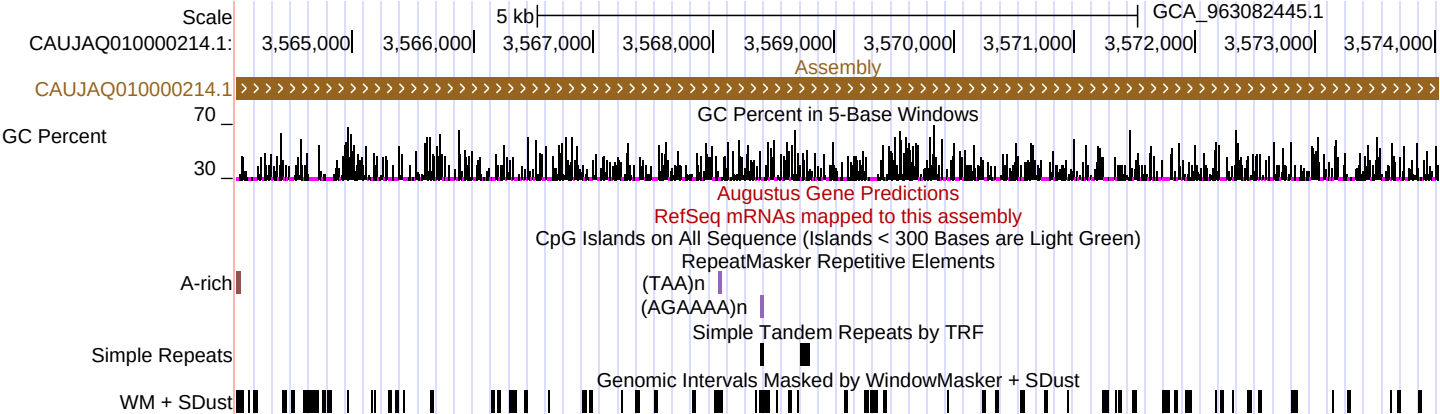
Scale
5 kb
GCA_963082445.1
CAUJAQ010000214.1:
3,565,000
3,566,000
3,567,000
3,568,000
3,569,000
3,570,000
3,571,000
3,572,000
3,573,000
3,574,000
Assembly
CAUJAQ010000214.1
GC Percent
70
30
GC Percent in 5-Base Windows
Augustus Gene Predictions
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
A-rich
(TAA)n
(AGAAAA)n
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust