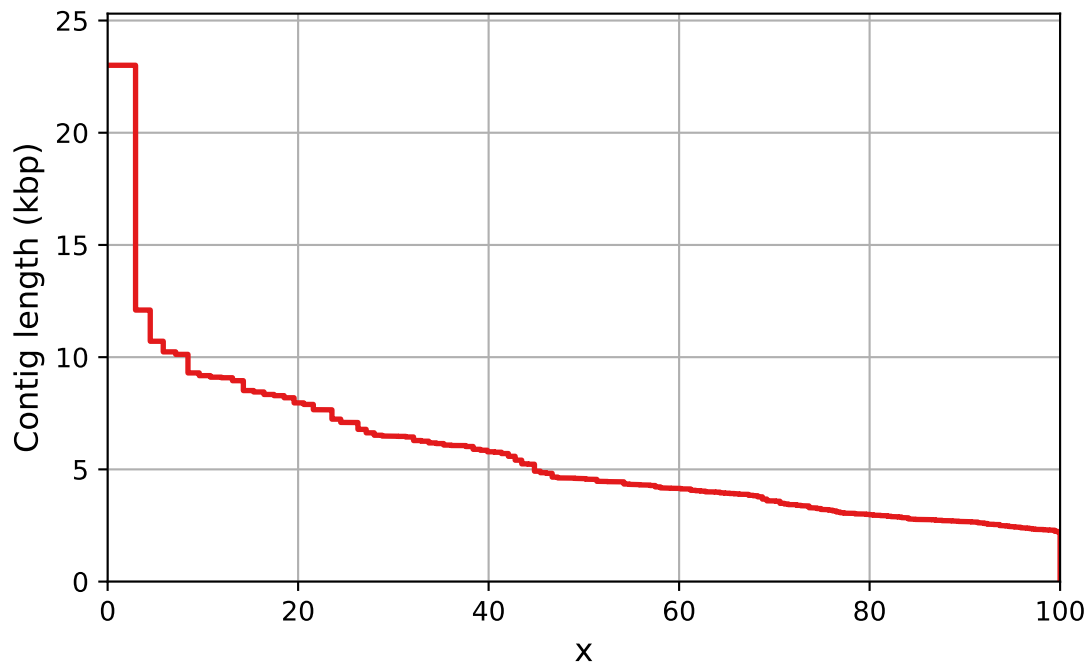


Report

	genome.nextpolish
# contigs (>= 0 bp)	177
# contigs (>= 1000 bp)	177
# contigs (>= 5000 bp)	46
# contigs (>= 10000 bp)	5
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	784446
Total length (>= 1000 bp)	784446
Total length (>= 5000 bp)	351522
Total length (>= 10000 bp)	66174
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	177
Largest contig	23006
Total length	784446
GC (%)	40.63
N50	4586
N90	2678
auN	5802.1
L50	55
L90	145
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 1 part

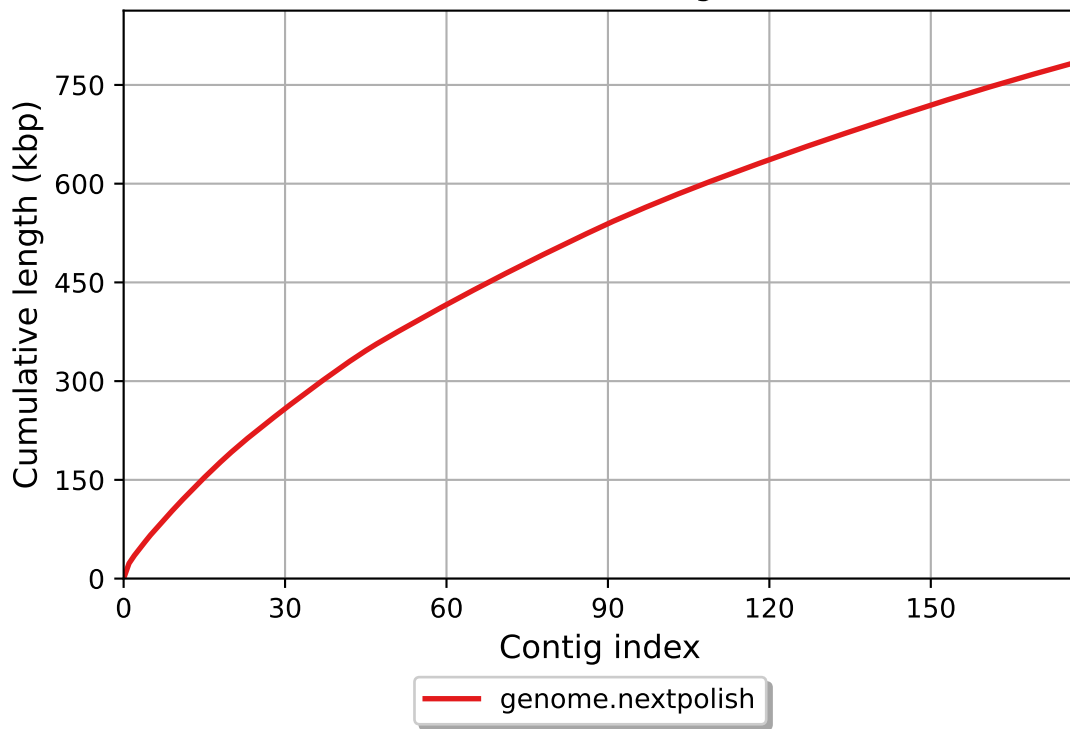
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

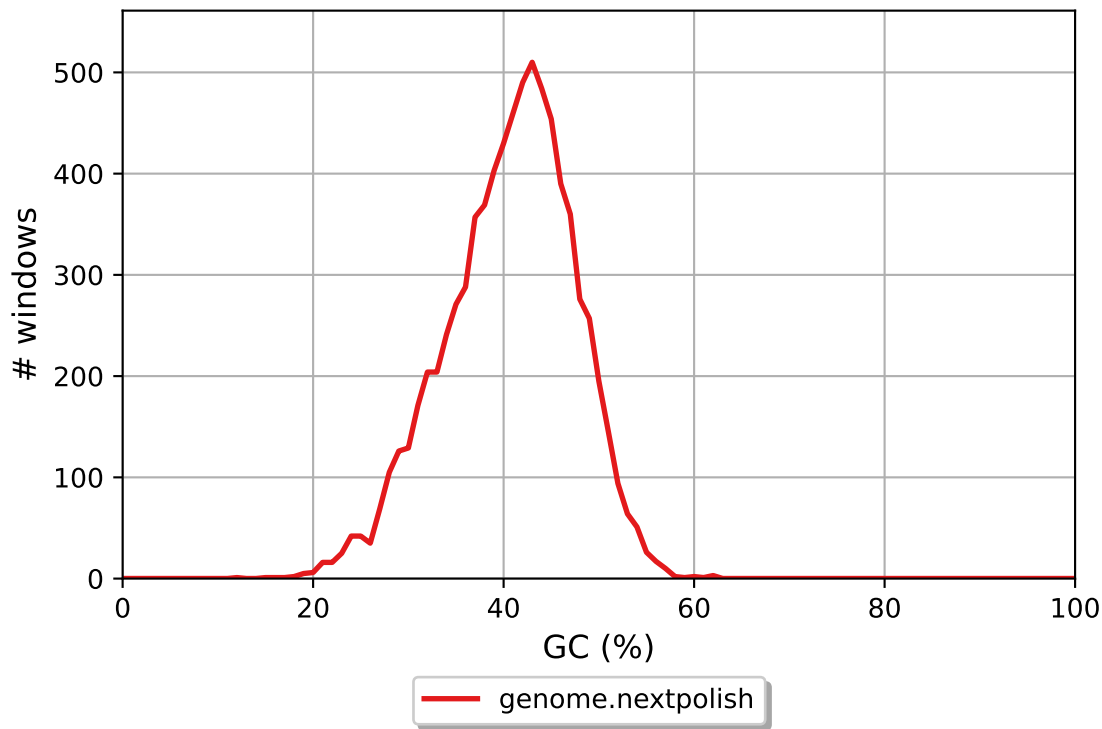


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

