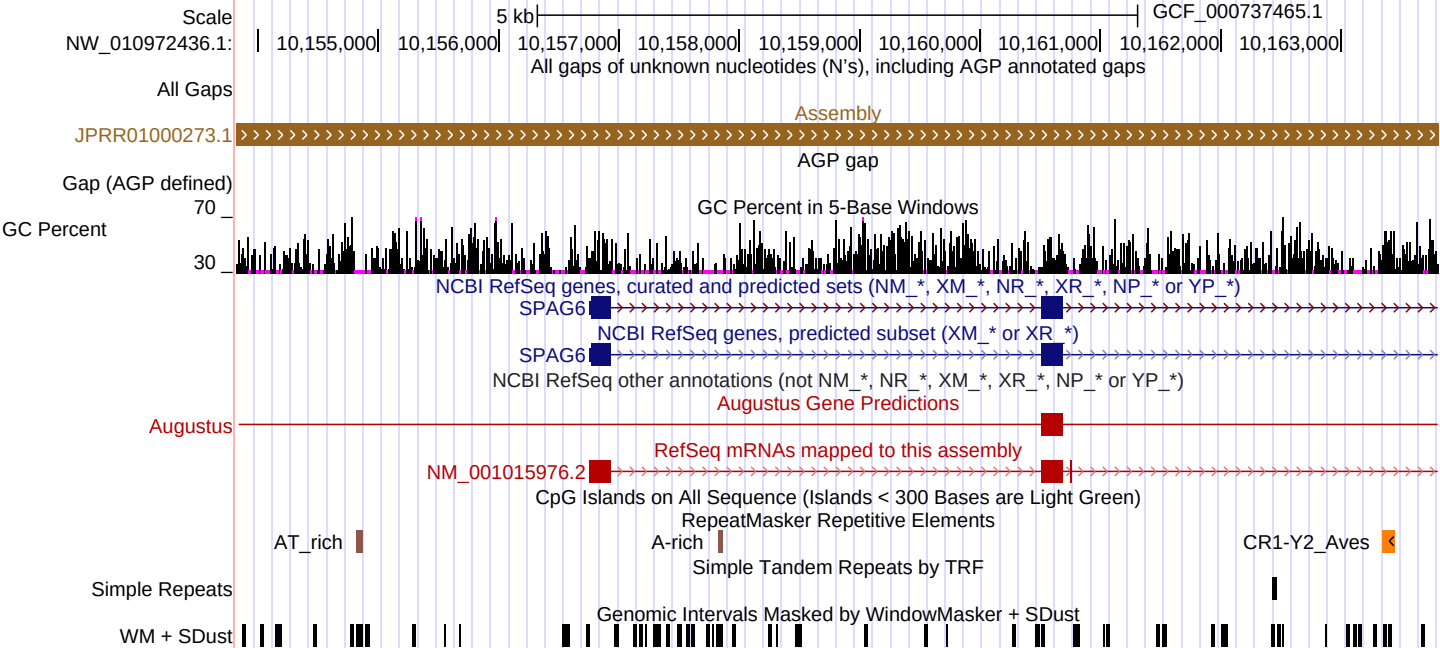
Scale
5 kb
GCF_000737465.1
NW_010972436.1:
10,155,000
10,156,000
10,157,000
10,158,000
10,159,000
10,160,000
10,161,000
10,162,000
10,163,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
JPRR01000273.1
AGP gap
Gap (AGP defined)
GC Percent in 5-Base Windows
70
GC Percent
30
NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)
SPAG6
NCBI RefSeq genes, predicted subset (XM_* or XR_*)
SPAG6
NCBI RefSeq other annotations (not NM_*, NR_*, XM_*, XR_*, NP_* or YP_*)
Augustus Gene Predictions
Augustus
RefSeq mRNAs mapped to this assembly
NM_001015976.2
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
AT_rich
A-rich
CR1-Y2_Aves
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust