

Scale
chr1:

5 kb
78,039,000 | 78,040,000 | 78,041,000 | 78,042,000 | 78,043,000 | 78,044,000 | 78,045,000 | 78,046,000 | 78,047,000 |

GC Percent

GC Percent in 5-Base Windows

XR 009450792.1

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

XM 059707757.1

XM 059707765.1

XM 059707749.1

XR 009450792.1

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

XM 059707757.1

XM 059707765.1

XM 059707749.1

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

Augustus

Augustus Gene Predictions

RefSeq mRNAs mapped to this assembly

NM 017815.3

CpG Islands (Islands < 300 Bases are Light Green)

Unmasked CpG

RepeatMasker Repetitive Elements

MIR1_Amn#SINE/MIR
(GTT)n#Simple_repeat
tRNA-Ile-ATT#tRNA

(TC)n

Charlie22a

(TC)n

L1ME4a
tRNA-Phe-TTY

(TC)n

(TC)n

(T)n

MIR1_Amn

L1MC5a

MER3

MER3

L1MA9

MamSINE1

(TAAAA)n

(TC)n

MIRb

HAL1b

Simple Repeats

Simple Tandem Repeats by TRF

Genomic Intervals Masked by WindowMasker + SDust

WM + SDust