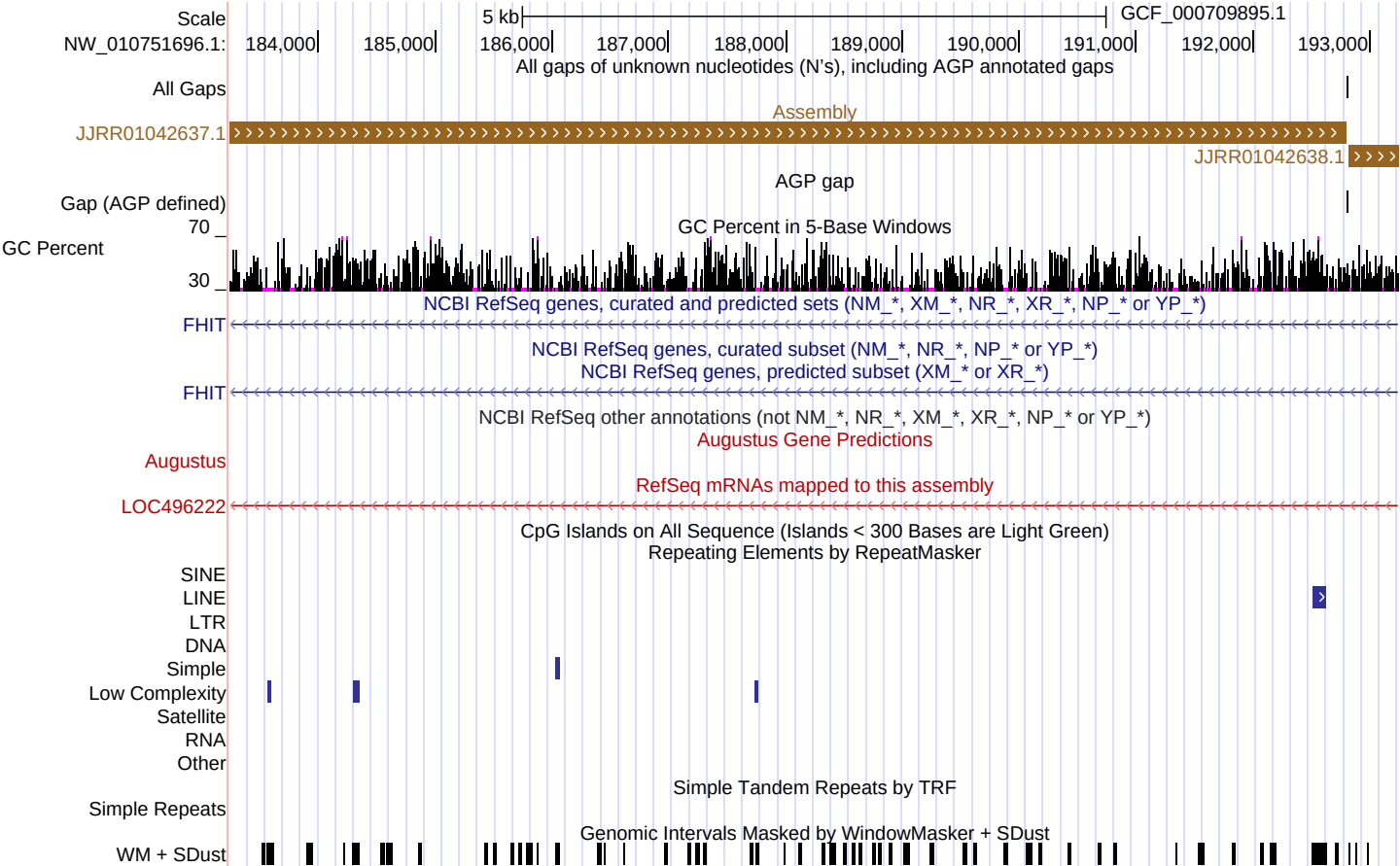
Scale
5 kb
GCF_000709895.1
NW_010751696.1:
184,000
185,000
186,000
187,000
188,000
189,000
190,000
191,000
192,000
193,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
JJRR01042637.1
JJRR01042638.1
AGP gap
Gap (AGP defined)
GC Percent in 5-Base Windows
GC Percent
70
30
NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)
FHIT
NCBI RefSeq genes, curated subset (NM_*, NR_*, NP_* or YP_*)
NCBI RefSeq genes, predicted subset (XM_* or XR_*)
FHIT
NCBI RefSeq other annotations (not NM_*, NR_*, XM_*, XR_*, NP_* or YP_*)
Augustus Gene Predictions
Augustus
RefSeq mRNAs mapped to this assembly
LOC496222
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
Repeating Elements by RepeatMasker
SINE
LINE
LTR
DNA
Simple
Low Complexity
Satellite
RNA
Other
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust