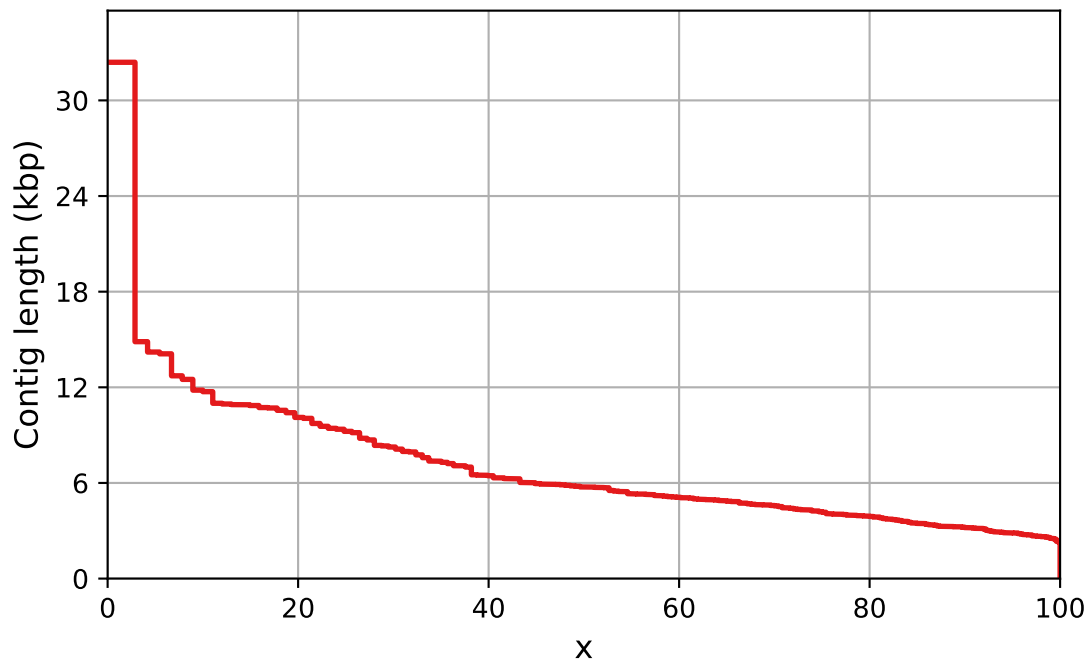


Report

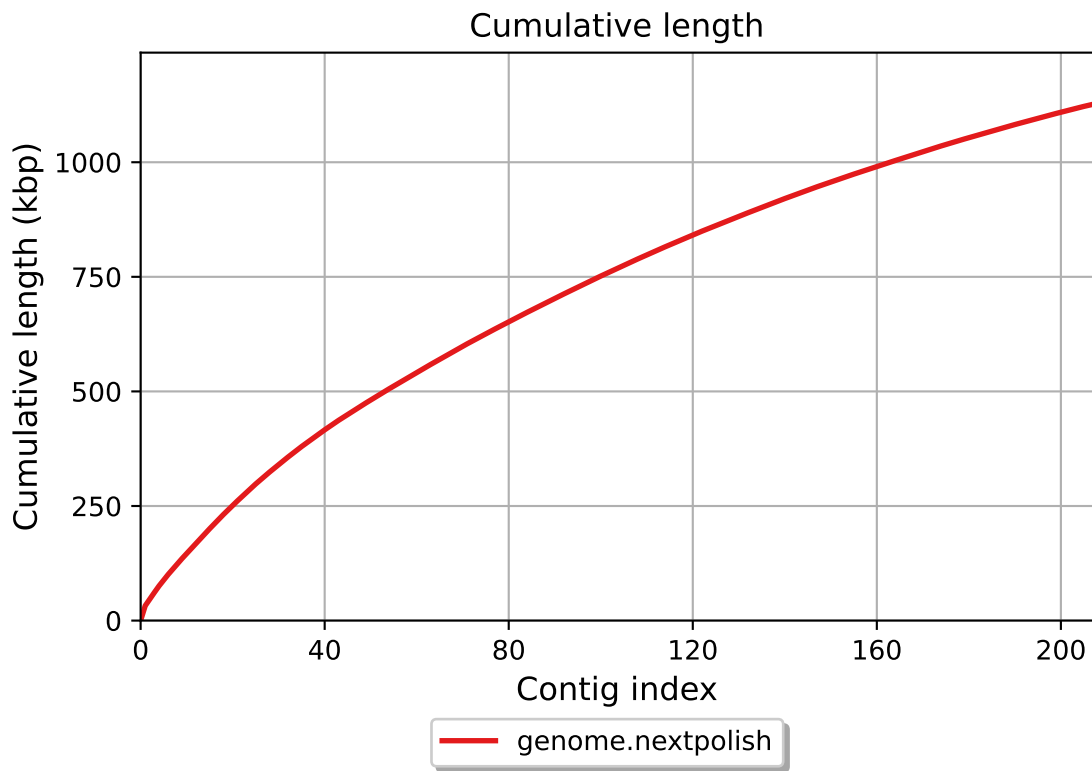
	genome.nextpolish
# contigs (>= 0 bp)	207
# contigs (>= 1000 bp)	207
# contigs (>= 5000 bp)	88
# contigs (>= 10000 bp)	19
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1126476
Total length (>= 1000 bp)	1126476
Total length (>= 5000 bp)	692827
Total length (>= 10000 bp)	241518
Total length (>= 25000 bp)	32391
Total length (>= 50000 bp)	0
# contigs	207
Largest contig	32391
Total length	1126476
GC (%)	37.43
N50	5750
N90	3201
auN	7311.8
L50	64
L90	168
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 0 part

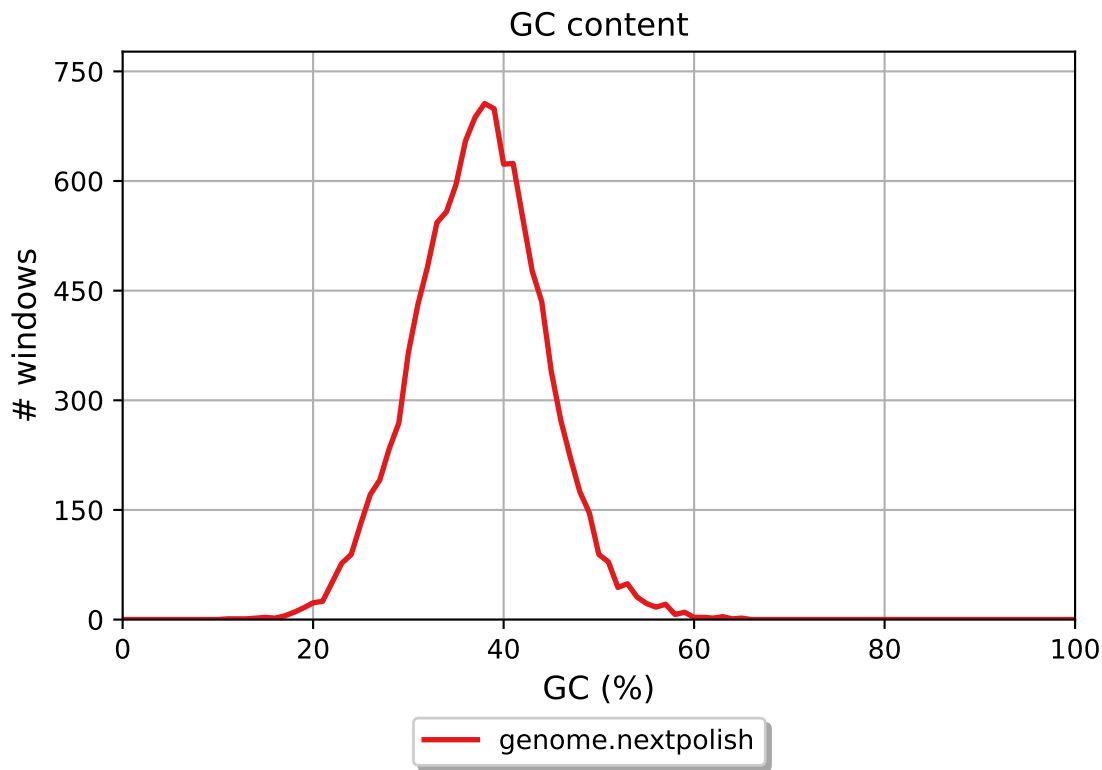
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

