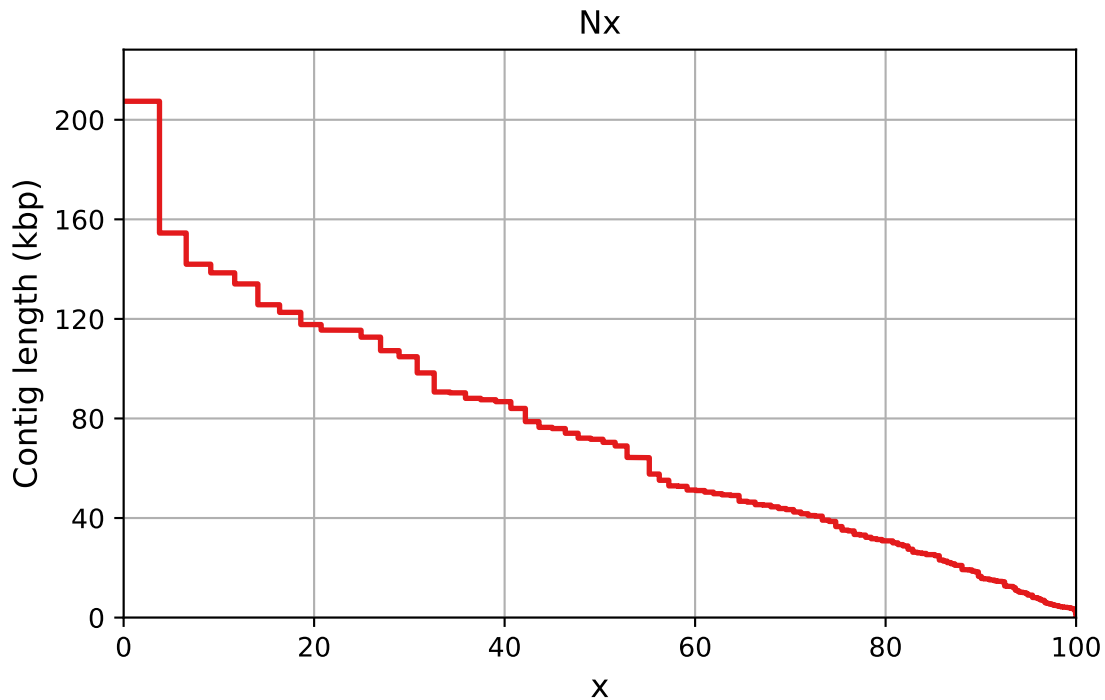


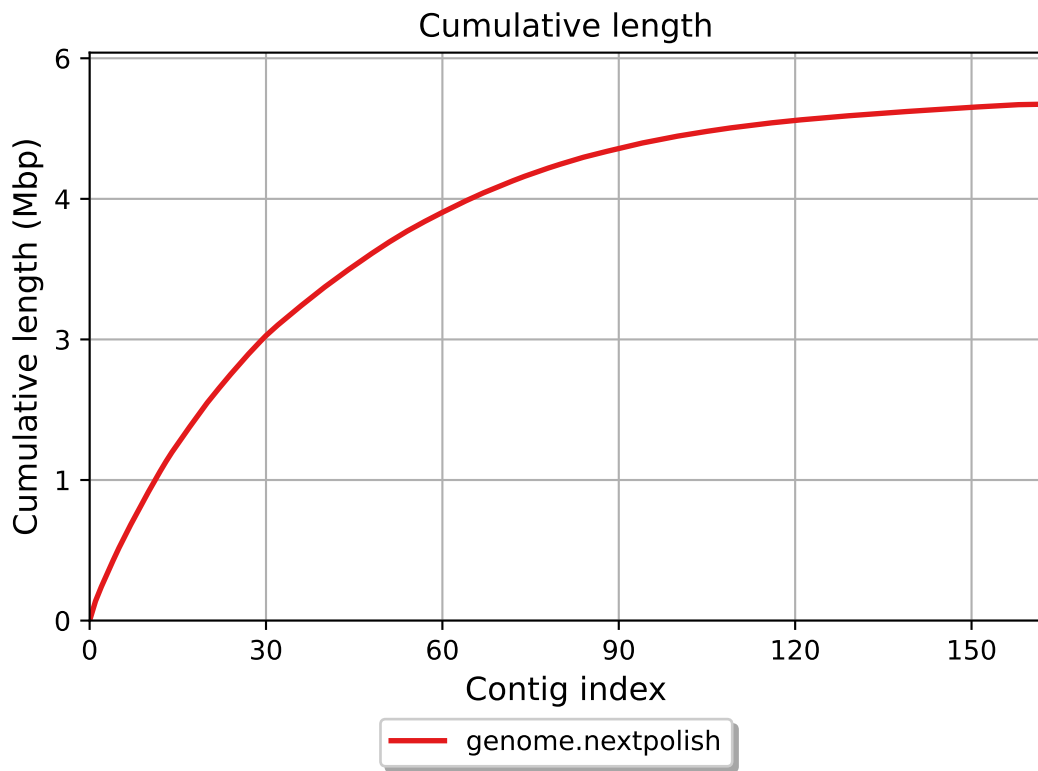
Report

	genome.nextpolish
# contigs (>= 0 bp)	162
# contigs (>= 1000 bp)	159
# contigs (>= 5000 bp)	129
# contigs (>= 10000 bp)	104
# contigs (>= 25000 bp)	72
# contigs (>= 50000 bp)	37
Total length (>= 0 bp)	5509214
Total length (>= 1000 bp)	5507045
Total length (>= 5000 bp)	5386137
Total length (>= 10000 bp)	5210233
Total length (>= 25000 bp)	4693029
Total length (>= 50000 bp)	3411725
# contigs	162
Largest contig	207436
Total length	5509214
GC (%)	57.08
N50	71634
N90	16572
auN	75841.7
L50	26
L90	85
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	15 + 3 part

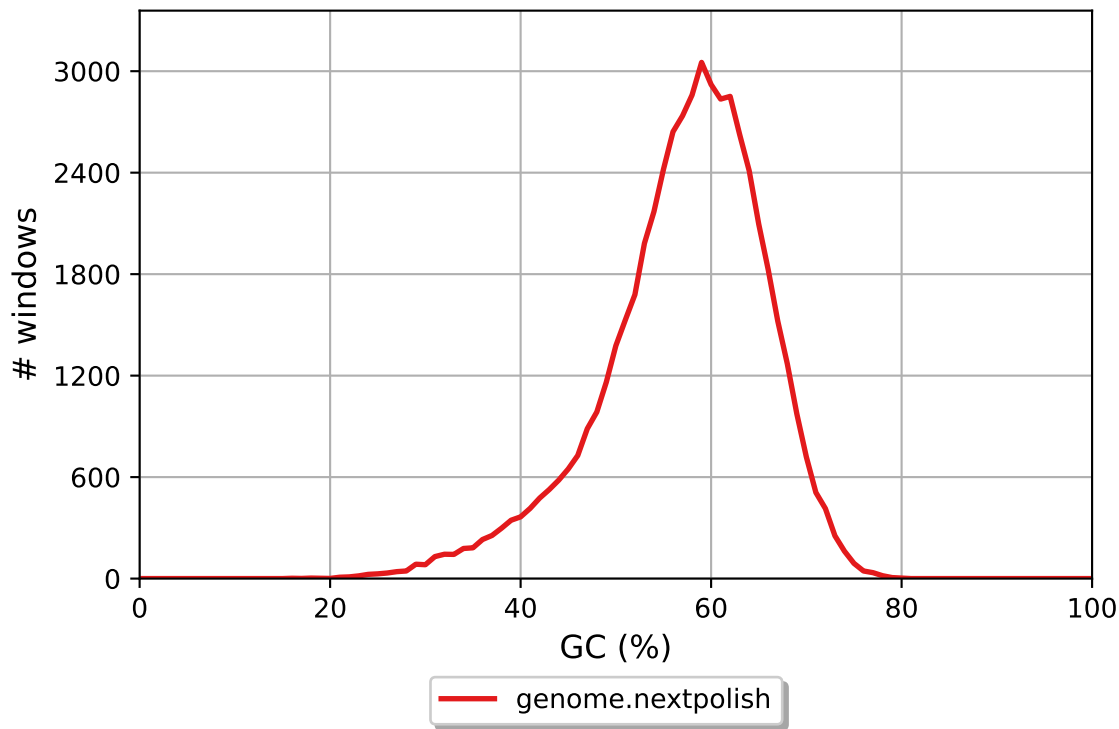
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— genome.nextpolish



GC content



genome.nextpolish GC content

