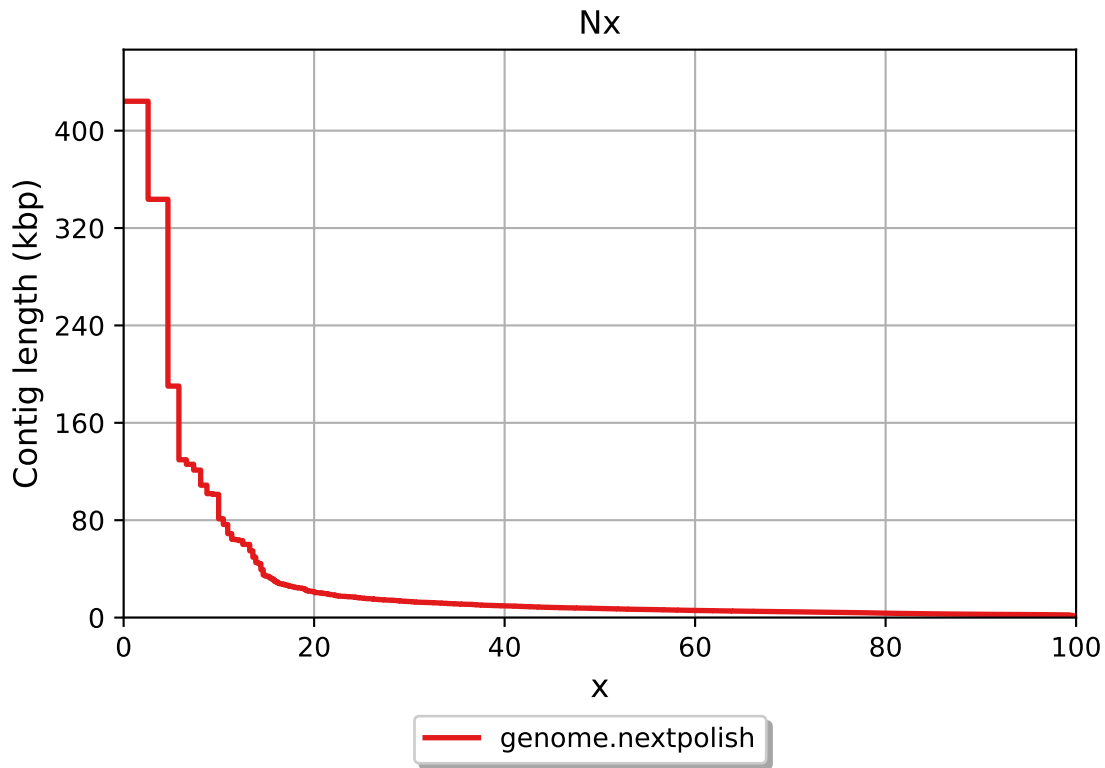


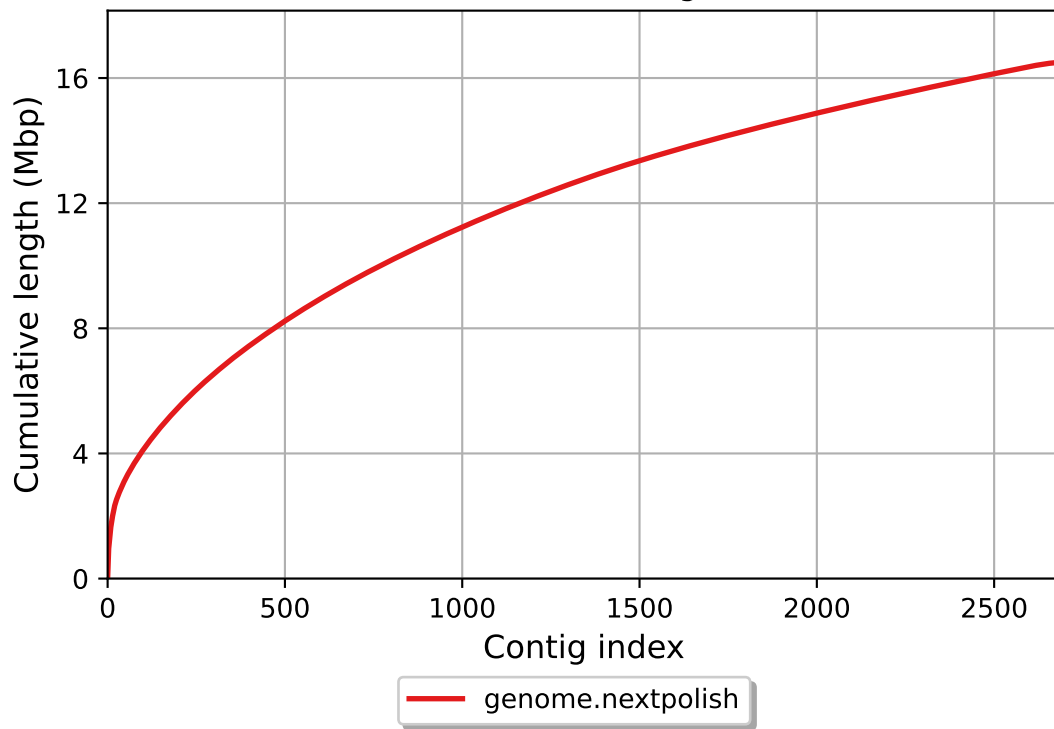
Report

	genome.nextpolish
# contigs (>= 0 bp)	2686
# contigs (>= 1000 bp)	2677
# contigs (>= 5000 bp)	985
# contigs (>= 10000 bp)	279
# contigs (>= 25000 bp)	41
# contigs (>= 50000 bp)	18
Total length (>= 0 bp)	16503761
Total length (>= 1000 bp)	16497668
Total length (>= 5000 bp)	11161227
Total length (>= 10000 bp)	6340435
Total length (>= 25000 bp)	2968372
Total length (>= 50000 bp)	2239621
# contigs	2686
Largest contig	424144
Total length	16503761
GC (%)	41.18
N50	7483
N90	2724
auN	35192.2
L50	503
L90	1992
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	28 + 2 part

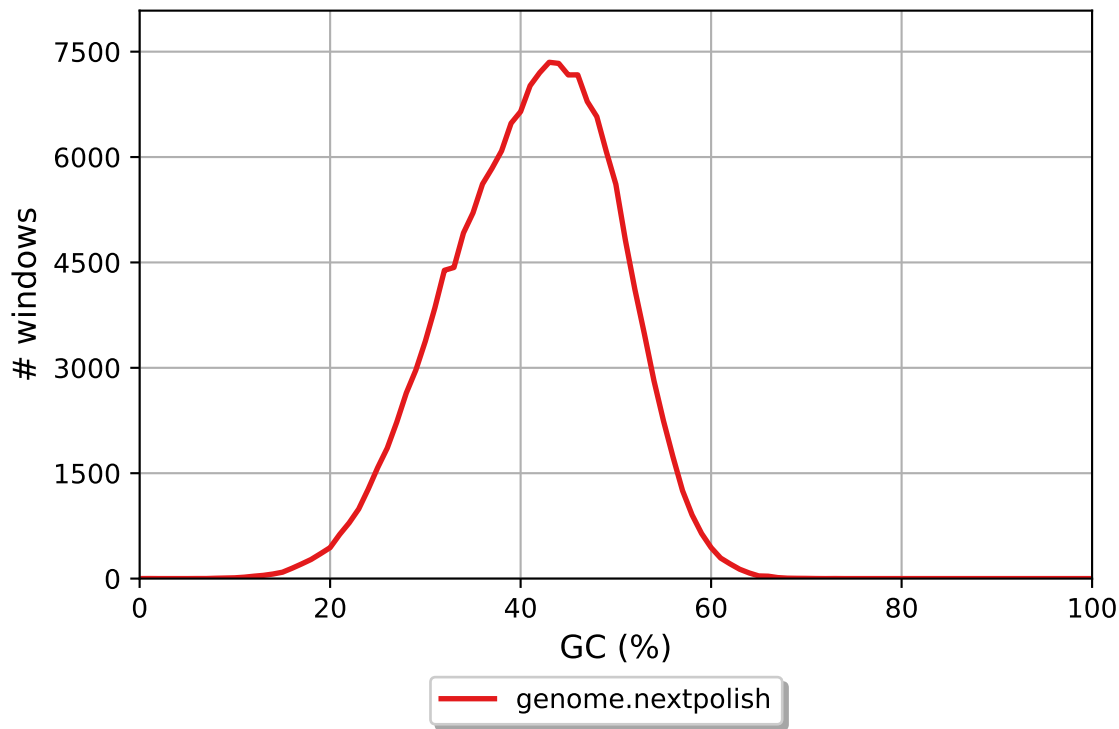
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

