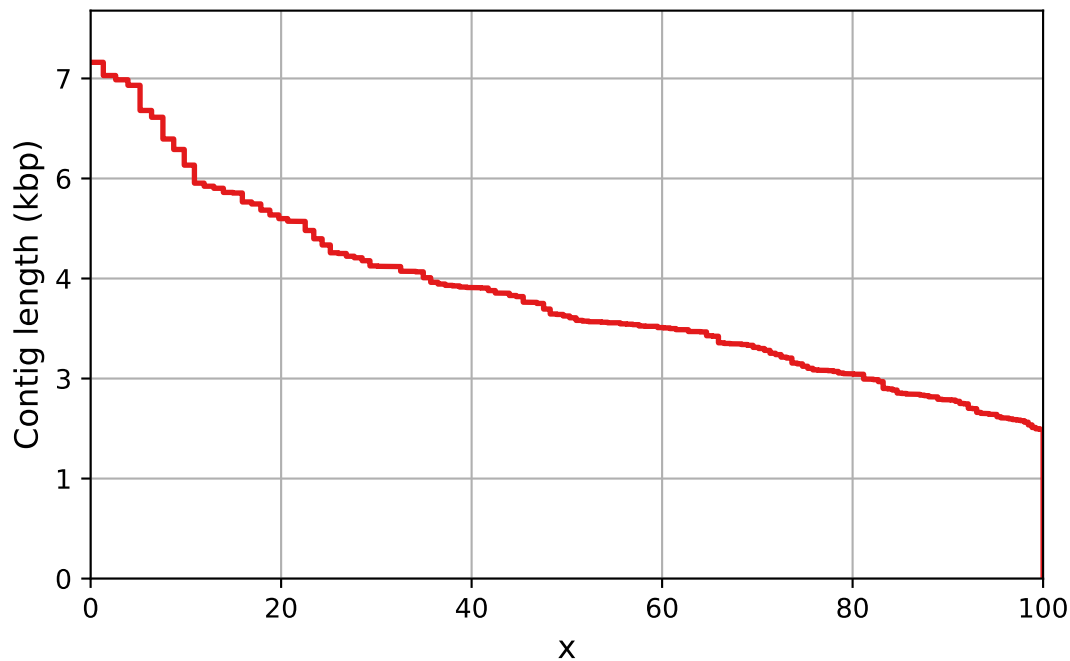


Report

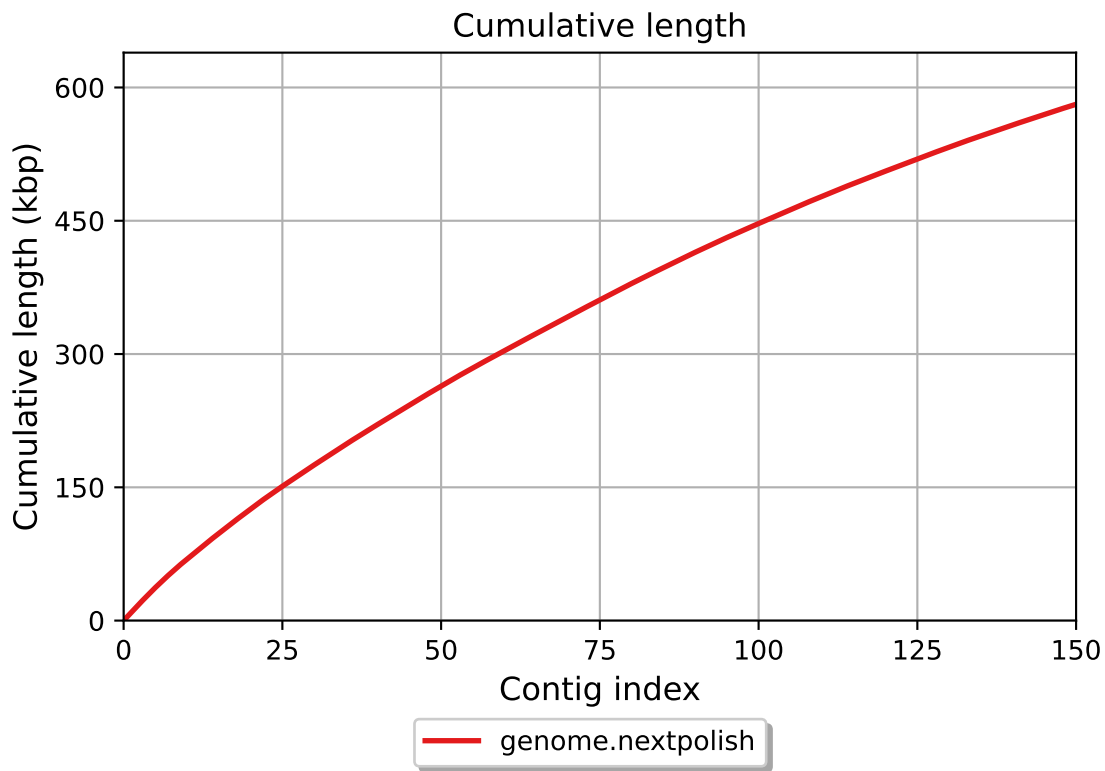
	genome.nextpolish
# contigs (>= 0 bp)	150
# contigs (>= 1000 bp)	150
# contigs (>= 5000 bp)	24
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	581105
Total length (>= 1000 bp)	581105
Total length (>= 5000 bp)	146244
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	150
Largest contig	7742
Total length	581105
GC (%)	48.08
N50	3938
N90	2681
auN	4264.6
L50	57
L90	127
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	1 + 0 part

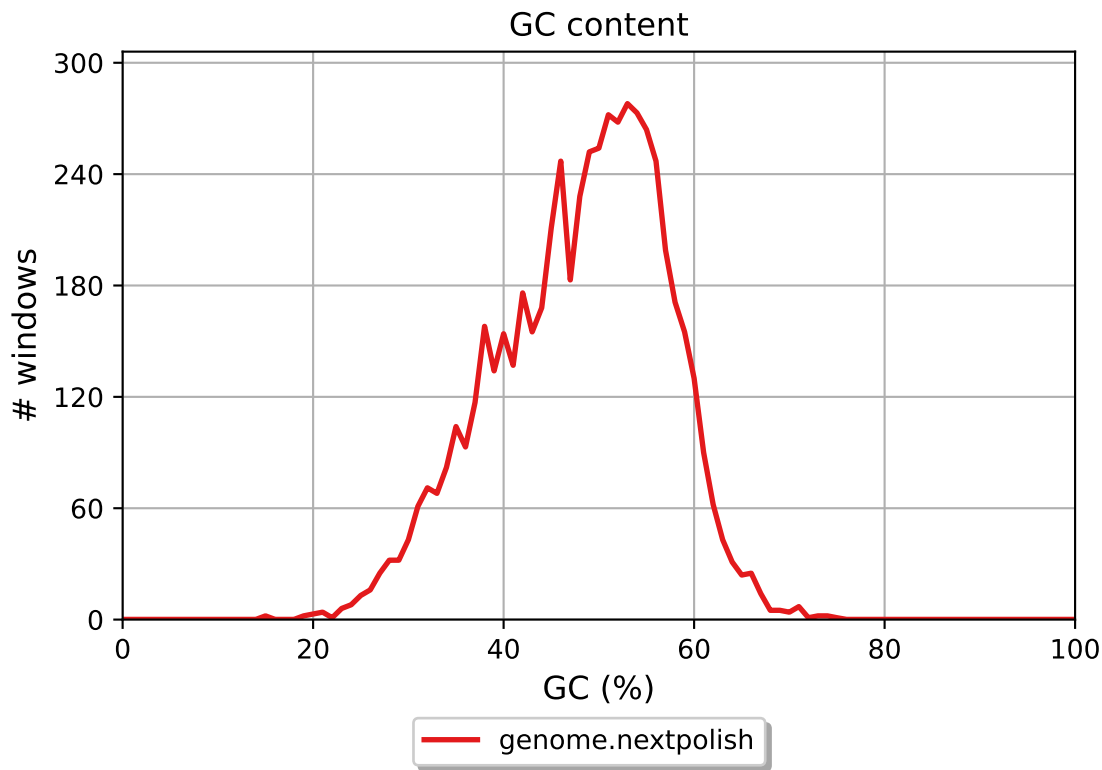
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

