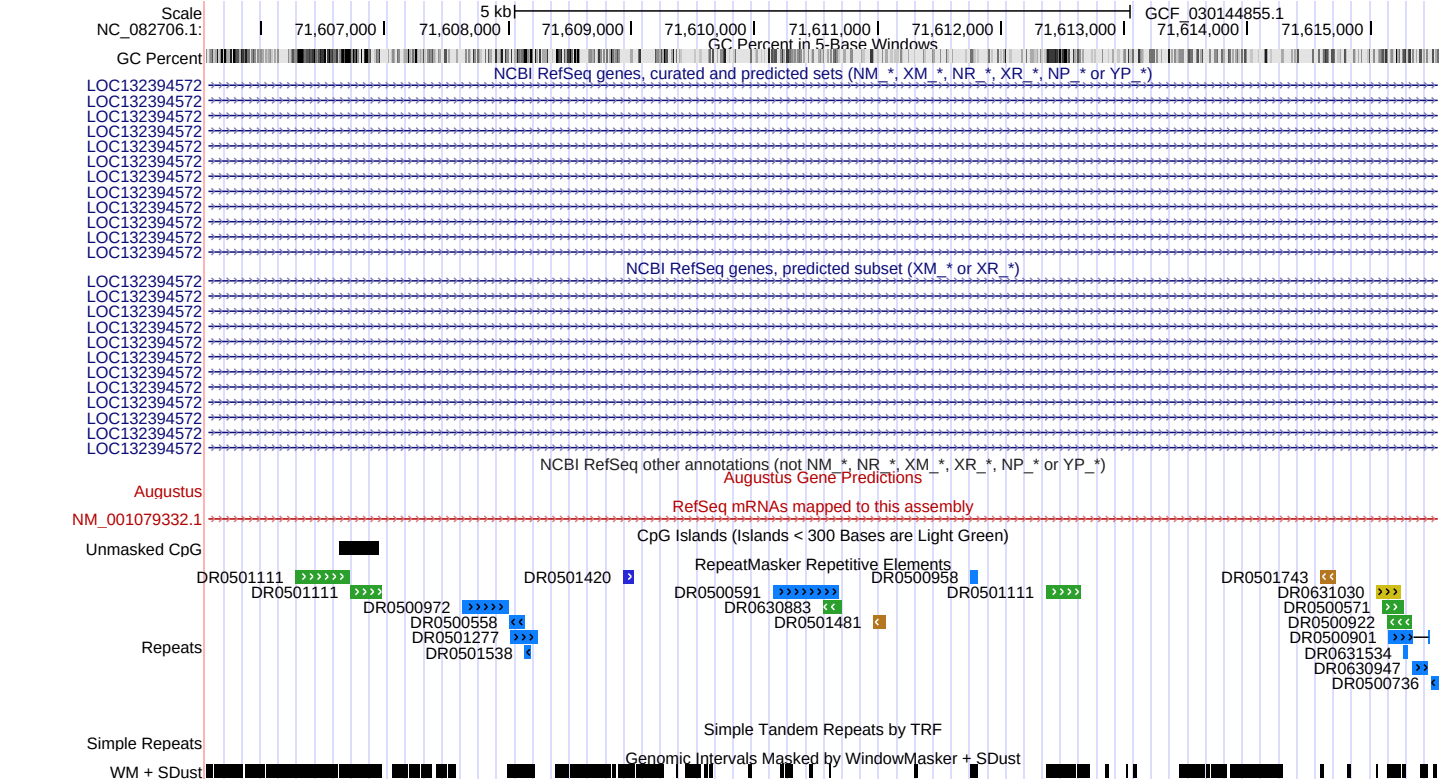
Scale
5 kb
GCF_030144855.1
NC_082706.1:
71,607,000
71,608,000
71,609,000
71,610,000
71,611,000
71,612,000
71,613,000
71,614,000
71,615,000
GC Percent
GC Percent in 5-Base Windows
NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)
LOC132394572
NCBI RefSeq genes, predicted 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
NM_001079332.1
CpG Islands (Islands < 300 Bases are Light Green)
Unmasked CpG
RepeatMasker Repetitive Elements
Repeats
DR0501111
DR0501420
DR0500958
DR0501743
DR0500591
DR0630883
DR0501481
DR0631030
DR0500571
DR0500972
DR0500558
DR0501277
DR0501538
DR0500922
DR0500901
DR0631534
DR0630947
DR0500736
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust