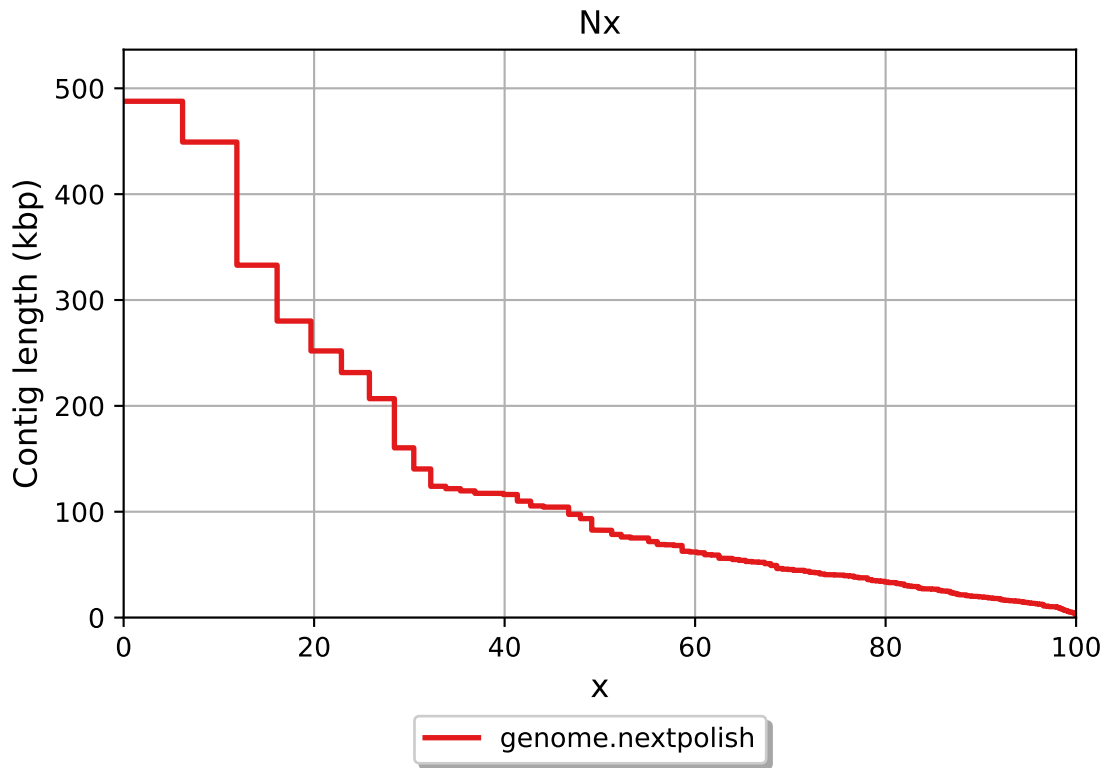


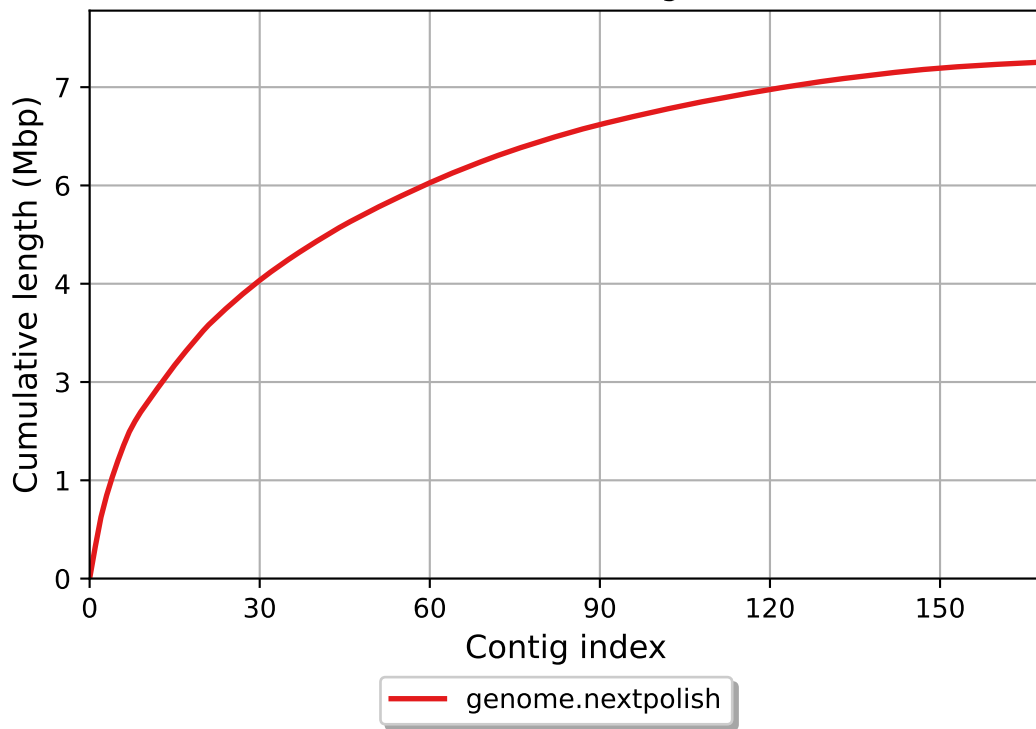
Report

	genome.nextpolish
# contigs (>= 0 bp)	168
# contigs (>= 1000 bp)	168
# contigs (>= 5000 bp)	159
# contigs (>= 10000 bp)	142
# contigs (>= 25000 bp)	85
# contigs (>= 50000 bp)	44
Total length (>= 0 bp)	7879973
Total length (>= 1000 bp)	7879973
Total length (>= 5000 bp)	7843232
Total length (>= 10000 bp)	7725058
Total length (>= 25000 bp)	6814820
Total length (>= 50000 bp)	5354245
# contigs	168
Largest contig	487751
Total length	7879973
GC (%)	57.95
N50	82656
N90	19662
auN	146104.7
L50	22
L90	98
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	20 + 2 part

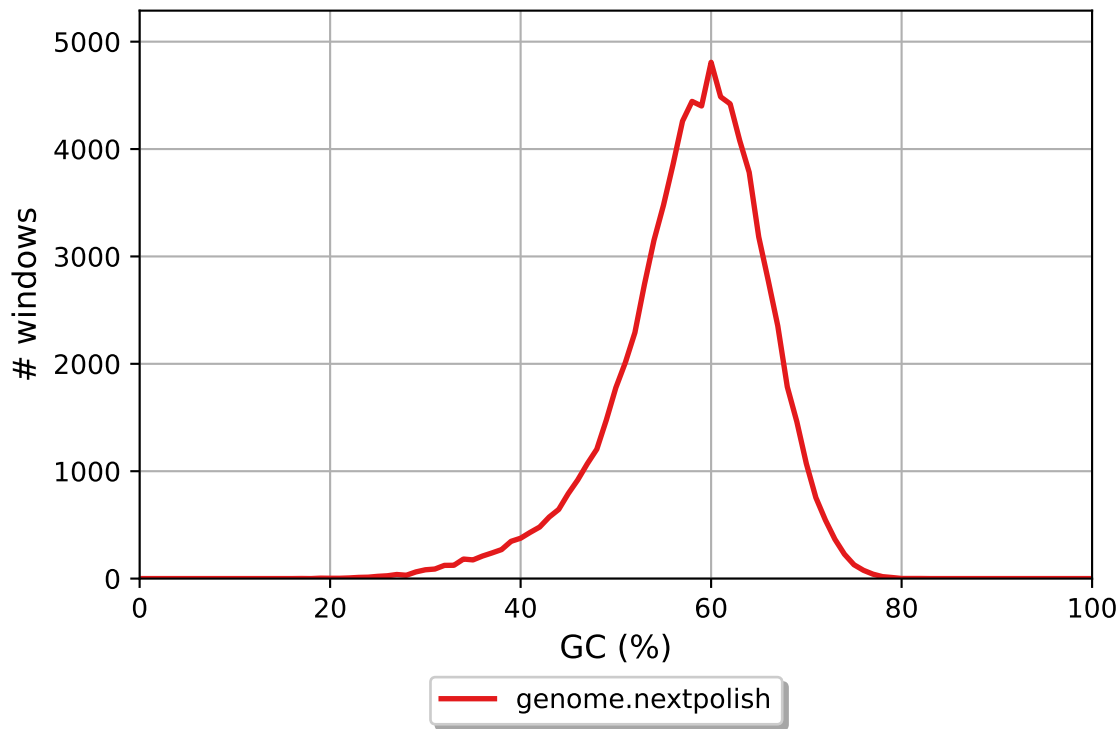
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

