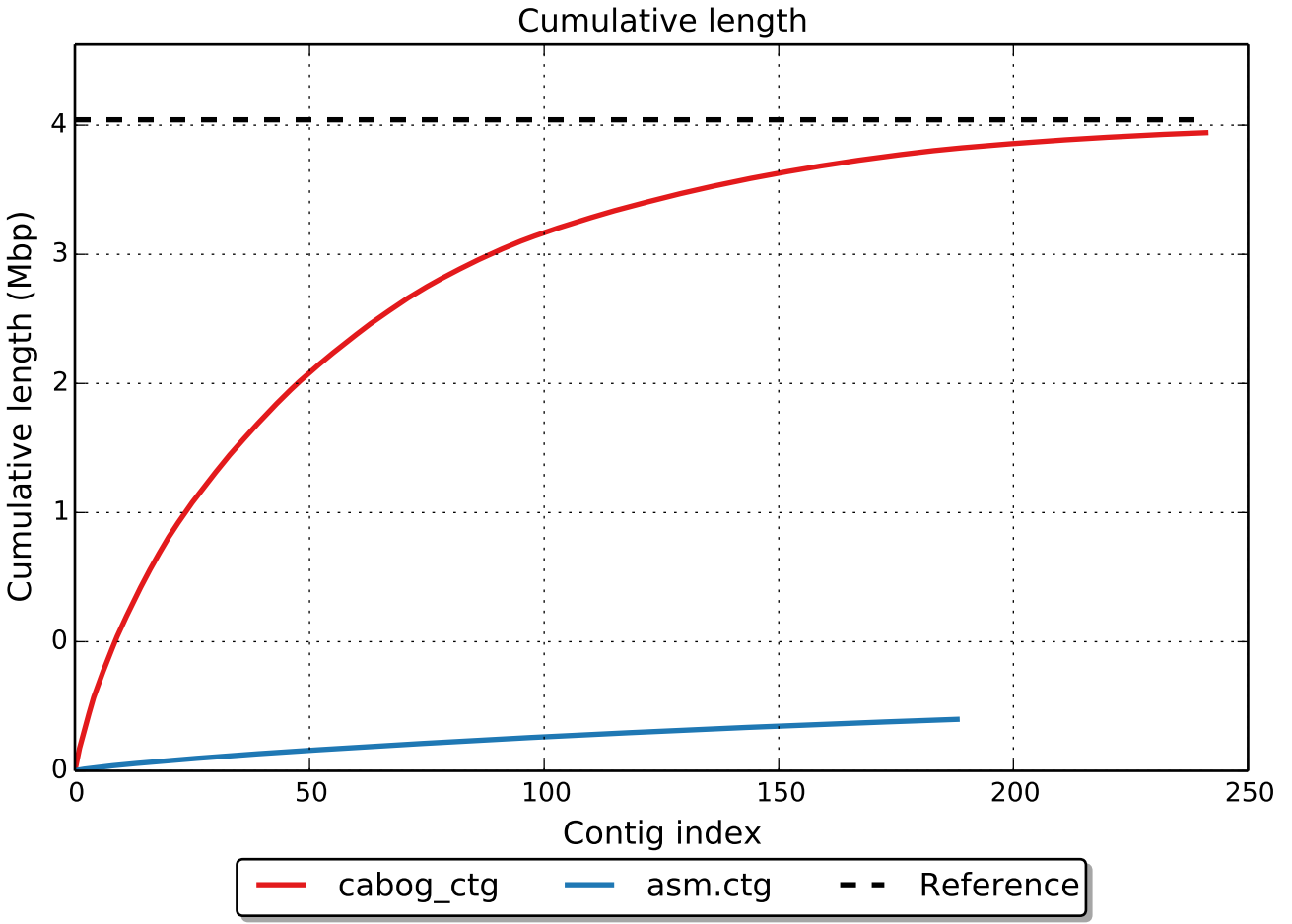
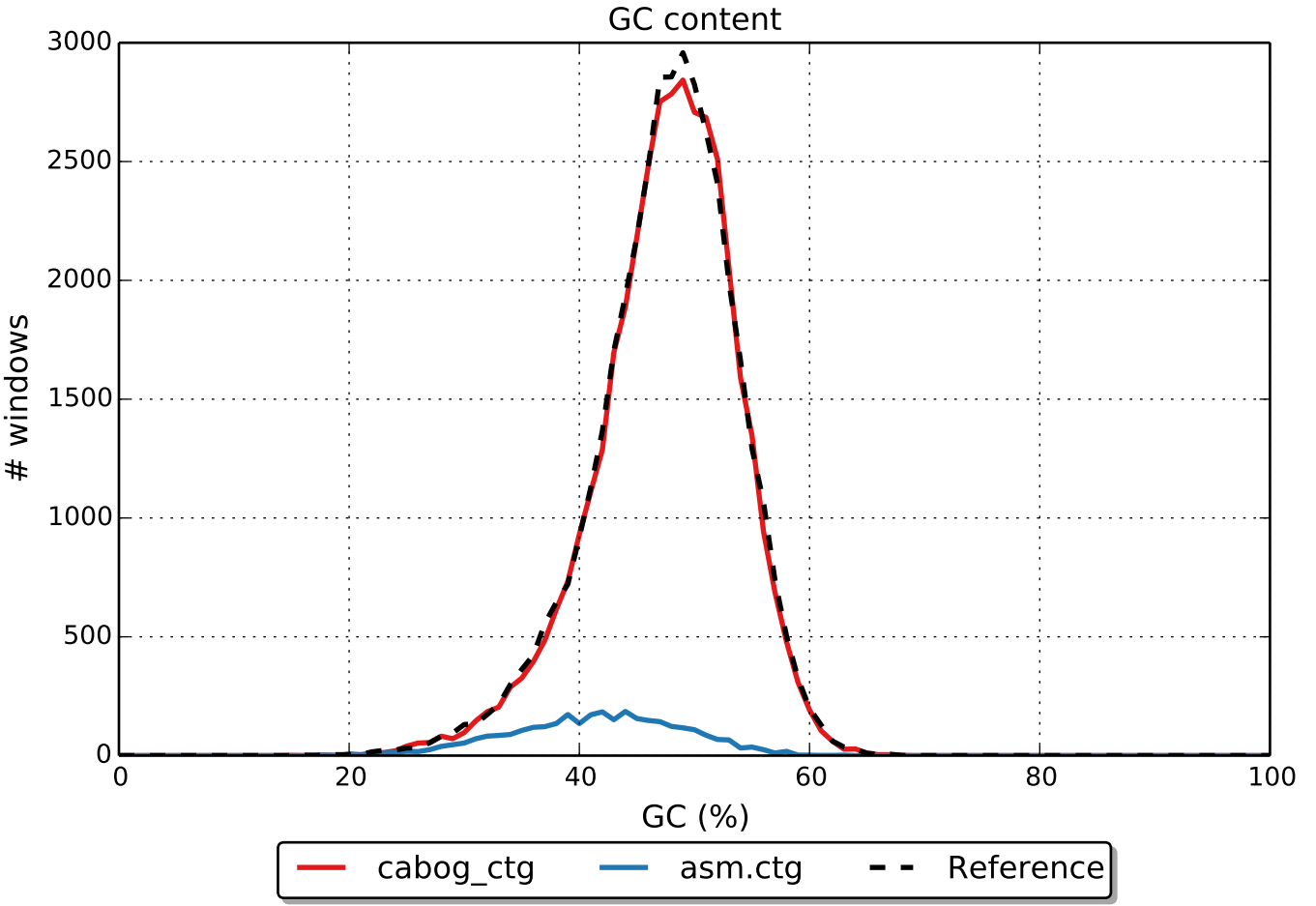
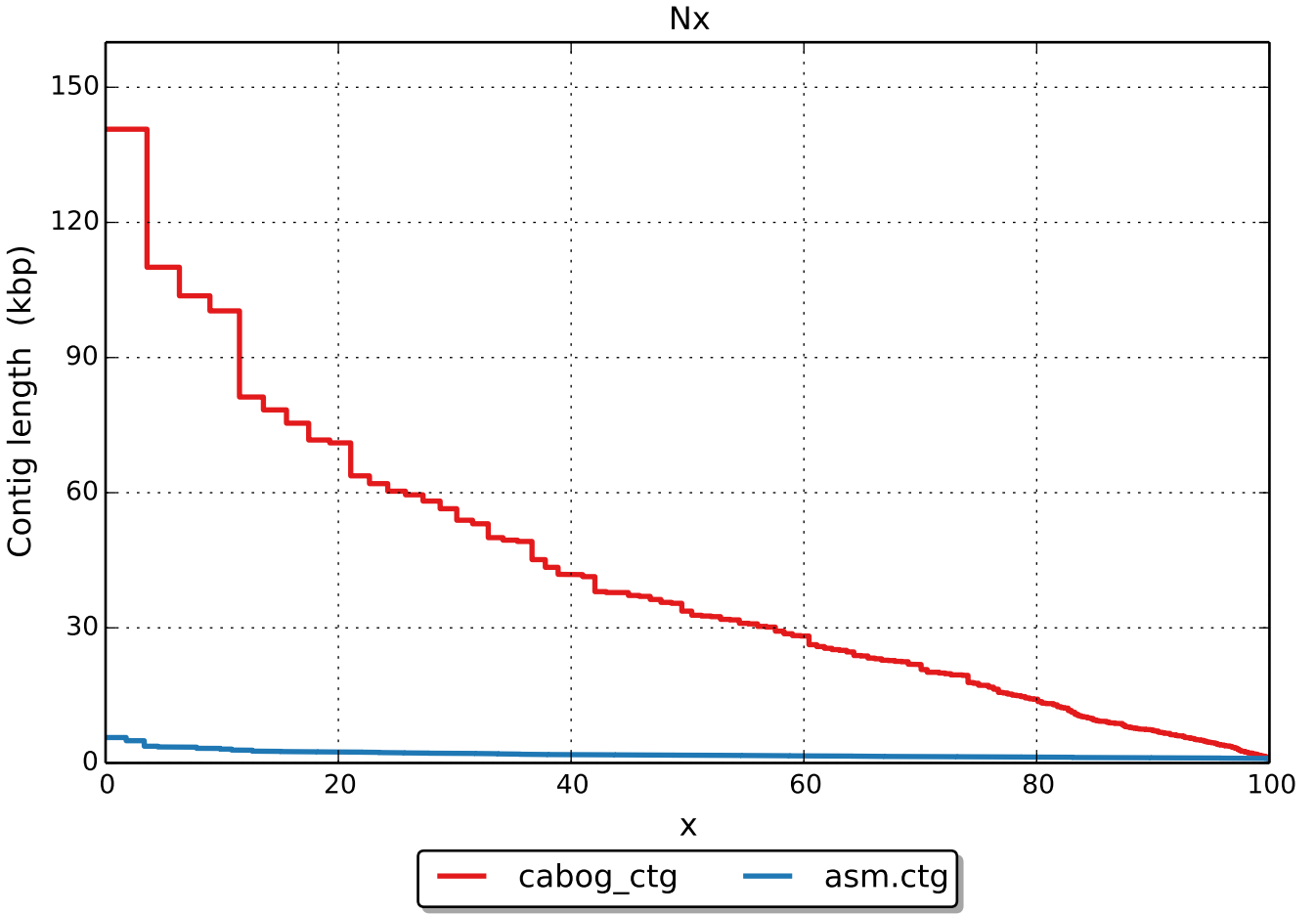
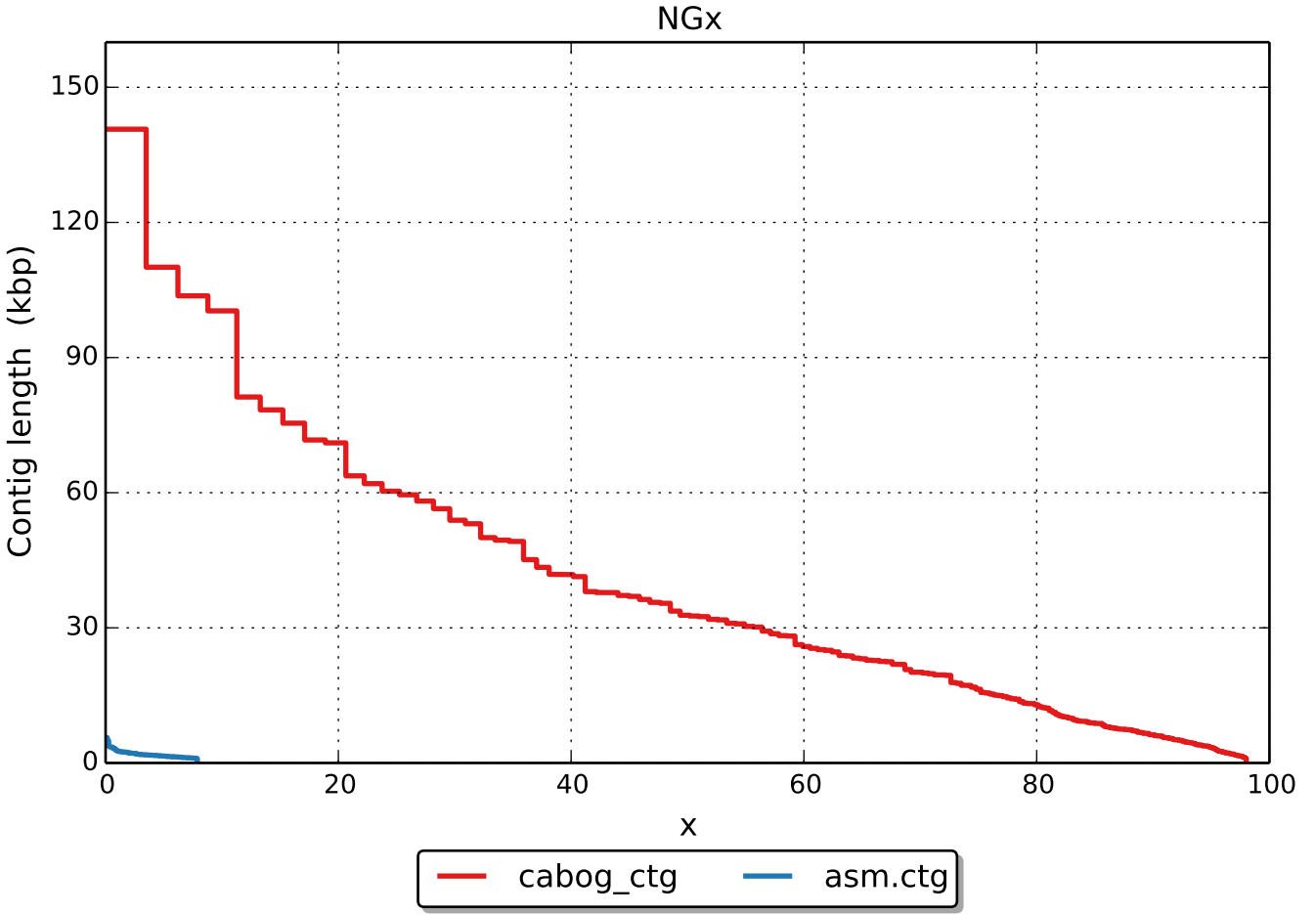
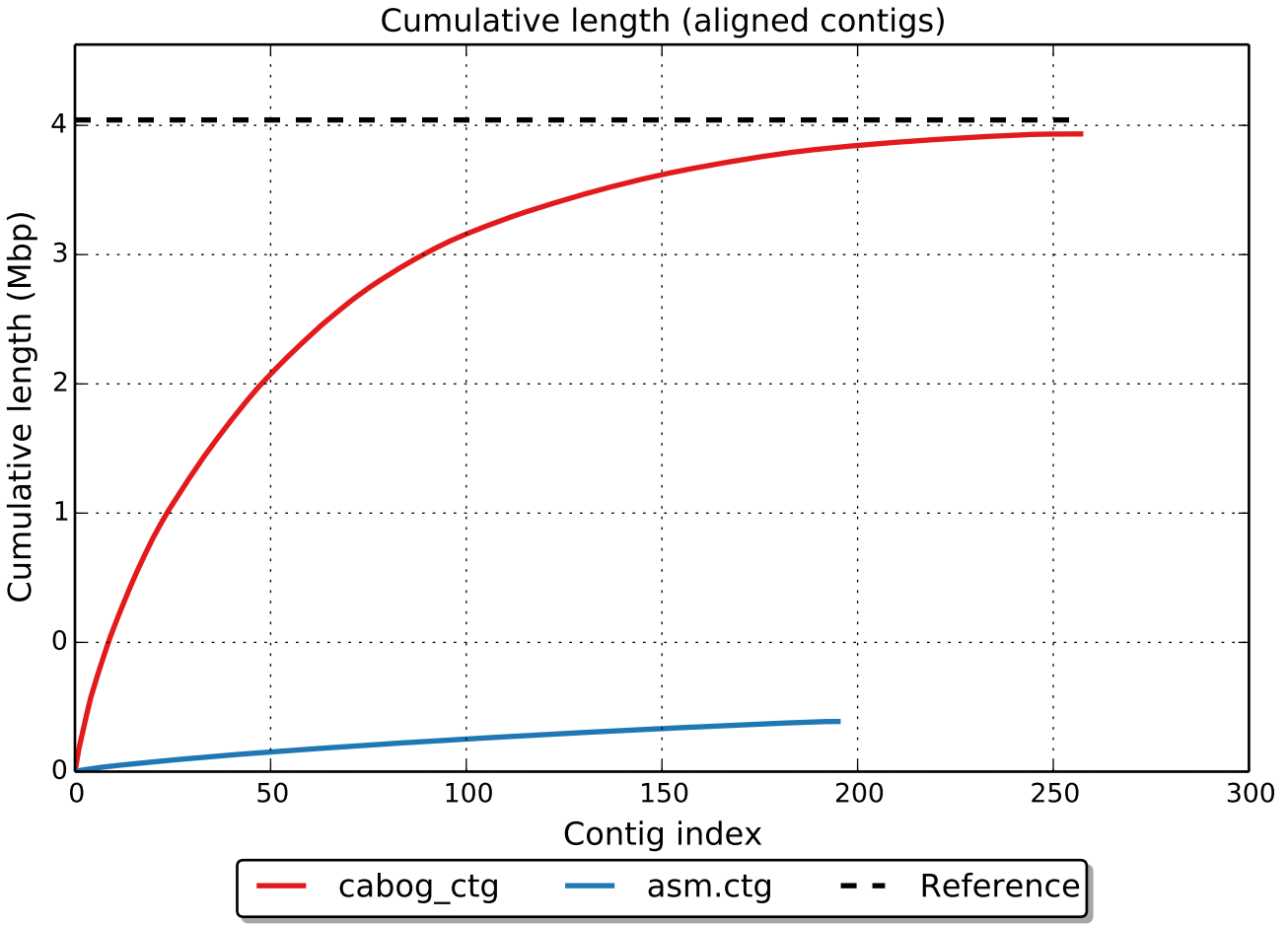
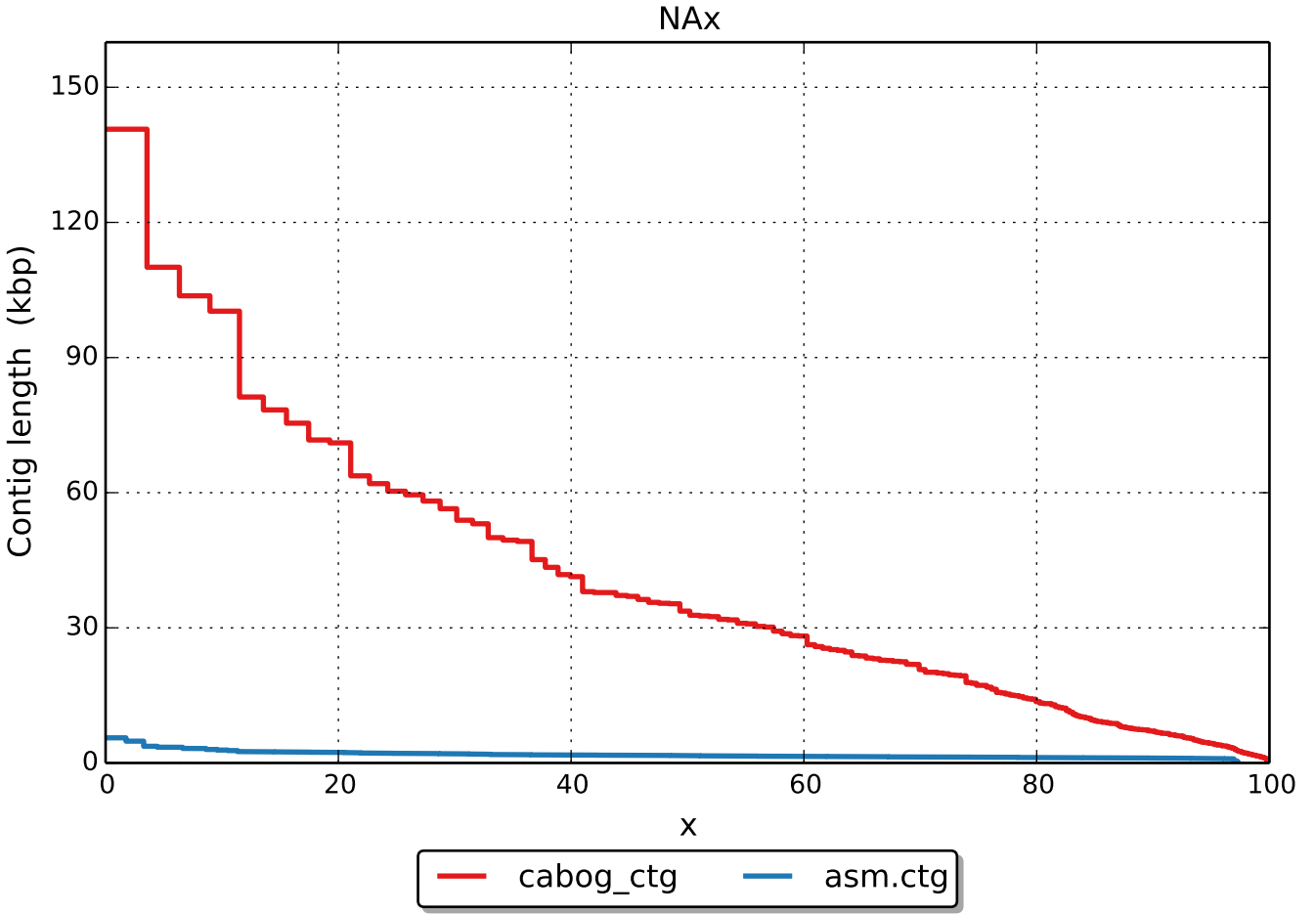

Cumulative length
Cumulative length (Mbp)
0
1
2
3
4
0
50
100
150
200
250
Contig index
cabog_ctg
asm.ctg
Reference

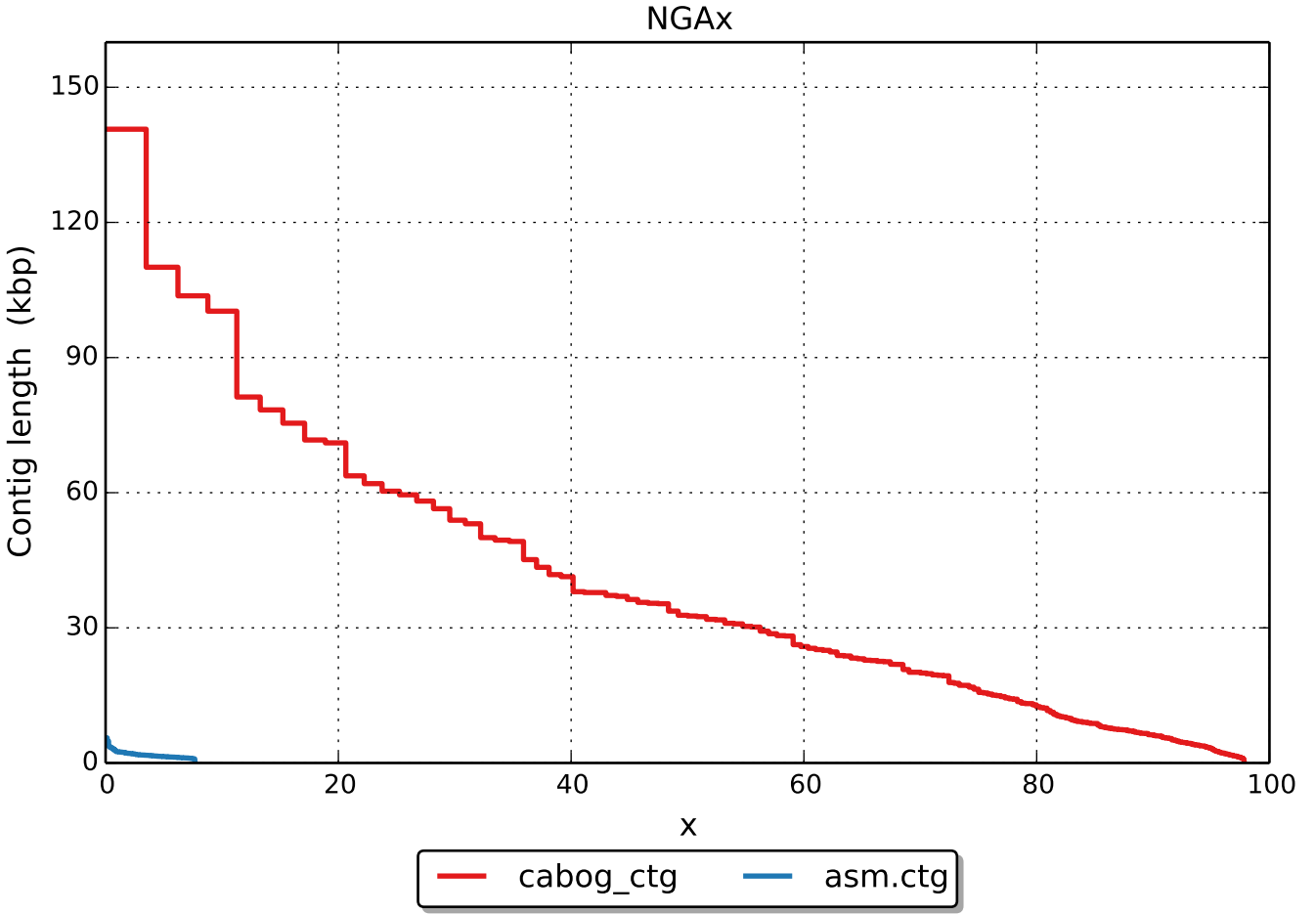

GC content
3000
2500
2000
1500
1000
500
0
windows
0
20
40
60
80
100
GC (%)
cabog_ctg
asm.ctg
Reference

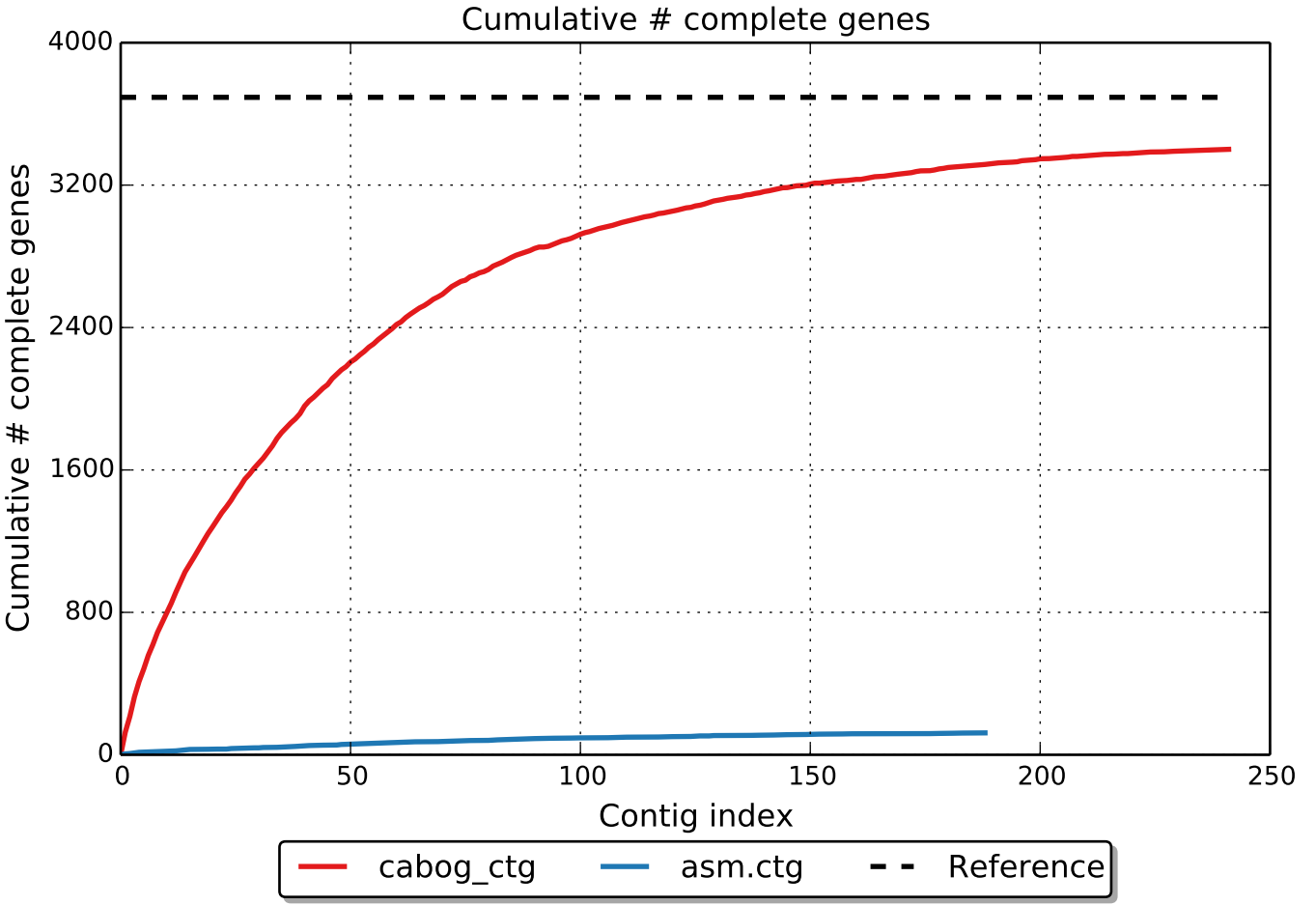

Nx
Contig length (kbp)
150
120
90
60
30
0
0
20
40
60
80
100
x
cabog_ctg
asm.ctg

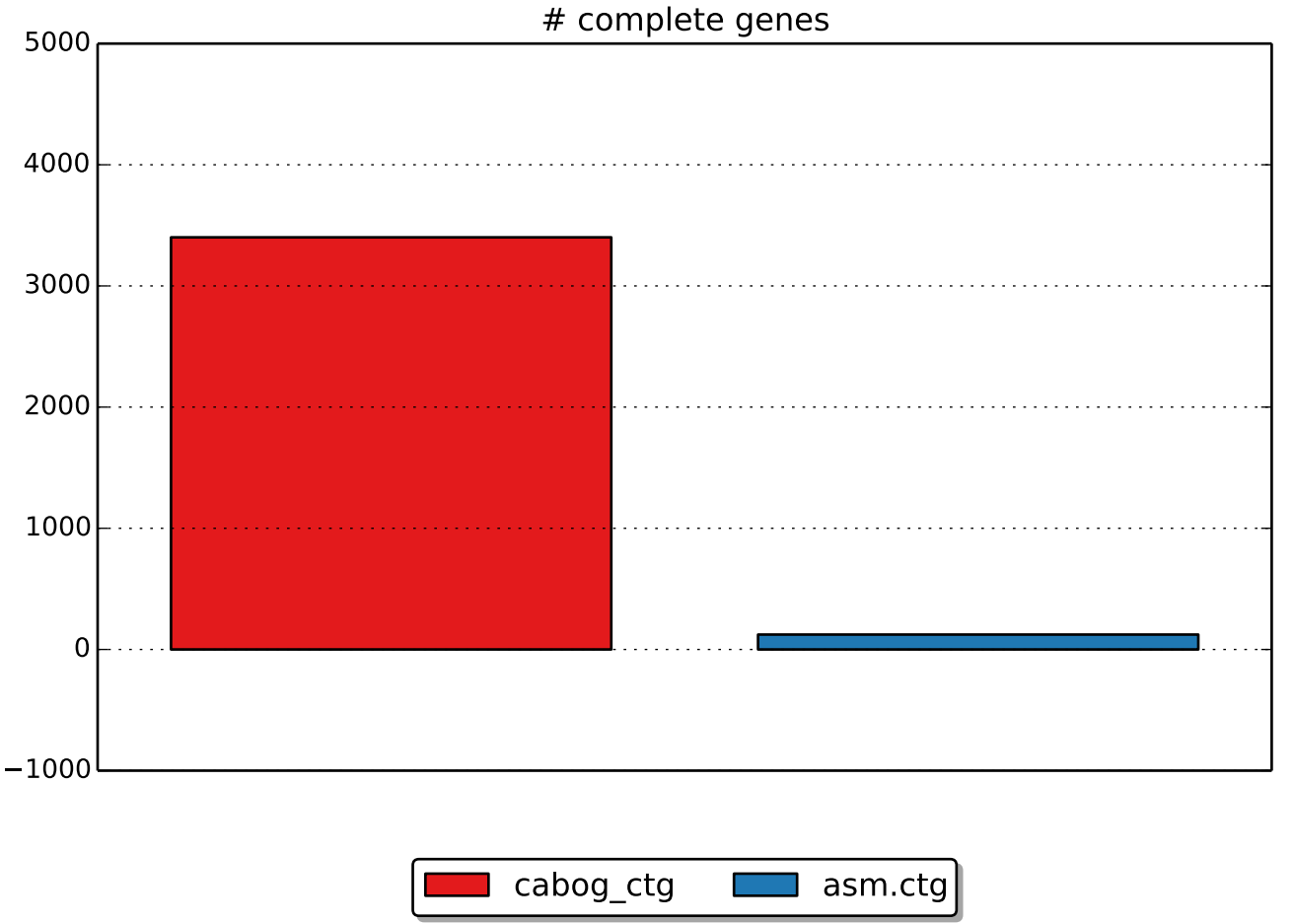

NGx
Contig length (kbp)
150
120
90
60
30
0
0
20
40
60
80
100
x
cabog_ctg
asm.ctg

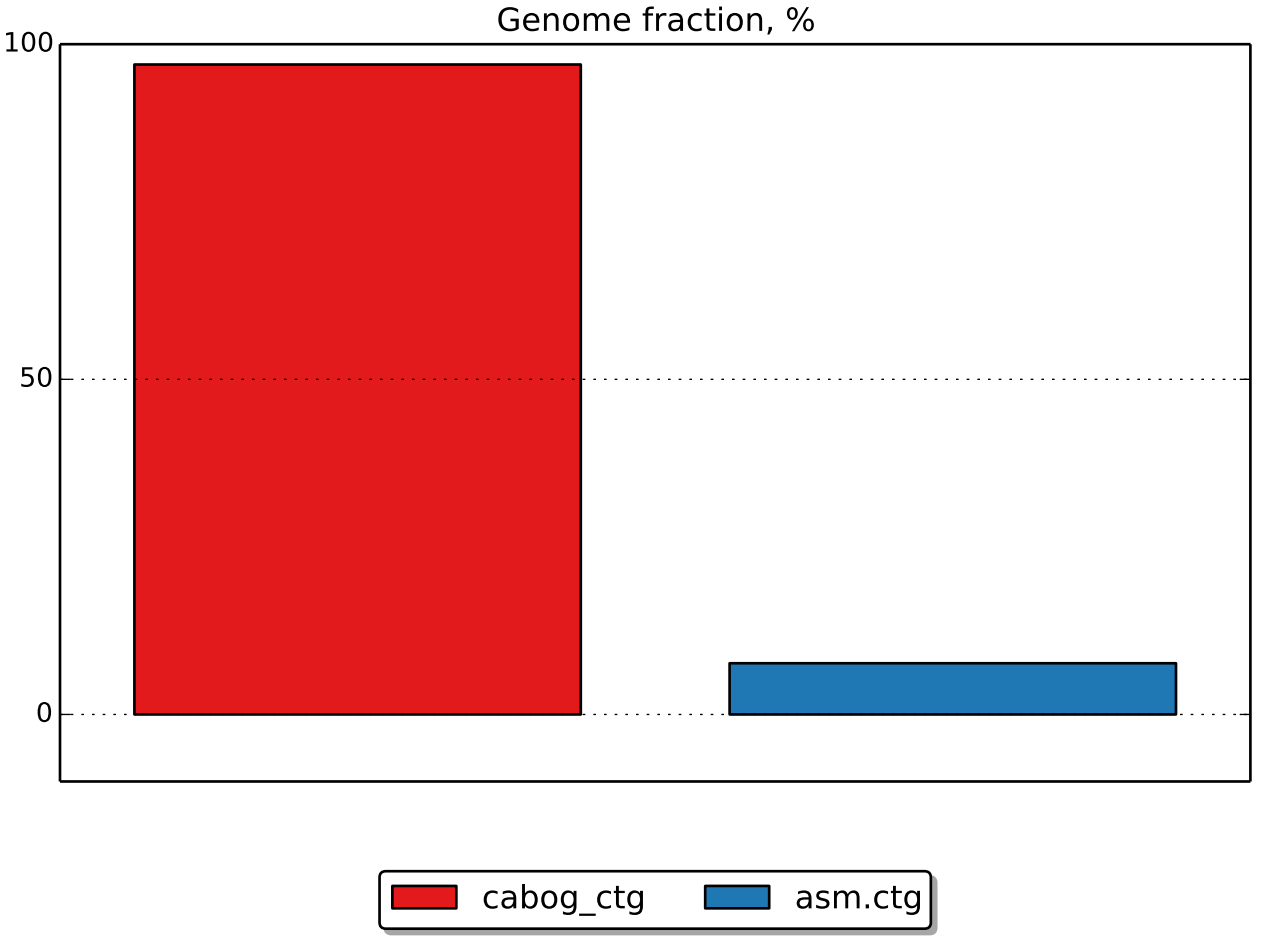

Cumulative length (aligned contigs)
Cumulative length (Mbp)
0
1
2
3
4
0
50
100
150
200
250
300
Contig index
cabog_ctg
asm.ctg
Reference


NAx
Contig length (kbp)
150
120
90
60
30
0
0
20
40
60
80
100
x
cabog_ctg
asm.ctg


NGAx
Contig length (kbp)
150
120
90
60
30
0
0
20
40
60
80
100
x
cabog_ctg
asm.ctg


Cumulative # complete genes
Cumulative # complete genes
4000
3200
2400
1600
800
0
0
50
100
150
200
250
Contig index
cabog_ctg
asm.ctg
Reference


complete genes
5000
4000
3000
2000
1000
0
−1000
cabog_ctg
asm.ctg


Genome fraction, %
100
50
0
cabog_ctg
asm.ctg