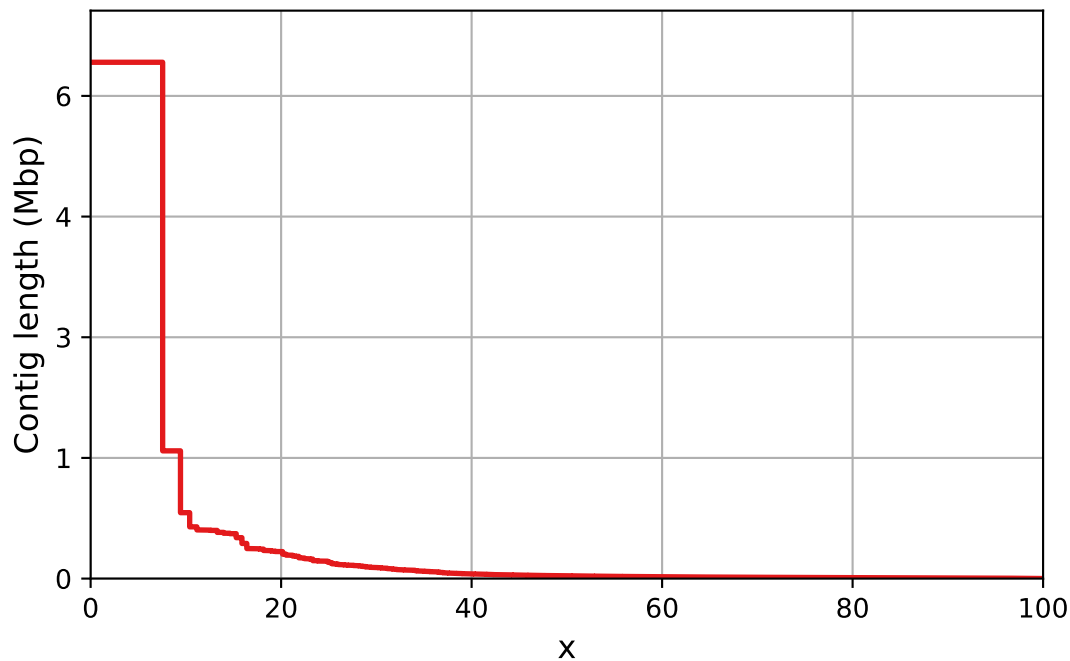


Report

	genome.nextpolish
# contigs (>= 0 bp)	4486
# contigs (>= 1000 bp)	4466
# contigs (>= 5000 bp)	3425
# contigs (>= 10000 bp)	2031
# contigs (>= 25000 bp)	569
# contigs (>= 50000 bp)	196
Total length (>= 0 bp)	84863268
Total length (>= 1000 bp)	84848565
Total length (>= 5000 bp)	81354001
Total length (>= 10000 bp)	71169638
Total length (>= 25000 bp)	48645566
Total length (>= 50000 bp)	35816252
# contigs	4486
Largest contig	6418108
Total length	84863268
GC (%)	54.56
N50	34691
N90	7614
auN	612384.1
L50	358
L90	2628
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	109 + 17 part

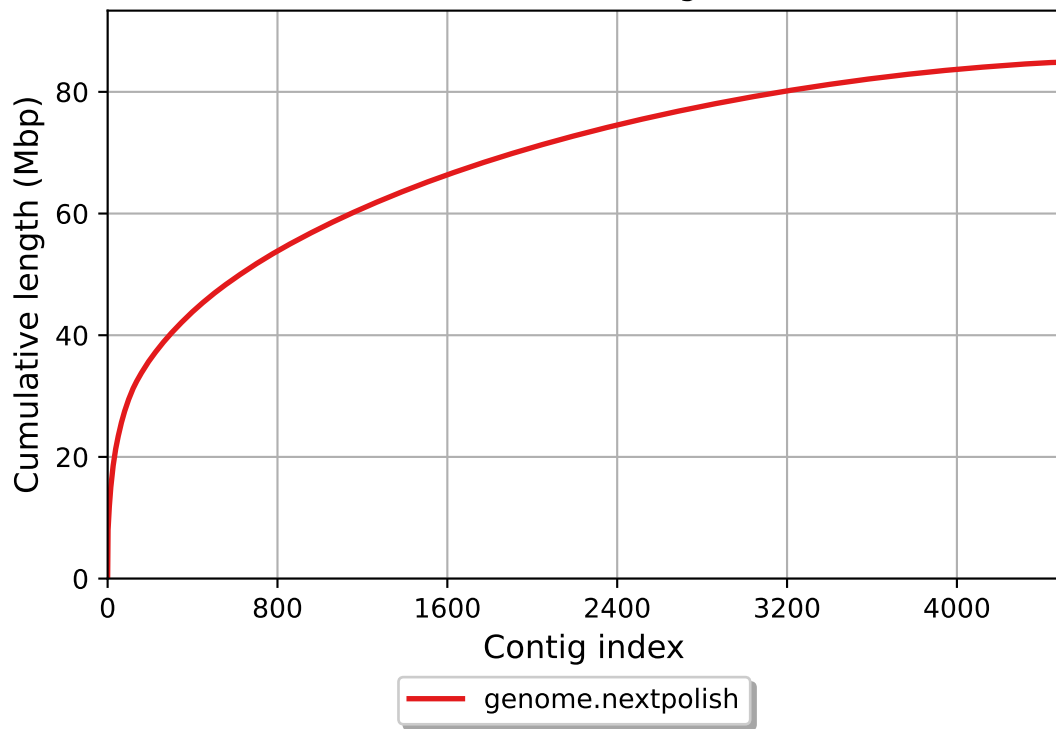
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

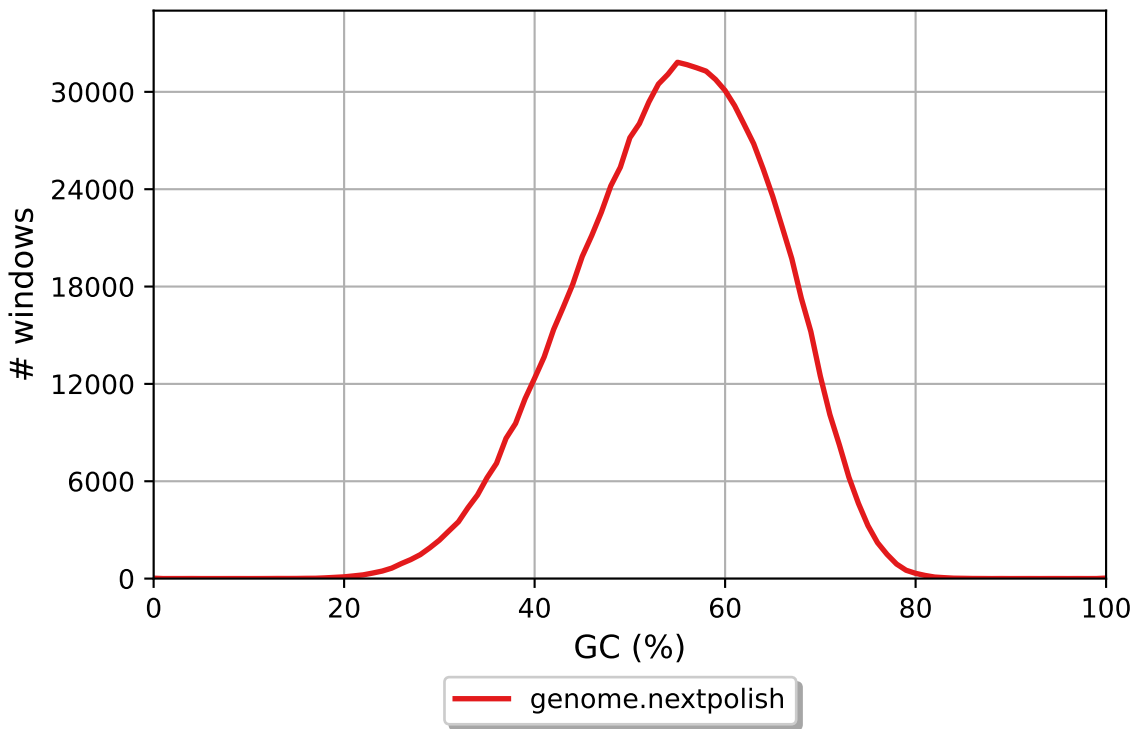


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

