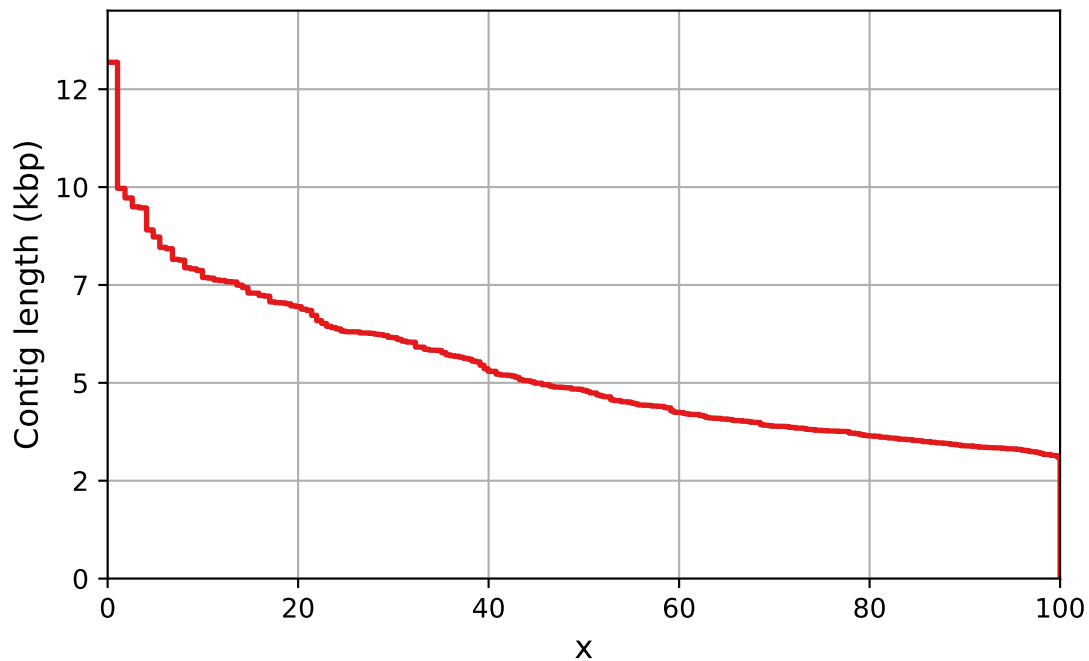


Report

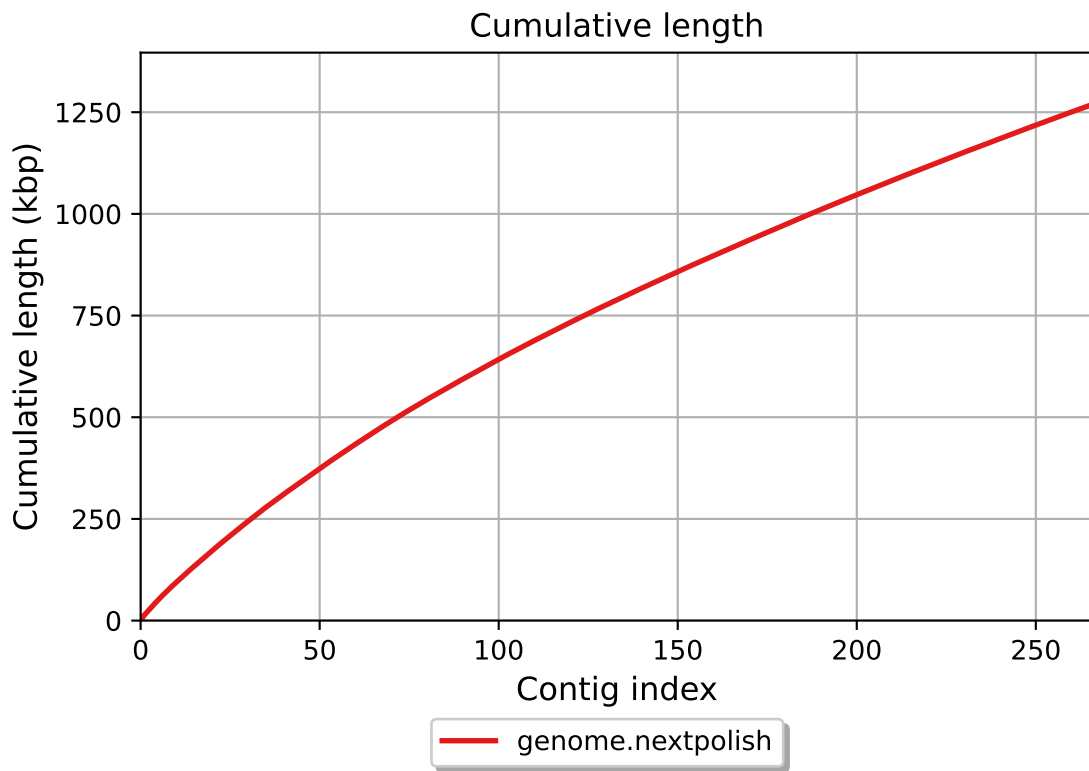
	genome.nextpolish
# contigs (>= 0 bp)	266
# contigs (>= 1000 bp)	266
# contigs (>= 5000 bp)	85
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1269513
Total length (>= 1000 bp)	1269513
Total length (>= 5000 bp)	569018
Total length (>= 10000 bp)	13188
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	266
Largest contig	13188
Total length	1269513
GC (%)	44.07
N50	4808
N90	3392
auN	5298.2
L50	99
L90	228
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	5 + 0 part

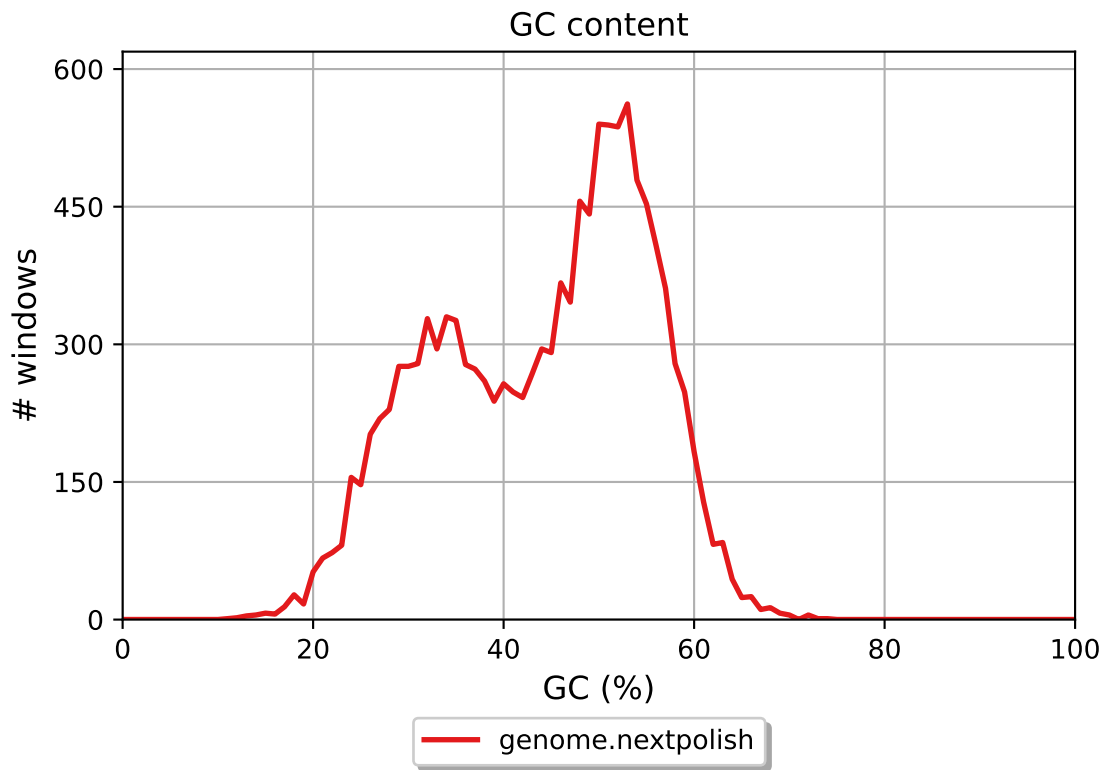
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

