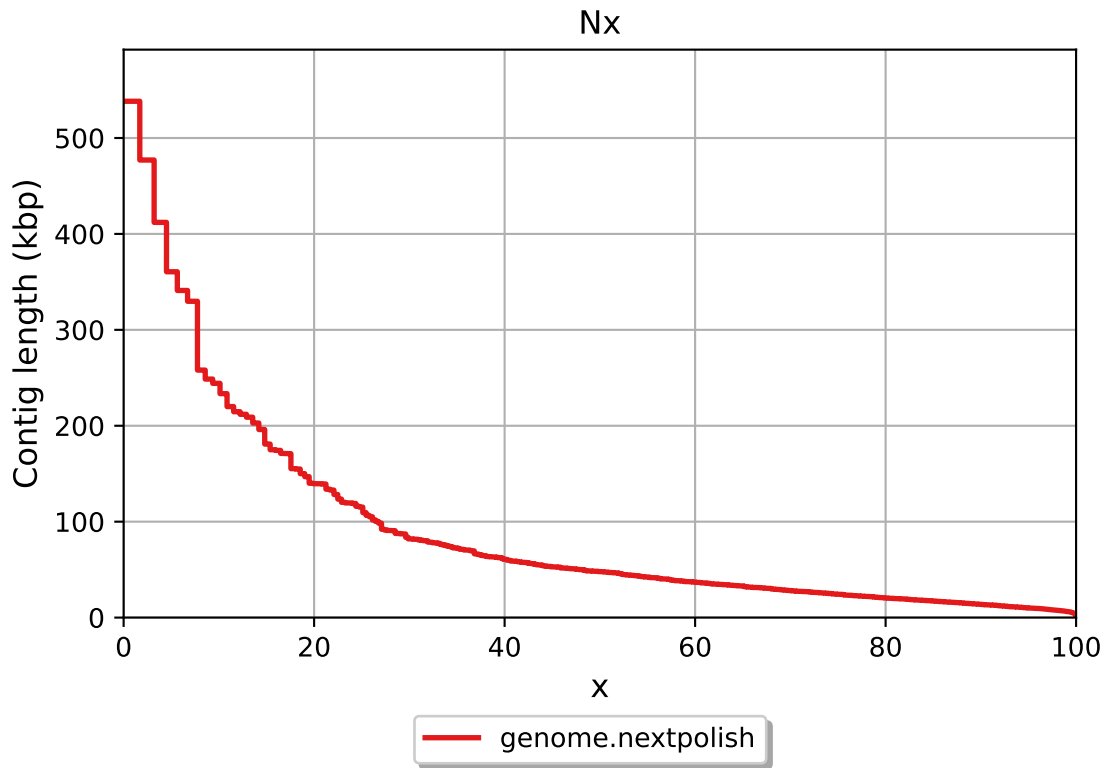


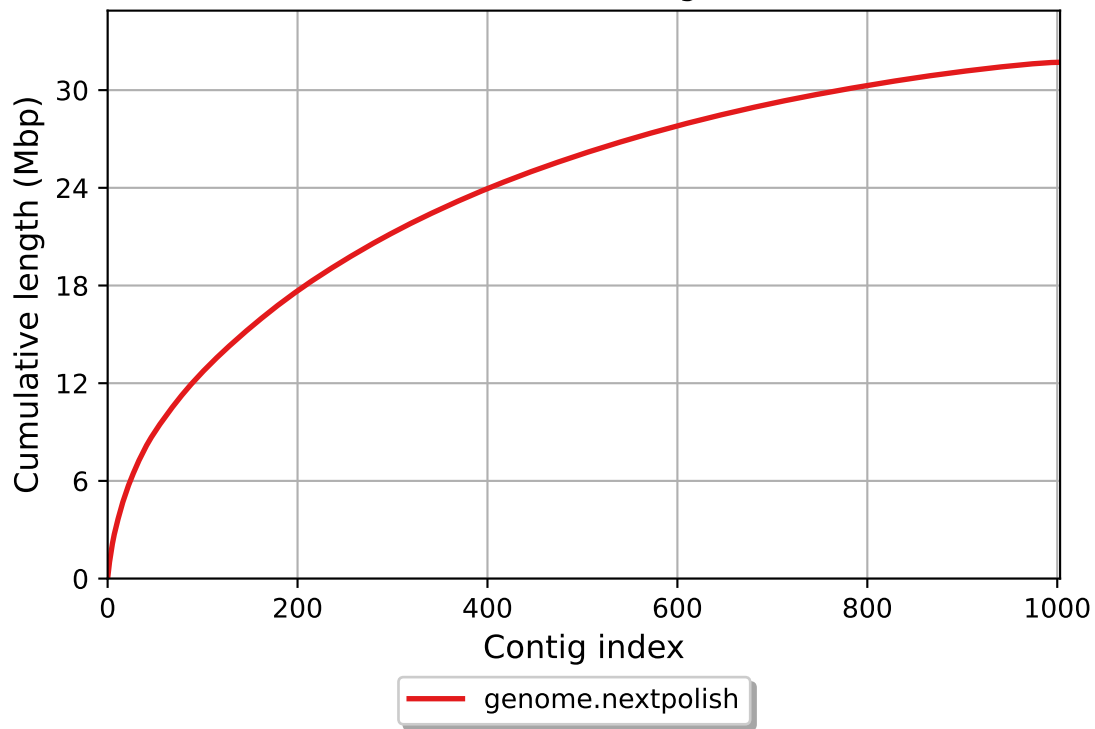
Report

	genome.nextpolish
# contigs (>= 0 bp)	1003
# contigs (>= 1000 bp)	1001
# contigs (>= 5000 bp)	974
# contigs (>= 10000 bp)	784
# contigs (>= 25000 bp)	383
# contigs (>= 50000 bp)	146
Total length (>= 0 bp)	31711460
Total length (>= 1000 bp)	31710017
Total length (>= 5000 bp)	31617209
Total length (>= 10000 bp)	30127033
Total length (>= 25000 bp)	23539216
Total length (>= 50000 bp)	15206148
# contigs	1003
Largest contig	538300
Total length	31711460
GC (%)	50.33
N50	47974
N90	13767
auN	93572.2
L50	160
L90	651
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	60 + 8 part

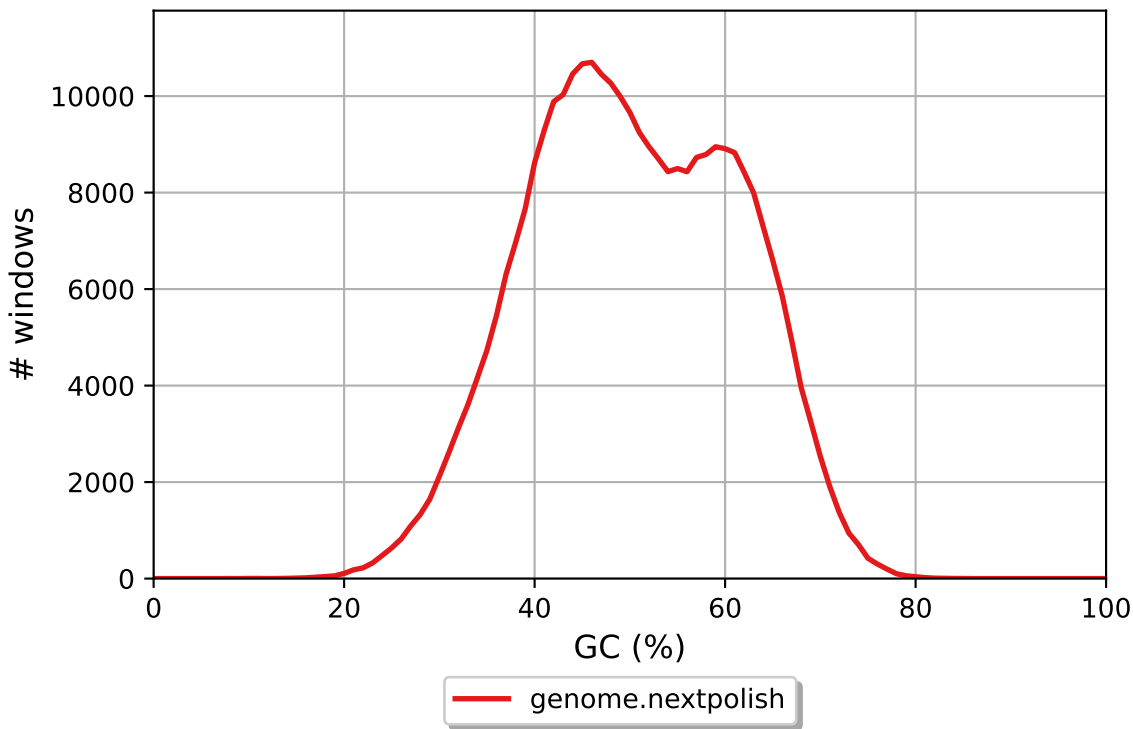
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

