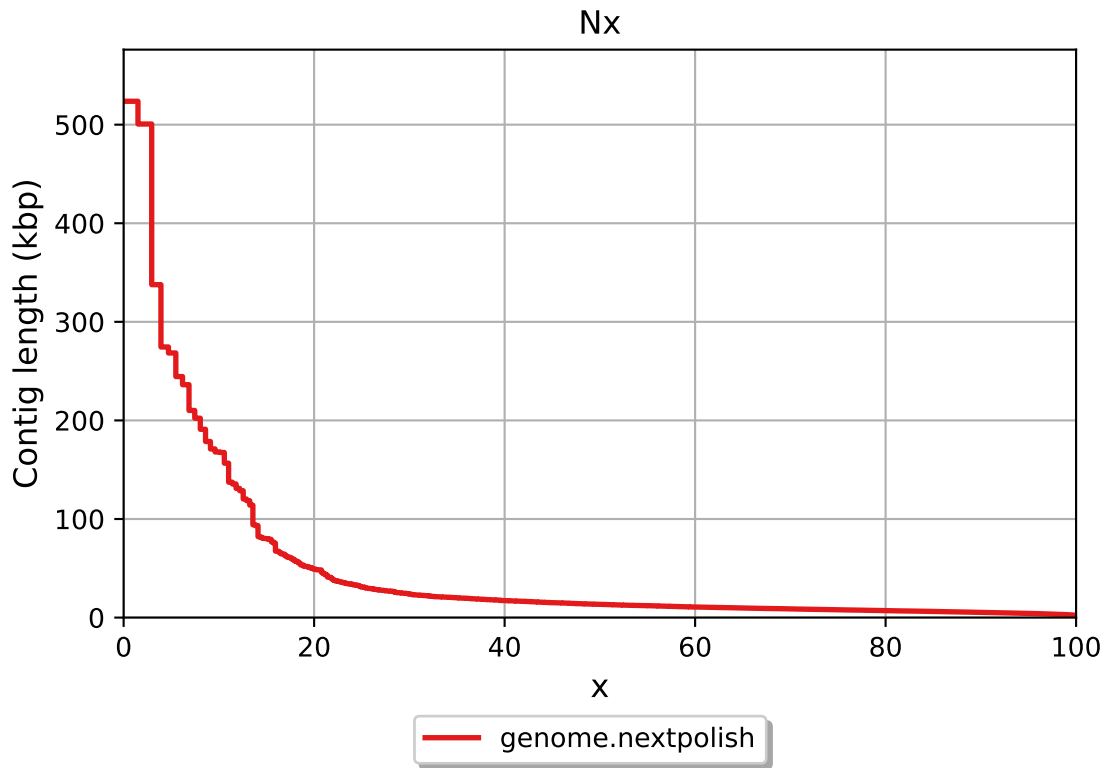


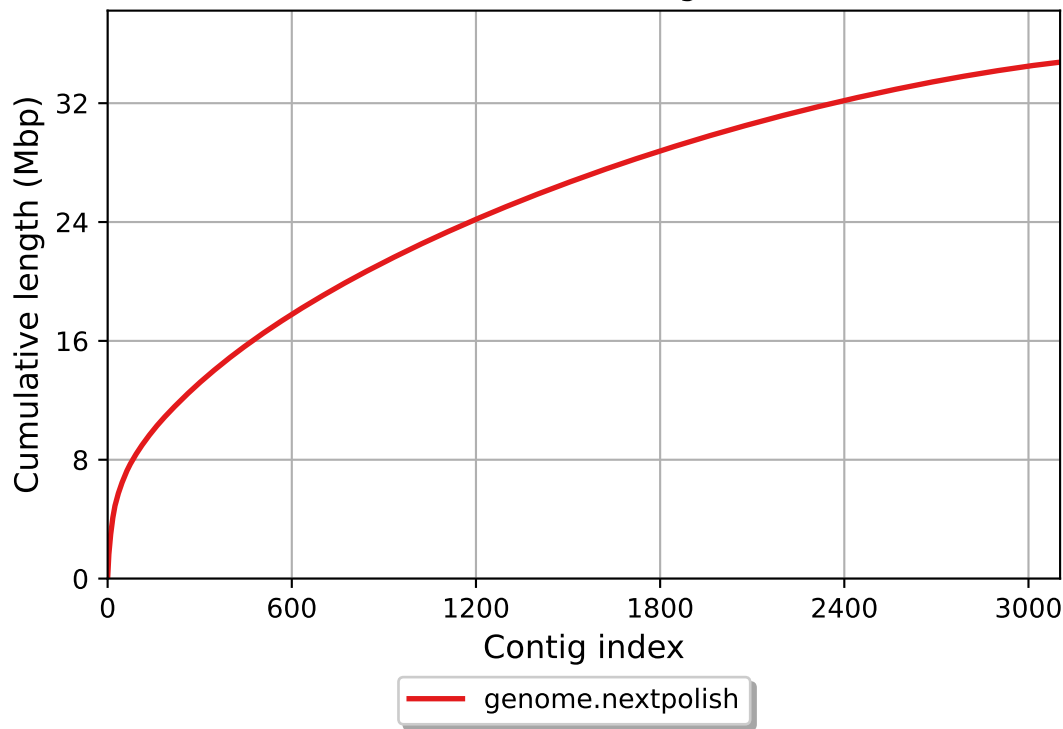
# Report

	genome.nextpolish
# contigs (>= 0 bp)	3103
# contigs (>= 1000 bp)	3100
# contigs (>= 5000 bp)	2310
# contigs (>= 10000 bp)	990
# contigs (>= 25000 bp)	154
# contigs (>= 50000 bp)	55
Total length (>= 0 bp)	34754551
Total length (>= 1000 bp)	34752580
Total length (>= 5000 bp)	31740174
Total length (>= 10000 bp)	22211576
Total length (>= 25000 bp)	10125073
Total length (>= 50000 bp)	6874583
# contigs	3103
Largest contig	523730
Total length	34754551
GC (%)	54.23
N50	13430
N90	5268
auN	51938.5
L50	570
L90	2221
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	73 + 11 part

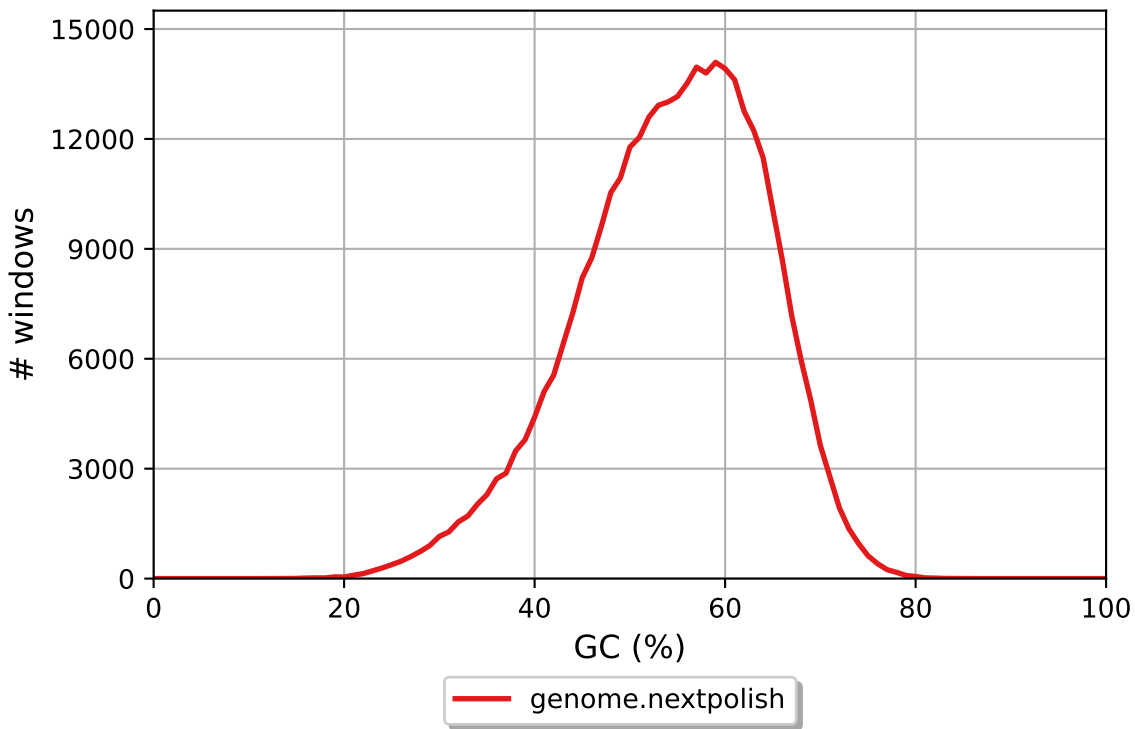
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

