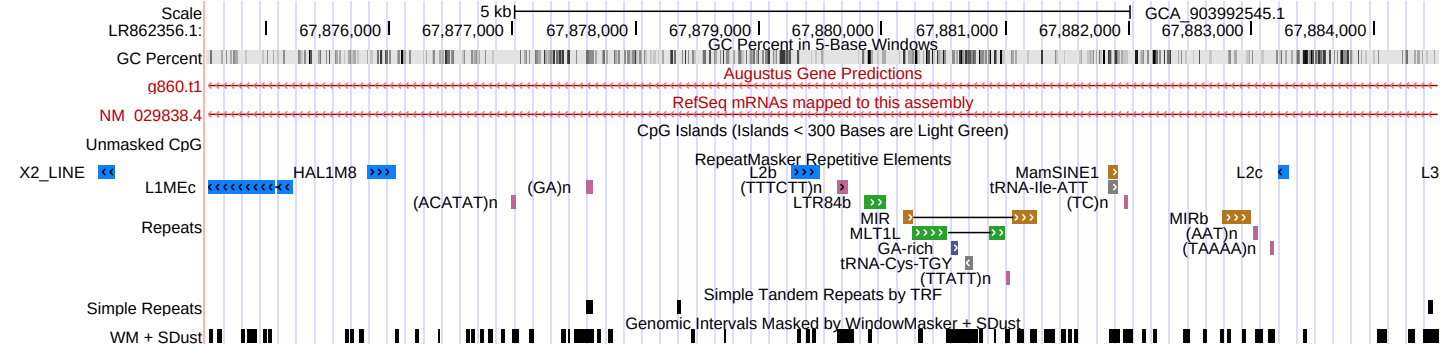
Scale
LR862356.1:
5 kb
GCA_903992545.1
67,876,000
67,877,000
67,878,000
67,879,000
67,880,000
67,881,000
67,882,000
67,883,000
67,884,000
GC Percent
GC Percent in 5-Base Windows
Augustus Gene Predictions
g860.t1
RefSeq mRNAs mapped to this assembly
NM_029838.4
Unmasked CpG
CpG Islands (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
Repeats
X2_LINE
L1MEc
HAL1M8
(ACATAT)n
(GA)n
L2b
(TTTCTT)n
LTR84b
MIR
MLT1L
GA-rich
tRNA-Cys-TGY
(TTATT)n
MamSINE1
tRNA-Ile-ATT
(TC)n
L2c
MIRb
(AAT)n
(TAAAA)n
Simple Repeats
Simple Tandem Repeats by TRF
WM + SDust
Genomic Intervals Masked by WindowMasker + SDust