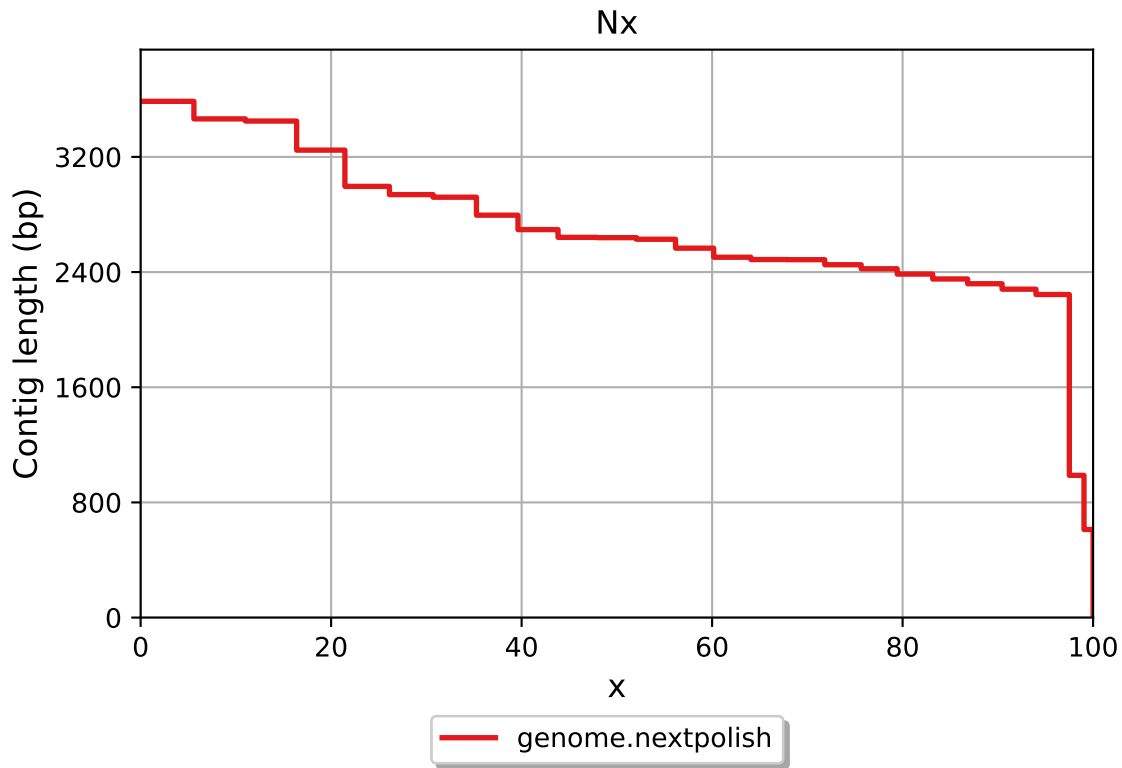


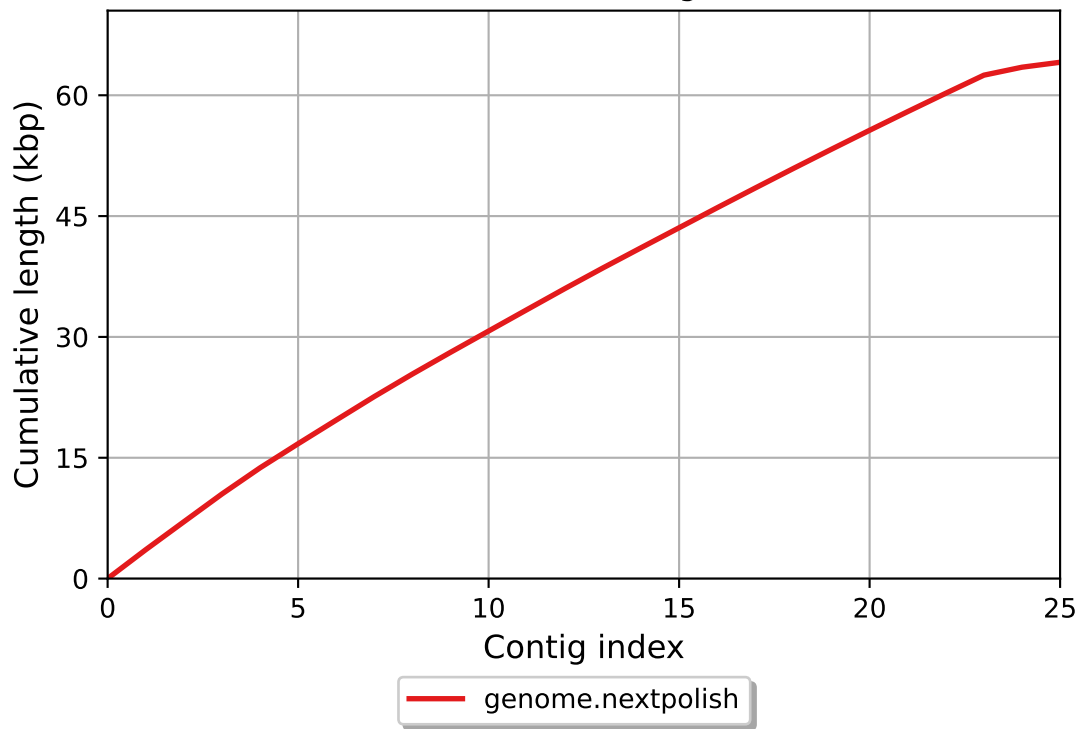
Report

	genome.nextpolish
# contigs (>= 0 bp)	25
# contigs (>= 1000 bp)	23
# contigs (>= 5000 bp)	0
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	64095
Total length (>= 1000 bp)	62495
Total length (>= 5000 bp)	0
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	25
Largest contig	3586
Total length	64095
GC (%)	39.85
N50	2639
N90	2319
auN	2724.6
L50	11
L90	21
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

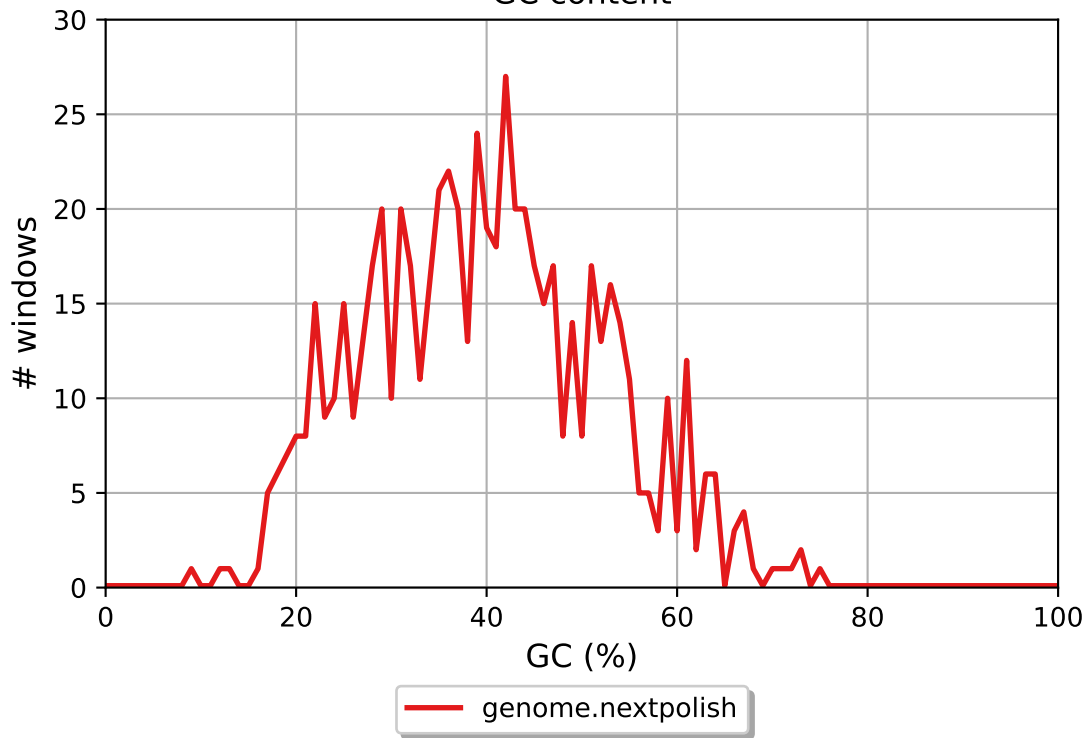
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

