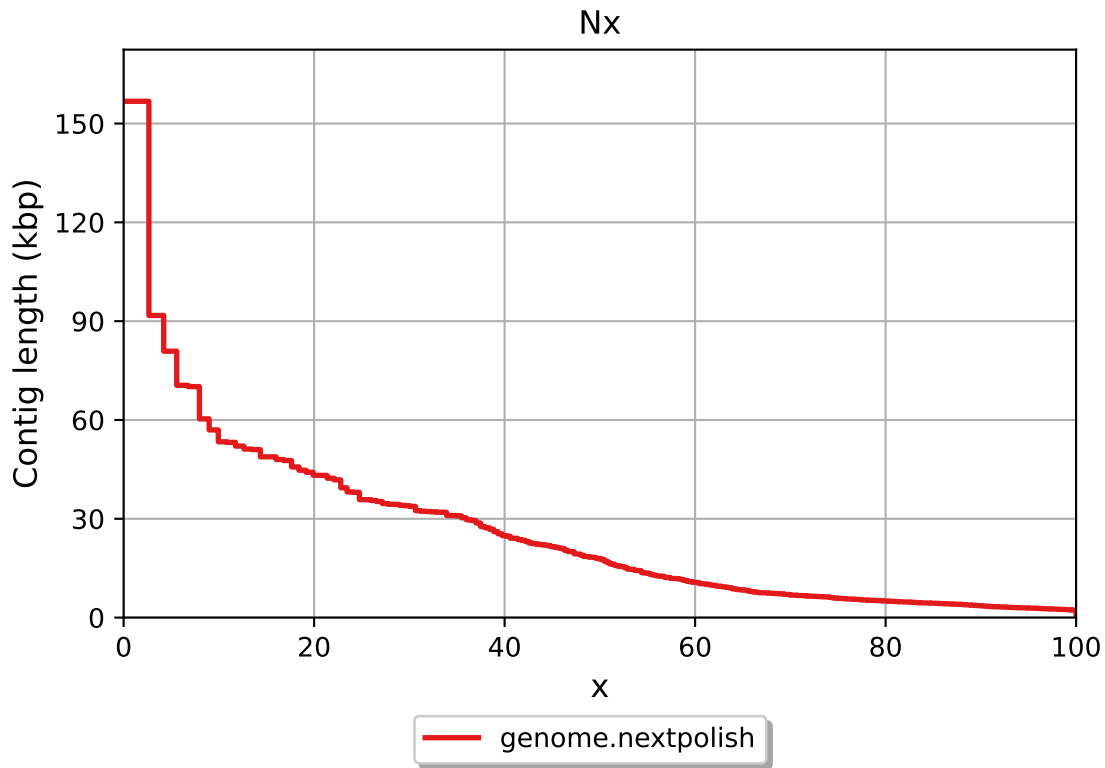


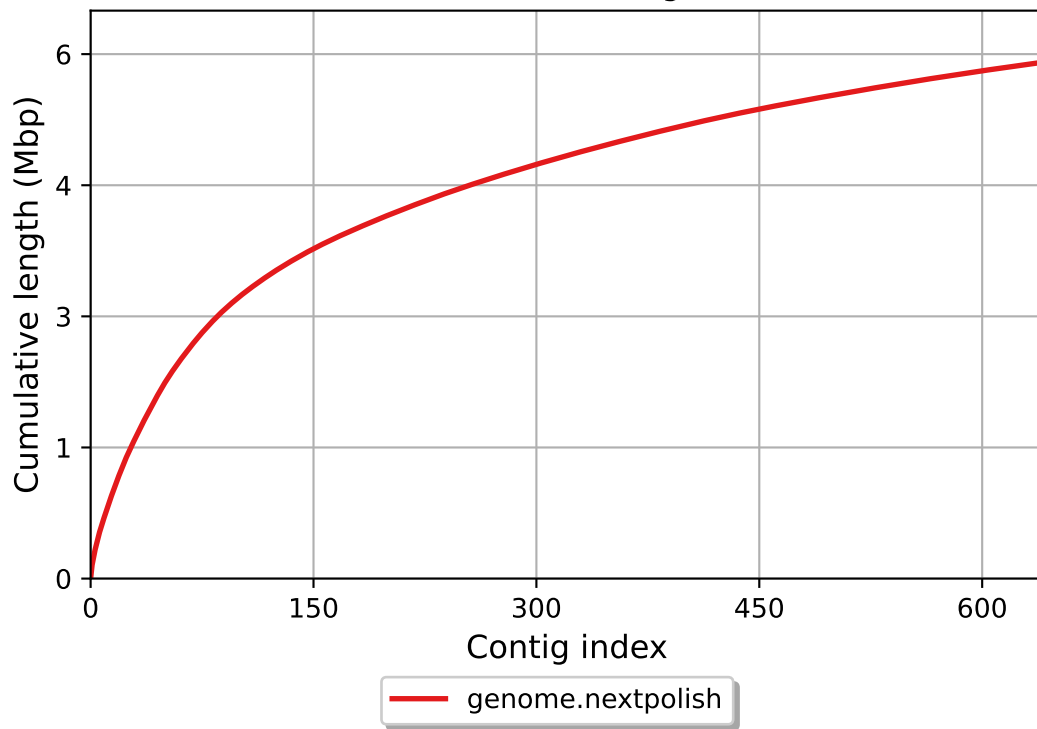
# Report

	genome.nextpolish
# contigs (>= 0 bp)	641
# contigs (>= 1000 bp)	641
# contigs (>= 5000 bp)	301
# contigs (>= 10000 bp)	135
# contigs (>= 25000 bp)	54
# contigs (>= 50000 bp)	12
Total length (>= 0 bp)	5906826
Total length (>= 1000 bp)	5906826
Total length (>= 5000 bp)	4743915
Total length (>= 10000 bp)	3631706
Total length (>= 25000 bp)	2347032
Total length (>= 50000 bp)	848129
# contigs	641
Largest contig	156762
Total length	5906826
GC (%)	46.78
N50	17823
N90	3586
auN	26947.0
L50	83
L90	435
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	7 + 1 part

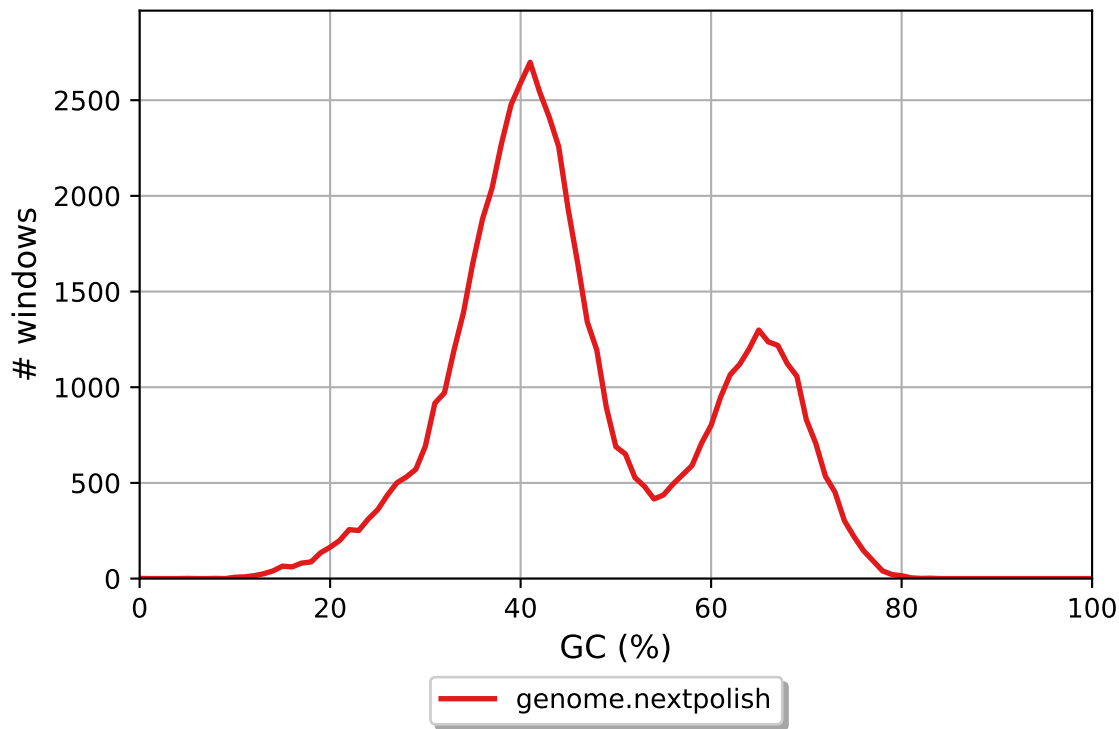
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

