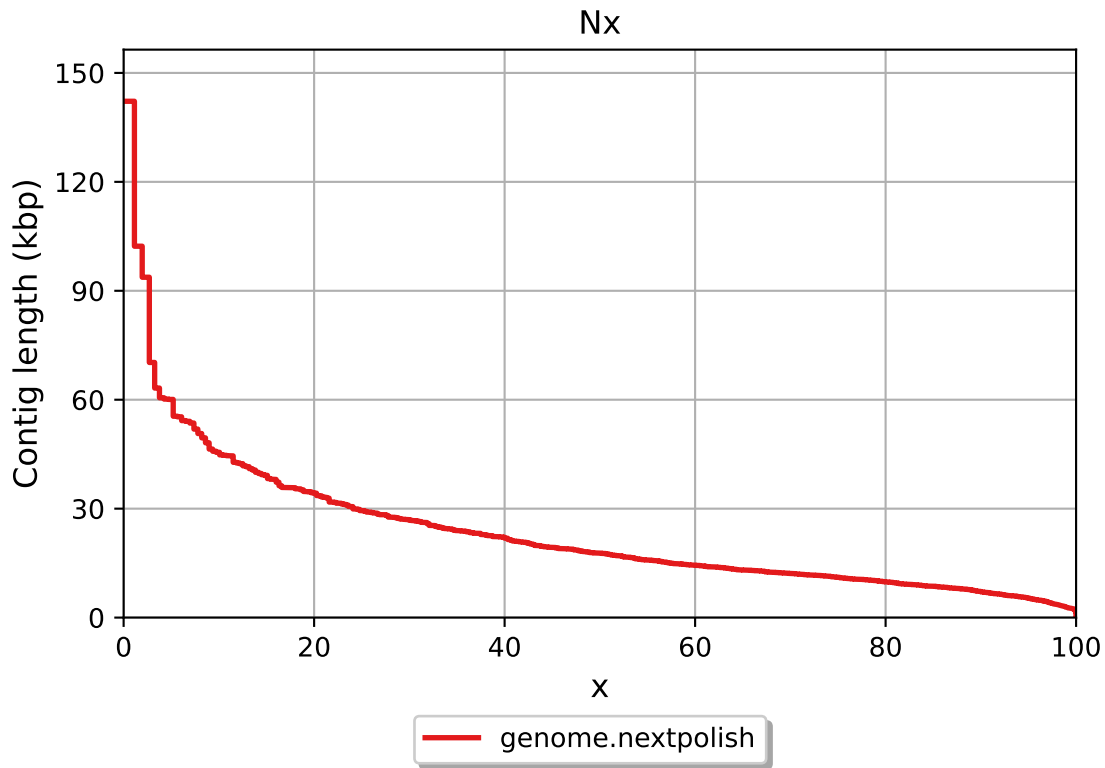


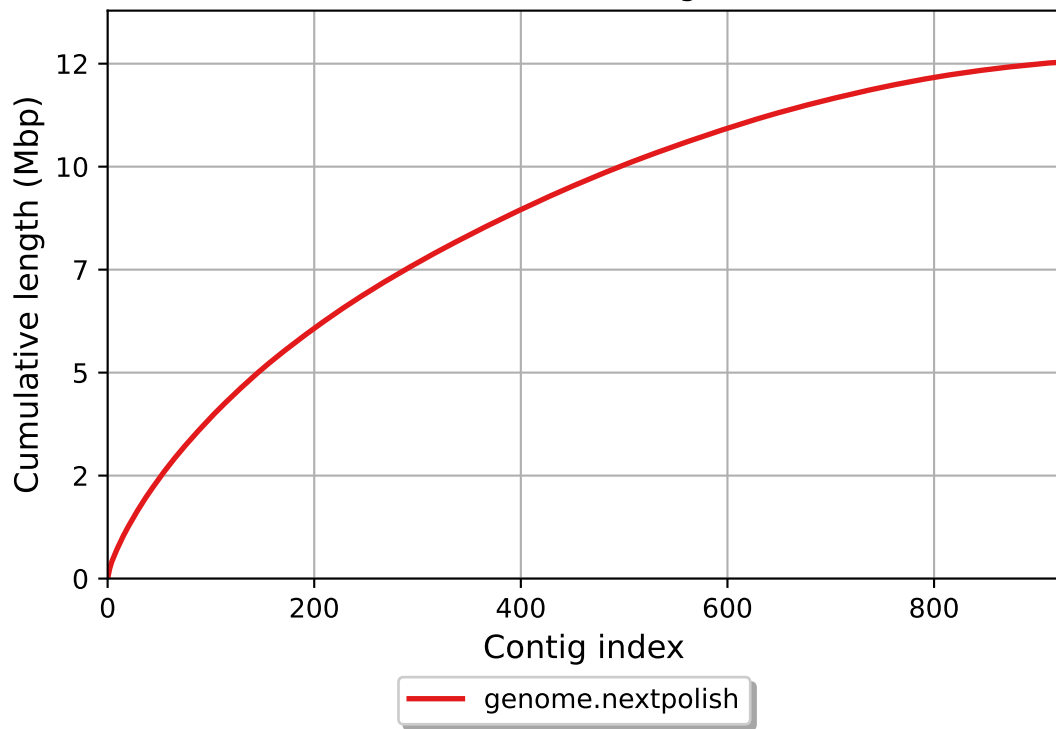
Report

	genome.nextpolish
# contigs (>= 0 bp)	922
# contigs (>= 1000 bp)	921
# contigs (>= 5000 bp)	763
# contigs (>= 10000 bp)	493
# contigs (>= 25000 bp)	109
# contigs (>= 50000 bp)	15
Total length (>= 0 bp)	12534181
Total length (>= 1000 bp)	12533619
Total length (>= 5000 bp)	11995155
Total length (>= 10000 bp)	9974404
Total length (>= 25000 bp)	4145719
Total length (>= 50000 bp)	1027668
# contigs	922
Largest contig	142192
Total length	12534181
GC (%)	48.24
N50	17736
N90	7166
auN	23847.2
L50	211
L90	646
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	19 + 6 part

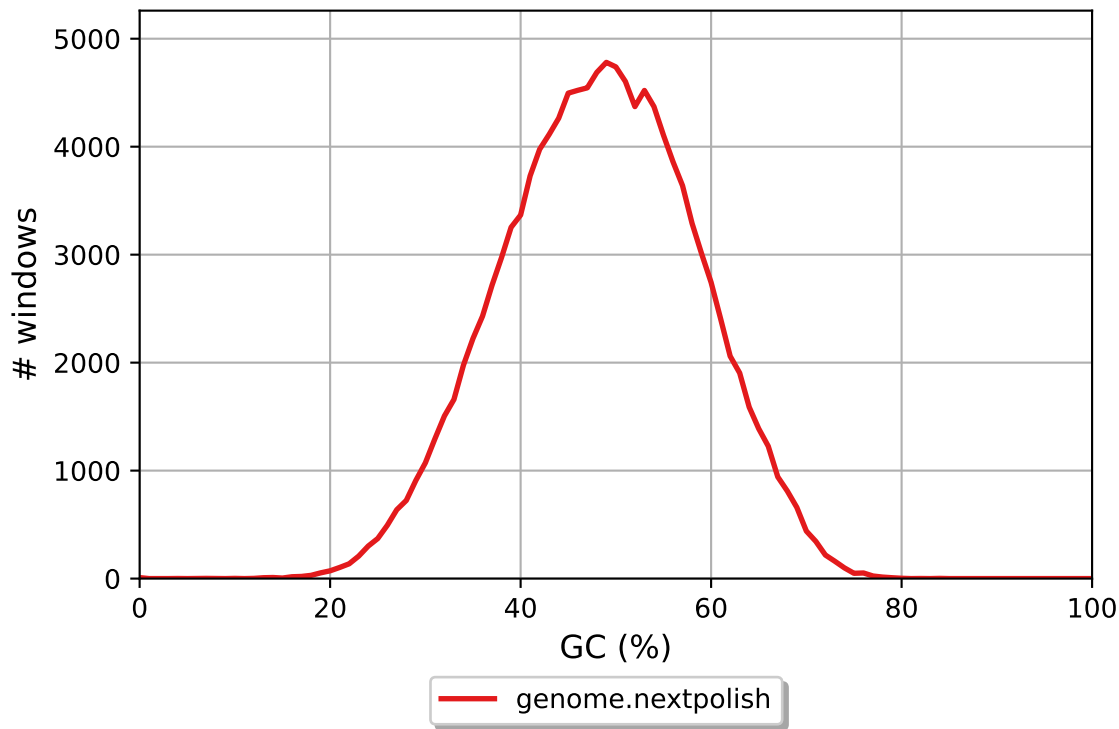
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



Cumulative length



GC content



genome.nextpolish GC content

