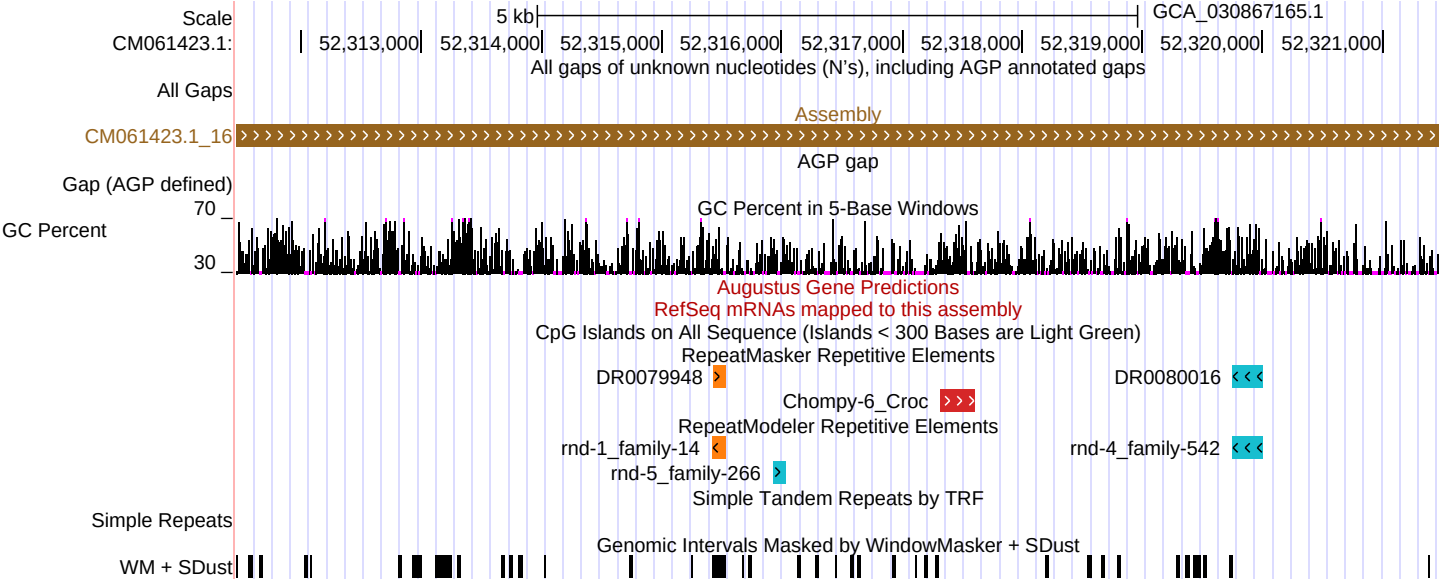
Scale
5 kb
GCA_030867165.1
CM061423.1:
52,313,000
52,314,000
52,315,000
52,316,000
52,317,000
52,318,000
52,319,000
52,320,000
52,321,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
CM061423.1_16
AGP gap
Gap (AGP defined)
GC Percent in 5-Base Windows
70
GC Percent
30
Augustus Gene Predictions
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
DR0079948
DR0080016
Chompy-6_Croc
RepeatModeler Repetitive Elements
rnd-1_family-14
rnd-4_family-542
rnd-5_family-266
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust