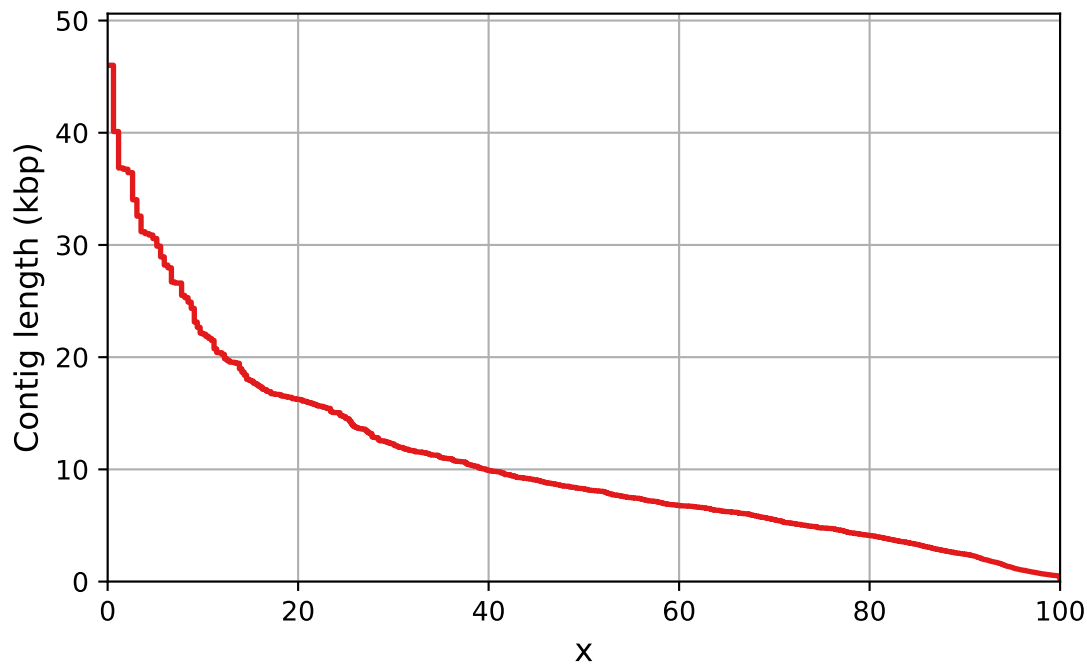


Report

	genome.nextpolish
# contigs (>= 0 bp)	1609
# contigs (>= 1000 bp)	1176
# contigs (>= 5000 bp)	537
# contigs (>= 10000 bp)	184
# contigs (>= 25000 bp)	20
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	7495668
Total length (>= 1000 bp)	7208032
Total length (>= 5000 bp)	5486645
Total length (>= 10000 bp)	2972679
Total length (>= 25000 bp)	632140
Total length (>= 50000 bp)	0
# contigs	1583
Largest contig	46013
Total length	7486427
GC (%)	65.23
N50	8275
N90	2449
auN	10718.0
L50	270
L90	882
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	16 + 1 part

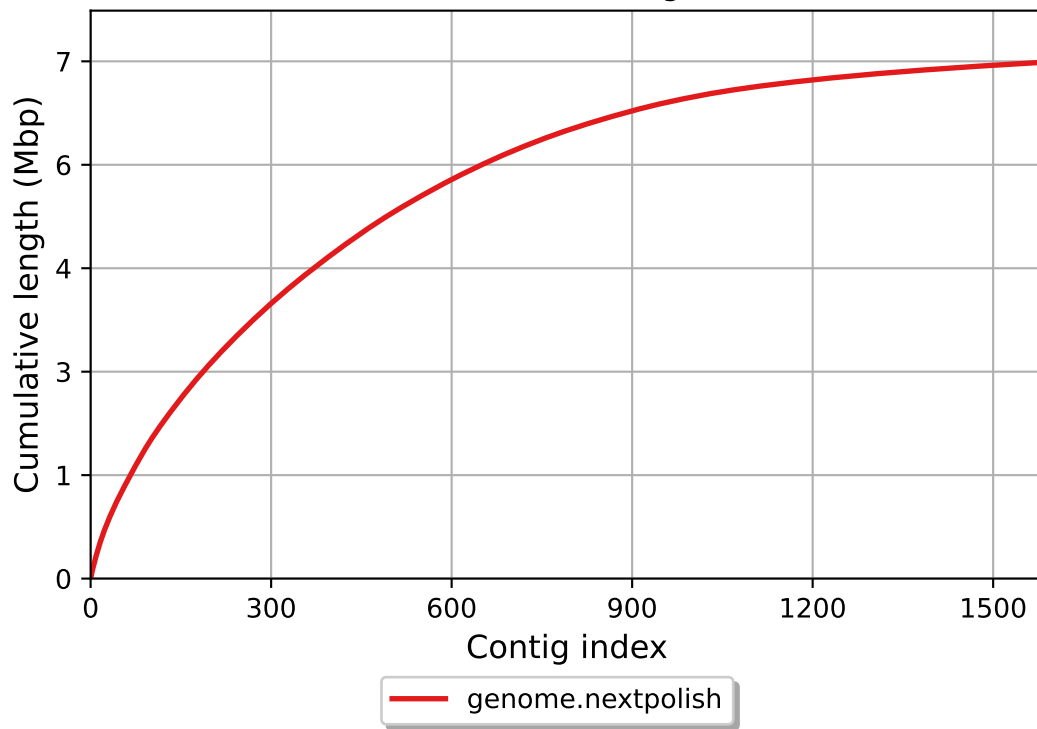
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

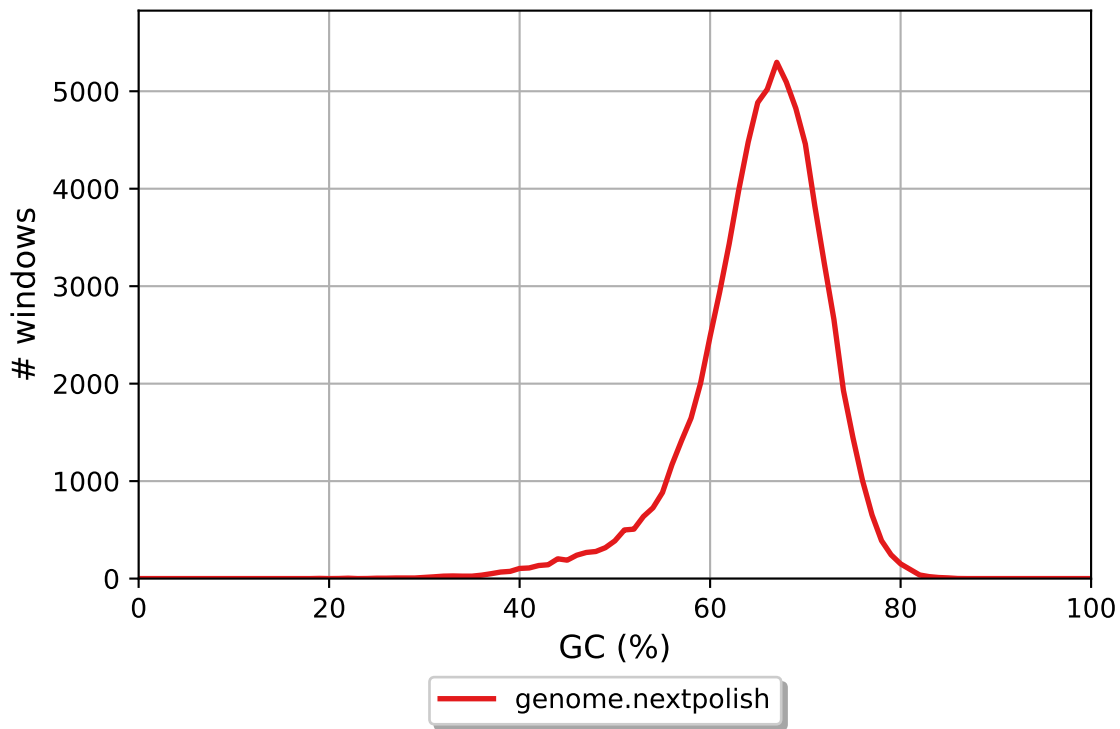


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

