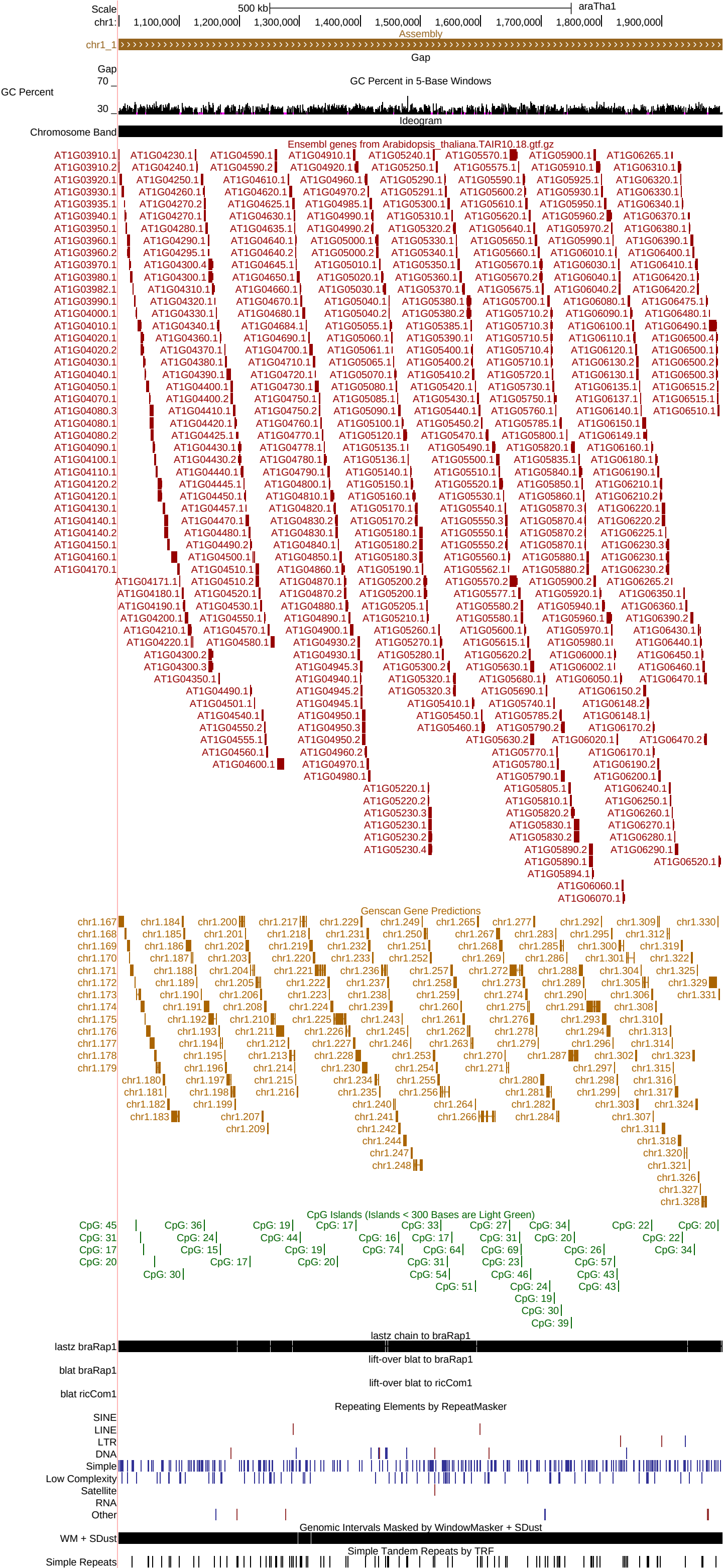
Scale
chr1:
500 kb
araTha1
1,100,000
1,200,000
1,300,000
1,400,000
1,500,000
1,600,000
1,700,000
1,800,000
1,900,000
Assembly
chr1_1
Gap
Gap
GC Percent in 5-Base Windows
GC Percent
70
30
Ideogram
Chromosome Band
Ensembl genes from Arabidopsis_thaliana.TAIR10.18.gtf.gz
Genscan Gene Predictions
CpG Islands (Islands < 300 Bases are Light Green)
lastz chain to braRap1
lastz braRap1
lift-over blat to braRap1
blat braRap1
lift-over blat to ricCom1
blat ricCom1
Repeating Elements by RepeatMasker
SINE
LINE
LTR
DNA
Simple
Low Complexity
Satellite
RNA
Other
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust
Simple Tandem Repeats by TRF
Simple Repeats