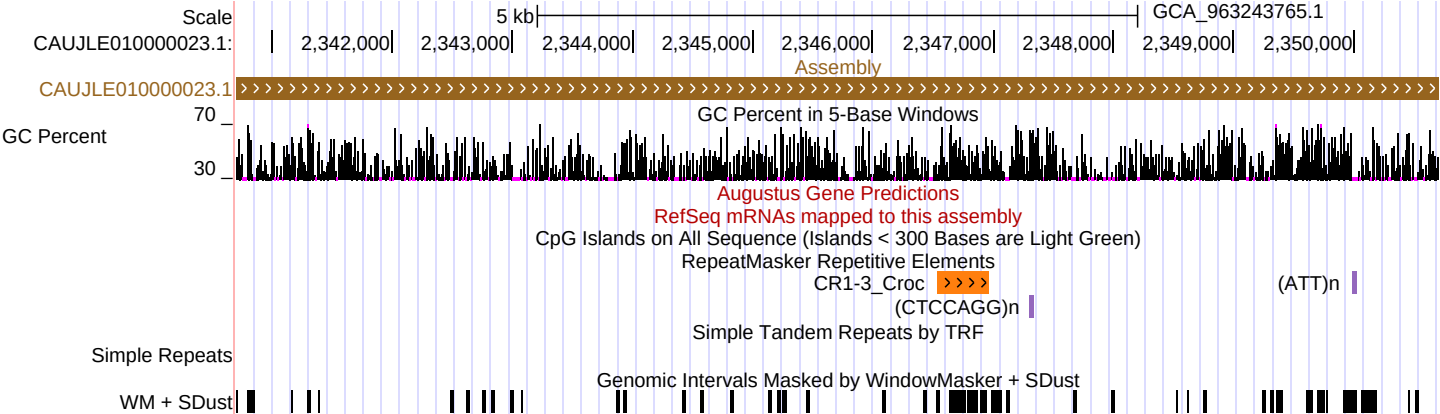
Scale
5 kb
GCA_963243765.1
CAUJLE010000023.1:
2,342,000
2,343,000
2,344,000
2,345,000
2,346,000
2,347,000
2,348,000
2,349,000
2,350,000
Assembly
CAUJLE010000023.1
70
GC Percent
30
GC Percent in 5-Base Windows
Augustus Gene Predictions
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
CR1-3_Croc
(ATT)n
(CTCCAGG)n
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust