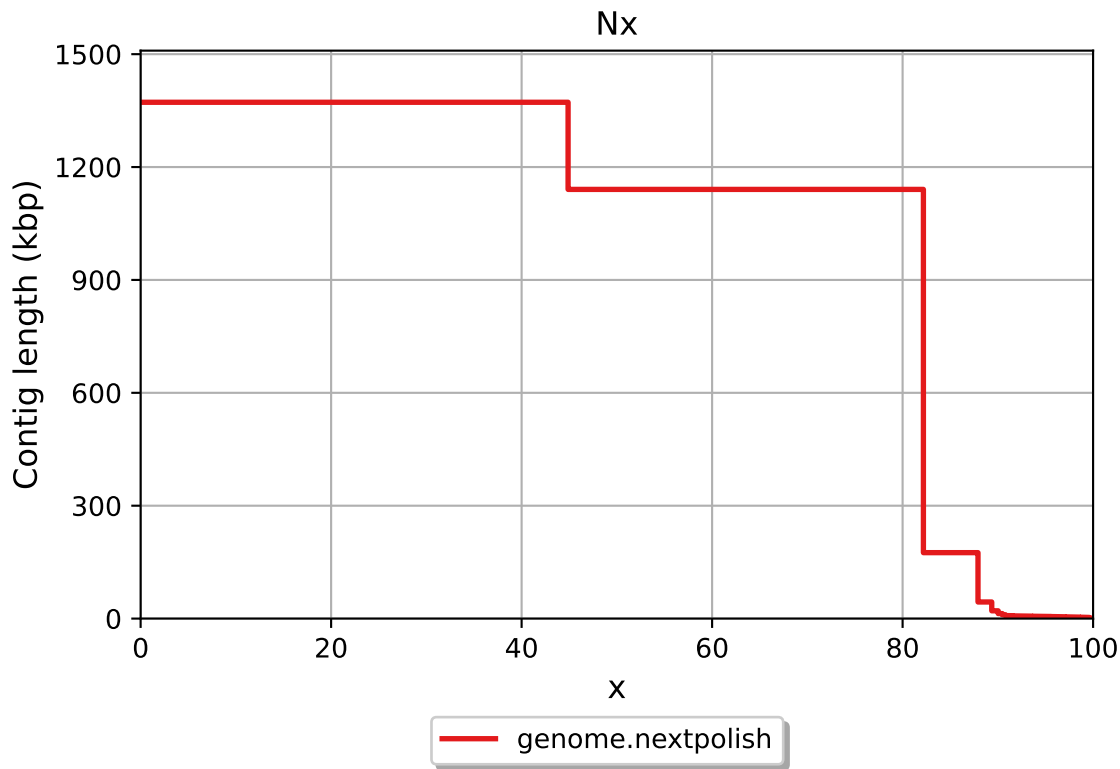


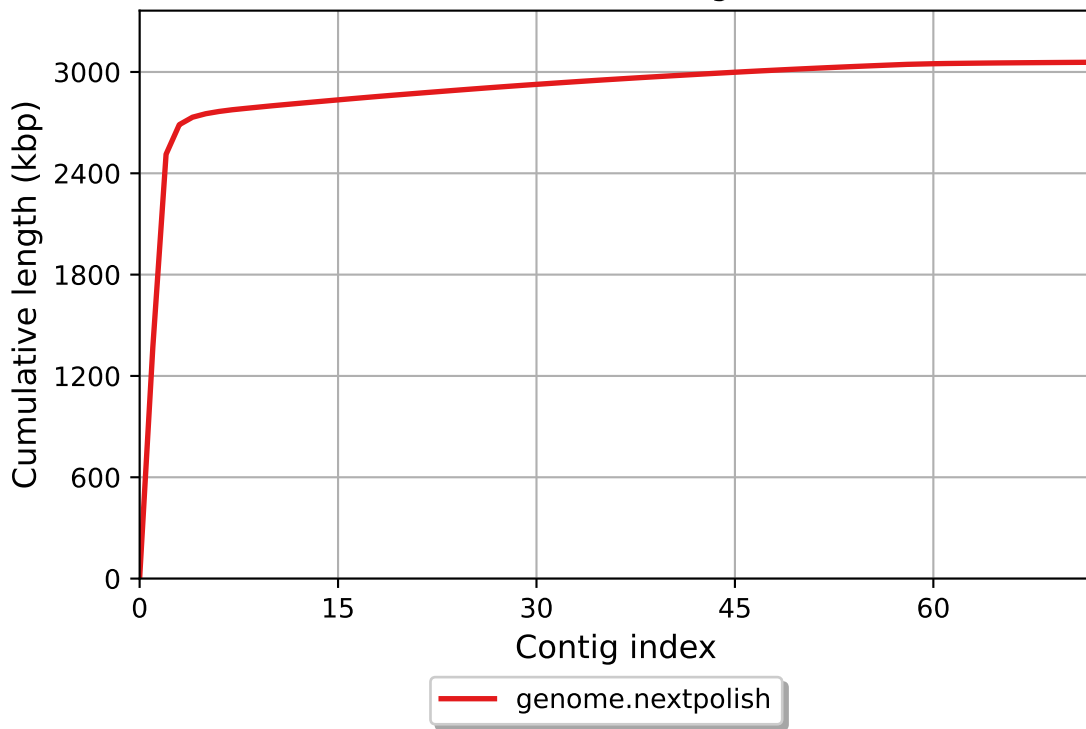
Report

	genome.nextpolish
# contigs (>= 0 bp)	72
# contigs (>= 1000 bp)	61
# contigs (>= 5000 bp)	35
# contigs (>= 10000 bp)	7
# contigs (>= 25000 bp)	4
# contigs (>= 50000 bp)	3
Total length (>= 0 bp)	3057453
Total length (>= 1000 bp)	3050123
Total length (>= 5000 bp)	2952700
Total length (>= 10000 bp)	2776249
Total length (>= 25000 bp)	2732110
Total length (>= 50000 bp)	2687987
# contigs	72
Largest contig	1372161
Total length	3057453
GC (%)	34.55
N50	1140594
N90	20673
auN	1052727.2
L50	2
L90	5
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	7 + 8 part

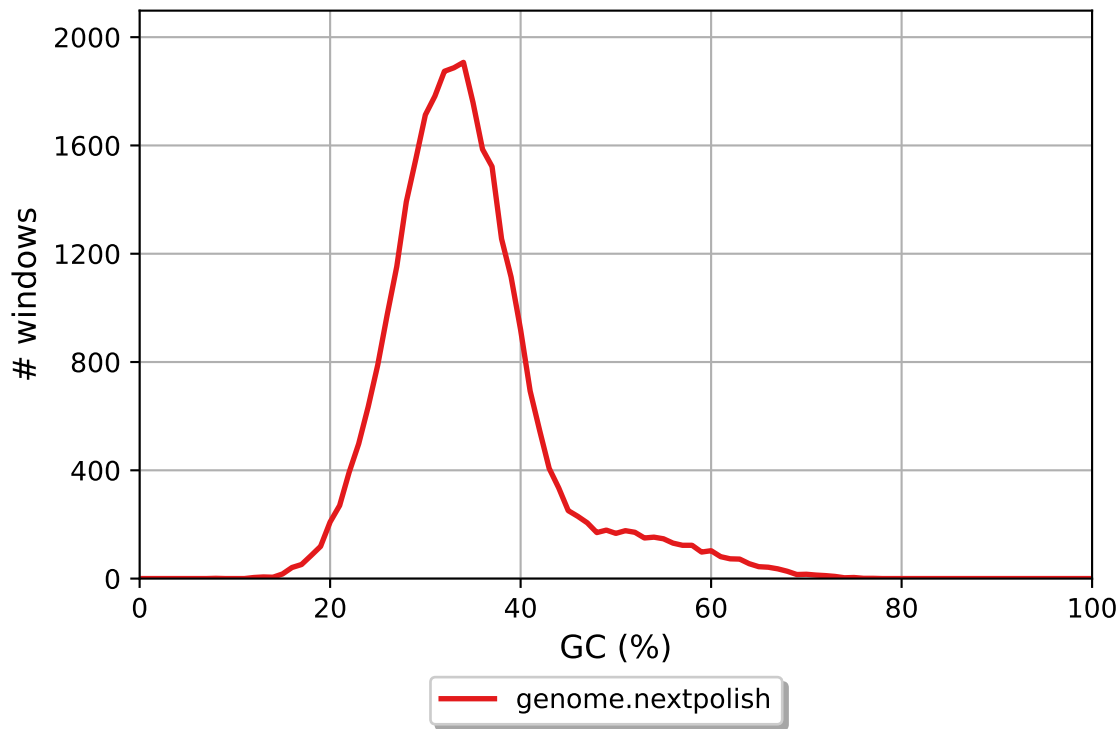
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

