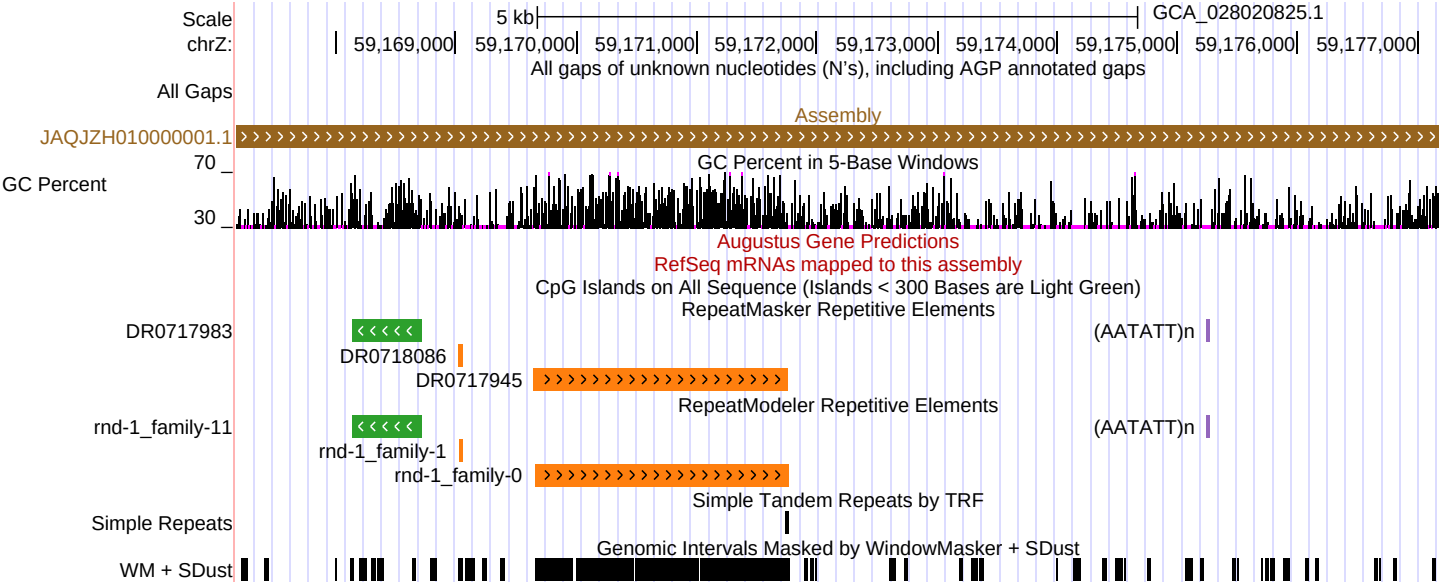
Scale
5 kb
GCA_028020825.1
chrZ:
59,169,000
59,170,000
59,171,000
59,172,000
59,173,000
59,174,000
59,175,000
59,176,000
59,177,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
JAQJZH010000001.1
70
GC Percent in 5-Base Windows
GC Percent
30
Augustus Gene Predictions
RefSeq mRNAs mapped to this assembly
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
DR0717983
(AATATT)n
DR0718086
DR0717945
RepeatModeler Repetitive Elements
rnd-1_family-11
(AATATT)n
rnd-1_family-1
rnd-1_family-0
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust