

Scale 5 kb | GCA_036417515.1

chr1: | 73,754,000 | 73,755,000 | 73,756,000 | 73,757,000 | 73,758,000 | 73,759,000 | 73,760,000 | 73,761,000 | 73,762,000 |

GC Percent in 5-Base Windows



Augustus Gene Predictions

RefSeq mRNAs mapped to this assembly

CpG Islands (Islands < 300 Bases are Light Green)

Unmasked CpG

RepeatMasker Repetitive Elements

(ATTTT)n

Eulor2C

GA-rich

(TC)n

(T)n

DR0017305

DR0017388

DR0017344

(A)n

DR0017353

Simple Repeats

Simple Tandem Repeats by TRF

Genomic Intervals Masked by WindowMasker + SDust

WM + SDust

