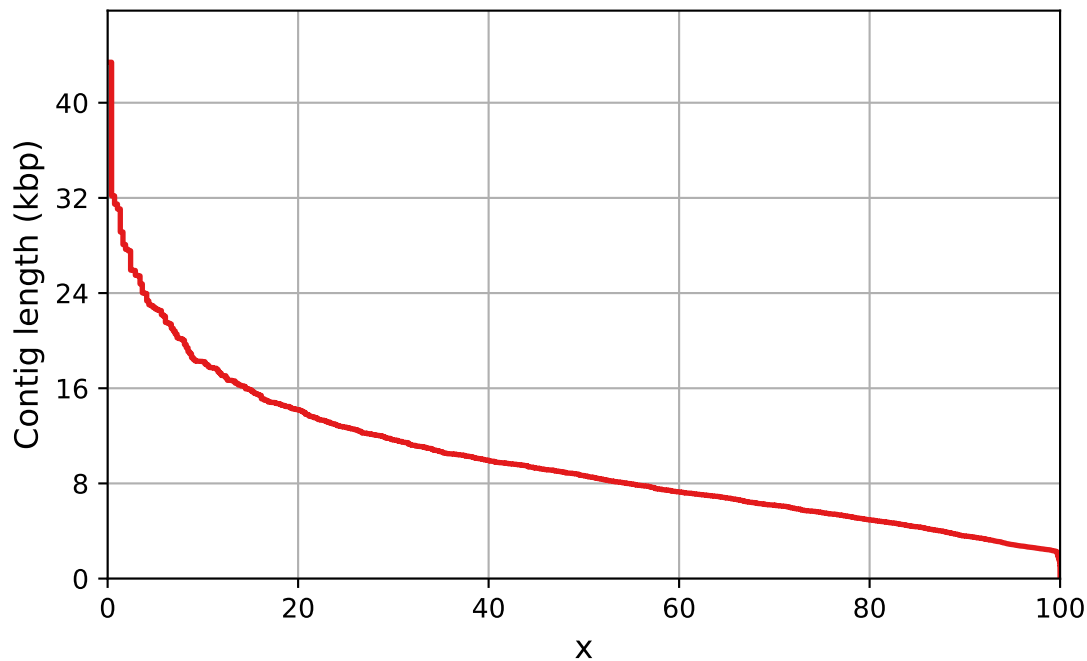


Report

	genome.nextpolish
# contigs (>= 0 bp)	1486
# contigs (>= 1000 bp)	1486
# contigs (>= 5000 bp)	863
# contigs (>= 10000 bp)	284
# contigs (>= 25000 bp)	12
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	10384694
Total length (>= 1000 bp)	10384694
Total length (>= 5000 bp)	8254304
Total length (>= 10000 bp)	4105554
Total length (>= 25000 bp)	353339
Total length (>= 50000 bp)	0
# contigs	1486
Largest contig	43401
Total length	10384694
GC (%)	50.79
N50	8661
N90	3585
auN	10037.5
L50	401
L90	1117
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	9 + 4 part

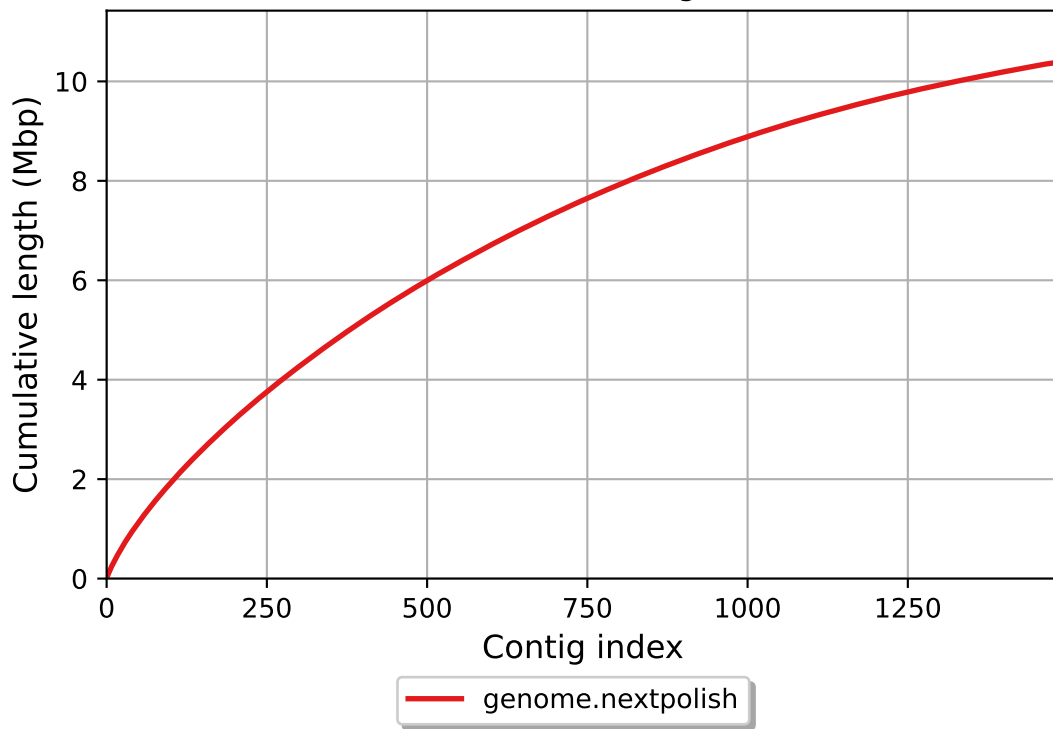
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

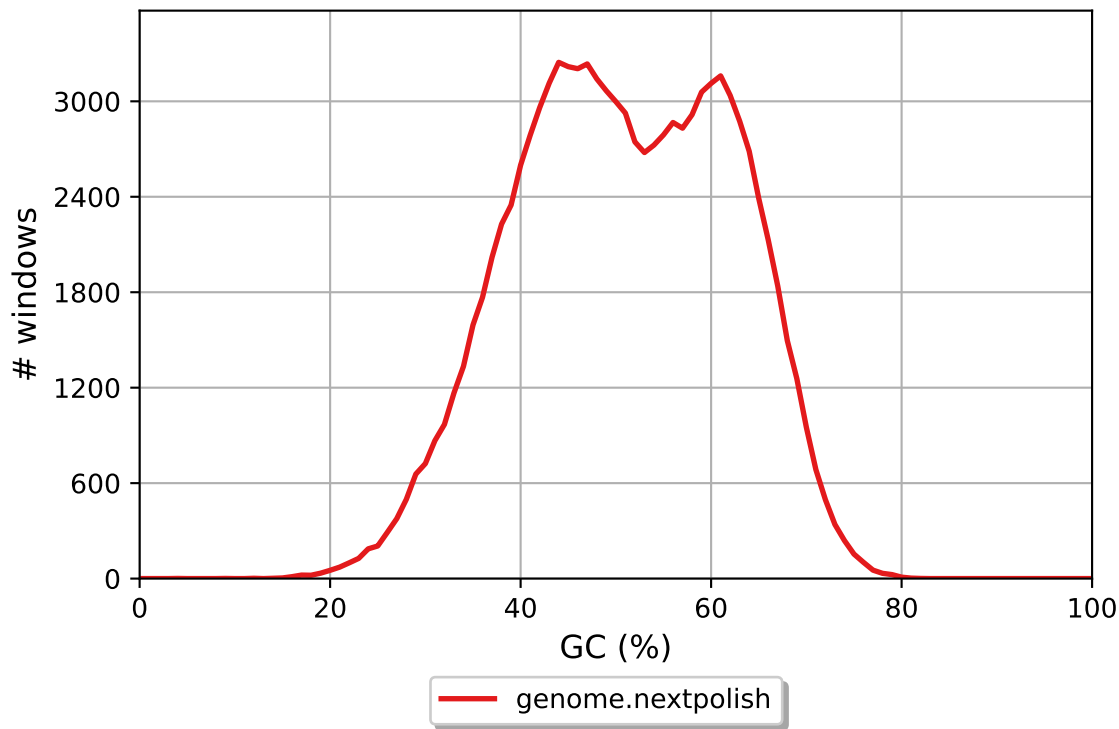


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

