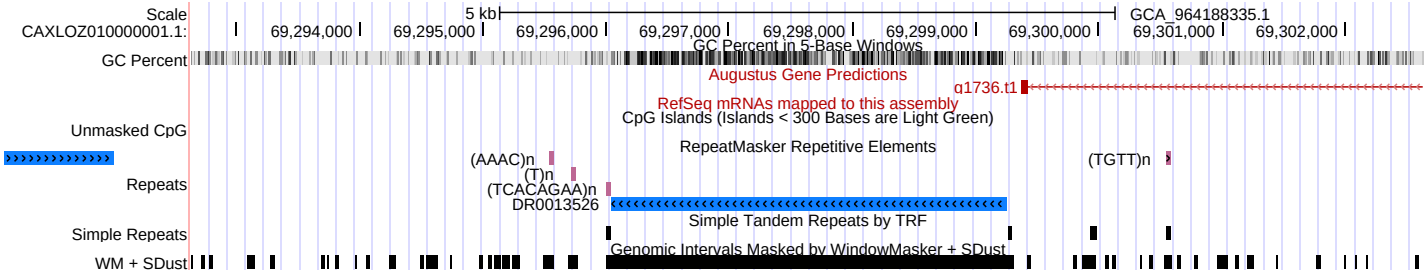
Scale
5 kb
GCA_964188335.1
CAXLOZ010000001.1:
69,294,000
69,295,000
69,296,000
69,297,000
69,298,000
69,299,000
69,300,000
69,301,000
69,302,000
GC Percent in 5-Base Windows
GC Percent
Augustus Gene Predictions
g1736.t1
RefSeq mRNAs mapped to this assembly
CpG Islands (Islands < 300 Bases are Light Green)
Unmasked CpG
RepeatMasker Repetitive Elements
(AAAC)n
(T)n
(TCACAGAA)n
DR0013526
(TGTT)n
Repeats
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust