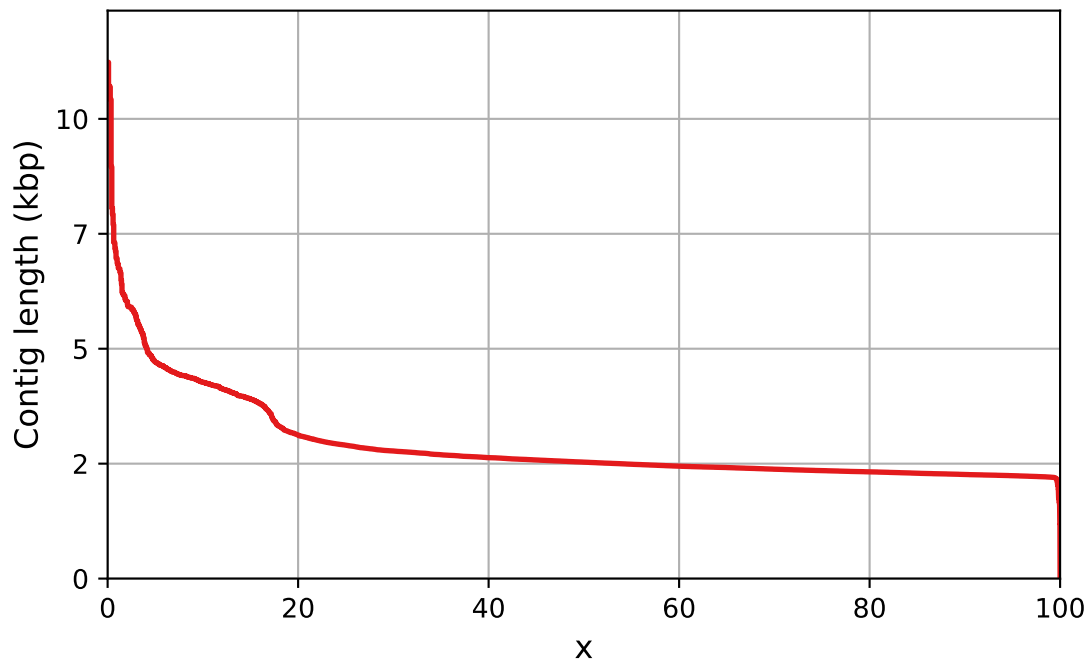


# Report

	genome.nextpolish
# contigs (>= 0 bp)	4376
# contigs (>= 1000 bp)	4376
# contigs (>= 5000 bp)	78
# contigs (>= 10000 bp)	4
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	11846934
Total length (>= 1000 bp)	11846934
Total length (>= 5000 bp)	490550
Total length (>= 10000 bp)	43069
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	4376
Largest contig	11230
Total length	11846934
GC (%)	48.11
N50	2533
N90	2265
auN	2910.8
L50	1854
L90	3842
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	12 + 10 part

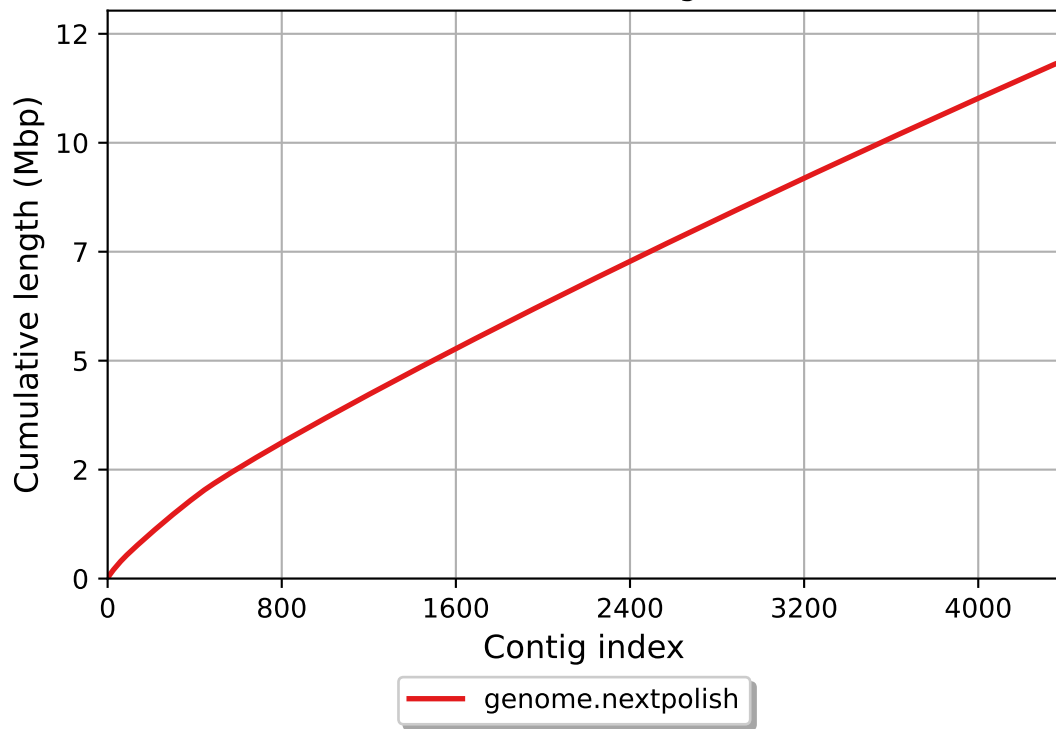
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

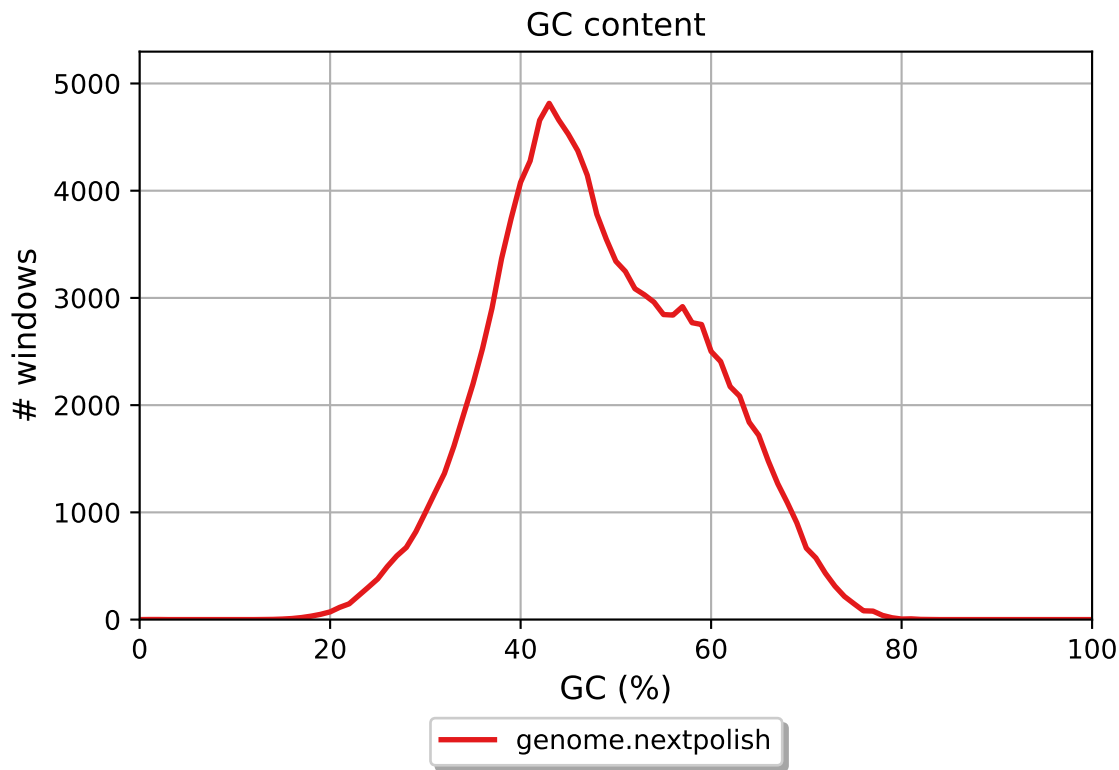
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

