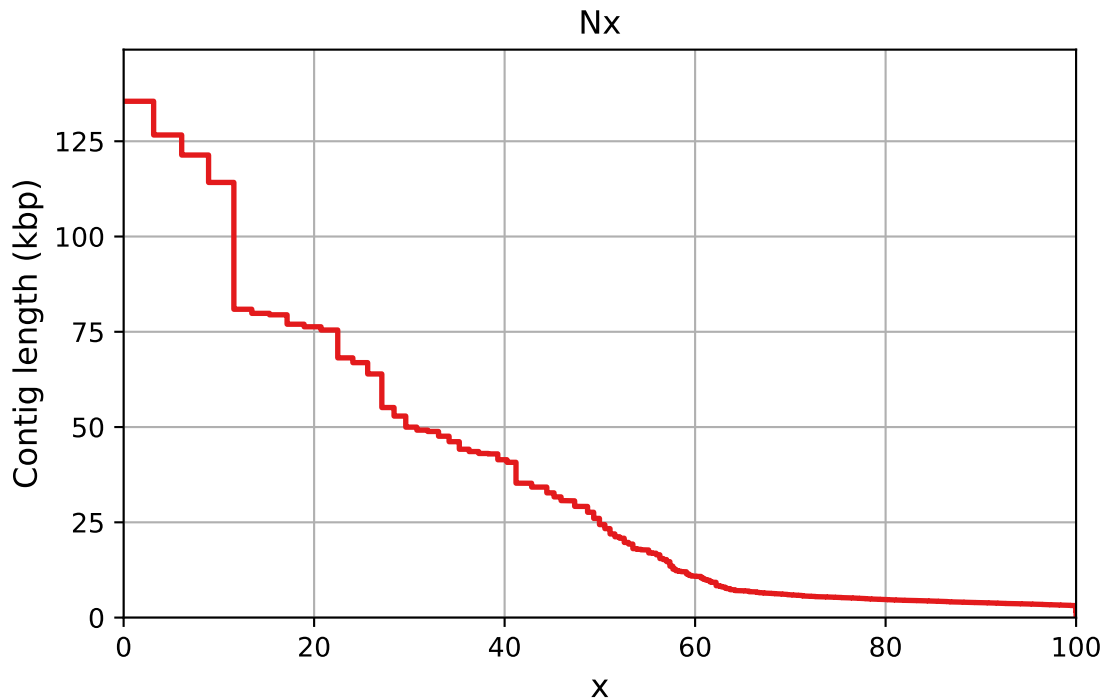


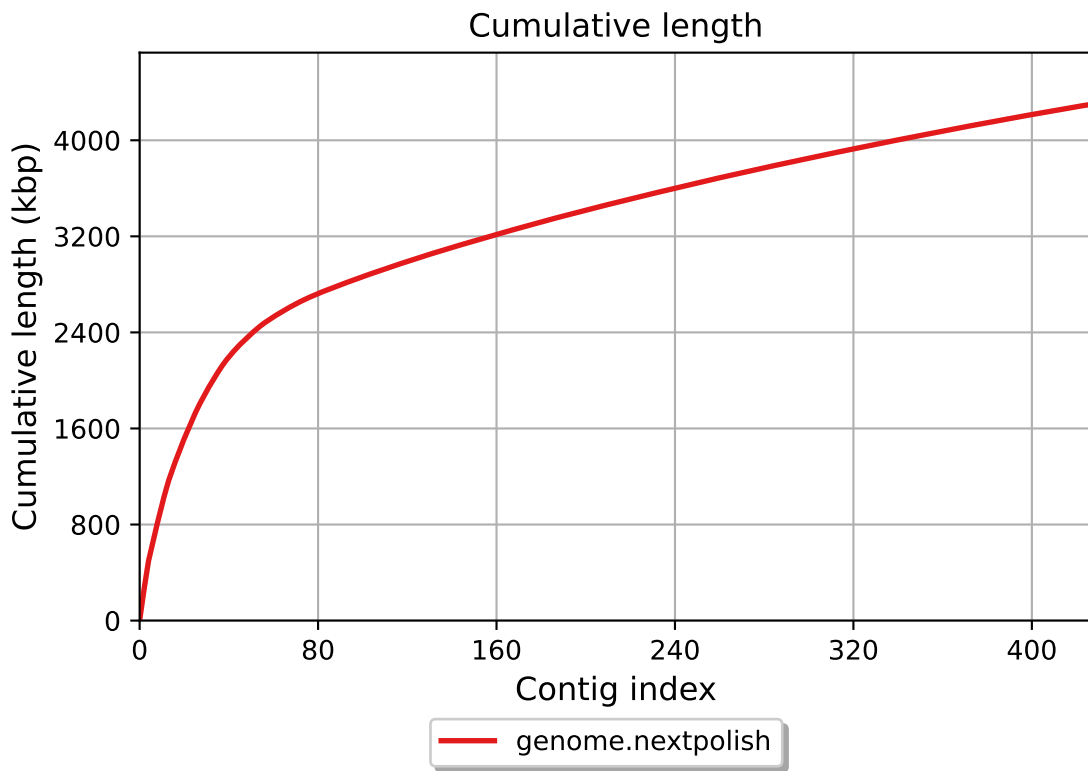
# Report

	genome.nextpolish
# contigs (>= 0 bp)	427
# contigs (>= 1000 bp)	427
# contigs (>= 5000 bp)	184
# contigs (>= 10000 bp)	69
# contigs (>= 25000 bp)	38
# contigs (>= 50000 bp)	15
Total length (>= 0 bp)	4299877
Total length (>= 1000 bp)	4299877
Total length (>= 5000 bp)	3339749
Total length (>= 10000 bp)	2627167
Total length (>= 25000 bp)	2148149
Total length (>= 50000 bp)	1273678
# contigs	427
Largest contig	135492
Total length	4299877
GC (%)	56.38
N50	24402
N90	3913
auN	39257.2
L50	39
L90	305
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	10 + 3 part

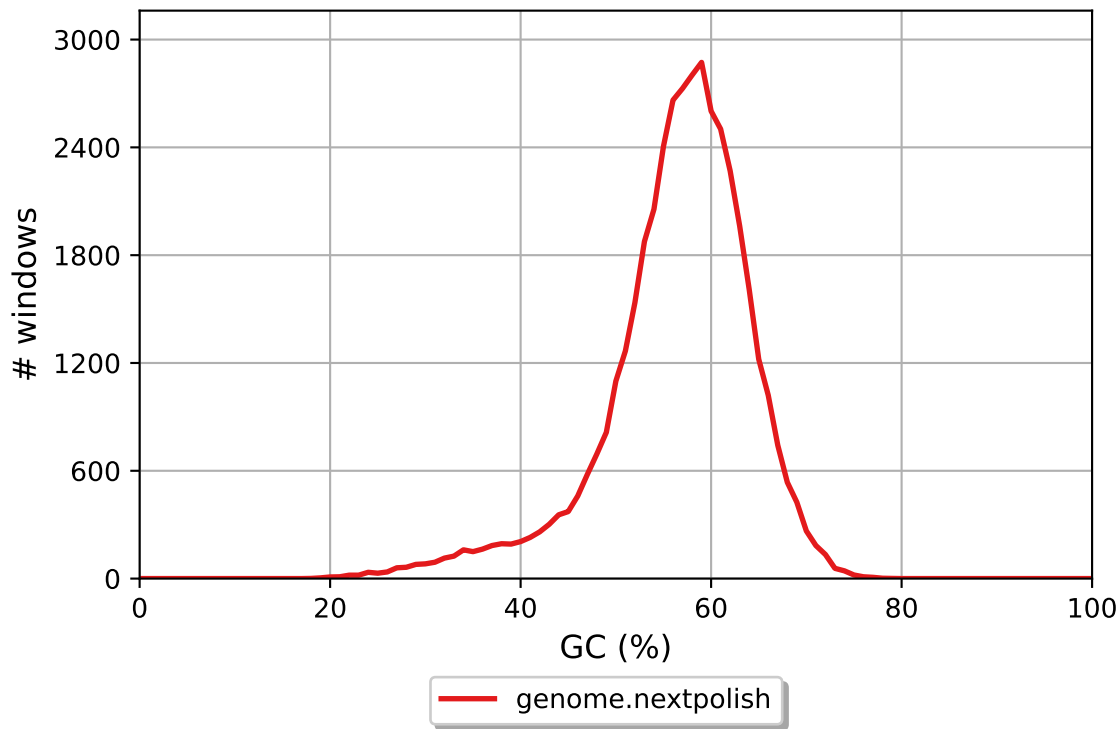
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— genome.nextpolish



## GC content



genome.nextpolish GC content

