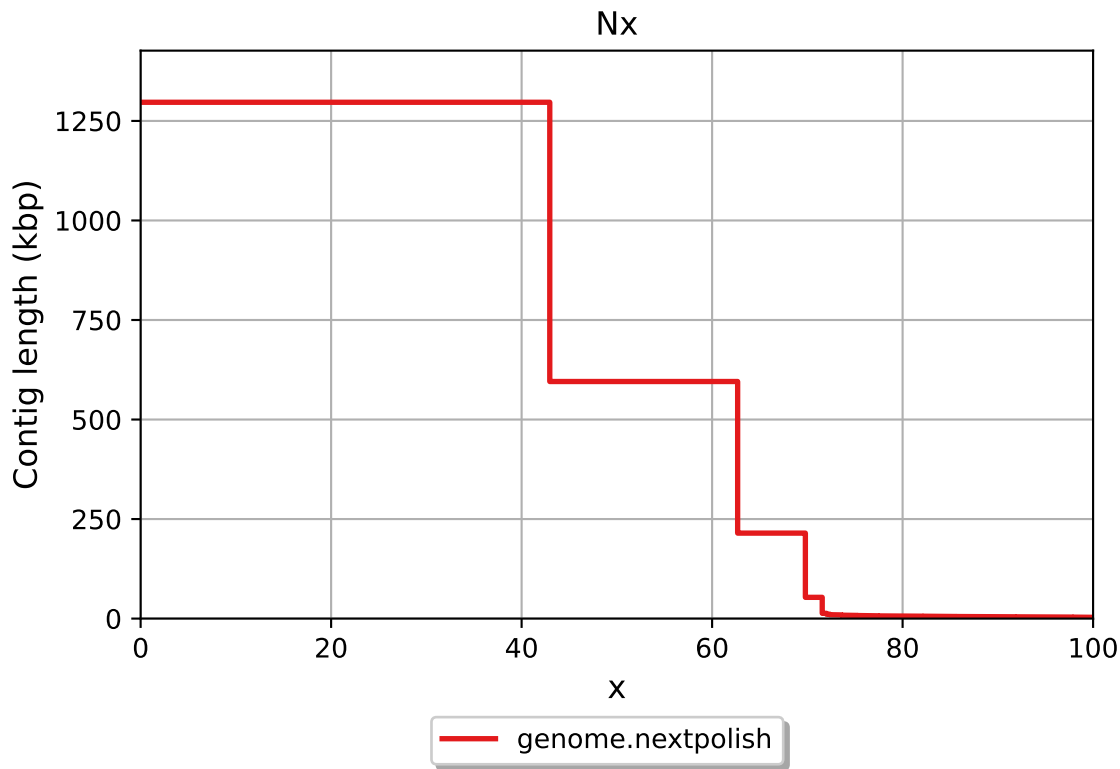
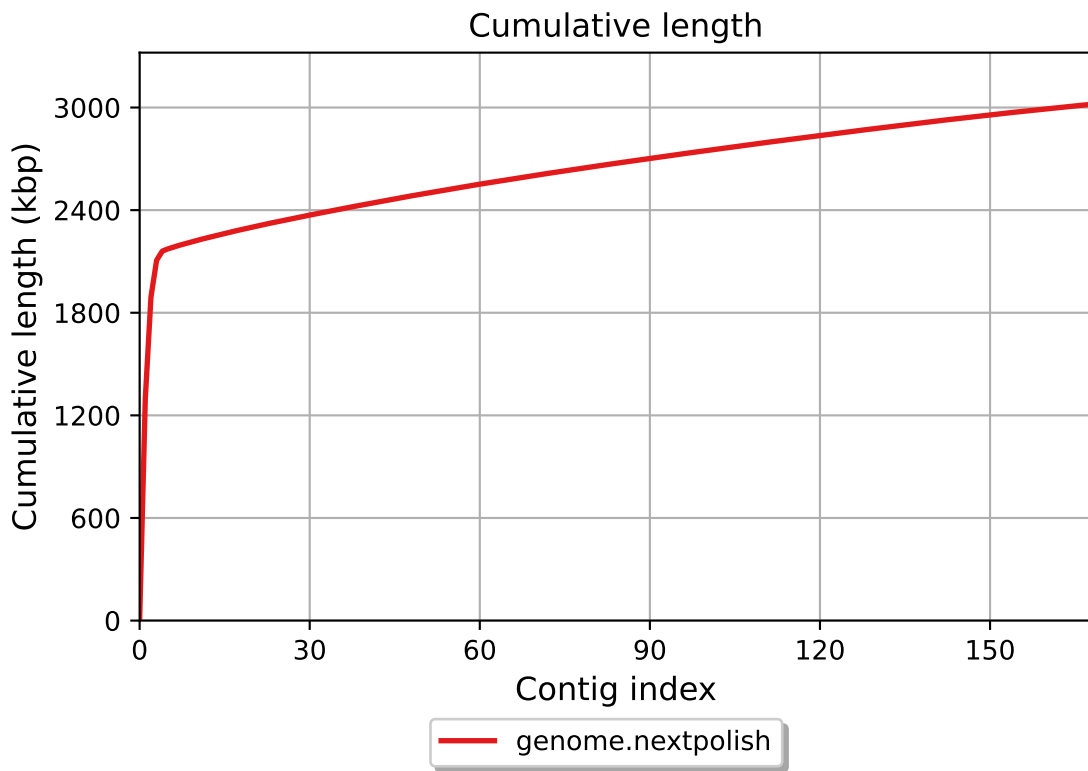


Report

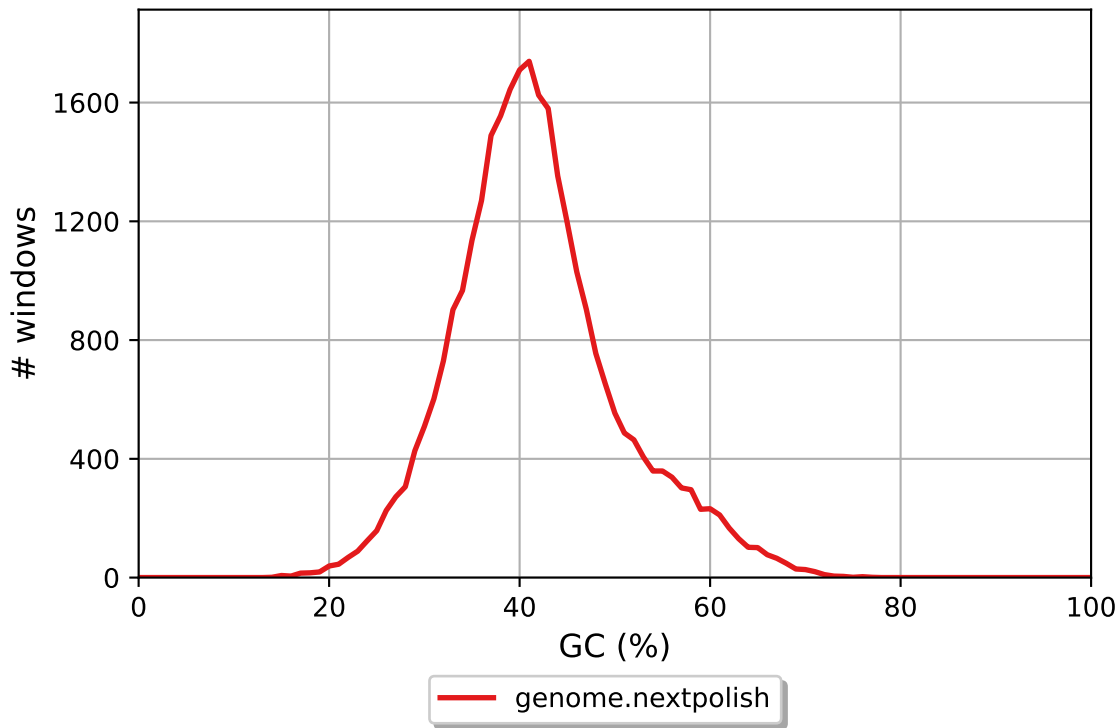
	genome.nextpolish
# contigs (>= 0 bp)	168
# contigs (>= 1000 bp)	168
# contigs (>= 5000 bp)	76
# contigs (>= 10000 bp)	6
# contigs (>= 25000 bp)	4
# contigs (>= 50000 bp)	4
Total length (>= 0 bp)	3019046
Total length (>= 1000 bp)	3019046
Total length (>= 5000 bp)	2634973
Total length (>= 10000 bp)	2184603
Total length (>= 25000 bp)	2160438
Total length (>= 50000 bp)	2160438
# contigs	168
Largest contig	1296885
Total length	3019046
GC (%)	41.76
N50	595496
N90	4630
auN	692395.4
L50	2
L90	94
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	14 + 1 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).





GC content



genome.nextpolish GC content

