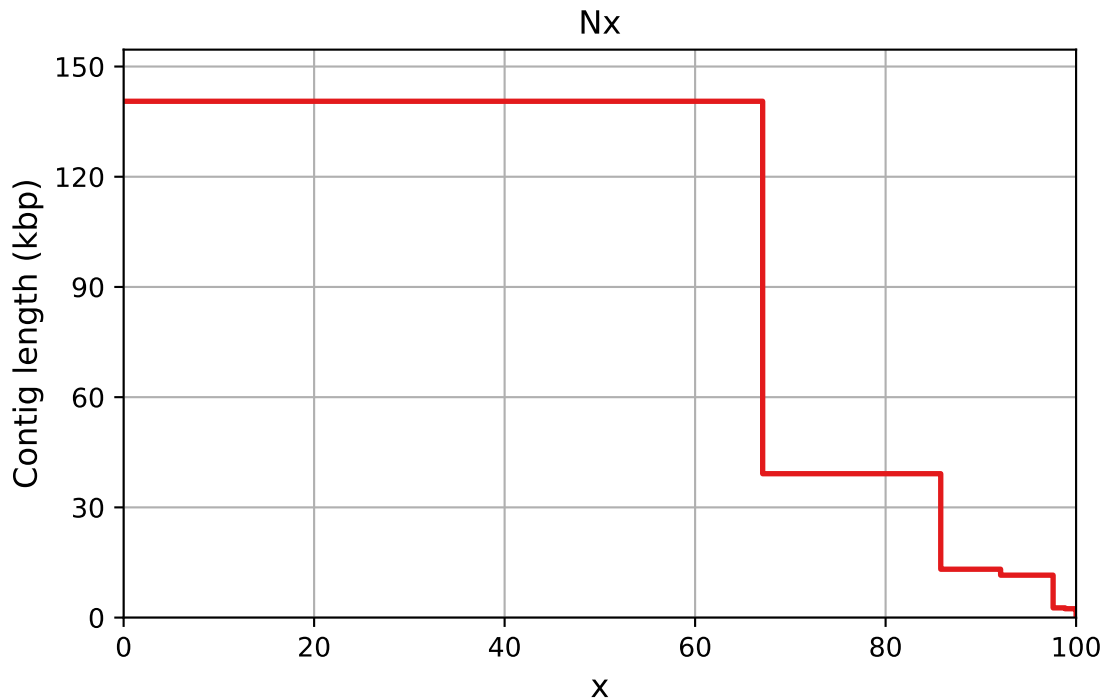


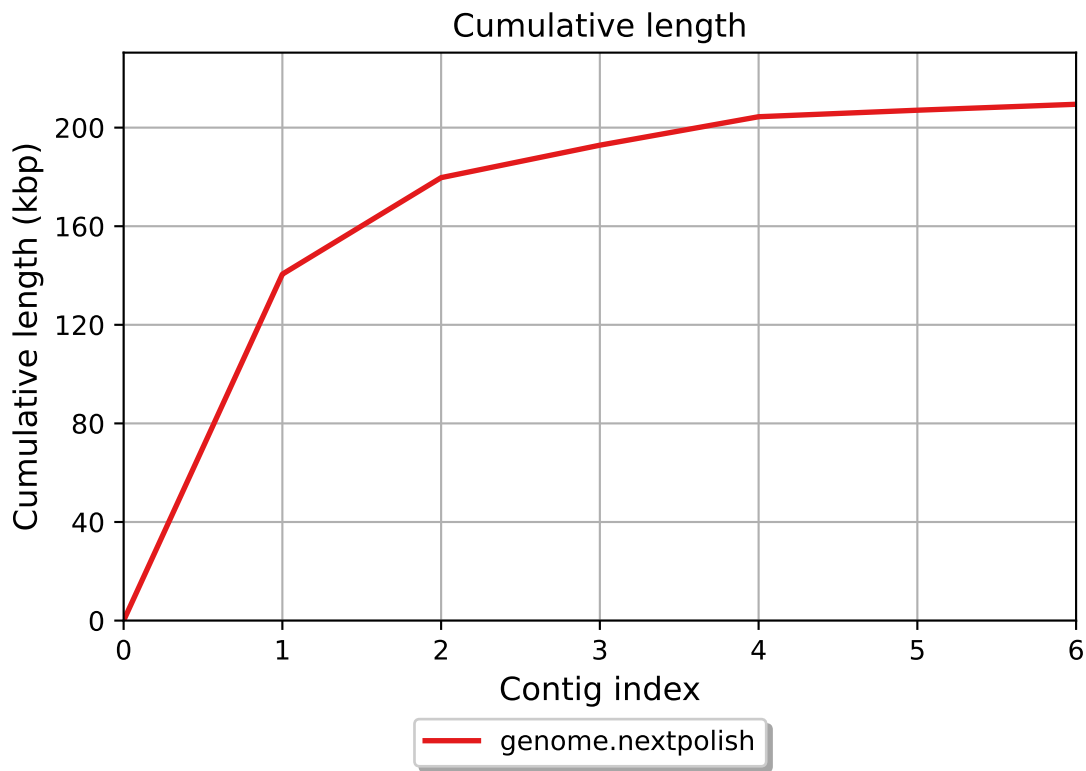
Report

	genome.nextpolish
# contigs (>= 0 bp)	6
# contigs (>= 1000 bp)	6
# contigs (>= 5000 bp)	4
# contigs (>= 10000 bp)	4
# contigs (>= 25000 bp)	2
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	209476
Total length (>= 1000 bp)	209476
Total length (>= 5000 bp)	204420
Total length (>= 10000 bp)	204420
Total length (>= 25000 bp)	179708
Total length (>= 50000 bp)	140544
# contigs	6
Largest contig	140544
Total length	209476
GC (%)	33.25
N50	140544
N90	13166
auN	103142.6
L50	1
L90	3
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

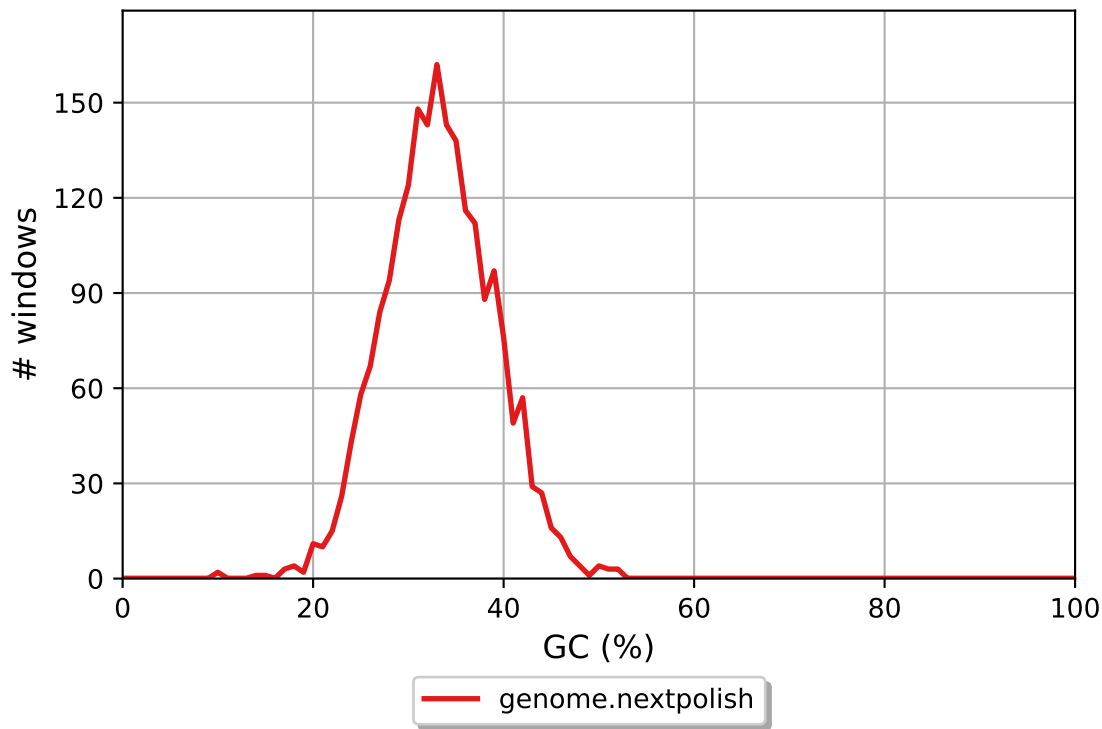
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— genome.nextpolish



GC content



genome.nextpolish GC content

