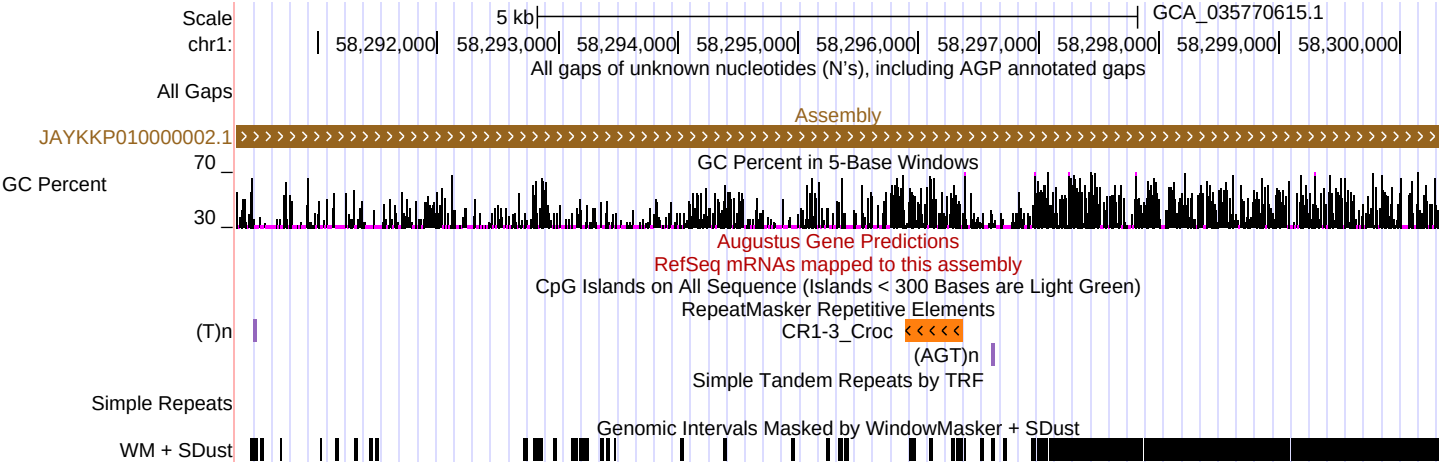
Scale
5 kb
GCA_035770615.1
chr1:
58,292,000
58,293,000
58,294,000
58,295,000
58,296,000
58,297,000
58,298,000
58,299,000
58,300,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
JAYKKP010000002.1
GC Percent in 5-Base Windows
70
GC Percent
30
Augustus Gene Predictions
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
(T)n
CR1-3_Croc
(AGT)n
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust