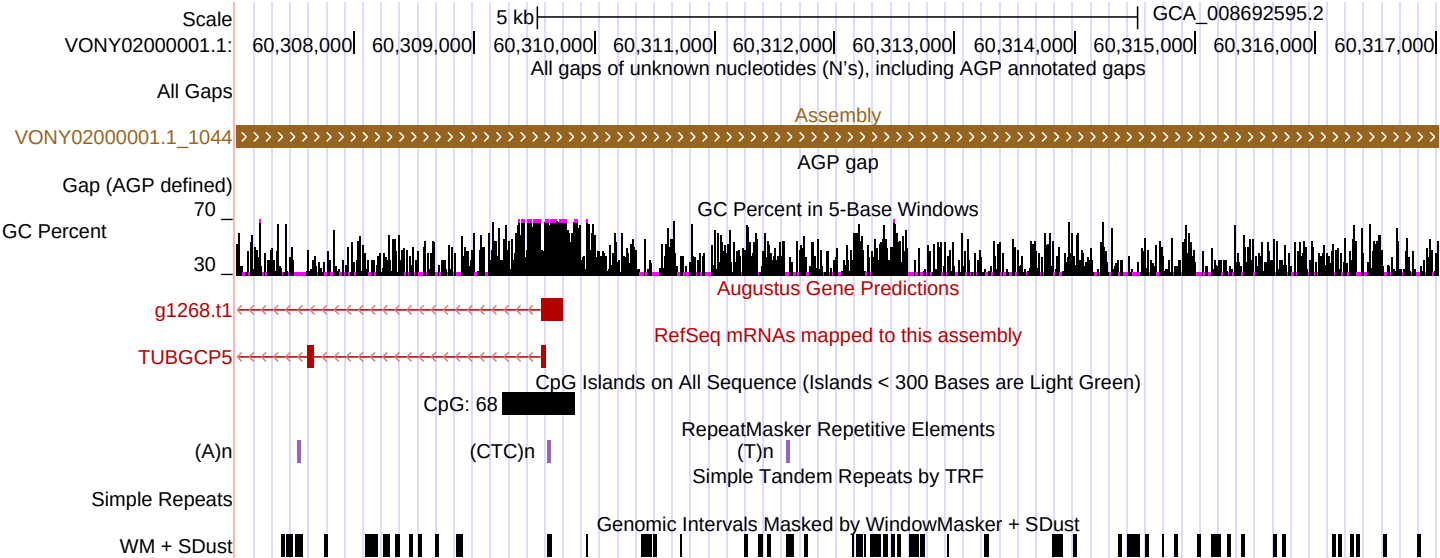
Scale
5 kb
GCA_008692595.2
VONY02000001.1:
60,308,000
60,309,000
60,310,000
60,311,000
60,312,000
60,313,000
60,314,000
60,315,000
60,316,000
60,317,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
VONY02000001.1_1044
AGP gap
Gap (AGP defined)
GC Percent in 5-Base Windows
70
GC Percent
30
Augustus Gene Predictions
g1268.t1
RefSeq mRNAs mapped to this assembly
TUBGCP5
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
CpG: 68
RepeatMasker Repetitive Elements
(A)n
(CTC)n
(T)n
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust