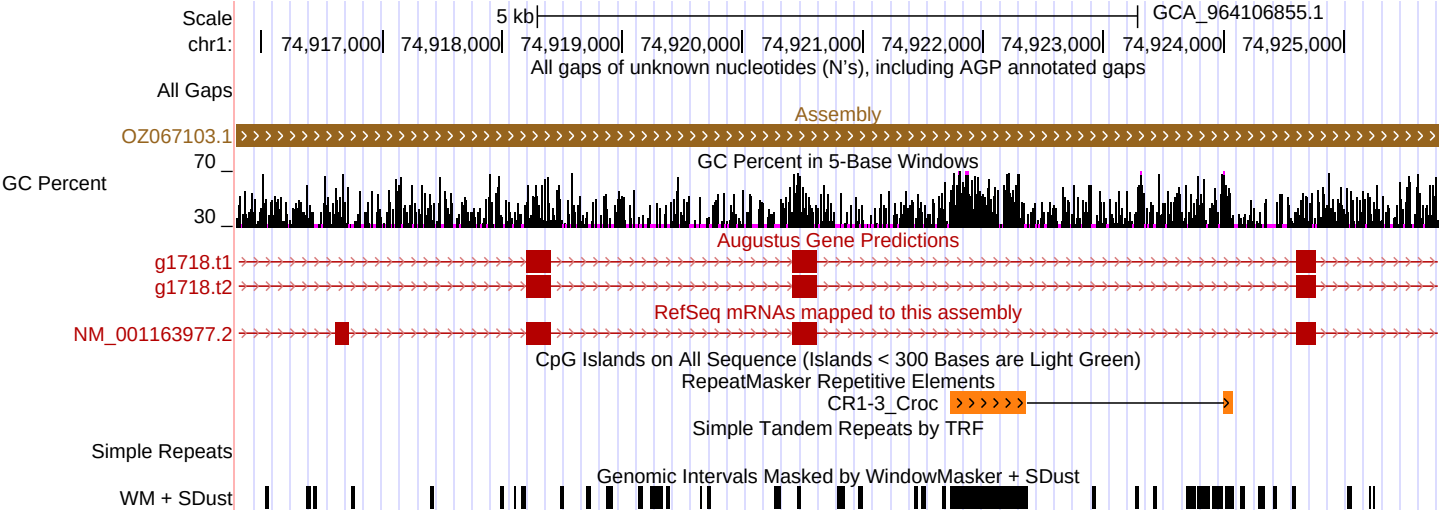
Scale
5 kb
GCA_964106855.1
chr1:
74,917,000
74,918,000
74,919,000
74,920,000
74,921,000
74,922,000
74,923,000
74,924,000
74,925,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
OZ067103.1
GC Percent in 5-Base Windows
70
GC Percent
30
Augustus Gene Predictions
g1718.t1
g1718.t2
RefSeq mRNAs mapped to this assembly
NM_001163977.2
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
CR1-3_Croc
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust