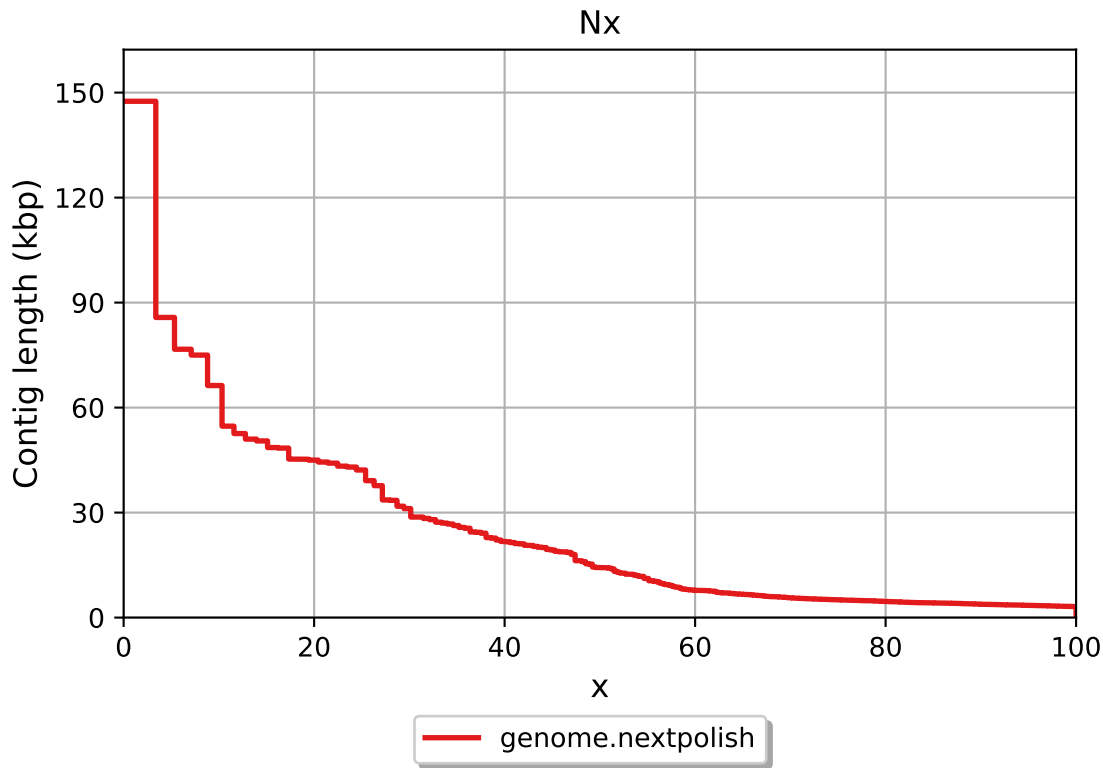


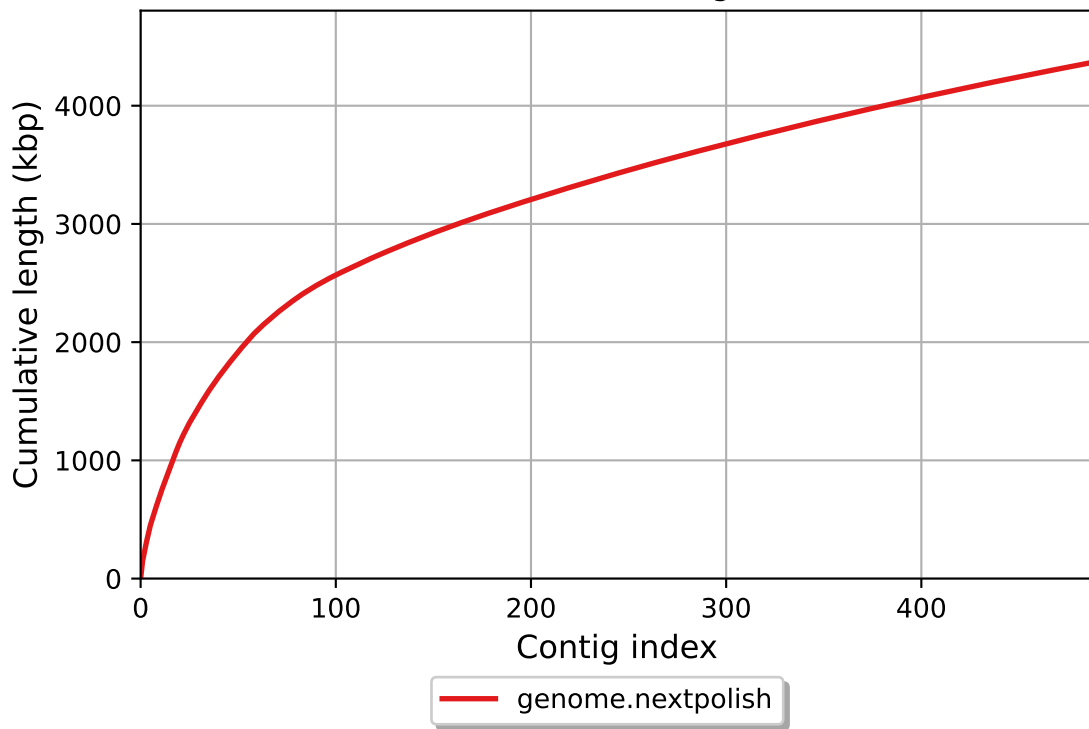
# Report

	genome.nextpolish
# contigs (>= 0 bp)	488
# contigs (>= 1000 bp)	488
# contigs (>= 5000 bp)	219
# contigs (>= 10000 bp)	88
# contigs (>= 25000 bp)	35
# contigs (>= 50000 bp)	9
Total length (>= 0 bp)	4367504
Total length (>= 1000 bp)	4367504
Total length (>= 5000 bp)	3304268
Total length (>= 10000 bp)	2459617
Total length (>= 25000 bp)	1588780
Total length (>= 50000 bp)	659991
# contigs	488
Largest contig	147522
Total length	4367504
GC (%)	57.32
N50	14310
N90	3811
auN	26561.5
L50	66
L90	363
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	6 + 0 part

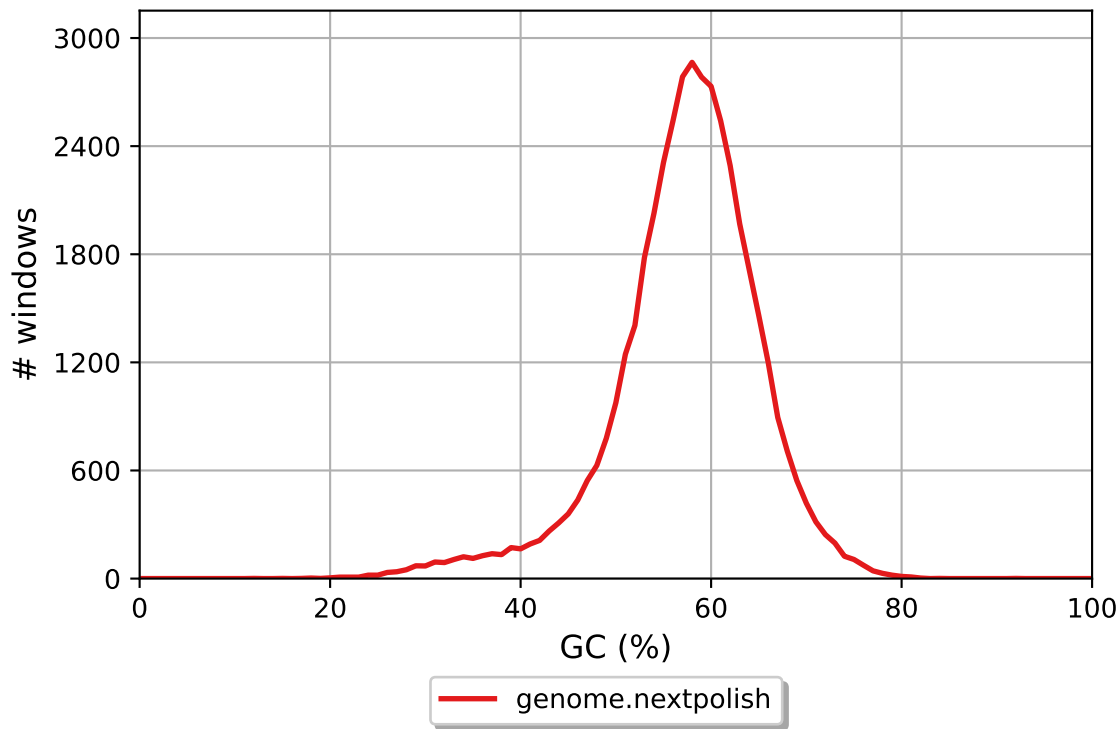
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



## GC content



genome.nextpolish GC content

