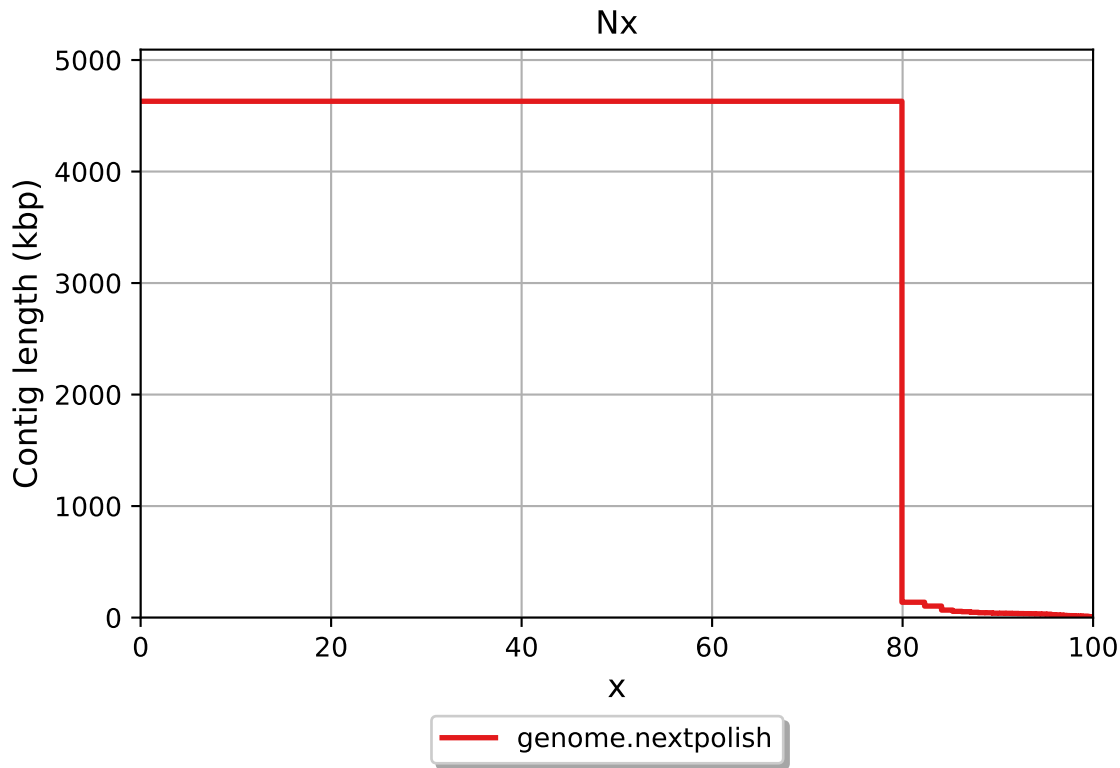


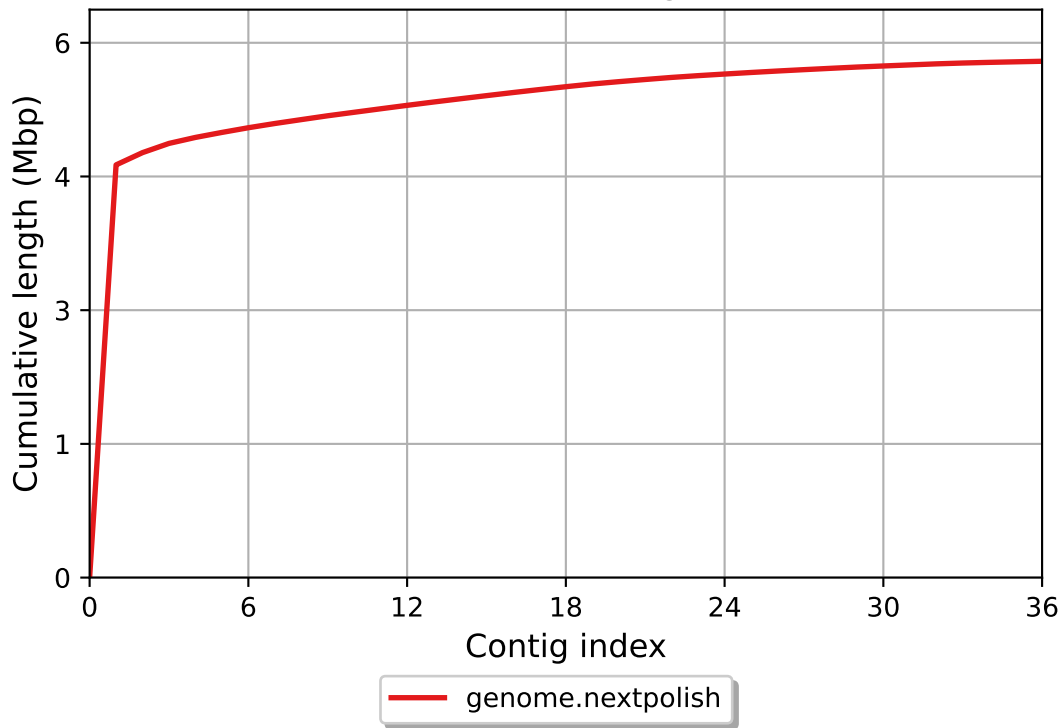
Report

	genome.nextpolish
# contigs (>= 0 bp)	36
# contigs (>= 1000 bp)	36
# contigs (>= 5000 bp)	36
# contigs (>= 10000 bp)	32
# contigs (>= 25000 bp)	20
# contigs (>= 50000 bp)	6
Total length (>= 0 bp)	5792551
Total length (>= 1000 bp)	5792551
Total length (>= 5000 bp)	5792551
Total length (>= 10000 bp)	5763497
Total length (>= 25000 bp)	5563824
Total length (>= 50000 bp)	5046922
# contigs	36
Largest contig	4630132
Total length	5792551
GC (%)	48.60
N50	4630132
N90	39194
auN	3711901.0
L50	1
L90	10
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	28 + 1 part

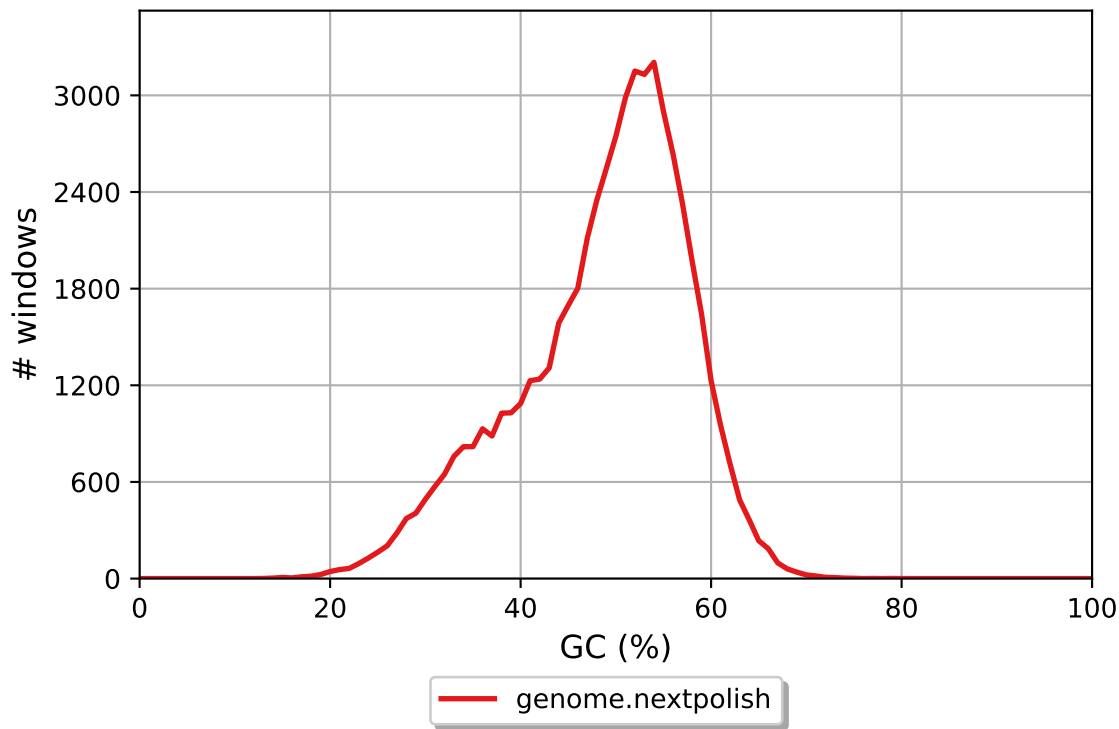
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

