

Scale 5 kb | GCF_037176945.1

chr1: 144,025,000 | 144,030,000

GC Percent in 5-Base Windows

GC Percent



NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)

LOC135360177



NCBI RefSeq genes, curated subset (NM_*, NR_*, NP_* or YP_*)

NCBI RefSeq genes, predicted subset (XM_* or XR_*)

LOC135360177



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

(ACG)n

(CACCAG)n

(ACA)n

(TG)n

GA-rich

(AC)n

(CA)n

Simple Repeats

Simple Tandem Repeats by TRF

Genomic Intervals Masked by WindowMasker + SDust

WM + SDust

