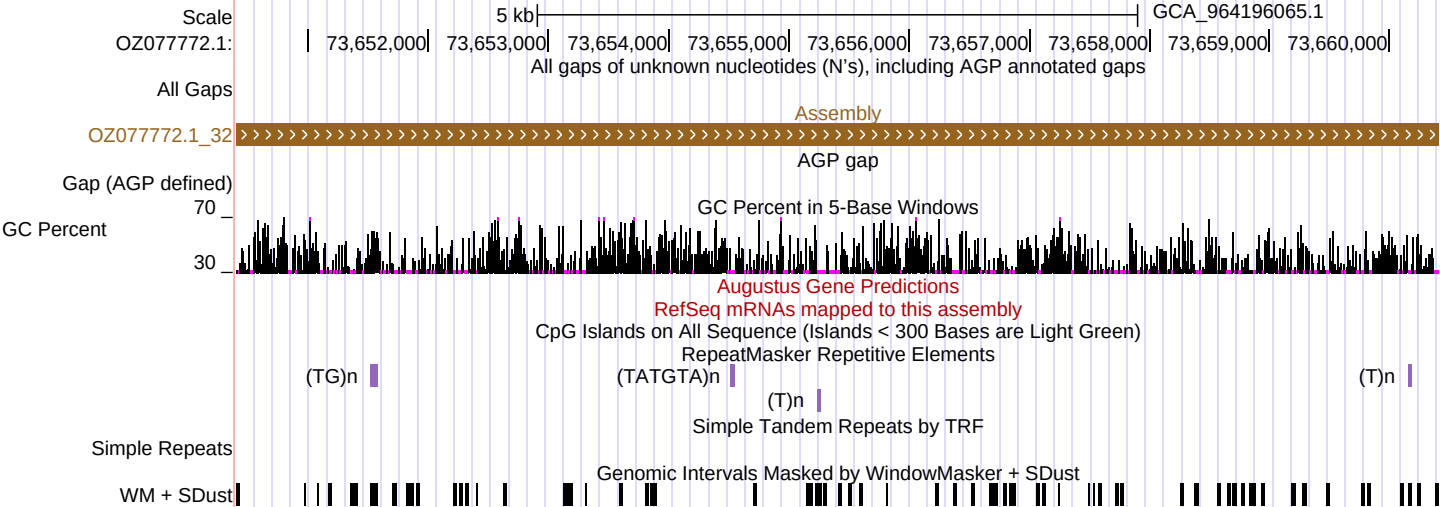
Scale
5 kb
GCA_964196065.1
OZ077772.1:
73,652,000
73,653,000
73,654,000
73,655,000
73,656,000
73,657,000
73,658,000
73,659,000
73,660,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
OZ077772.1_32
AGP gap
Gap (AGP defined)
GC Percent in 5-Base Windows
70
GC Percent
30
Augustus Gene Predictions
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
(TG)n
(TATGTA)n
(T)n
(T)n
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust