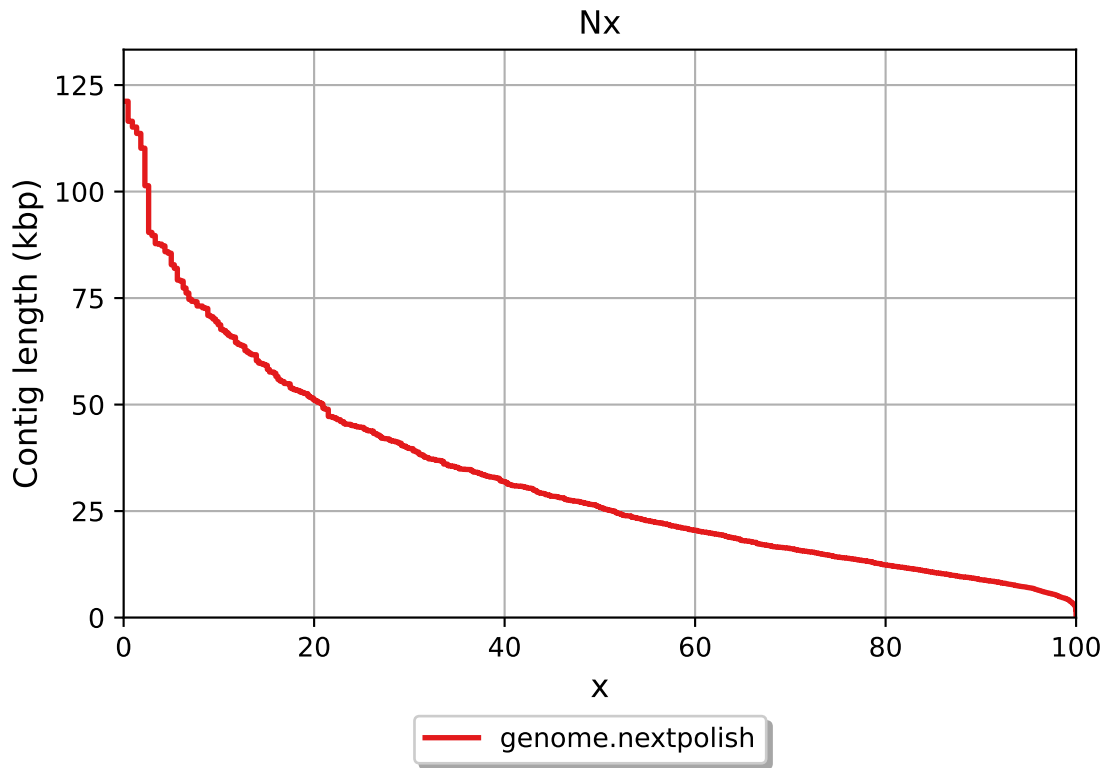


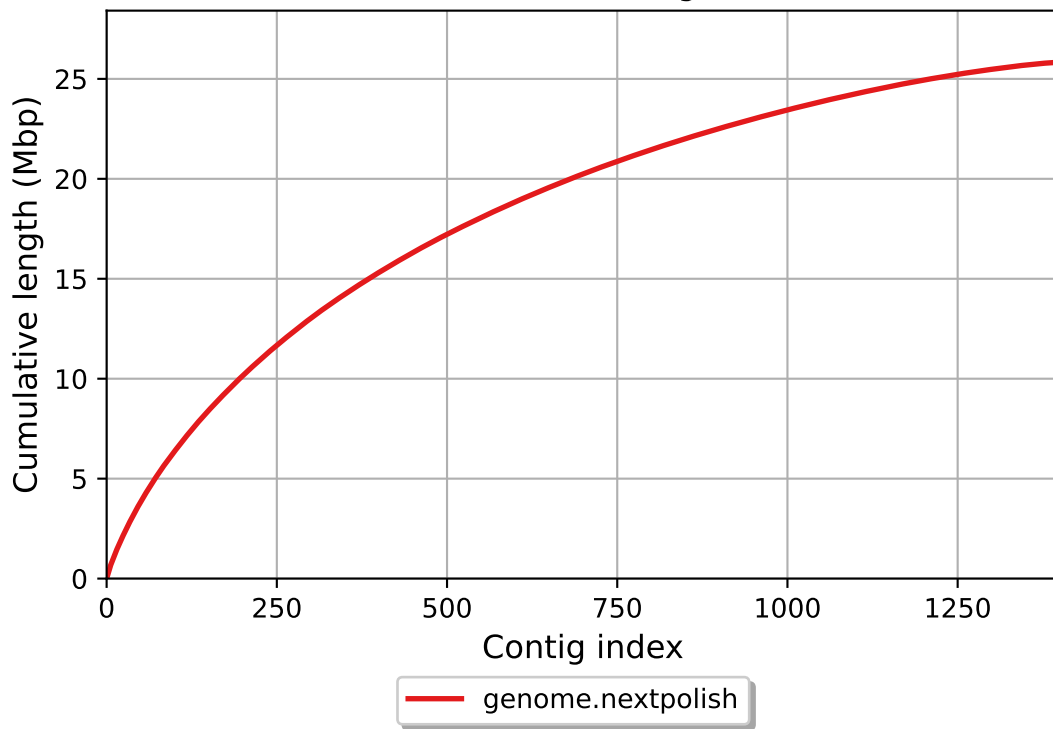
Report

	genome.nextpolish
# contigs (>= 0 bp)	1399
# contigs (>= 1000 bp)	1395
# contigs (>= 5000 bp)	1276
# contigs (>= 10000 bp)	889
# contigs (>= 25000 bp)	310
# contigs (>= 50000 bp)	79
Total length (>= 0 bp)	25831273
Total length (>= 1000 bp)	25828351
Total length (>= 5000 bp)	25364110
Total length (>= 10000 bp)	22411303
Total length (>= 25000 bp)	13280899
Total length (>= 50000 bp)	5403873
# contigs	1399
Largest contig	121197
Total length	25831273
GC (%)	43.98
N50	25932
N90	8887
auN	32885.8
L50	296
L90	978
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	67 + 8 part

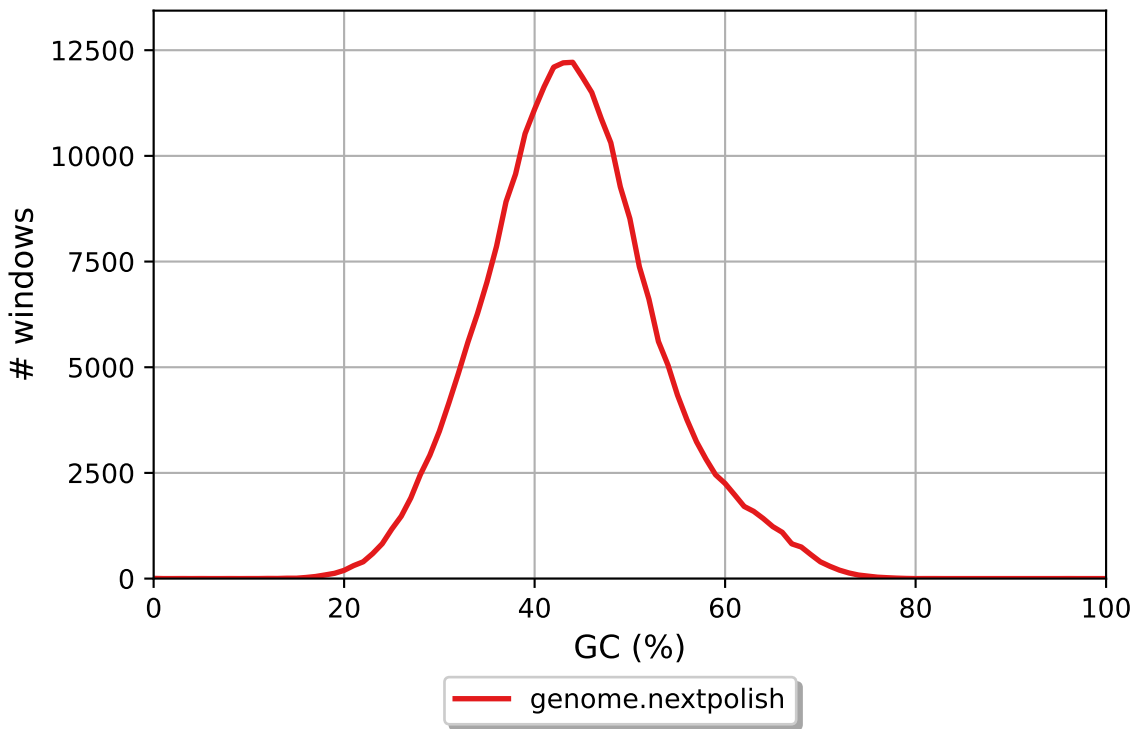
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

