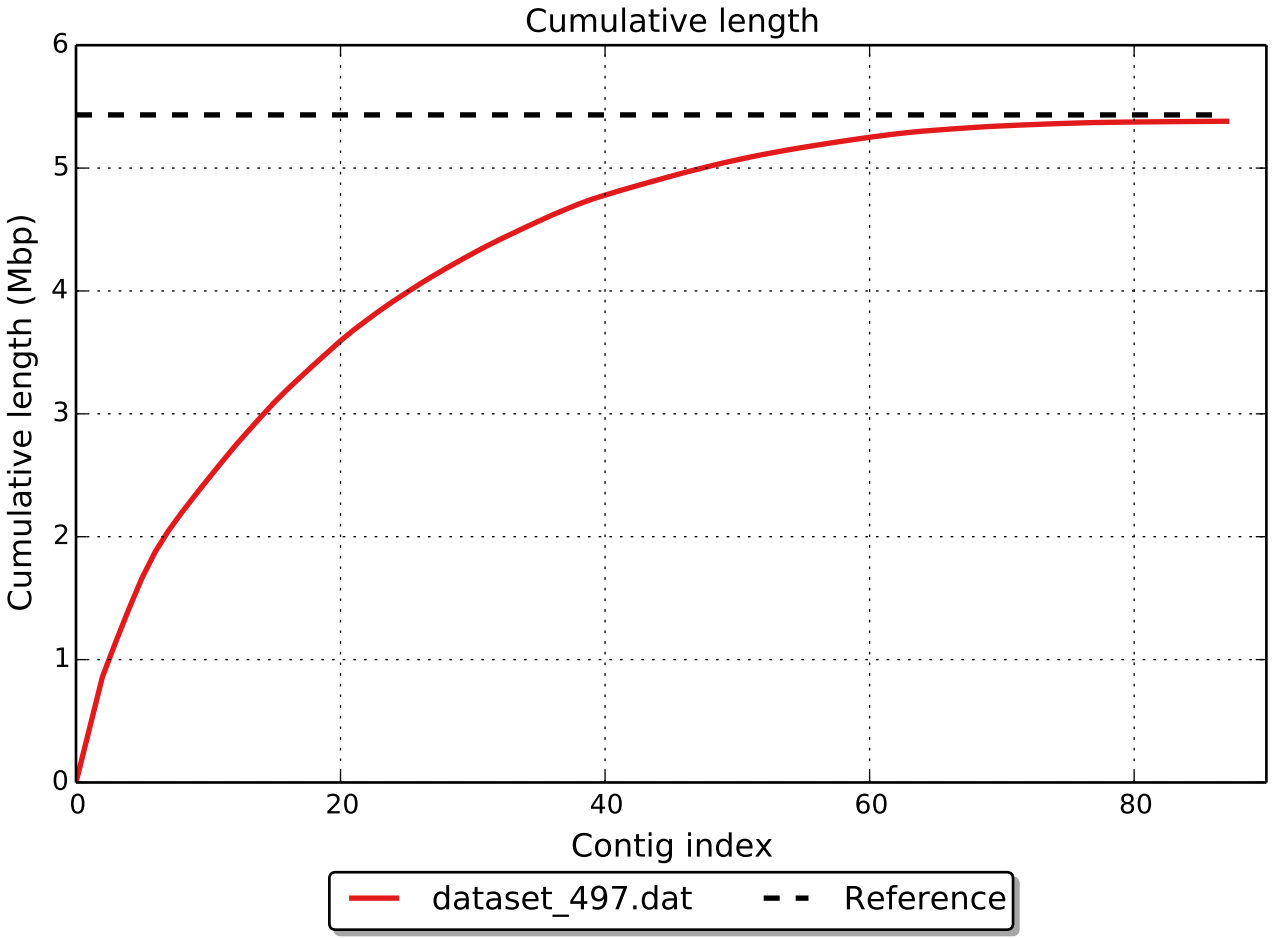
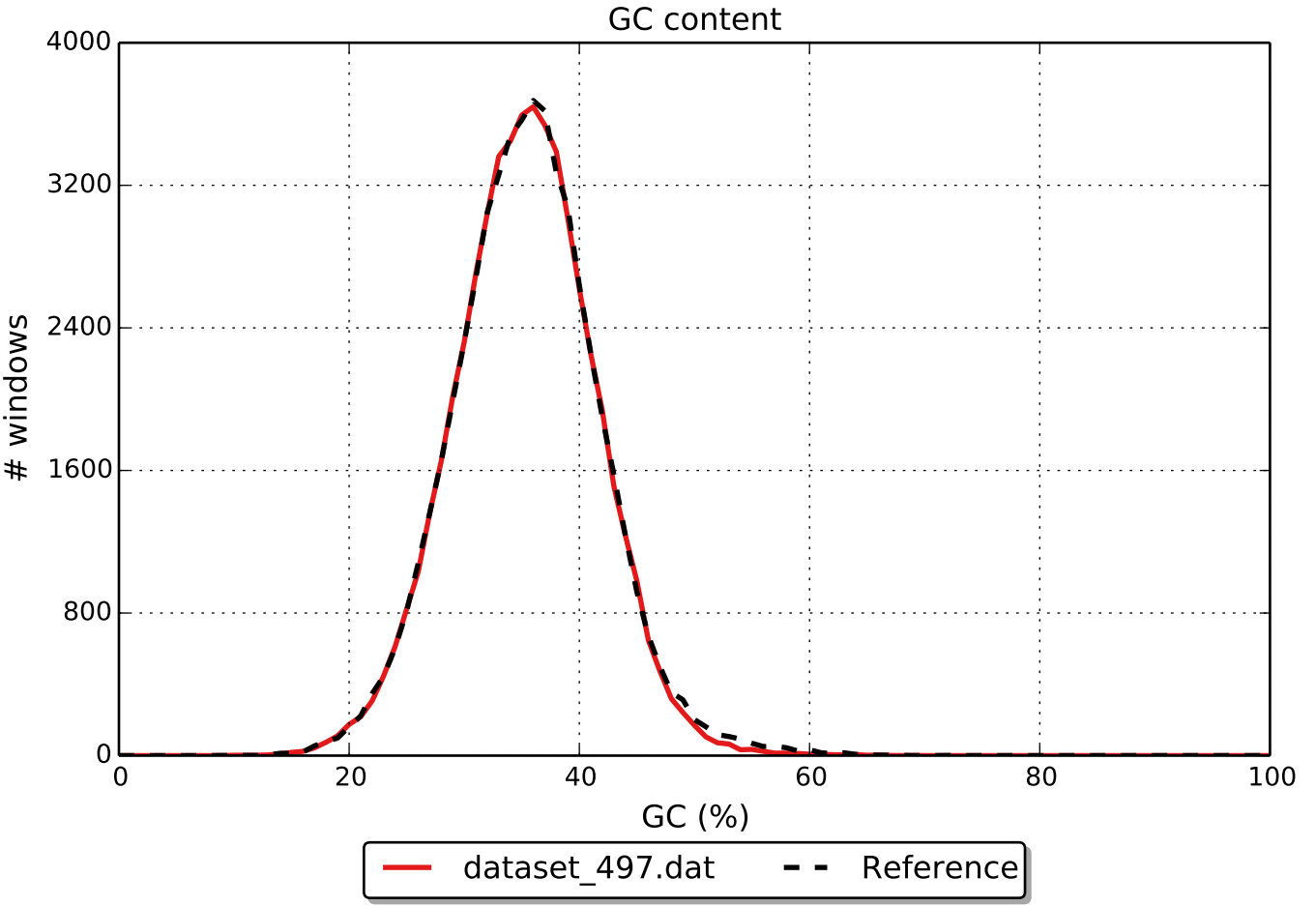
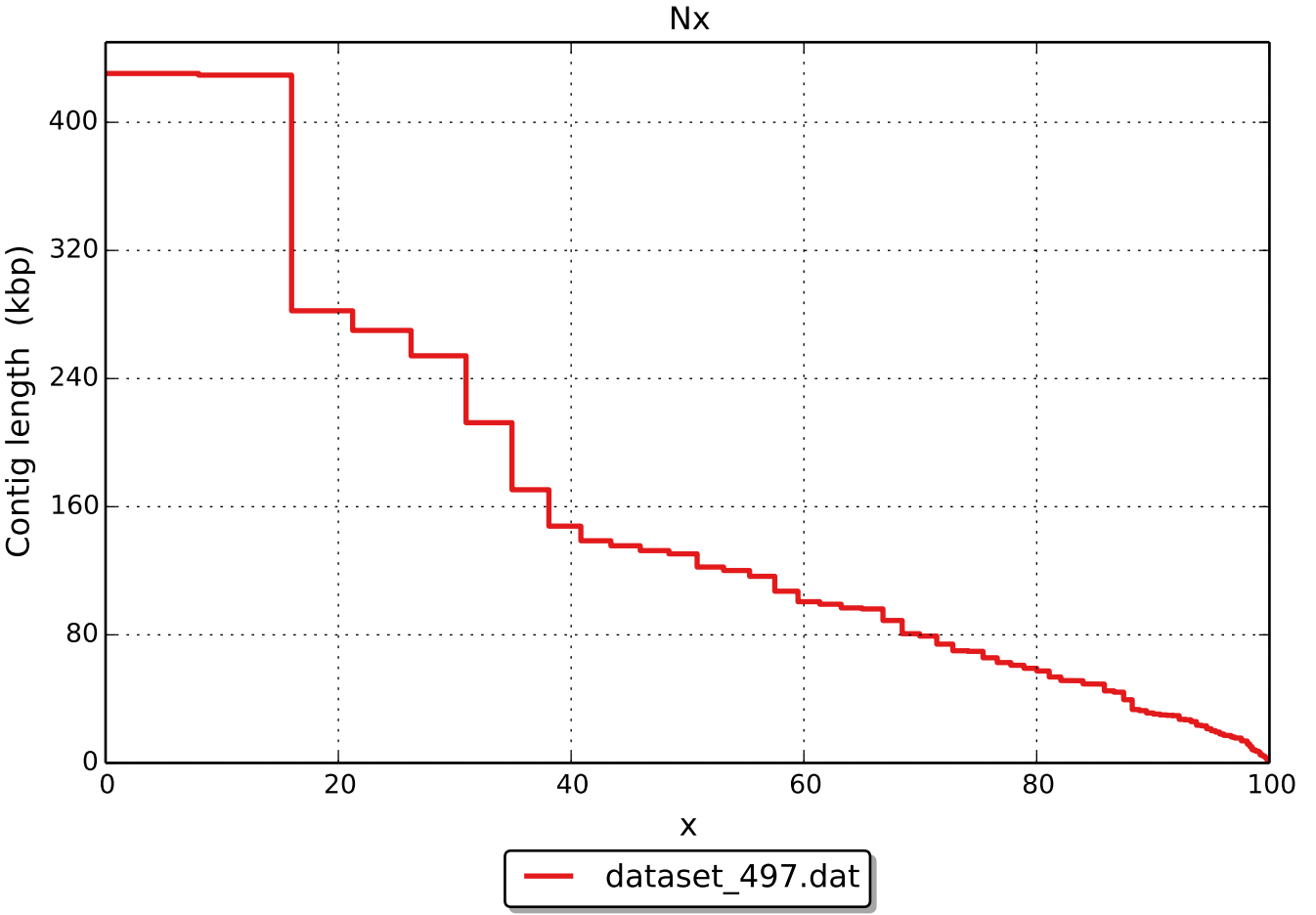
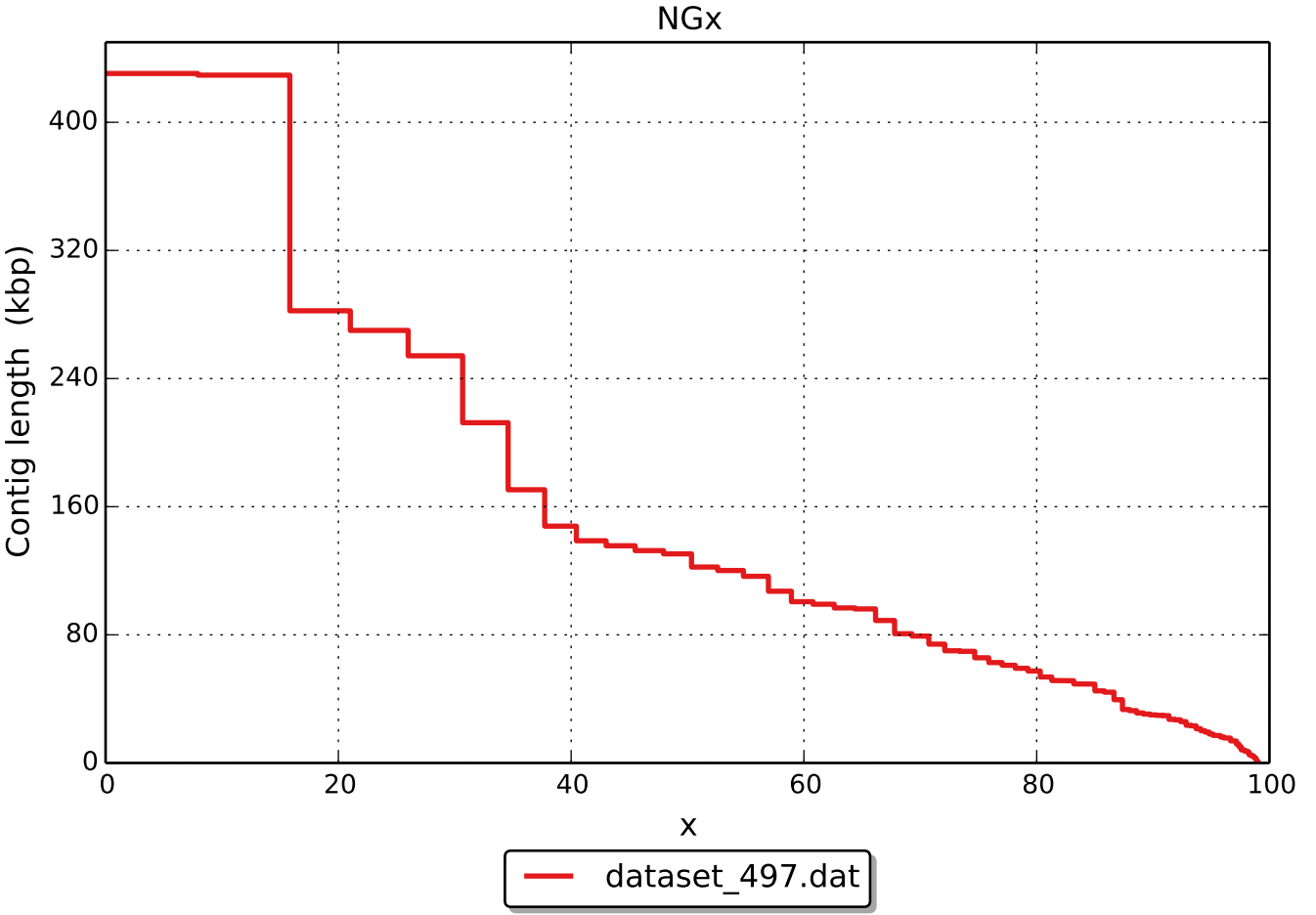
Combined_reference

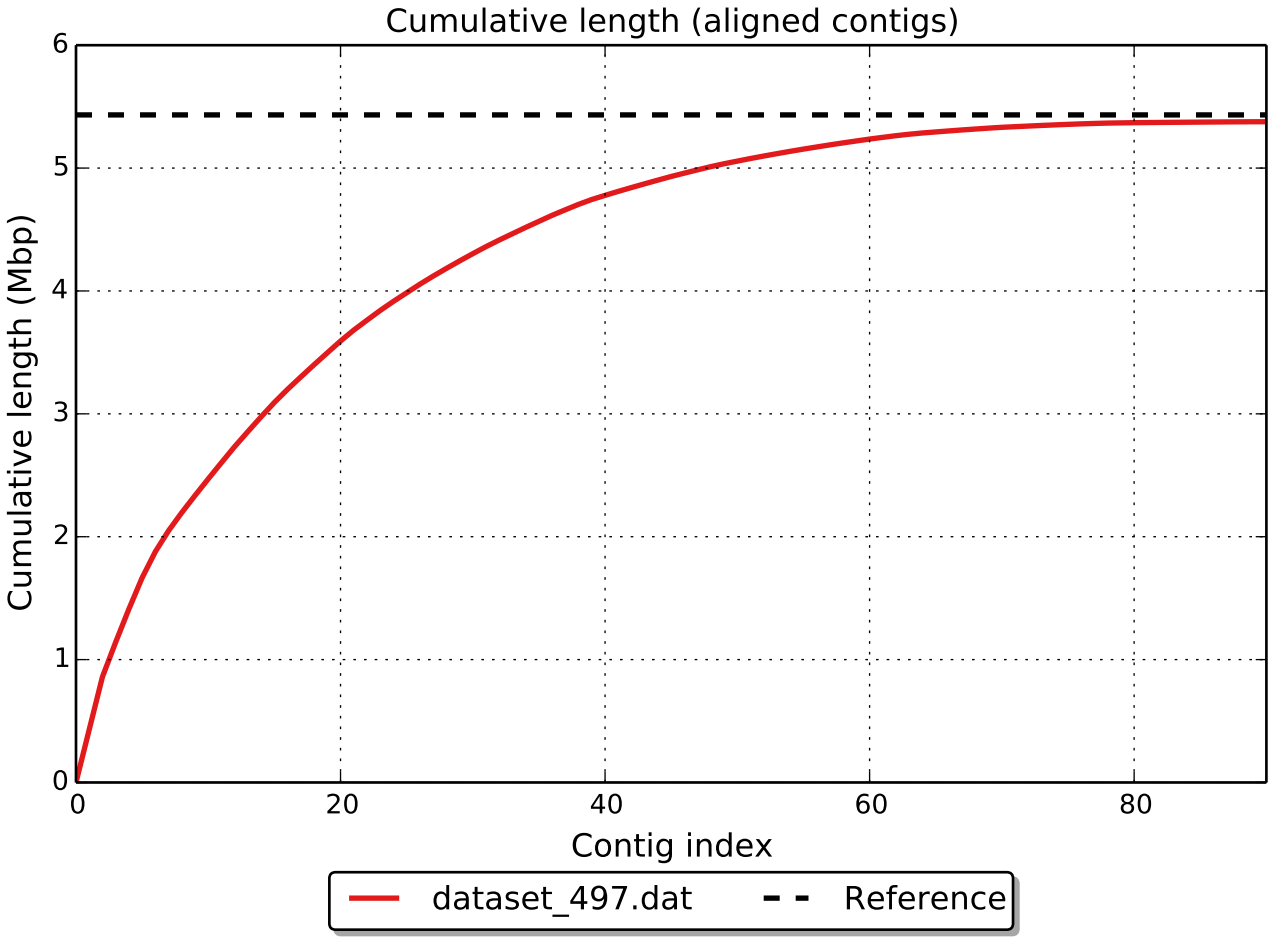
B_cereus_ATCC_1098

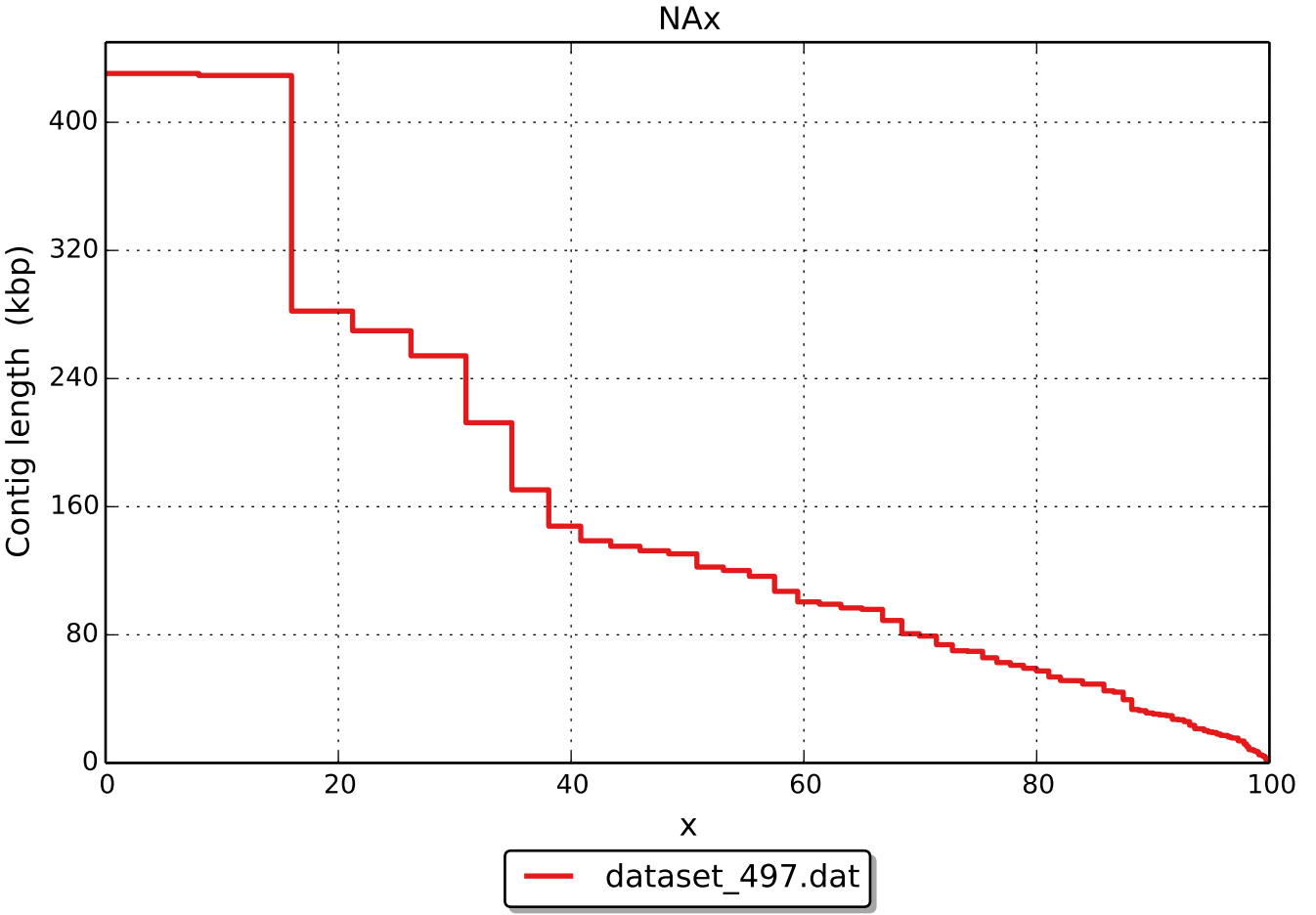

Cumulative length
Cumulative length (Mbp)
Contig index
0
1
2
3
4
5
6
0
20
40
60
80
dataset_497.dat
Reference

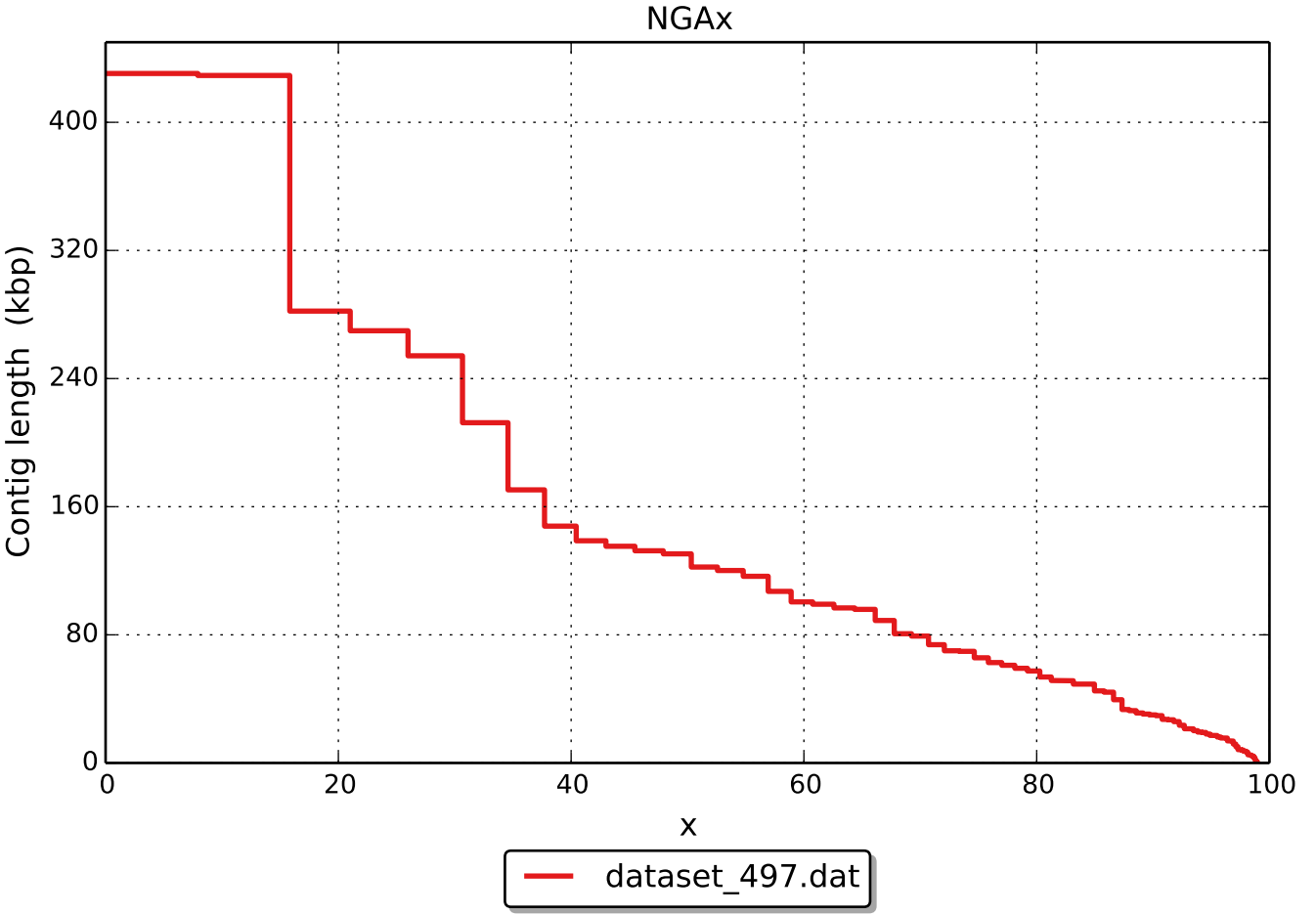

GC content
4000
3200
2400
1600
800
0
windows
0
20
40
60
80
100
GC (%)
dataset_497.dat
Reference

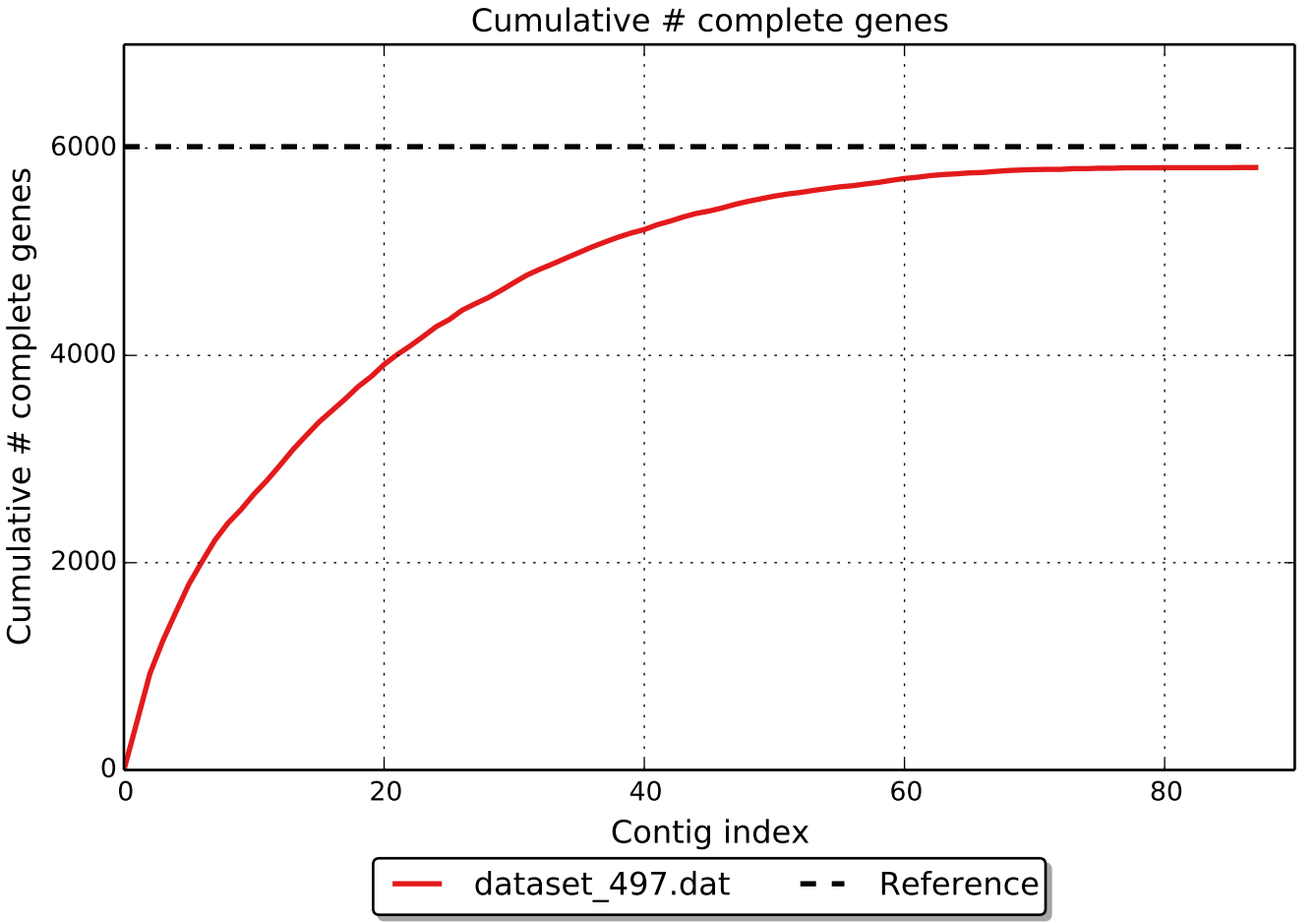

Nx
Contig length (kbp)
400
320
240
160
80
0
0
20
40
60
80
100
x
dataset_497.dat

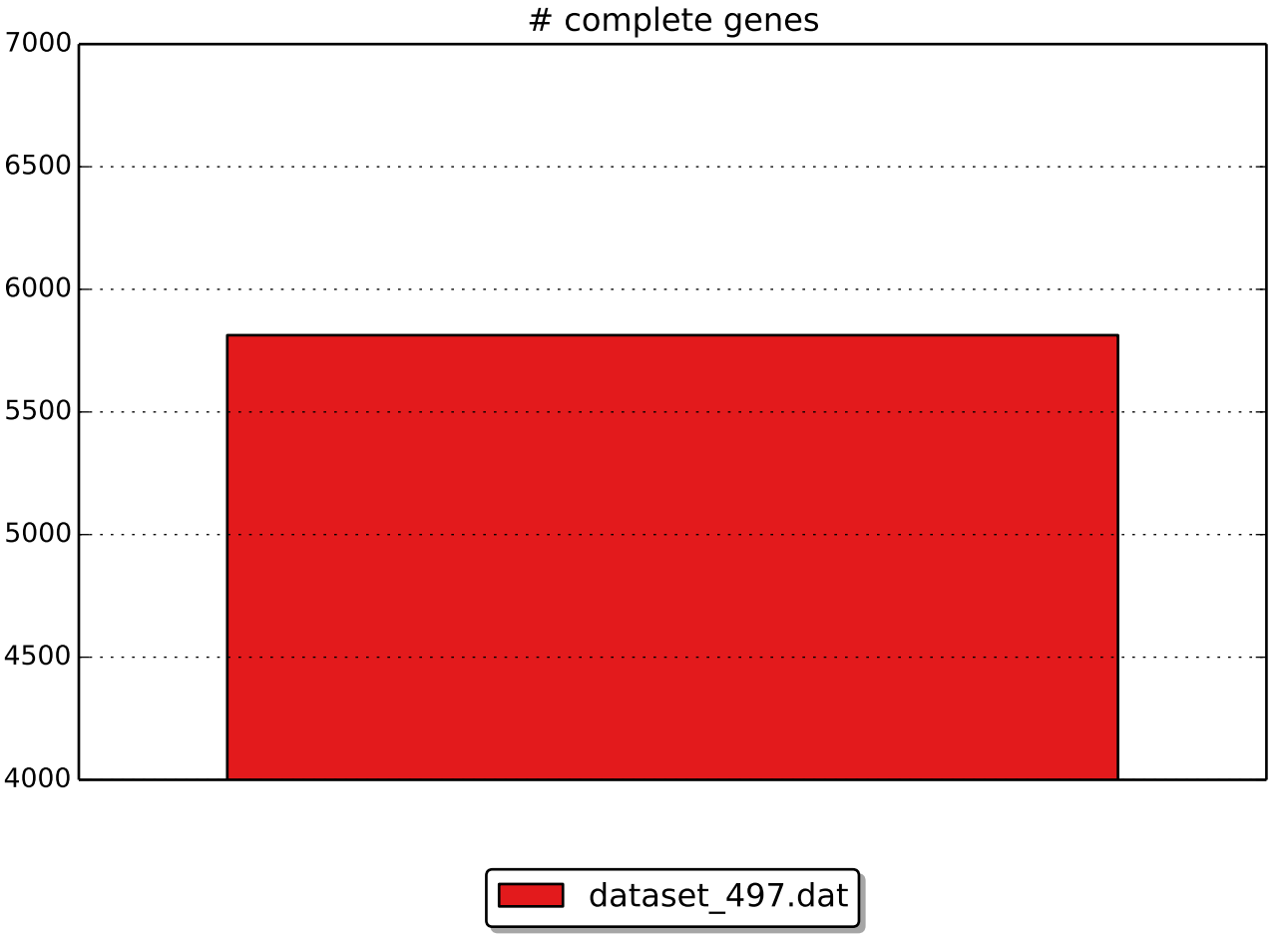

NGx
Contig length (kbp)
400
320
240
160
80
0
0
20
40
60
80
100
x
dataset_497.dat

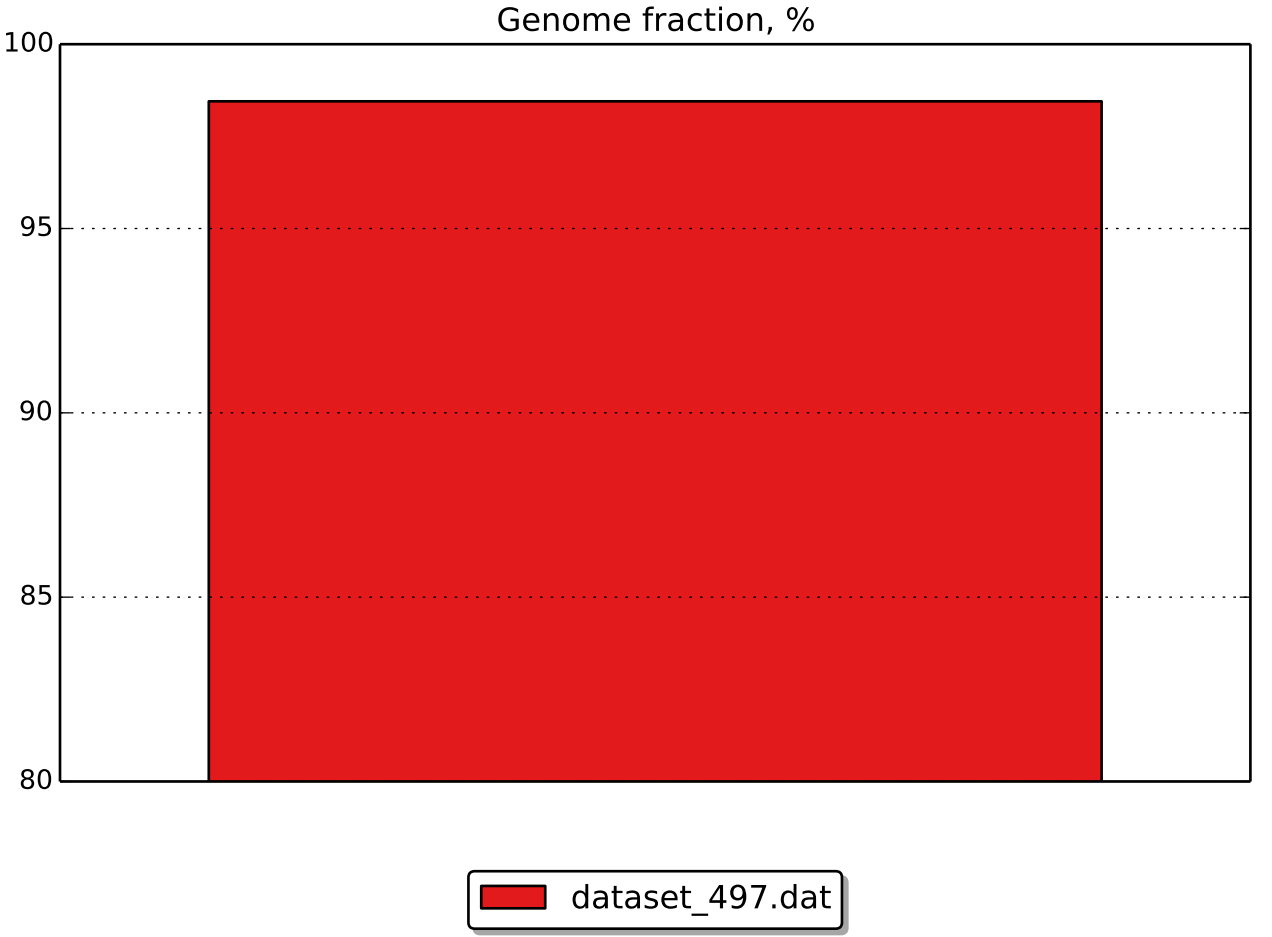

Cumulative length (aligned contigs)
Cumulative length (Mbp)
0
1
2
3
4
5
6
0
20
40
60
80
Contig index
dataset_497.dat
Reference


NAx
Contig length (kbp)
400
320
240
160
80
0
0
20
40
60
80
100
x
dataset_497.dat

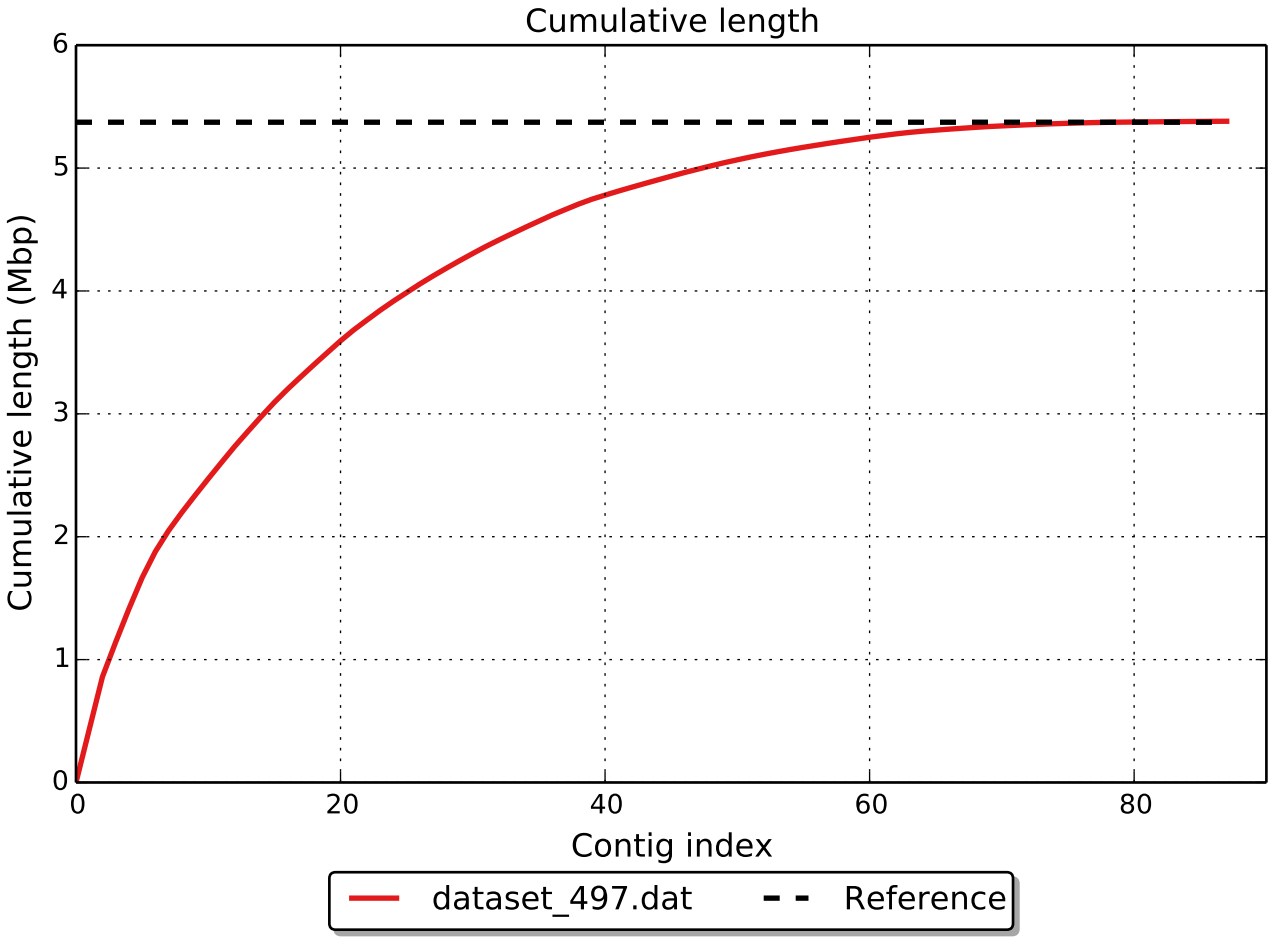

NGAx
Contig length (kbp)
400
320
240
160
80
0
0
20
40
60
80
100
x
dataset_497.dat

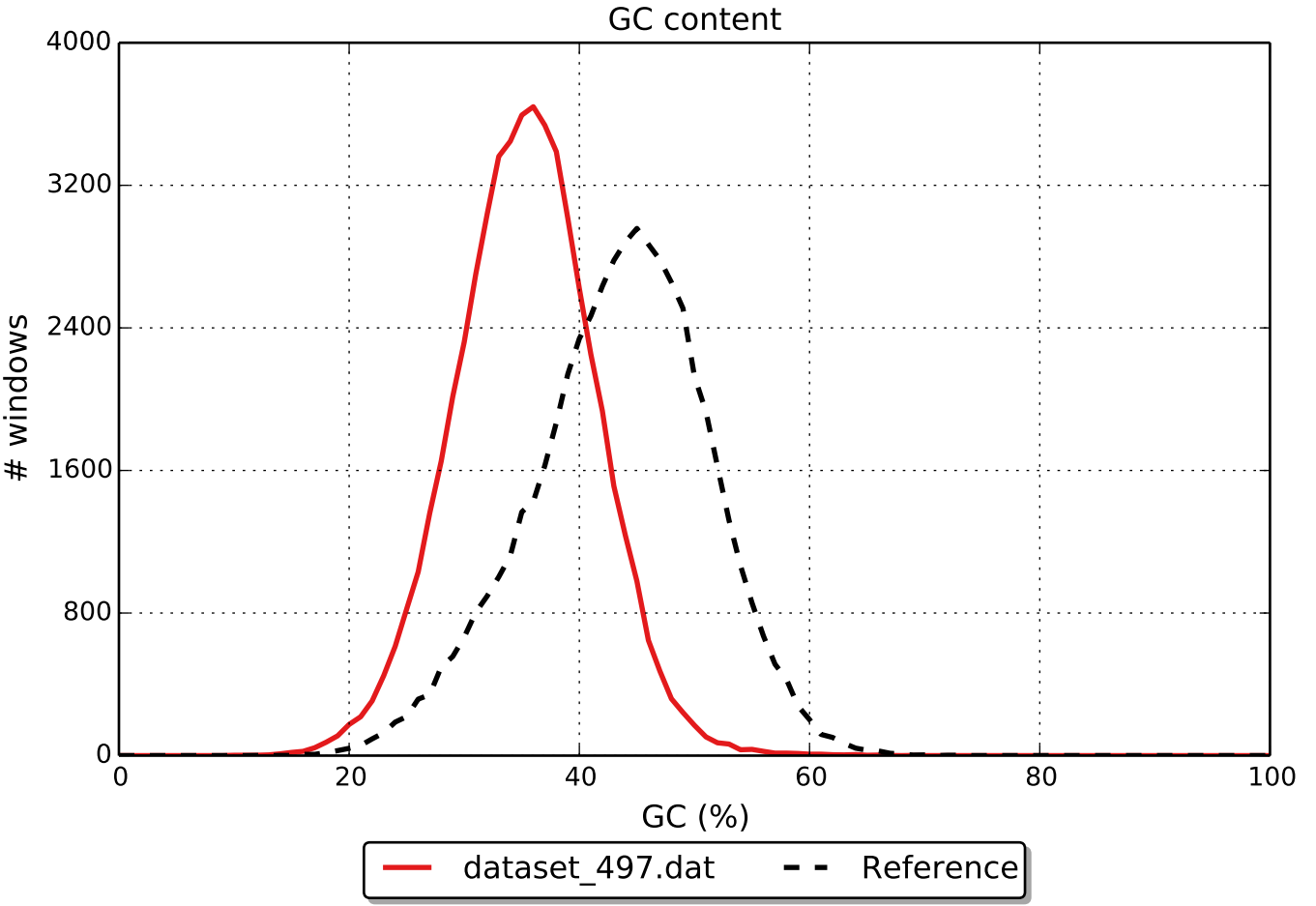

Cumulative # complete genes
Cumulative # complete genes
6000
4000
2000
0
0
20
40
60
80
Contig index
dataset_497.dat
Reference

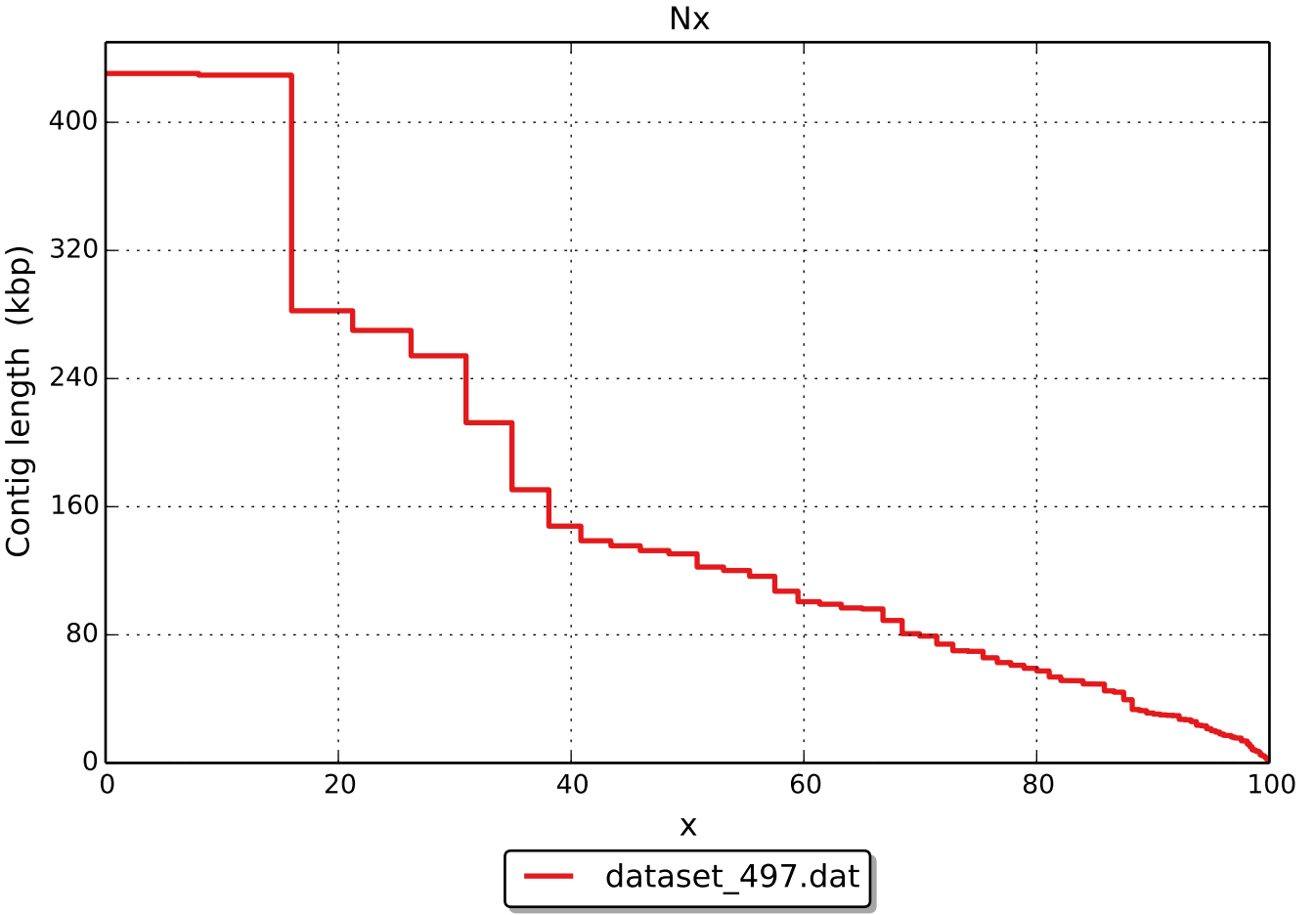

complete genes
7000
6500
6000
5500
5000
4500
4000
dataset_497.dat

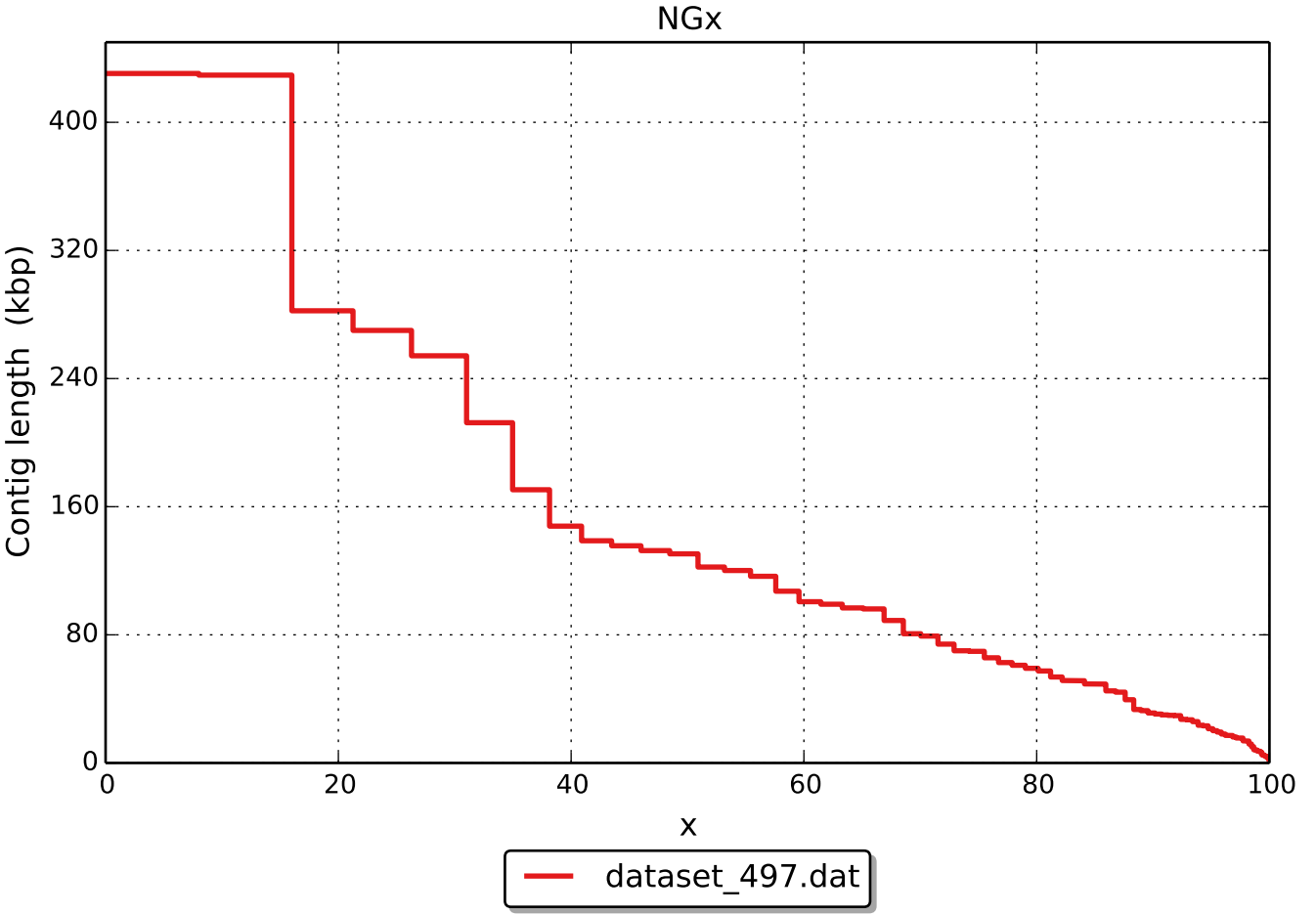

Genome fraction, %
100
95
90
85
80
dataset_497.dat

B_fragilis_638R


Cumulative length
Cumulative length (Mbp)
Contig index
0
1
2
3
4
5
6
0
20
40
60
80
dataset_497.dat
Reference

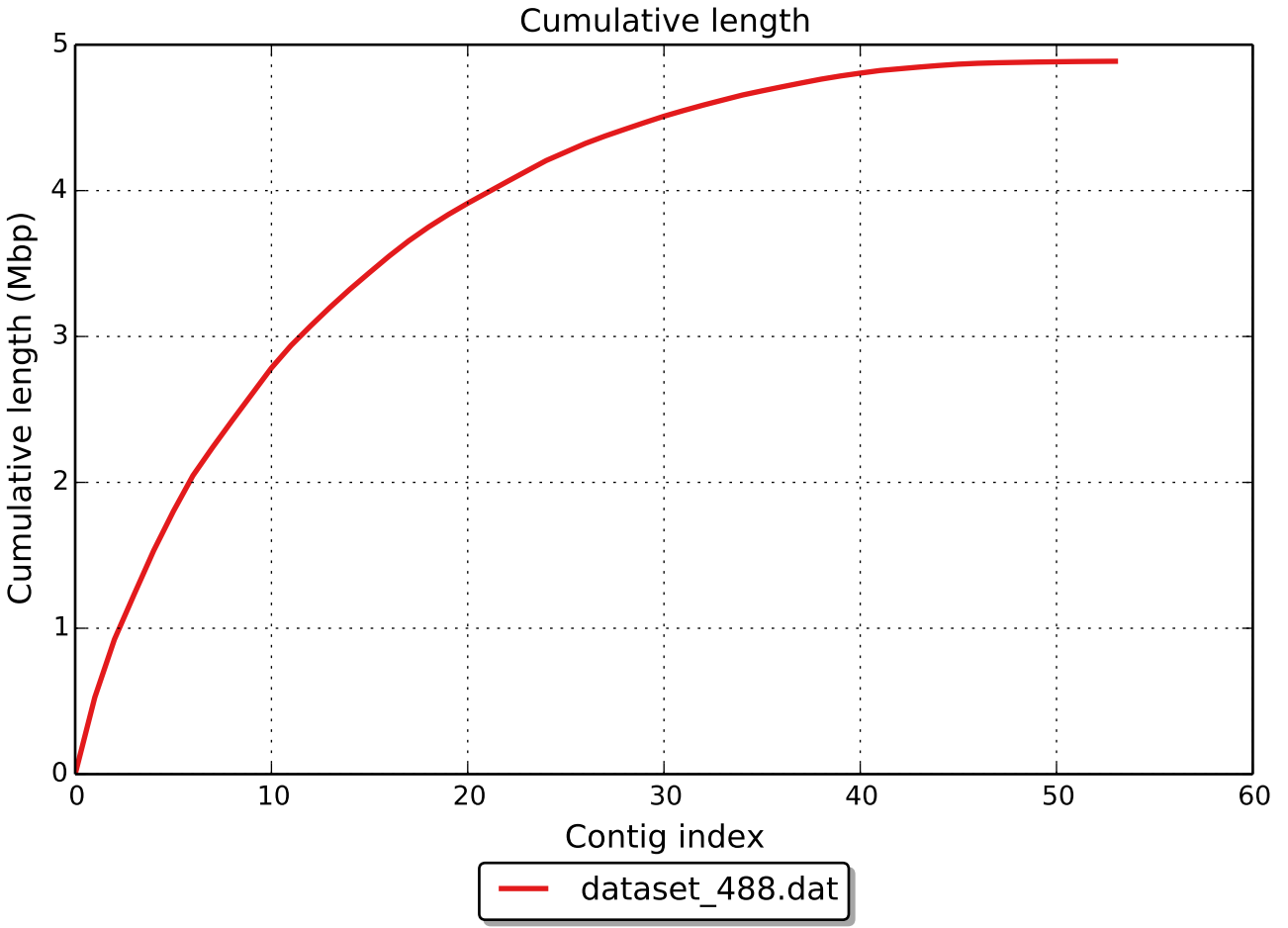

GC content
4000
3200
2400
1600
800
0
windows
0
20
40
60
80
100
GC (%)
dataset_497.dat
Reference

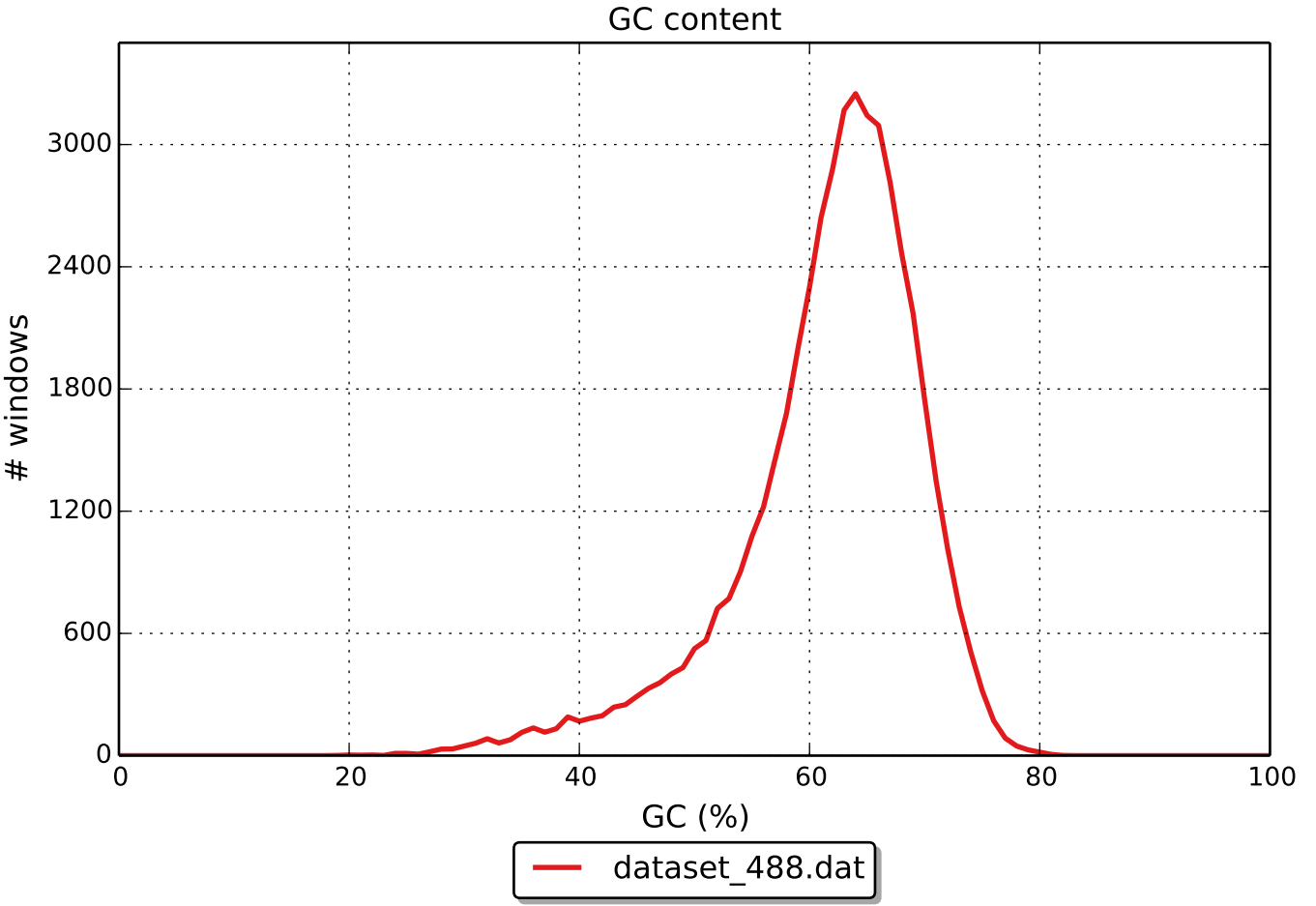

Nx
Contig length (kbp)
400
320
240
160
80
0
0
20
40
60
80
100
x
dataset_497.dat

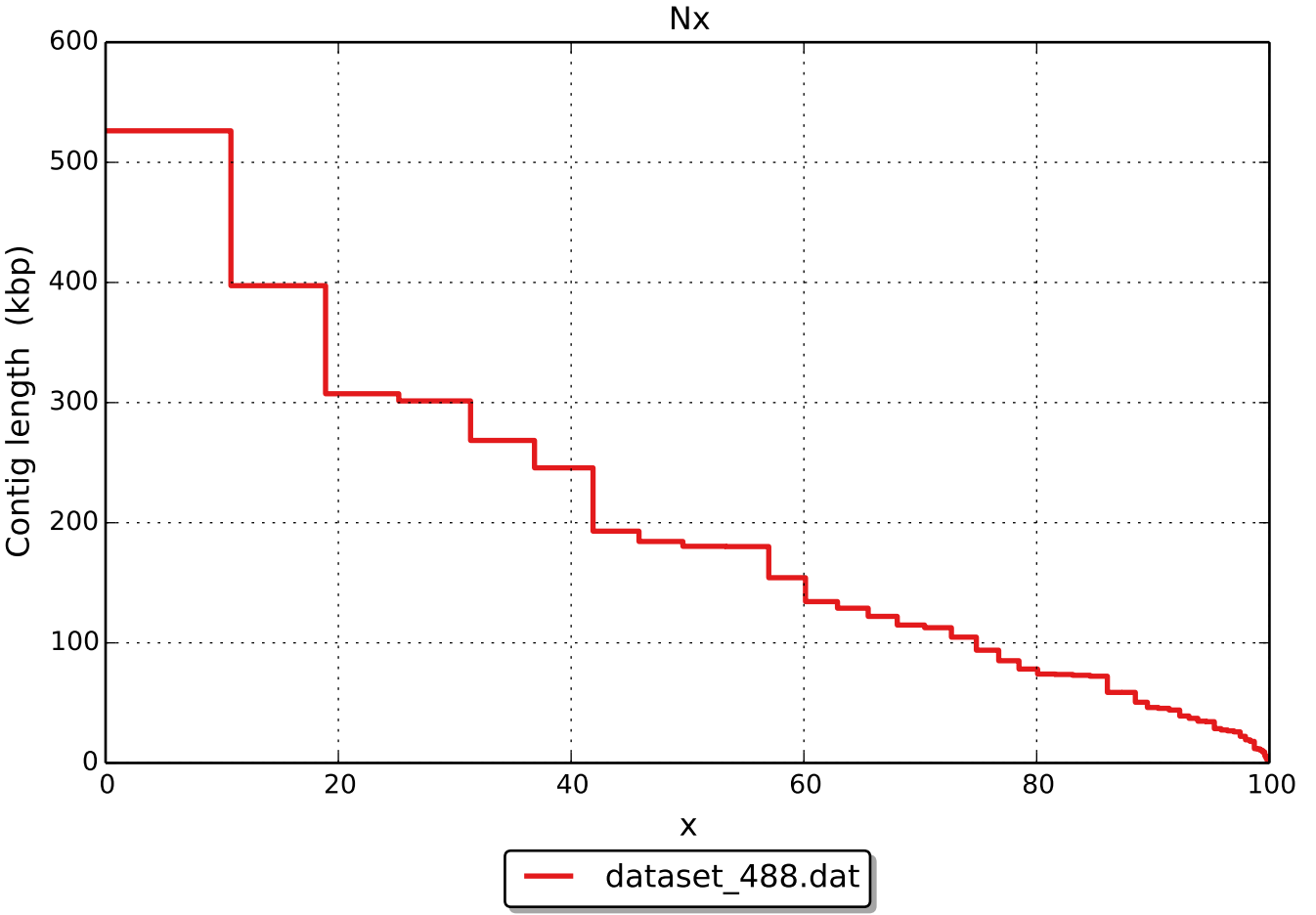

NGx
Contig length (kbp)
400
320
240
160
80
0
0
20
40
60
80
100
x
dataset_497.dat

Combined_history

Contig

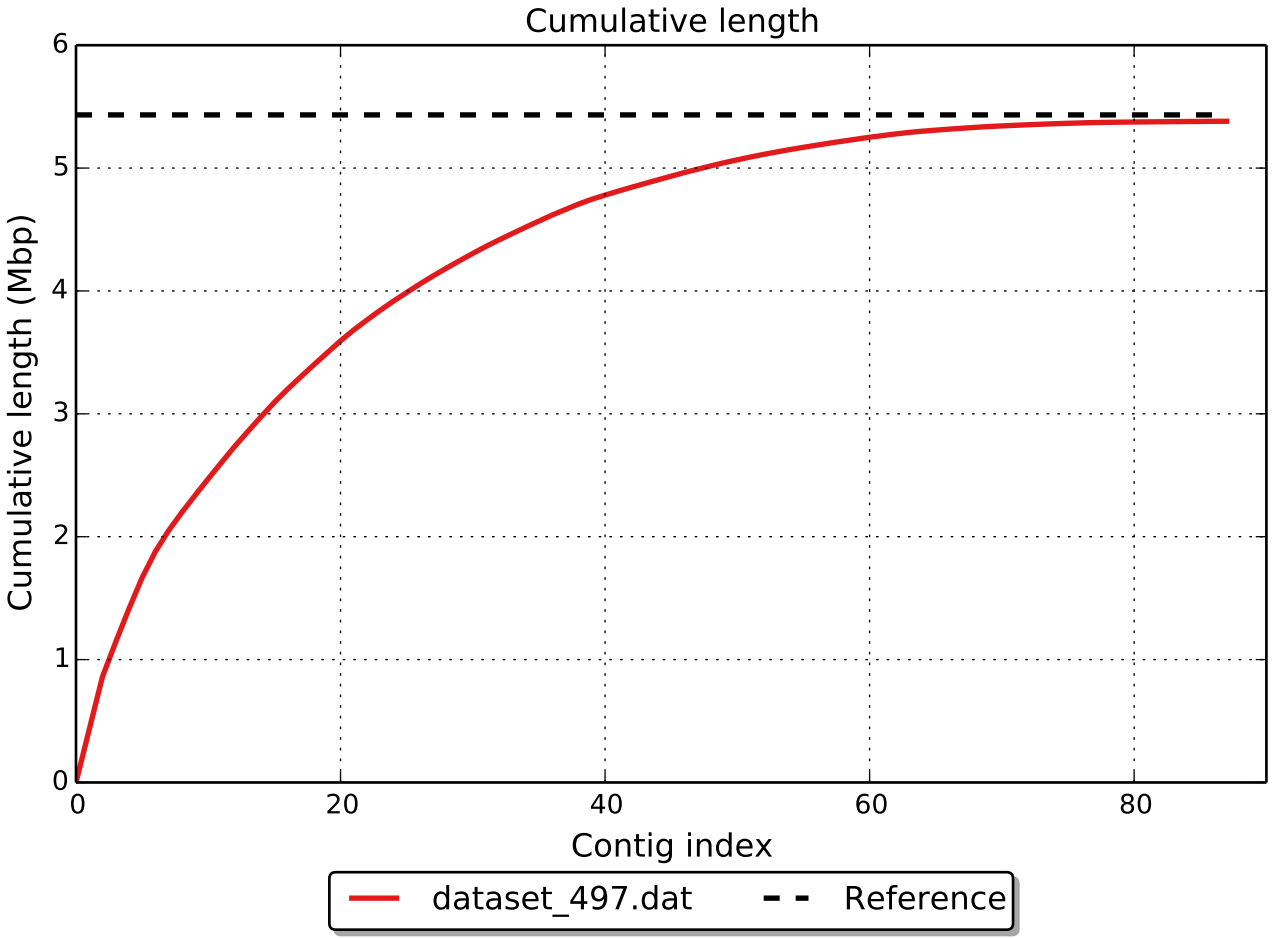

Cumulative length
Cumulative length (Mbp)
0
1
2
3
4
5
0
10
20
30
40
50
60
Contig index
dataset_488.dat

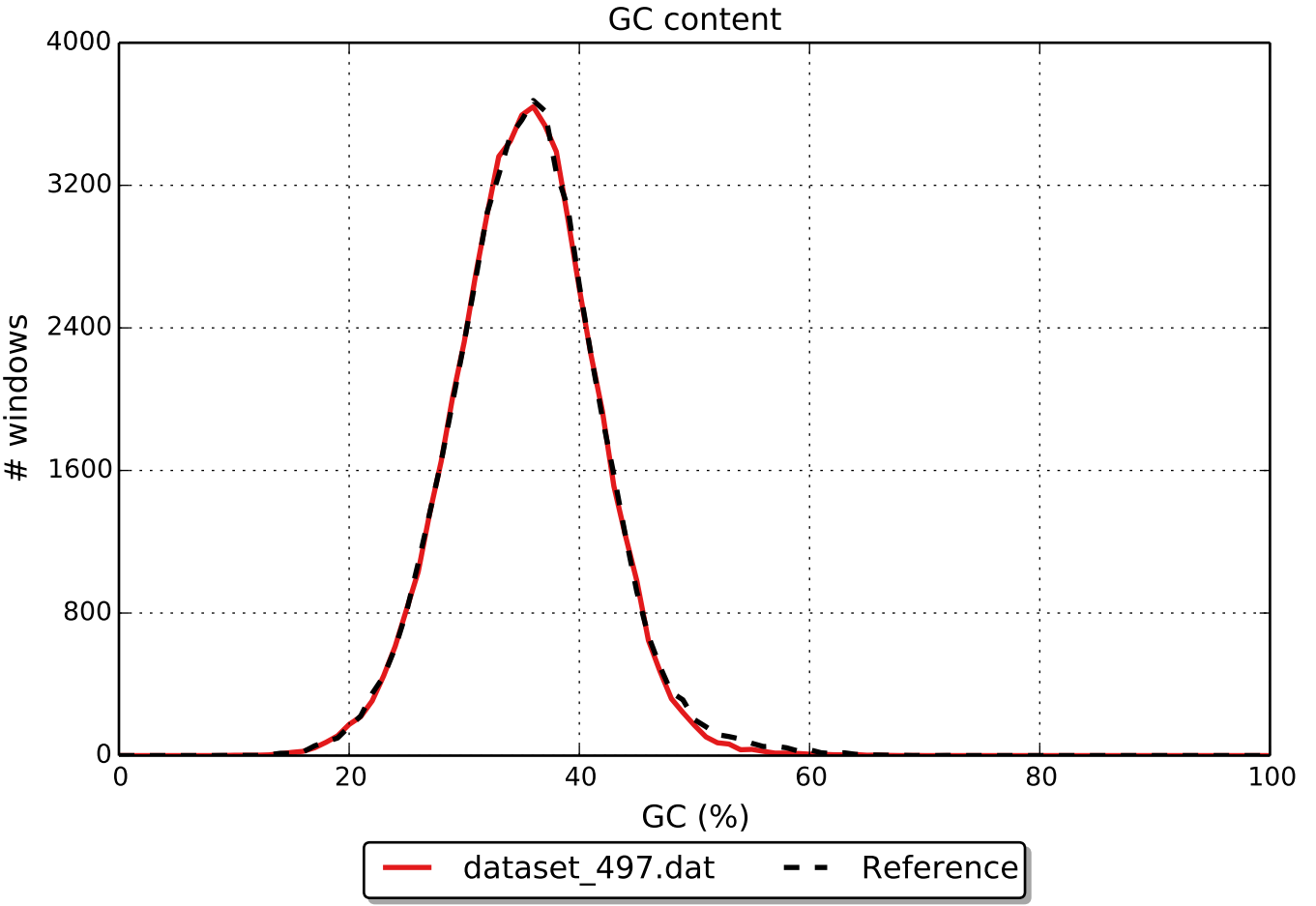

GC content
windows
3000
2400
1800
1200
600
0
0
20
40
60
80
100
GC (%)
dataset_488.dat

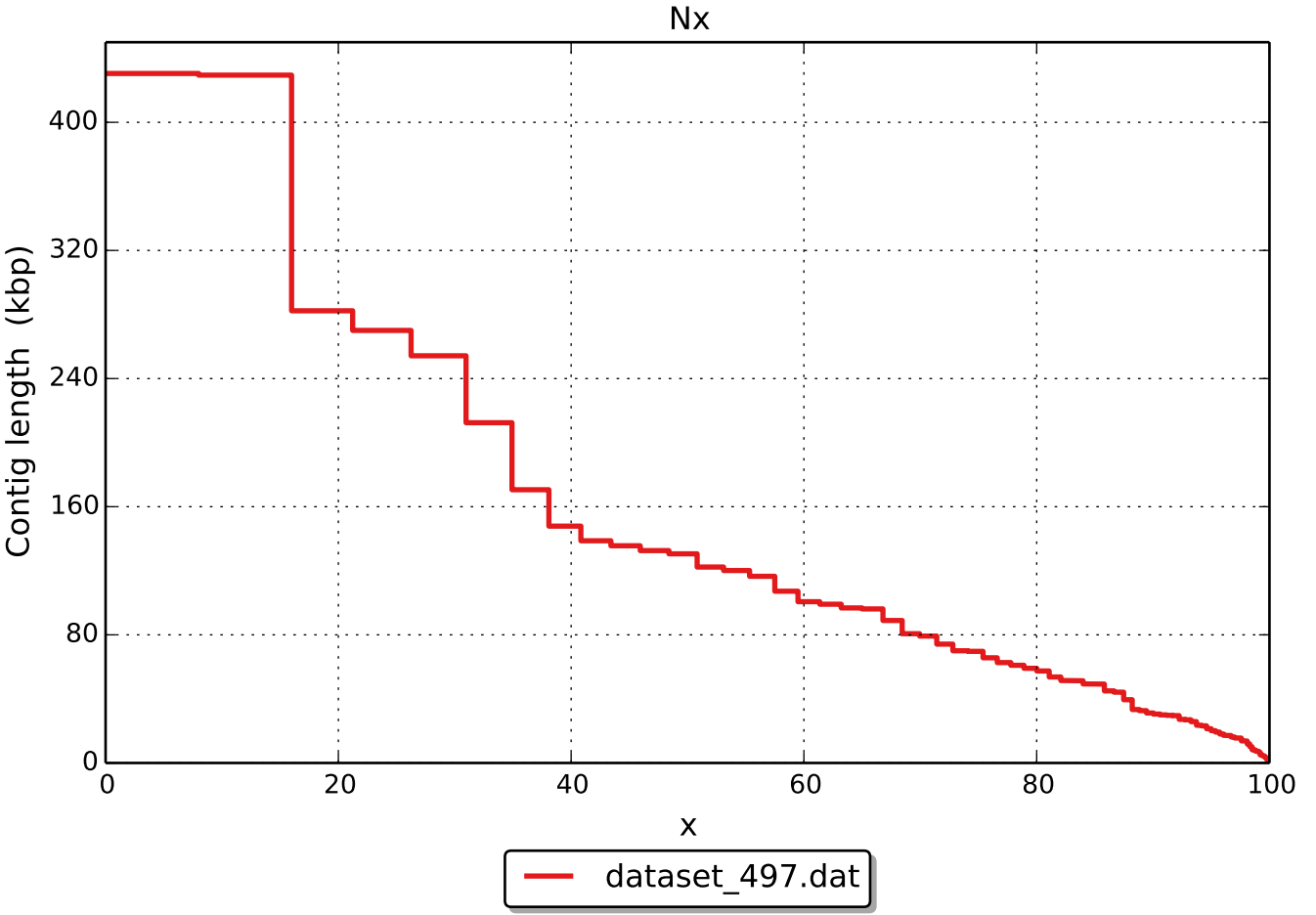

Nx
Contig length (kbp)
600
500
400
300
200
100
0
0
20
40
60
80
100
x
dataset_488.dat

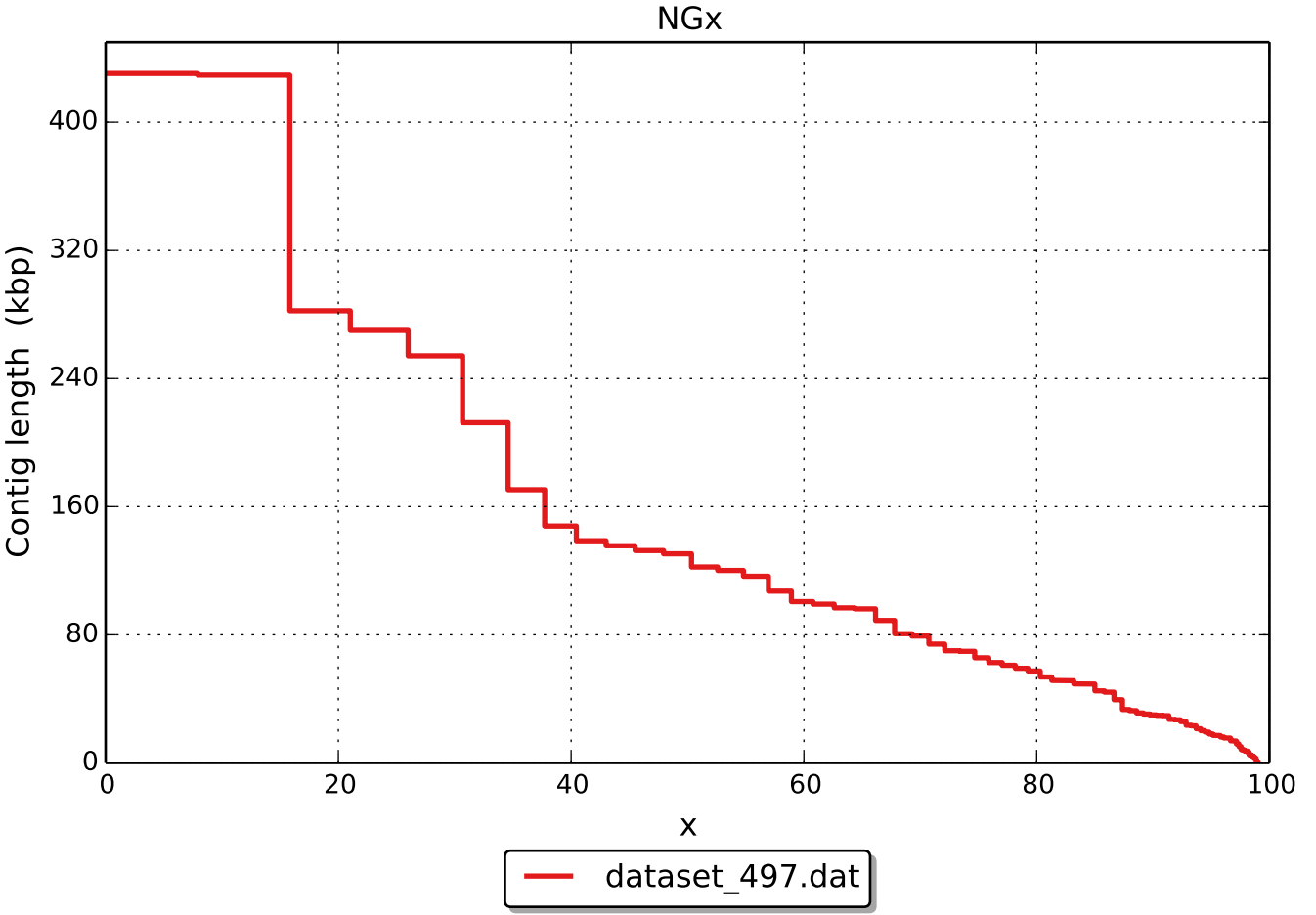
Combined_reference

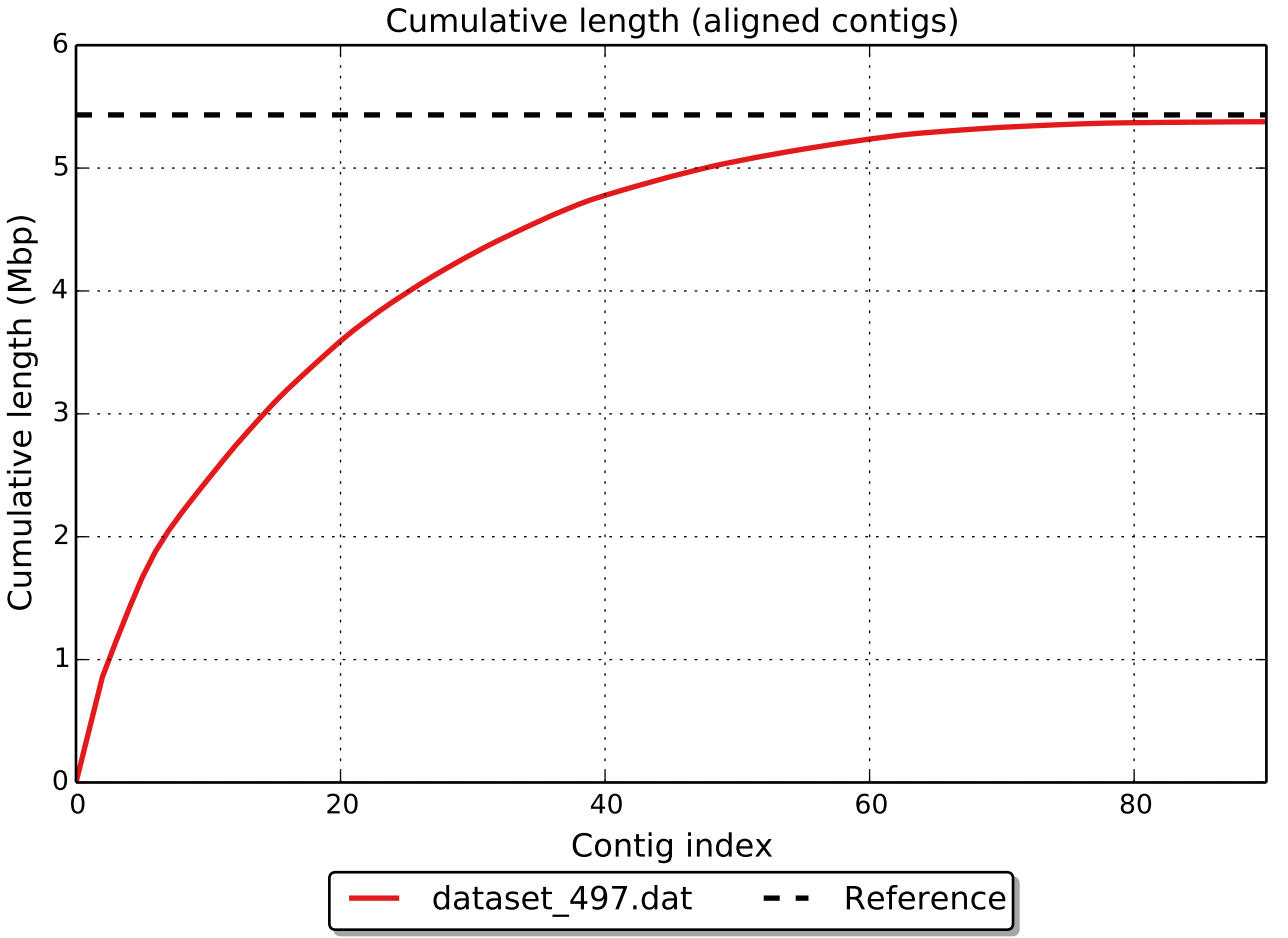
B_cereus_ATCC_1098

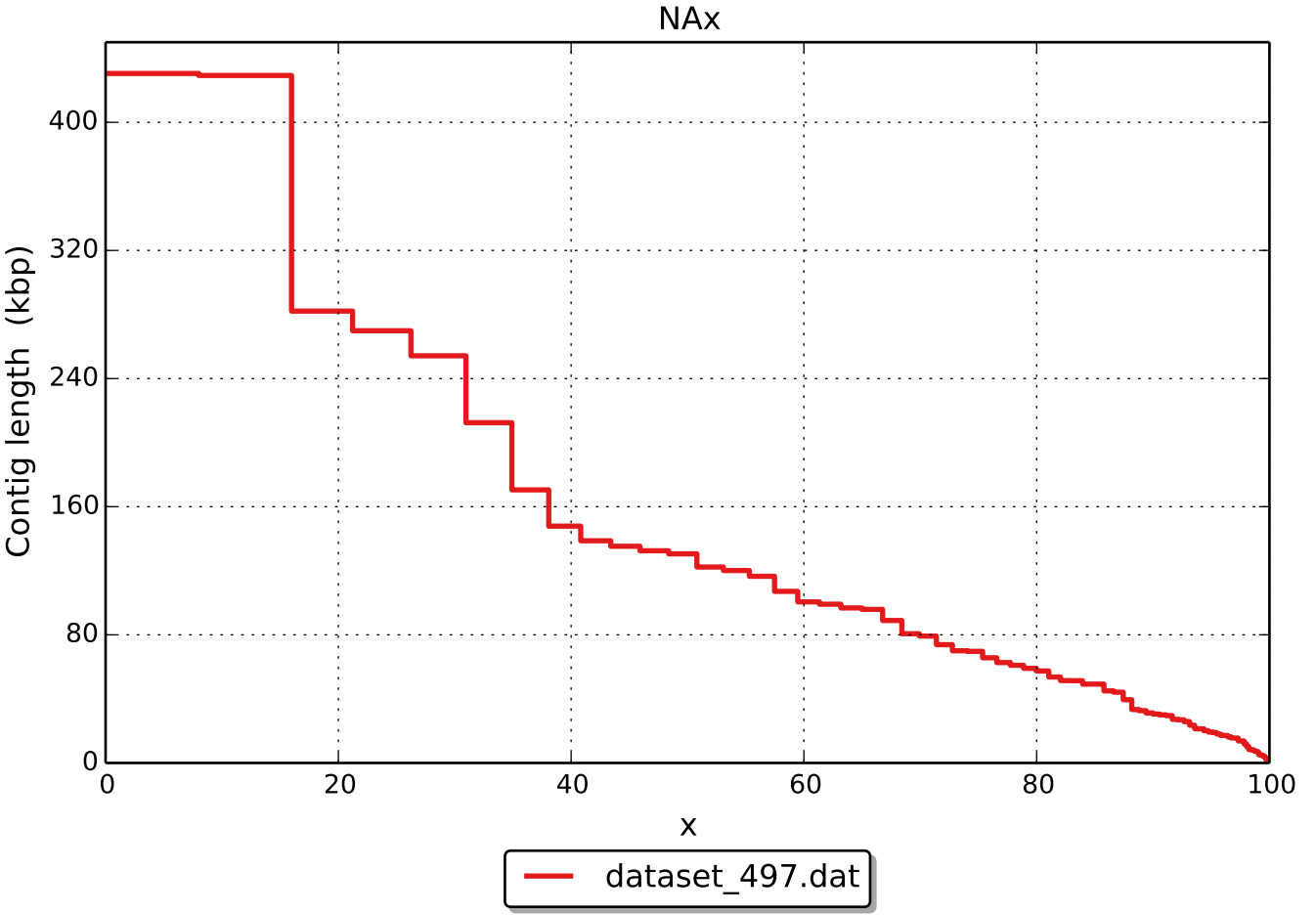

Cumulative length
Cumulative length (Mbp)
Contig index
0
1
2
3
4
5
6
0
20
40
60
80
dataset_497.dat
Reference

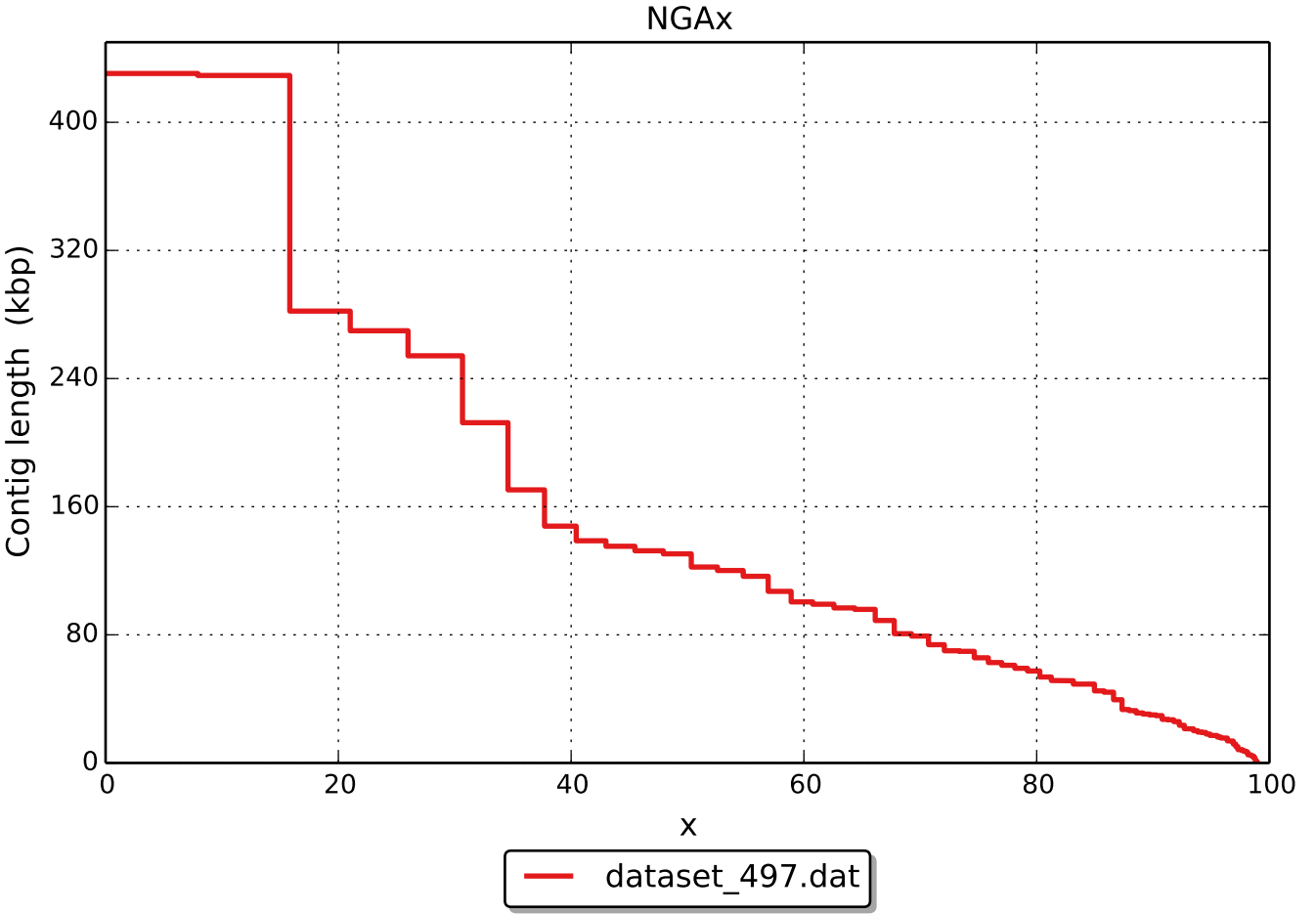

GC content
4000
3200
2400
1600
800
0
windows
0
20
40
60
80
100
GC (%)
dataset_497.dat
Reference

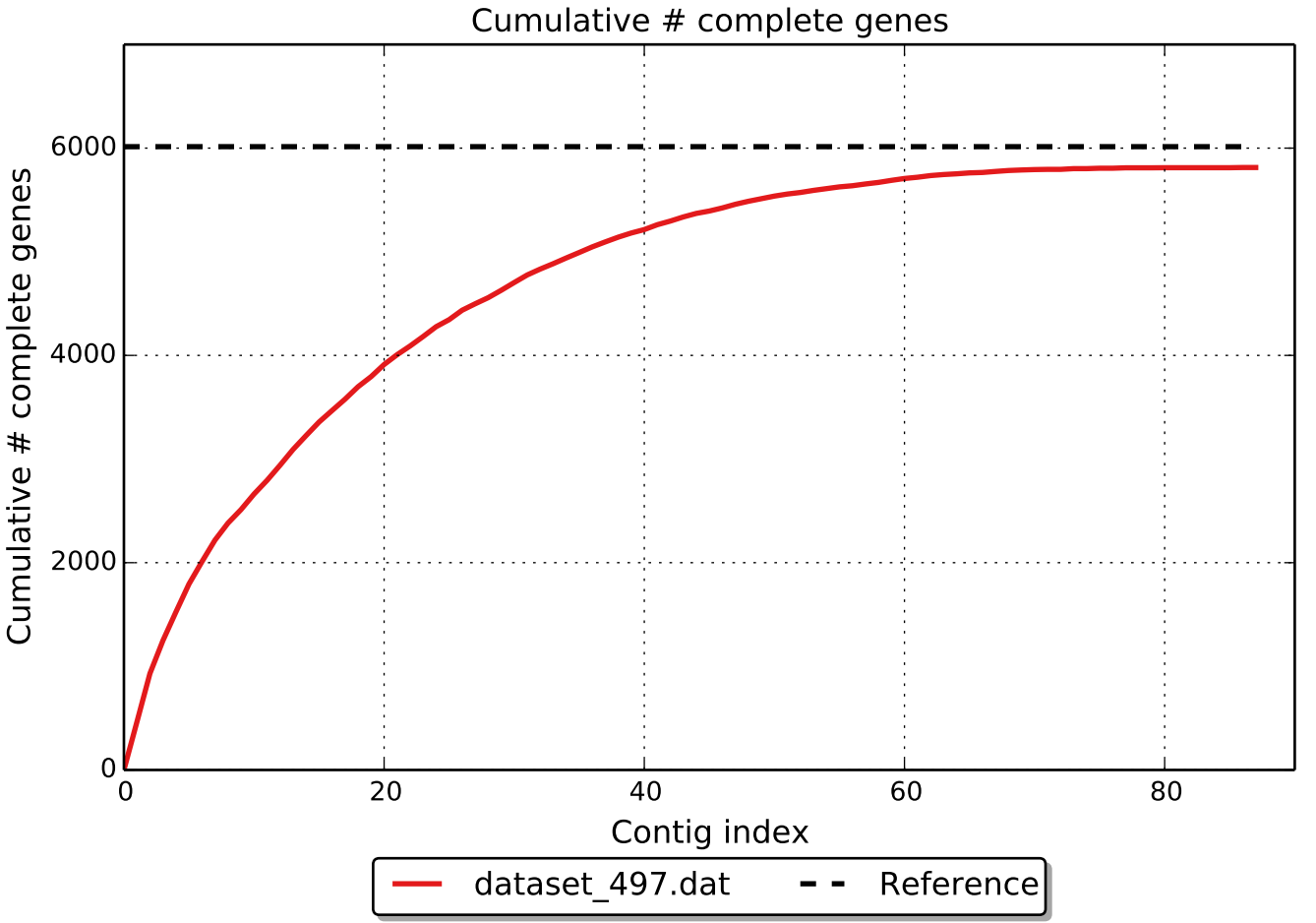

Nx
Contig length (kbp)
400
320
240
160
80
0
0
20
40
60
80
100
x
dataset_497.dat

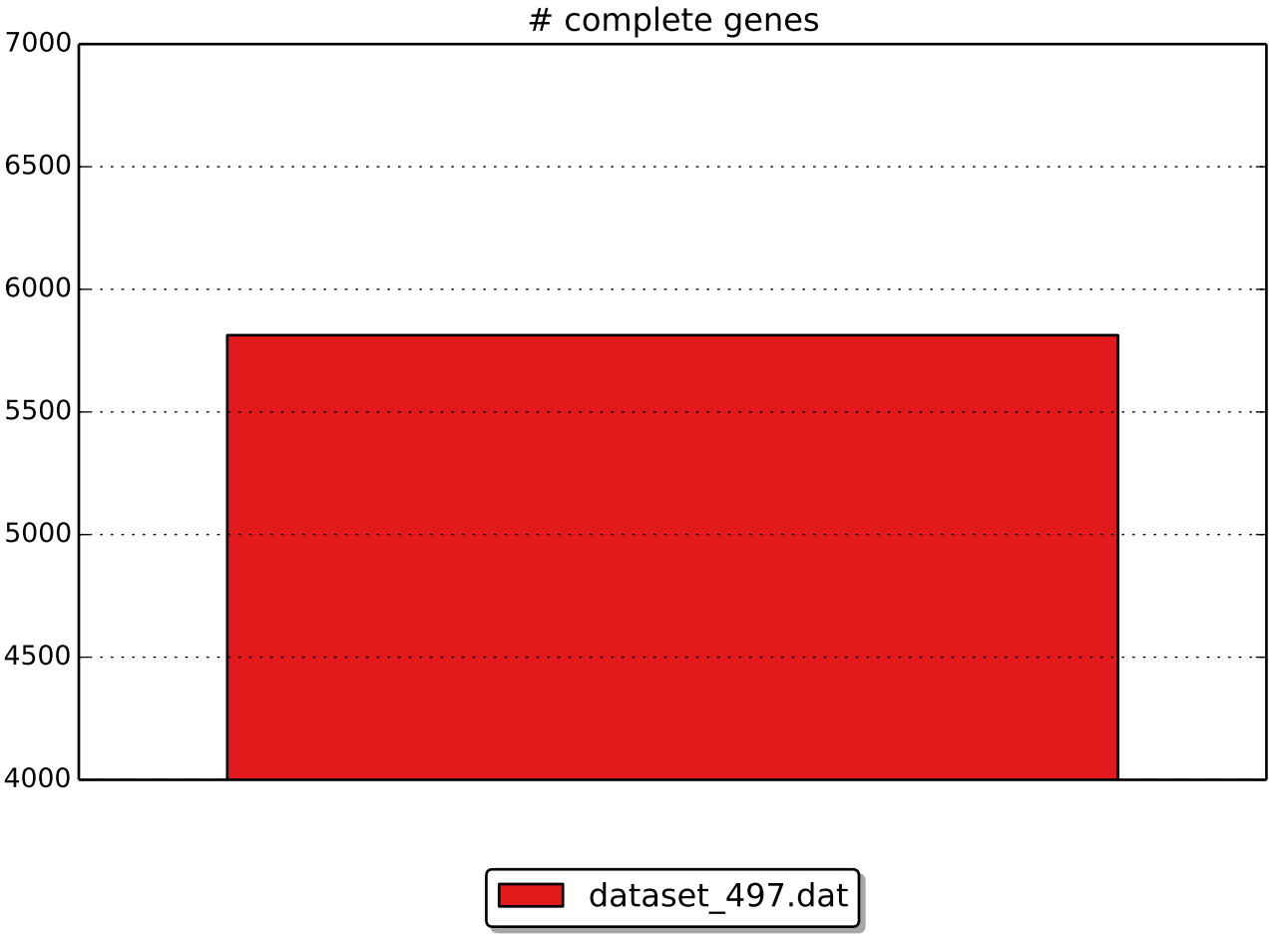

NGx
Contig length (kbp)
400
320
240
160
80
0
0
20
40
60
80
100
x
dataset_497.dat

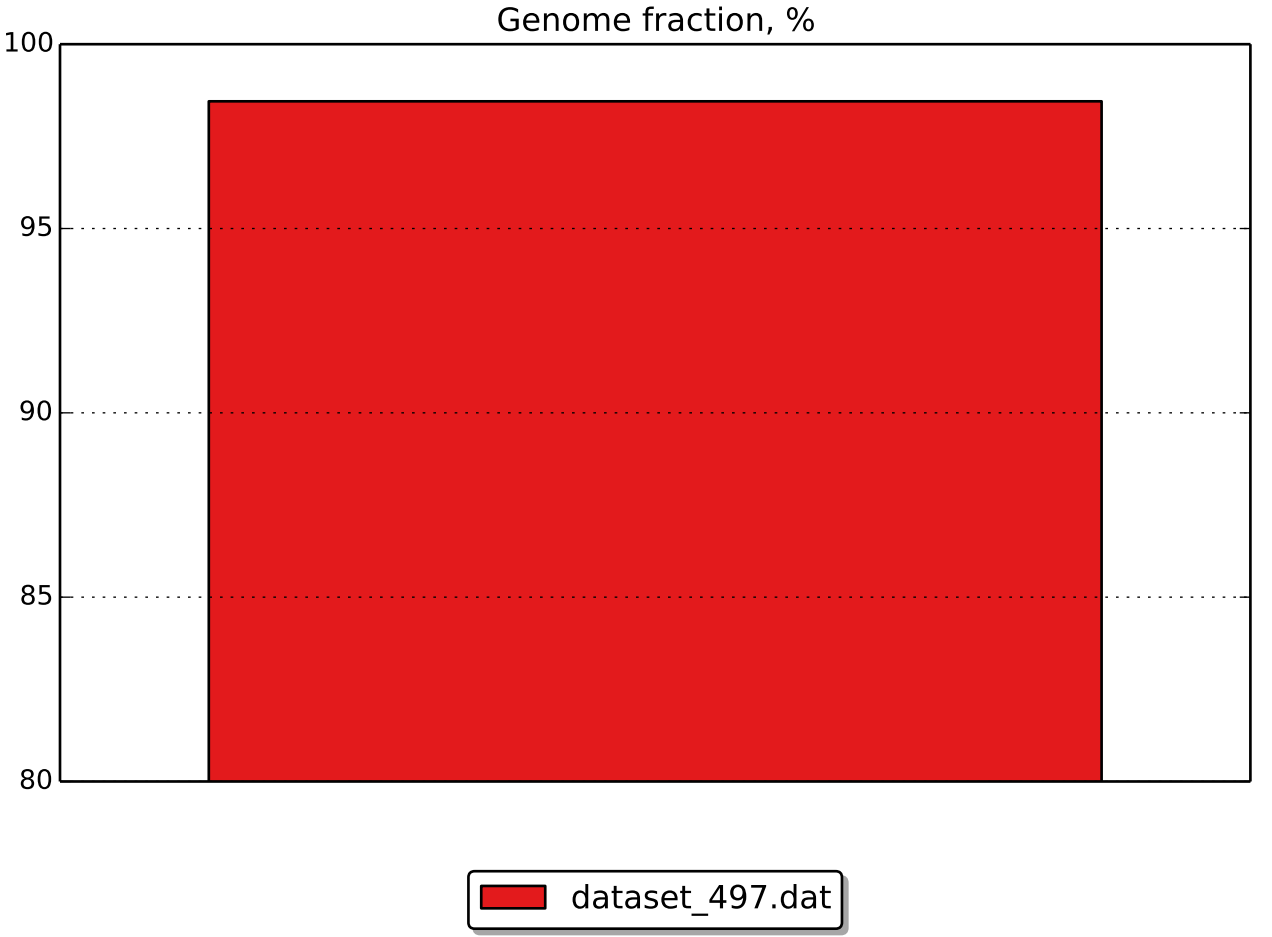

Cumulative length (aligned contigs)
Cumulative length (Mbp)
0
1
2
3
4
5
6
0
20
40
60
80
Contig index
dataset_497.dat
Reference


NAx
Contig length (kbp)
400
320
240
160
80
0
0
20
40
60
80
100
x
dataset_497.dat

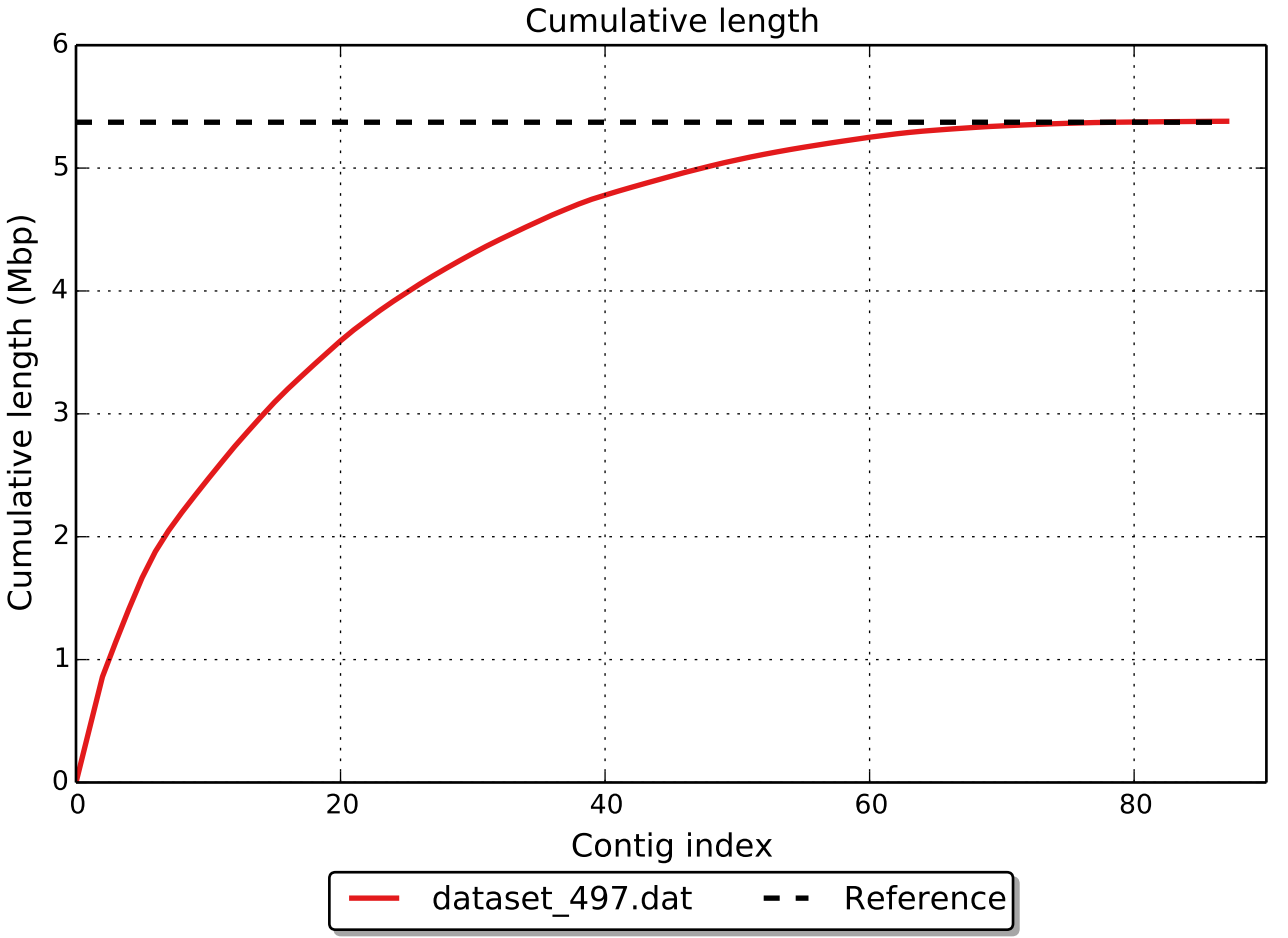

NGAx
Contig length (kbp)
400
320
240
160
80
0
0
20
40
60
80
100
x
dataset_497.dat

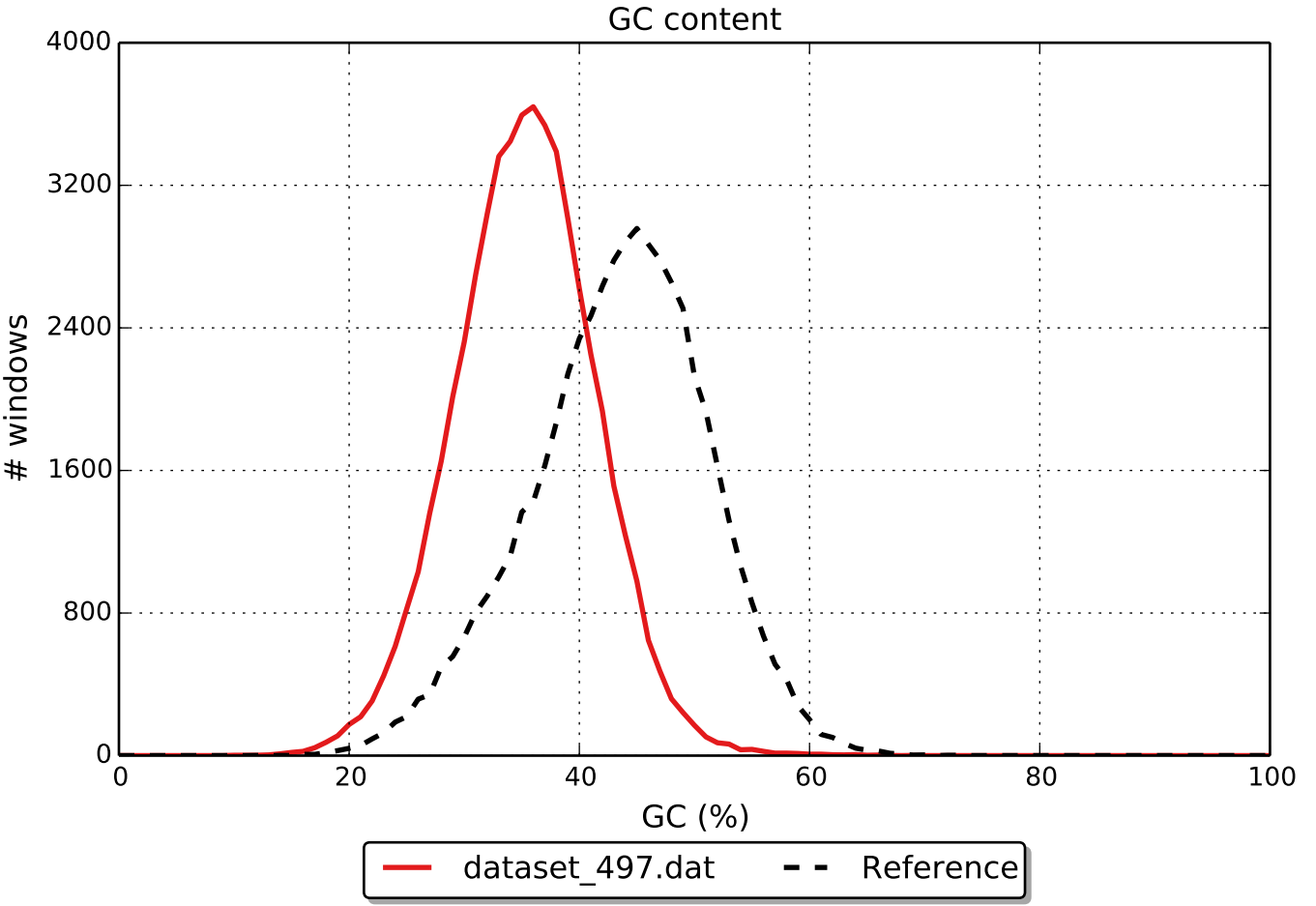

Cumulative # complete genes
Cumulative # complete genes
6000
4000
2000
0
0
20
40
60
80
Contig index
dataset_497.dat
Reference

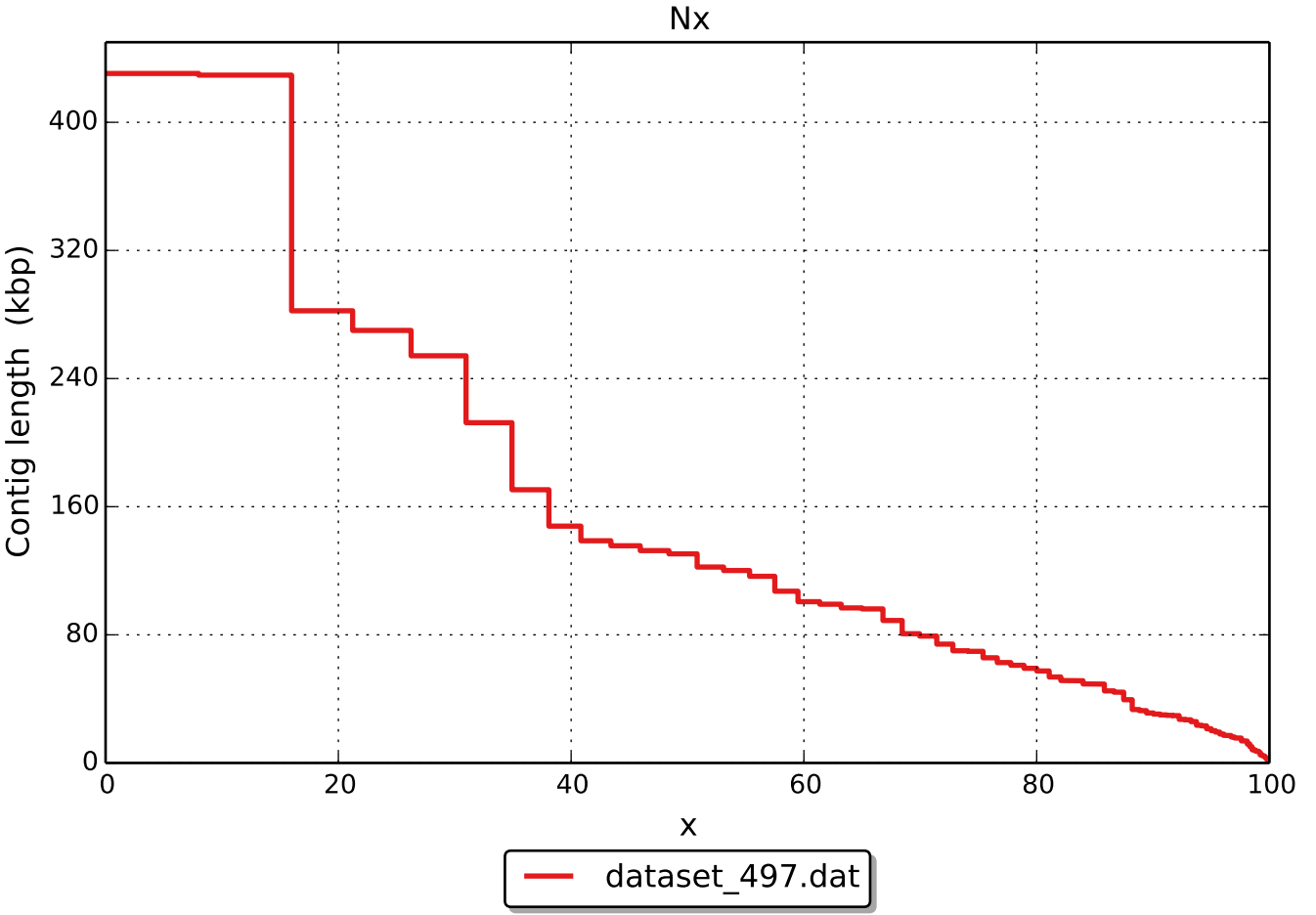

complete genes
7000
6500
6000
5500
5000
4500
4000
dataset_497.dat

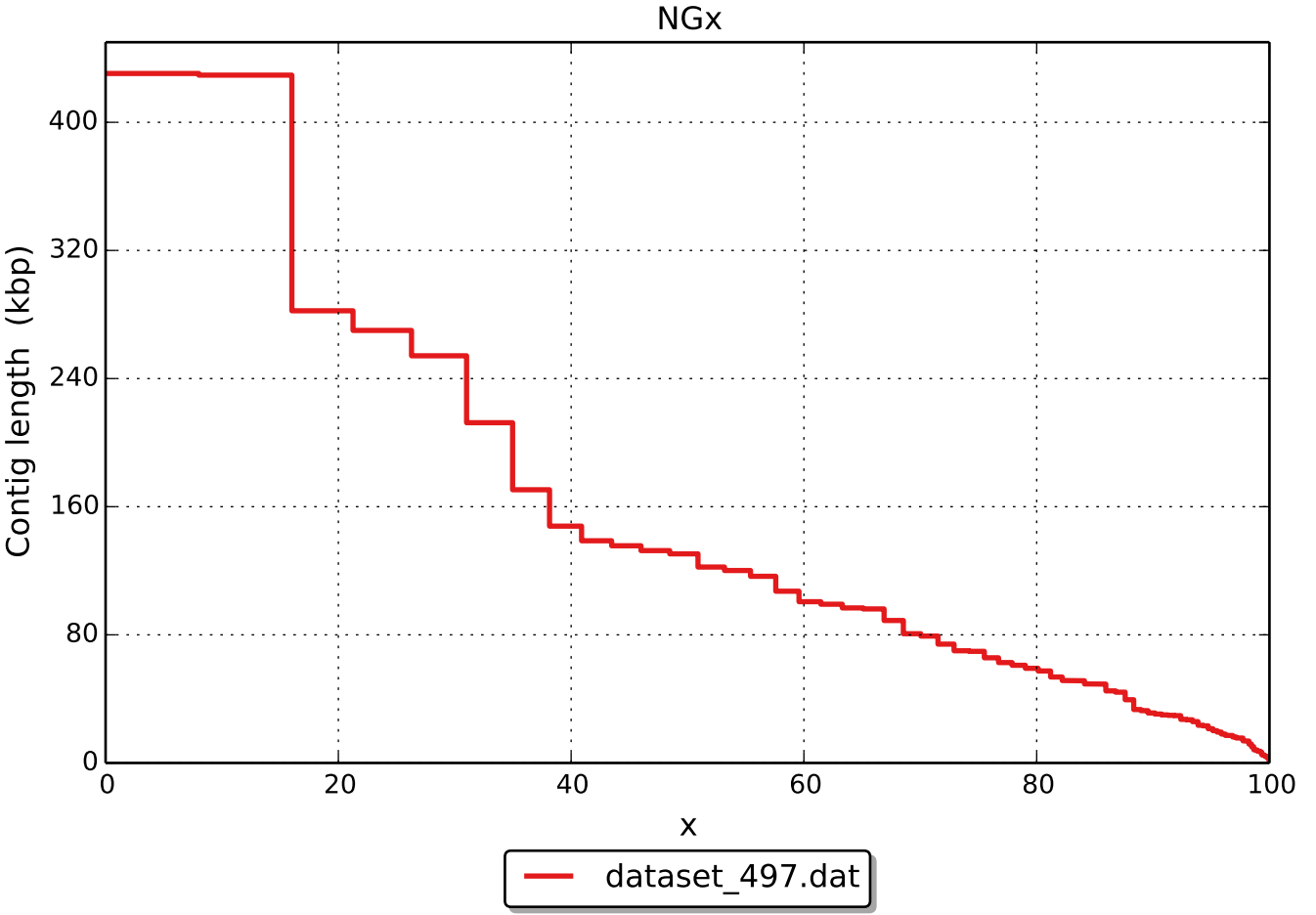

Genome fraction, %
100
95
90
85
80
dataset_497.dat

B_fragilis_638R


Cumulative length
Cumulative length (Mbp)
Contig index
0
1
2
3
4
5
6
0
20
40
60
80
dataset_497.dat
Reference


GC content
4000
3200
2400
1600
800
0
windows
0
20
40
60
80
100
GC (%)
dataset_497.dat
Reference


Nx
Contig length (kbp)
400
320
240
160
80
0
0
20
40
60
80
100
x
dataset_497.dat


NGx
Contig length (kbp)
400
320
240
160
80
0
0
20
40
60
80
100
x
dataset_497.dat