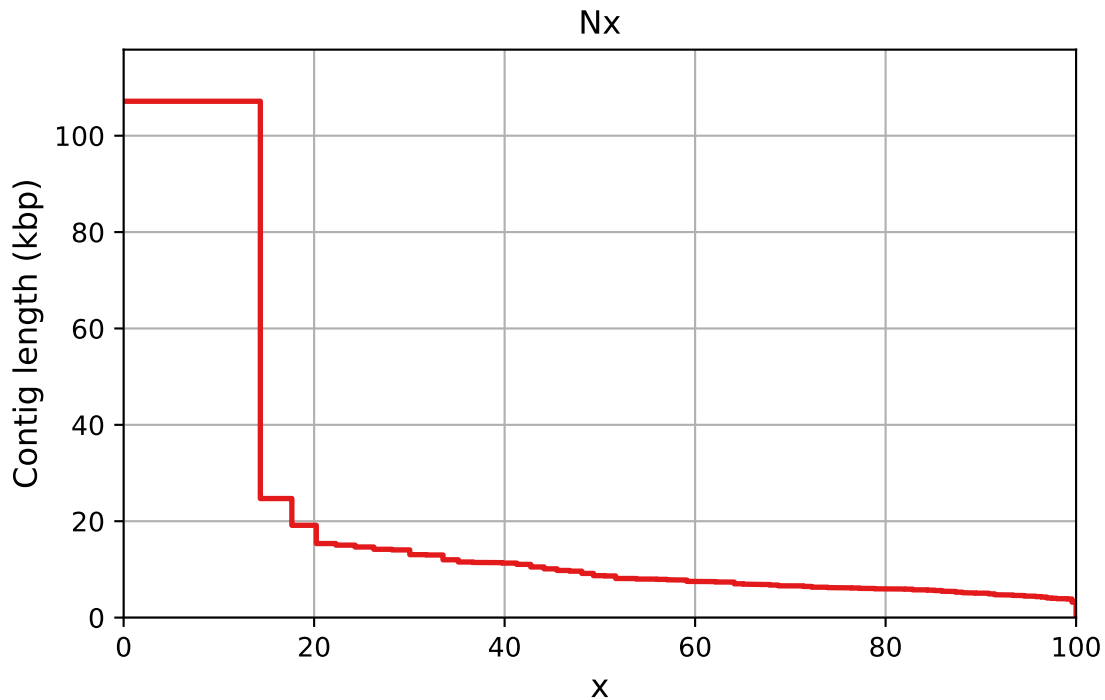


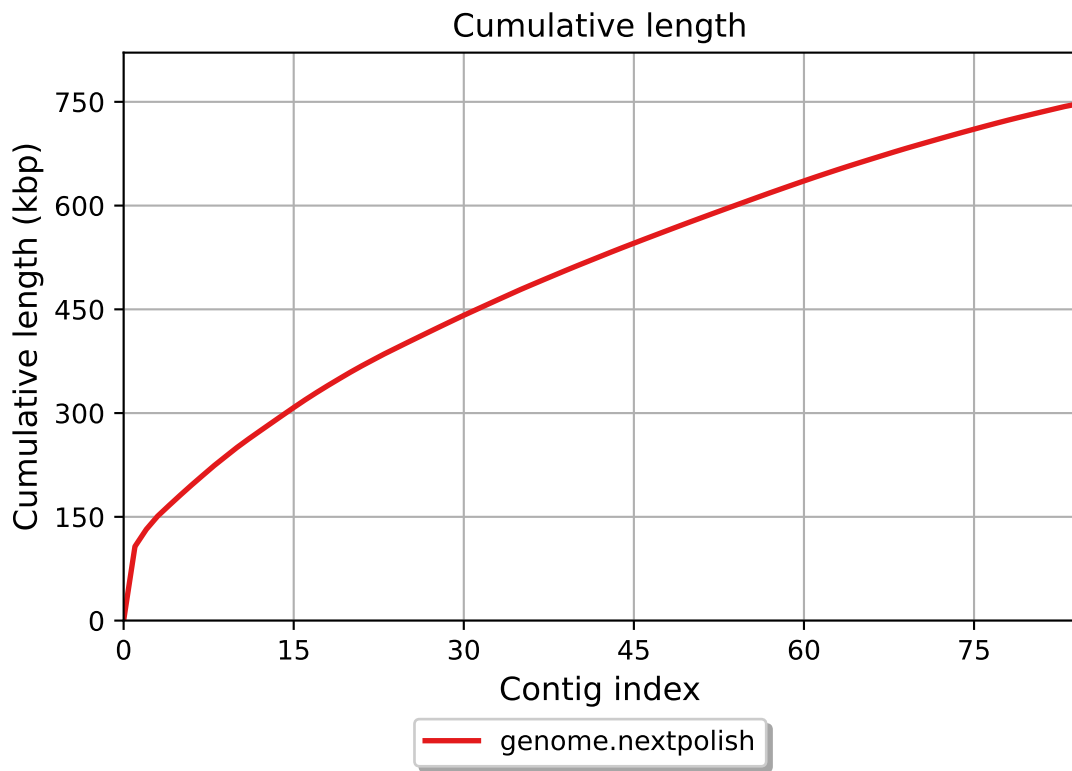
# Report

	genome.nextpolish
# contigs (>= 0 bp)	84
# contigs (>= 1000 bp)	84
# contigs (>= 5000 bp)	68
# contigs (>= 10000 bp)	18
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	746485
Total length (>= 1000 bp)	746485
Total length (>= 5000 bp)	677792
Total length (>= 10000 bp)	339673
Total length (>= 25000 bp)	107148
Total length (>= 50000 bp)	107148
# contigs	84
Largest contig	107148
Total length	746485
GC (%)	60.16
N50	8705
N90	5069
auN	23480.1
L50	22
L90	67
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 1 part

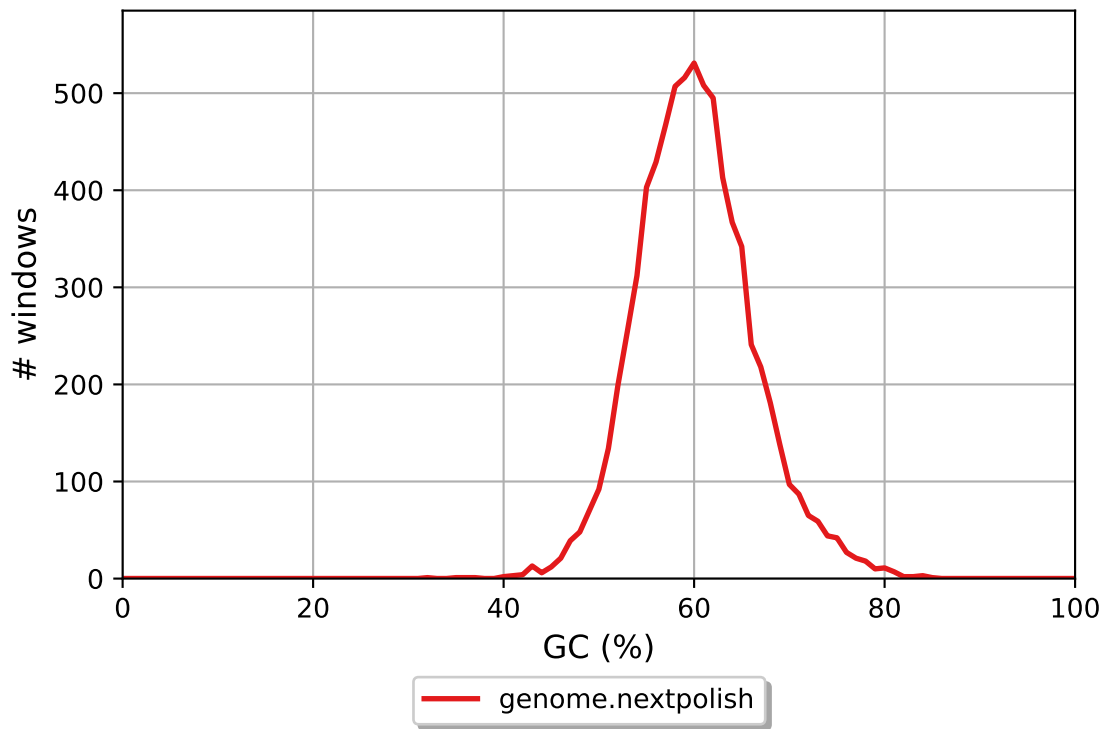
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— genome.nextpolish



## GC content



genome.nextpolish GC content

