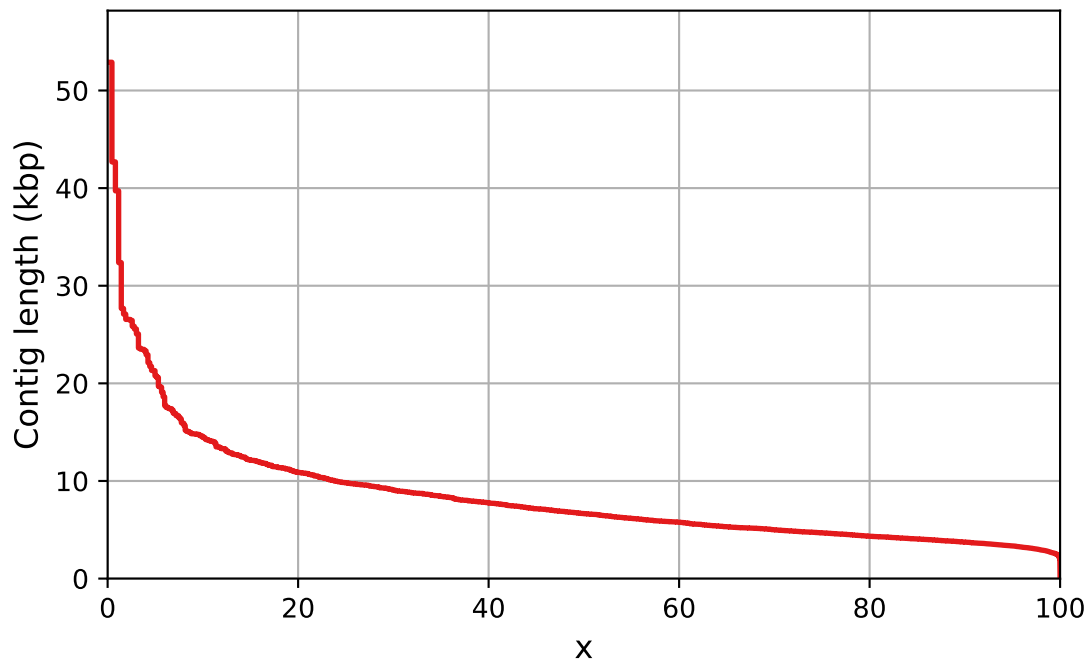


# Report

	genome.nextpolish
# contigs (>= 0 bp)	1886
# contigs (>= 1000 bp)	1886
# contigs (>= 5000 bp)	979
# contigs (>= 10000 bp)	194
# contigs (>= 25000 bp)	12
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	11694908
Total length (>= 1000 bp)	11694908
Total length (>= 5000 bp)	8174840
Total length (>= 10000 bp)	2781377
Total length (>= 25000 bp)	378466
Total length (>= 50000 bp)	52891
# contigs	1886
Largest contig	52891
Total length	11694908
GC (%)	39.74
N50	6655
N90	3735
auN	8465.1
L50	573
L90	1521
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	20 + 2 part

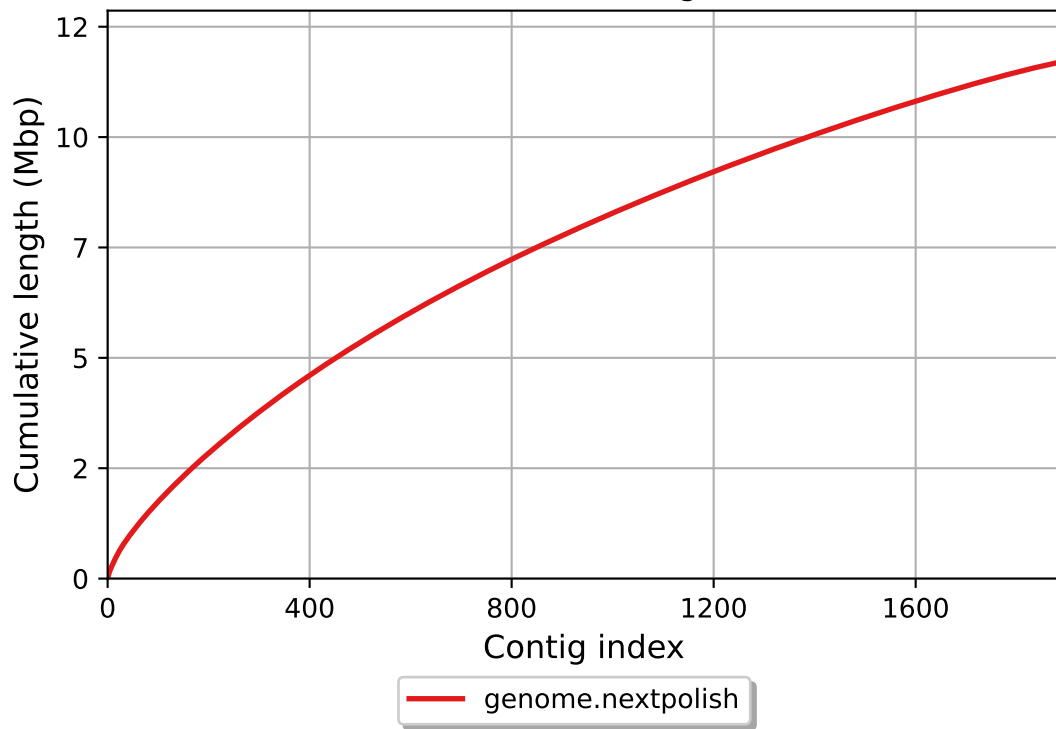
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

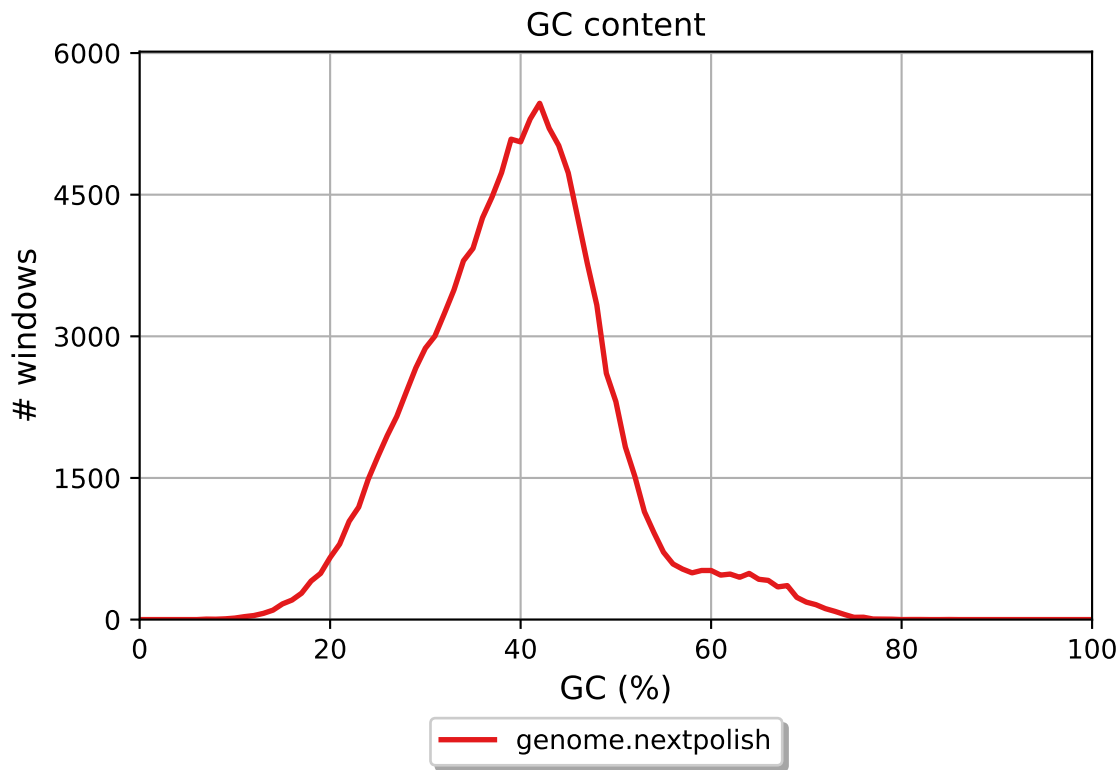
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

