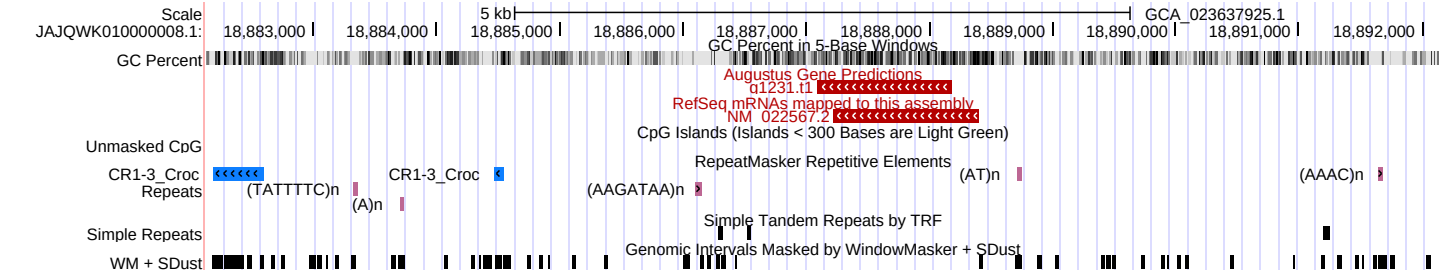
Scale
5 kb
GCA_023637925.1
JAJQWK010000008.1:
18,883,000
18,884,000
18,885,000
18,886,000
18,887,000
18,888,000
18,889,000
18,890,000
18,891,000
18,892,000
GC Percent in 5-Base Windows
GC Percent
Augustus Gene Predictions
q1231.t1
RefSeq mRNAs mapped to this assembly
NM_022567.2
CpG Islands (Islands < 300 Bases are Light Green)
Unmasked CpG
RepeatMasker Repetitive Elements
CR1-3_Croc
CR1-3_Croc
(AT)n
(AAAC)n
Repeats
(TATTTTC)n
(A)n
(AAGATAA)n
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust