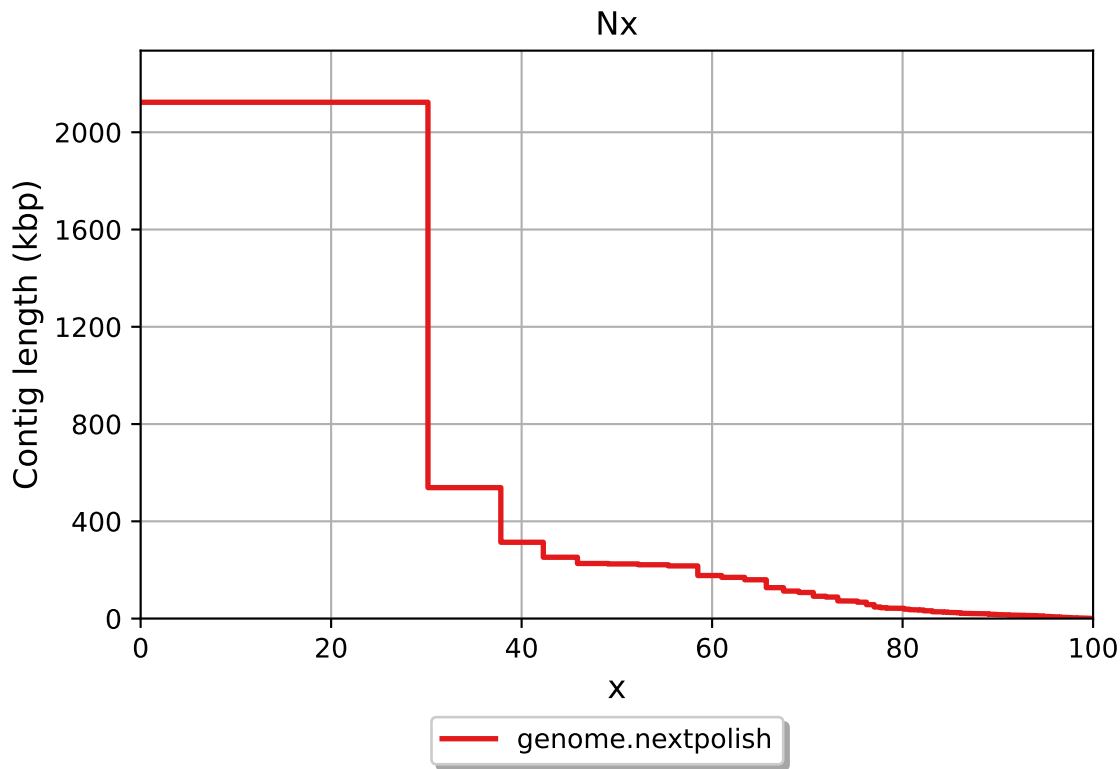


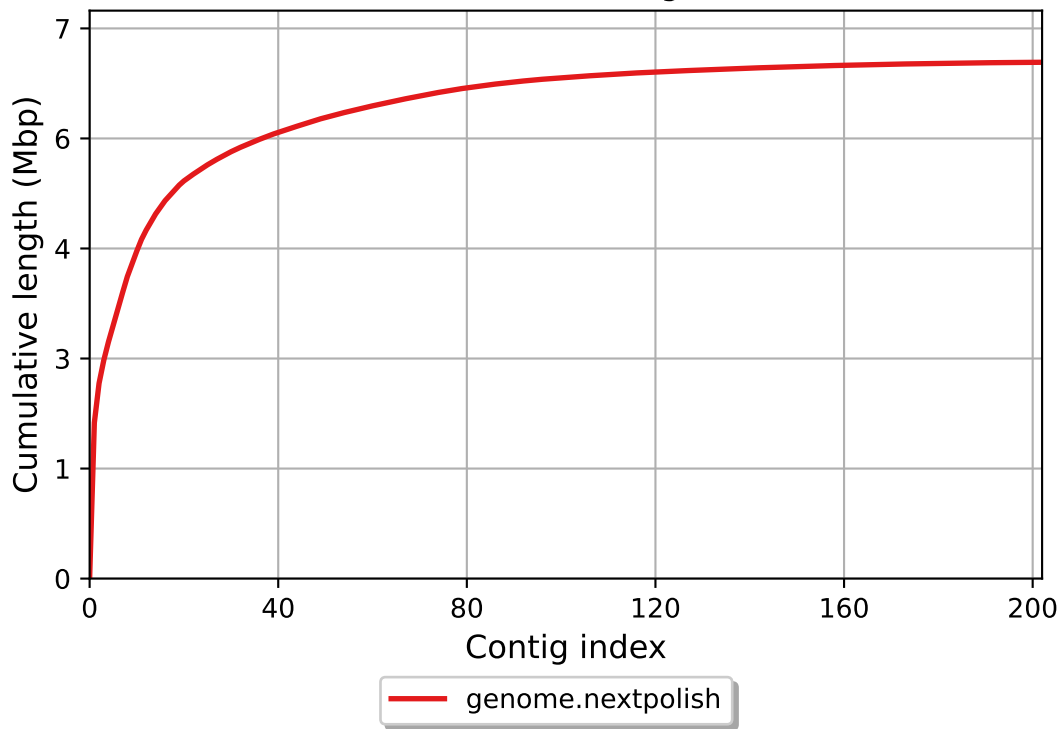
# Report

	genome.nextpolish
# contigs (>= 0 bp)	206
# contigs (>= 1000 bp)	179
# contigs (>= 5000 bp)	99
# contigs (>= 10000 bp)	78
# contigs (>= 25000 bp)	37
# contigs (>= 50000 bp)	20
Total length (>= 0 bp)	7040069
Total length (>= 1000 bp)	7021932
Total length (>= 5000 bp)	6821974
Total length (>= 10000 bp)	6671120
Total length (>= 25000 bp)	6012611
Total length (>= 50000 bp)	5420995
# contigs	202
Largest contig	2123331
Total length	7038468
GC (%)	39.97
N50	224734
N90	16375
auN	760600.3
L50	6
L90	53
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	32 + 0 part

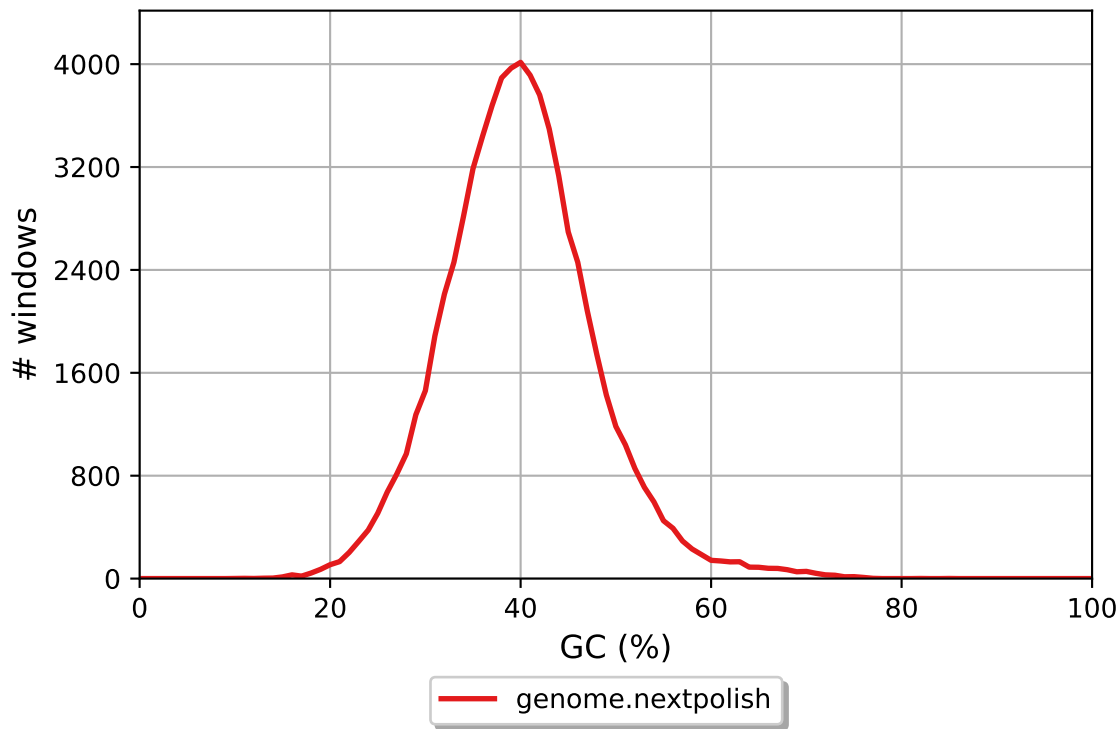
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



## GC content



genome.nextpolish GC content

