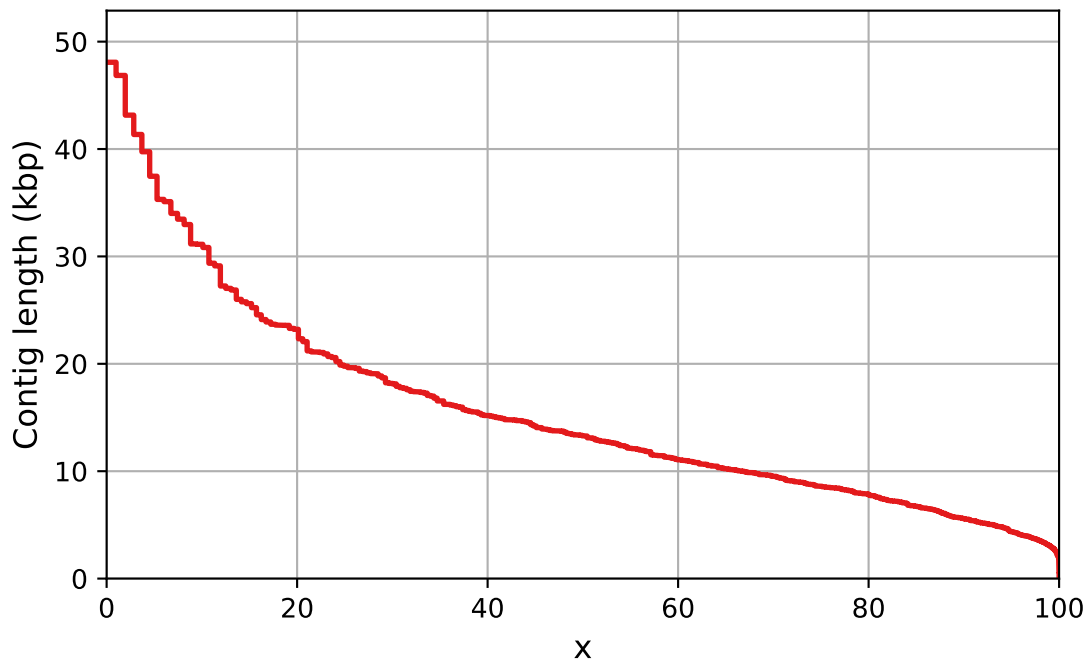


Report

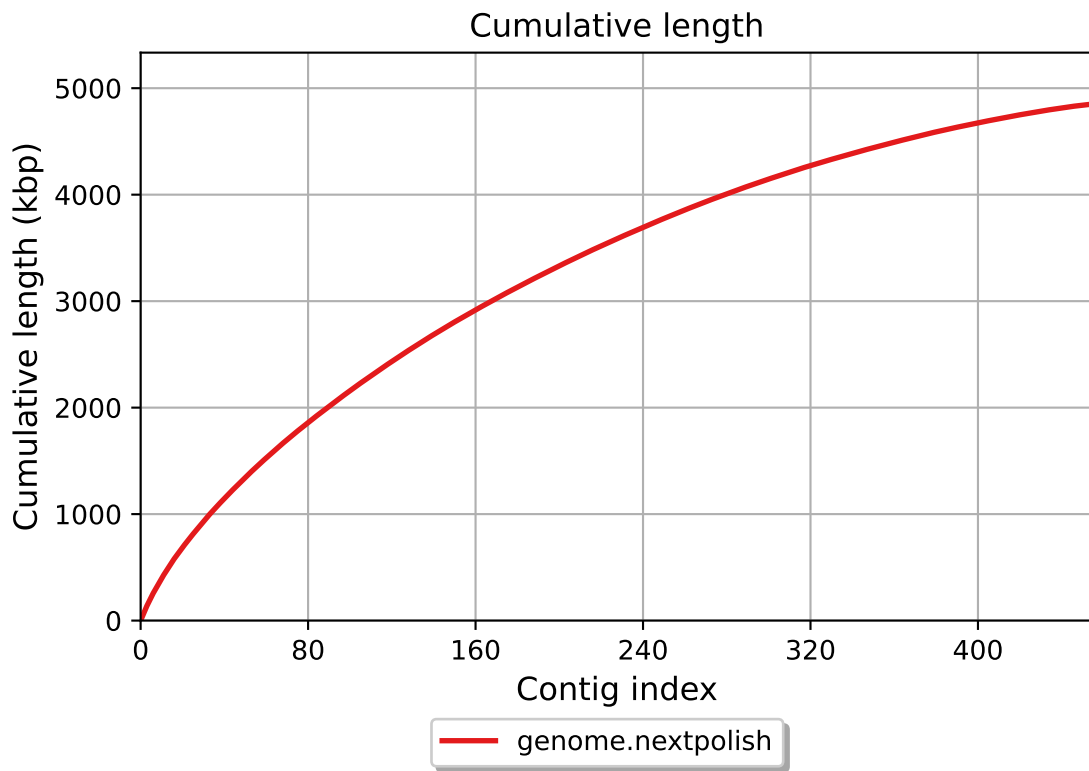
	genome.nextpolish
# contigs (>= 0 bp)	455
# contigs (>= 1000 bp)	454
# contigs (>= 5000 bp)	364
# contigs (>= 10000 bp)	190
# contigs (>= 25000 bp)	23
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	4849000
Total length (>= 1000 bp)	4848393
Total length (>= 5000 bp)	4512888
Total length (>= 10000 bp)	3233040
Total length (>= 25000 bp)	762860
Total length (>= 50000 bp)	0
# contigs	455
Largest contig	48079
Total length	4849000
GC (%)	43.28
N50	13298
N90	5589
auN	15750.2
L50	120
L90	336
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	15 + 3 part

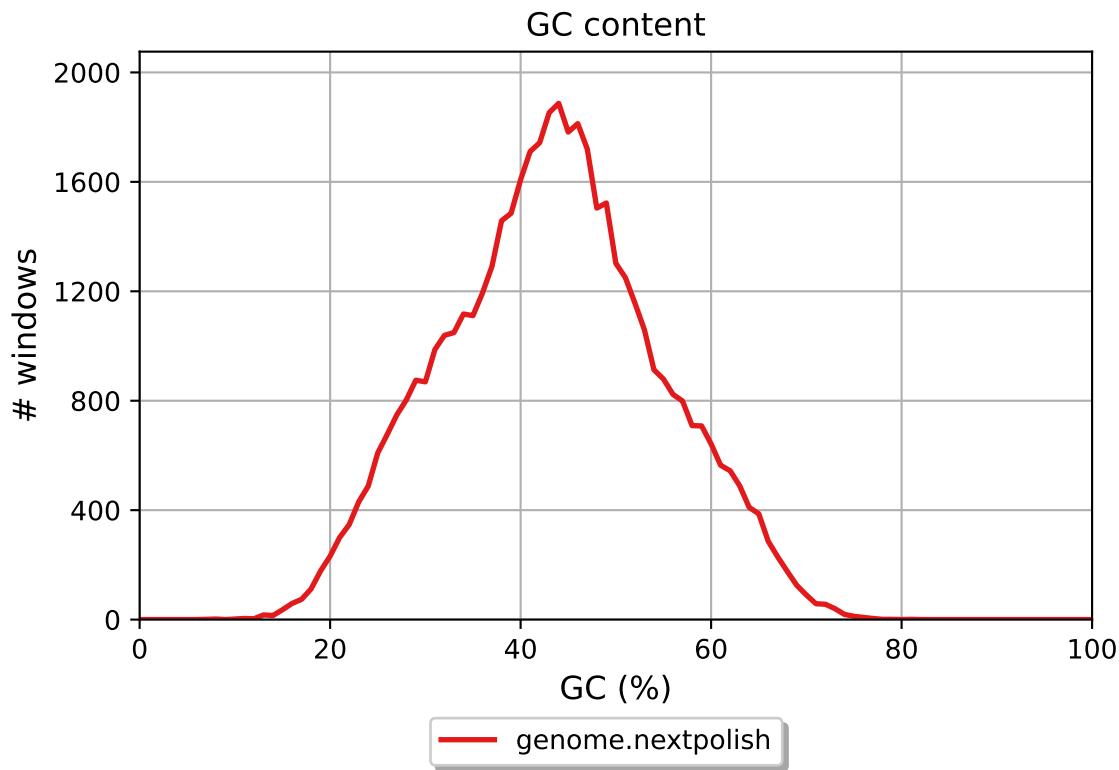
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

