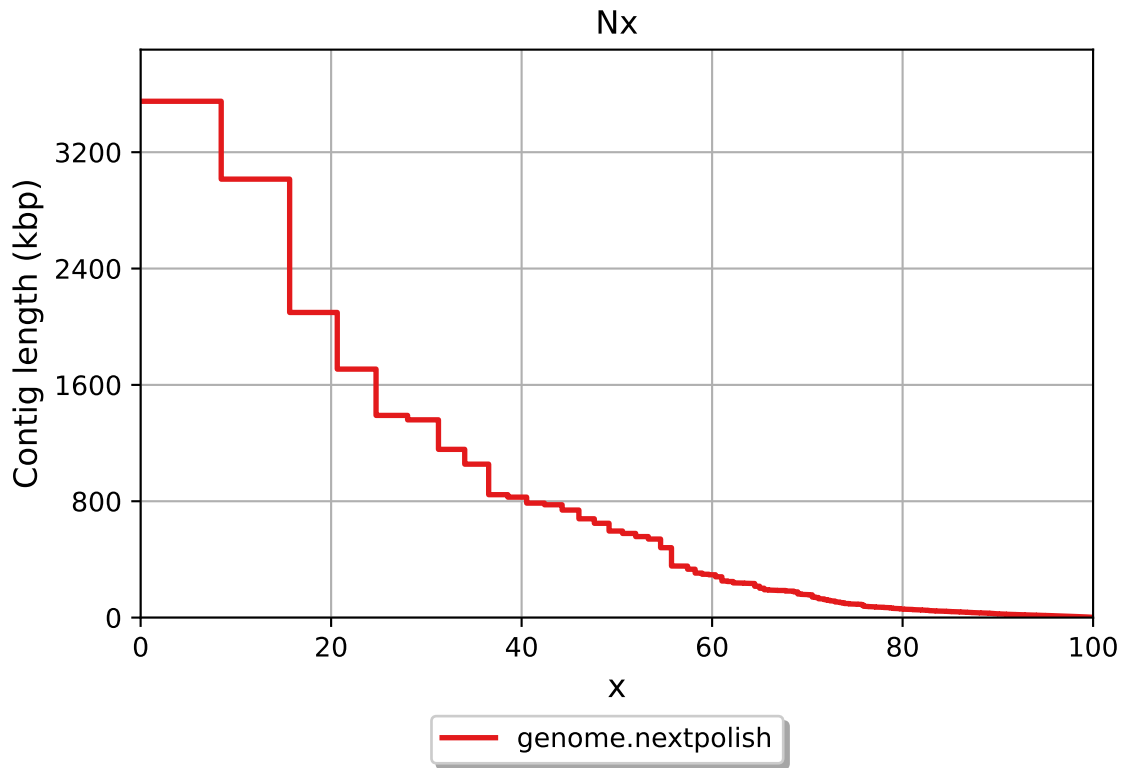


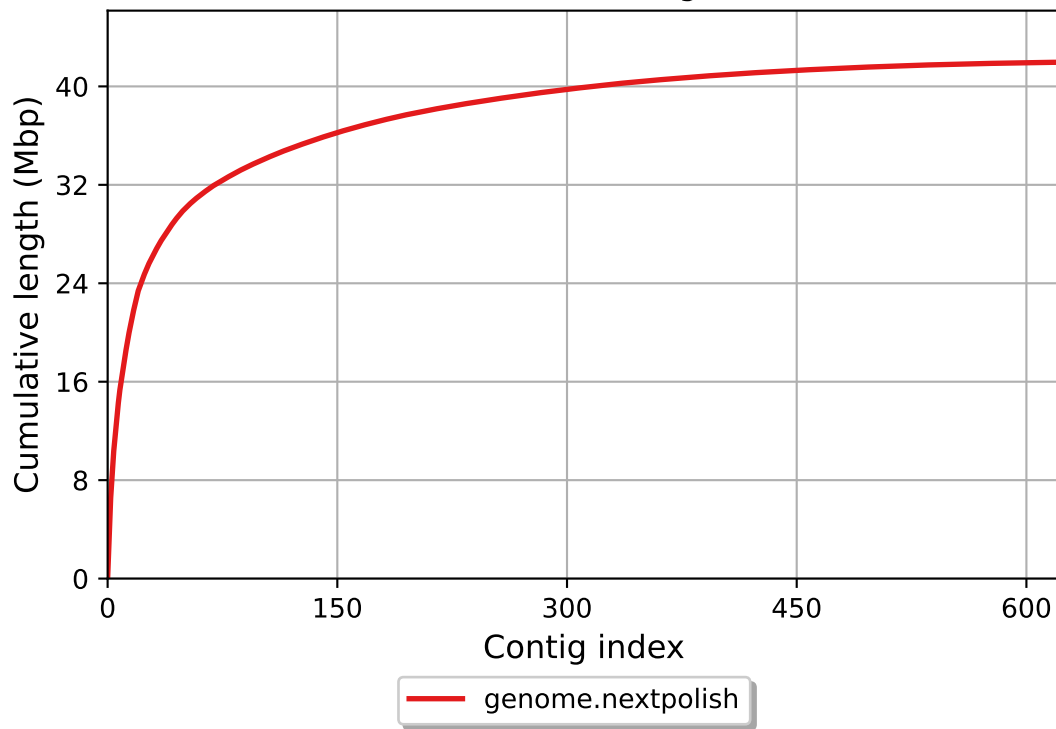
# Report

	genome.nextpolish
# contigs (>= 0 bp)	622
# contigs (>= 1000 bp)	620
# contigs (>= 5000 bp)	496
# contigs (>= 10000 bp)	375
# contigs (>= 25000 bp)	201
# contigs (>= 50000 bp)	113
Total length (>= 0 bp)	41961042
Total length (>= 1000 bp)	41959453
Total length (>= 5000 bp)	41569027
Total length (>= 10000 bp)	40689862
Total length (>= 25000 bp)	37845341
Total length (>= 50000 bp)	34654770
# contigs	622
Largest contig	3550210
Total length	41961042
GC (%)	45.93
N50	595110
N90	25642
auN	1023400.0
L50	16
L90	198
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	122 + 0 part

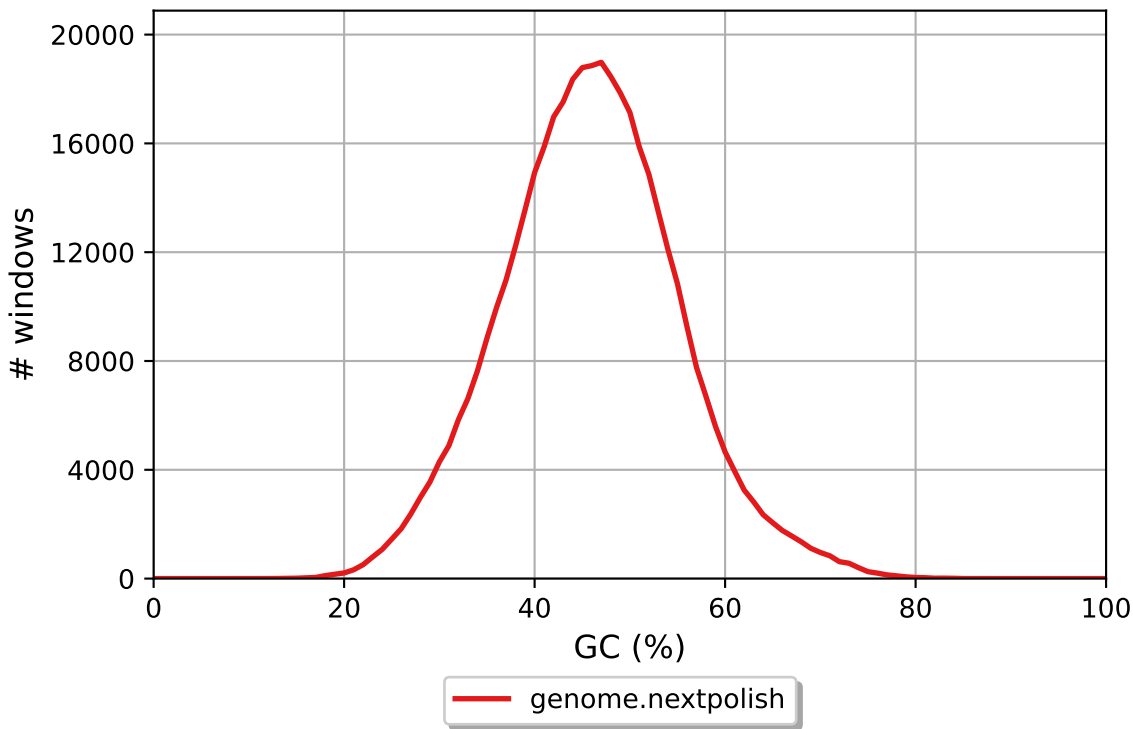
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

