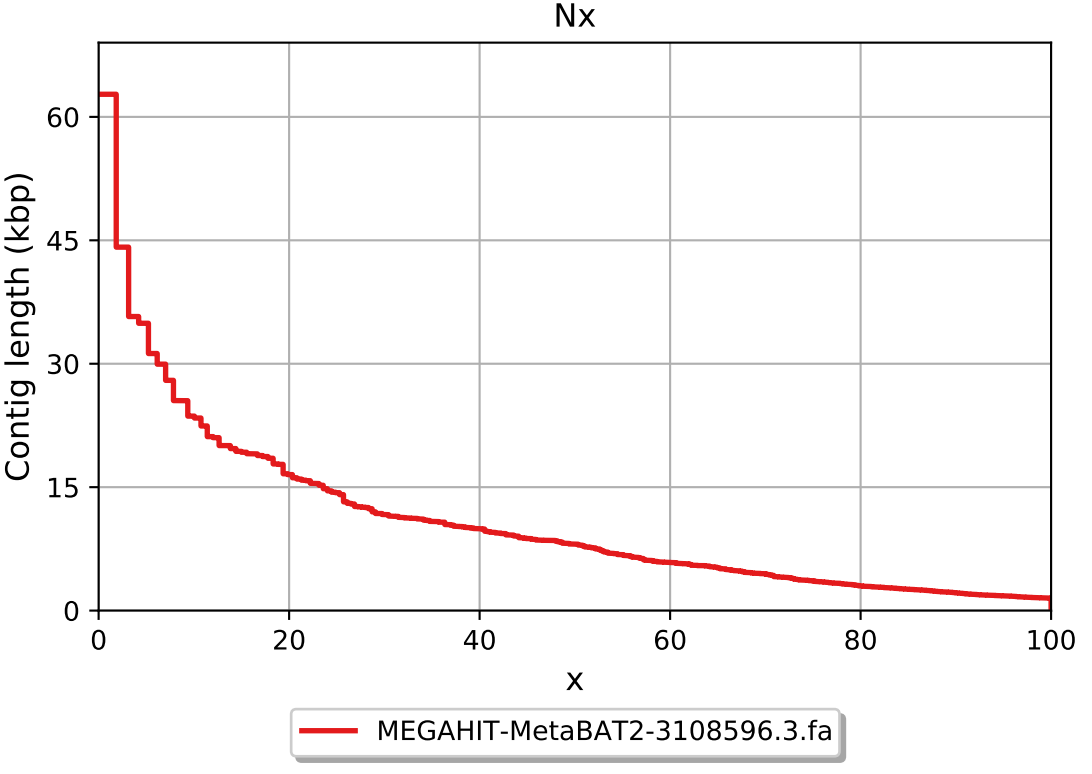
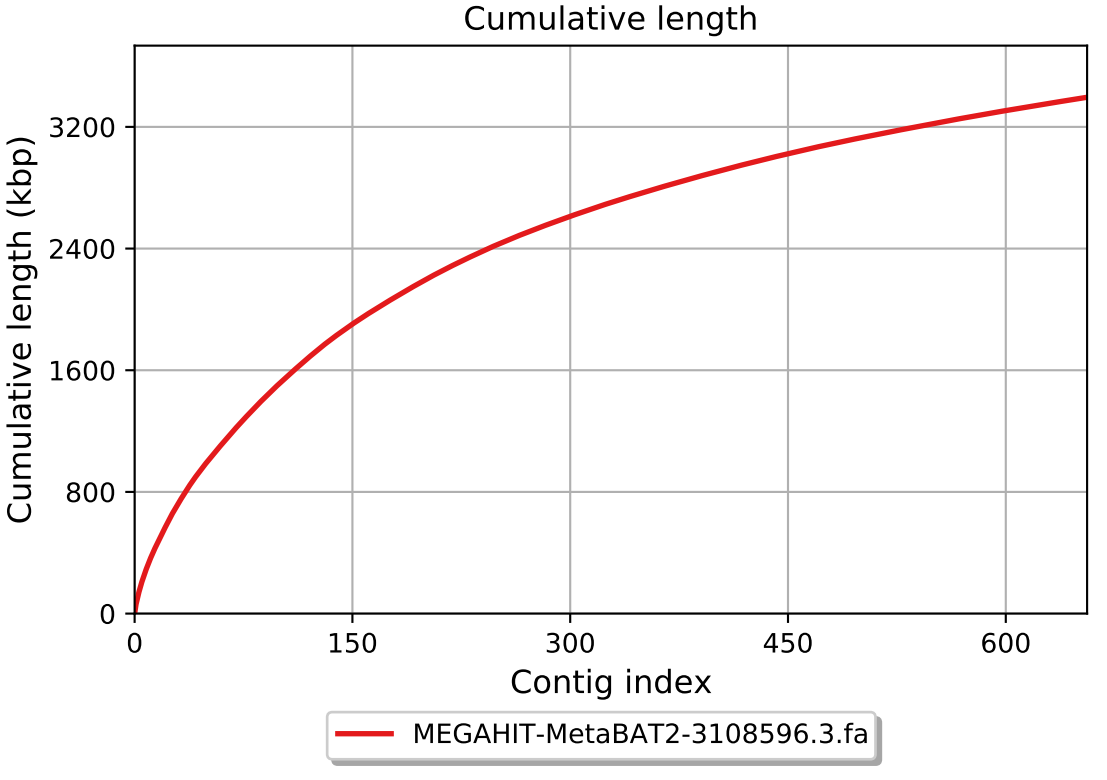
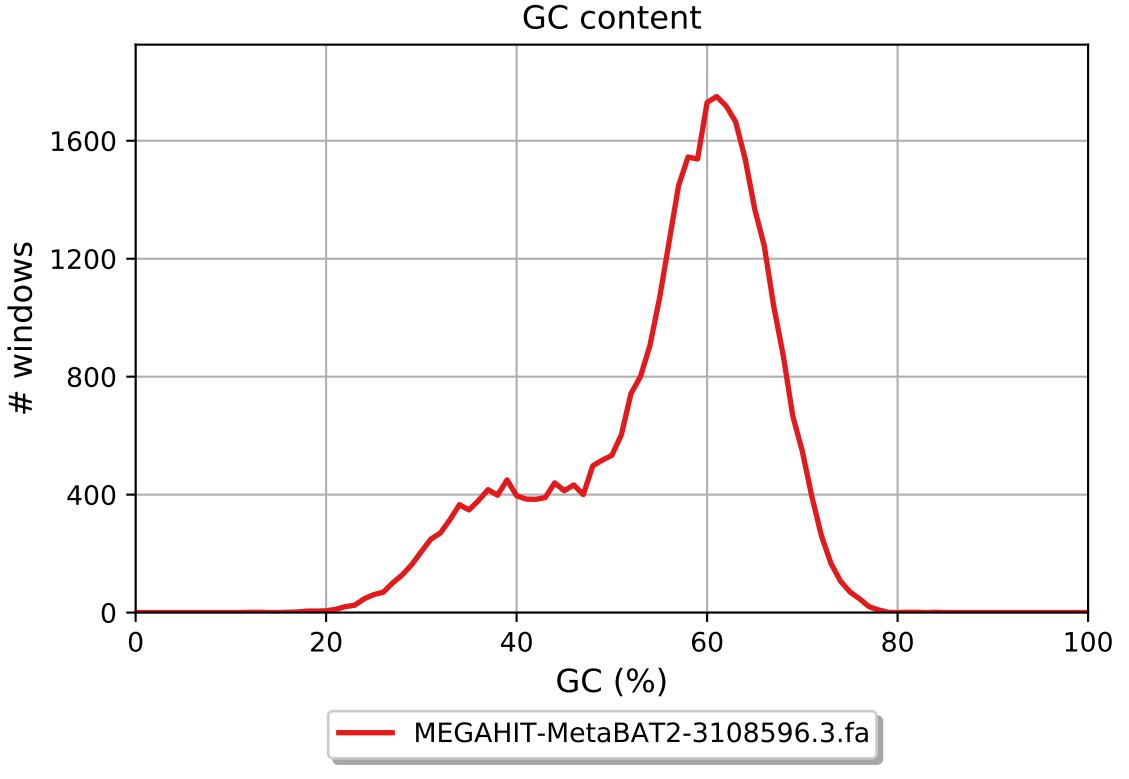
Report

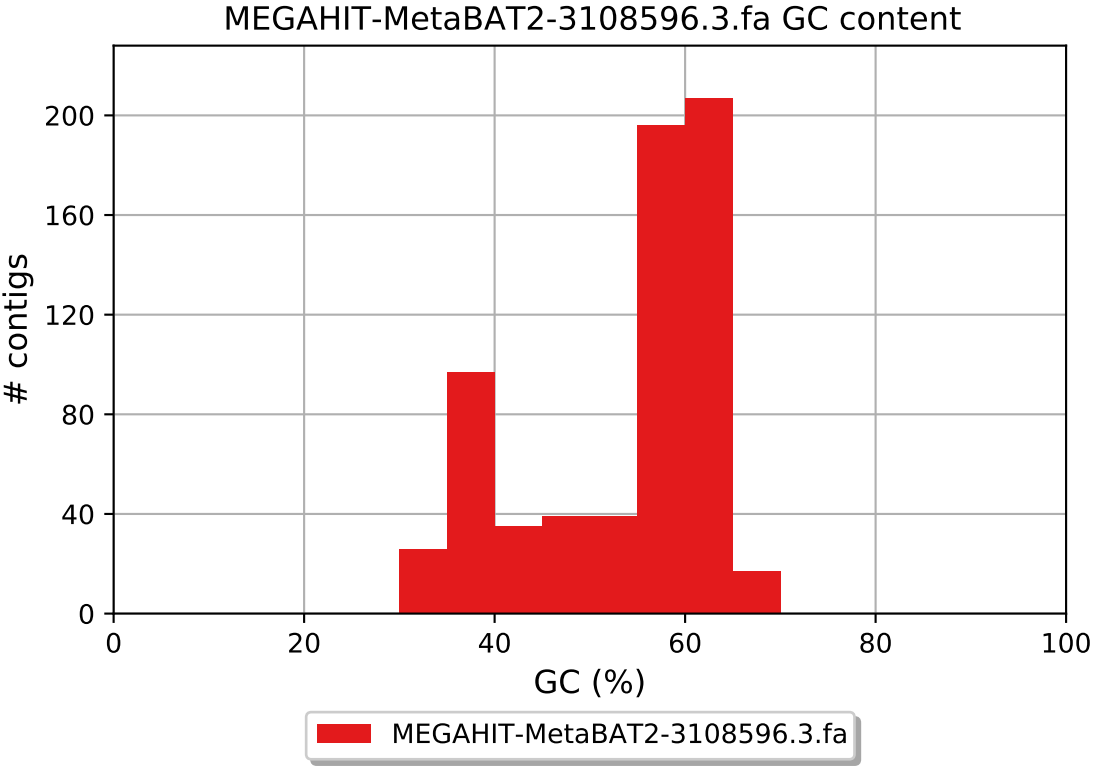
	MEGAHIT-MetaBAT2-3108596.3.fa
# contigs (>= 0 bp)	656
# contigs (>= 1000 bp)	656
# contigs (>= 5000 bp)	208
# contigs (>= 10000 bp)	81
# contigs (>= 25000 bp)	9
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	3394758
Total length (>= 1000 bp)	3394758
Total length (>= 5000 bp)	2234986
Total length (>= 10000 bp)	1336147
Total length (>= 25000 bp)	317773
Total length (>= 50000 bp)	62749
# contigs	656
Largest contig	62749
Total length	3394758
GC (%)	55.63
N50	8073
N75	3600
L50	122
L75	282
# N's per 100 kbp	0.00
# predicted rRNA genes	0 + 1 part

All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).


Nx
Contig length (kbp)
60
45
30
15
0
0
20
40
60
80
100
x
MEGAHIT-MetaBAT2-3108596.3.fa


Cumulative length
Cumulative length (kbp)
3200
2400
1600
800
0
0
150
300
450
600
Contig index
MEGAHIT-MetaBAT2-3108596.3.fa


GC content
windows
0
400
800
1200
1600
0
20
40
60
80
100
GC (%)
MEGAHIT-MetaBAT2-3108596.3.fa


MEGAHIT-MetaBAT2-3108596.3.fa GC content
contigs
0
40
80
120
160
200
0
20
40
60
80
100
GC (%)
MEGAHIT-MetaBAT2-3108596.3.fa