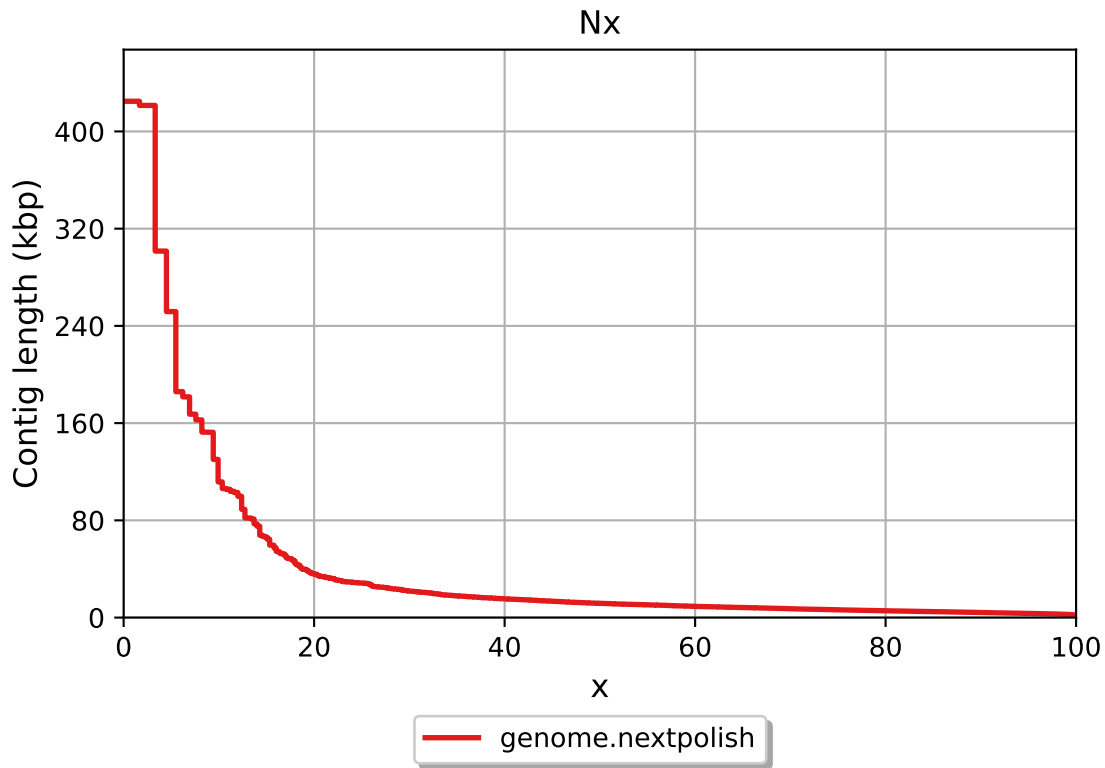


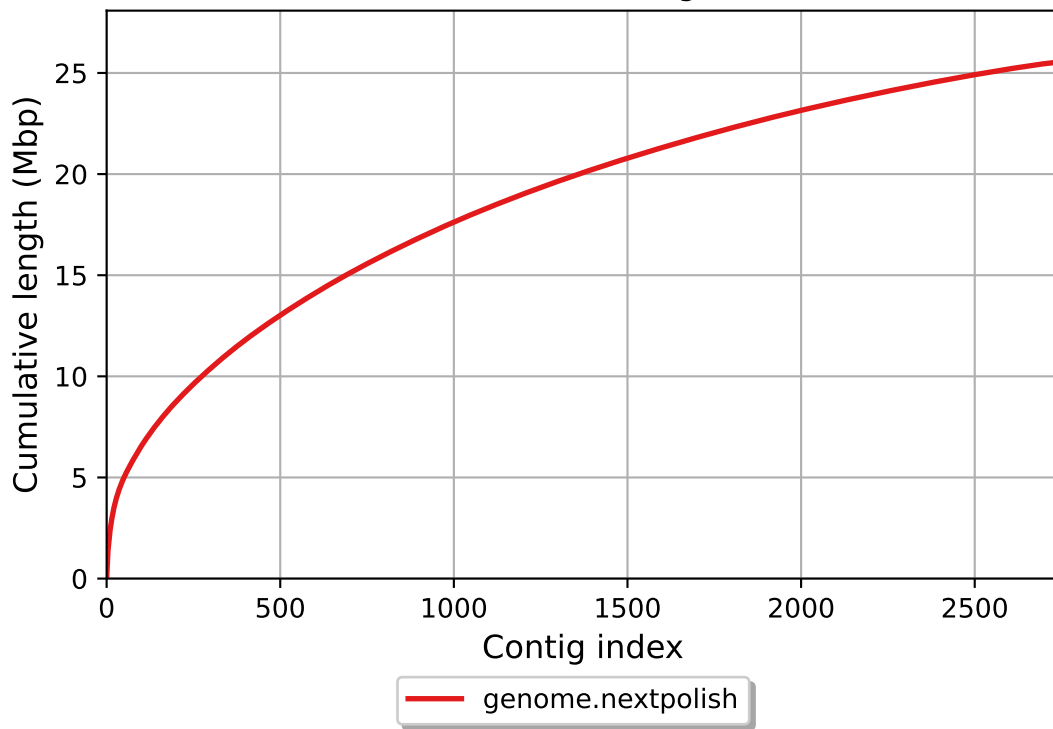
# Report

	genome.nextpolish
# contigs (>= 0 bp)	2743
# contigs (>= 1000 bp)	2732
# contigs (>= 5000 bp)	1637
# contigs (>= 10000 bp)	645
# contigs (>= 25000 bp)	114
# contigs (>= 50000 bp)	35
Total length (>= 0 bp)	25529101
Total length (>= 1000 bp)	25521539
Total length (>= 5000 bp)	21499847
Total length (>= 10000 bp)	14572336
Total length (>= 25000 bp)	6917037
Total length (>= 50000 bp)	4355093
# contigs	2743
Largest contig	424814
Total length	25529101
GC (%)	42.90
N50	11694
N90	4152
auN	43689.8
L50	478
L90	1958
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	51 + 9 part

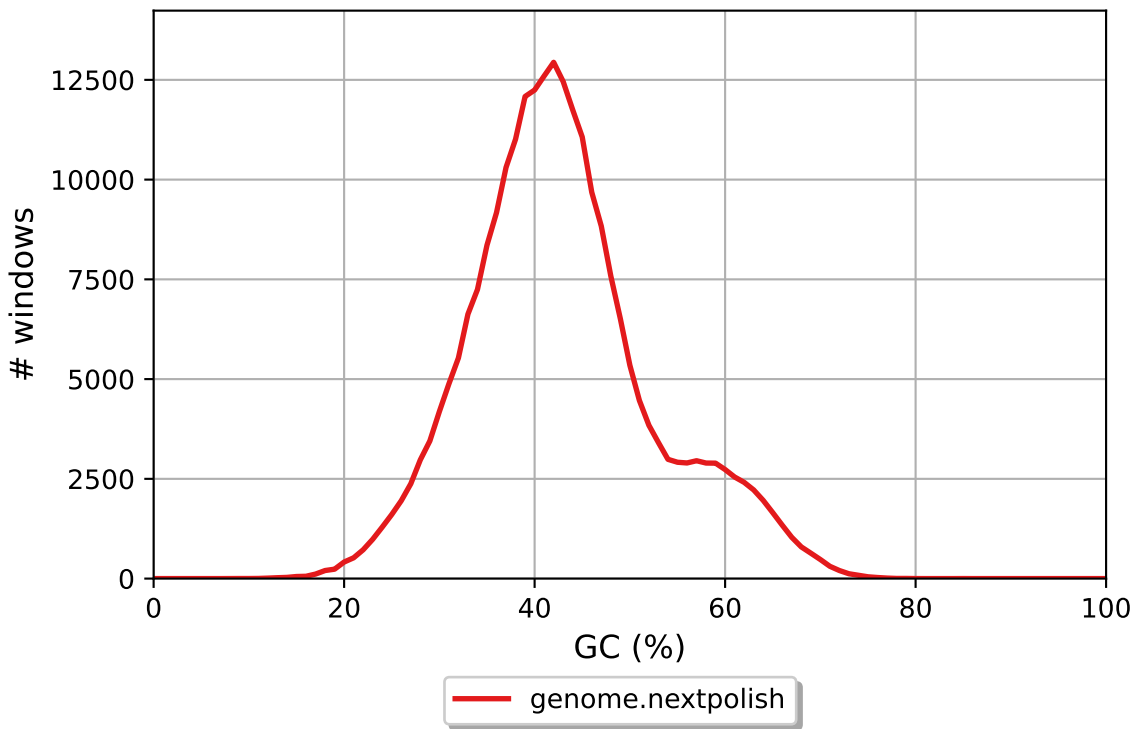
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



## GC content



genome.nextpolish GC content

