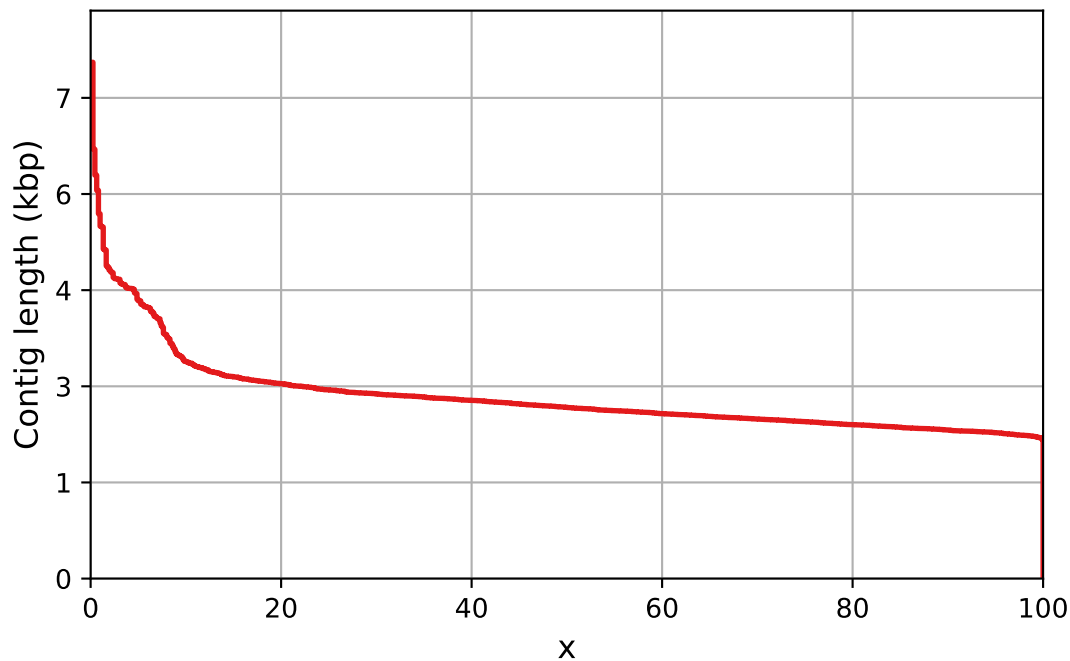


Report

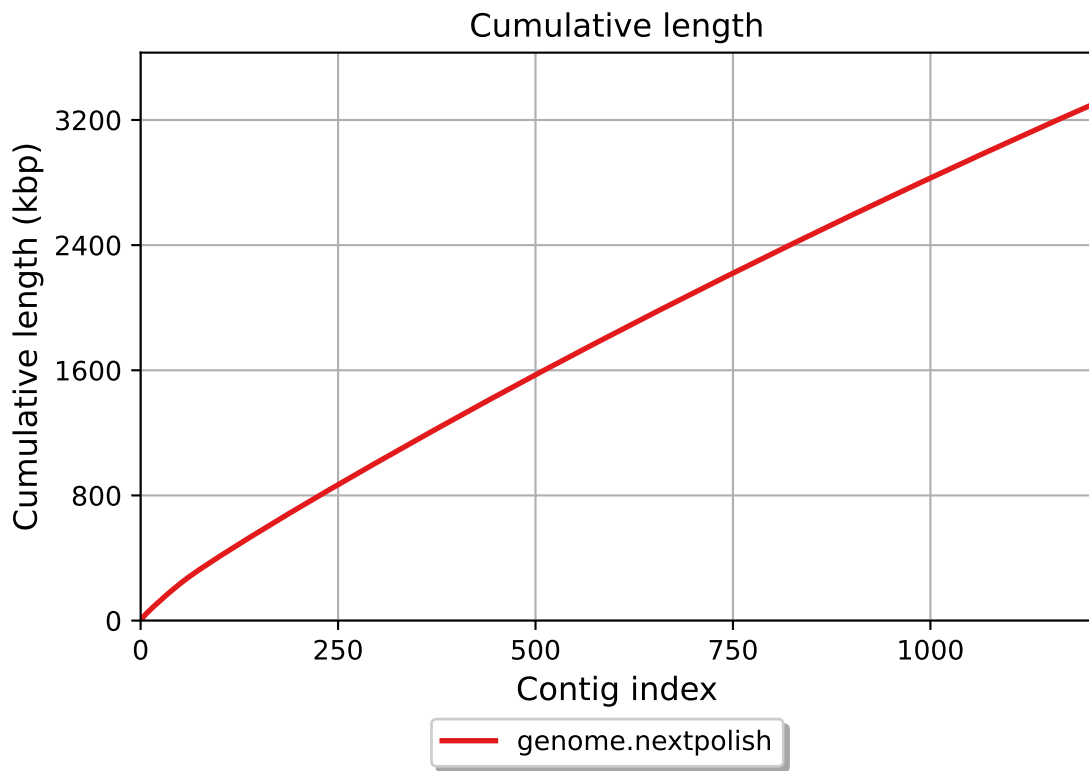
	genome.nextpolish
# contigs (>= 0 bp)	1206
# contigs (>= 1000 bp)	1206
# contigs (>= 5000 bp)	9
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	3300388
Total length (>= 1000 bp)	3300388
Total length (>= 5000 bp)	54044
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	1206
Largest contig	8055
Total length	3300388
GC (%)	45.34
N50	2671
N90	2318
auN	2841.2
L50	530
L90	1061
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	5 + 2 part

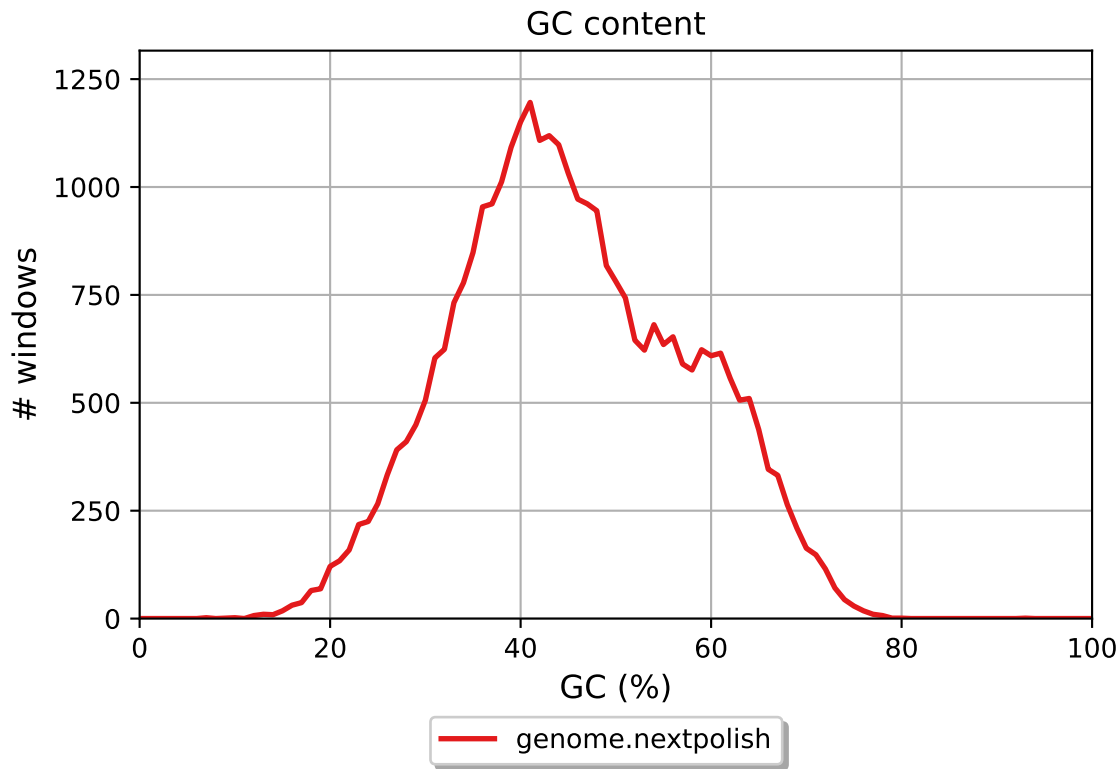
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

