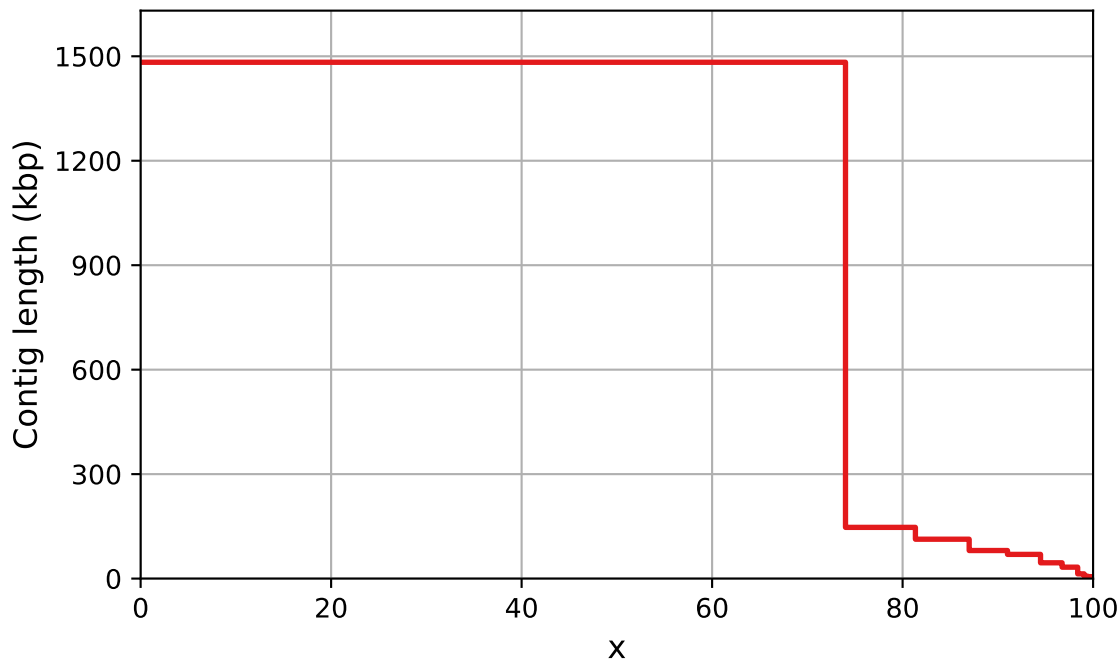


# Report

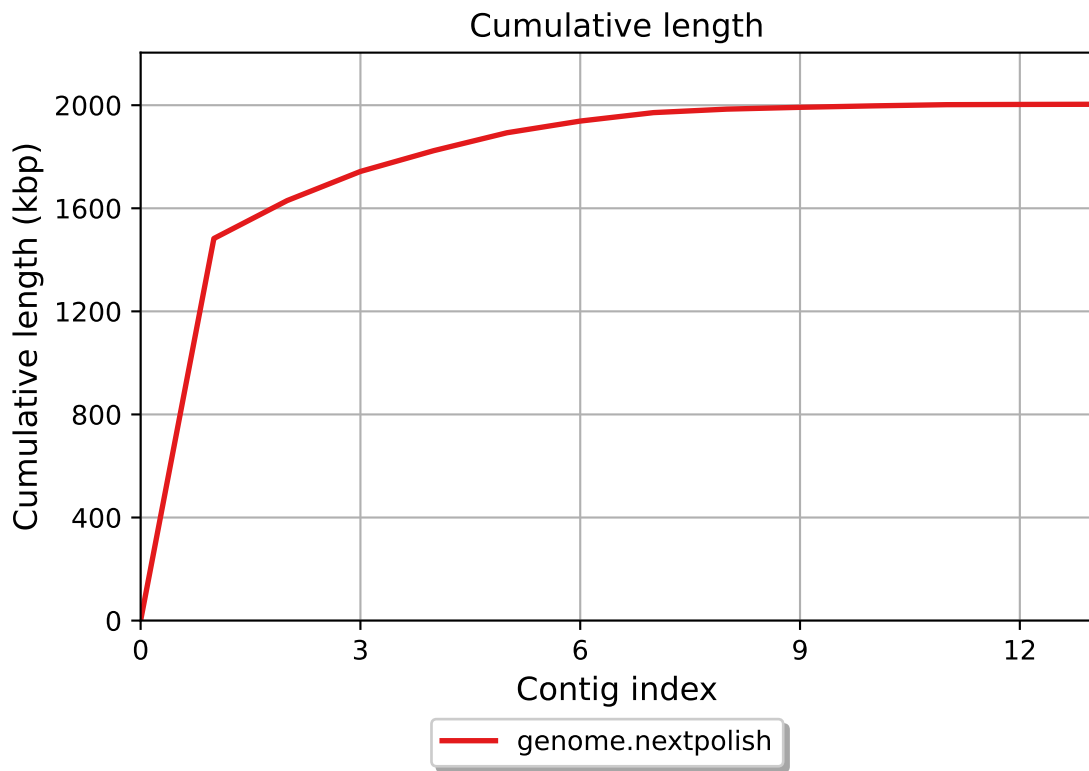
	genome.nextpolish
# contigs (>= 0 bp)	13
# contigs (>= 1000 bp)	11
# contigs (>= 5000 bp)	10
# contigs (>= 10000 bp)	8
# contigs (>= 25000 bp)	7
# contigs (>= 50000 bp)	5
Total length (>= 0 bp)	2003638
Total length (>= 1000 bp)	2002063
Total length (>= 5000 bp)	1997426
Total length (>= 10000 bp)	1984786
Total length (>= 25000 bp)	1971226
Total length (>= 50000 bp)	1893071
# contigs	13
Largest contig	1482783
Total length	2003638
GC (%)	41.30
N50	1482783
N90	80606
auN	1121867.7
L50	1
L90	4
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	8 + 1 part

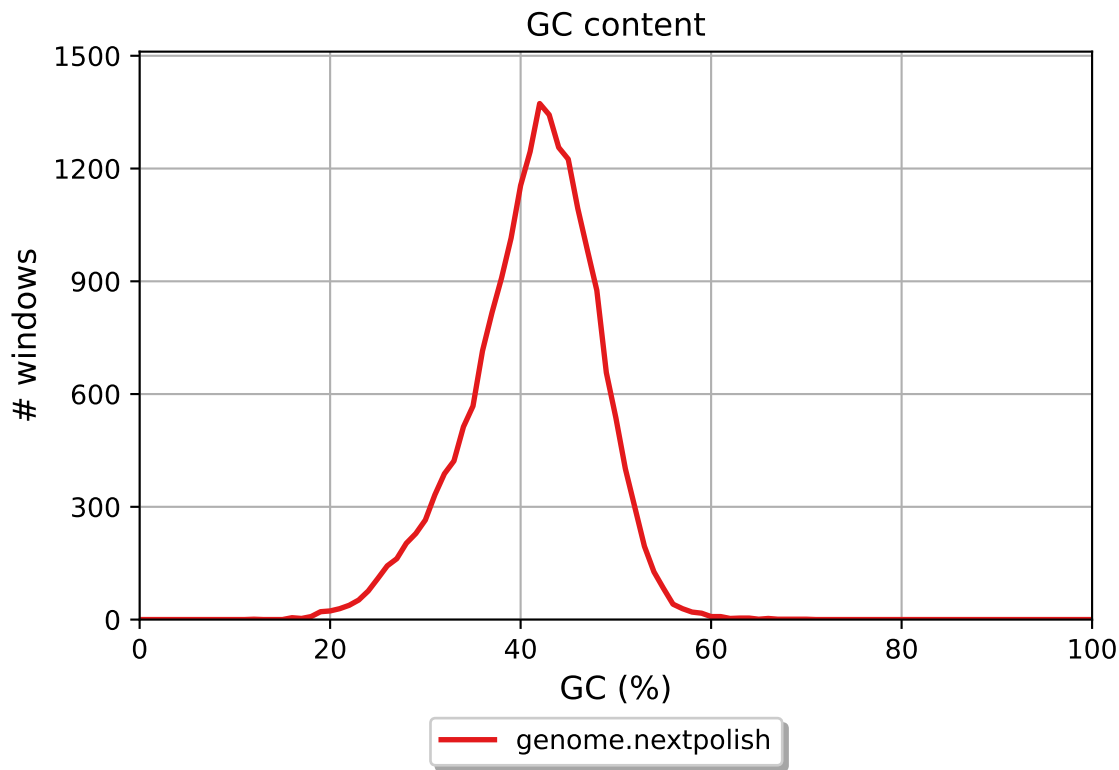
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

