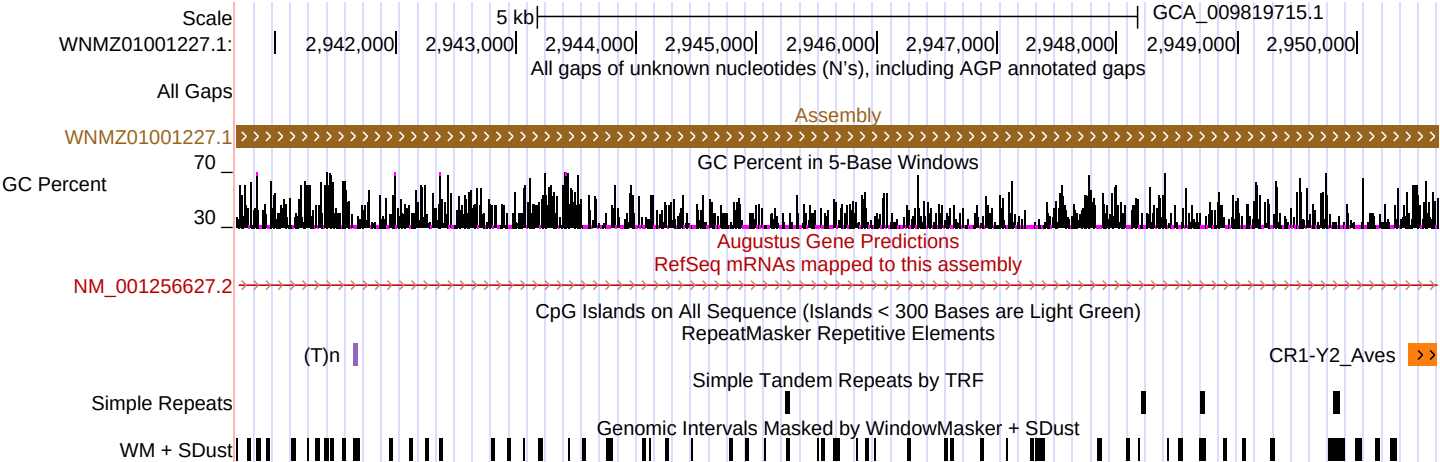
Scale
5 kb
GCA_009819715.1
WNMZ01001227.1:
2,942,000
2,943,000
2,944,000
2,945,000
2,946,000
2,947,000
2,948,000
2,949,000
2,950,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
WNMZ01001227.1
GC Percent in 5-Base Windows
GC Percent
70
30
Augustus Gene Predictions
RefSeq mRNAs mapped to this assembly
NM_001256627.2
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
(T)n
CR1-Y2_Aves
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust