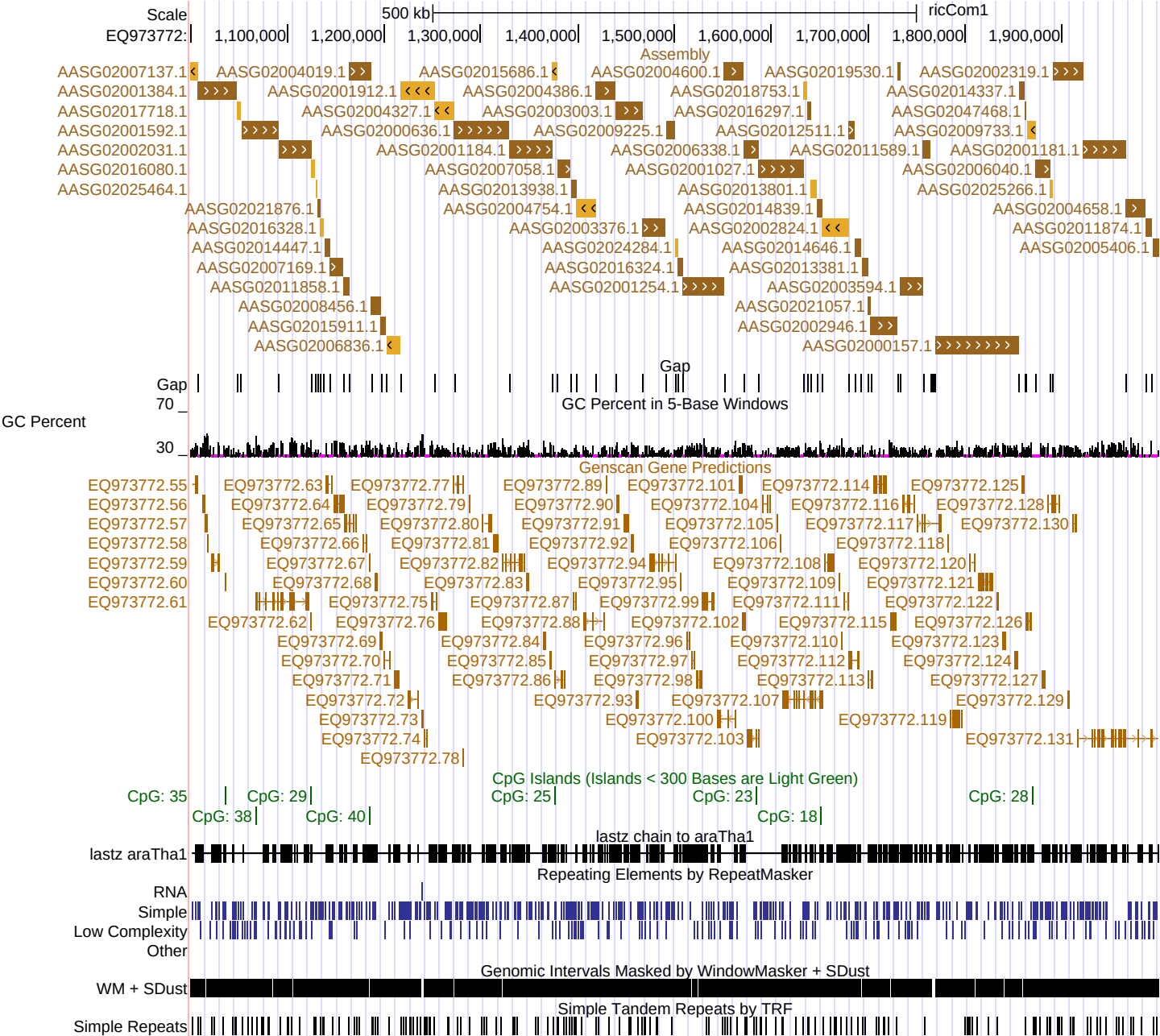
Scale
500 kb
ricCom1
EQ973772:
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1,200,000
1,300,000
1,400,000
1,500,000
1,600,000
1,700,000
1,800,000
1,900,000
Assembly
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AASG02004019.1
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AASG02019530.1
AASG02002319.1
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AASG02004386.1
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AASG02002824.1
AASG02011874.1
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AASG02006836.1
AASG02000157.1
Gap
Gap
GC Percent
70
30
GC Percent in 5-Base Windows
Genscan Gene Predictions
EQ973772.55
EQ973772.63
EQ973772.77
EQ973772.89
EQ973772.101
EQ973772.114
EQ973772.125
EQ973772.56
EQ973772.64
EQ973772.79
EQ973772.90
EQ973772.104
EQ973772.116
EQ973772.128
EQ973772.57
EQ973772.65
EQ973772.80
EQ973772.91
EQ973772.105
EQ973772.117
EQ973772.130
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EQ973772.66
EQ973772.81
EQ973772.92
EQ973772.106
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EQ973772.59
EQ973772.67
EQ973772.82
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EQ973772.122
EQ973772.62
EQ973772.76
EQ973772.88
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EQ973772.126
EQ973772.69
EQ973772.84
EQ973772.96
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EQ973772.112
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EQ973772.107
EQ973772.129
EQ973772.73
EQ973772.100
EQ973772.119
EQ973772.74
EQ973772.103
EQ973772.131
EQ973772.78
CpG Islands (Islands < 300 Bases are Light Green)
CpG: 35
CpG: 29
CpG: 25
CpG: 23
CpG: 28
CpG: 38
CpG: 40
CpG: 18
lastz chain to araTha1
lastz araTha1
Repeating Elements by RepeatMasker
RNA
Simple
Low Complexity
Other
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust
Simple Tandem Repeats by TRF
Simple Repeats