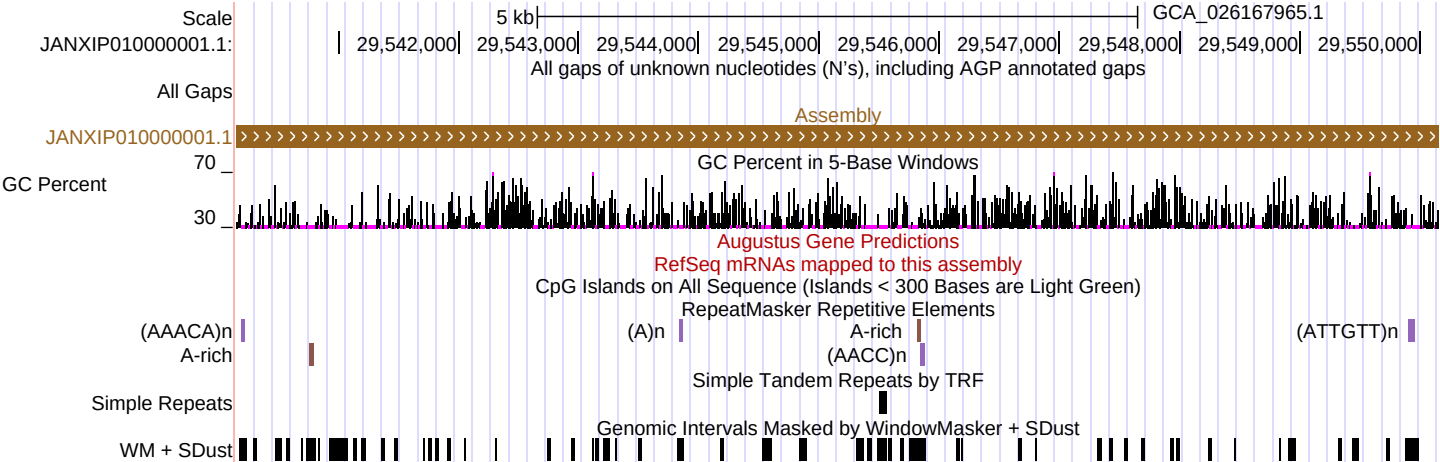
Scale
5 kb
GCA_026167965.1
JANXIP010000001.1:
29,542,000
29,543,000
29,544,000
29,545,000
29,546,000
29,547,000
29,548,000
29,549,000
29,550,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
JANXIP010000001.1
GC Percent in 5-Base Windows
70
GC Percent
30
Augustus Gene Predictions
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
(AAACA)n
A-rich
(A)n
A-rich
(AACC)n
(ATTGTT)n
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust