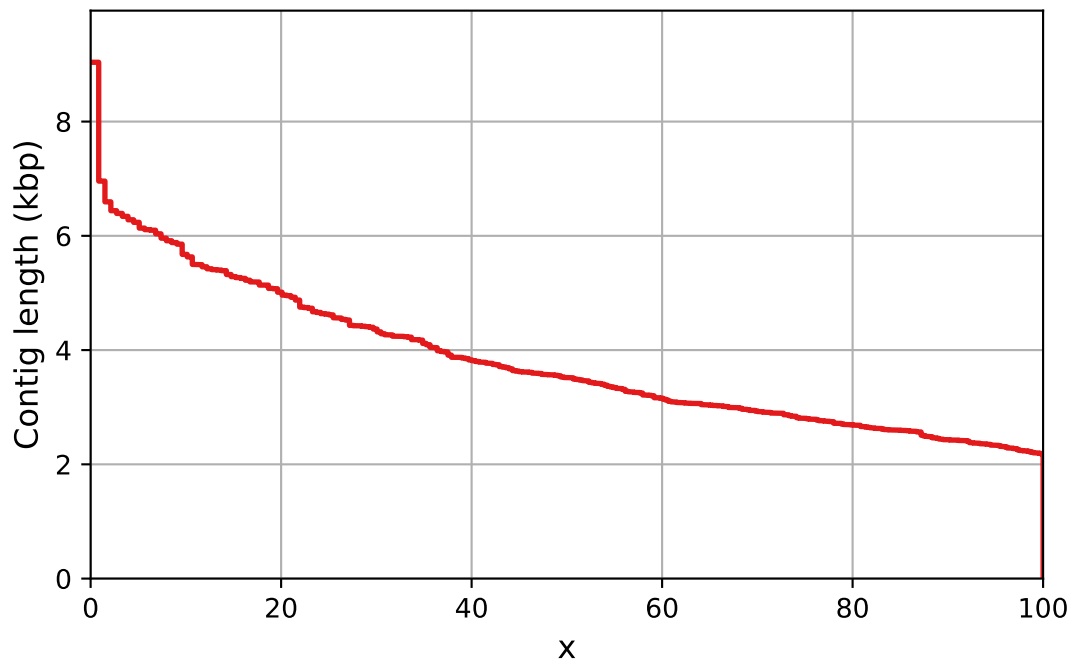


# Report

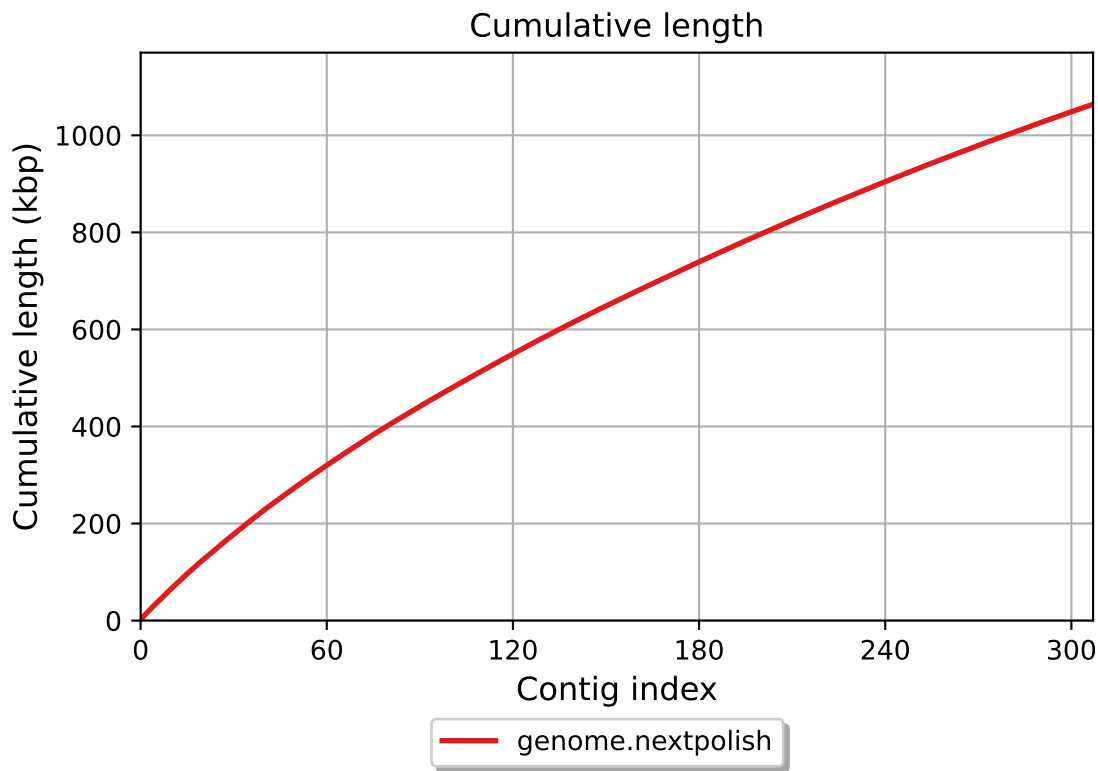
	genome.nextpolish
# contigs (>= 0 bp)	307
# contigs (>= 1000 bp)	307
# contigs (>= 5000 bp)	37
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1064022
Total length (>= 1000 bp)	1064022
Total length (>= 5000 bp)	213830
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	307
Largest contig	9038
Total length	1064022
GC (%)	40.40
N50	3520
N90	2431
auN	3822.1
L50	115
L90	262
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 2 part

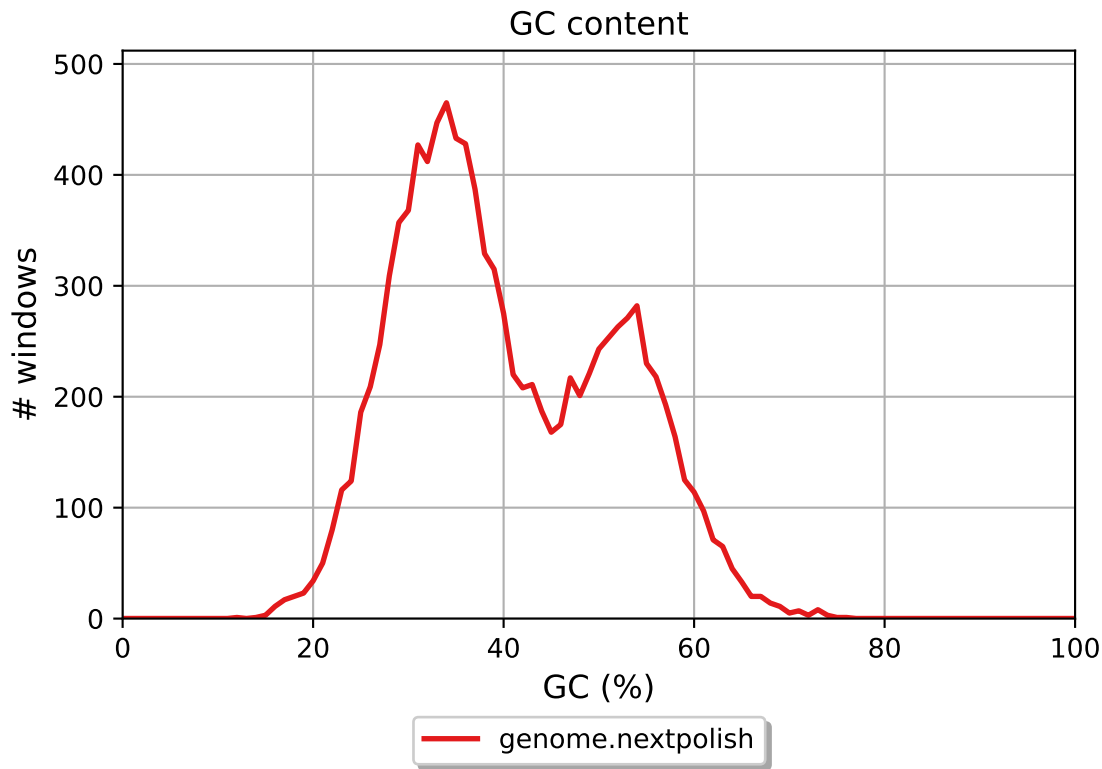
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

