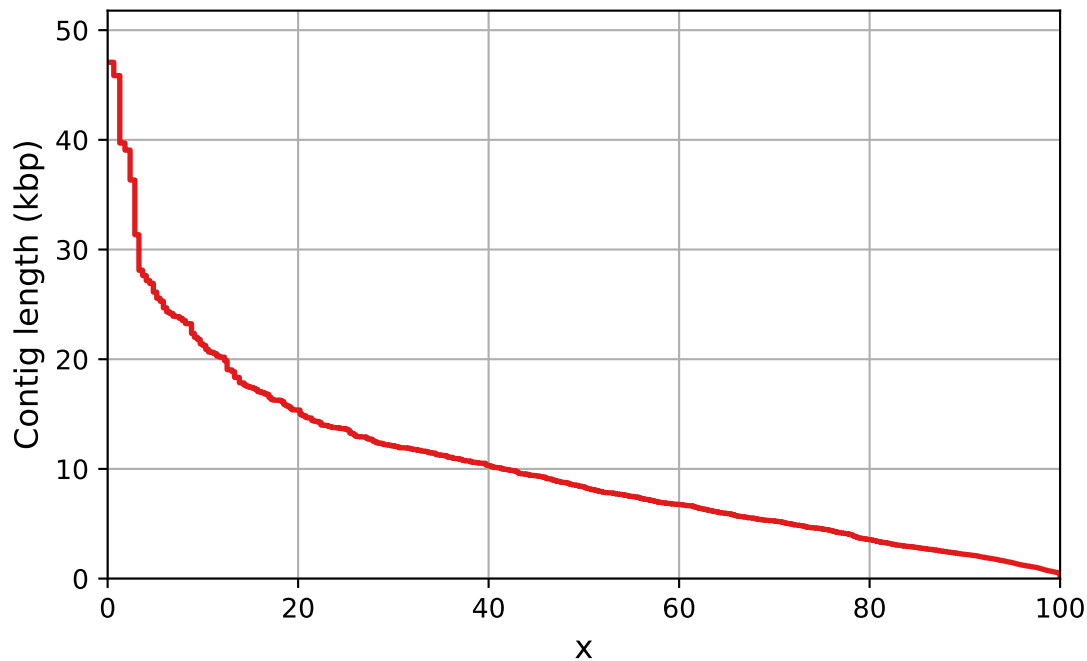


Report

	genome.nextpolish
# contigs (>= 0 bp)	1522
# contigs (>= 1000 bp)	1268
# contigs (>= 5000 bp)	508
# contigs (>= 10000 bp)	196
# contigs (>= 25000 bp)	13
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	7278966
Total length (>= 1000 bp)	7101484
Total length (>= 5000 bp)	5211156
Total length (>= 10000 bp)	3010797
Total length (>= 25000 bp)	426160
Total length (>= 50000 bp)	0
# contigs	1518
Largest contig	47072
Total length	7277088
GC (%)	54.19
N50	8359
N90	2214
auN	10395.1
L50	265
L90	912
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	1 + 0 part

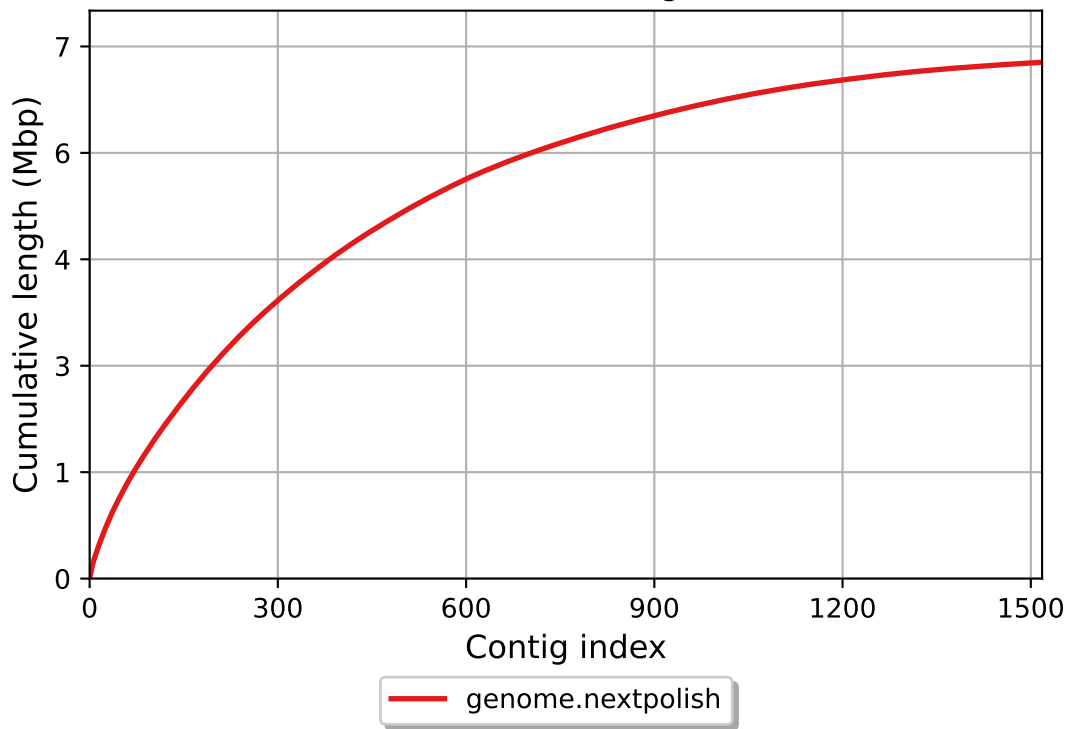
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

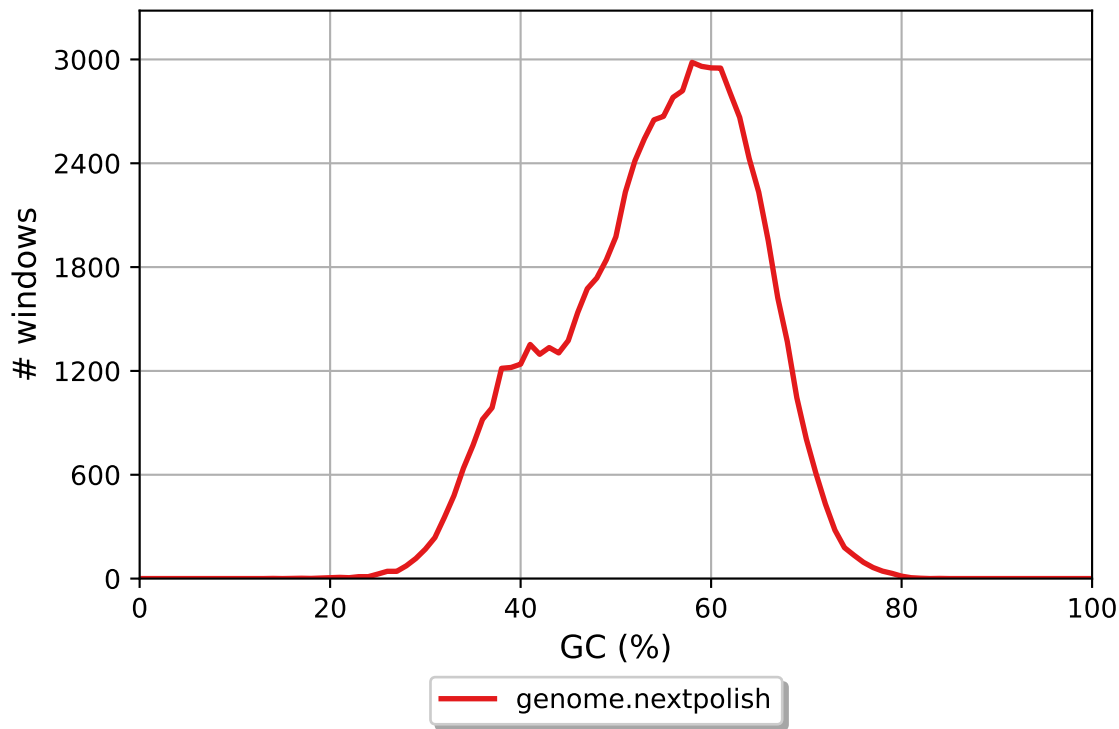


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

