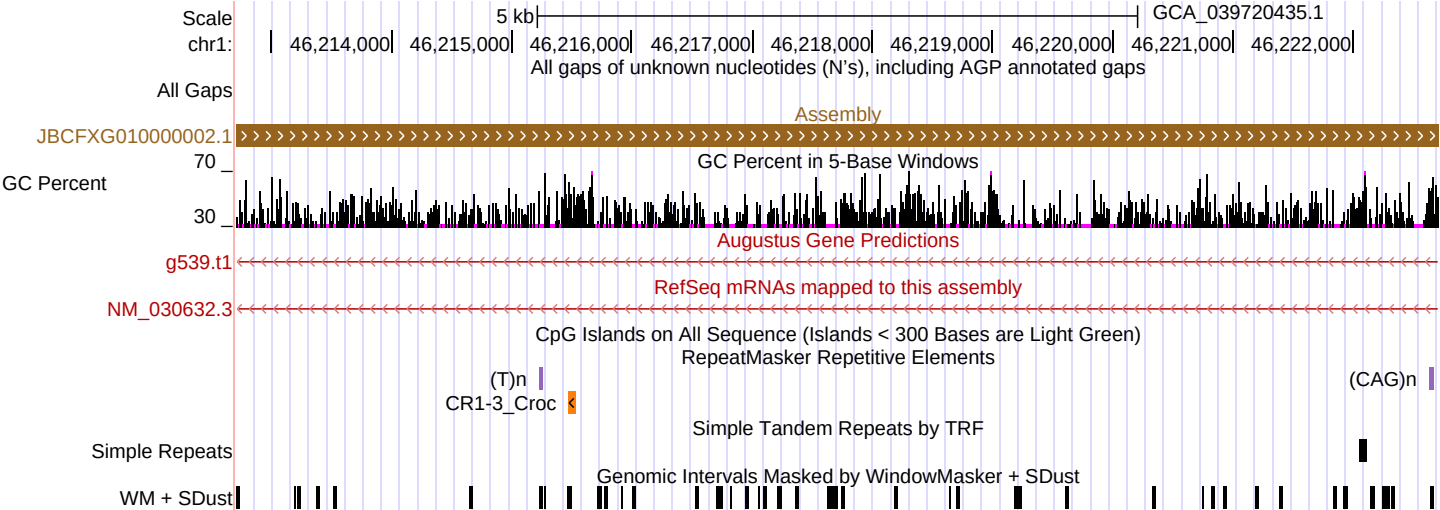
Scale
5 kb
GCA_039720435.1
chr1:
46,214,000
46,215,000
46,216,000
46,217,000
46,218,000
46,219,000
46,220,000
46,221,000
46,222,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
JBCFXG010000002.1
GC Percent in 5-Base Windows
70
GC Percent
30
Augustus Gene Predictions
g539.t1
RefSeq mRNAs mapped to this assembly
NM_030632.3
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
(T)n
(CAG)n
CR1-3_Croc
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust