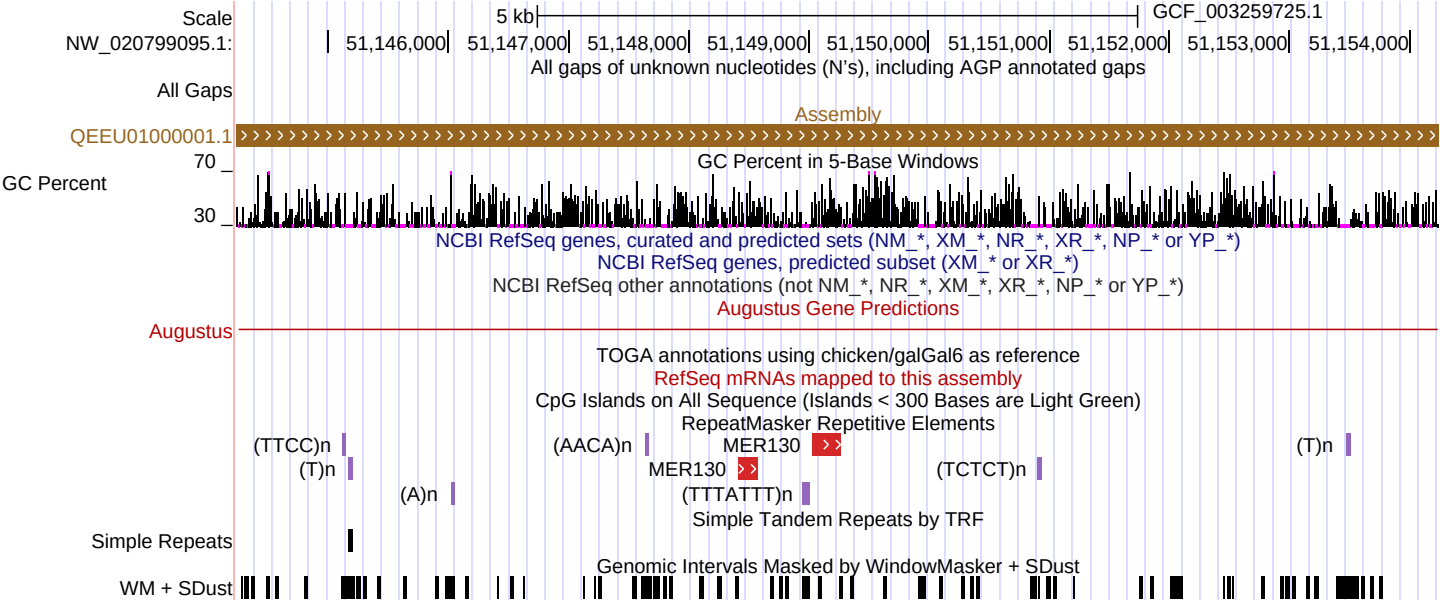
Scale
5 kb
GCF_003259725.1
NW_020799095.1:
51,146,000
51,147,000
51,148,000
51,149,000
51,150,000
51,151,000
51,152,000
51,153,000
51,154,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
QEEU01000001.1
70
GC Percent
30
GC Percent in 5-Base Windows
NCBI RefSeq 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
TOGA annotations using chicken/galGal6 as reference
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
(TTCC)n
(AACA)n
MER130
(T)n
(T)n
MER130
(TCTCT)n
(A)n
(TTTATTT)n
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust