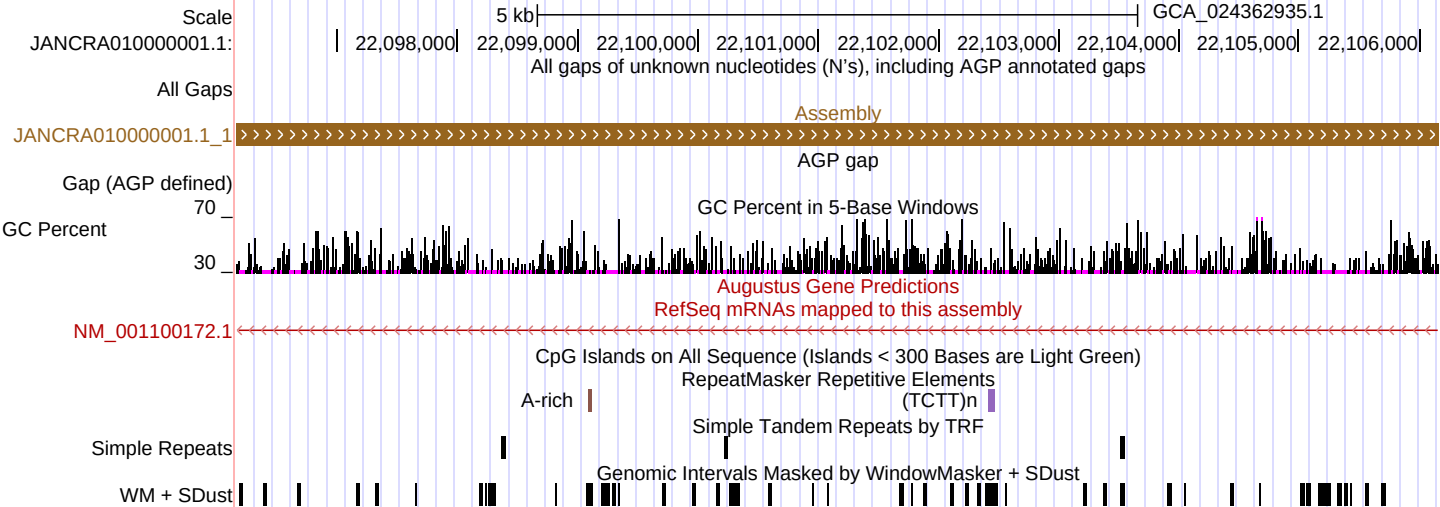
Scale
5 kb
GCA_024362935.1
JANCRA010000001.1:
22,098,000
22,099,000
22,100,000
22,101,000
22,102,000
22,103,000
22,104,000
22,105,000
22,106,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
JANCRA010000001.1_1
AGP gap
Gap (AGP defined)
GC Percent in 5-Base Windows
70
GC Percent
30
Augustus Gene Predictions
RefSeq mRNAs mapped to this assembly
NM_001100172.1
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
A-rich
(TCTT)n
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust