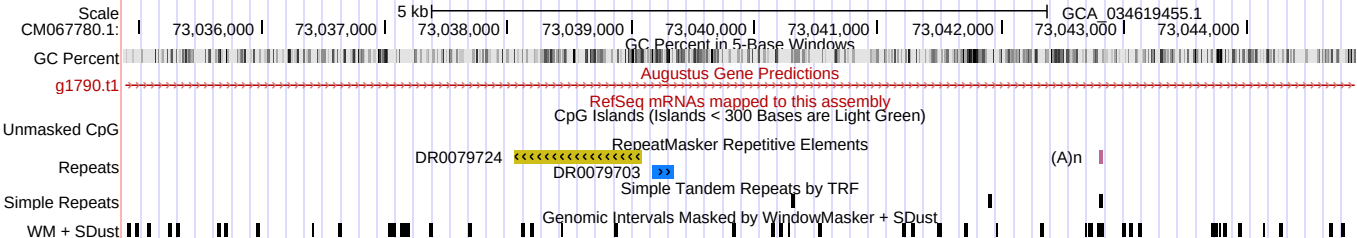
Scale
5 kb
GCA_034619455.1
CM067780.1:
73,036,000
73,037,000
73,038,000
73,039,000
73,040,000
73,041,000
73,042,000
73,043,000
73,044,000
GC Percent
GC Percent in 5-Base Windows
Augustus Gene Predictions
g1790.t1
RefSeq mRNAs mapped to this assembly
Unmasked CpG
CpG Islands (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
Repeats
DR0079724
DR0079703
(A)n
Simple Repeats
Simple Tandem Repeats by TRF
WM + SDust
Genomic Intervals Masked by WindowMasker + SDust