

Scale 5 kb | GCA_019023105.1

chr2: | 51,926,000 | 51,927,000 | 51,928,000 | 51,929,000 | 51,930,000 | 51,931,000 | 51,932,000 | 51,933,000 | 51,934,000 |

GC Percent in 5-Base Windows



Augustus Gene Predictions

TOGA annotations using chicken/galGal6 as reference

RefSeq mRNAs mapped to this assembly

CpG Islands (Islands < 300 Bases are Light Green)

Unmasked CpG

RepeatMasker Repetitive Elements

CR1AVI >

(T)_n

CR1-8_Crp <<

(TTGG)_n

(CTT)_n

Simple Tandem Repeats by TRF

Simple Repeats

Genomic Intervals Masked by WindowMasker + SDust

WM + SDust

