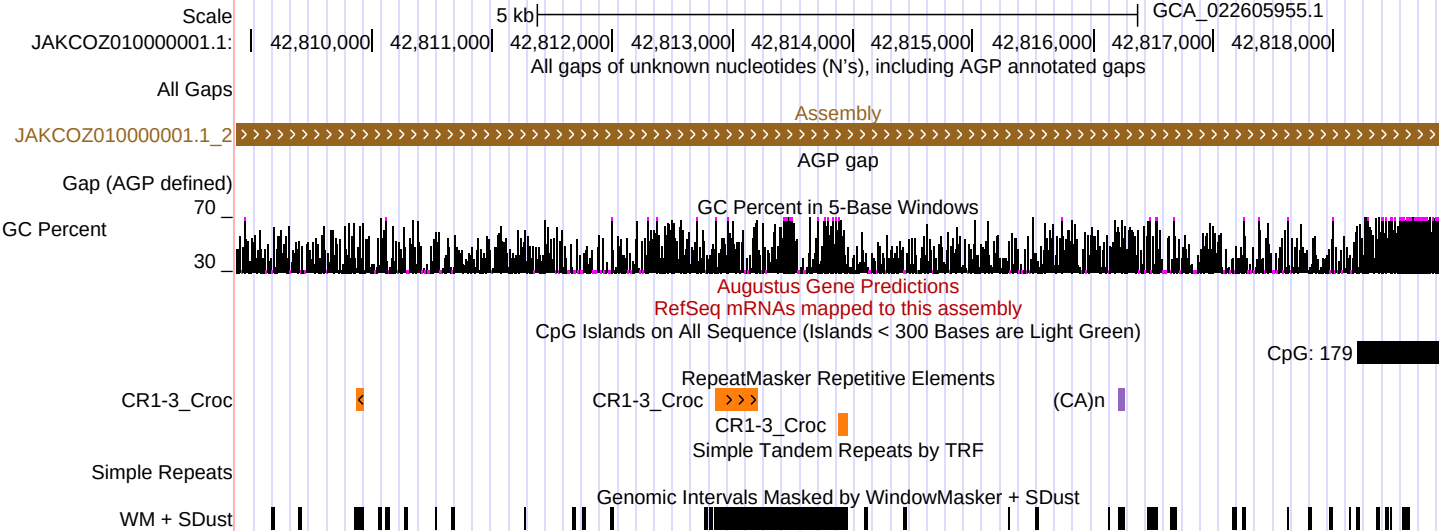
Scale
5 kb
GCA_022605955.1
JAKCOZ010000001.1:
42,810,000
42,811,000
42,812,000
42,813,000
42,814,000
42,815,000
42,816,000
42,817,000
42,818,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
JAKCOZ010000001.1_2
AGP gap
Gap (AGP defined)
GC Percent in 5-Base Windows
70
GC Percent
30
Augustus Gene Predictions
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
CpG: 179
RepeatMasker Repetitive Elements
CR1-3_Croc
CR1-3_Croc
(CA)n
CR1-3_Croc
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust