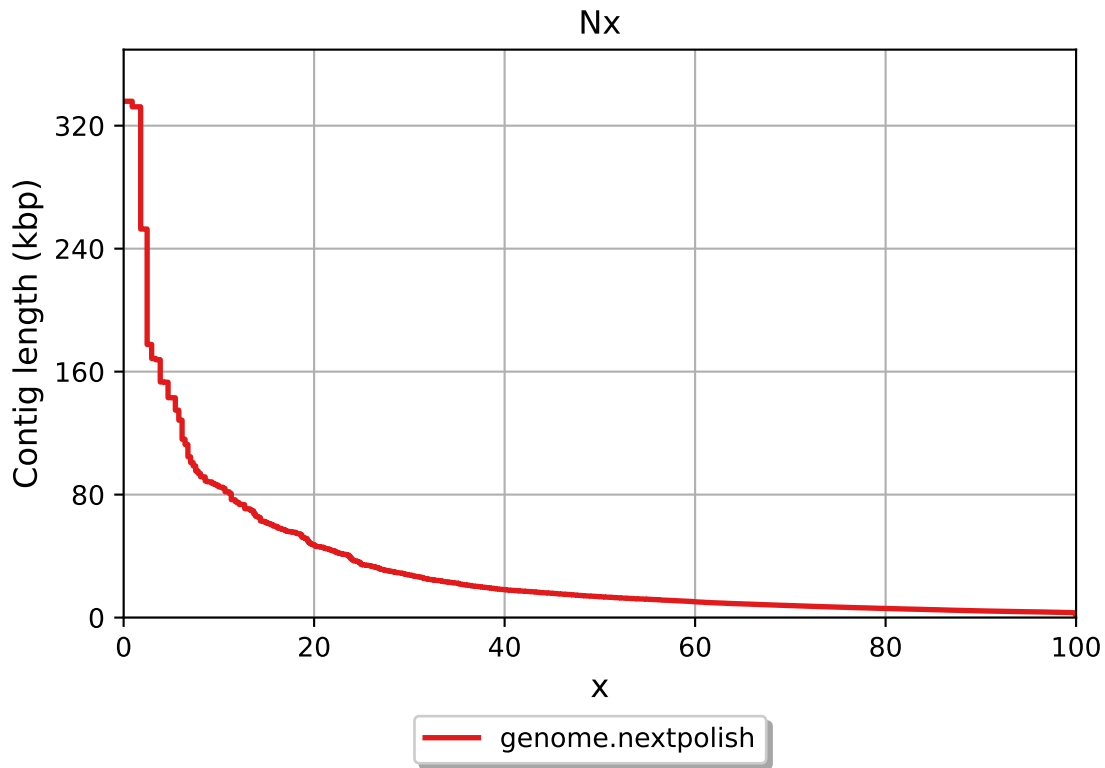


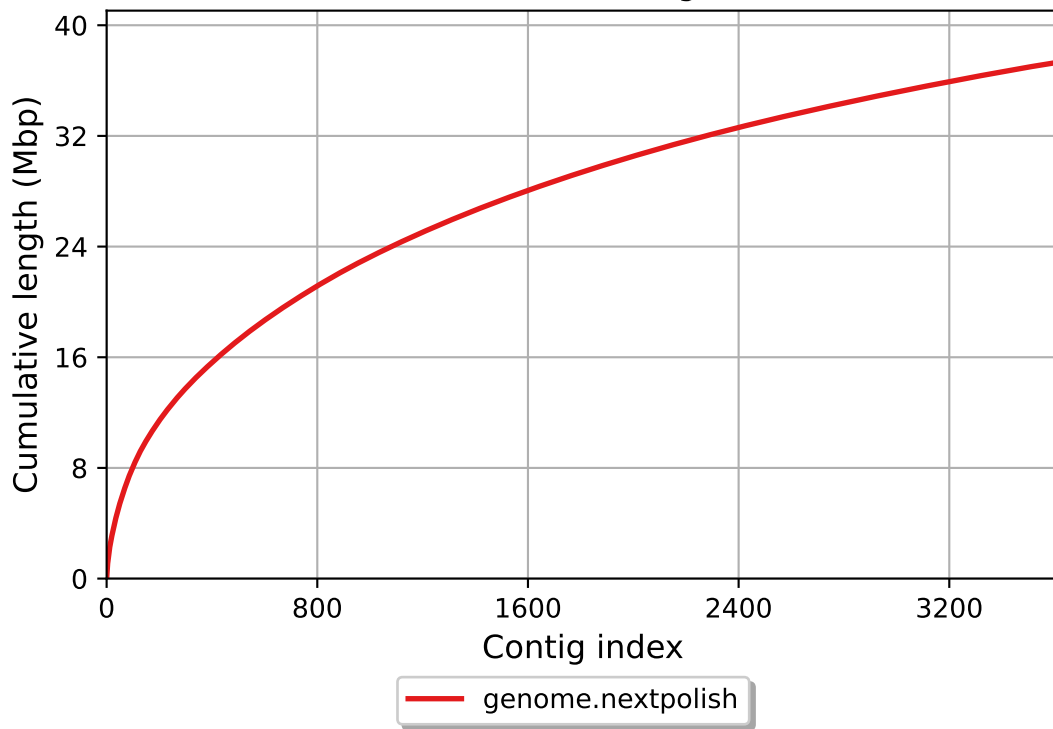
Report

	genome.nextpolish
# contigs (>= 0 bp)	3617
# contigs (>= 1000 bp)	3613
# contigs (>= 5000 bp)	2294
# contigs (>= 10000 bp)	943
# contigs (>= 25000 bp)	219
# contigs (>= 50000 bp)	82
Total length (>= 0 bp)	37323502
Total length (>= 1000 bp)	37320759
Total length (>= 5000 bp)	32095604
Total length (>= 10000 bp)	22690188
Total length (>= 25000 bp)	11917203
Total length (>= 50000 bp)	7242004
# contigs	3617
Largest contig	335877
Total length	37323502
GC (%)	39.86
N50	13581
N90	4353
auN	34621.2
L50	599
L90	2616
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	16 + 5 part

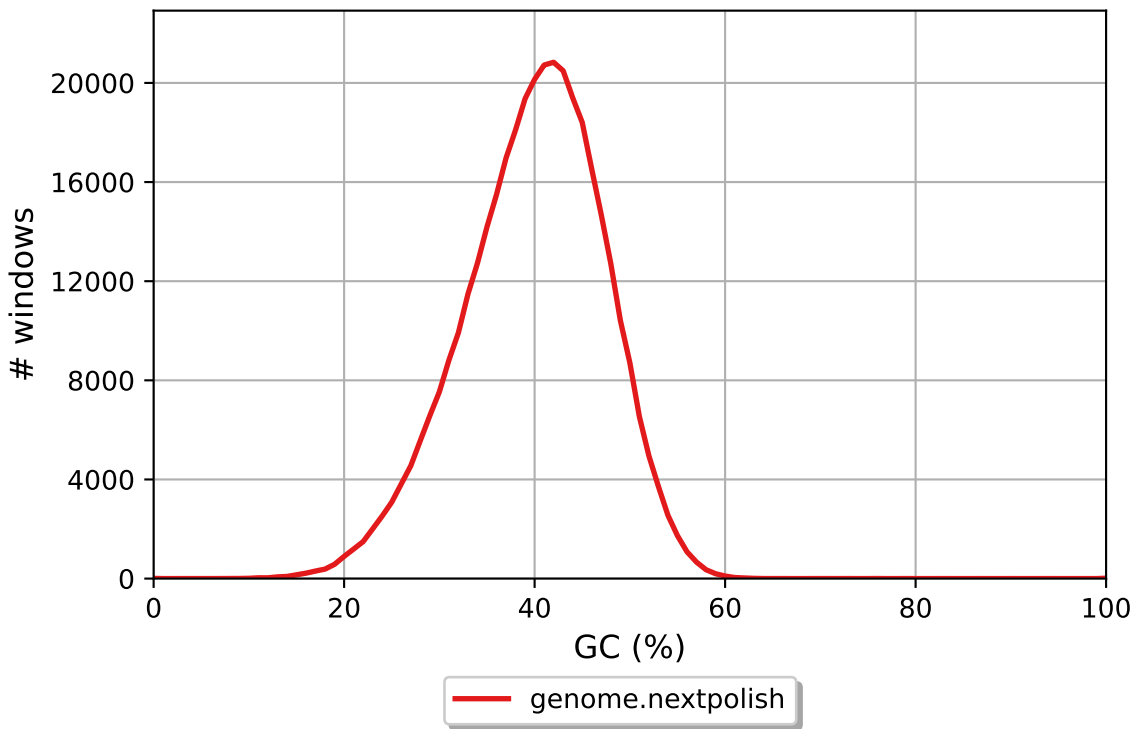
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

