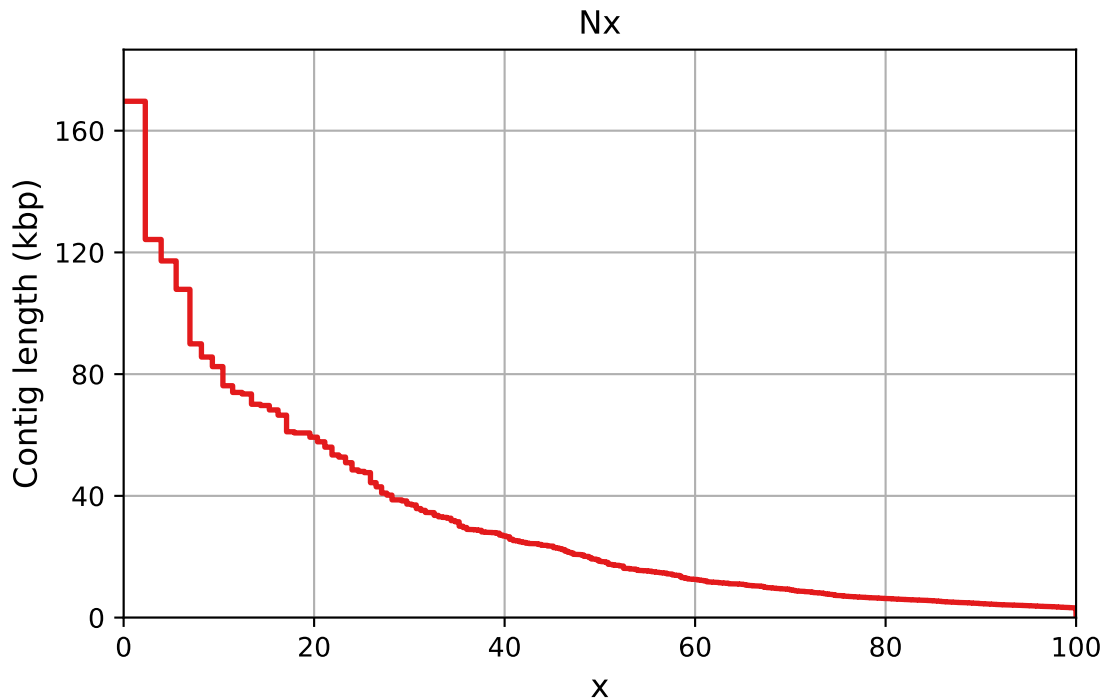


Report

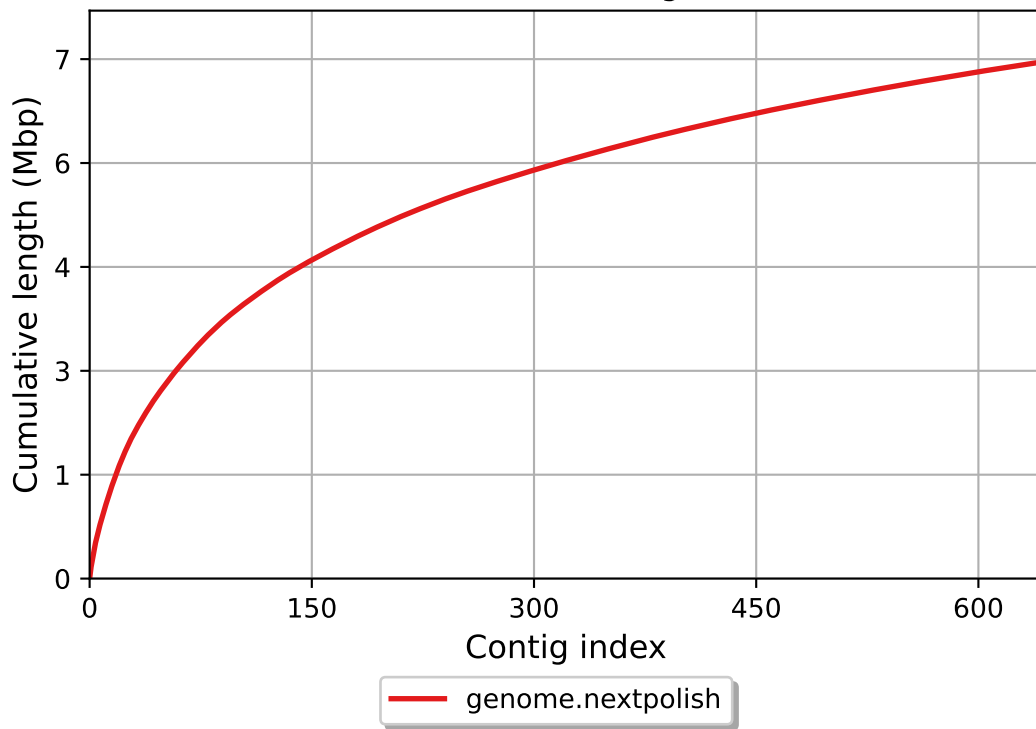
	genome.nextpolish
# contigs (>= 0 bp)	643
# contigs (>= 1000 bp)	641
# contigs (>= 5000 bp)	409
# contigs (>= 10000 bp)	188
# contigs (>= 25000 bp)	62
# contigs (>= 50000 bp)	23
Total length (>= 0 bp)	7454780
Total length (>= 1000 bp)	7452967
Total length (>= 5000 bp)	6522201
Total length (>= 10000 bp)	5015341
Total length (>= 25000 bp)	3097668
Total length (>= 50000 bp)	1787762
# contigs	643
Largest contig	169667
Total length	7454780
GC (%)	45.50
N50	18516
N90	4576
auN	33318.2
L50	91
L90	448
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	18 + 4 part

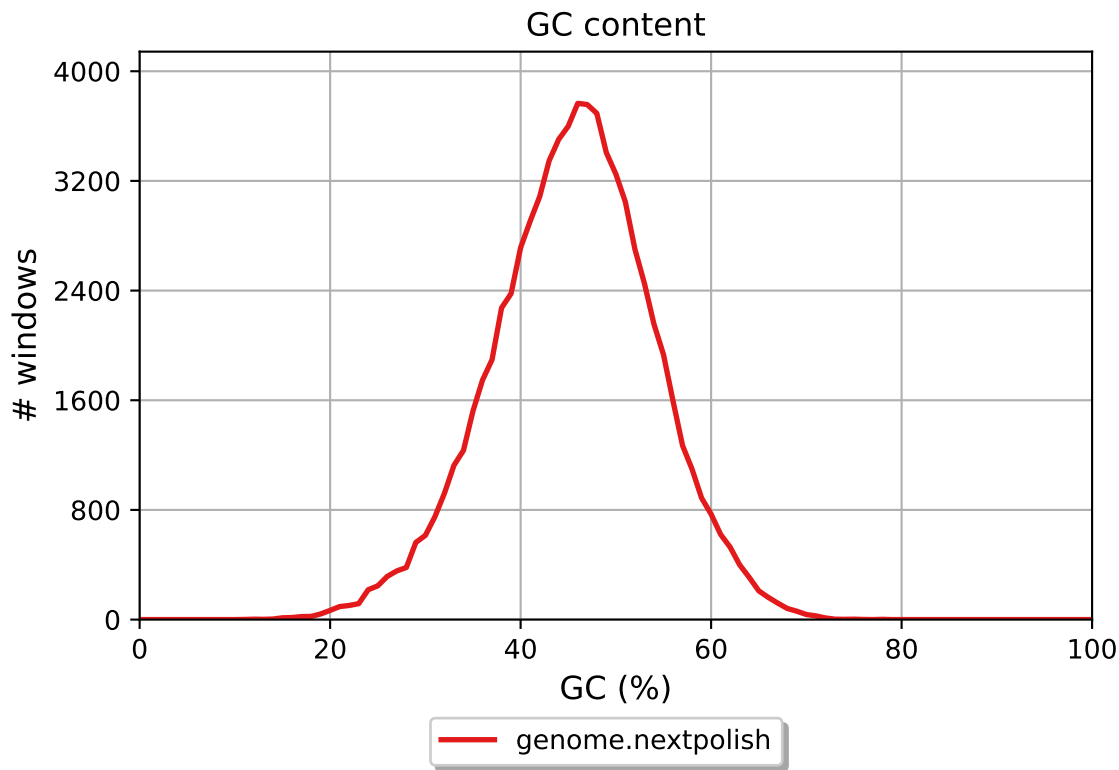
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

