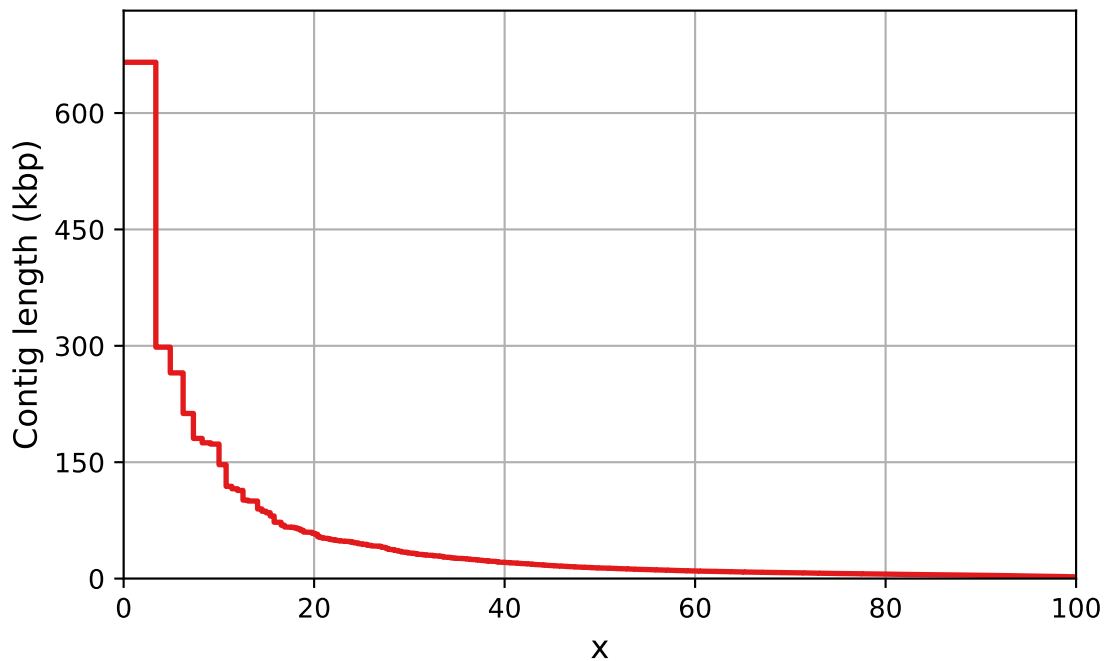


Report

	genome.nextpolish
# contigs (>= 0 bp)	1982
# contigs (>= 1000 bp)	1980
# contigs (>= 5000 bp)	1137
# contigs (>= 10000 bp)	425
# contigs (>= 25000 bp)	120
# contigs (>= 50000 bp)	38
Total length (>= 0 bp)	19674713
Total length (>= 1000 bp)	19673254
Total length (>= 5000 bp)	16635218
Total length (>= 10000 bp)	11661085
Total length (>= 25000 bp)	7158580
Total length (>= 50000 bp)	4320273
# contigs	1982
Largest contig	665436
Total length	19674713
GC (%)	38.80
N50	13686
N90	4264
auN	58870.8
L50	270
L90	1369
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	11 + 4 part

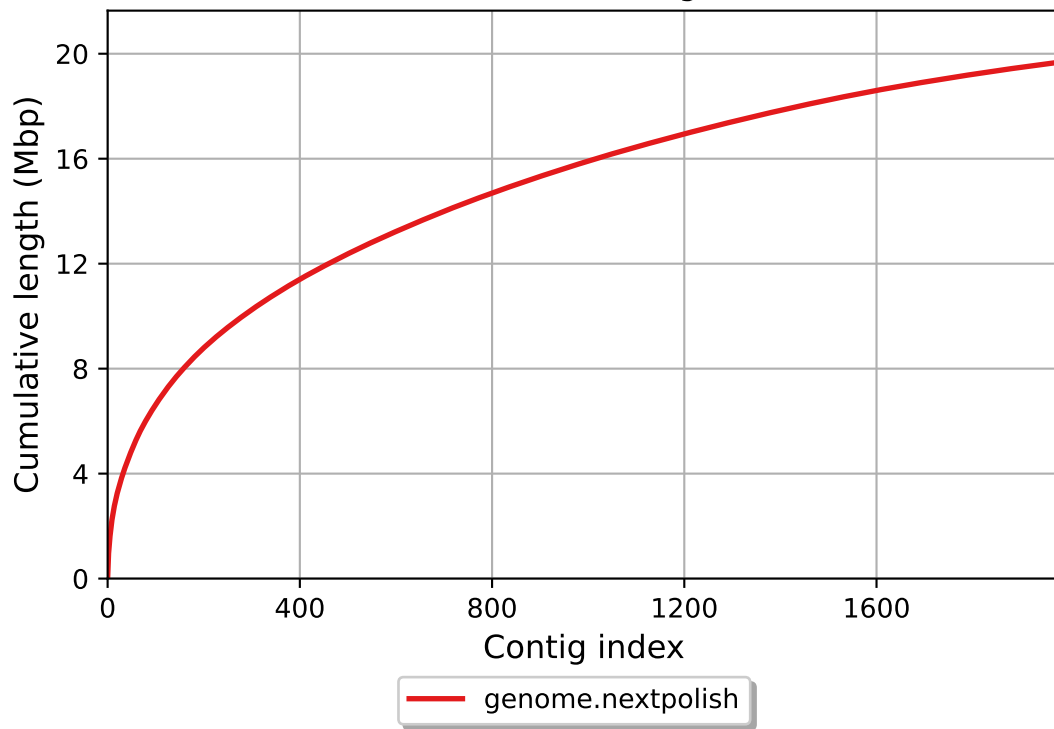
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

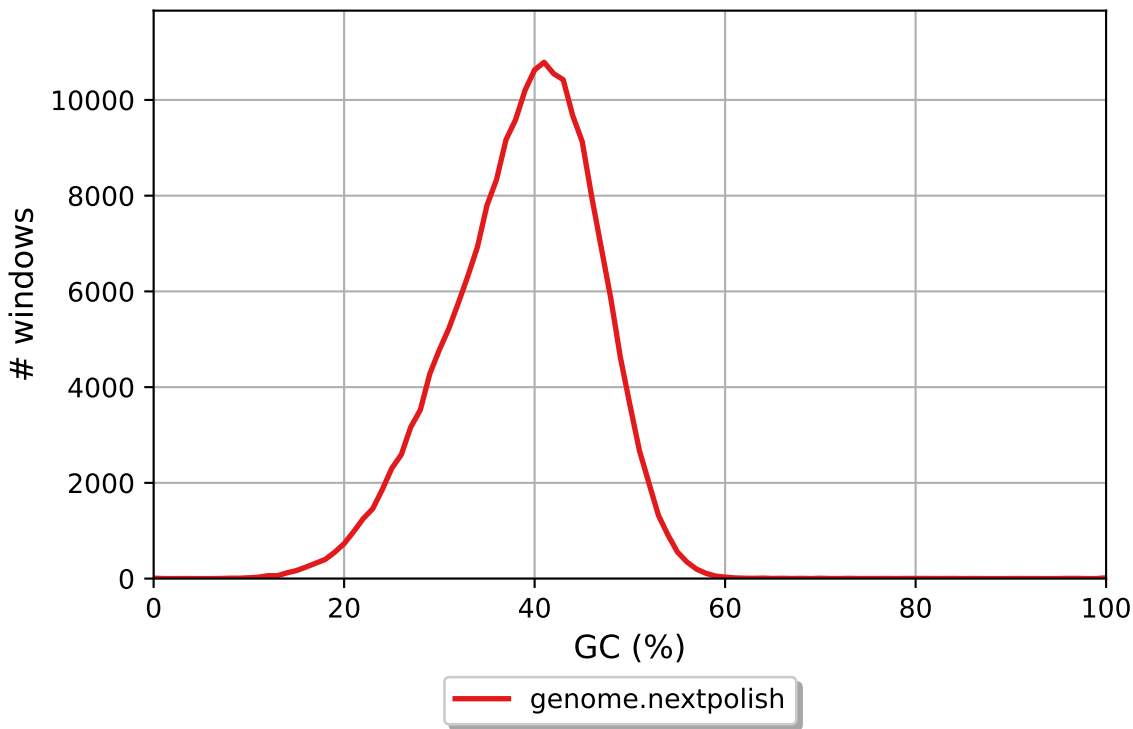


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

