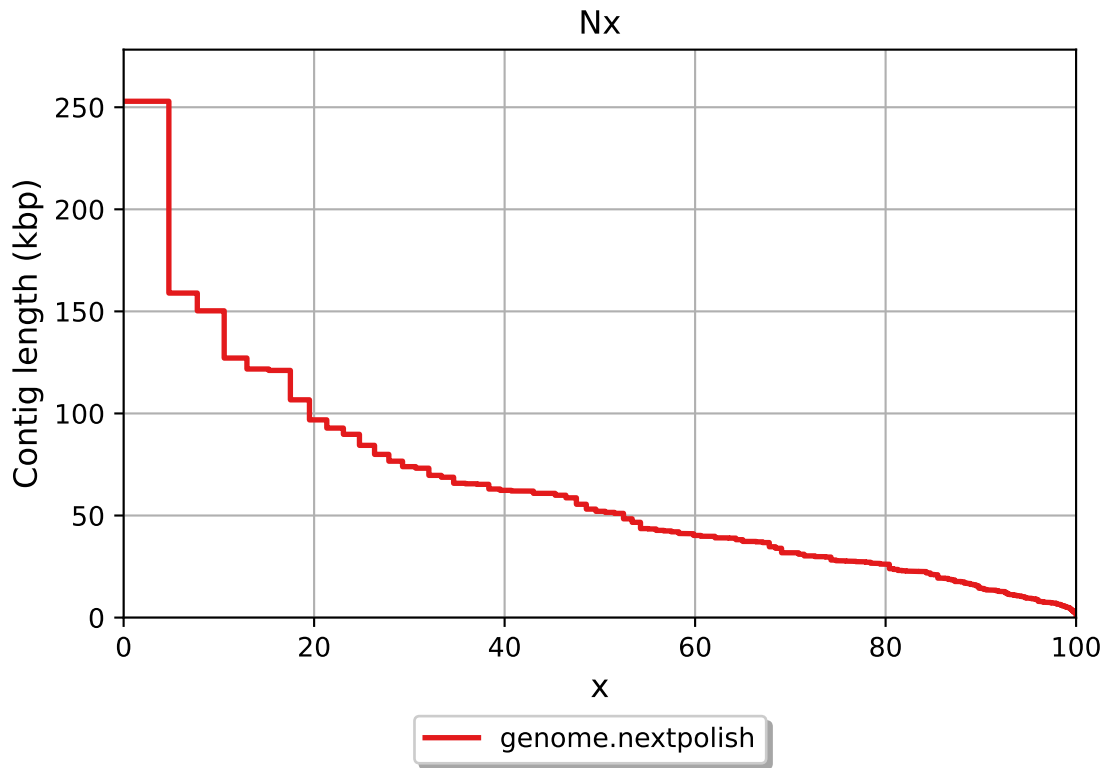


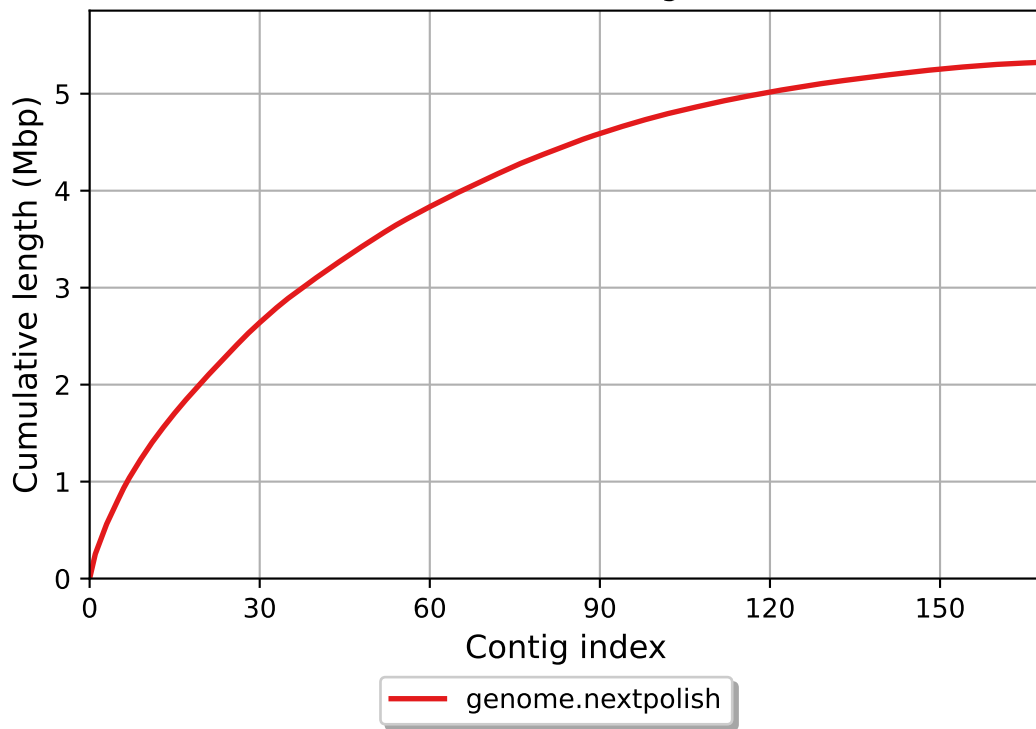
# Report

	genome.nextpolish
# contigs (>= 0 bp)	168
# contigs (>= 1000 bp)	168
# contigs (>= 5000 bp)	154
# contigs (>= 10000 bp)	122
# contigs (>= 25000 bp)	76
# contigs (>= 50000 bp)	33
Total length (>= 0 bp)	5324276
Total length (>= 1000 bp)	5324276
Total length (>= 5000 bp)	5275324
Total length (>= 10000 bp)	5037876
Total length (>= 25000 bp)	4280384
Total length (>= 50000 bp)	2794261
# contigs	168
Largest contig	252926
Total length	5324276
GC (%)	57.49
N50	52066
N90	14399
auN	67601.1
L50	31
L90	102
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	11 + 0 part

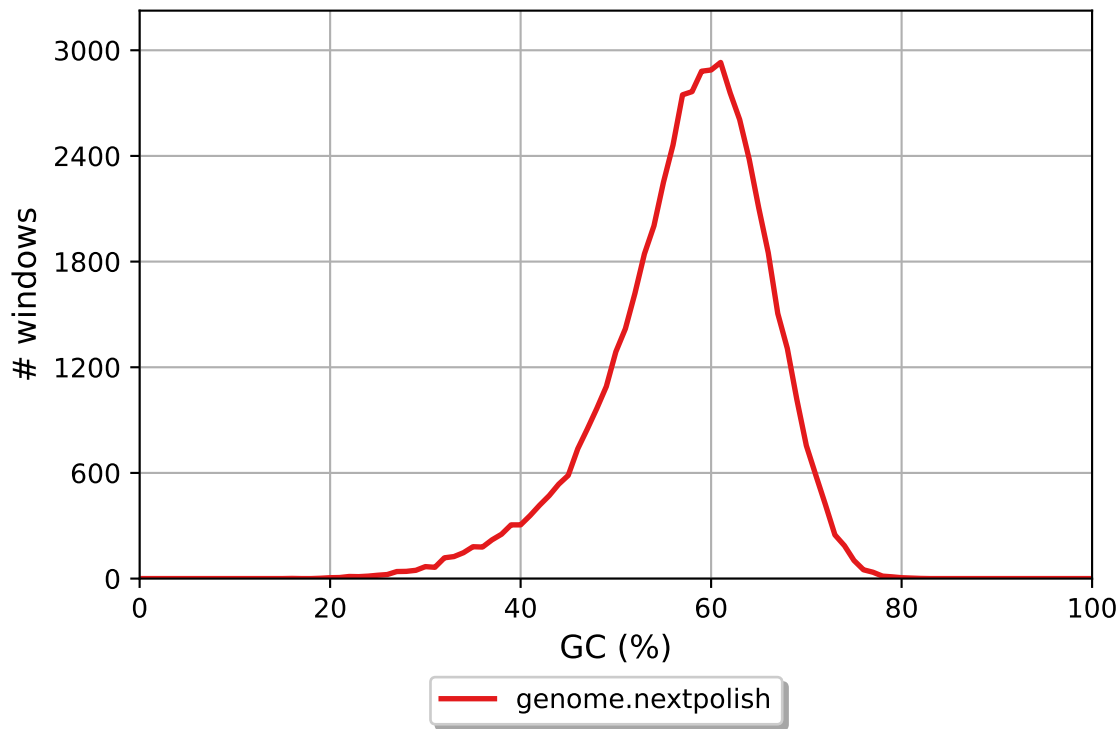
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



## GC content



genome.nextpolish GC content

