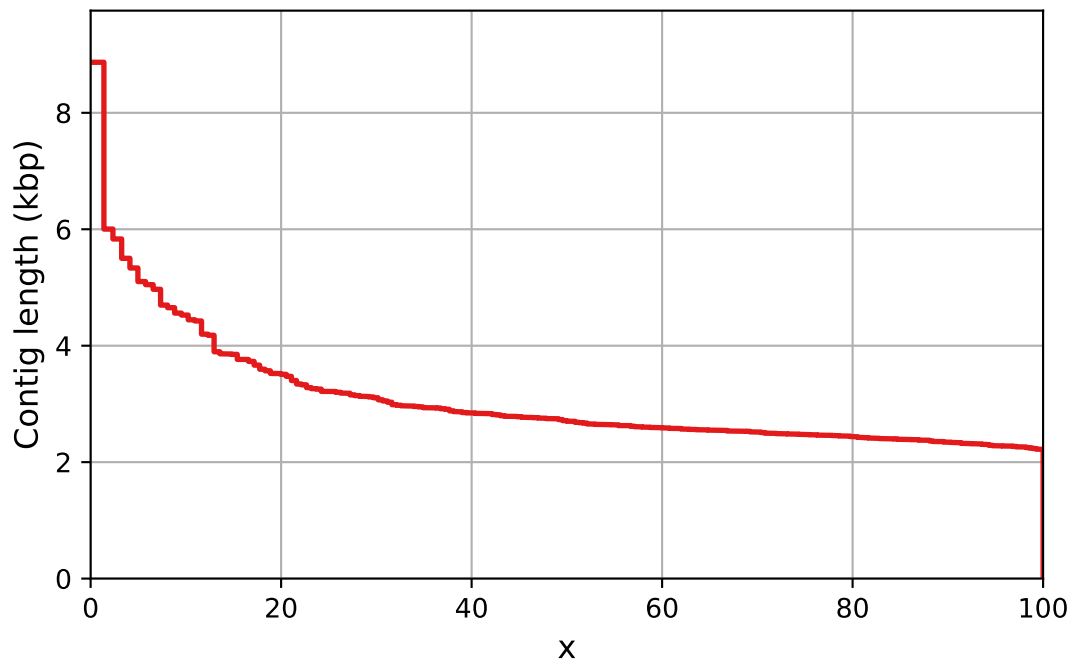


Report

	genome.nextpolish
# contigs (>= 0 bp)	221
# contigs (>= 1000 bp)	221
# contigs (>= 5000 bp)	7
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	635266
Total length (>= 1000 bp)	635266
Total length (>= 5000 bp)	41691
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	221
Largest contig	8868
Total length	635266
GC (%)	37.89
N50	2703
N90	2342
auN	3091.5
L50	93
L90	194
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 1 part

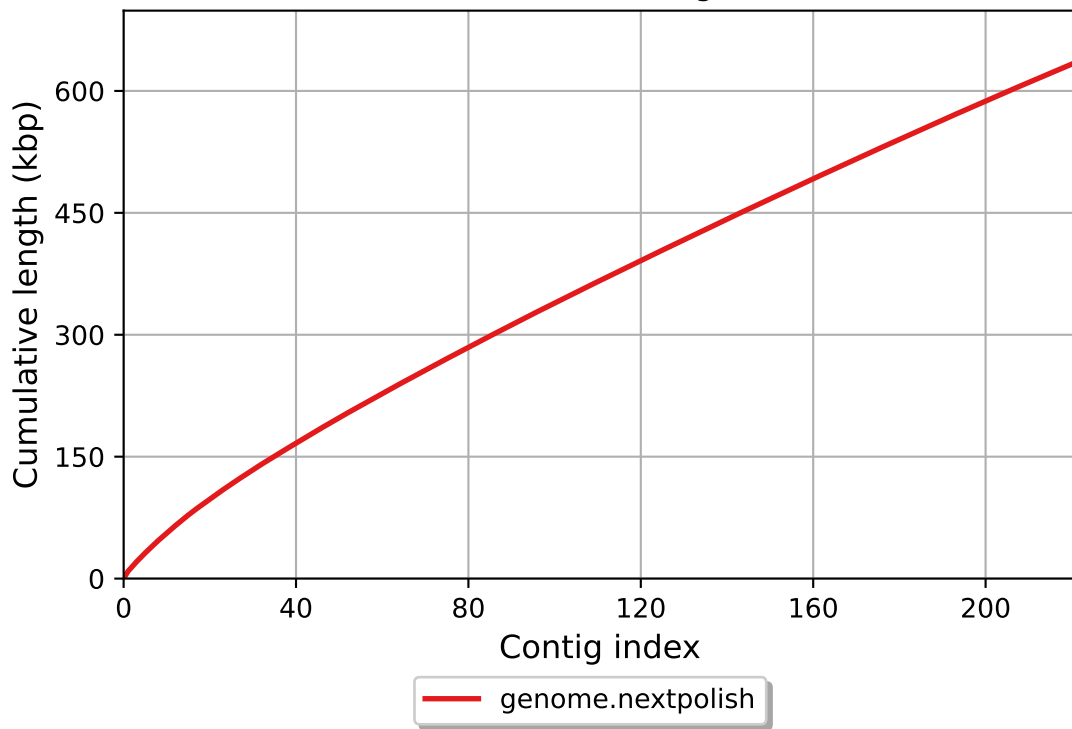
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

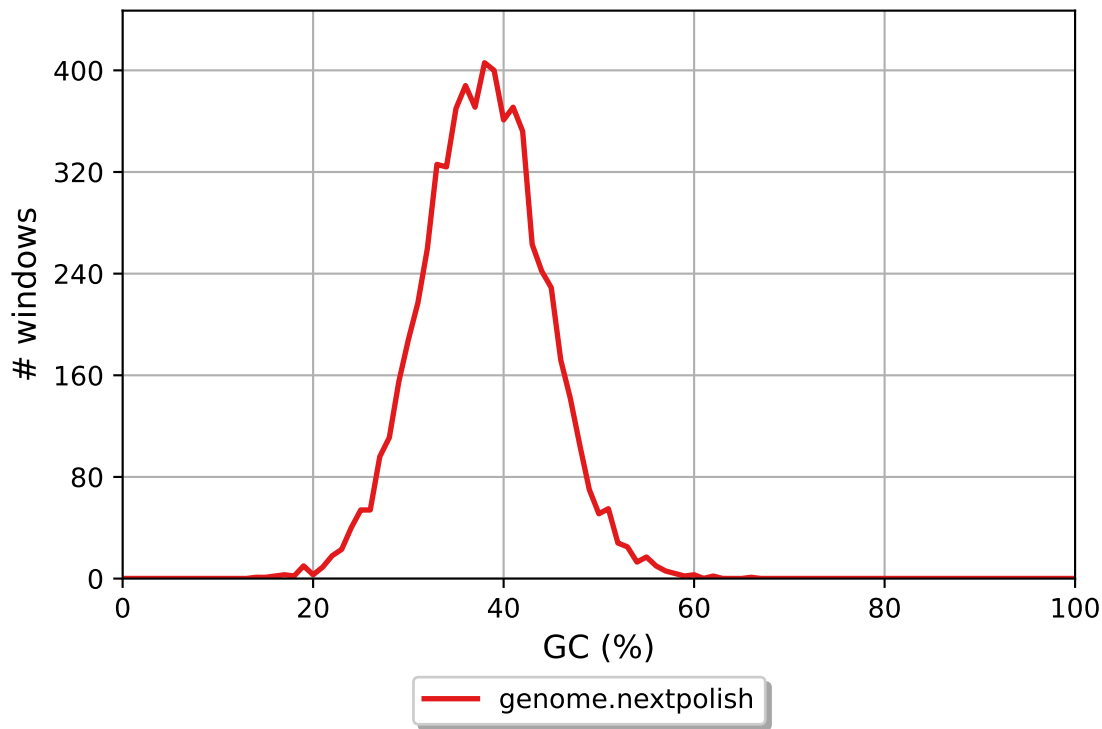


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

