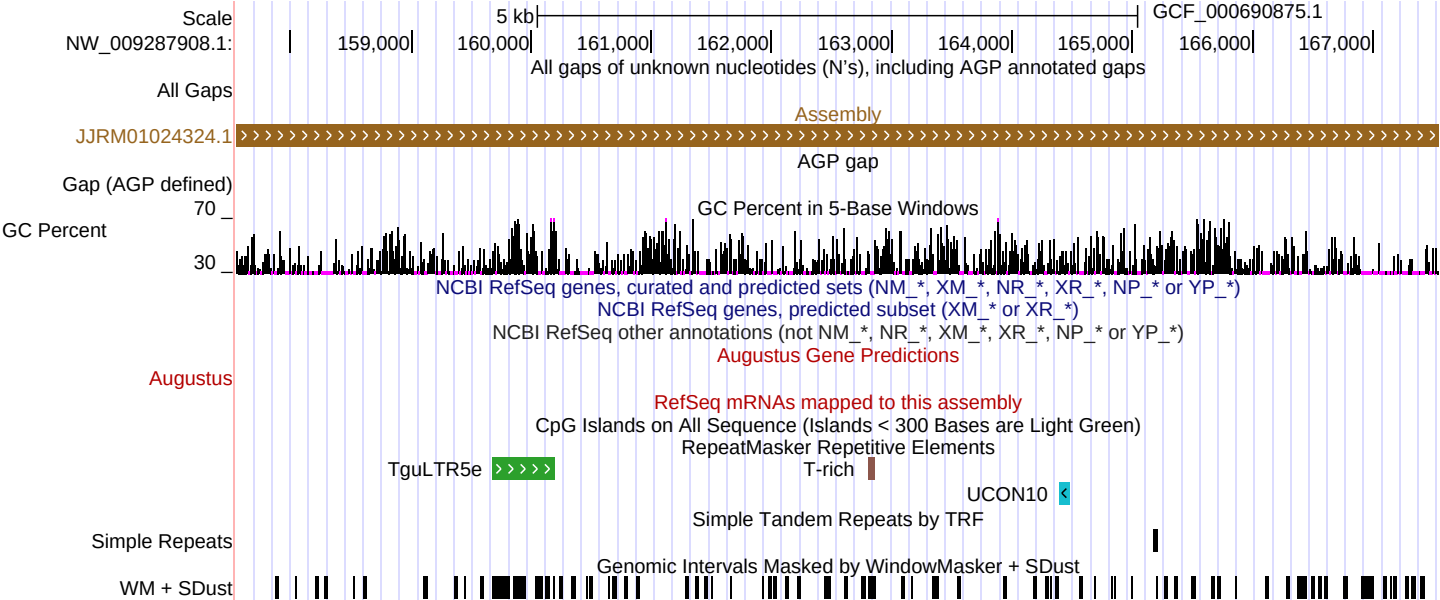
Scale
5 kb
GCF_000690875.1
NW_009287908.1:
159,000
160,000
161,000
162,000
163,000
164,000
165,000
166,000
167,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
JJRM01024324.1
AGP gap
Gap (AGP defined)
70
GC Percent in 5-Base Windows
GC Percent
30
NCBI RefSeq genes, 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
TguLTR5e
T-rich
UCON10
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust