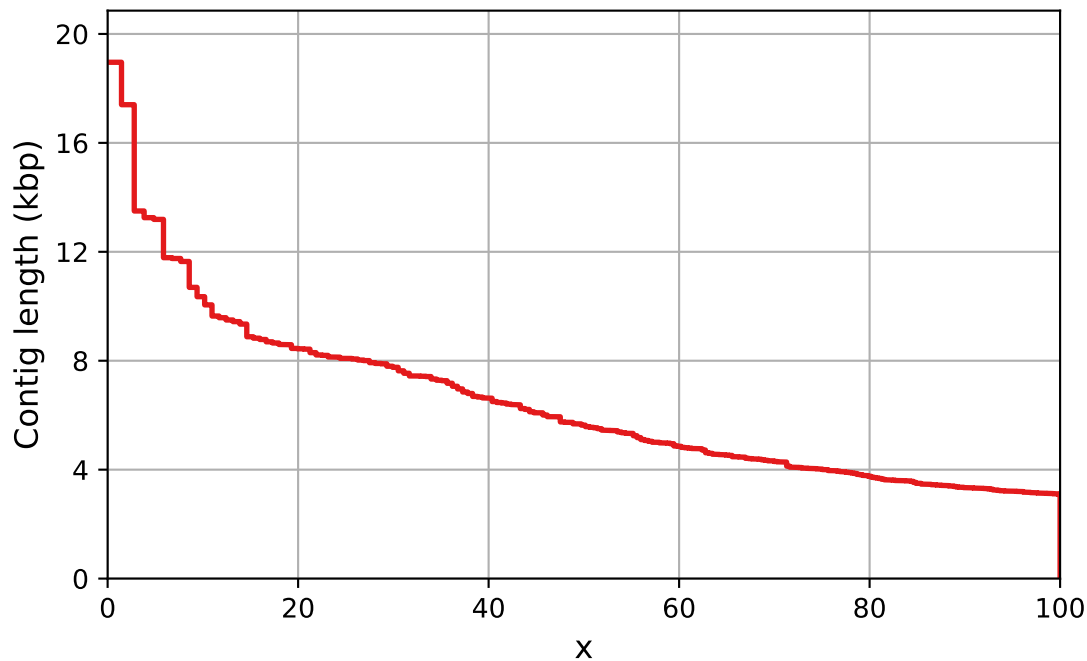


Report

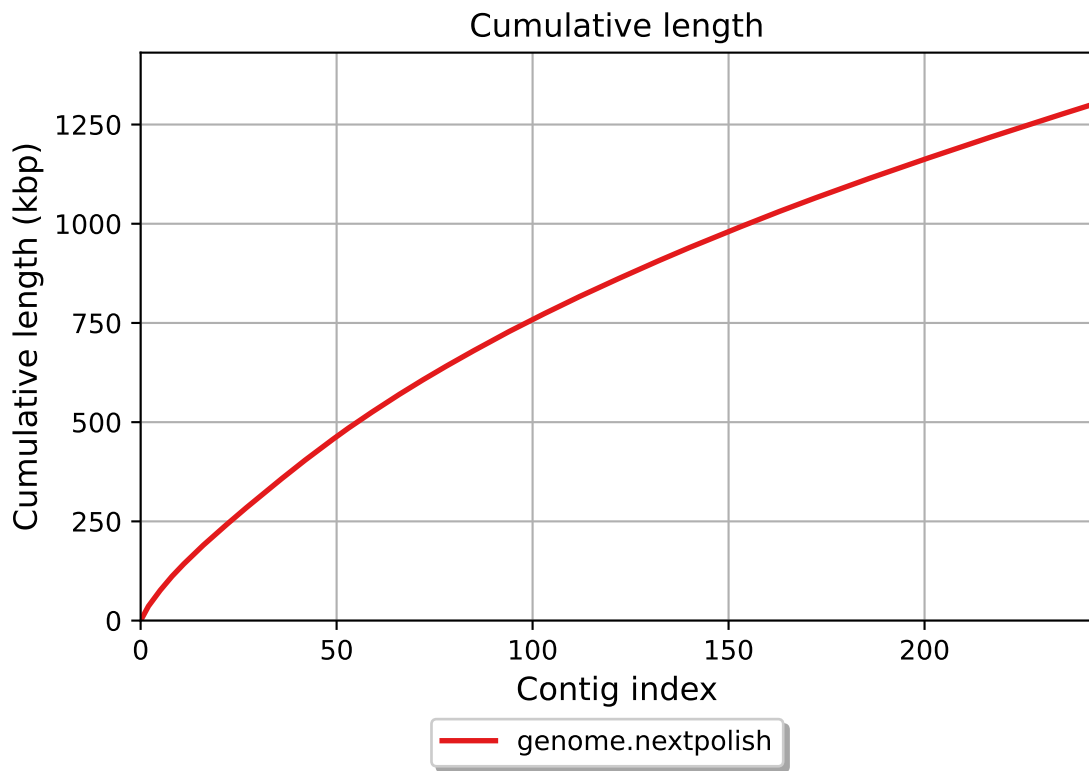
	genome.nextpolish
# contigs (>= 0 bp)	243
# contigs (>= 1000 bp)	243
# contigs (>= 5000 bp)	99
# contigs (>= 10000 bp)	11
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1301059
Total length (>= 1000 bp)	1301059
Total length (>= 5000 bp)	753579
Total length (>= 10000 bp)	142585
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	243
Largest contig	18960
Total length	1301059
GC (%)	48.04
N50	5635
N90	3341
auN	6492.8
L50	80
L90	203
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 0 part

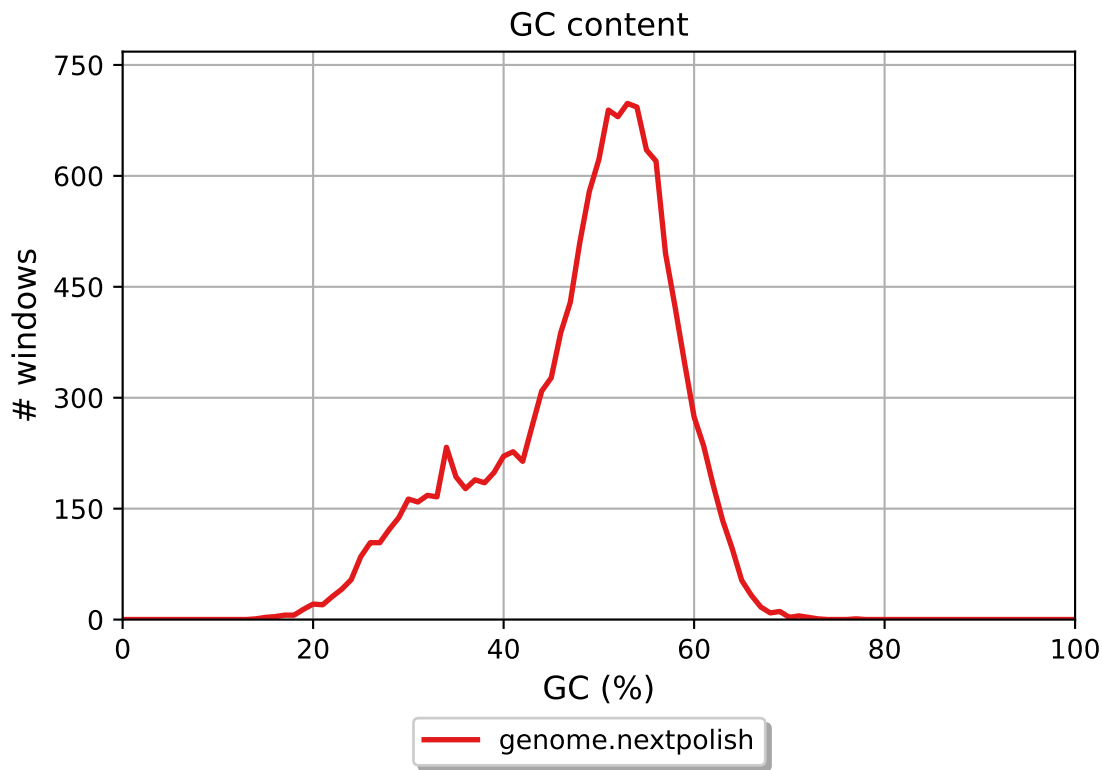
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

