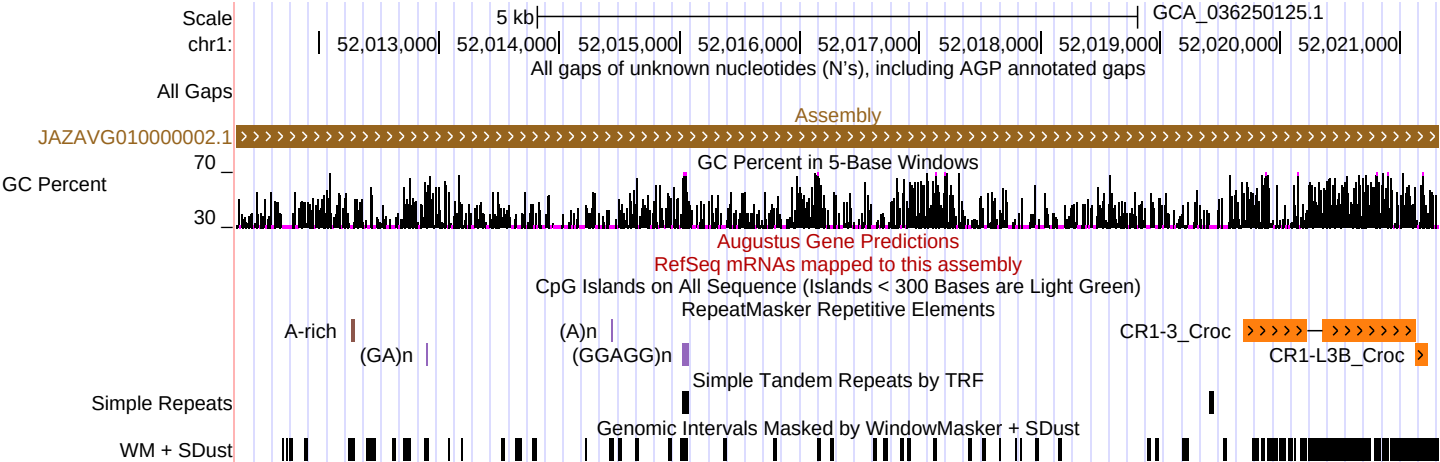
Scale
5 kb
GCA_036250125.1
chr1:
52,013,000
52,014,000
52,015,000
52,016,000
52,017,000
52,018,000
52,019,000
52,020,000
52,021,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
JAZAVG010000002.1
70
GC Percent in 5-Base Windows
GC Percent
30
Augustus Gene Predictions
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
A-rich
(A)n
CR1-3_Croc
(GA)n
(GGAGG)n
CR1-L3B_Croc
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust