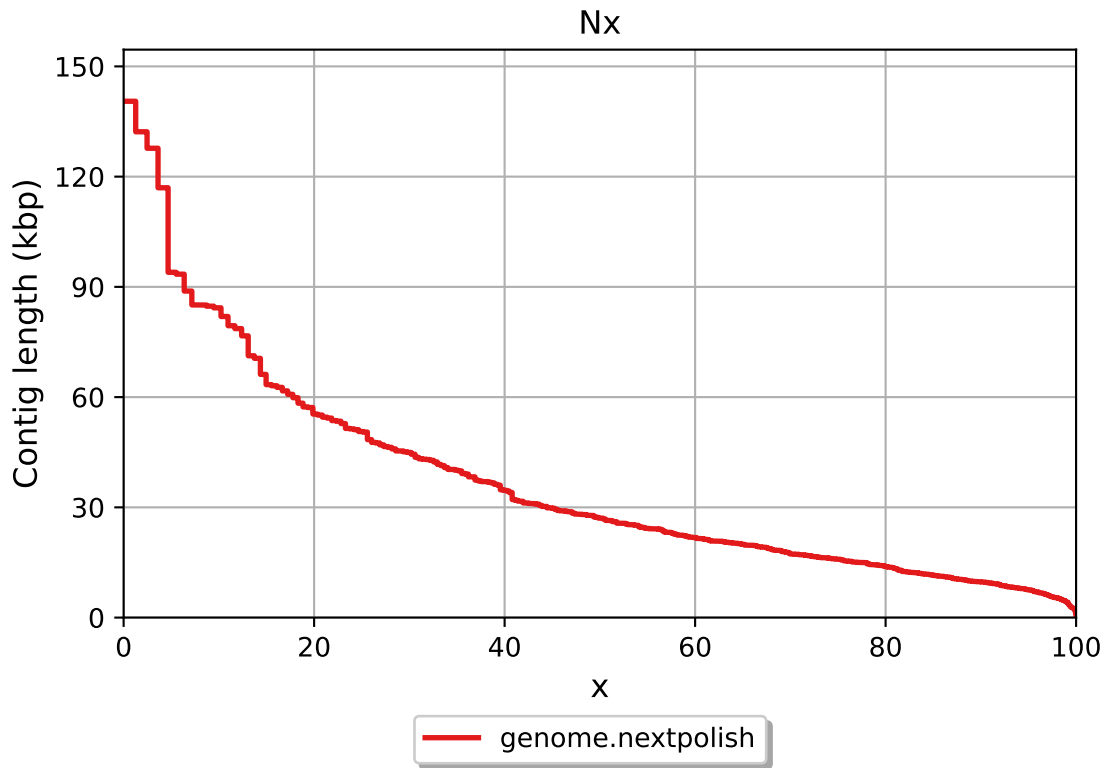


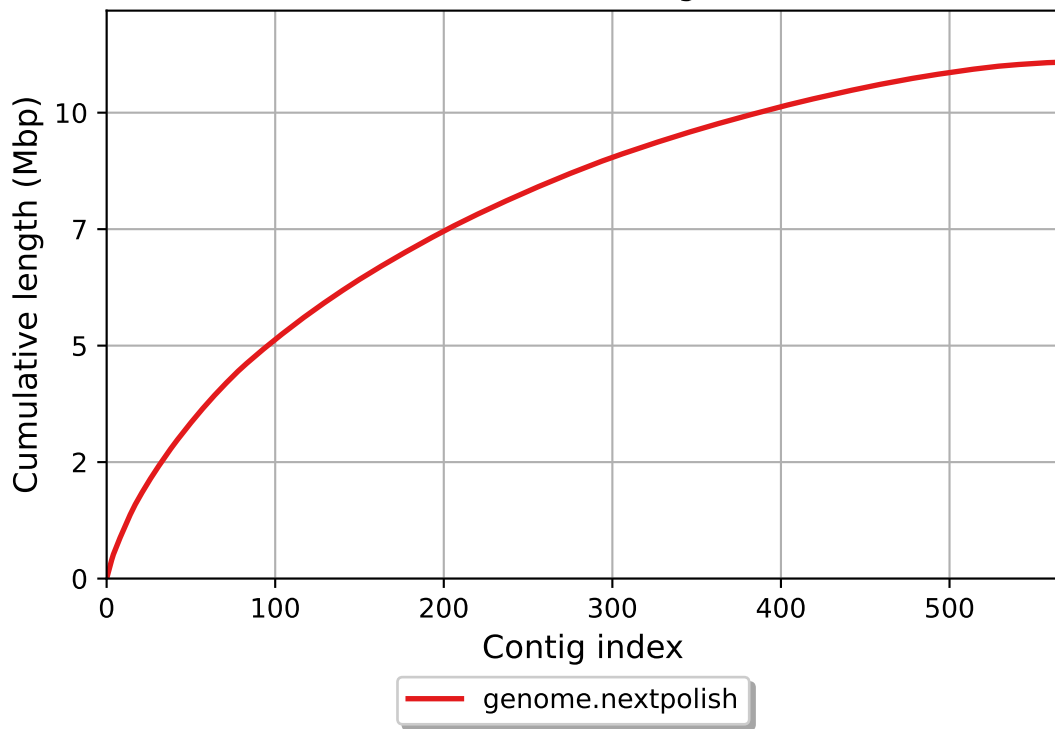
Report

	genome.nextpolish
# contigs (>= 0 bp)	565
# contigs (>= 1000 bp)	564
# contigs (>= 5000 bp)	508
# contigs (>= 10000 bp)	369
# contigs (>= 25000 bp)	132
# contigs (>= 50000 bp)	39
Total length (>= 0 bp)	11081984
Total length (>= 1000 bp)	11081394
Total length (>= 5000 bp)	10903807
Total length (>= 10000 bp)	9828375
Total length (>= 25000 bp)	5991686
Total length (>= 50000 bp)	2836119
# contigs	565
Largest contig	140509
Total length	11081984
GC (%)	48.14
N50	27067
N90	9731
auN	37417.0
L50	115
L90	384
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	35 + 3 part

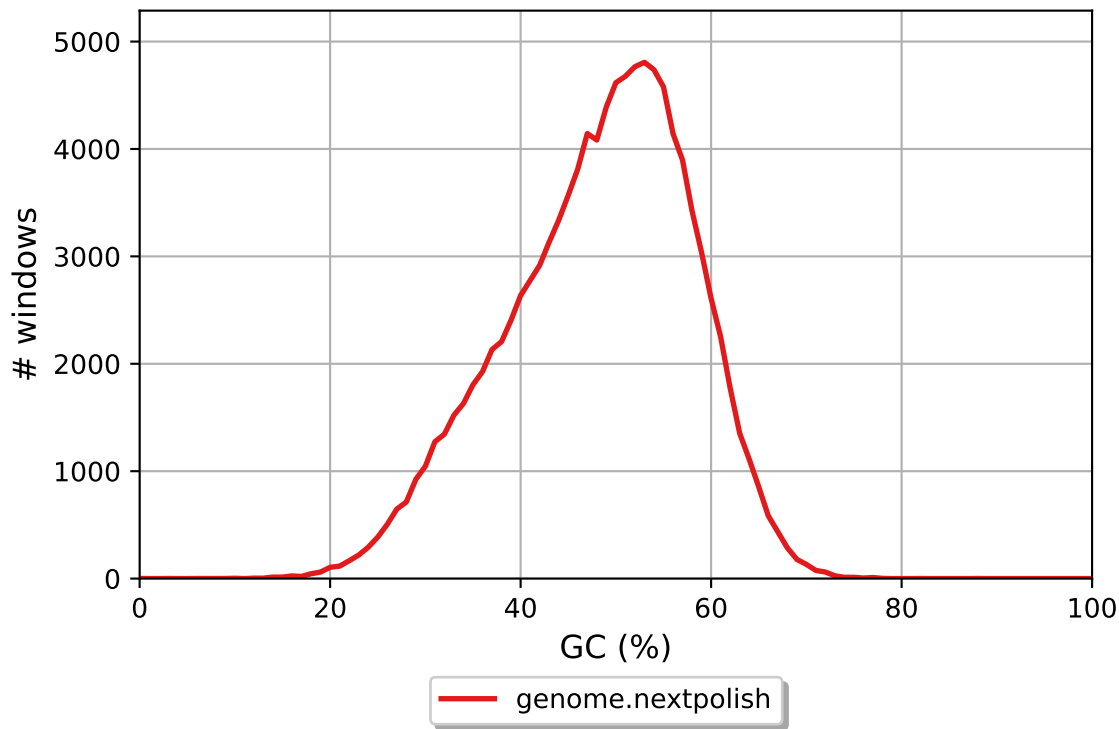
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

