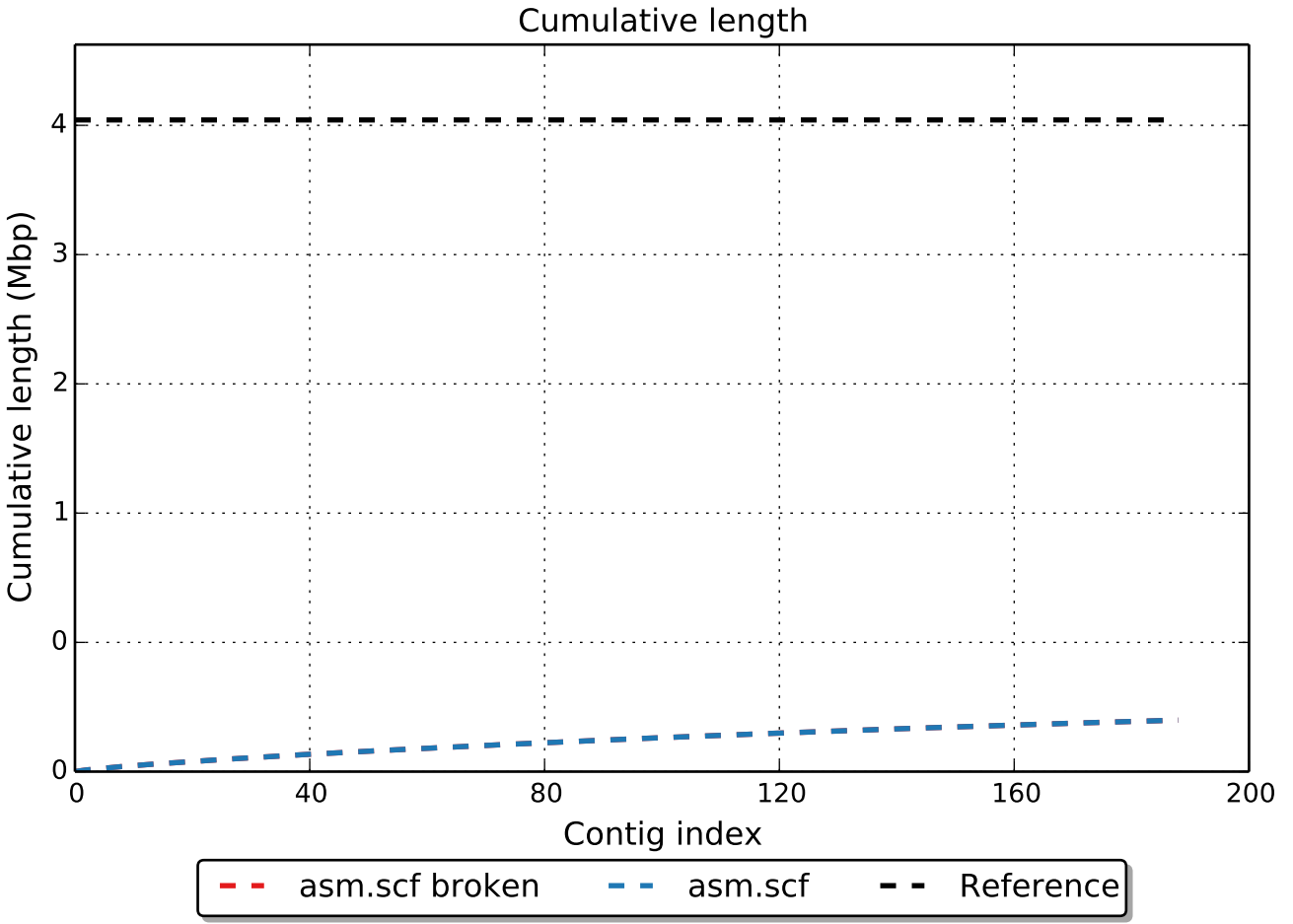
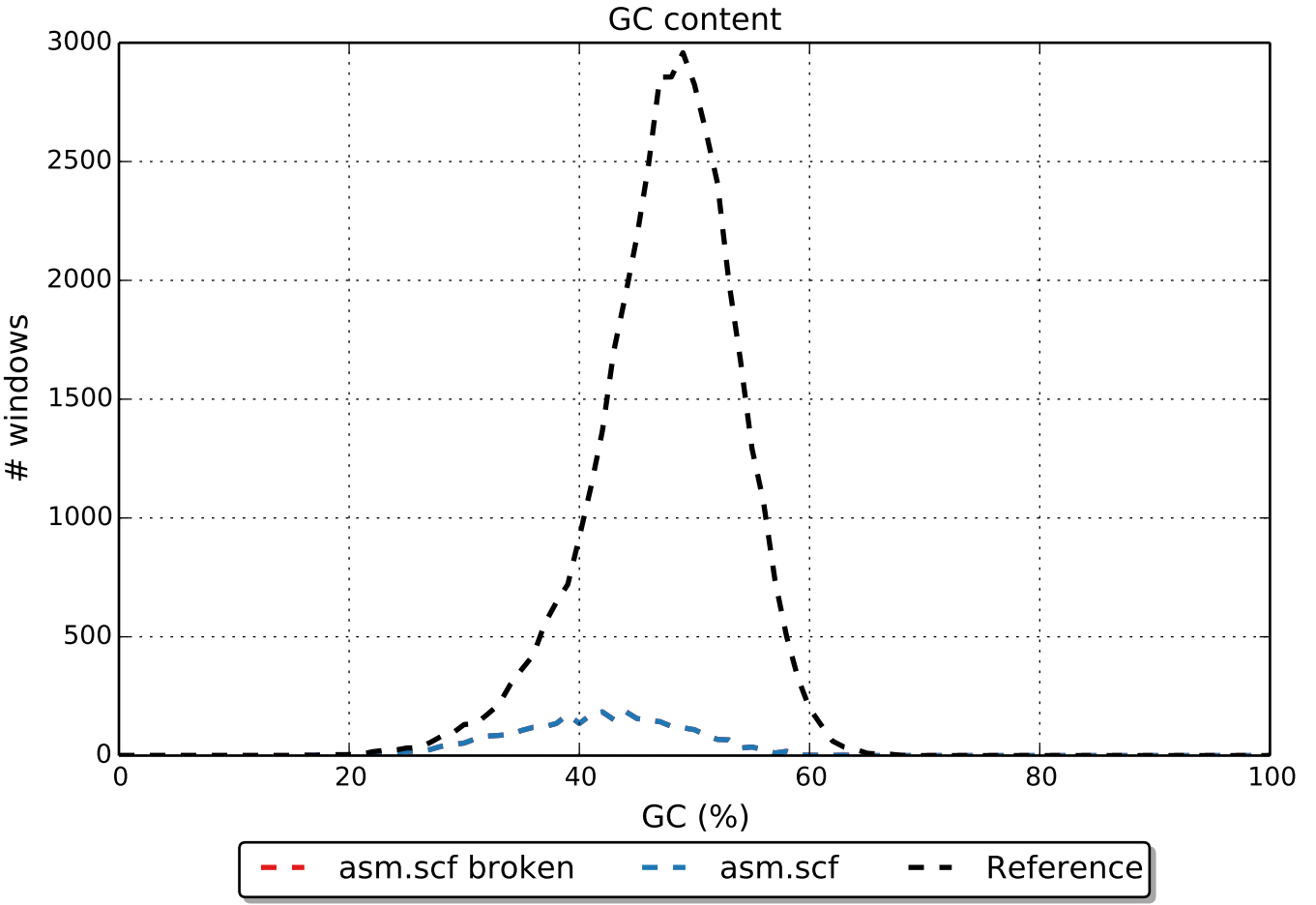
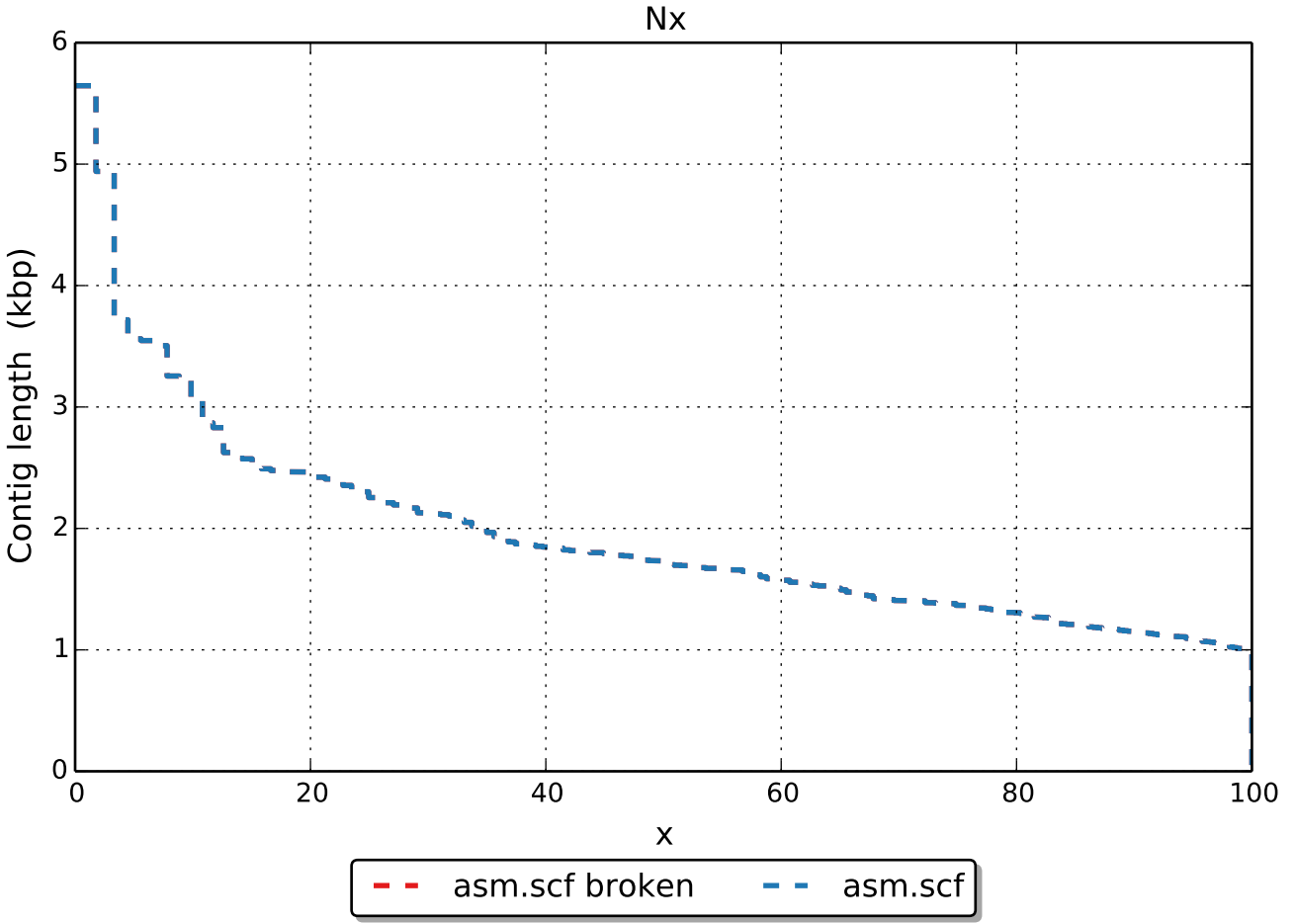
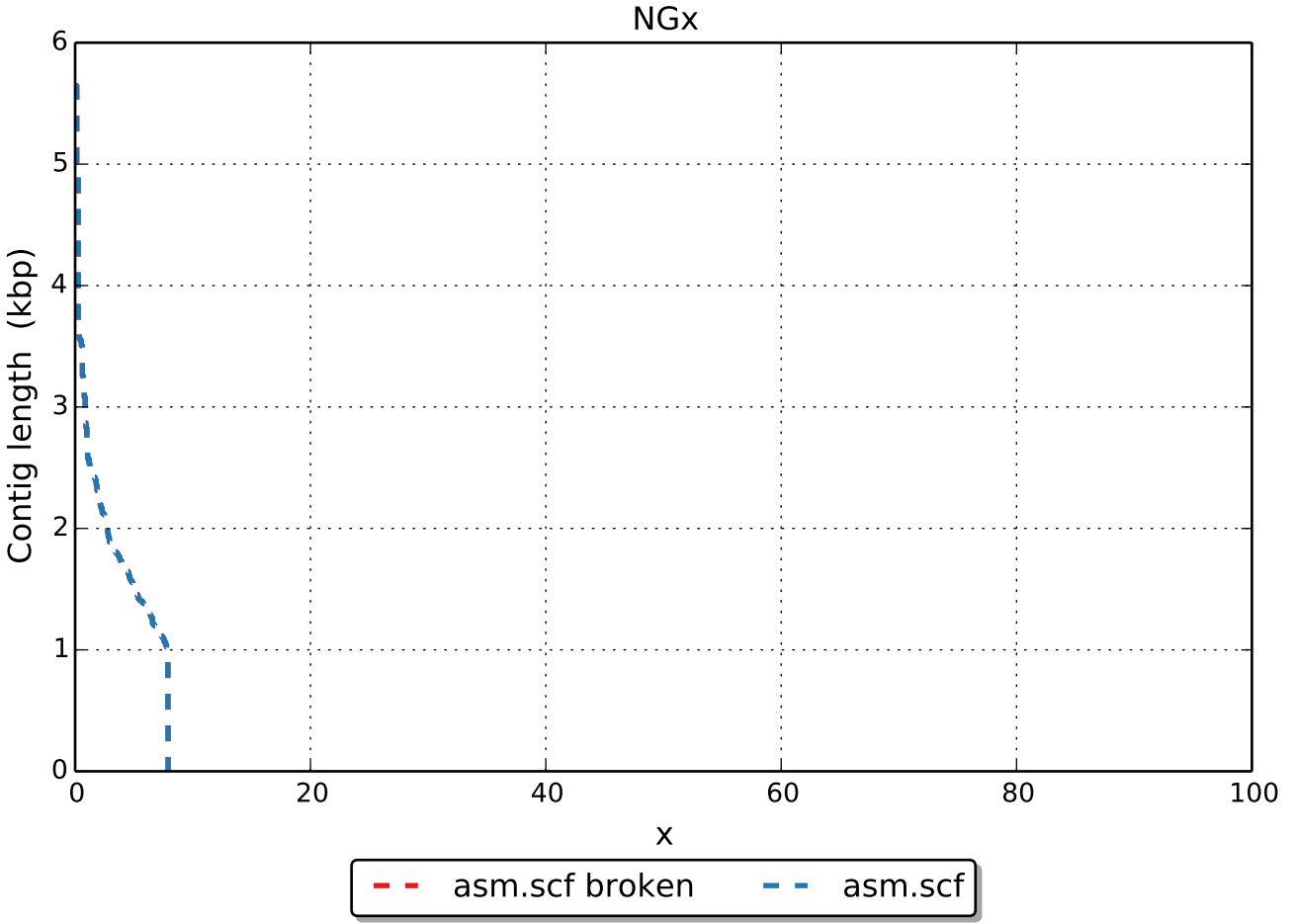
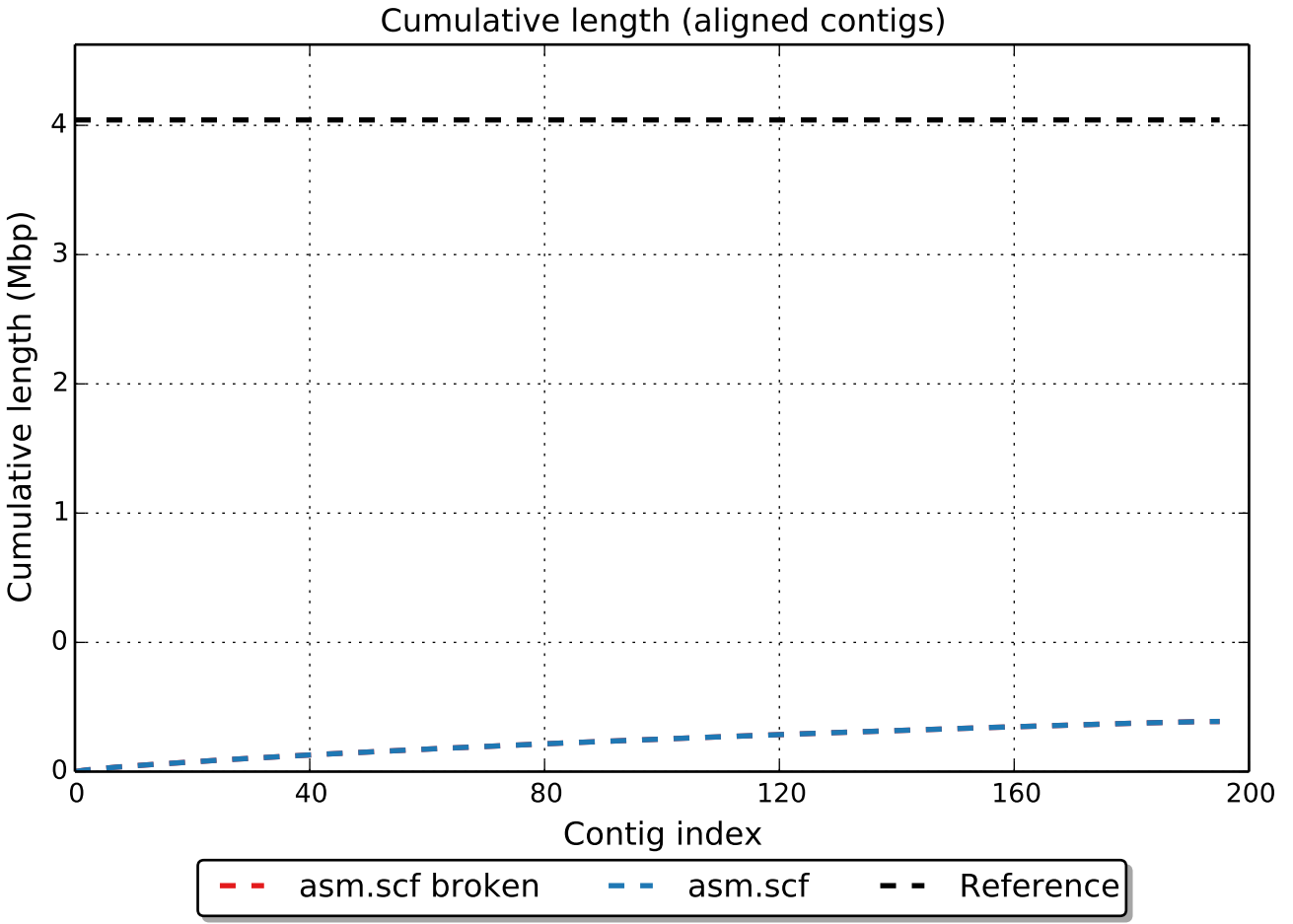
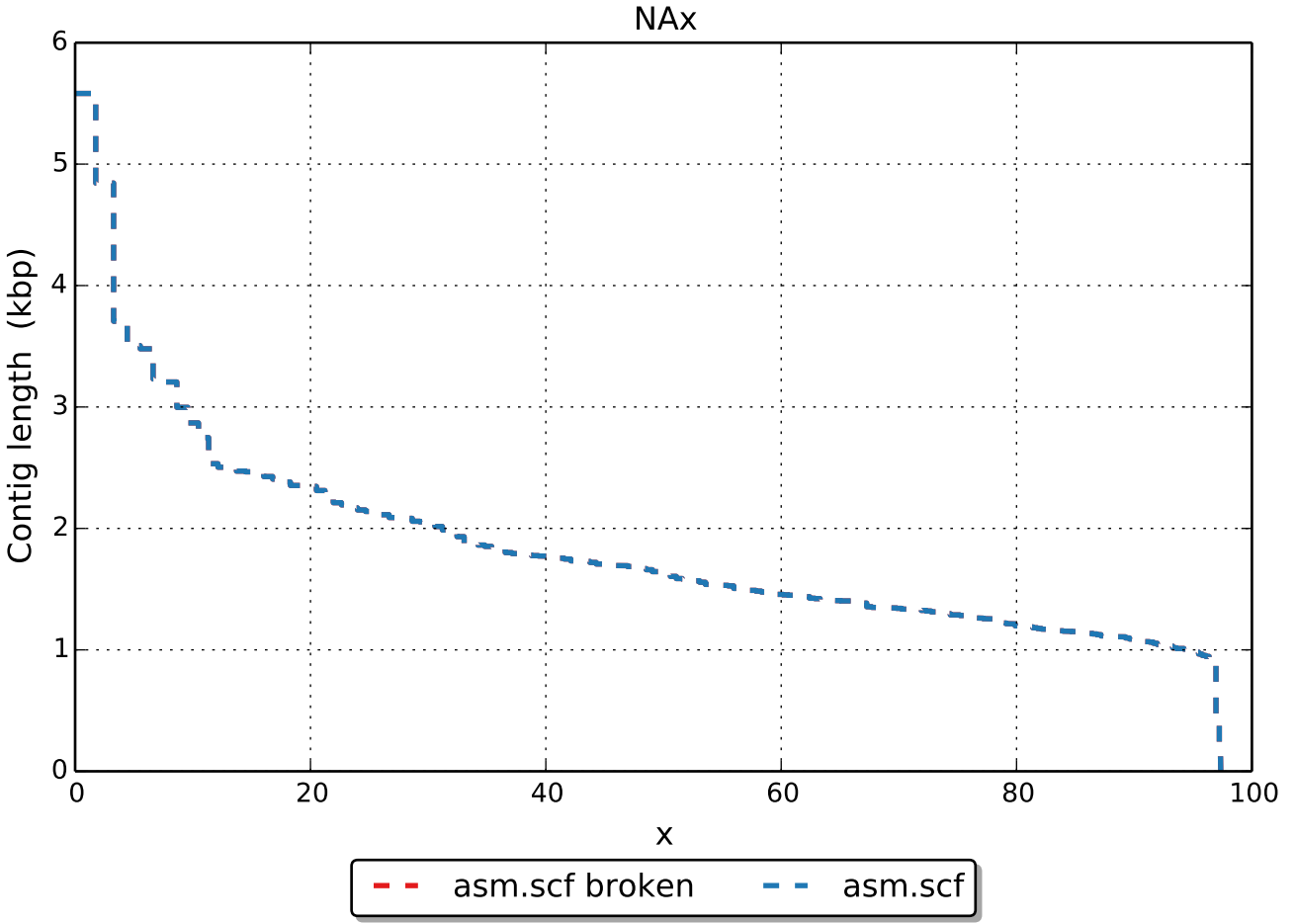

Cumulative length
Cumulative length (Mbp)
4
3
2
1
0
0
40
80
120
160
200
Contig index
asm.scf broken
asm.scf
Reference

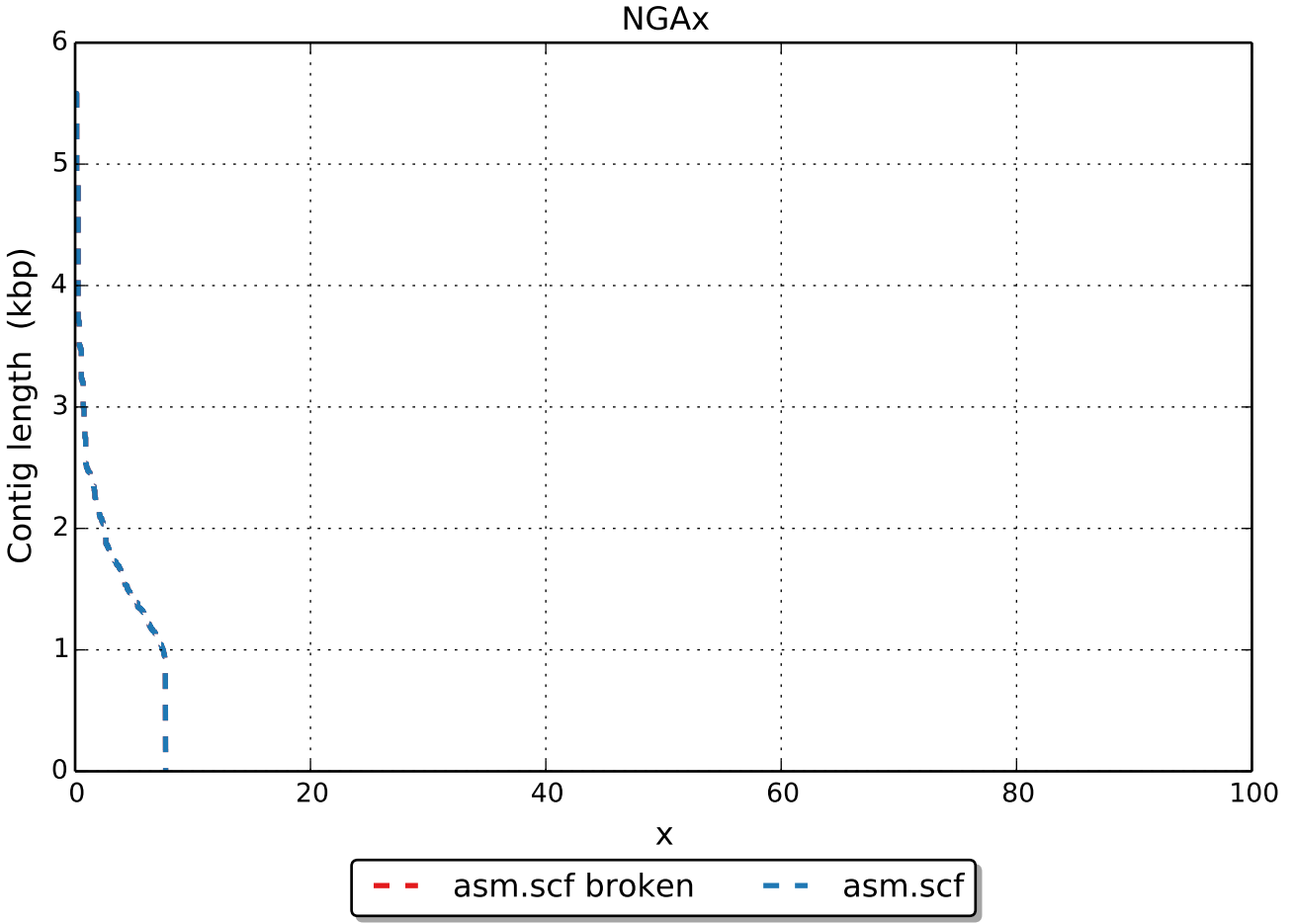

GC content
3000
2500
2000
1500
1000
500
0
windows
0
20
40
60
80
100
GC (%)
asm.scf broken
asm.scf
Reference

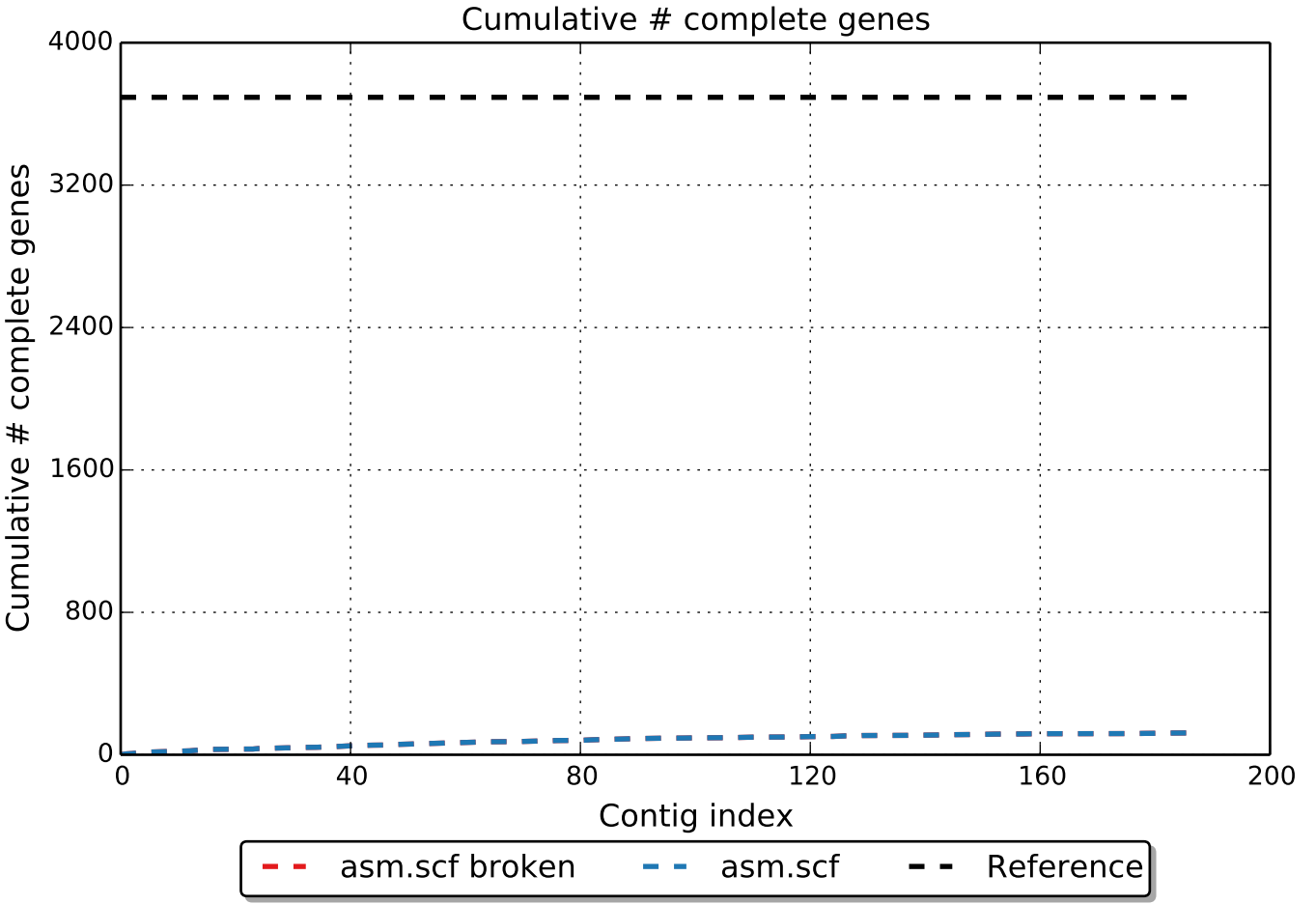

Nx
Contig length (kbp)
0
1
2
3
4
5
6
0
20
40
60
80
100
x
asm.scf broken
asm.scf

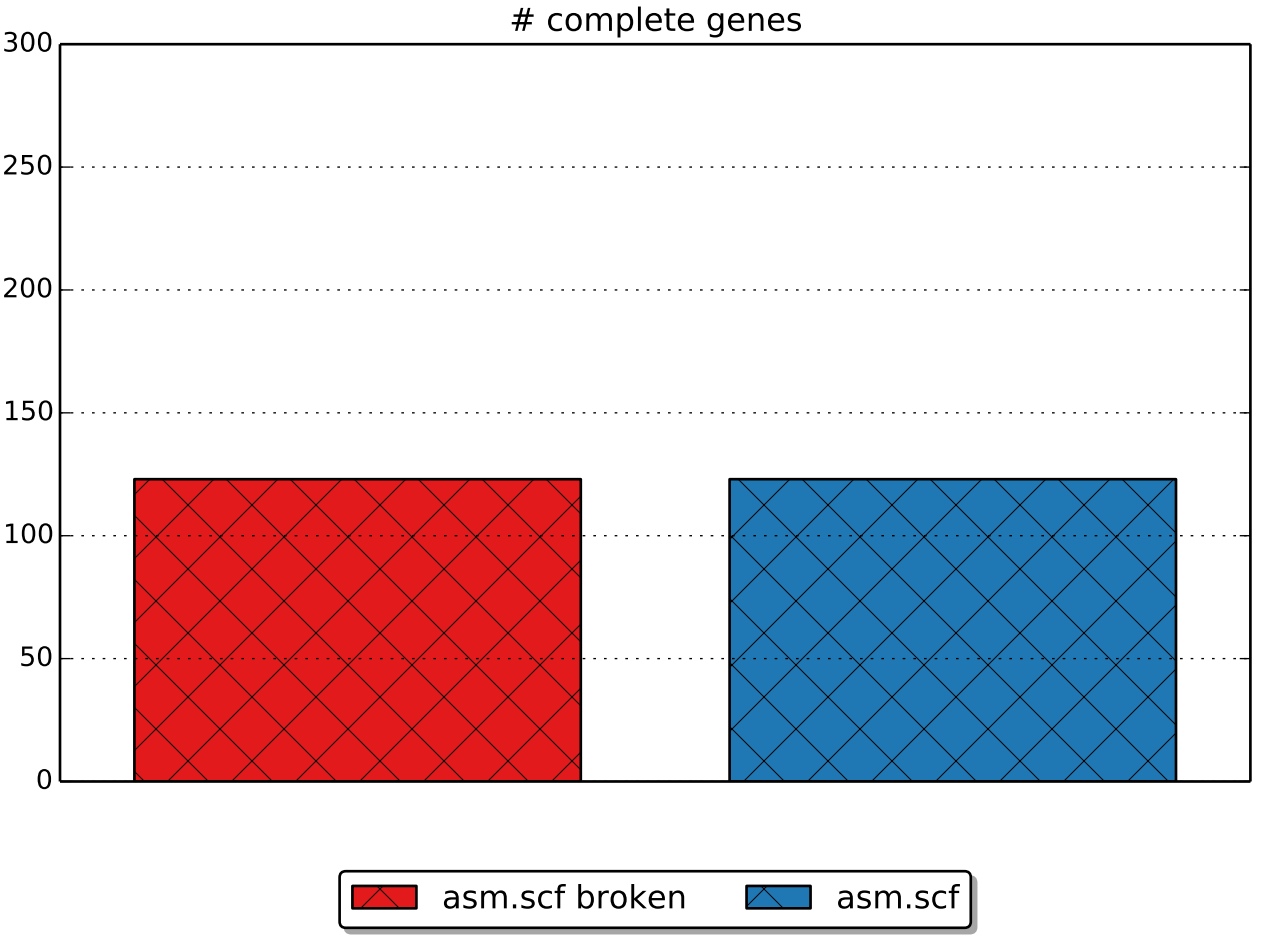

NGx
Contig length (kbp)
0
1
2
3
4
5
6
0
20
40
60
80
100
x
asm.scf broken
asm.scf

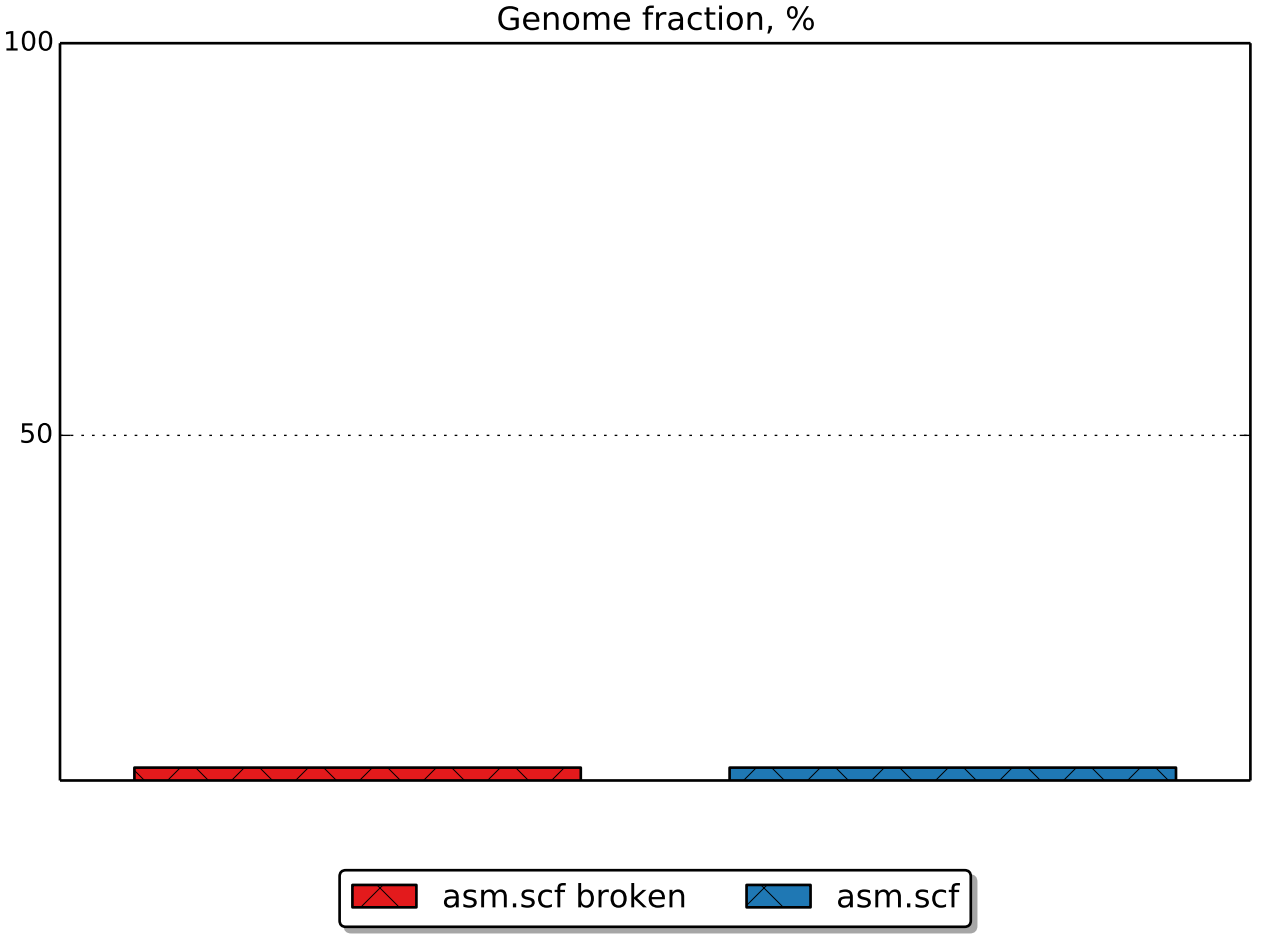

Cumulative length (aligned contigs)
Cumulative length (Mbp)
0
1
2
3
4
0
40
80
120
160
200
Contig index
asm.scf broken
asm.scf
Reference


NAx
Contig length (kbp)
0
1
2
3
4
5
6
0
20
40
60
80
100
x
asm.scf broken
asm.scf


NGAx
Contig length (kbp)
0
1
2
3
4
5
6
0
20
40
60
80
100
x
asm.scf broken
asm.scf


Cumulative # complete genes
Cumulative # complete genes
4000
3200
2400
1600
800
0
0
40
80
120
160
200
Contig index
asm.scf broken
asm.scf
Reference


complete genes
300
250
200
150
100
50
0
asm.scf broken
asm.scf


Genome fraction, %
100
50
asm.scf broken
asm.scf