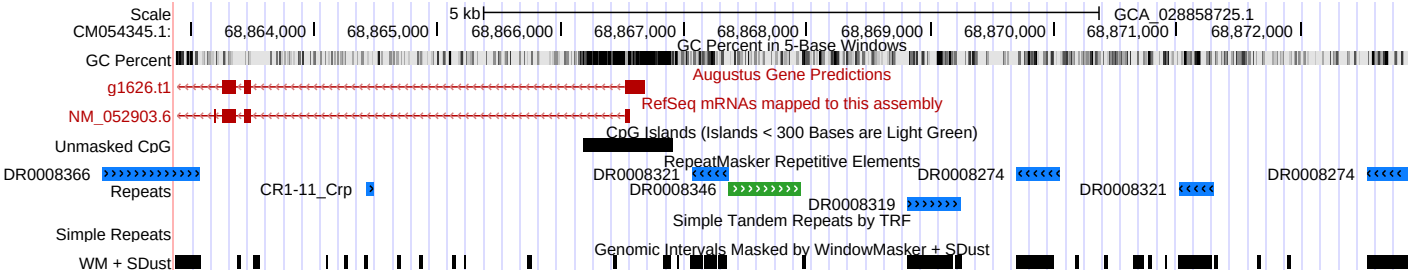
Scale
5 kb
GCA_028858725.1
CM054345.1:
68,864,000
68,865,000
68,866,000
68,867,000
68,868,000
68,869,000
68,870,000
68,871,000
68,872,000
GC Percent
GC Percent in 5-Base Windows
Augustus Gene Predictions
g1626.t1
RefSeq mRNAs mapped to this assembly
NM_052903.6
Unmasked CpG
CpG Islands (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
DR0008366
DR0008321
DR0008274
DR0008274
Repeats
CR1-11_Crp
DR0008346
DR0008321
DR0008319
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust