

Scale 5 kb | GCF\_020740725.1

chr2: | 41,152,000 | 41,153,000 | 41,154,000 | 41,155,000 | 41,156,000 | 41,157,000 | 41,158,000 | 41,159,000 | 41,160,000 |

GC Percent in 5-Base Windows



NCBI RefSeq genes, curated and predicted sets (NM\_\*, XM\_\*, NR\_\*, XR\_\*, NP\_\* or YP\_\*)

NCBI RefSeq genes, predicted subset (XM\_\* or XR\_\*)

NCBI RefSeq other annotations (not NM\_\*, NR\_\*, XM\_\*, XR\_\*, NP\_\* or YP\_\*)

Augustus Gene Predictions

Augustus

RefSeq mRNAs mapped to this assembly

CpG Islands (Islands < 300 Bases are Light Green)

Unmasked CpG

RepeatMasker Repetitive Elements

CR1-3\_Croc

CR1-1\_Amn

CR1\_Amni-1

CR1-11\_Crp

CR1-L3

Simple Tandem Repeats by TRF

Simple Repeats

Genomic Intervals Masked by WindowMasker + SDust

WM + SDust

