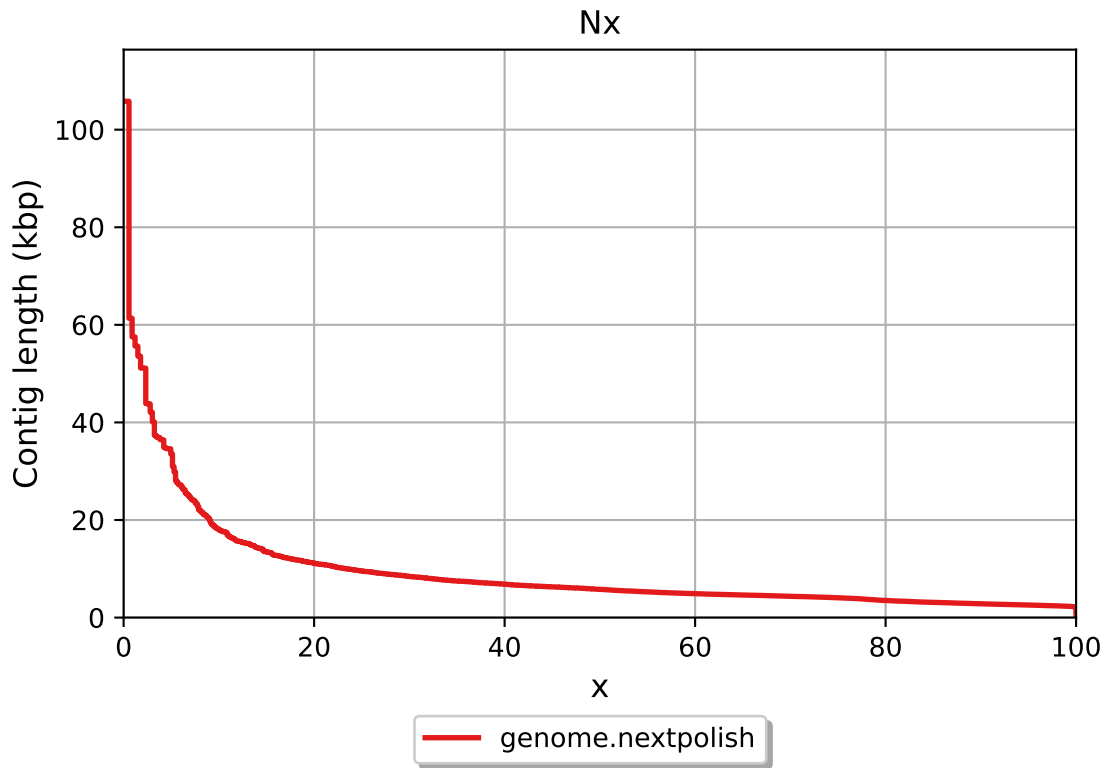


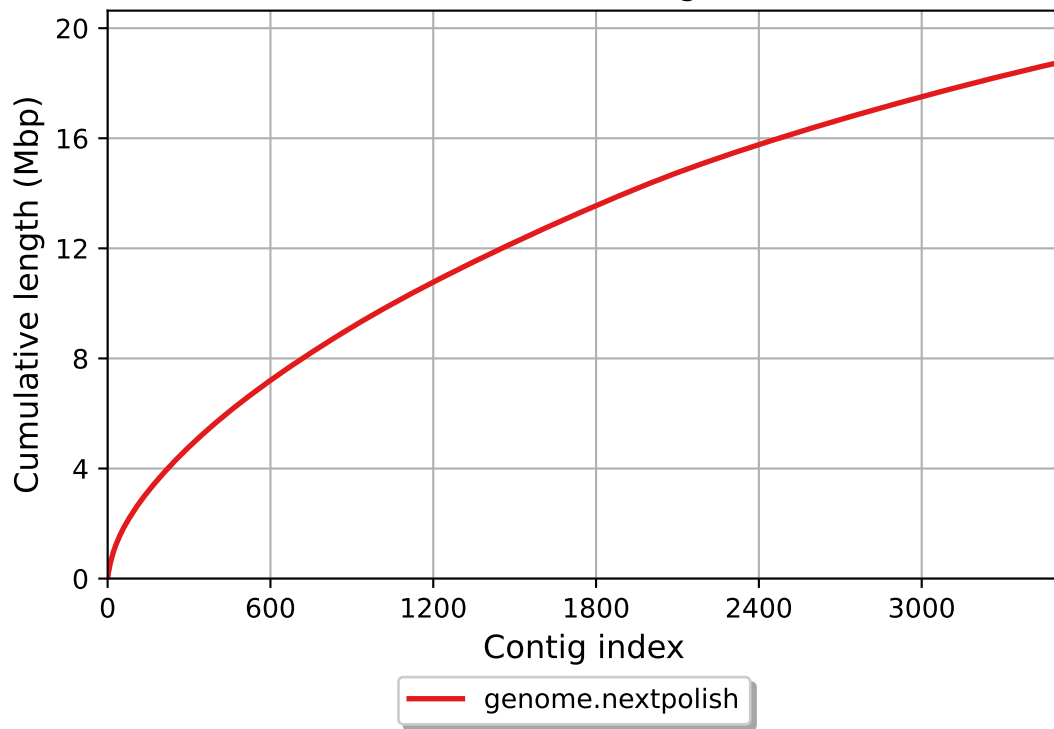
Report

	genome.nextpolish
# contigs (>= 0 bp)	3510
# contigs (>= 1000 bp)	3509
# contigs (>= 5000 bp)	1243
# contigs (>= 10000 bp)	257
# contigs (>= 25000 bp)	33
# contigs (>= 50000 bp)	7
Total length (>= 0 bp)	18761667
Total length (>= 1000 bp)	18761090
Total length (>= 5000 bp)	10989708
Total length (>= 10000 bp)	4369473
Total length (>= 25000 bp)	1288535
Total length (>= 50000 bp)	436182
# contigs	3510
Largest contig	105820
Total length	18761667
GC (%)	45.82
N50	5796
N90	2830
auN	9477.6
L50	942
L90	2773
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	33 + 6 part

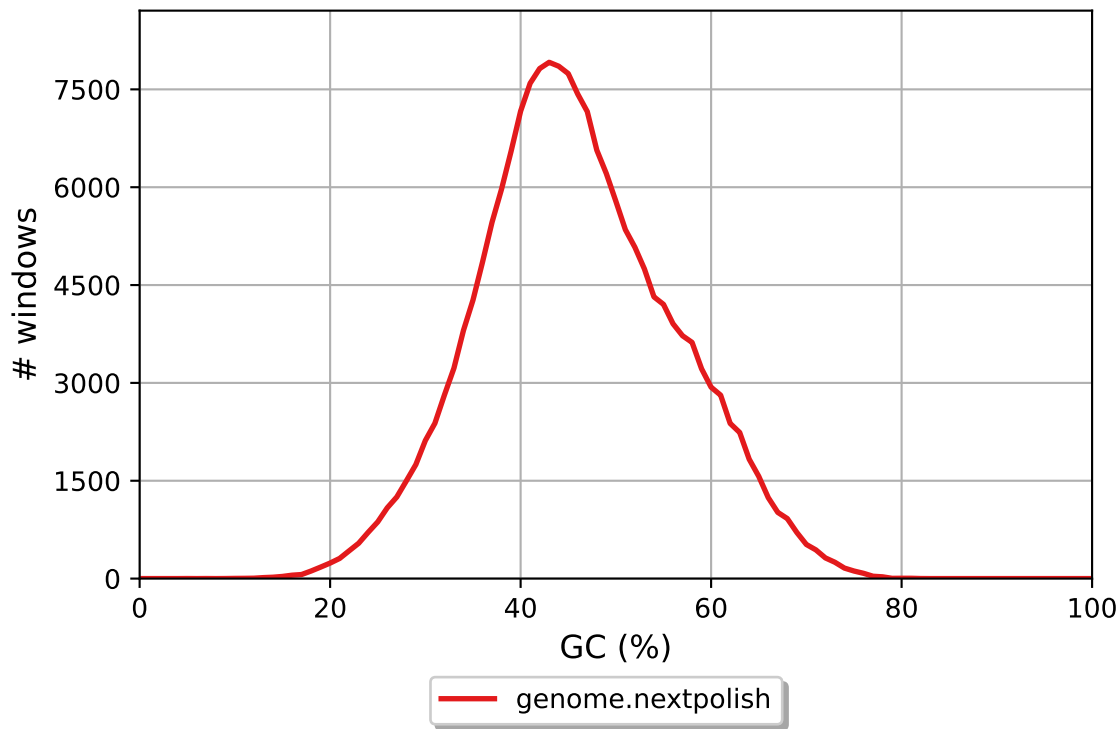
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

