

Scale 5 kb | GCF_018139145.2

chr1: | 73,006,000 | 73,007,000 | 73,008,000 | 73,009,000 | 73,010,000 | 73,011,000 | 73,012,000 | 73,013,000 | 73,014,000 |

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

(TTCCC)n

CR1-11_Crp <<<

Simple Tandem Repeats by TRF

Simple Repeats

Genomic Intervals Masked by WindowMasker + SDust

WM + SDust