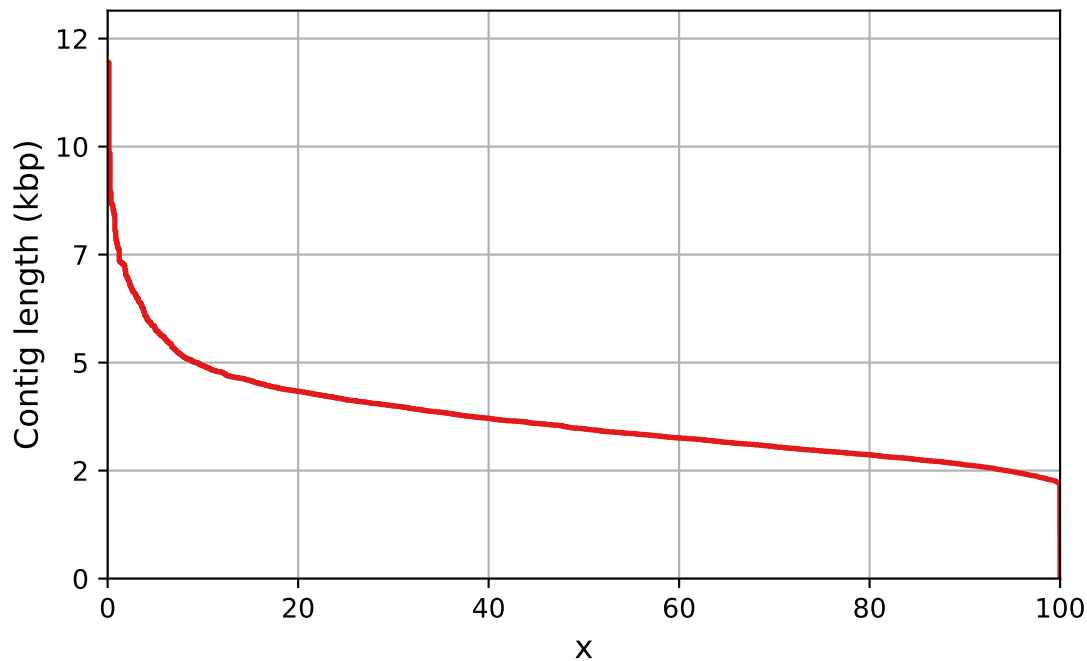


Report

	genome.nextpolish
# contigs (>= 0 bp)	2467
# contigs (>= 1000 bp)	2467
# contigs (>= 5000 bp)	131
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	8539679
Total length (>= 1000 bp)	8539679
Total length (>= 5000 bp)	794923
Total length (>= 10000 bp)	11951
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	2467
Largest contig	11951
Total length	8539679
GC (%)	46.43
N50	3468
N90	2631
auN	3704.6
L50	995
L90	2120
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	8 + 5 part

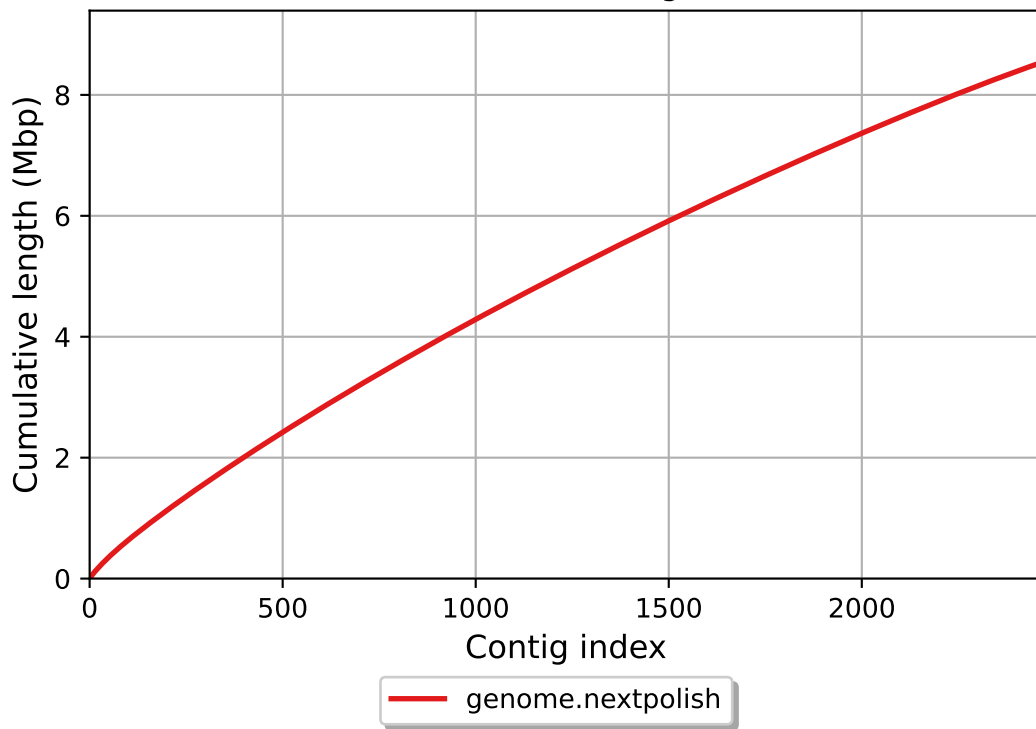
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

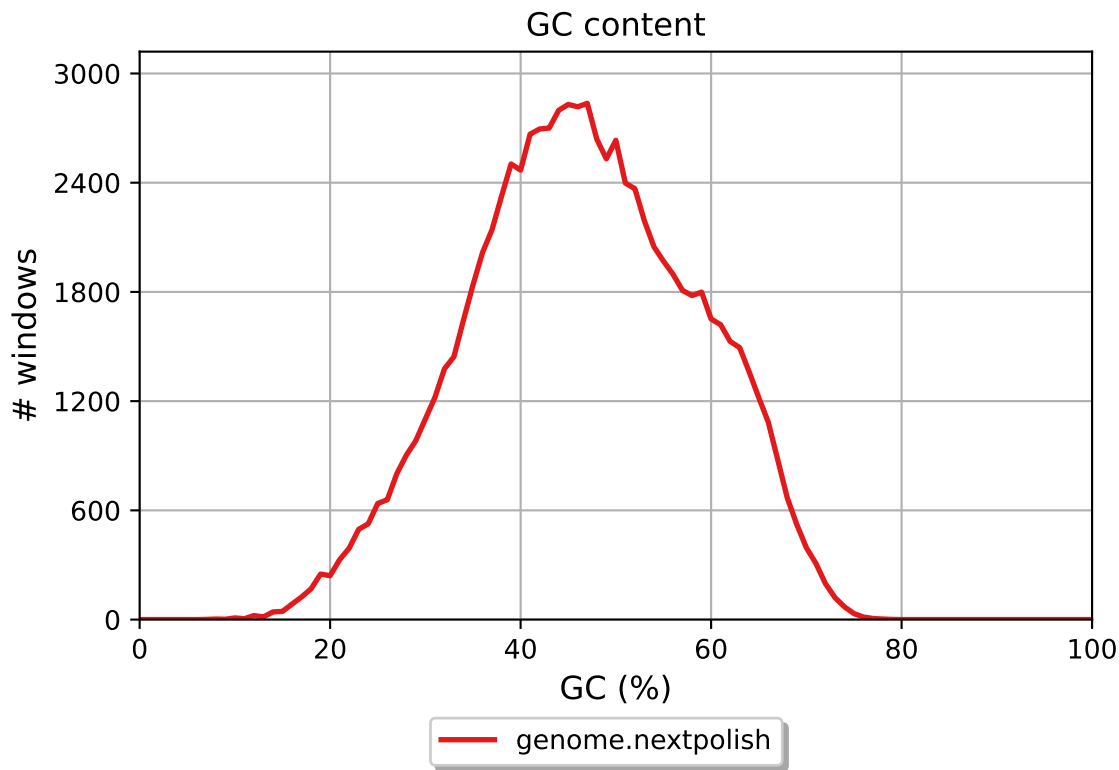
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

