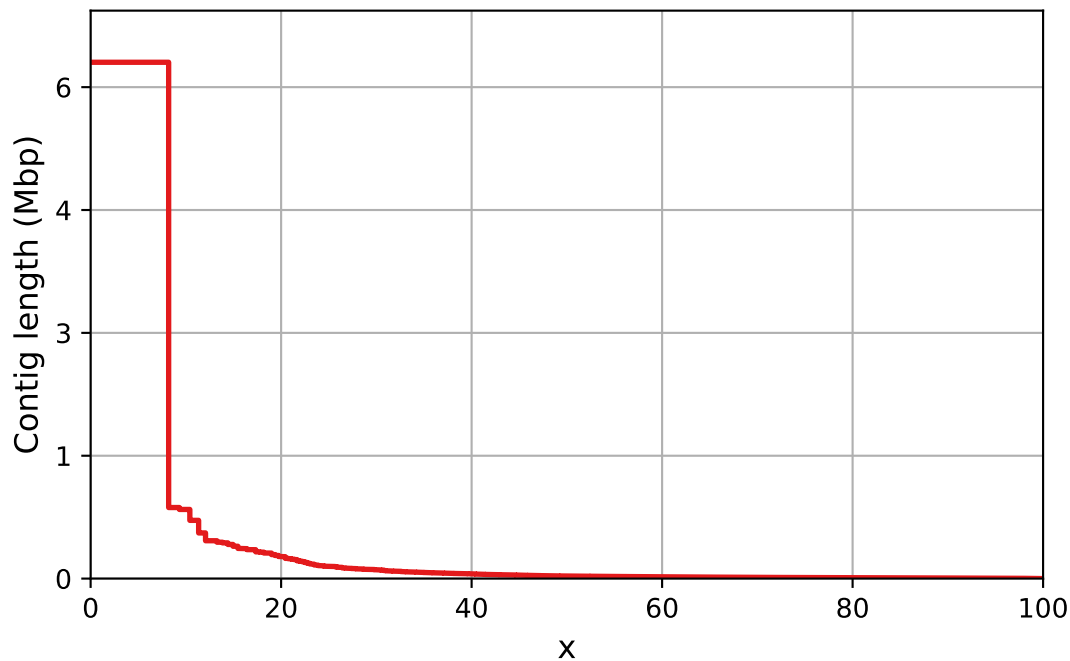


# Report

	genome.nextpolish
# contigs (>= 0 bp)	4439
# contigs (>= 1000 bp)	4418
# contigs (>= 5000 bp)	3197
# contigs (>= 10000 bp)	1856
# contigs (>= 25000 bp)	532
# contigs (>= 50000 bp)	202
Total length (>= 0 bp)	76946664
Total length (>= 1000 bp)	76932539
Total length (>= 5000 bp)	72962967
Total length (>= 10000 bp)	63263079
Total length (>= 25000 bp)	43182279
Total length (>= 50000 bp)	32205690
# contigs	4439
Largest contig	6303783
Total length	76946664
GC (%)	53.07
N50	31227
N90	6923
auN	611209.2
L50	363
L90	2571
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	118 + 15 part

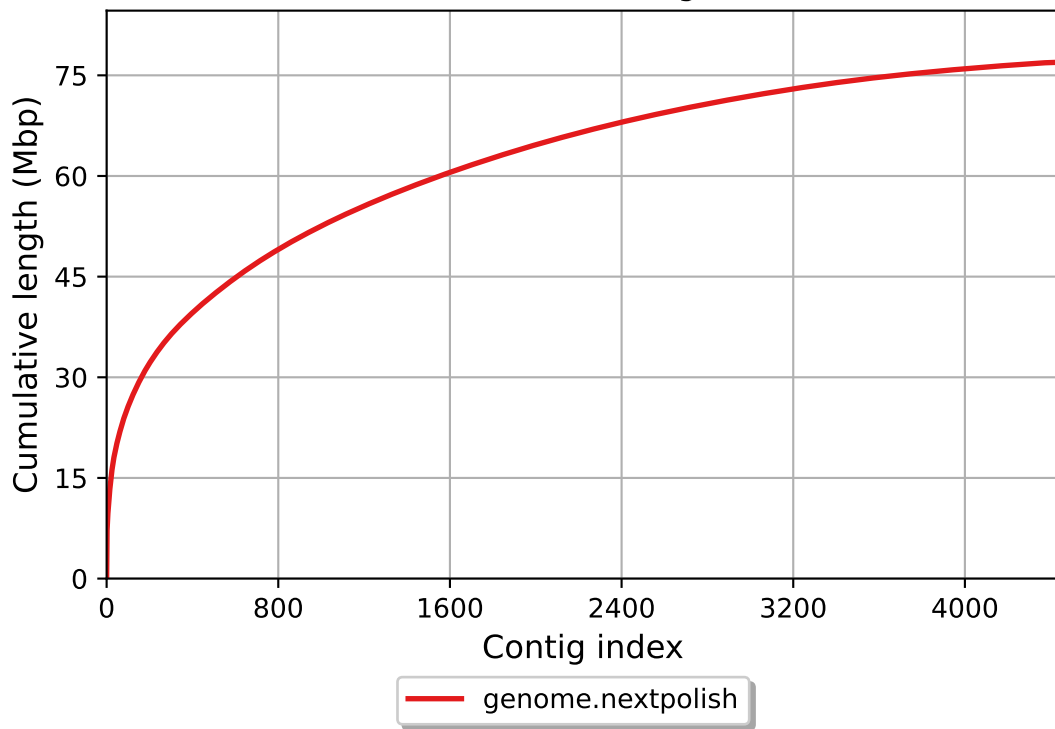
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

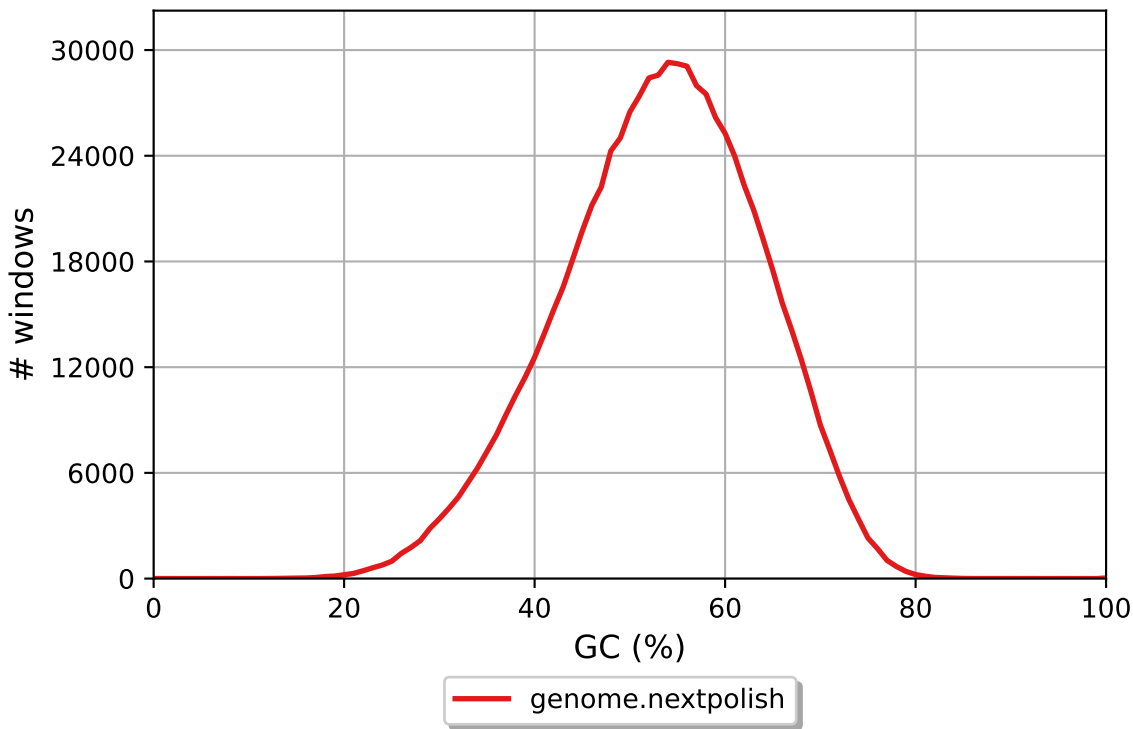


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

