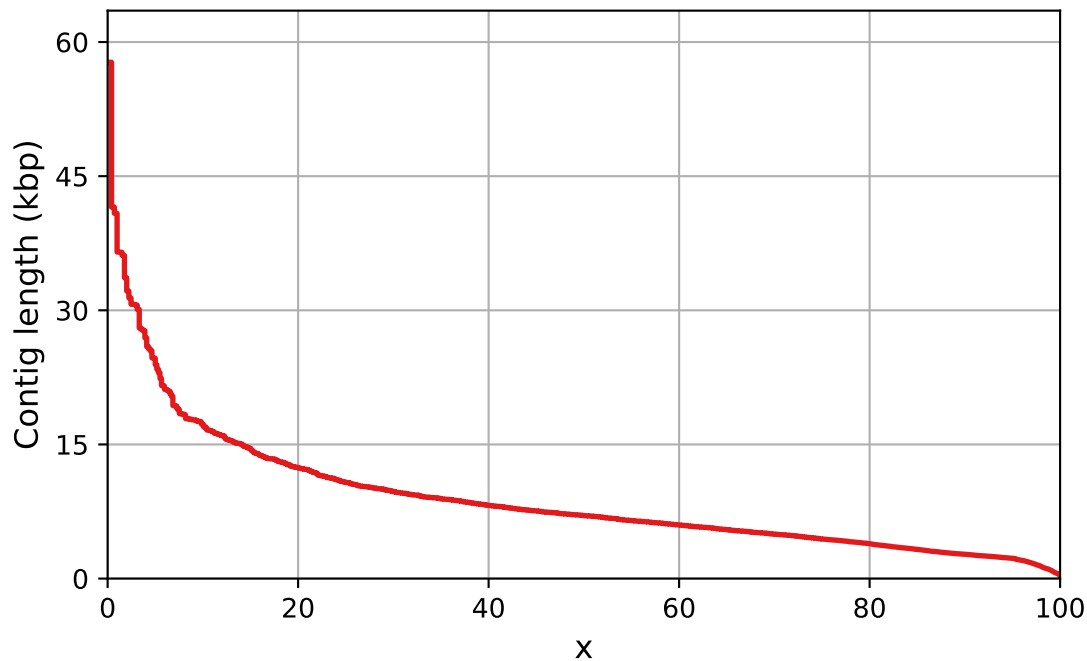


Report

	genome.nextpolish
# contigs (>= 0 bp)	2722
# contigs (>= 1000 bp)	2487
# contigs (>= 5000 bp)	1099
# contigs (>= 10000 bp)	271
# contigs (>= 25000 bp)	20
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	14113096
Total length (>= 1000 bp)	13956301
Total length (>= 5000 bp)	9838639
Total length (>= 10000 bp)	4077422
Total length (>= 25000 bp)	656101
Total length (>= 50000 bp)	57737
# contigs	2715
Largest contig	57737
Total length	14110282
GC (%)	46.71
N50	7040
N90	2720
auN	9027.2
L50	632
L90	1863
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	26 + 7 part

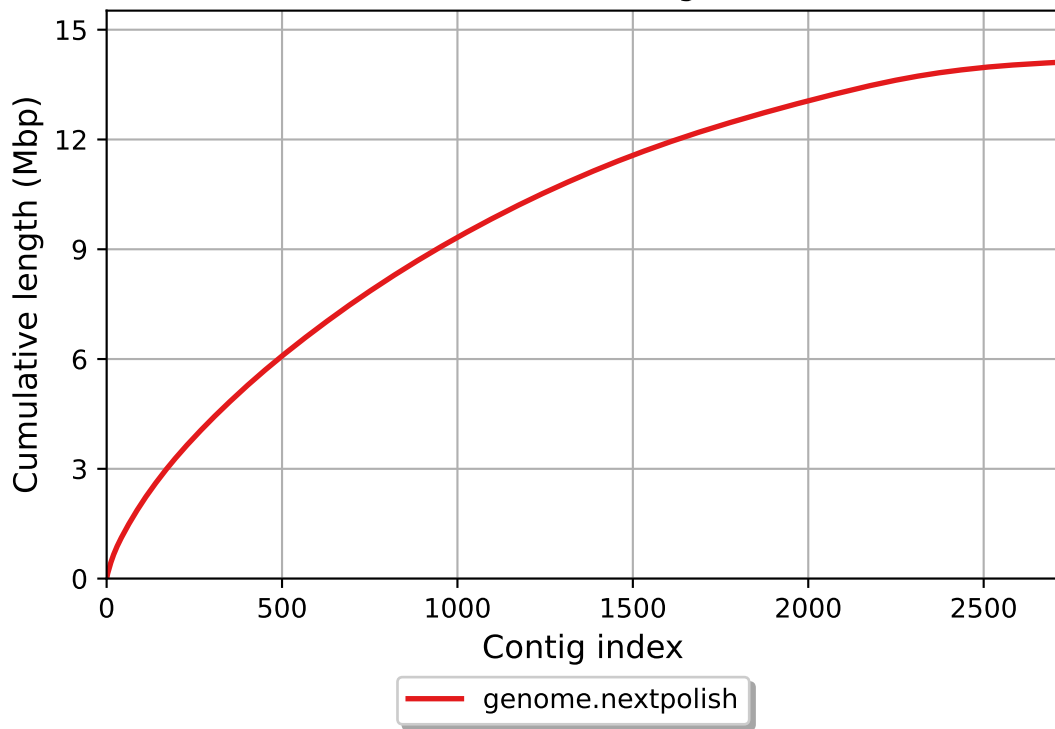
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

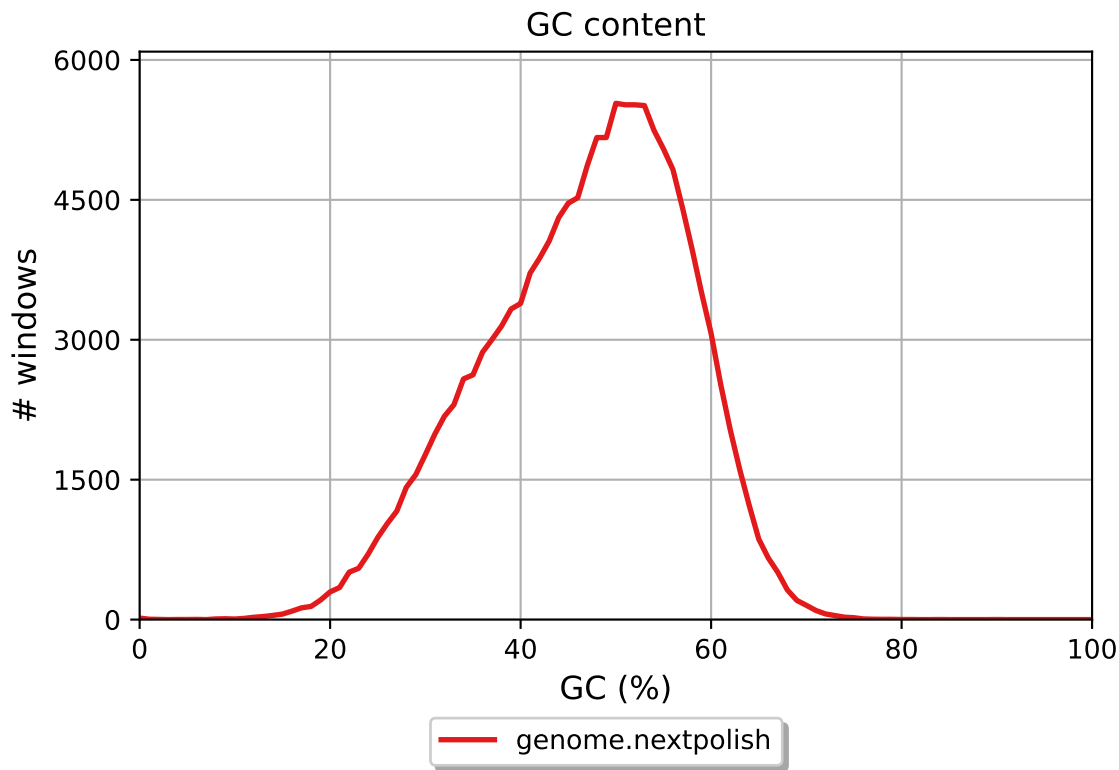
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

