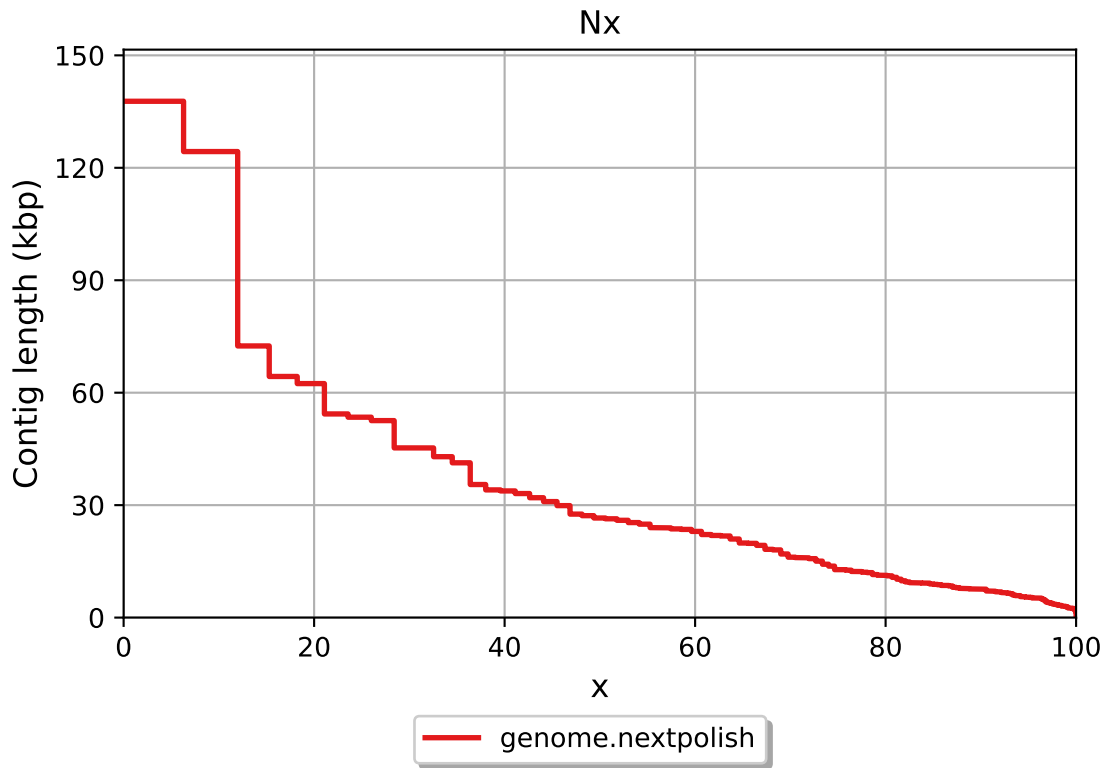


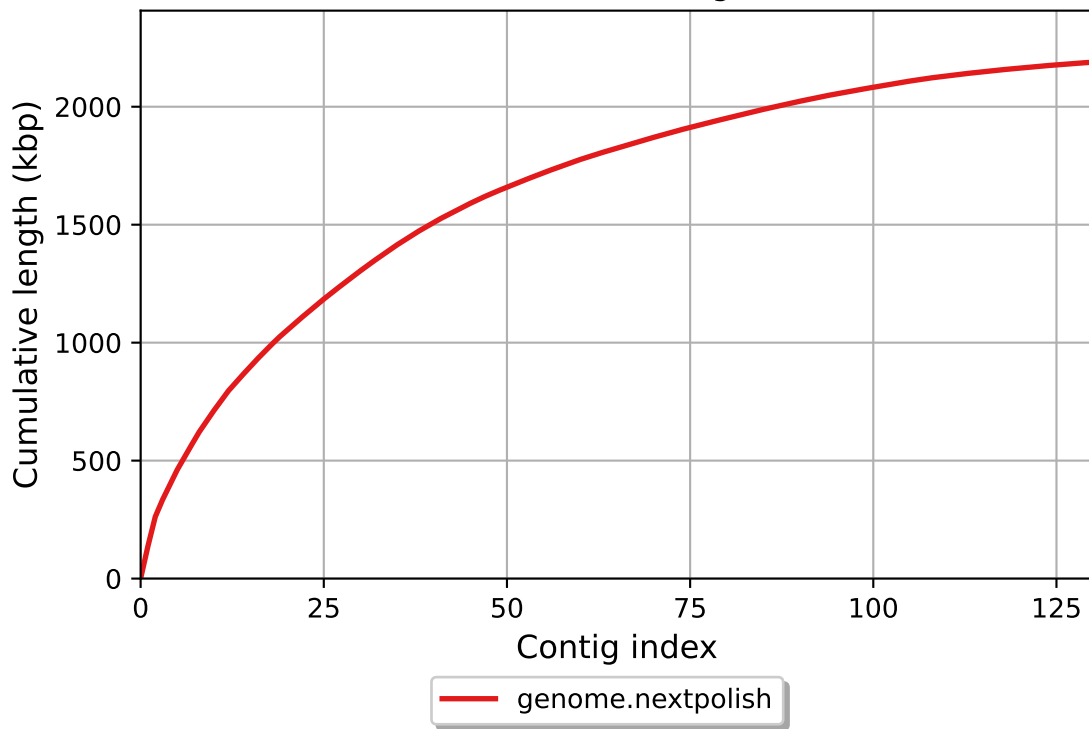
Report

	genome.nextpolish
# contigs (>= 0 bp)	130
# contigs (>= 1000 bp)	130
# contigs (>= 5000 bp)	105
# contigs (>= 10000 bp)	61
# contigs (>= 25000 bp)	25
# contigs (>= 50000 bp)	8
Total length (>= 0 bp)	2188446
Total length (>= 1000 bp)	2188446
Total length (>= 5000 bp)	2108838
Total length (>= 10000 bp)	1786597
Total length (>= 25000 bp)	1184687
Total length (>= 50000 bp)	621655
# contigs	130
Largest contig	137749
Total length	2188446
GC (%)	35.01
N50	26568
N90	7576
auN	40782.2
L50	22
L90	83
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	5 + 2 part

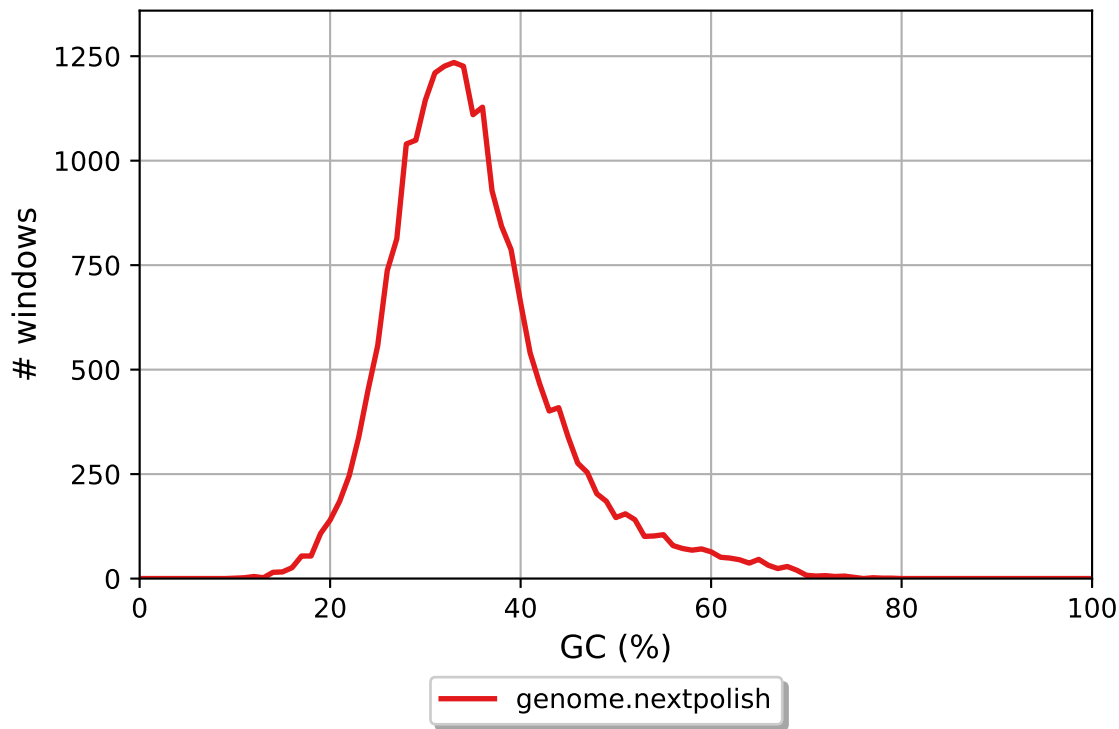
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

