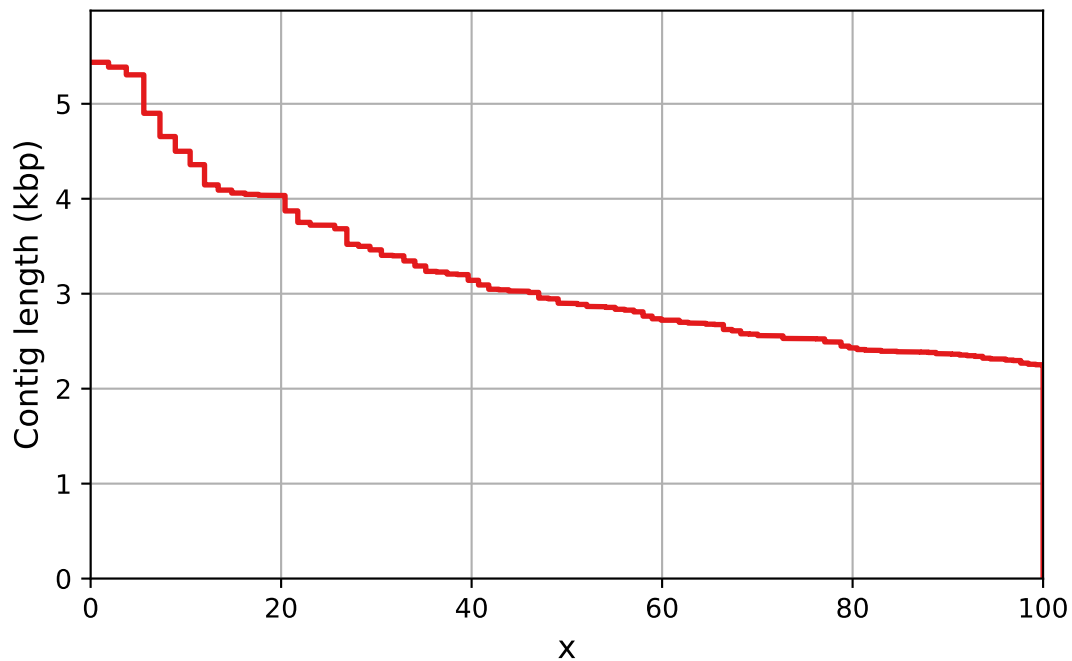


Report

	genome.nextpolish
# contigs (>= 0 bp)	96
# contigs (>= 1000 bp)	96
# contigs (>= 5000 bp)	3
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	288848
Total length (>= 1000 bp)	288848
Total length (>= 5000 bp)	16128
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	96
Largest contig	5437
Total length	288848
GC (%)	37.83
N50	2900
N90	2366
auN	3193.1
L50	39
L90	84
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 0 part

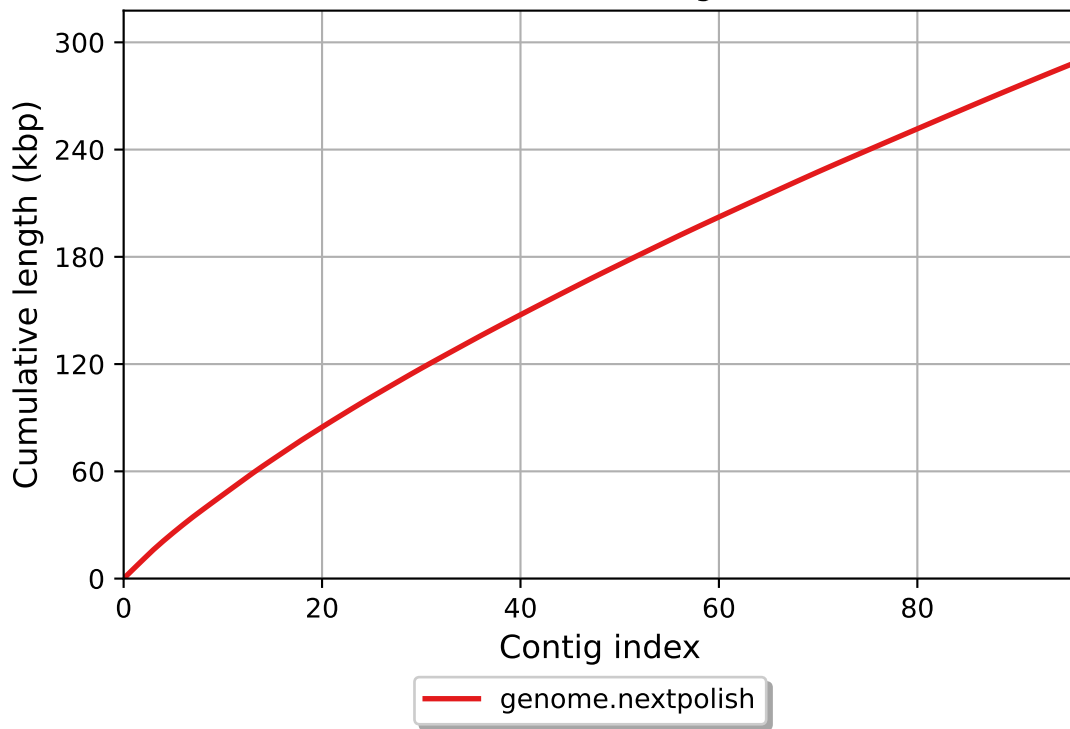
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

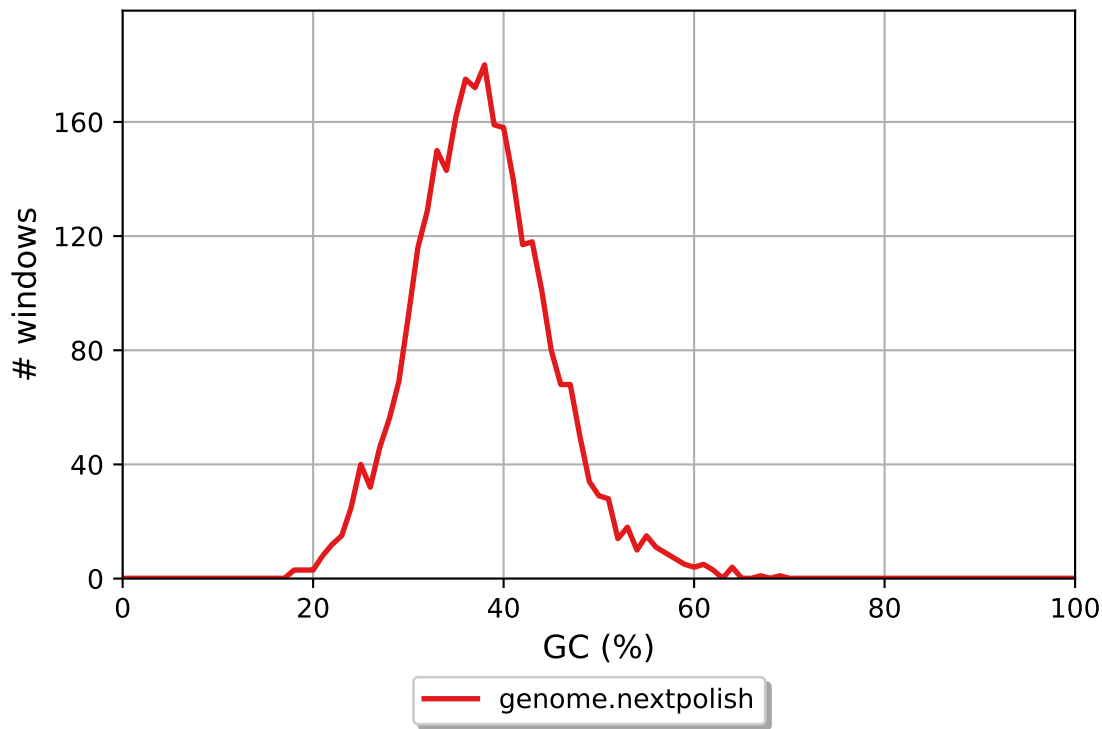


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

