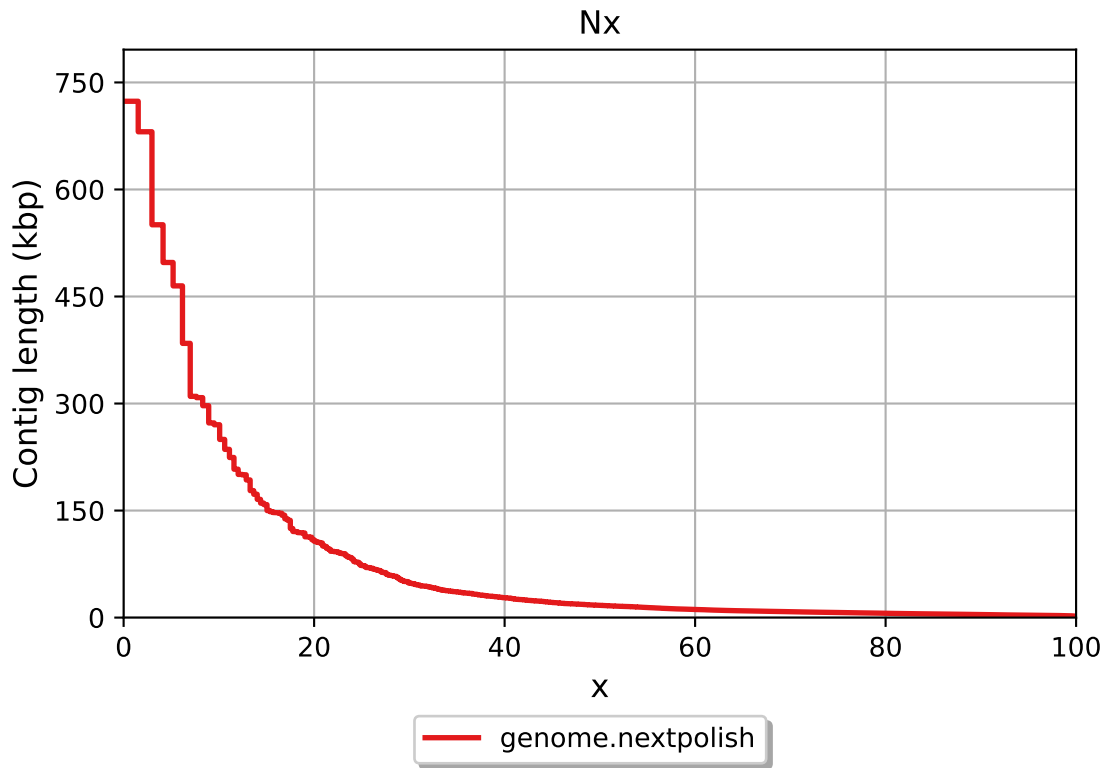


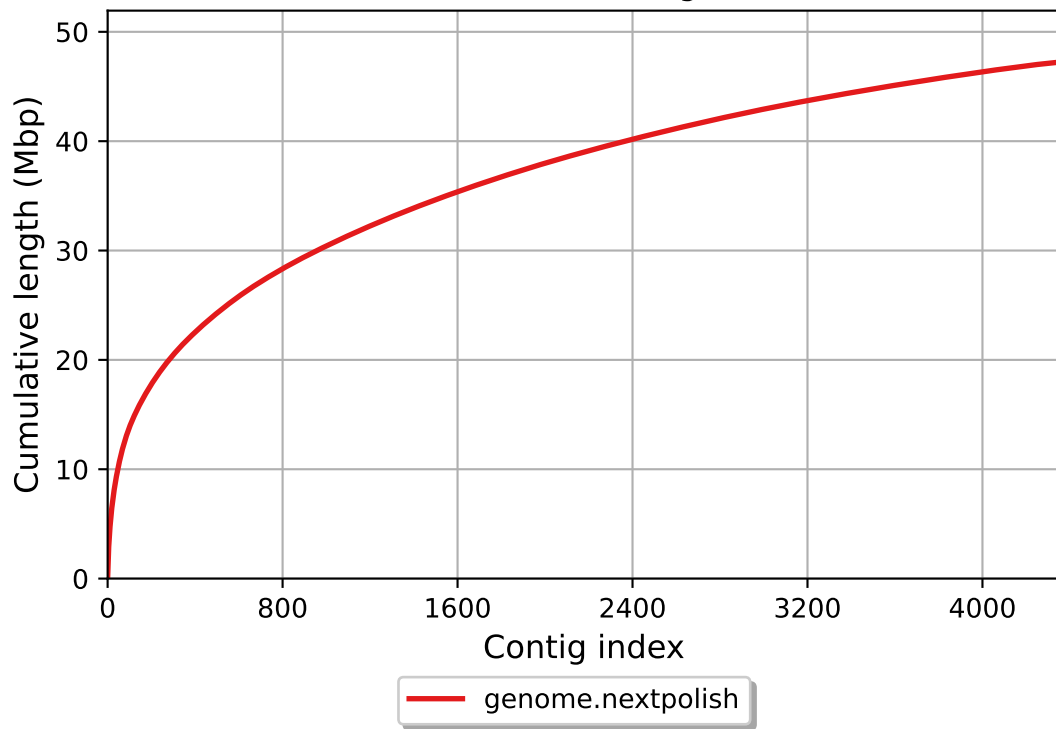
Report

	genome.nextpolish
# contigs (>= 0 bp)	4356
# contigs (>= 1000 bp)	4343
# contigs (>= 5000 bp)	2455
# contigs (>= 10000 bp)	945
# contigs (>= 25000 bp)	273
# contigs (>= 50000 bp)	105
Total length (>= 0 bp)	47216659
Total length (>= 1000 bp)	47206607
Total length (>= 5000 bp)	40456036
Total length (>= 10000 bp)	29878827
Total length (>= 25000 bp)	19811716
Total length (>= 50000 bp)	14115050
# contigs	4355
Largest contig	723617
Total length	47216243
GC (%)	39.65
N50	17147
N90	4262
auN	83144.7
L50	460
L90	2896
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	68 + 9 part

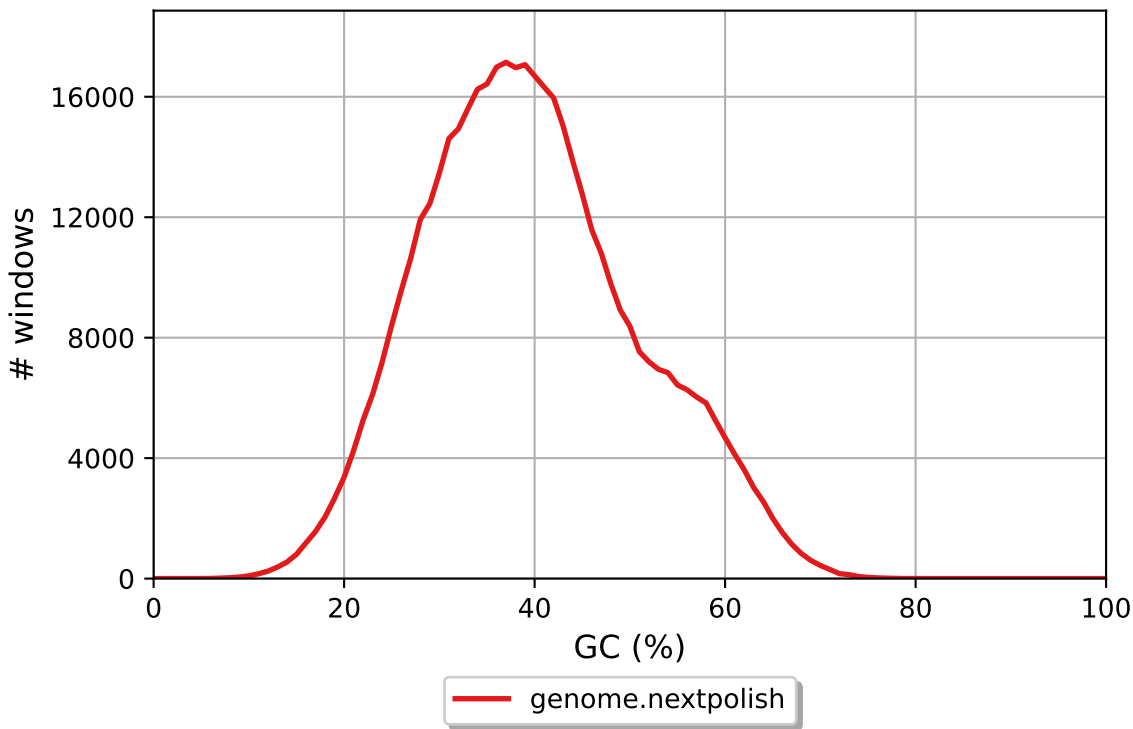
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

