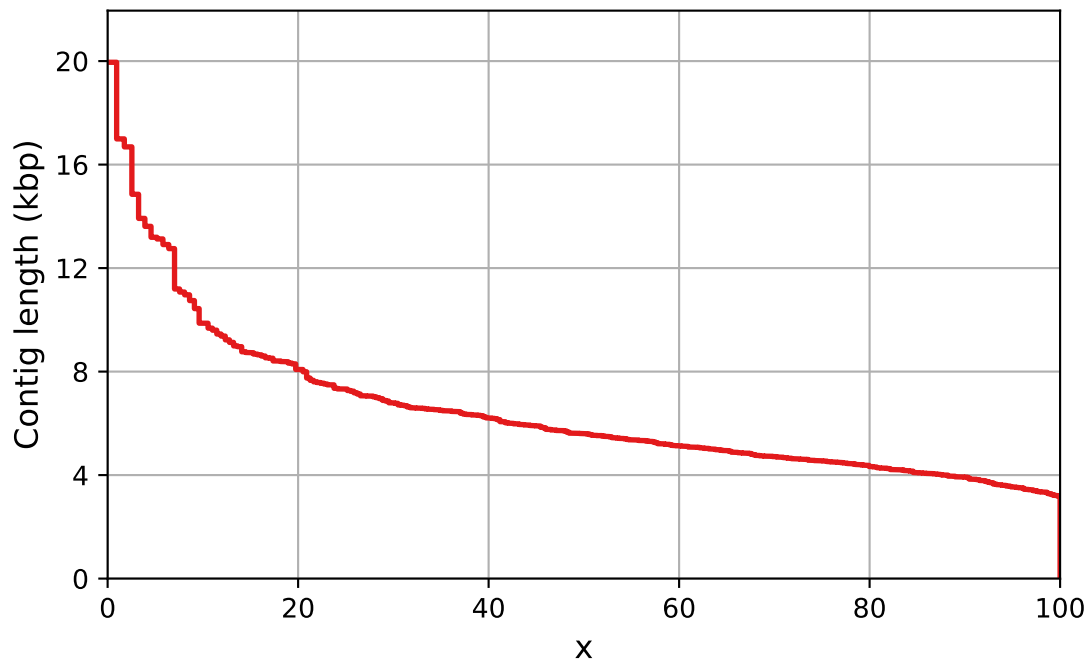


Report

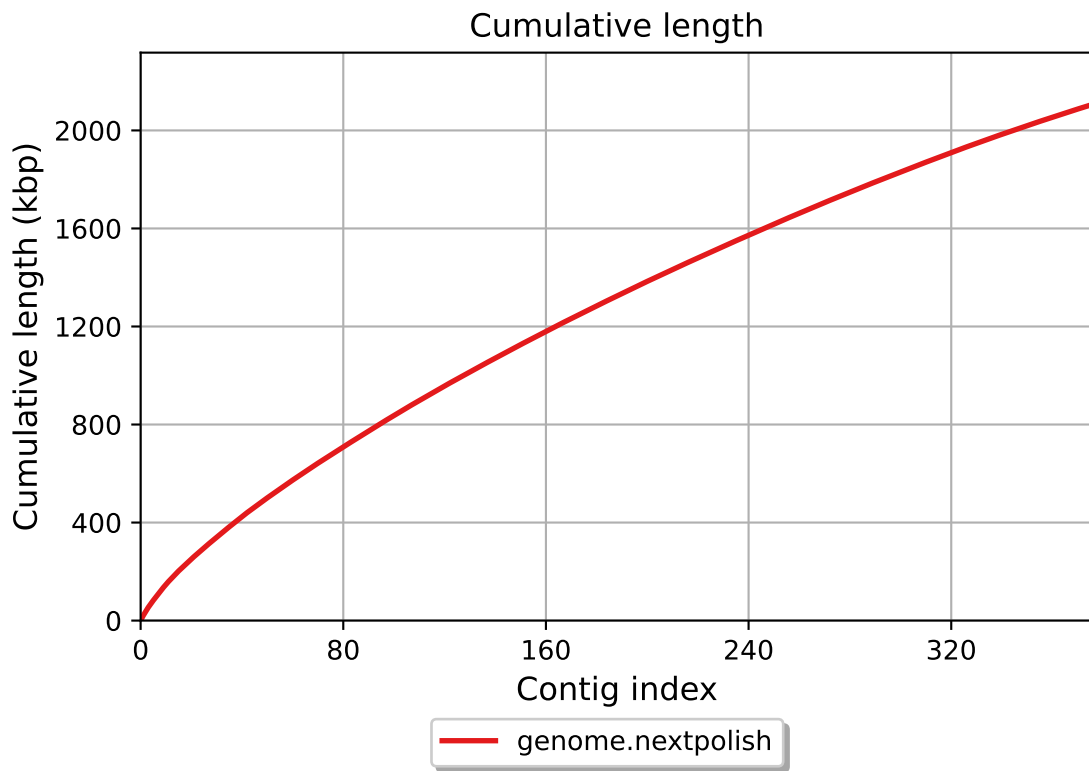
	genome.nextpolish
# contigs (>= 0 bp)	376
# contigs (>= 1000 bp)	376
# contigs (>= 5000 bp)	192
# contigs (>= 10000 bp)	15
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2106723
Total length (>= 1000 bp)	2106723
Total length (>= 5000 bp)	1344544
Total length (>= 10000 bp)	202411
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	376
Largest contig	19954
Total length	2106723
GC (%)	55.20
N50	5608
N90	3922
auN	6490.5
L50	137
L90	317
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 0 part

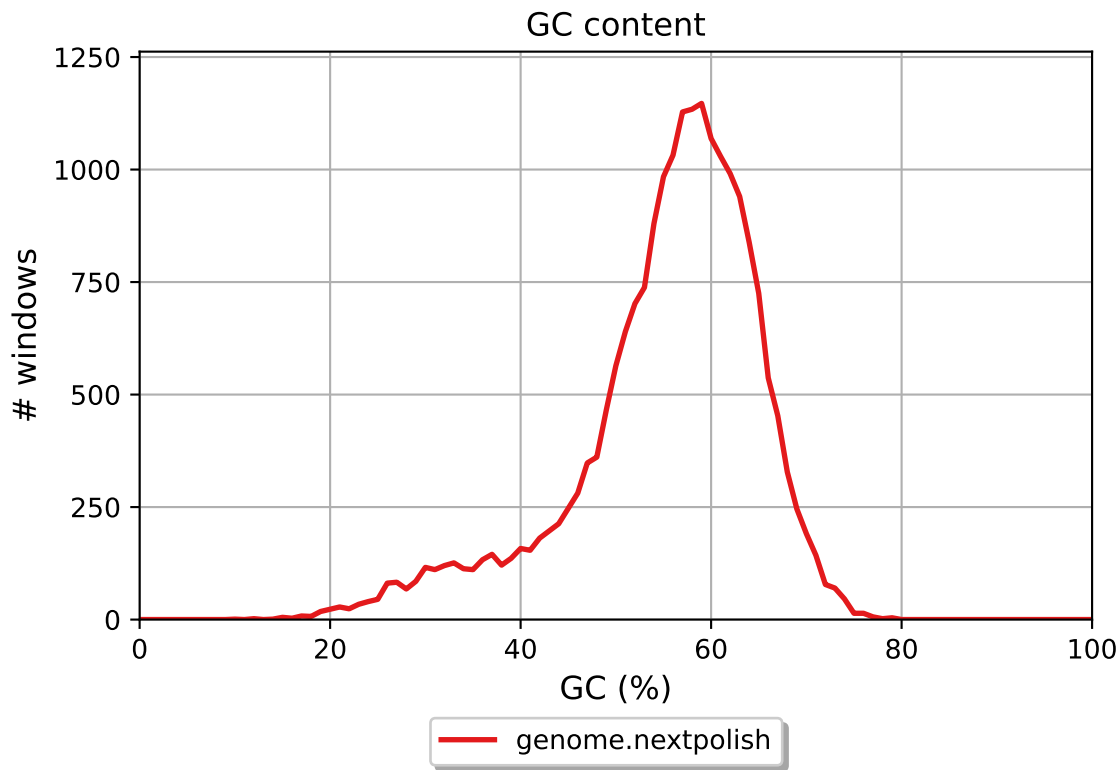
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

