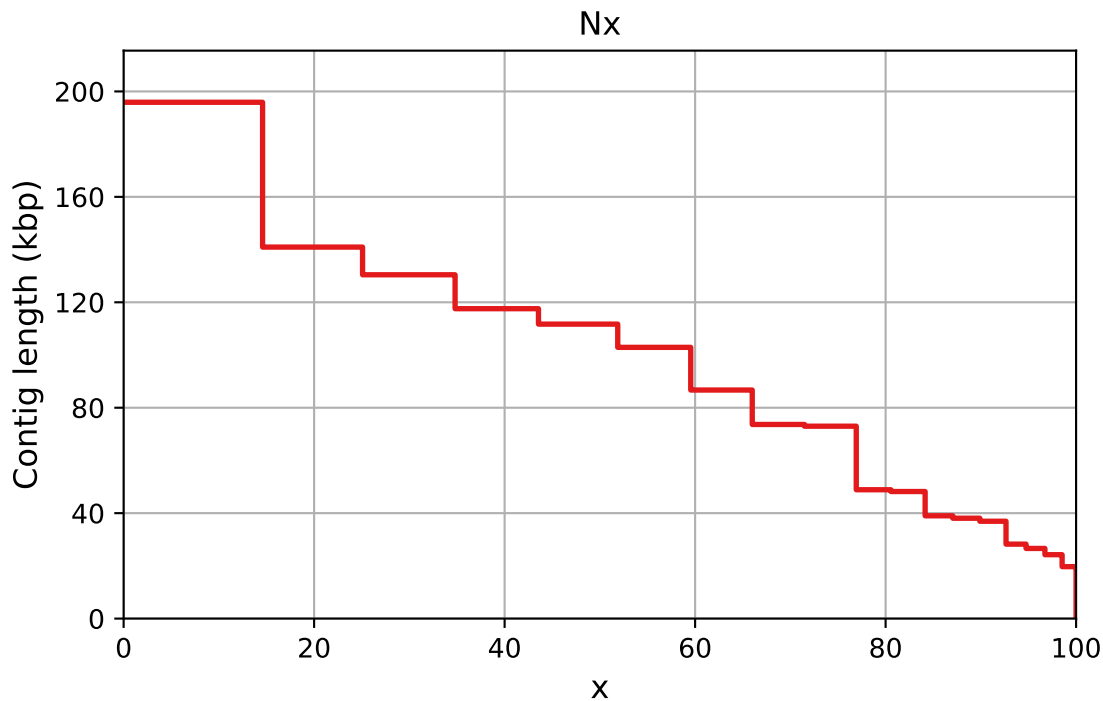


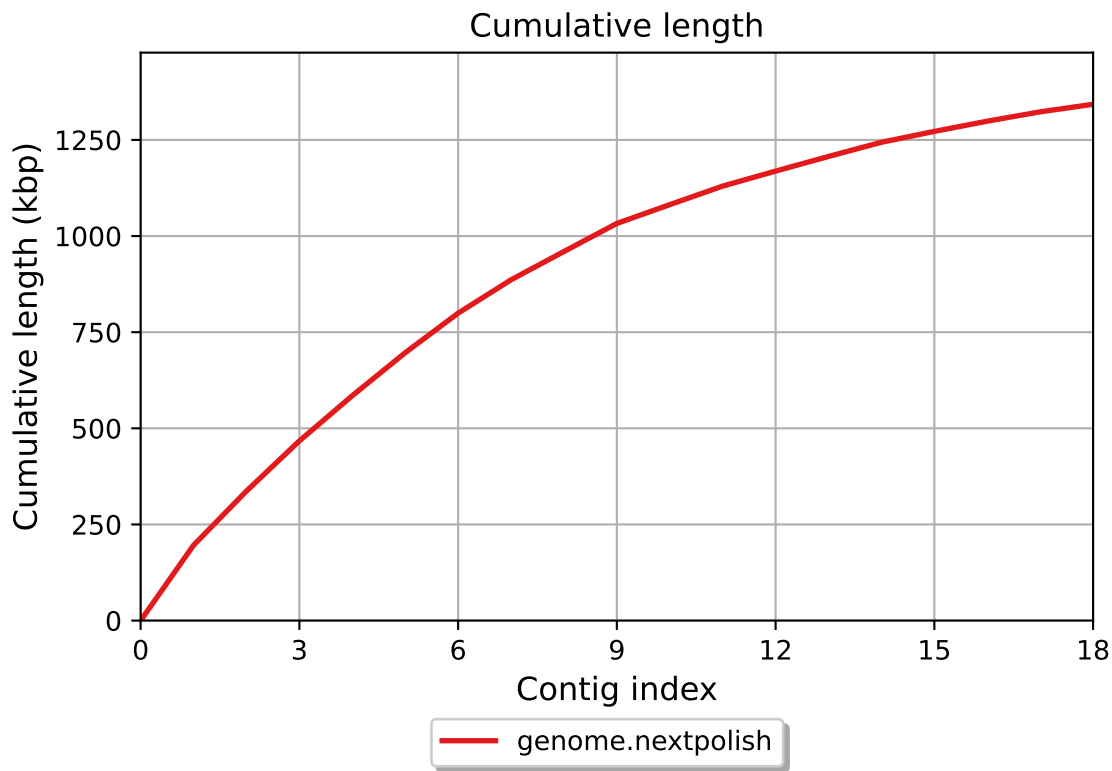
# Report

	genome.nextpolish
# contigs (>= 0 bp)	18
# contigs (>= 1000 bp)	18
# contigs (>= 5000 bp)	18
# contigs (>= 10000 bp)	18
# contigs (>= 25000 bp)	16
# contigs (>= 50000 bp)	9
Total length (>= 0 bp)	1342721
Total length (>= 1000 bp)	1342721
Total length (>= 5000 bp)	1342721
Total length (>= 10000 bp)	1342721
Total length (>= 25000 bp)	1298775
Total length (>= 50000 bp)	1032820
# contigs	18
Largest contig	195888
Total length	1342721
GC (%)	58.96
N50	111715
N90	36932
auN	105715.4
L50	5
L90	14
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 0 part

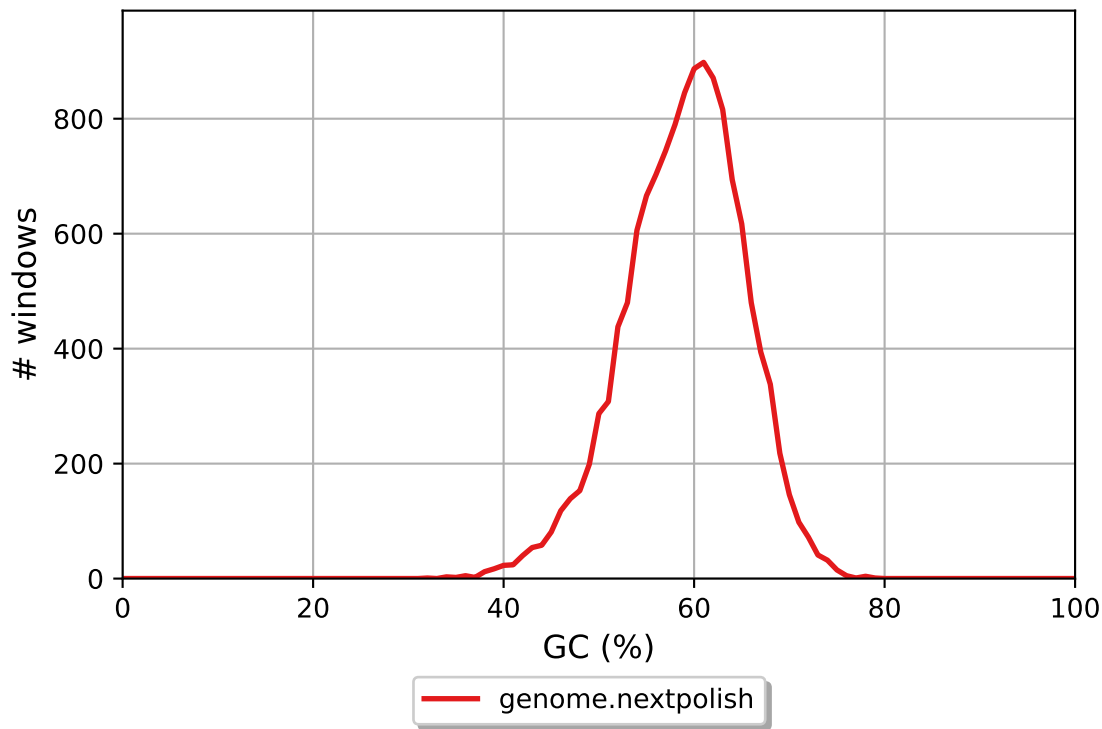
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— genome.nextpolish



## GC content



genome.nextpolish GC content

