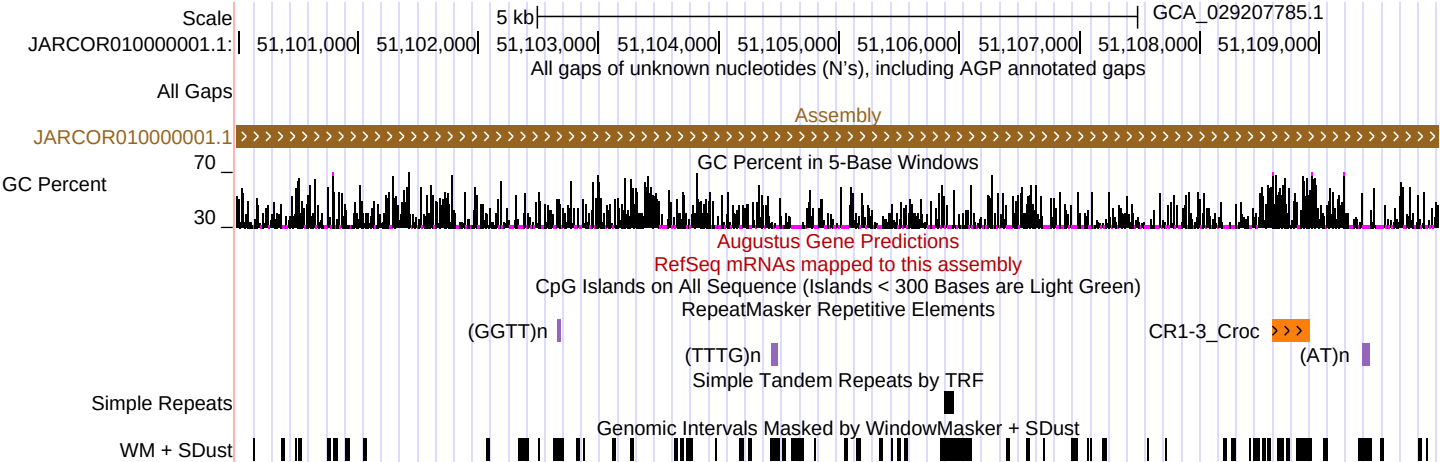
Scale
5 kb
GCA_029207785.1
JARCOR010000001.1:
51,101,000
51,102,000
51,103,000
51,104,000
51,105,000
51,106,000
51,107,000
51,108,000
51,109,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
JARCOR010000001.1
GC Percent in 5-Base Windows
70
GC Percent
30
Augustus Gene Predictions
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
(GGTT)n
CR1-3_Croc
(TTTG)n
(AT)n
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust