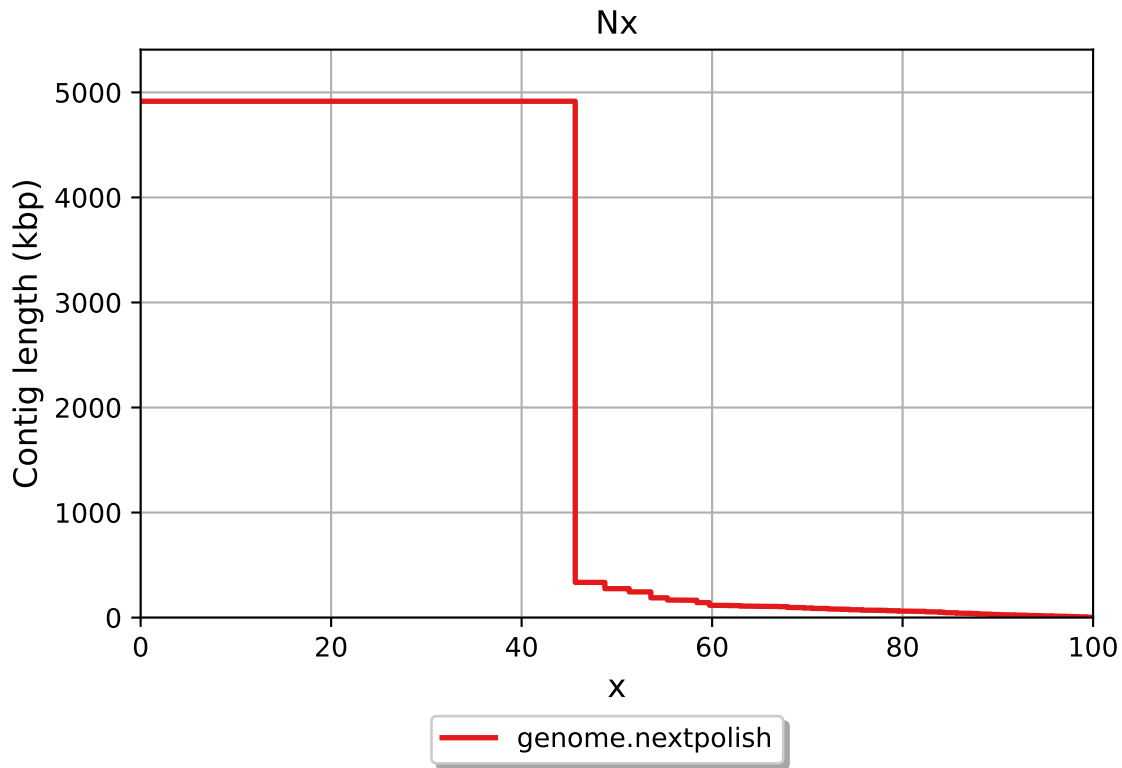


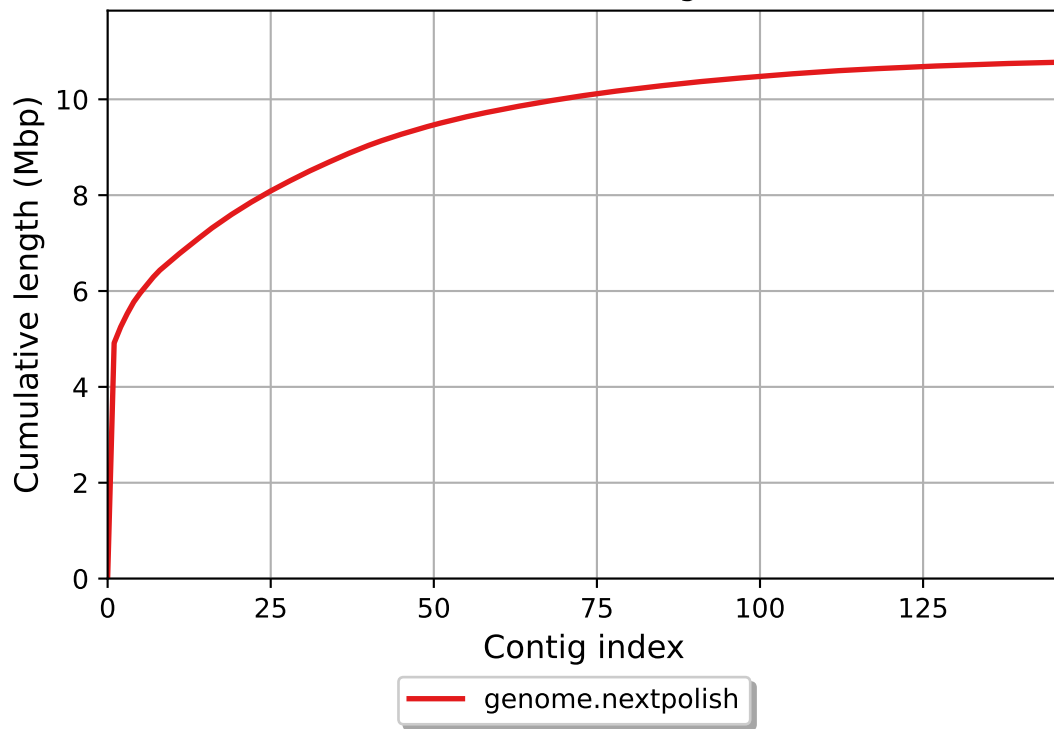
Report

	genome.nextpolish
# contigs (>= 0 bp)	146
# contigs (>= 1000 bp)	146
# contigs (>= 5000 bp)	130
# contigs (>= 10000 bp)	104
# contigs (>= 25000 bp)	63
# contigs (>= 50000 bp)	41
Total length (>= 0 bp)	10772753
Total length (>= 1000 bp)	10772753
Total length (>= 5000 bp)	10710369
Total length (>= 10000 bp)	10522195
Total length (>= 25000 bp)	9854786
Total length (>= 50000 bp)	9089804
# contigs	146
Largest contig	4915547
Total length	10772753
GC (%)	53.84
N50	275430
N90	28266
auN	2301172.4
L50	3
L90	58
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	31 + 0 part

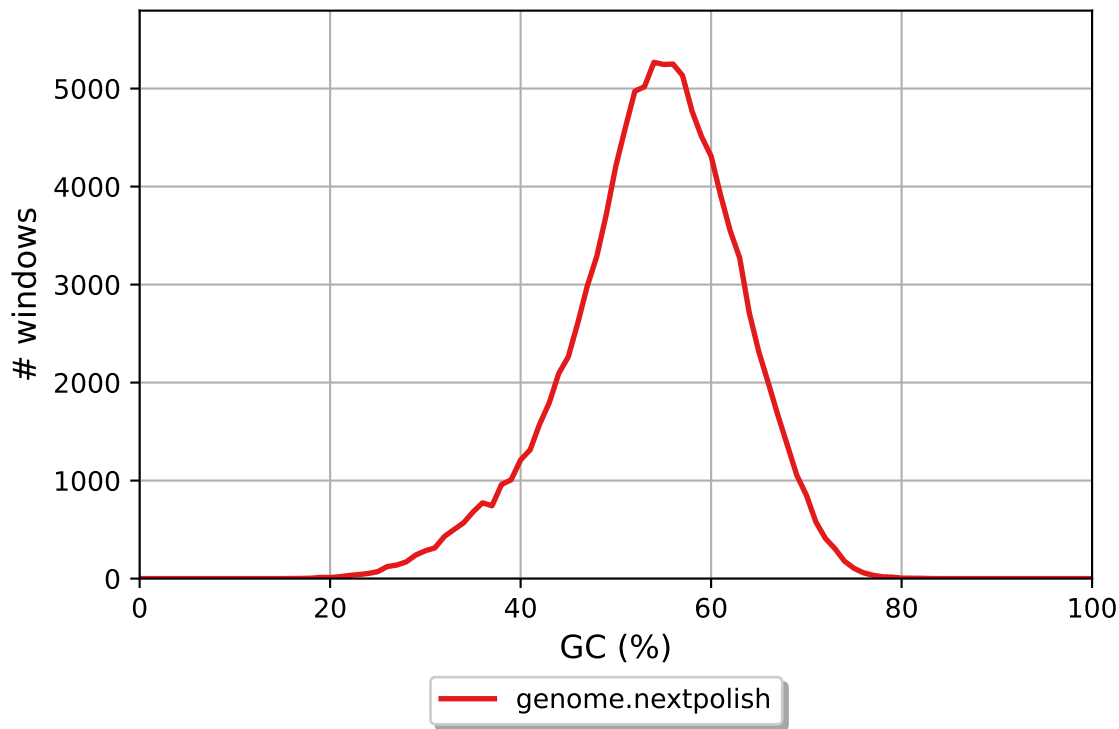
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

