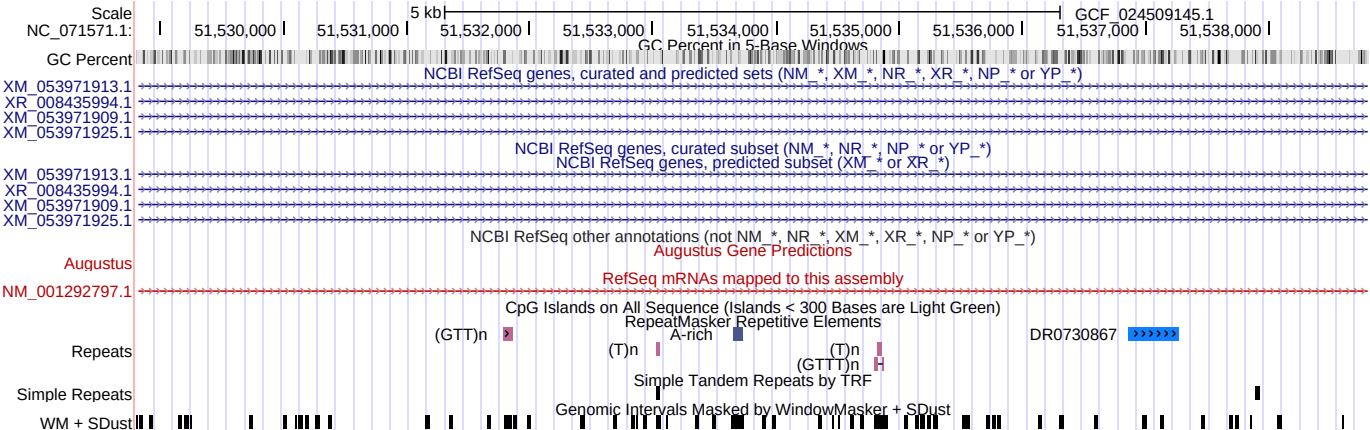
Scale
5 kb
GCF_024509145.1
NC_071571.1:
51,530,000
51,531,000
51,532,000
51,533,000
51,534,000
51,535,000
51,536,000
51,537,000
51,538,000
GC Percent in 5-Base Windows
GC Percent
NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)
XM_053971913.1
XR_008435994.1
XM_053971909.1
XM_053971925.1
NCBI RefSeq genes, curated subset (NM_*, NR_*, NP_* or YP_*)
NCBI RefSeq genes, predicted subset (XM_* or XR_*)
XM_053971913.1
XR_008435994.1
XM_053971909.1
XM_053971925.1
NCBI RefSeq other annotations (not NM_*, NR_*, XM_*, XR_*, NP_* or YP_*)
Augustus Gene Predictions
Augustus
RefSeq mRNAs mapped to this assembly
NM_001292797.1
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMaster Repetitive Elements
(GTT)n
A-rich
DR0730867
Repeats
(T)n
(T)n
(GTTT)n
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust