

Scale

5 kb

GCF\_026419915.1

chrZ:

| 39,019,000 | 39,020,000 | 39,021,000 | 39,022,000 | 39,023,000 | 39,024,000 | 39,025,000 | 39,026,000 | 39,027,000 |

GC Percent in 5-Base Windows

GC Percent



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

LOC128136651



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

LOC128136651



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

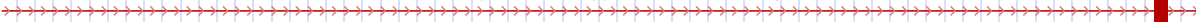
Augustus Gene Predictions

Augustus



RefSeq mRNAs mapped to this assembly

NM\_204108.1



CpG Islands (Islands < 300 Bases are Light Green)

Unmasked CpG

RepeatMasker Repetitive Elements

Simple Tandem Repeats by TRF

Simple Repeats

Genomic Intervals Masked by WindowMasker + SDust

WM + SDust

