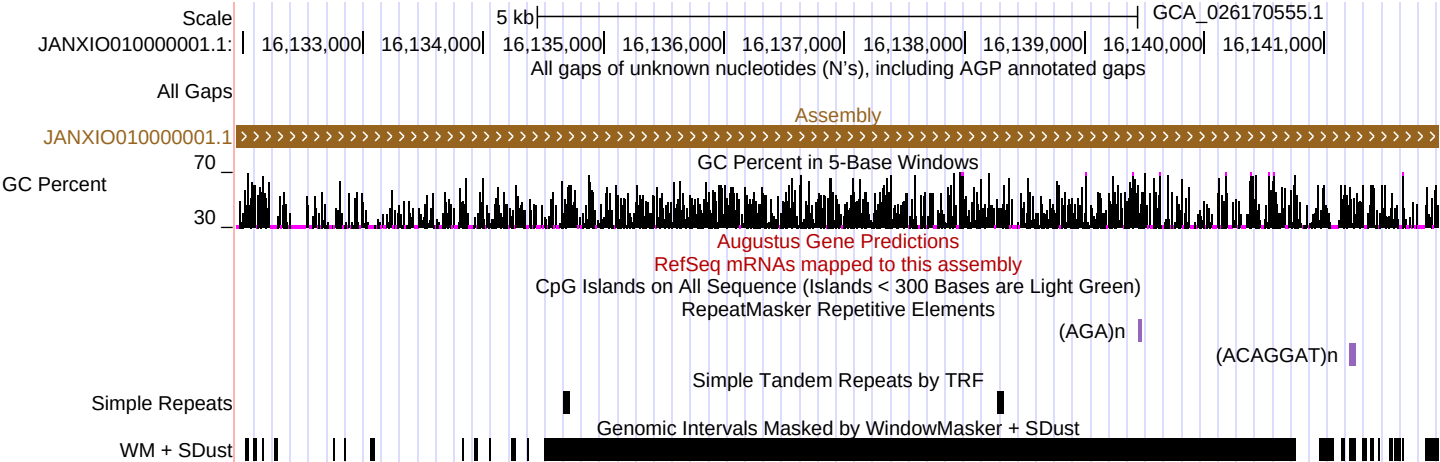
Scale
5 kb
GCA_026170555.1
JANXIO010000001.1:
16,133,000
16,134,000
16,135,000
16,136,000
16,137,000
16,138,000
16,139,000
16,140,000
16,141,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
JANXIO010000001.1
GC Percent in 5-Base Windows
70
GC Percent
30
Augustus Gene Predictions
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
(AGA)n
(ACAGGAT)n
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust