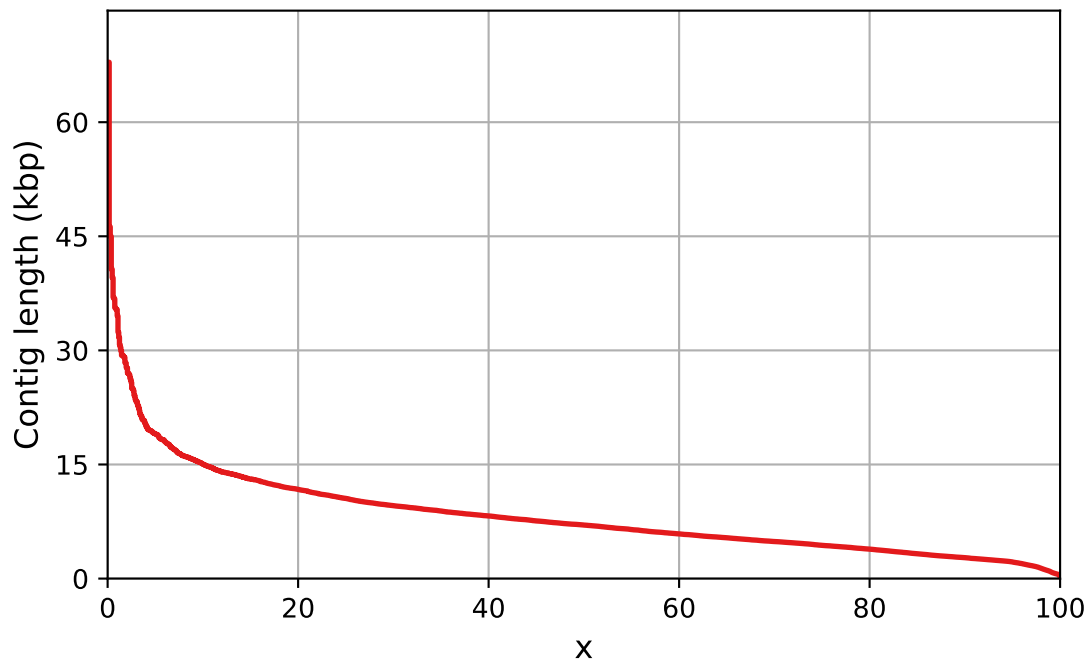


Report

	genome.nextpolish
# contigs (>= 0 bp)	8197
# contigs (>= 1000 bp)	7452
# contigs (>= 5000 bp)	3249
# contigs (>= 10000 bp)	807
# contigs (>= 25000 bp)	33
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	41701314
Total length (>= 1000 bp)	41205333
Total length (>= 5000 bp)	28592300
Total length (>= 10000 bp)	11395893
Total length (>= 25000 bp)	1078506
Total length (>= 50000 bp)	67877
# contigs	8162
Largest contig	67877
Total length	41689578
GC (%)	53.05
N50	7039
N90	2750
auN	8412.8
L50	1941
L90	5608
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	20 + 4 part

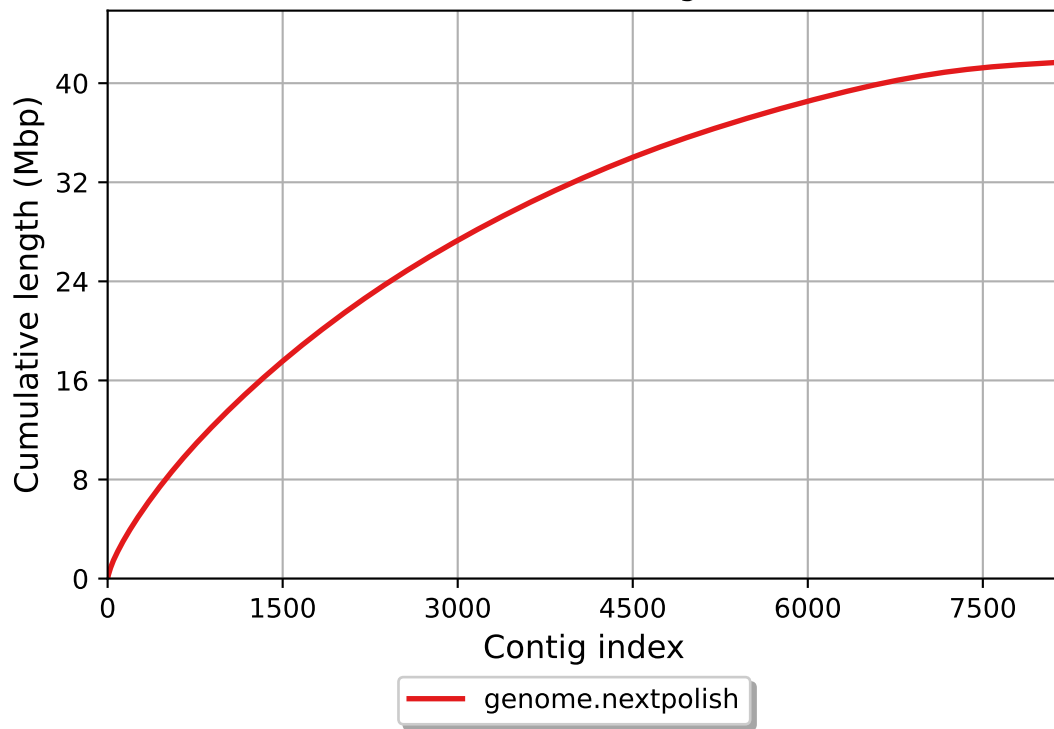
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

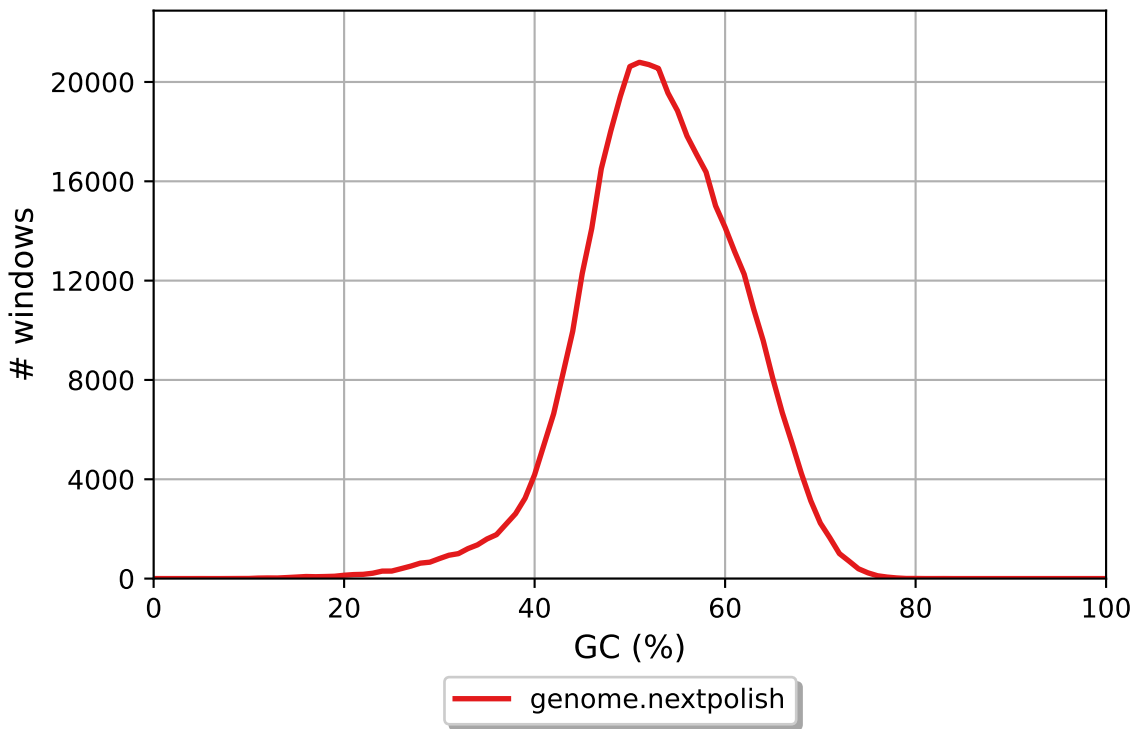


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

