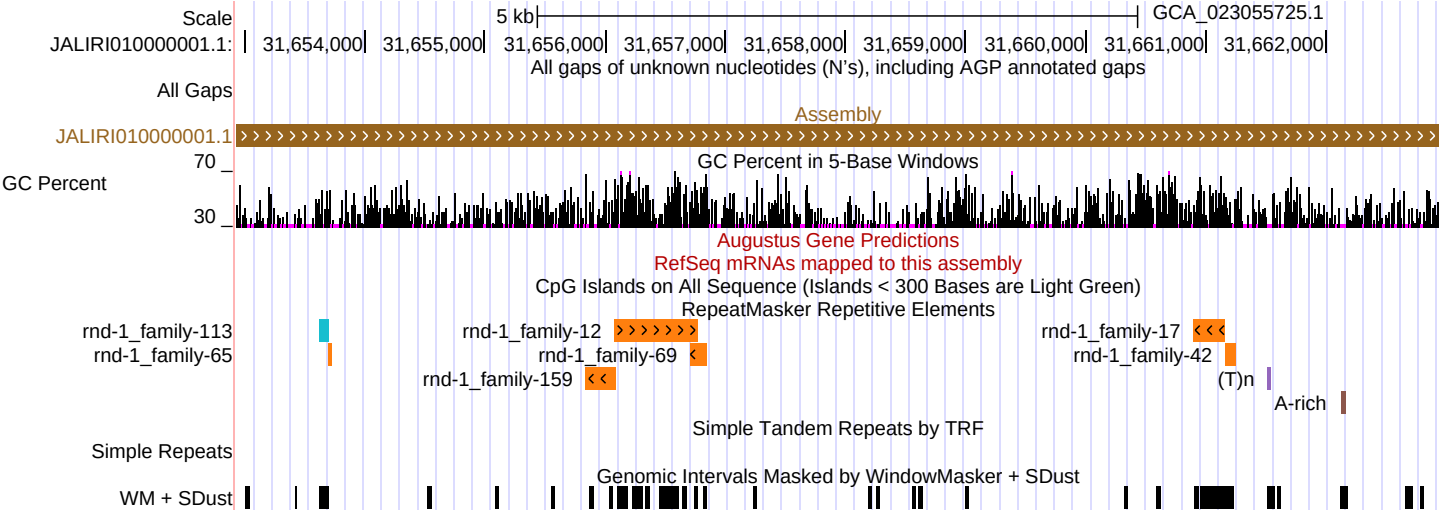
Scale
5 kb
GCA_023055725.1
JALIRI010000001.1:
31,654,000
31,655,000
31,656,000
31,657,000
31,658,000
31,659,000
31,660,000
31,661,000
31,662,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
JALIRI010000001.1
70
GC Percent in 5-Base Windows
GC Percent
30
Augustus Gene Predictions
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
rnd-1_family-113
rnd-1_family-12
rnd-1_family-17
rnd-1_family-65
rnd-1_family-69
rnd-1_family-42
rnd-1_family-159
(T)n
A-rich
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust