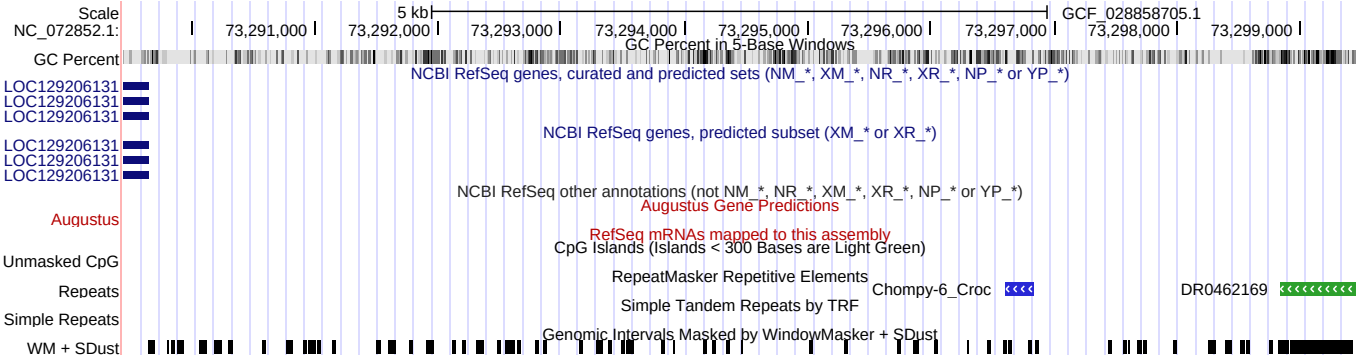
Scale
5 kb
GCF_028858705.1
NC_072852.1:
73,291,000
73,292,000
73,293,000
73,294,000
73,295,000
73,296,000
73,297,000
73,298,000
73,299,000
GC Percent in 5-Base Windows
GC Percent
NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)
LOC129206131
LOC129206131
LOC129206131
NCBI RefSeq genes, predicted subset (XM_* or XR_*)
LOC129206131
LOC129206131
LOC129206131
NCBI RefSeq other annotations (not NM_*, NR_*, XM_*, XR_*, NP_* or YP_*)
Augustus Gene Predictions
Augustus
RefSeq mRNAs mapped to this assembly
CpG Islands (Islands < 300 Bases are Light Green)
Unmasked CpG
RepeatMasker Repetitive Elements
Repeats
Chompy-6_Croc
DR0462169
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust