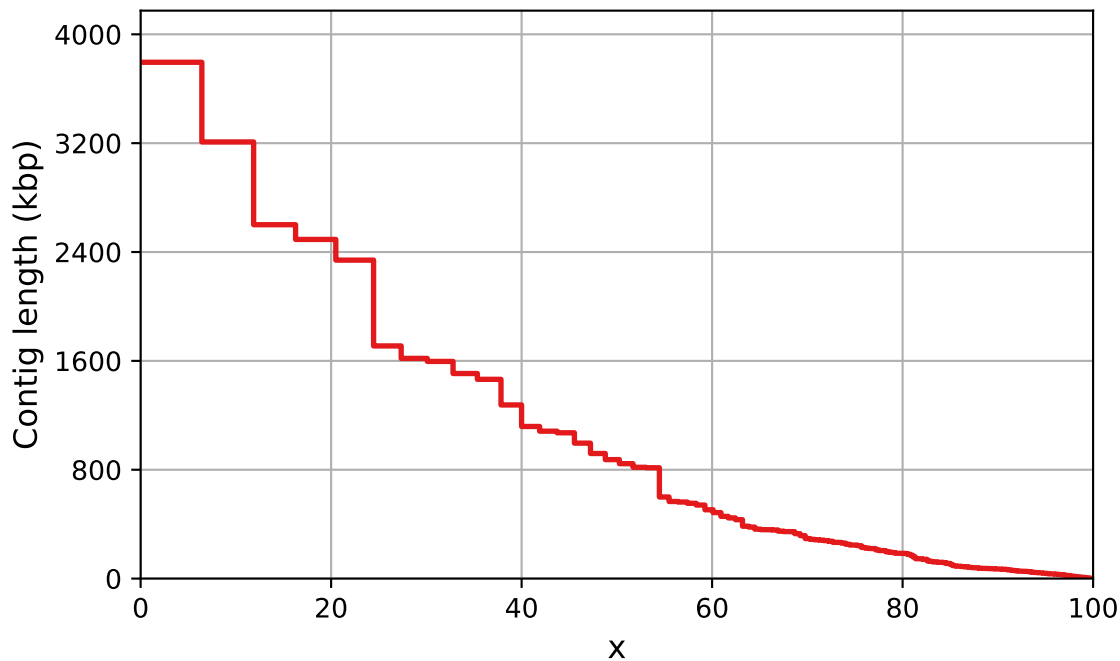


# Report

	genome.nextpolish
# contigs (>= 0 bp)	462
# contigs (>= 1000 bp)	424
# contigs (>= 5000 bp)	341
# contigs (>= 10000 bp)	280
# contigs (>= 25000 bp)	218
# contigs (>= 50000 bp)	158
Total length (>= 0 bp)	59026672
Total length (>= 1000 bp)	59003500
Total length (>= 5000 bp)	58778705
Total length (>= 10000 bp)	58324488
Total length (>= 25000 bp)	57301104
Total length (>= 50000 bp)	55133039
# contigs	458
Largest contig	3794579
Total length	59025281
GC (%)	43.57
N50	874365
N90	70271
auN	1218258.7
L50	17
L90	125
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	151 + 2 part

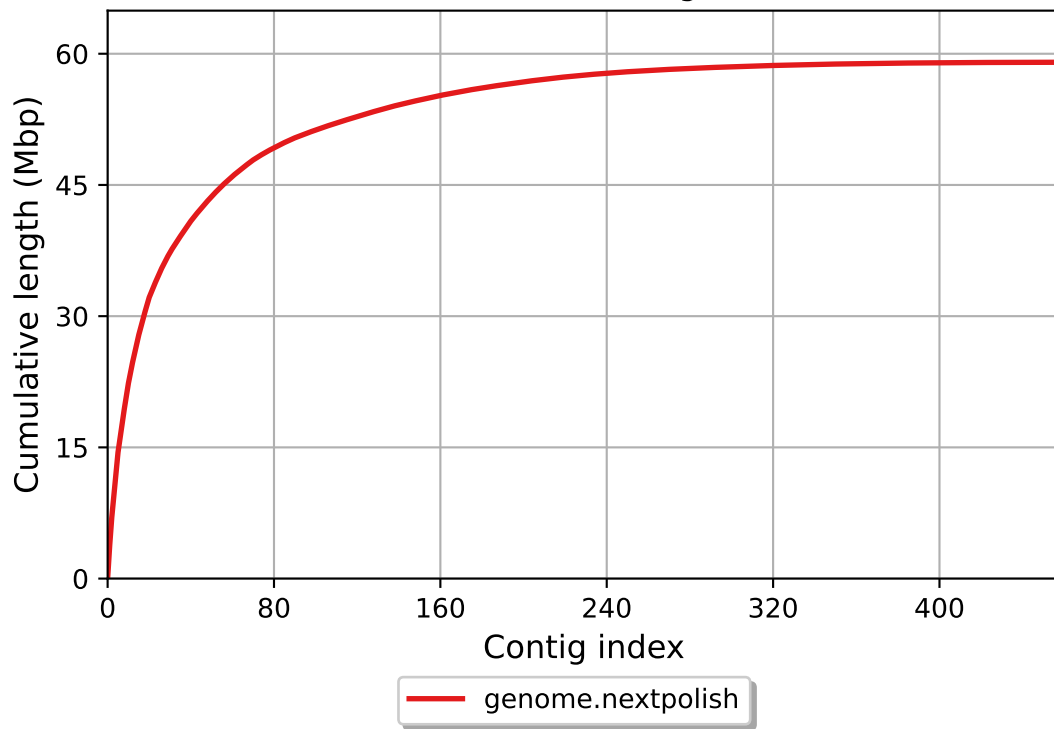
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

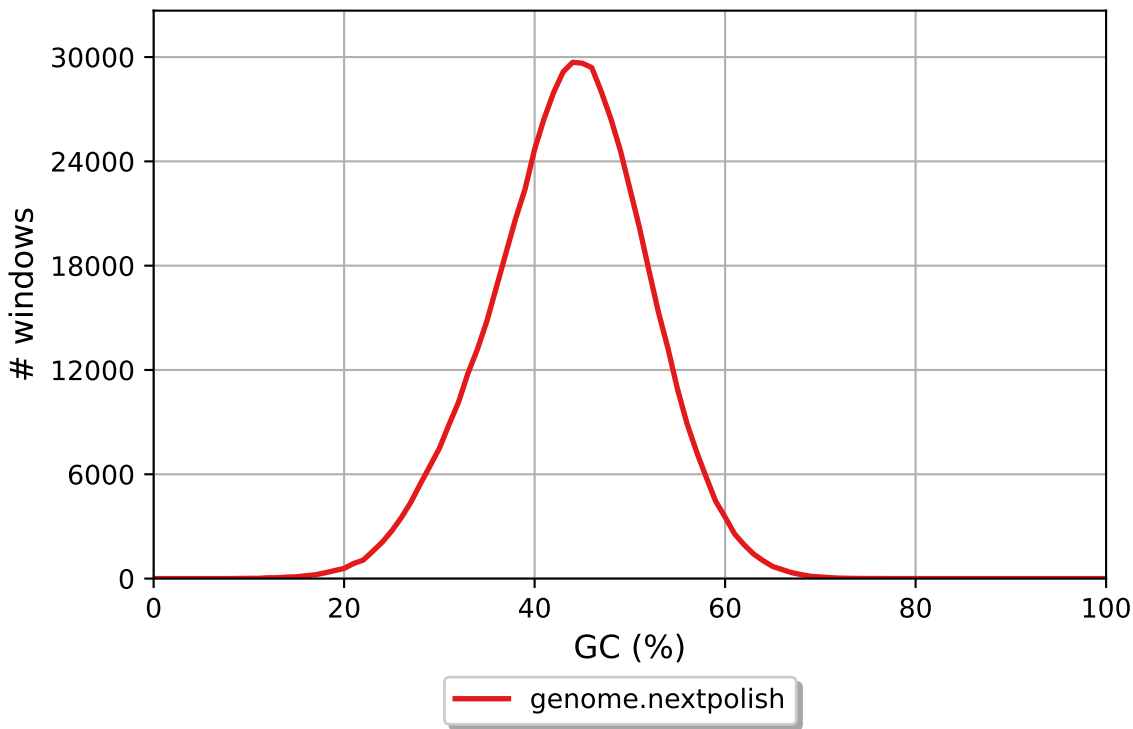


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

