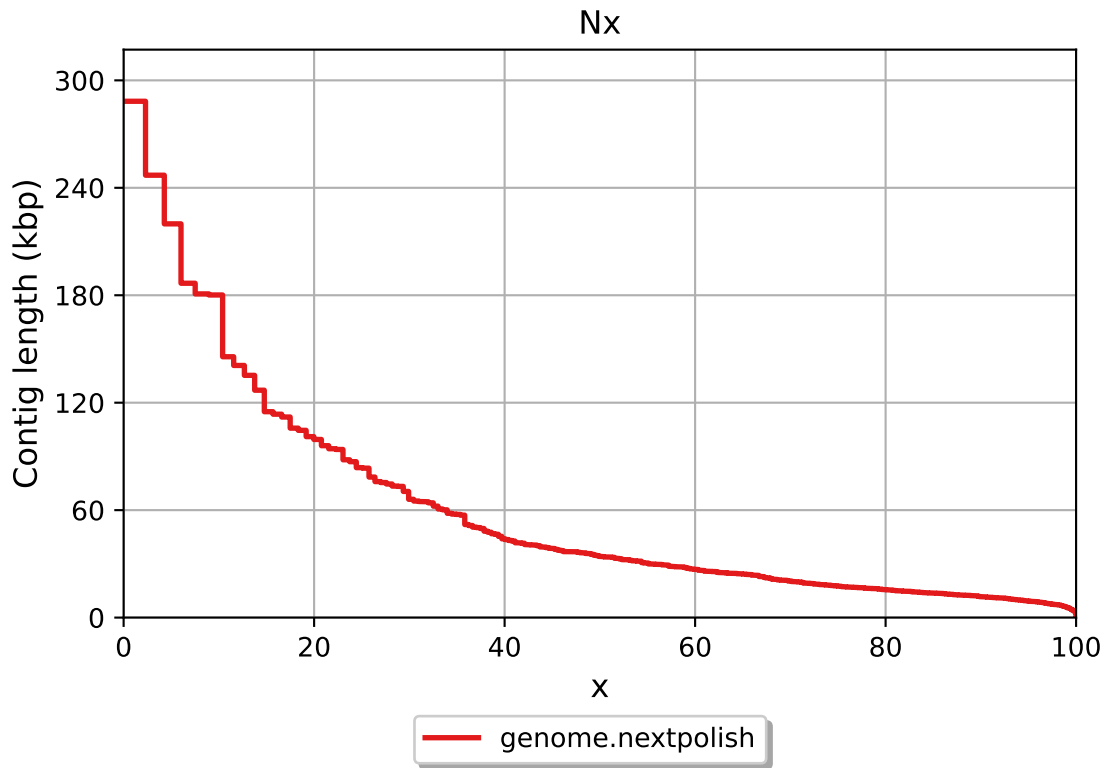


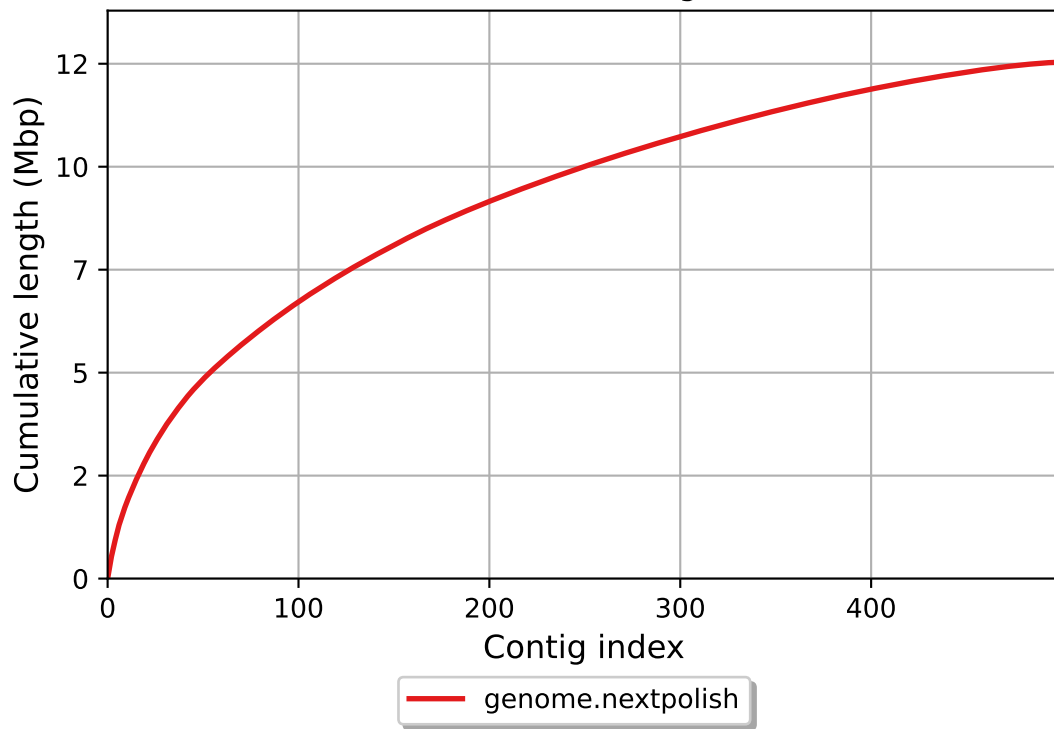
Report

	genome.nextpolish
# contigs (>= 0 bp)	499
# contigs (>= 1000 bp)	498
# contigs (>= 5000 bp)	477
# contigs (>= 10000 bp)	386
# contigs (>= 25000 bp)	143
# contigs (>= 50000 bp)	47
Total length (>= 0 bp)	12535777
Total length (>= 1000 bp)	12534870
Total length (>= 5000 bp)	12462504
Total length (>= 10000 bp)	11748096
Total length (>= 25000 bp)	7922816
Total length (>= 50000 bp)	4693731
# contigs	499
Largest contig	288322
Total length	12535777
GC (%)	46.43
N50	34173
N90	11691
auN	62422.3
L50	87
L90	344
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	34 + 8 part

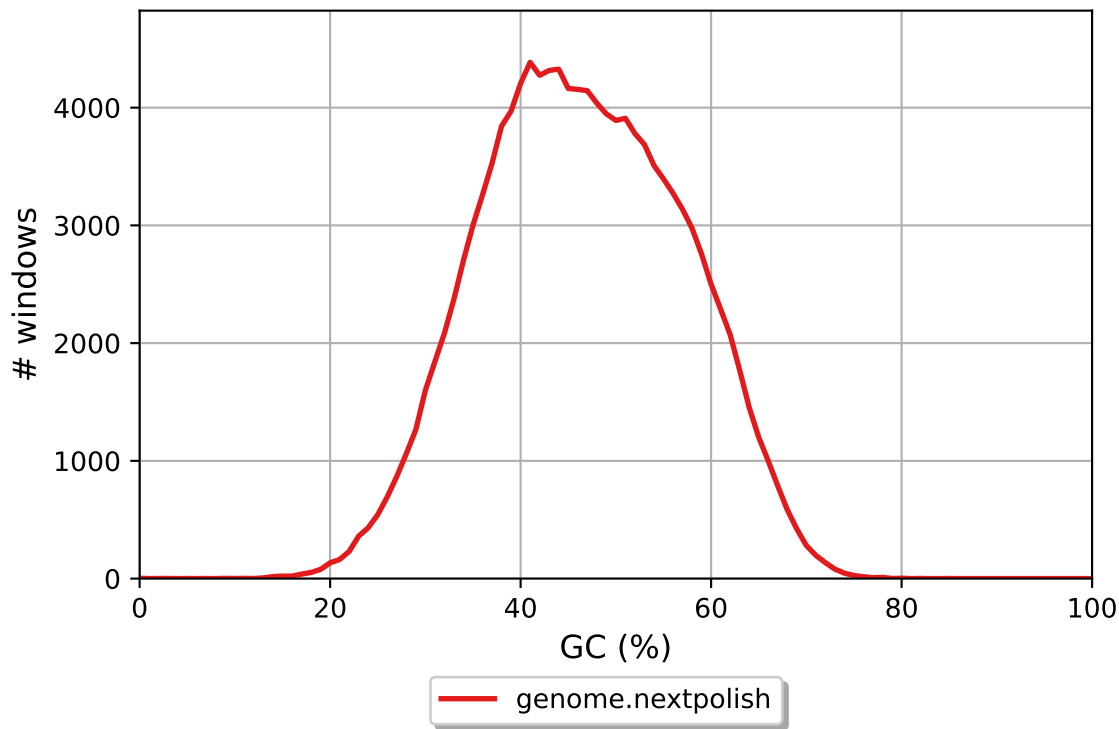
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

