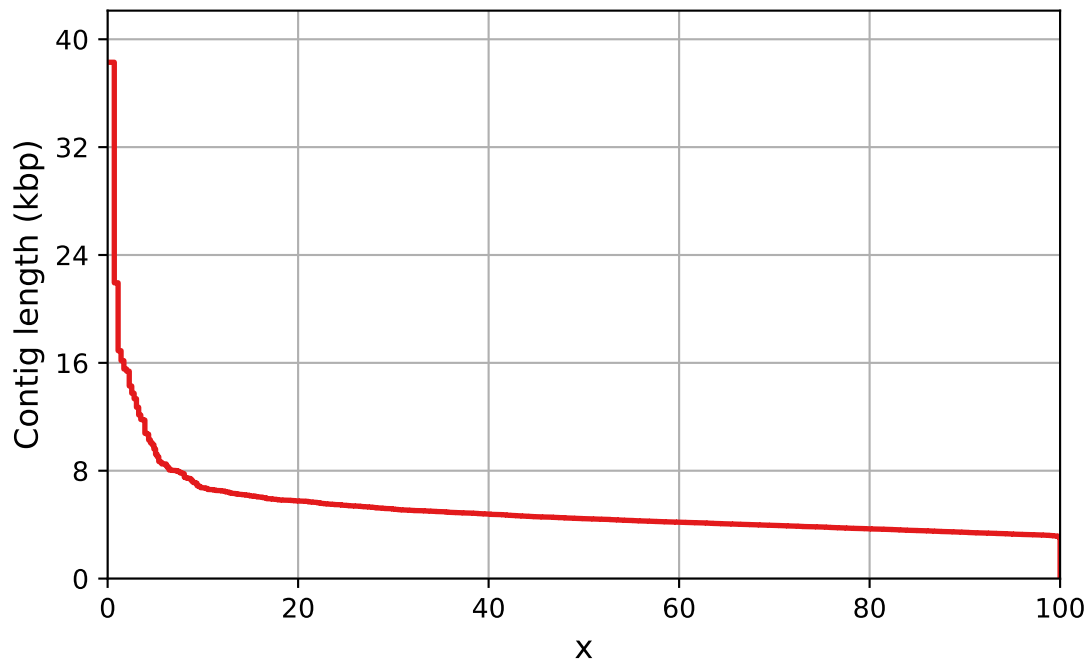


# Report

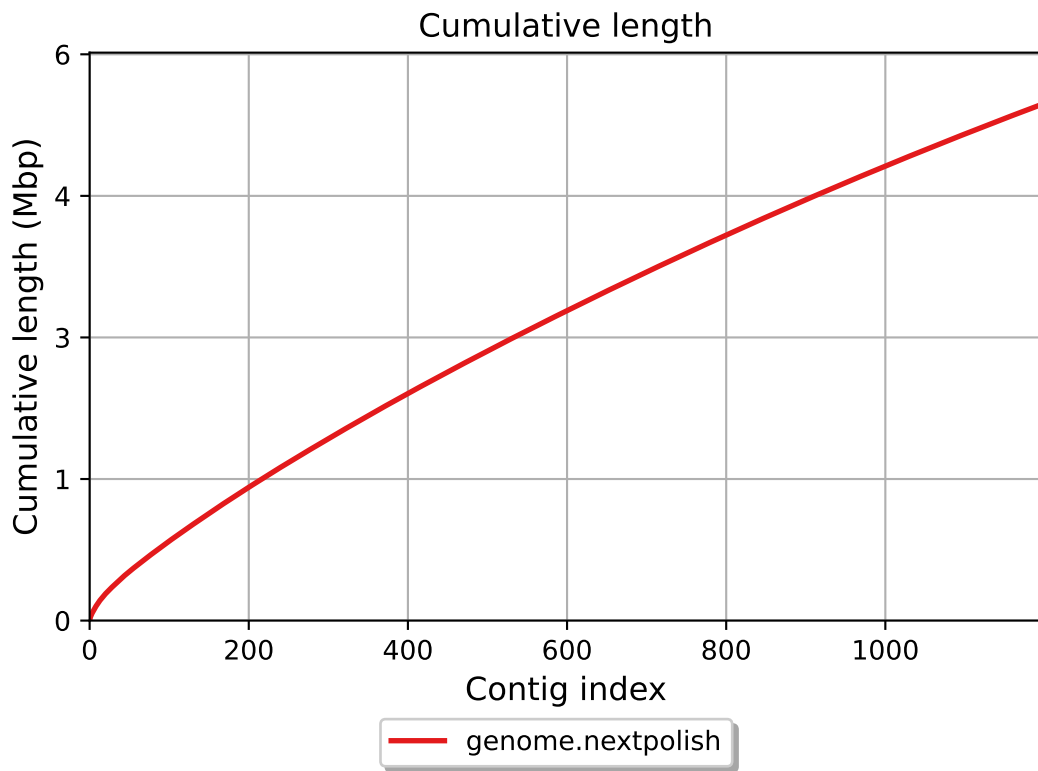
	genome.nextpolish
# contigs (>= 0 bp)	1197
# contigs (>= 1000 bp)	1197
# contigs (>= 5000 bp)	287
# contigs (>= 10000 bp)	17
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	5470608
Total length (>= 1000 bp)	5470608
Total length (>= 5000 bp)	1862331
Total length (>= 10000 bp)	255833
Total length (>= 25000 bp)	38293
Total length (>= 50000 bp)	0
# contigs	1197
Largest contig	38293
Total length	5470608
GC (%)	39.31
N50	4446
N90	3433
auN	5282.4
L50	473
L90	1032
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	15 + 7 part

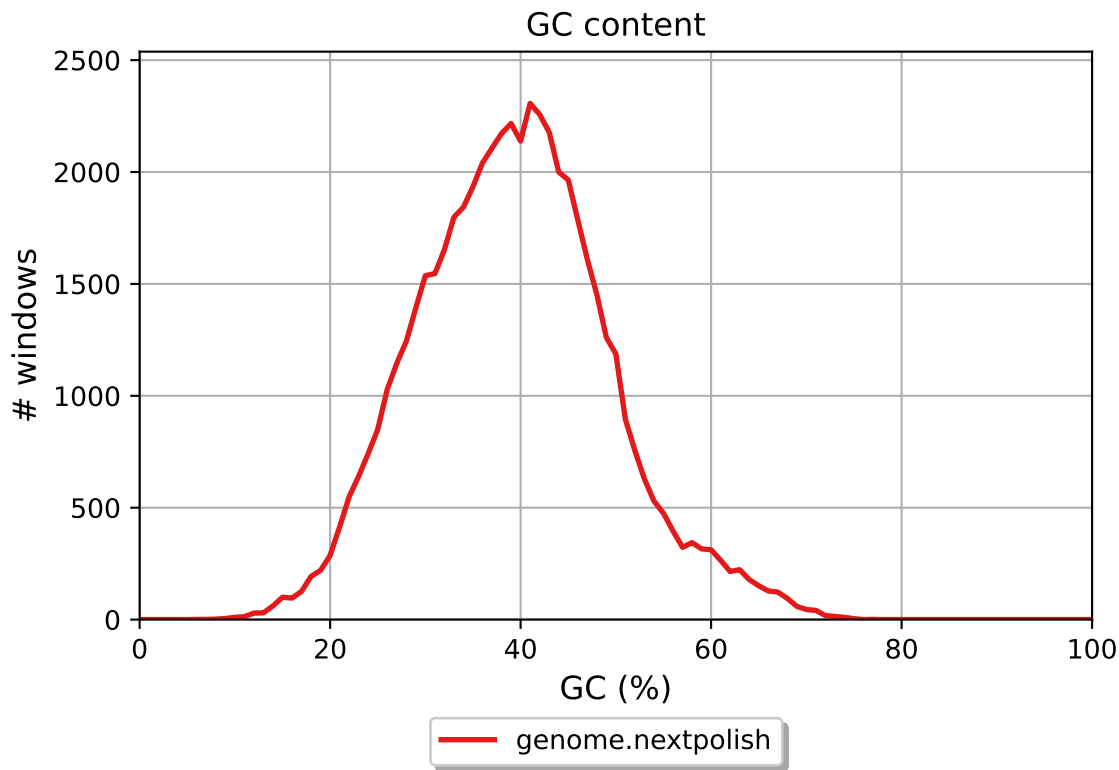
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

