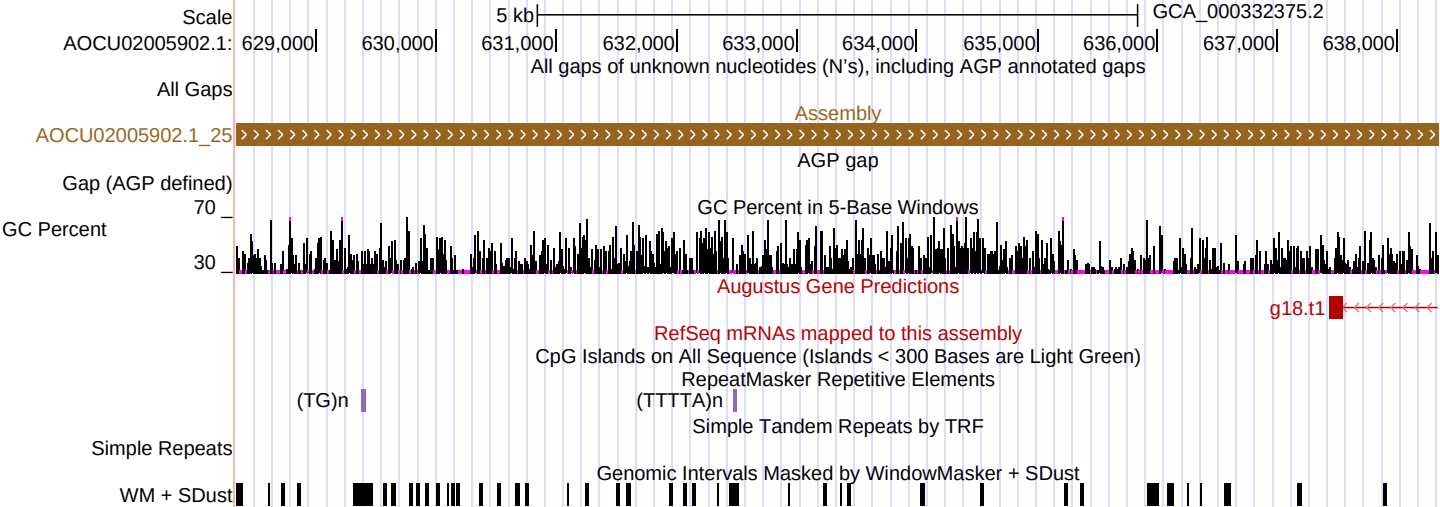
Scale
5 kb
GCA_000332375.2
AOCU02005902.1:
629,000
630,000
631,000
632,000
633,000
634,000
635,000
636,000
637,000
638,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
AOCU02005902.1_25
AGP gap
Gap (AGP defined)
GC Percent in 5-Base Windows
70
GC Percent
30
Augustus Gene Predictions
g18.t1
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
(TG)n
(TTTTA)n
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust