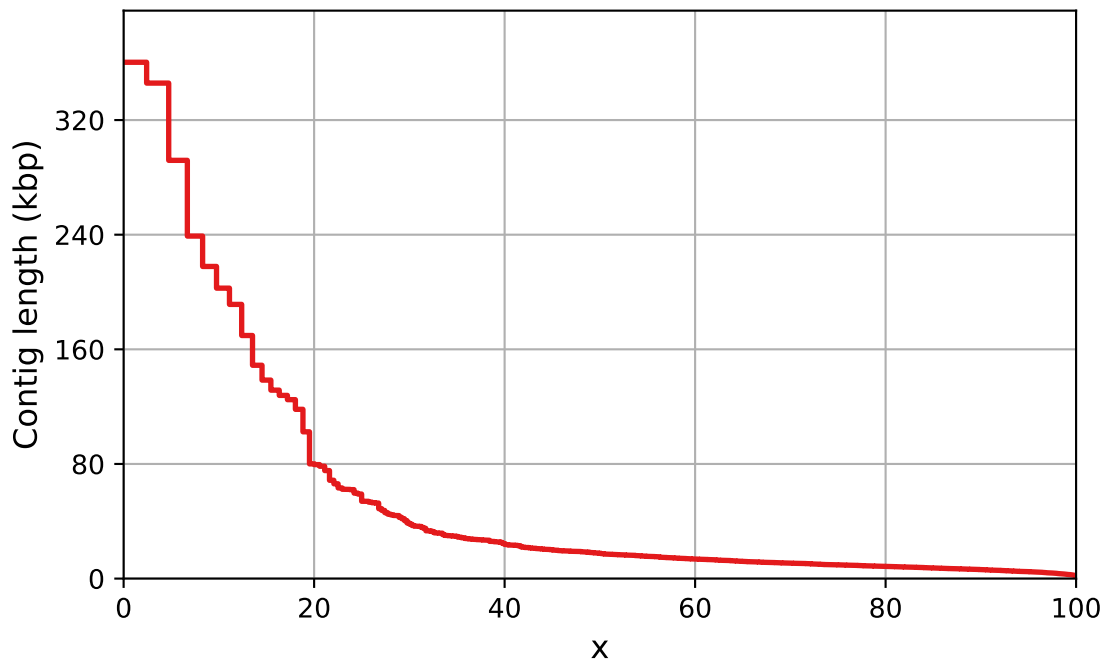


Report

	genome.nextpolish
# contigs (>= 0 bp)	1094
# contigs (>= 1000 bp)	1089
# contigs (>= 5000 bp)	853
# contigs (>= 10000 bp)	432
# contigs (>= 25000 bp)	92
# contigs (>= 50000 bp)	32
Total length (>= 0 bp)	14916491
Total length (>= 1000 bp)	14913066
Total length (>= 5000 bp)	14030671
Total length (>= 10000 bp)	10907460
Total length (>= 25000 bp)	5934005
Total length (>= 50000 bp)	3994308
# contigs	1094
Largest contig	360330
Total length	14916491
GC (%)	43.76
N50	17650
N90	6297
auN	60418.1
L50	168
L90	746
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	34 + 5 part

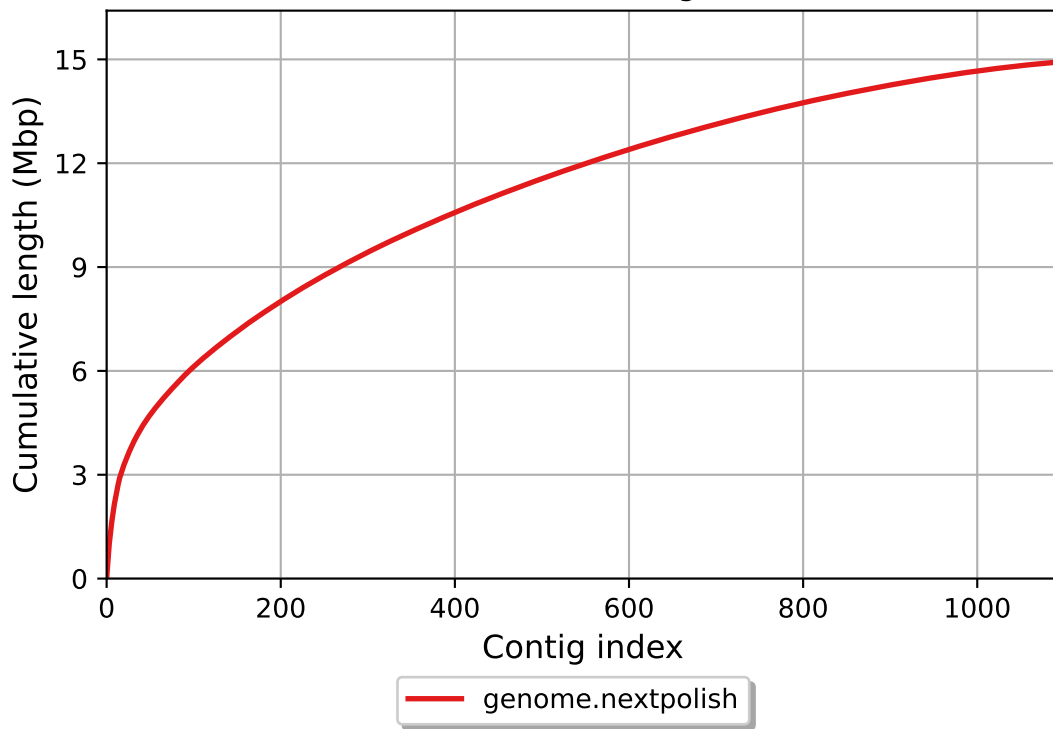
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

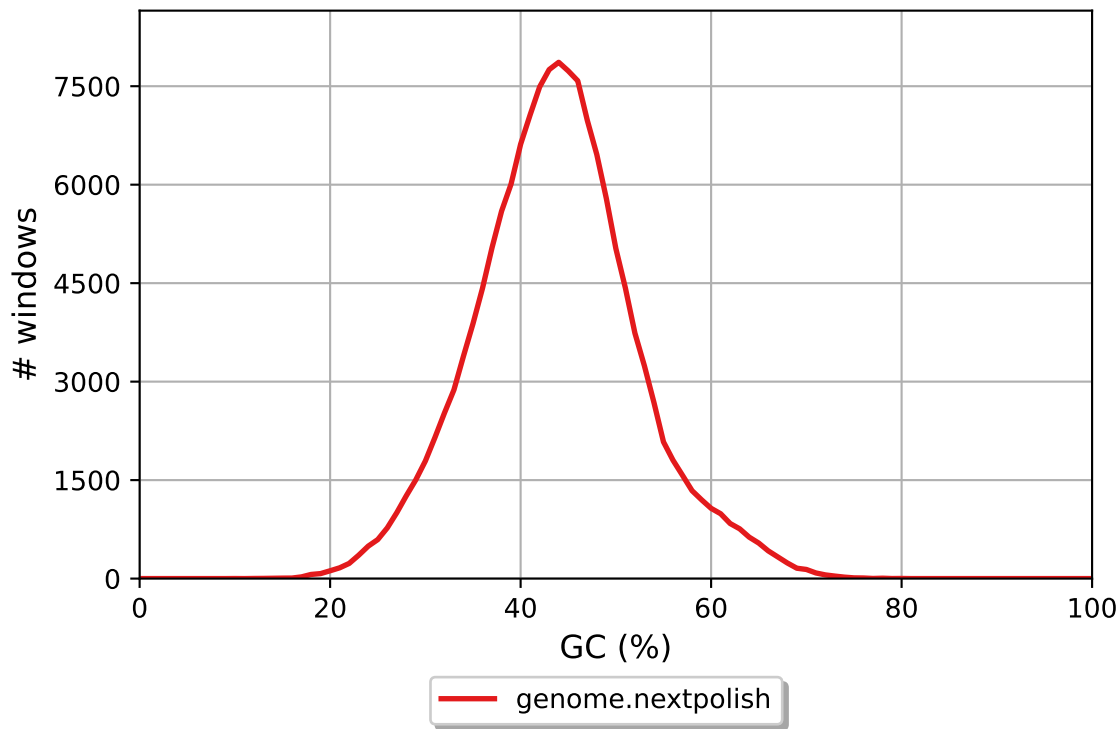


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

