

Scale 5 kb | GCF\_029582105.1  
chr2: | 49,466,000 | 49,467,000 | 49,468,000 | 49,469,000 | 49,470,000 | 49,471,000 | 49,472,000 | 49,473,000 | 49,474,000 |

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  
(CTTC)n

DR0722820  
(CTCTCCT)n

DR0723023 <<

DR0722774

DR0722844 <

DR0722882 <

DR0722824 <<

(TTTCTGC)n

DR0723144 <<

(AC)n

Simple Tandem Repeats by TRF

Simple Repeats

Genomic Intervals Masked by WindowMasker + SDust

WM + SDust

