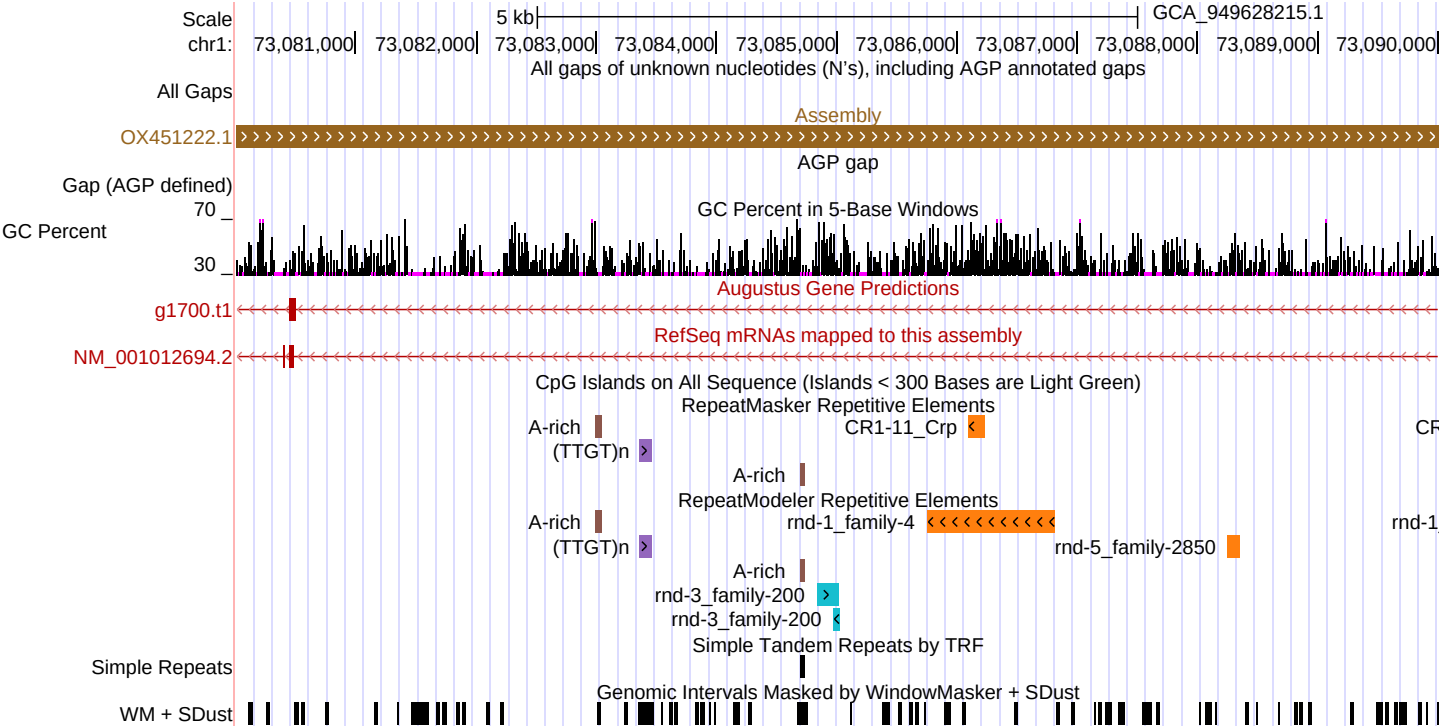
Scale
5 kb
GCA_949628215.1
chr1:
73,081,000
73,082,000
73,083,000
73,084,000
73,085,000
73,086,000
73,087,000
73,088,000
73,089,000
73,090,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
OX451222.1
AGP gap
Gap (AGP defined)
GC Percent in 5-Base Windows
70
GC Percent
30
Augustus Gene Predictions
g1700.t1
RefSeq mRNAs mapped to this assembly
NM_001012694.2
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
A-rich
CR1-11_Crp
(TTGT)n
A-rich
RepeatModeler Repetitive Elements
A-rich
rnd-1_family-4
(TTGT)n
rnd-5_family-2850
A-rich
rnd-3_family-200
rnd-3_family-200
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust