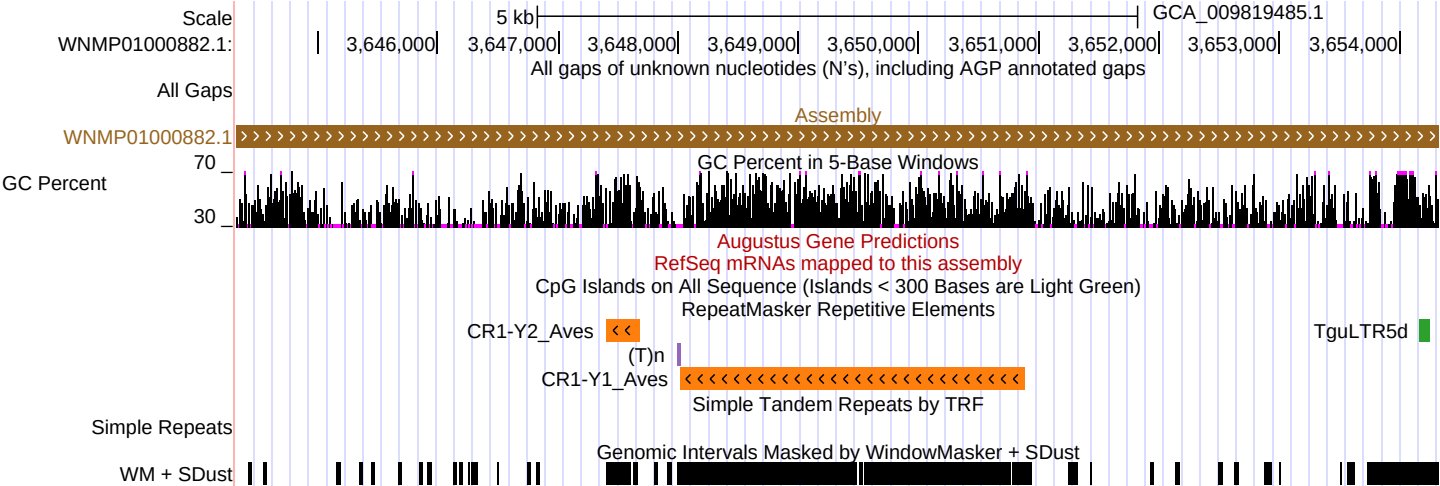
Scale
5 kb
GCA_009819485.1
WNMP01000882.1:
3,646,000
3,647,000
3,648,000
3,649,000
3,650,000
3,651,000
3,652,000
3,653,000
3,654,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
WNMP01000882.1
GC Percent in 5-Base Windows
GC Percent
70
30
Augustus Gene Predictions
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
CR1-Y2_Aves
TguLTR5d
(T)n
CR1-Y1_Aves
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust