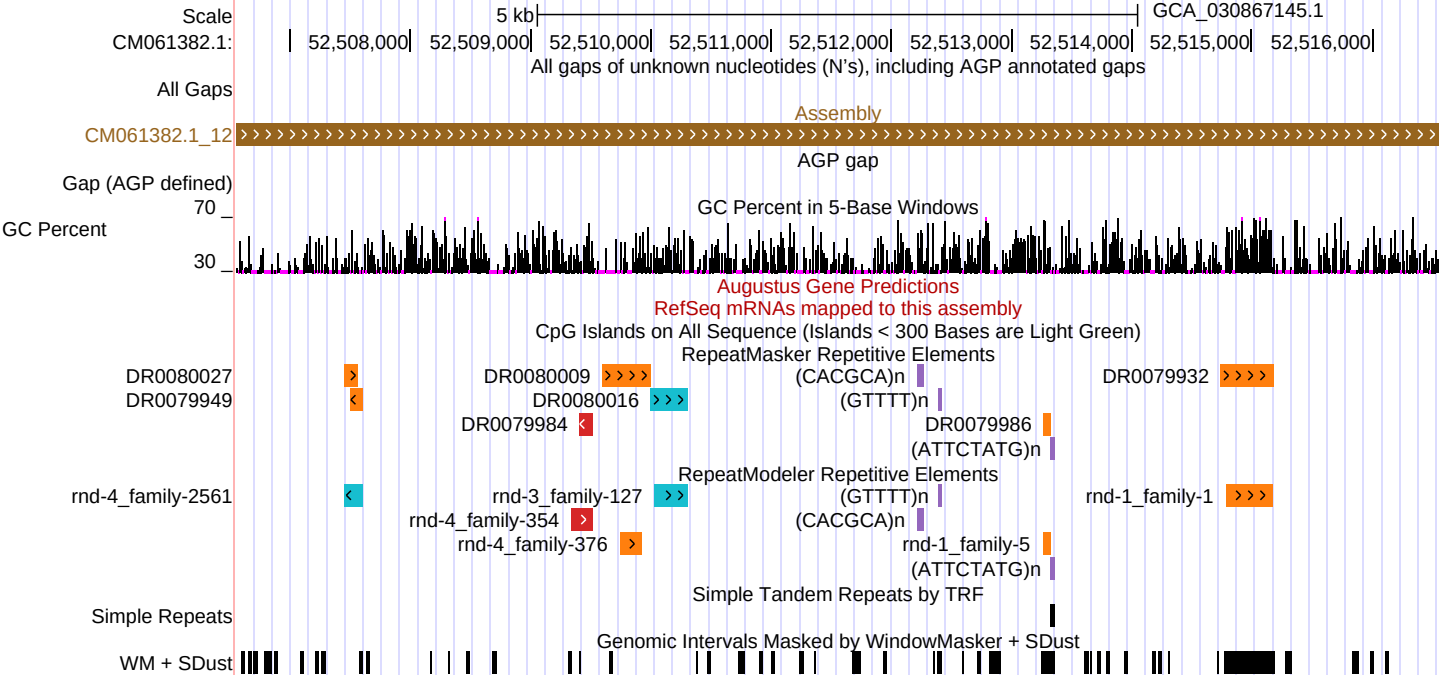
Scale
5 kb
GCA_030867145.1
CM061382.1:
52,508,000
52,509,000
52,510,000
52,511,000
52,512,000
52,513,000
52,514,000
52,515,000
52,516,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
CM061382.1_12
AGP gap
Gap (AGP defined)
GC Percent in 5-Base Windows
70
GC Percent
30
Augustus Gene Predictions
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
DR0080027
DR0079949
DR0080009
DR0080016
DR0079984
(CACGCA)n
(GTTTT)n
DR0079986
(ATTCTATG)n
DR0079932
RepeatModeler Repetitive Elements
rnd-4_family-2561
rnd-3_family-127
rnd-4_family-354
rnd-4_family-376
(GTTTT)n
(CACGCA)n
rnd-1_family-5
(ATTCTATG)n
rnd-1_family-1
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust