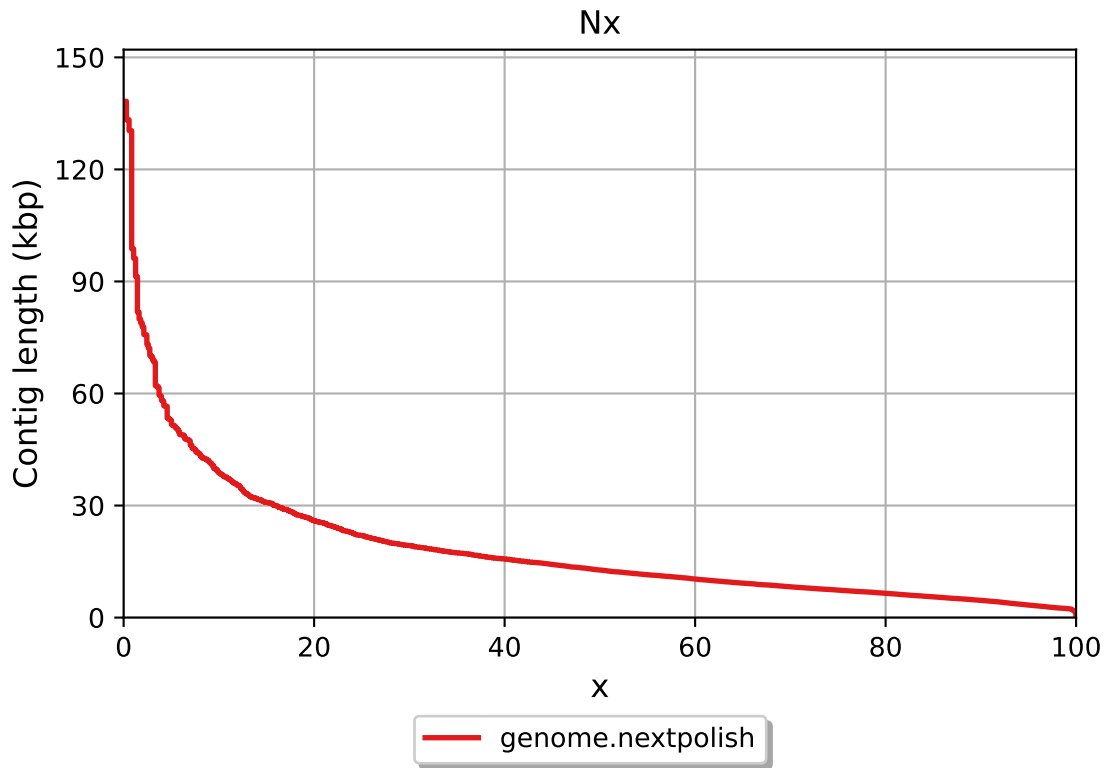


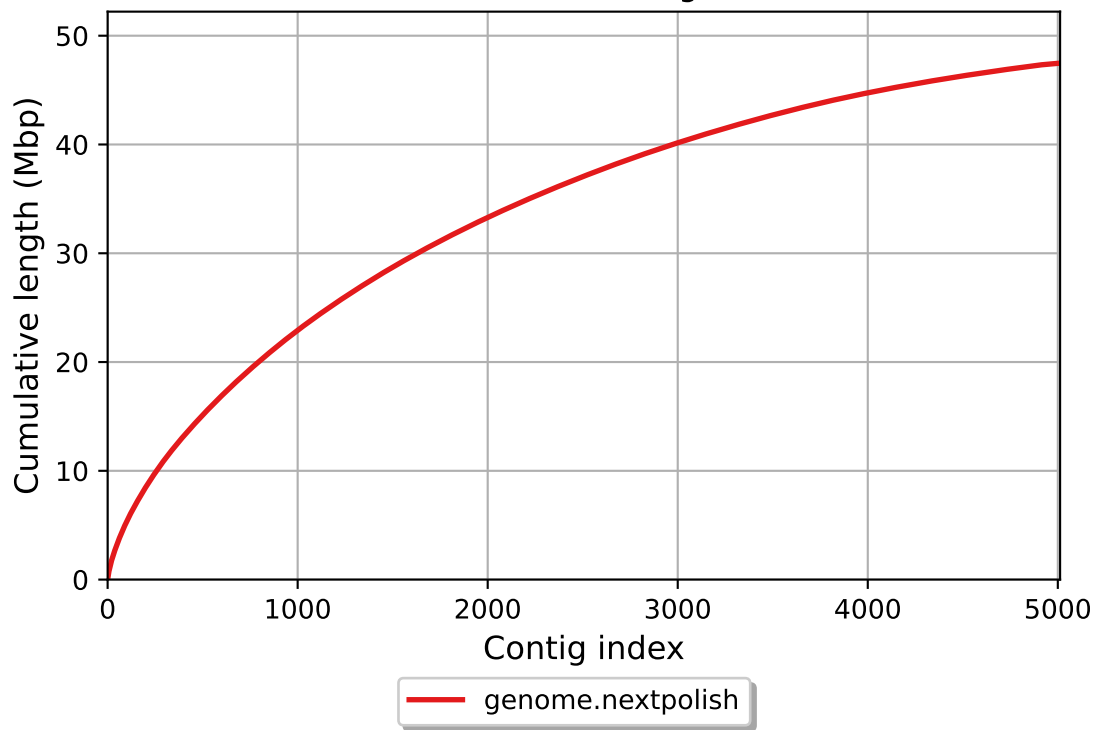
Report

	genome.nextpolish
# contigs (>= 0 bp)	5012
# contigs (>= 1000 bp)	4997
# contigs (>= 5000 bp)	3312
# contigs (>= 10000 bp)	1541
# contigs (>= 25000 bp)	262
# contigs (>= 50000 bp)	39
Total length (>= 0 bp)	47466536
Total length (>= 1000 bp)	47455952
Total length (>= 5000 bp)	41807377
Total length (>= 10000 bp)	29135482
Total length (>= 25000 bp)	10117514
Total length (>= 50000 bp)	2740958
# contigs	5012
Largest contig	138246
Total length	47466536
GC (%)	43.83
N50	12721
N90	4612
auN	18566.1
L50	1063
L90	3502
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	48 + 14 part

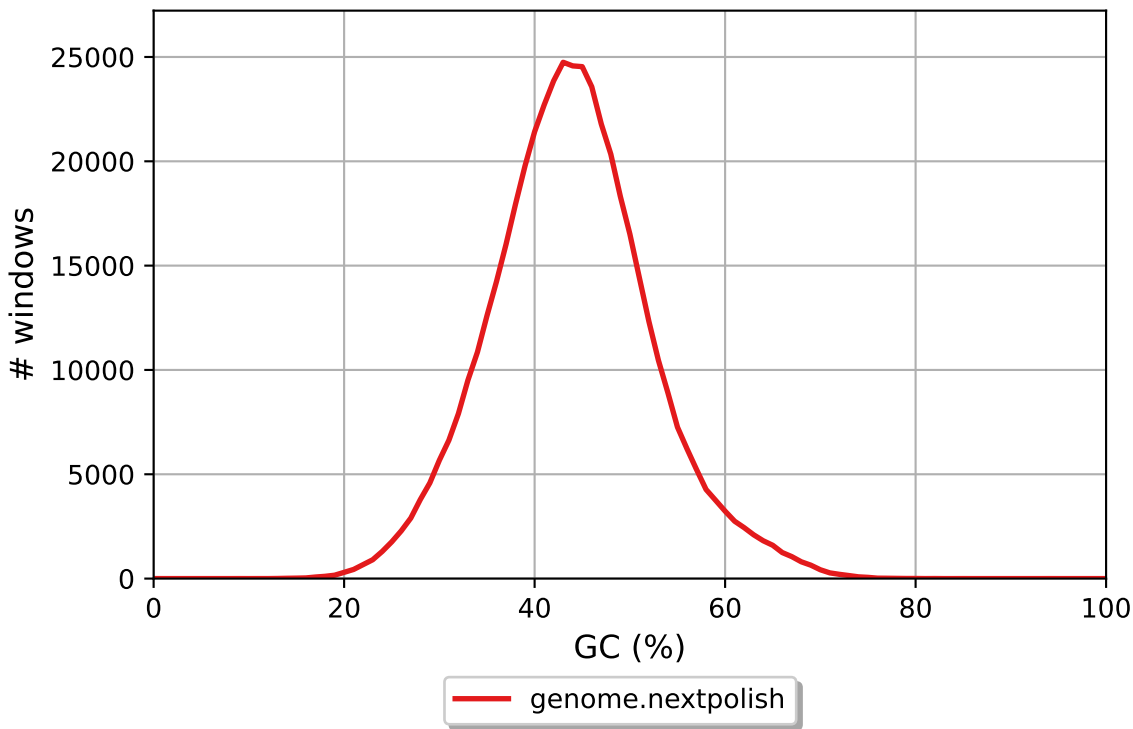
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

