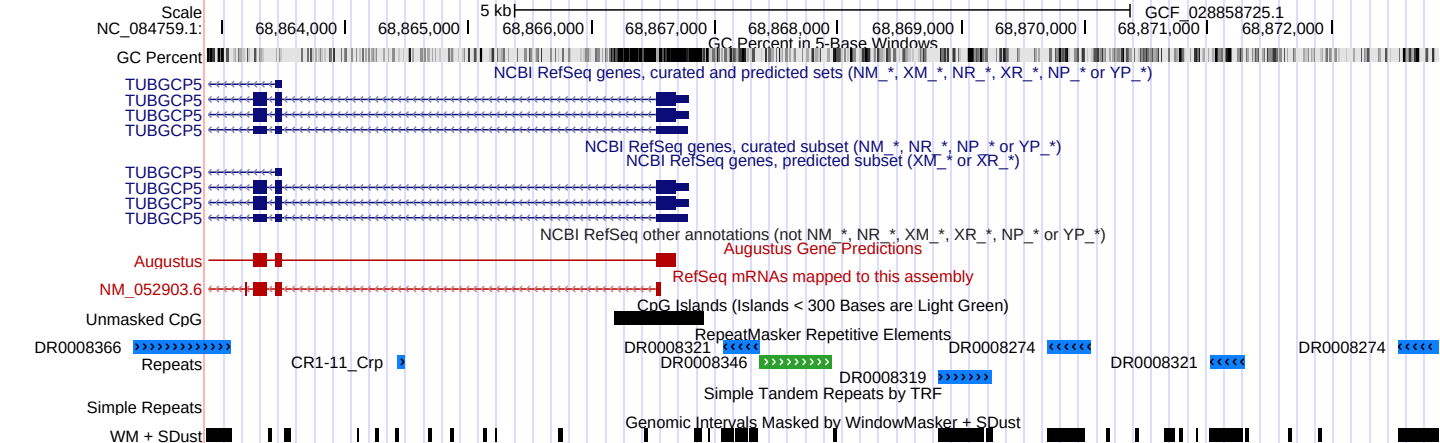
Scale
5 kb
GCF_028858725.1
NC_084759.1:
68,864,000
68,865,000
68,866,000
68,867,000
68,868,000
68,869,000
68,870,000
68,871,000
68,872,000
GC Percent
GC Percent in 5-Base Windows
NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)
TUBGCP5
TUBGCP5
TUBGCP5
TUBGCP5
NCBI RefSeq genes, curated subset (NM_*, NR_*, NP_* or YP_*)
NCBI RefSeq genes, predicted subset (XM_* or XR_*)
TUBGCP5
TUBGCP5
TUBGCP5
TUBGCP5
NCBI RefSeq other annotations (not NM_*, NR_*, XM_*, XR_*, NP_* or YP_*)
Augustus
Augustus Gene Predictions
NM_052903.6
RefSeq mRNAs mapped to this assembly
Unmasked CpG
CpG Islands (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
DR0008366
Repeats
CR1-11_Crp
DR0008321
DR0008346
DR0008319
DR0008274
DR0008321
DR0008274
Simple Repeats
Simple Tandem Repeats by TRF
WM + SDust
Genomic Intervals Masked by WindowMasker + SDust