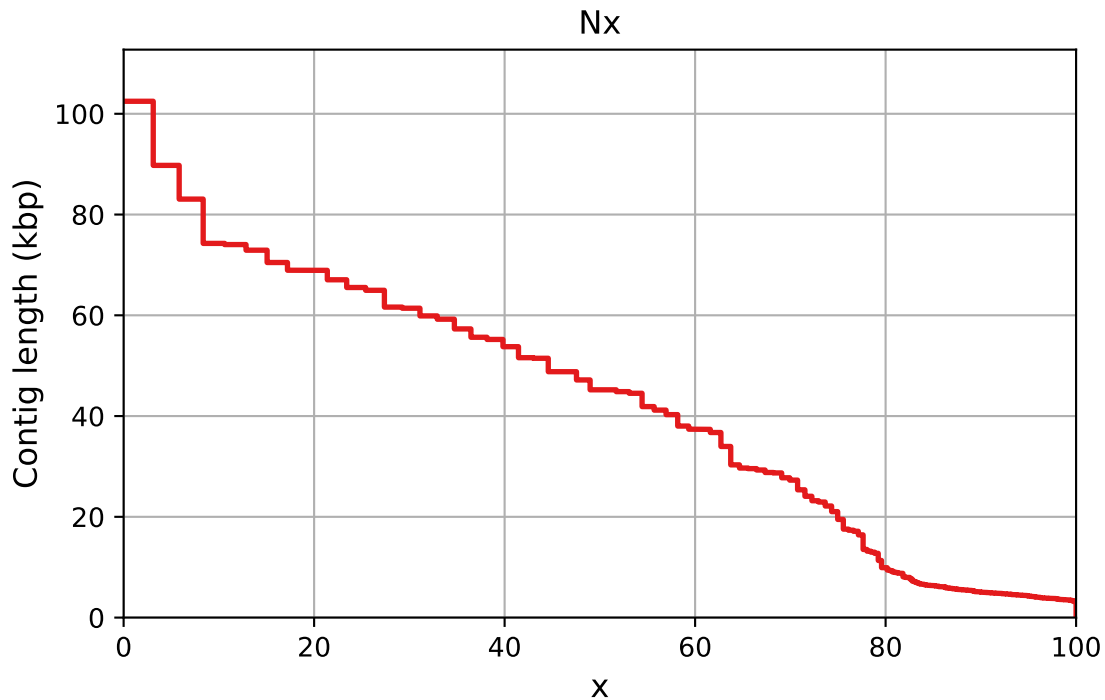


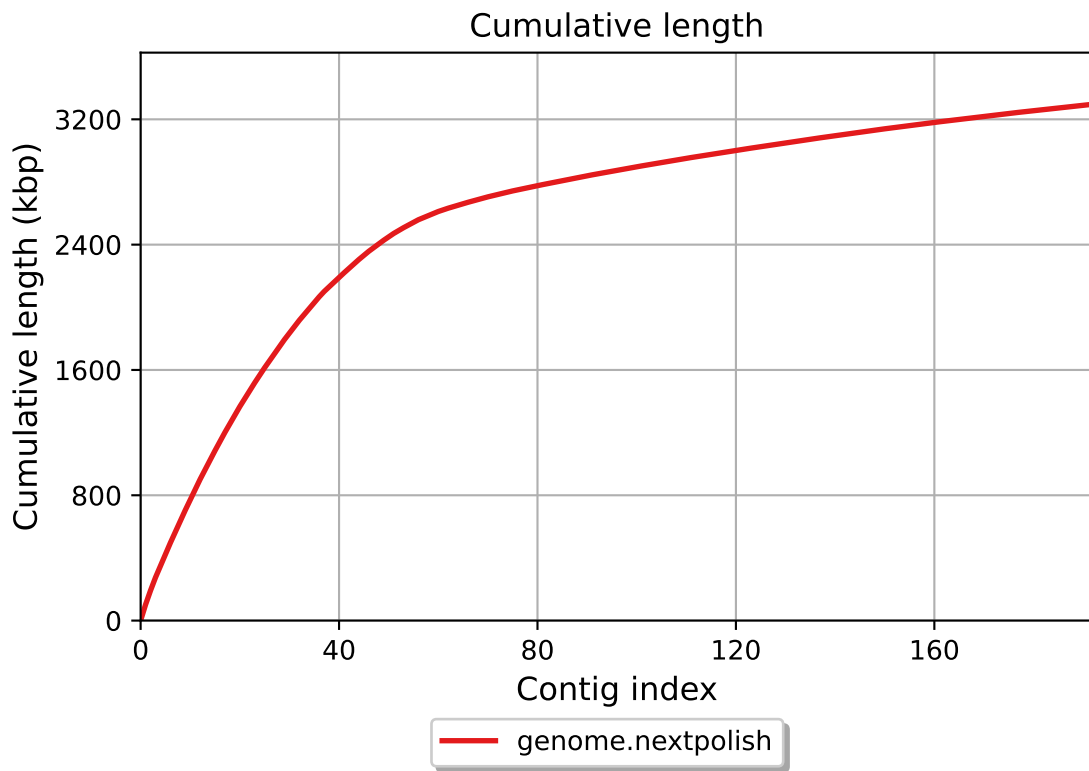
Report

	genome.nextpolish
# contigs (>= 0 bp)	192
# contigs (>= 1000 bp)	192
# contigs (>= 5000 bp)	116
# contigs (>= 10000 bp)	61
# contigs (>= 25000 bp)	46
# contigs (>= 50000 bp)	22
Total length (>= 0 bp)	3296102
Total length (>= 1000 bp)	3296102
Total length (>= 5000 bp)	2981411
Total length (>= 10000 bp)	2622800
Total length (>= 25000 bp)	2357699
Total length (>= 50000 bp)	1469539
# contigs	192
Largest contig	102483
Total length	3296102
GC (%)	58.35
N50	45213
N90	5041
auN	43835.1
L50	26
L90	114
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	5 + 0 part

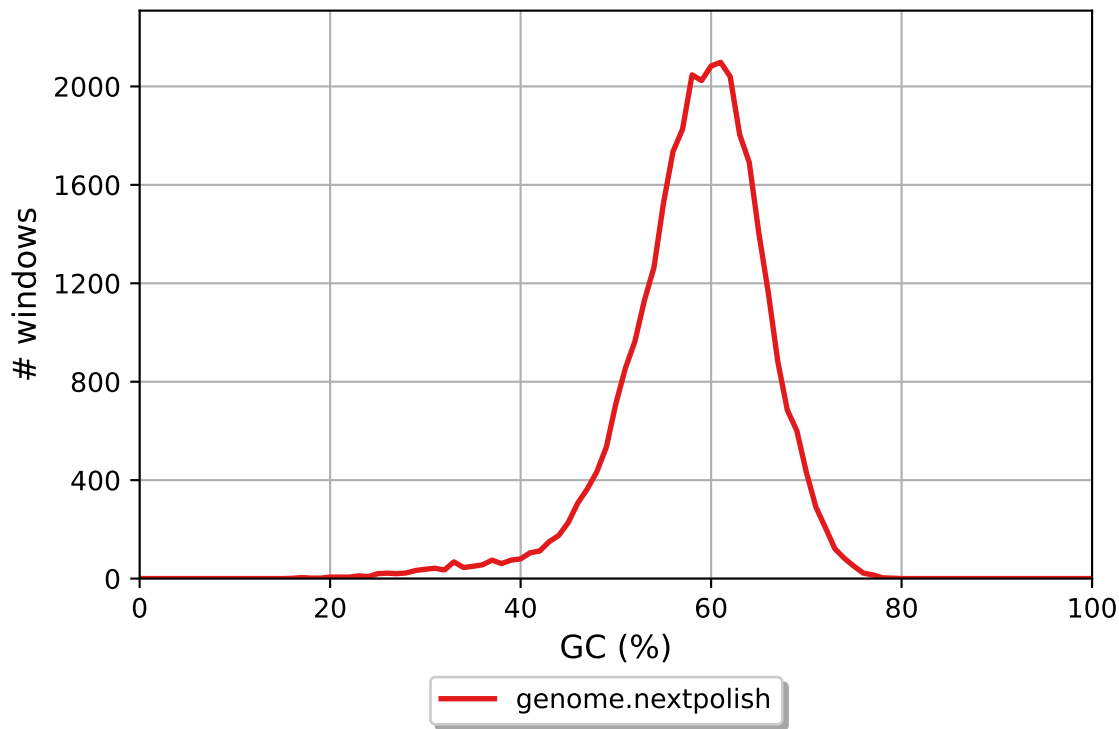
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— genome.nextpolish



GC content



genome.nextpolish GC content

