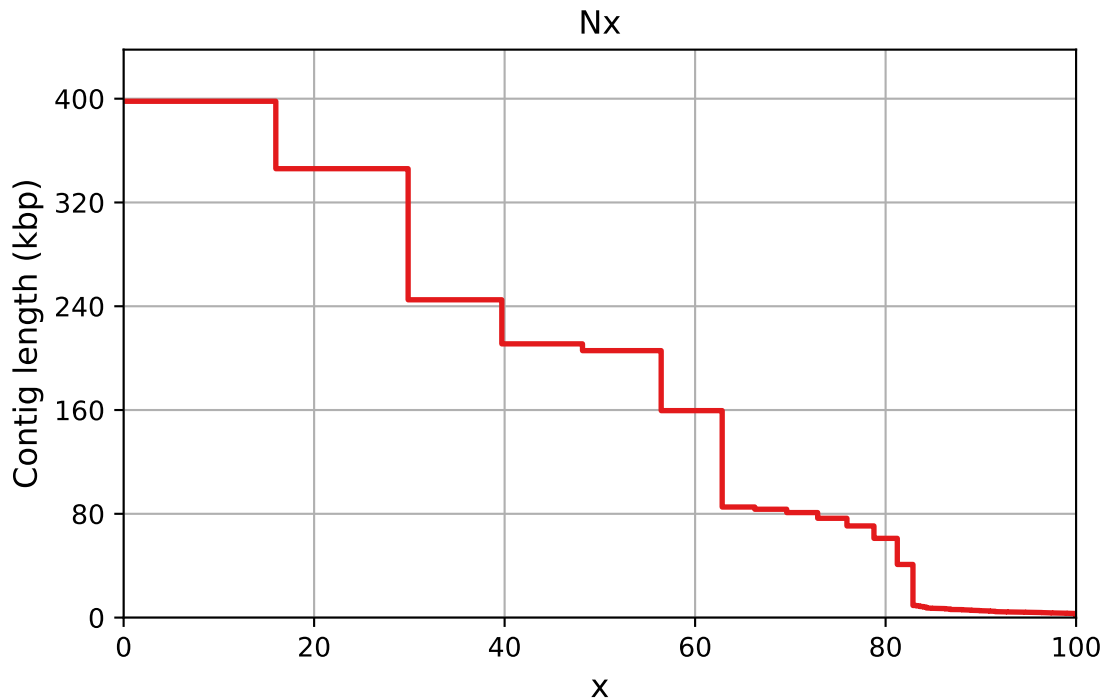


Report

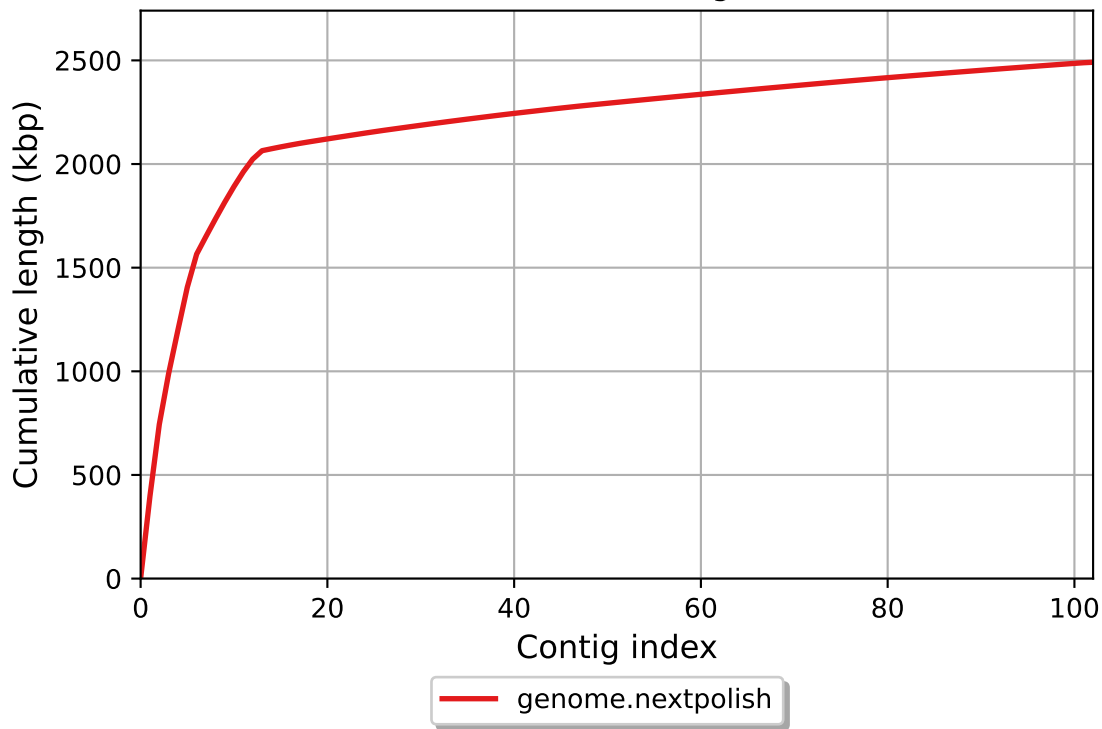
	genome.nextpolish
# contigs (>= 0 bp)	102
# contigs (>= 1000 bp)	102
# contigs (>= 5000 bp)	44
# contigs (>= 10000 bp)	13
# contigs (>= 25000 bp)	13
# contigs (>= 50000 bp)	12
Total length (>= 0 bp)	2490931
Total length (>= 1000 bp)	2490931
Total length (>= 5000 bp)	2264899
Total length (>= 10000 bp)	2064296
Total length (>= 25000 bp)	2064296
Total length (>= 50000 bp)	2023342
# contigs	102
Largest contig	398012
Total length	2490931
GC (%)	42.56
N50	205781
N90	5305
auN	196625.3
L50	5
L90	40
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	10 + 1 part

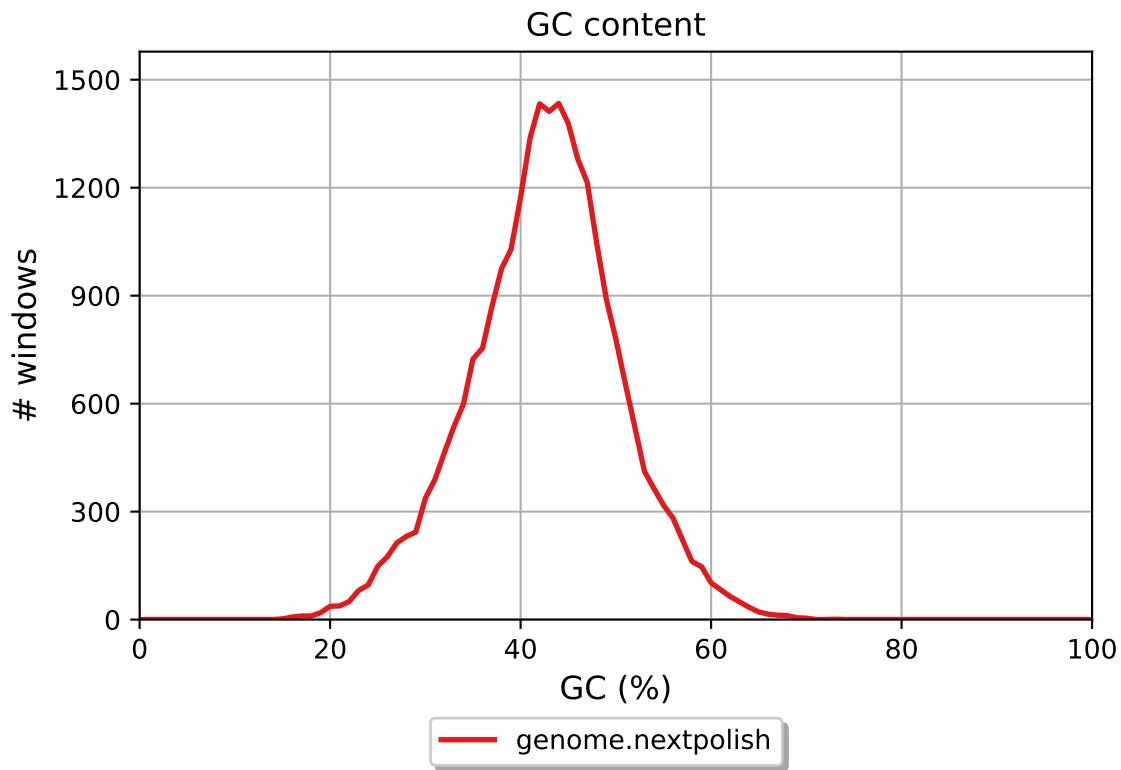
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

