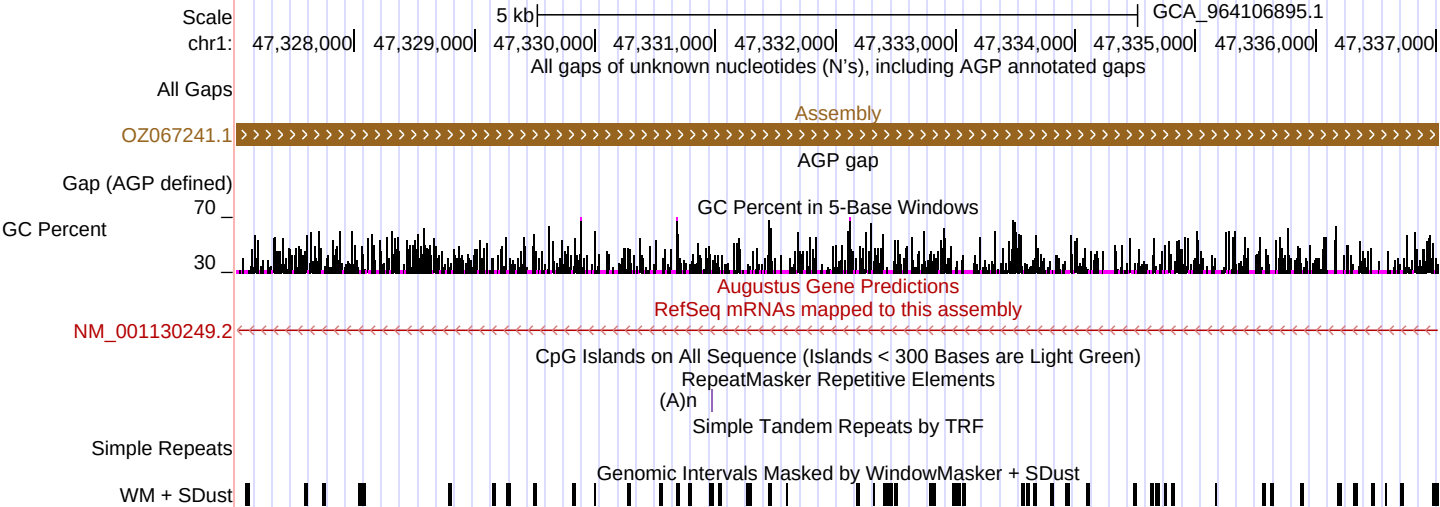
Scale
5 kb
GCA_964106895.1
chr1:
47,328,000
47,329,000
47,330,000
47,331,000
47,332,000
47,333,000
47,334,000
47,335,000
47,336,000
47,337,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
OZ067241.1
AGP gap
Gap (AGP defined)
GC Percent in 5-Base Windows
70
GC Percent
30
Augustus Gene Predictions
RefSeq mRNAs mapped to this assembly
NM_001130249.2
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
(A)n
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust