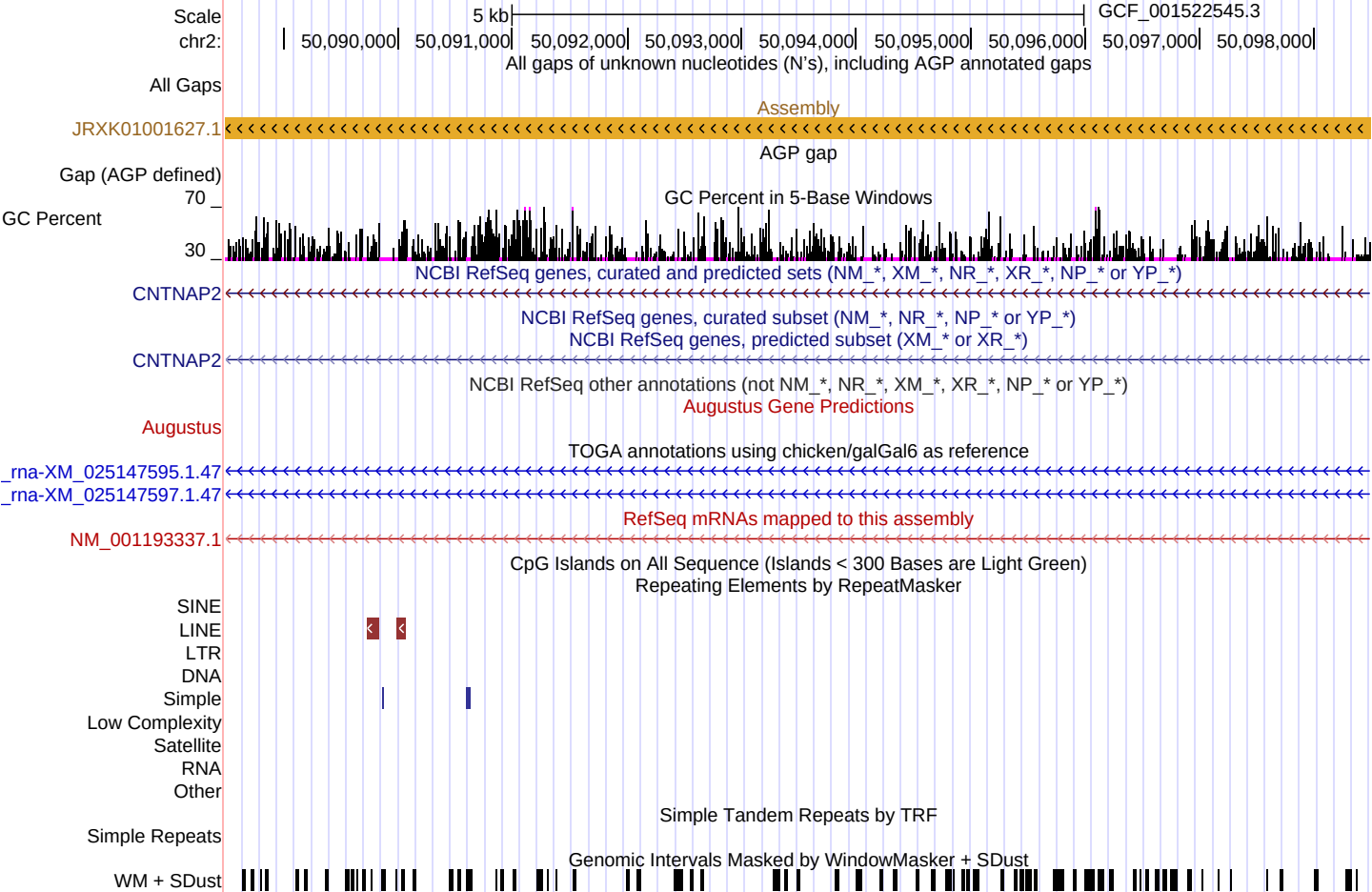
Scale
chr2:
5 kb
GCF_001522545.3
50,090,000
50,091,000
50,092,000
50,093,000
50,094,000
50,095,000
50,096,000
50,097,000
50,098,000
All gaps of unknown nucleotides (N's), including AGP annotated gaps
All Gaps
Assembly
JRXK01001627.1
AGP gap
Gap (AGP defined)
GC Percent in 5-Base Windows
70
GC Percent
30
NCBI RefSeq genes, curated and predicted sets (NM_*, XM_*, NR_*, XR_*, NP_* or YP_*)
CNTNAP2
NCBI RefSeq genes, curated subset (NM_*, NR_*, NP_* or YP_*)
NCBI RefSeq genes, predicted subset (XM_* or XR_*)
CNTNAP2
NCBI RefSeq other annotations (not NM_*, NR_*, XM_*, XR_*, NP_* or YP_*)
Augustus Gene Predictions
Augustus
TOGA annotations using chicken/galGal6 as reference
_rna-XM_025147595.1.47
_rna-XM_025147597.1.47
RefSeq mRNAs mapped to this assembly
NM_001193337.1
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
Repeating Elements by RepeatMasker
SINE
LINE
LTR
DNA
Simple
Low Complexity
Satellite
RNA
Other
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust