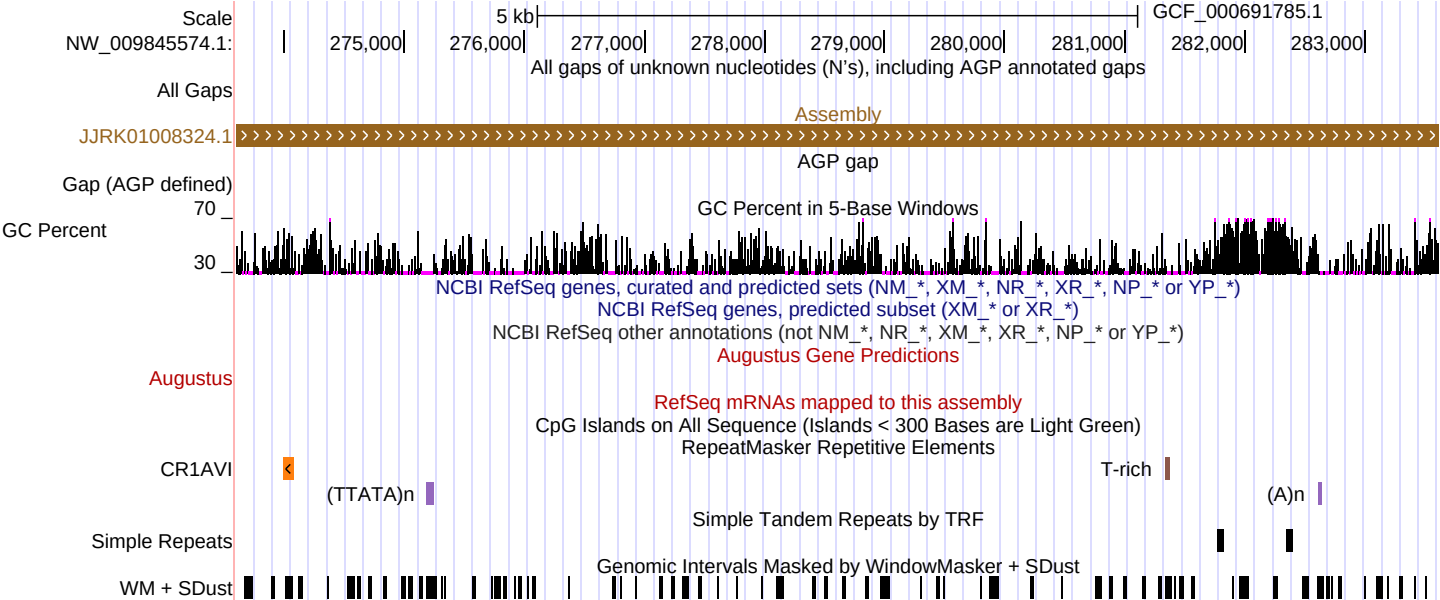
Scale
5 kb
GCF_000691785.1
NW_009845574.1:
275,000
276,000
277,000
278,000
279,000
280,000
281,000
282,000
283,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
JJRK01008324.1
AGP gap
Gap (AGP defined)
70
GC Percent
30
GC Percent in 5-Base Windows
NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)
NCBI RefSeq genes, predicted subset (XM_* or XR_*)
NCBI RefSeq other annotations (not NM_*, NR_*, XM_*, XR_*, NP_* or YP_*)
Augustus Gene Predictions
Augustus
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
CR1AVI
T-rich
(TTATA)n
(A)n
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust