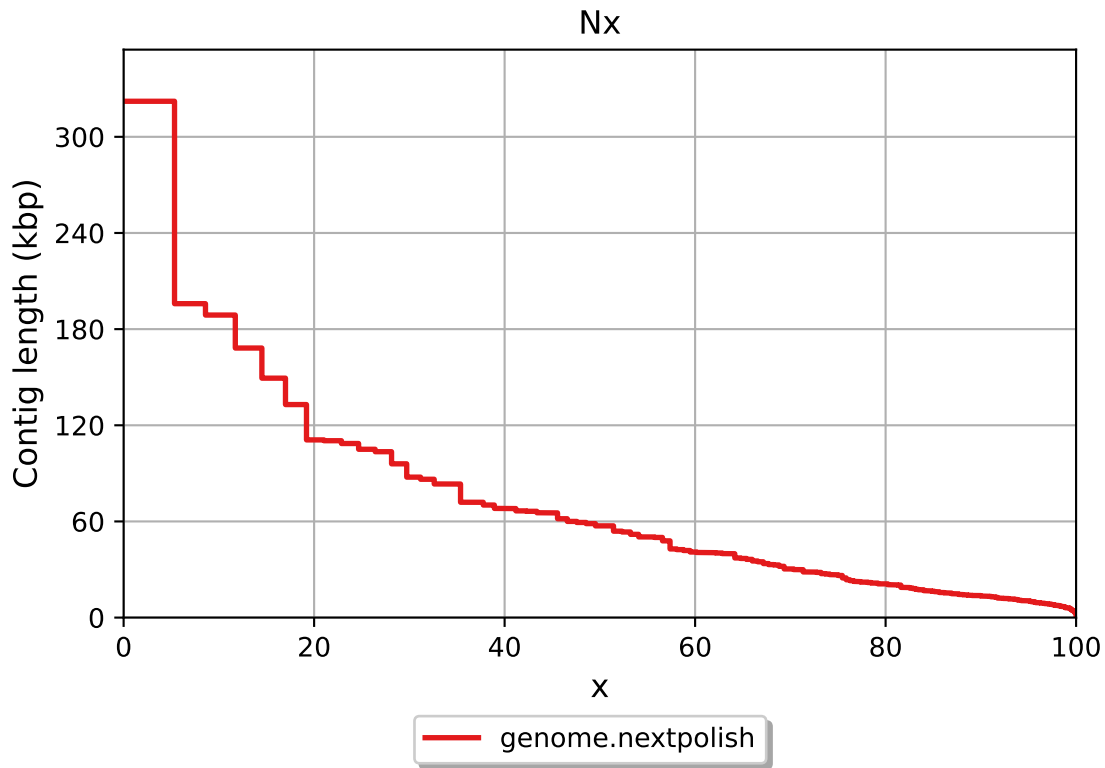


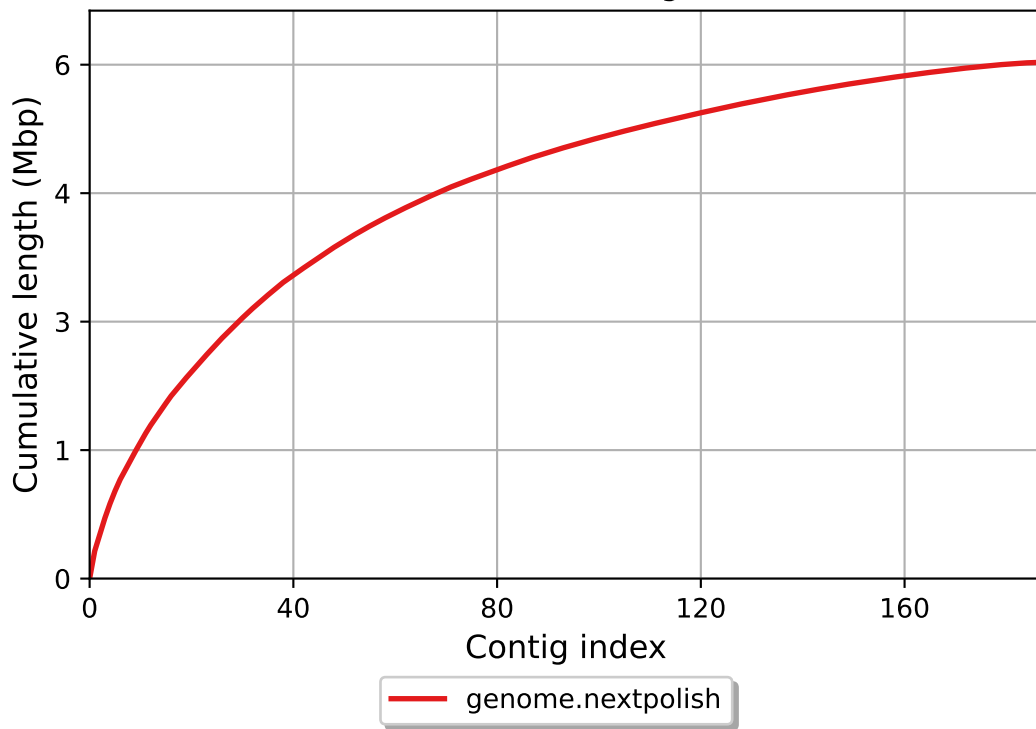
Report

	genome.nextpolish
# contigs (>= 0 bp)	187
# contigs (>= 1000 bp)	186
# contigs (>= 5000 bp)	178
# contigs (>= 10000 bp)	146
# contigs (>= 25000 bp)	70
# contigs (>= 50000 bp)	37
Total length (>= 0 bp)	6028344
Total length (>= 1000 bp)	6027756
Total length (>= 5000 bp)	5996133
Total length (>= 10000 bp)	5744256
Total length (>= 25000 bp)	4548846
Total length (>= 50000 bp)	3410485
# contigs	187
Largest contig	322186
Total length	6028344
GC (%)	41.85
N50	57240
N90	13441
auN	80254.0
L50	30
L90	120
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	13 + 0 part

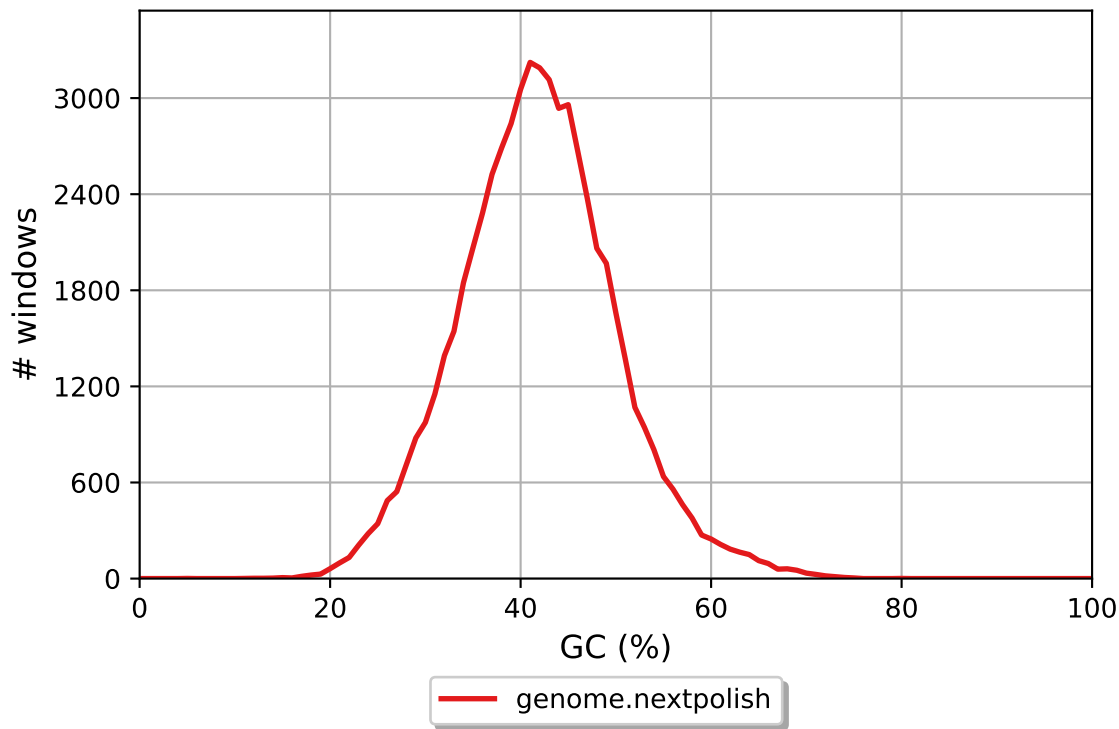
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

