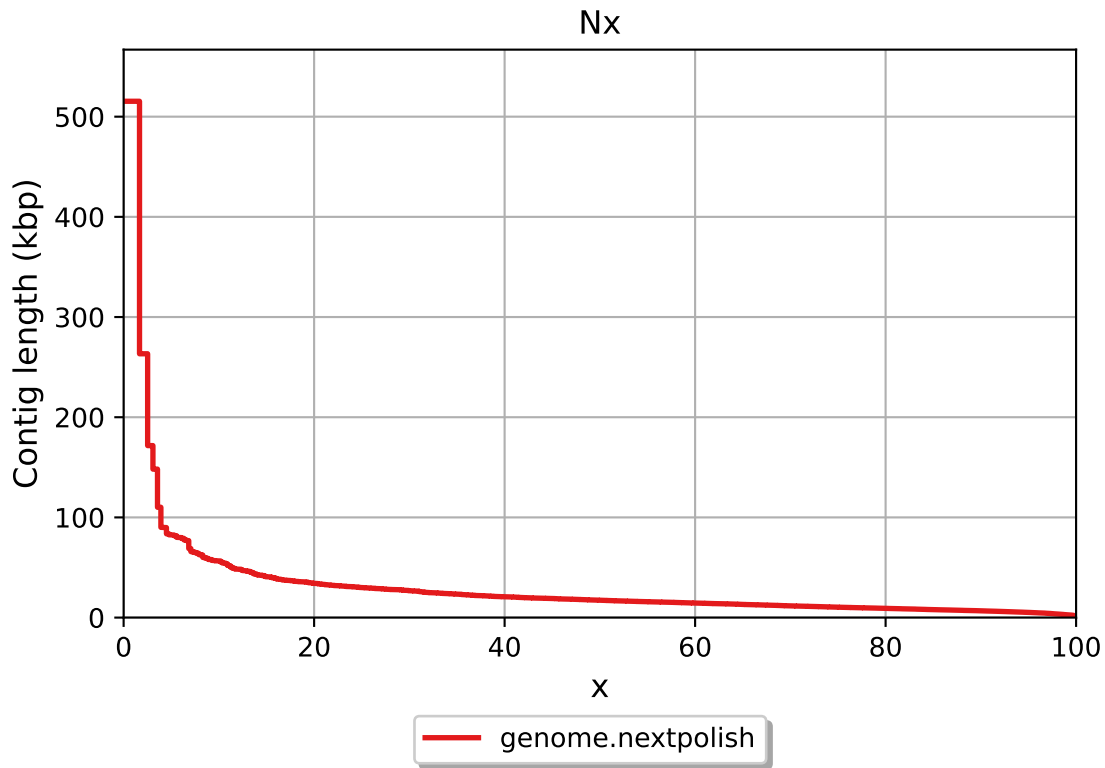


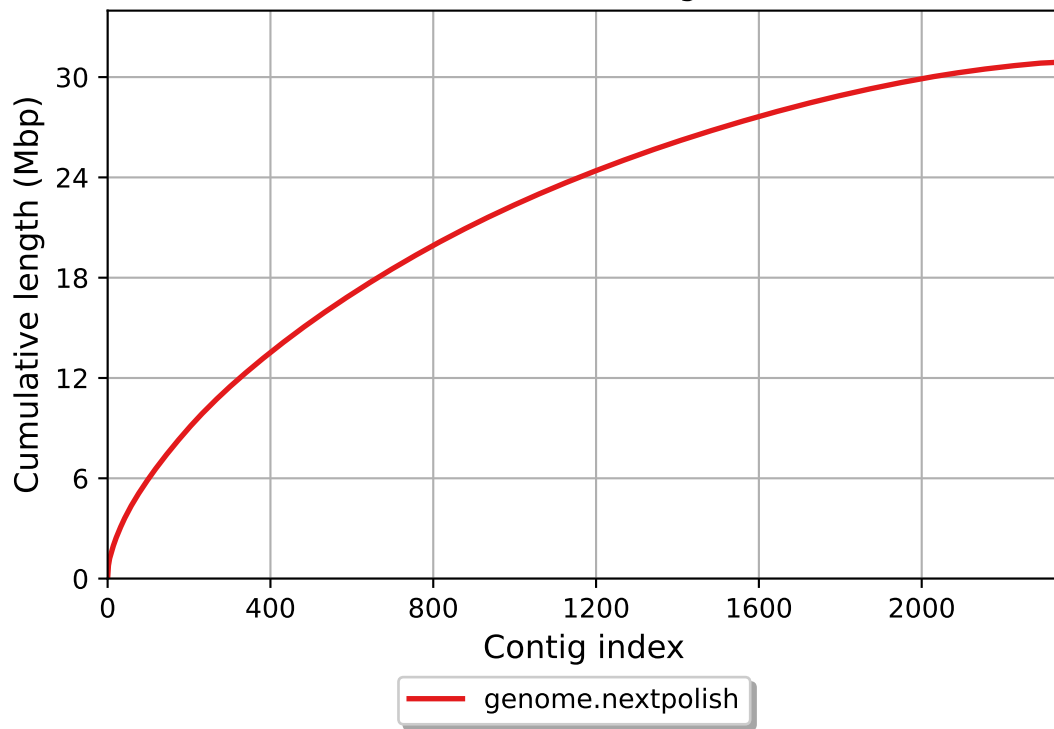
Report

	genome.nextpolish
# contigs (>= 0 bp)	2340
# contigs (>= 1000 bp)	2321
# contigs (>= 5000 bp)	1906
# contigs (>= 10000 bp)	1120
# contigs (>= 25000 bp)	231
# contigs (>= 50000 bp)	40
Total length (>= 0 bp)	30887344
Total length (>= 1000 bp)	30874728
Total length (>= 5000 bp)	29464283
Total length (>= 10000 bp)	23626589
Total length (>= 25000 bp)	9841873
Total length (>= 50000 bp)	3517369
# contigs	2340
Largest contig	515374
Total length	30887344
GC (%)	45.86
N50	17346
N90	6746
auN	33991.8
L50	505
L90	1623
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	67 + 12 part

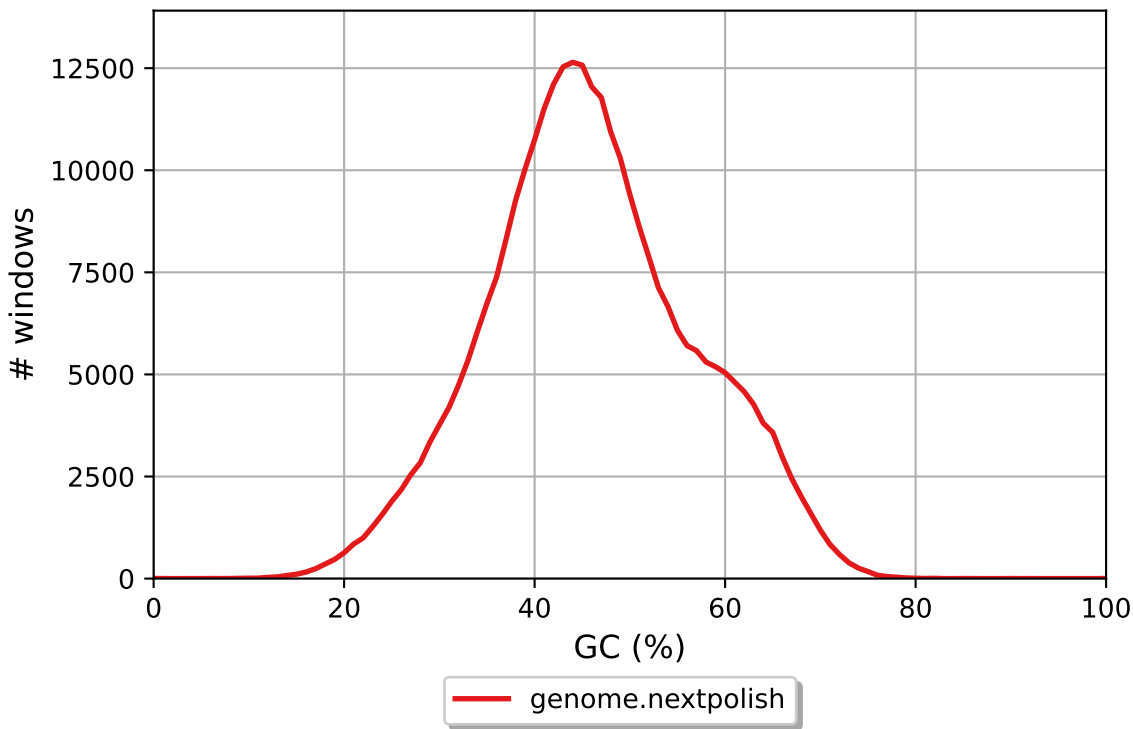
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

