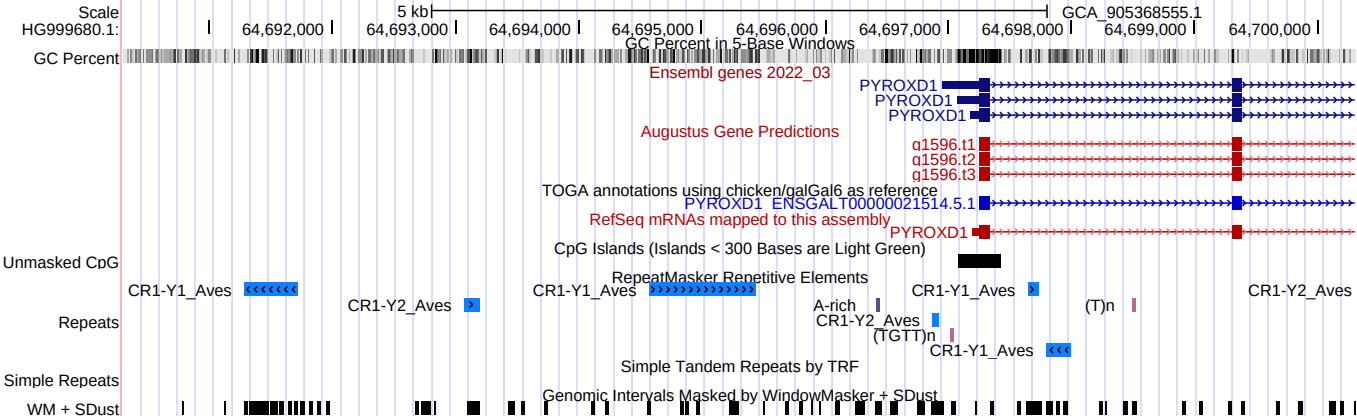
Scale
HG999680.1:
5 kb
GCA_905368555.1
64,692,000
64,693,000
64,694,000
64,695,000
64,696,000
64,697,000
64,698,000
64,699,000
64,700,000
GC Percent
GC Percent in 5-Base Windows
Ensembl genes 2022_03
PYROXD1
PYROXD1
PYROXD1
Augustus Gene Predictions
g1596.t1
g1596.t2
g1596.t3
TOGA annotations using chicken/galGal6 as reference
PYROXD1 ENSGALT00000021514.5.1
RefSeq mRNAs mapped to this assembly
PYROXD1
Unmasked CpG
CpG Islands (Islands < 300 Bases are Light Green)
RepeatMasker Repetitive Elements
CR1-Y1_Aves
CR1-Y2_Aves
CR1-Y1_Aves
CR1-Y1_Aves
CR1-Y2_Aves
A-rich
(T)n
CR1-Y2_Aves
(TGTT)n
CR1-Y1_Aves
Repeats
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust