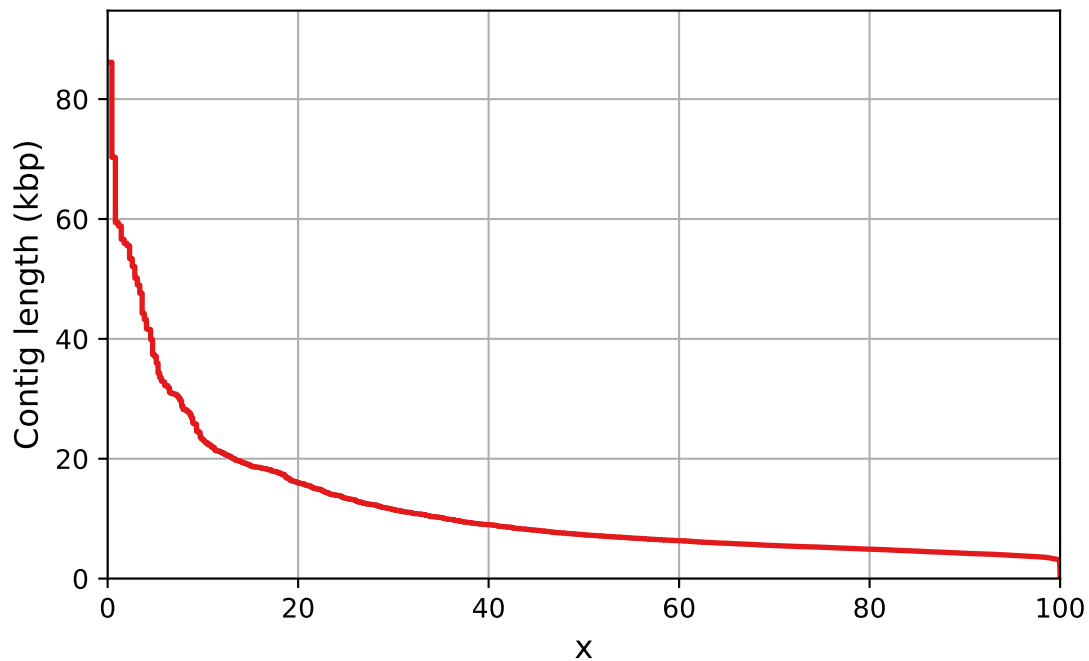


Report

	genome.nextpolish
# contigs (>= 0 bp)	2627
# contigs (>= 1000 bp)	2627
# contigs (>= 5000 bp)	1645
# contigs (>= 10000 bp)	391
# contigs (>= 25000 bp)	46
# contigs (>= 50000 bp)	10
Total length (>= 0 bp)	19185991
Total length (>= 1000 bp)	19185991
Total length (>= 5000 bp)	15072790
Total length (>= 10000 bp)	6788746
Total length (>= 25000 bp)	1790637
Total length (>= 50000 bp)	598393
# contigs	2627
Largest contig	86147
Total length	19185991
GC (%)	36.71
N50	7310
N90	4228
auN	11939.4
L50	724
L90	2122
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

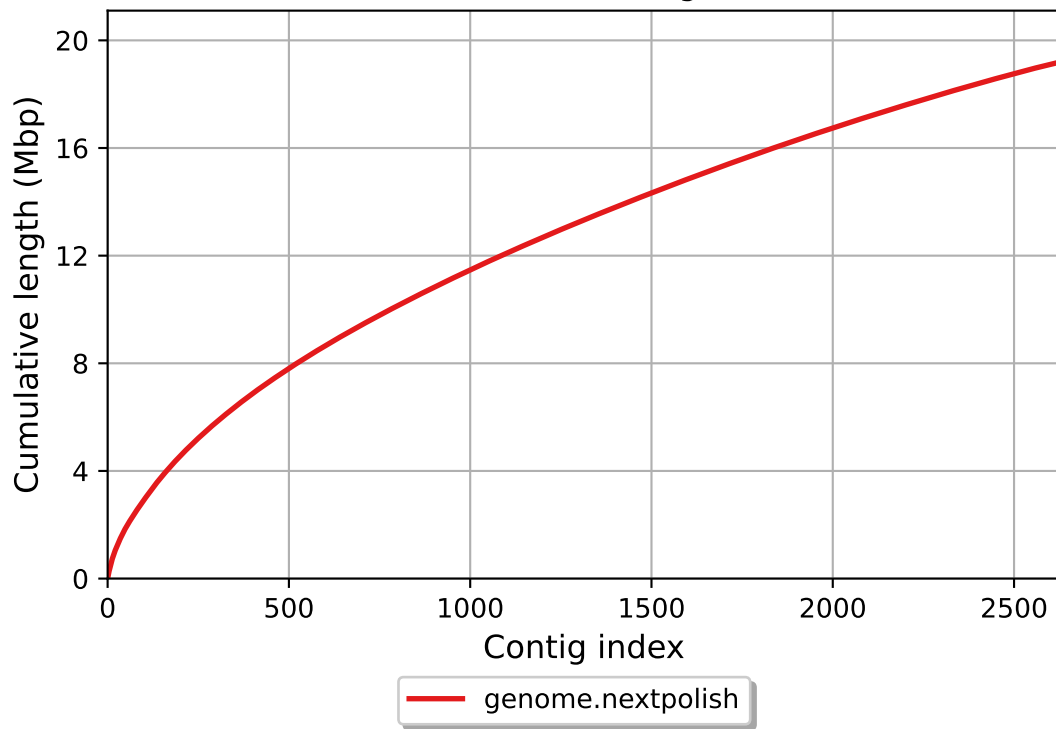
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

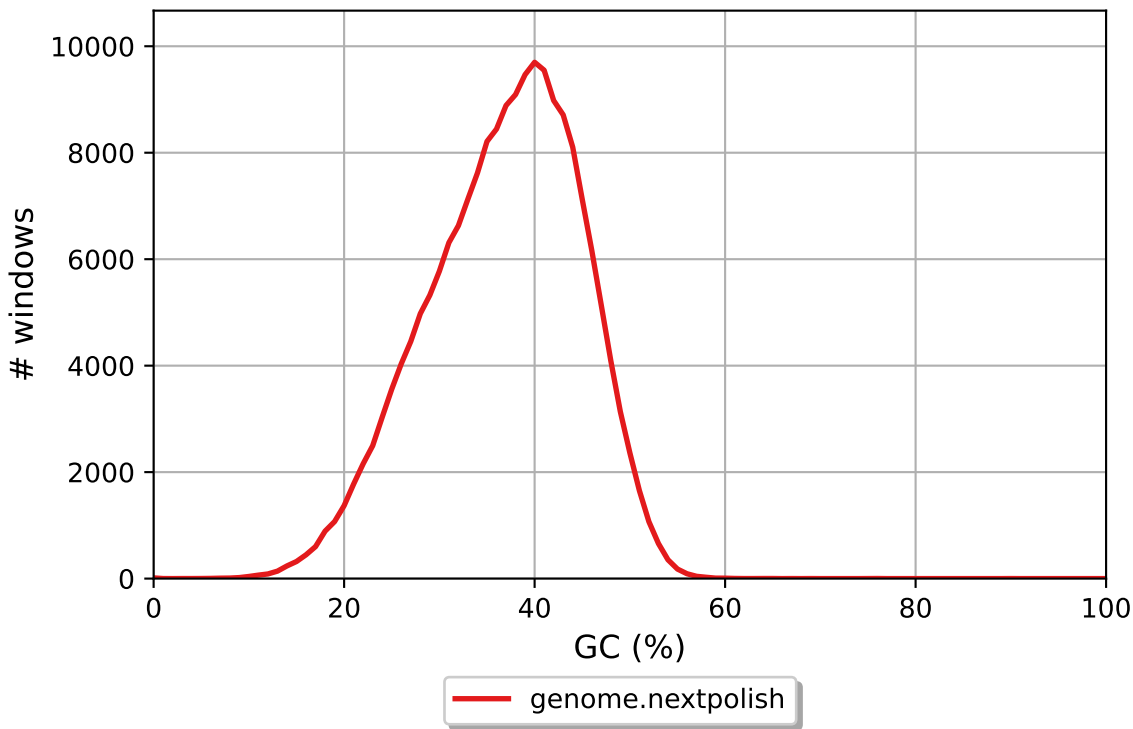


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

