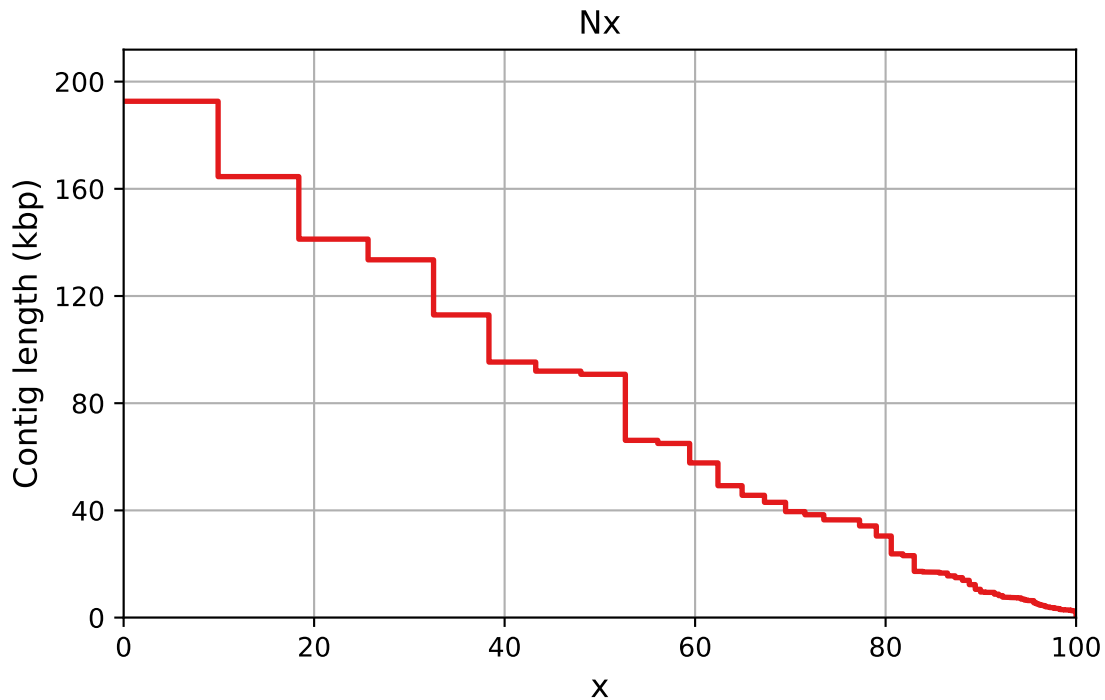


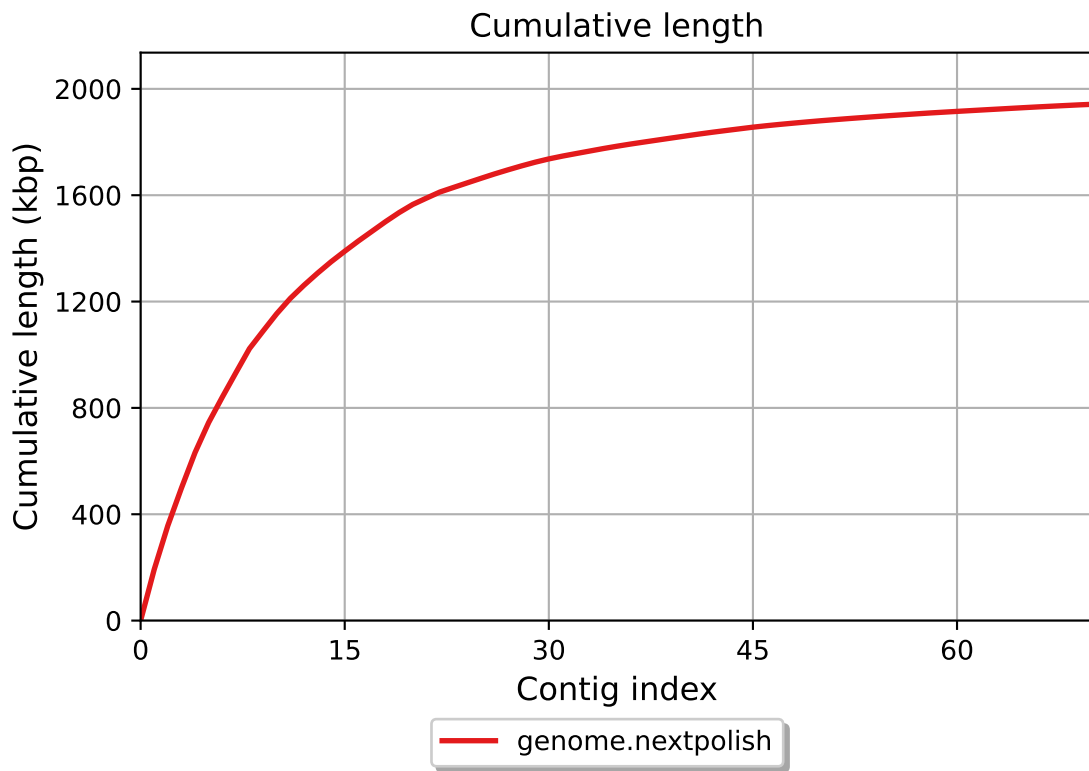
# Report

	genome.nextpolish
# contigs (>= 0 bp)	70
# contigs (>= 1000 bp)	70
# contigs (>= 5000 bp)	47
# contigs (>= 10000 bp)	31
# contigs (>= 25000 bp)	20
# contigs (>= 50000 bp)	11
Total length (>= 0 bp)	1942064
Total length (>= 1000 bp)	1942064
Total length (>= 5000 bp)	1866491
Total length (>= 10000 bp)	1747085
Total length (>= 25000 bp)	1565187
Total length (>= 50000 bp)	1211836
# contigs	70
Largest contig	192669
Total length	1942064
GC (%)	55.70
N50	90786
N90	9547
auN	88017.8
L50	8
L90	32
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	9 + 1 part

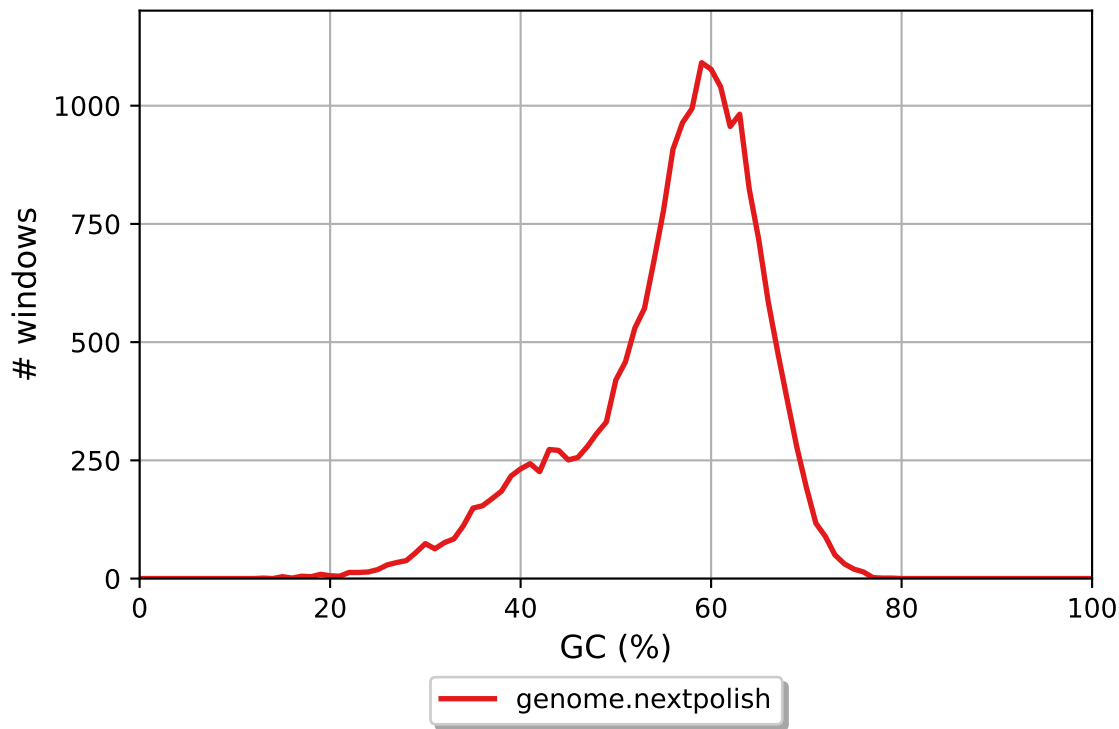
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— genome.nextpolish



## GC content



genome.nextpolish GC content

