

Scale

5 kb

GCF_003850225.1

chr1:

| 67,616,000 | 67,617,000 | 67,618,000 | 67,619,000 | 67,620,000 | 67,621,000 | 67,622,000 | 67,623,000 | 67,624,000 |

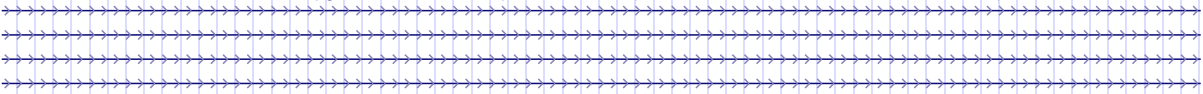
GC Percent in 5-Base Windows

GC Percent



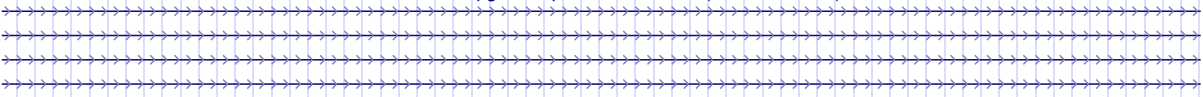
NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)

GABRG3
GABRG3
GABRG3
GABRG3



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

GABRG3
GABRG3
GABRG3
GABRG3



NCBI RefSeq other annotations (not NM_*, NR_*, XM_*, XR_*, NP_* or YP_*)
Augustus Gene Predictions

Augustus



RefSeq mRNAs mapped to this assembly

NM_024370.3



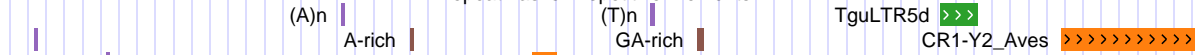
CpG Islands (Islands < 300 Bases are Light Green)

Unmasked CpG



RepeatMasker Repetitive Elements

(TTTTA)n
(AAAAC)n



CR1-Y2_Aves



Simple Repeats



WM + SDust

