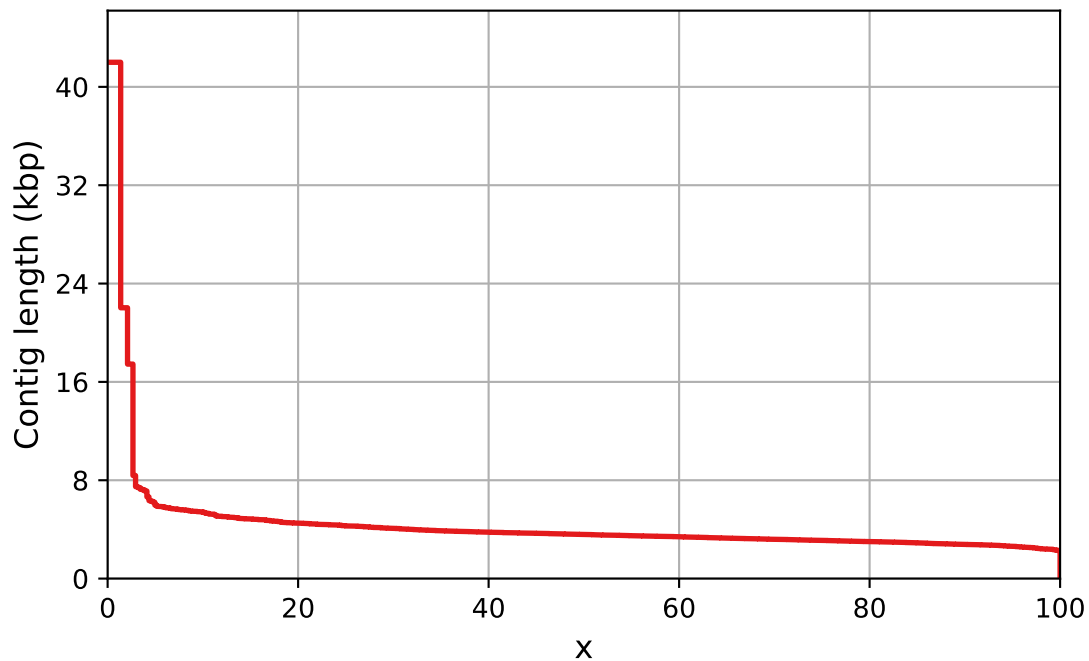


# Report

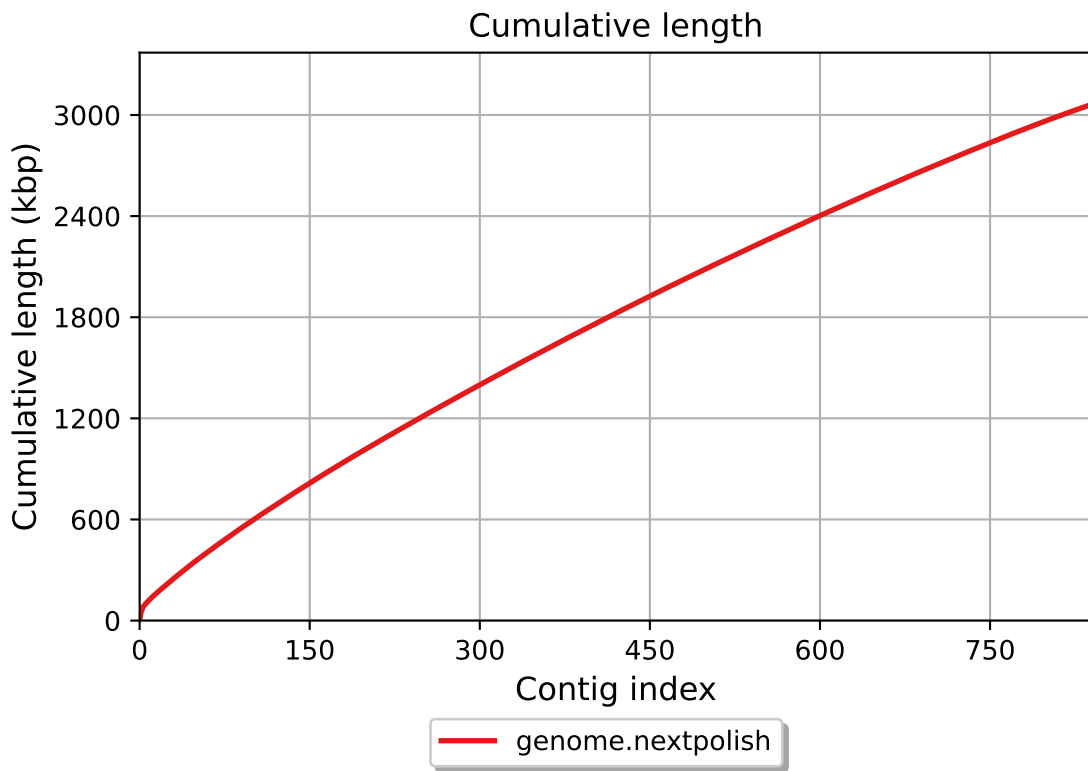
	genome.nextpolish
# contigs (>= 0 bp)	840
# contigs (>= 1000 bp)	840
# contigs (>= 5000 bp)	56
# contigs (>= 10000 bp)	3
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	3063522
Total length (>= 1000 bp)	3063522
Total length (>= 5000 bp)	387040
Total length (>= 10000 bp)	81478
Total length (>= 25000 bp)	42000
Total length (>= 50000 bp)	0
# contigs	840
Largest contig	42000
Total length	3063522
GC (%)	43.51
N50	3593
N90	2780
auN	4495.5
L50	337
L90	722
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	8 + 4 part

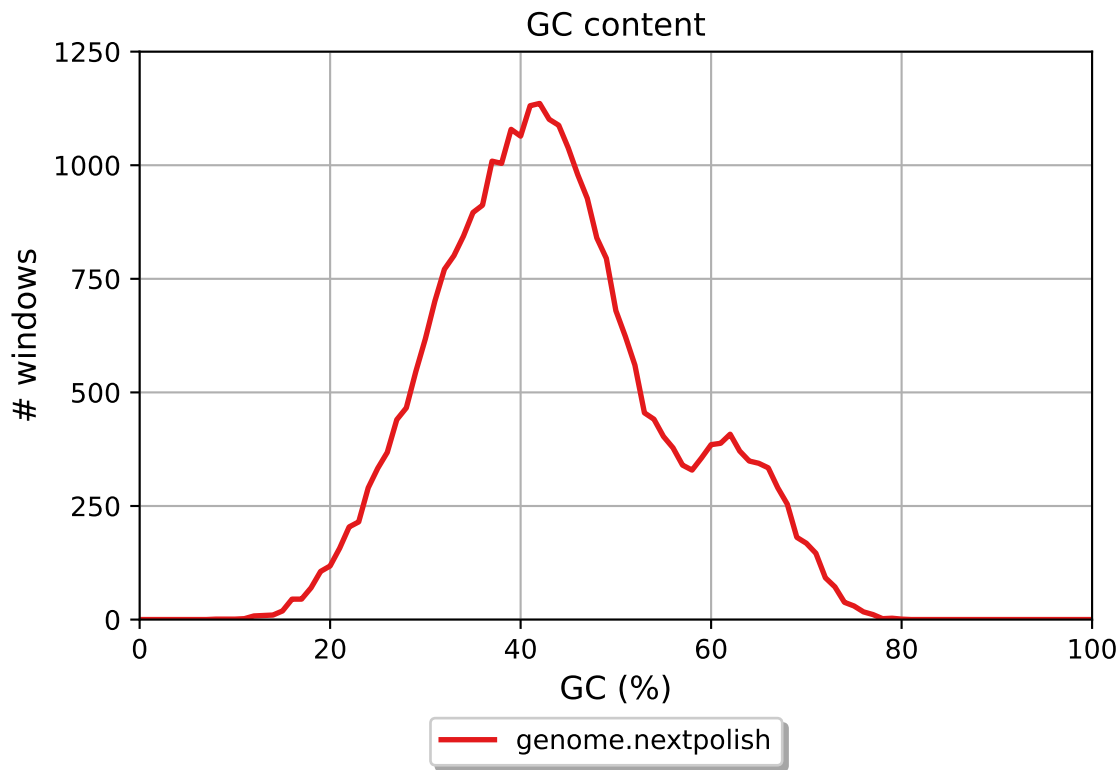
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

