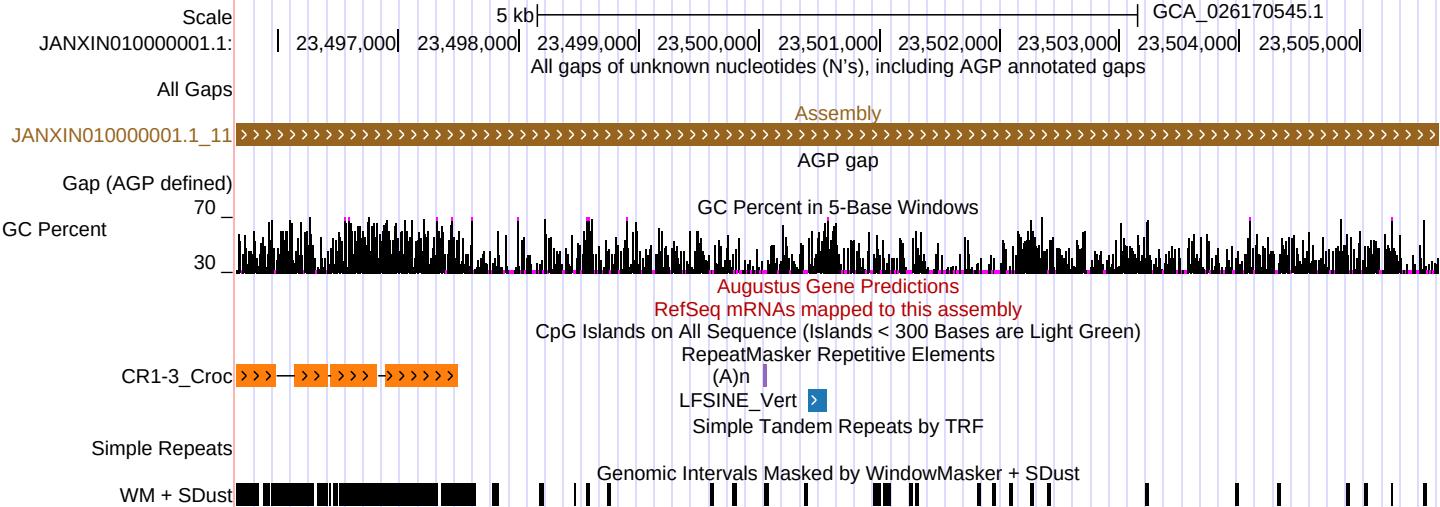
Scale
5 kb
GCA_026170545.1
JANXIN010000001.1:
23,497,000
23,498,000
23,499,000
23,500,000
23,501,000
23,502,000
23,503,000
23,504,000
23,505,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
JANXIN010000001.1_11
AGP gap
Gap (AGP defined)
GC Percent in 5-Base Windows
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
CR1-3_Croc
(A)n
LFSINE_Vert
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust