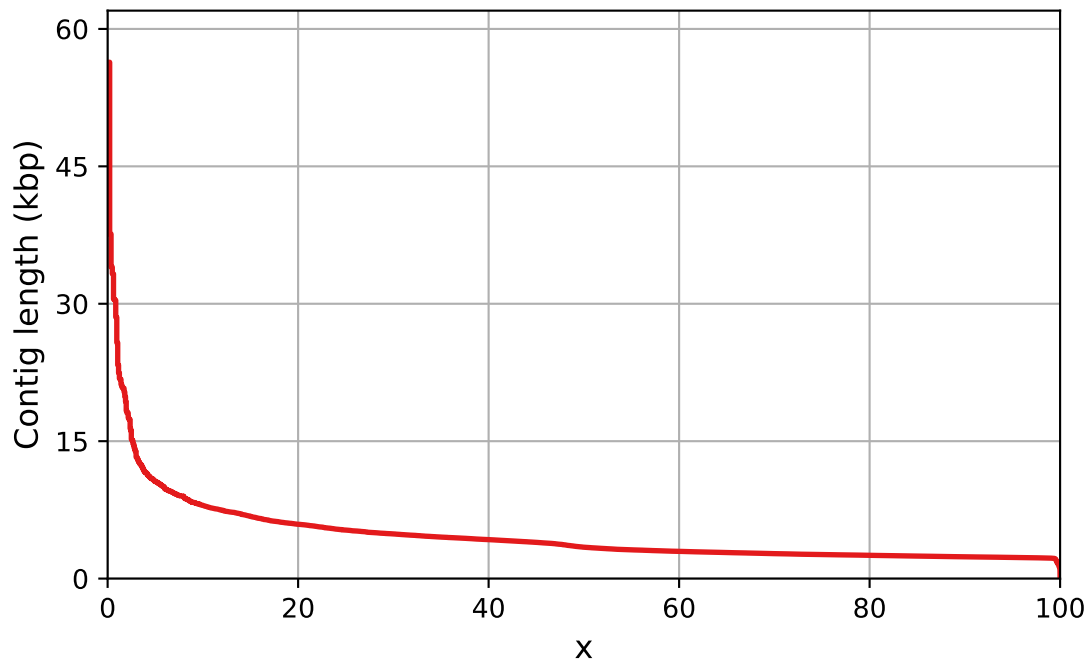


Report

	genome.nextpolish
# contigs (>= 0 bp)	7300
# contigs (>= 1000 bp)	7294
# contigs (>= 5000 bp)	1004
# contigs (>= 10000 bp)	104
# contigs (>= 25000 bp)	8
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	26071488
Total length (>= 1000 bp)	26067613
Total length (>= 5000 bp)	7352224
Total length (>= 10000 bp)	1542182
Total length (>= 25000 bp)	276420
Total length (>= 50000 bp)	56360
# contigs	7300
Largest contig	56360
Total length	26071488
GC (%)	43.10
N50	3421
N90	2378
auN	4837.9
L50	2343
L90	6141
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	28 + 9 part

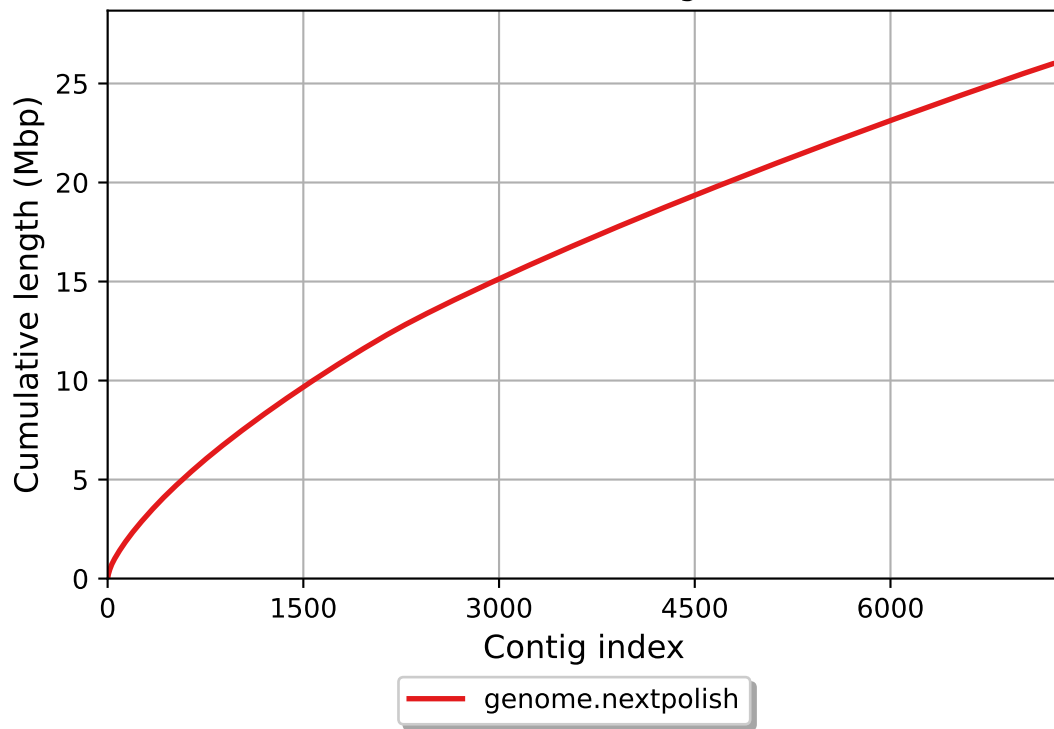
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

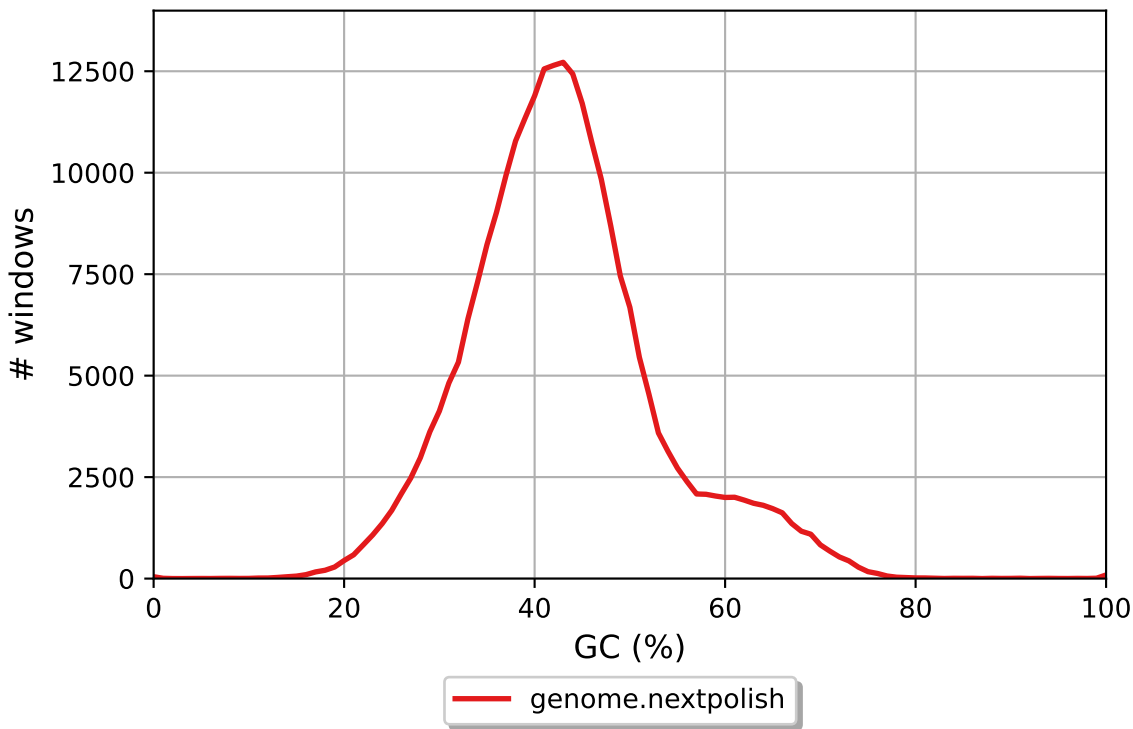


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

