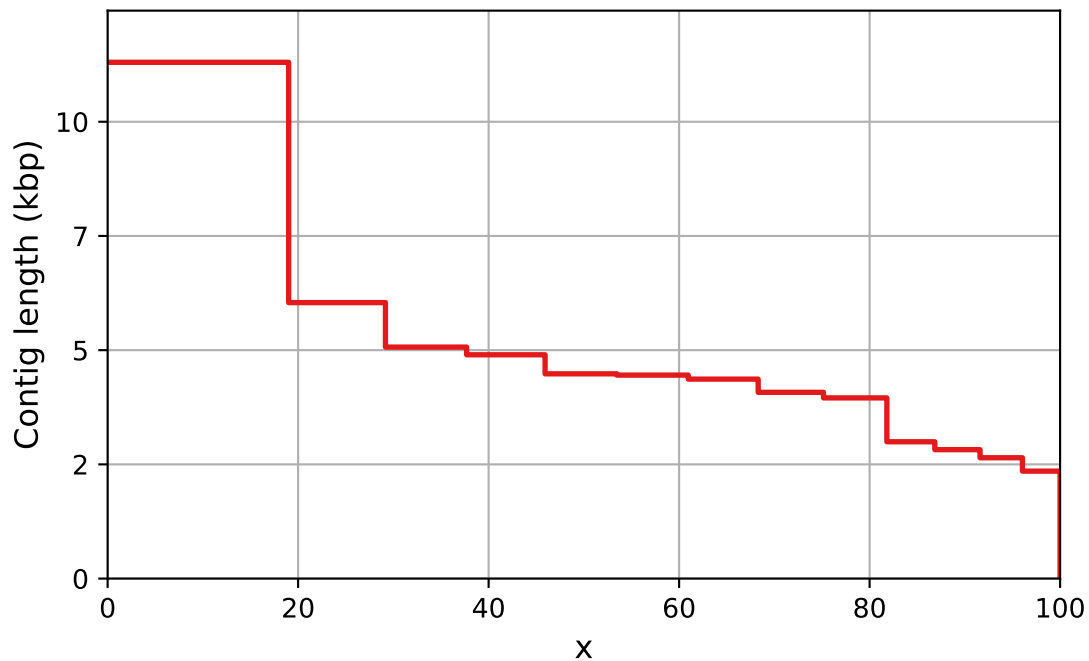


Report

	genome.nextpolish
# contigs (>= 0 bp)	13
# contigs (>= 1000 bp)	13
# contigs (>= 5000 bp)	3
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	59456
Total length (>= 1000 bp)	59456
Total length (>= 5000 bp)	22409
Total length (>= 10000 bp)	11301
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	13
Largest contig	11301
Total length	59456
GC (%)	57.77
N50	4482
N90	2822
auN	5627.6
L50	5
L90	11
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	1 + 1 part

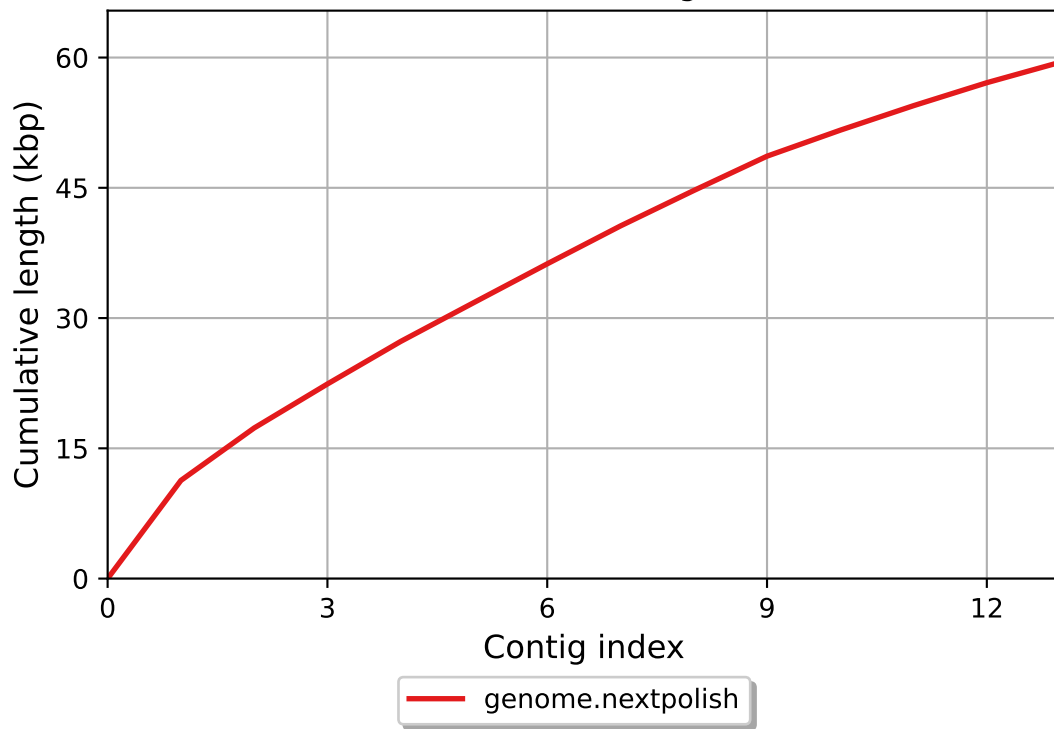
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

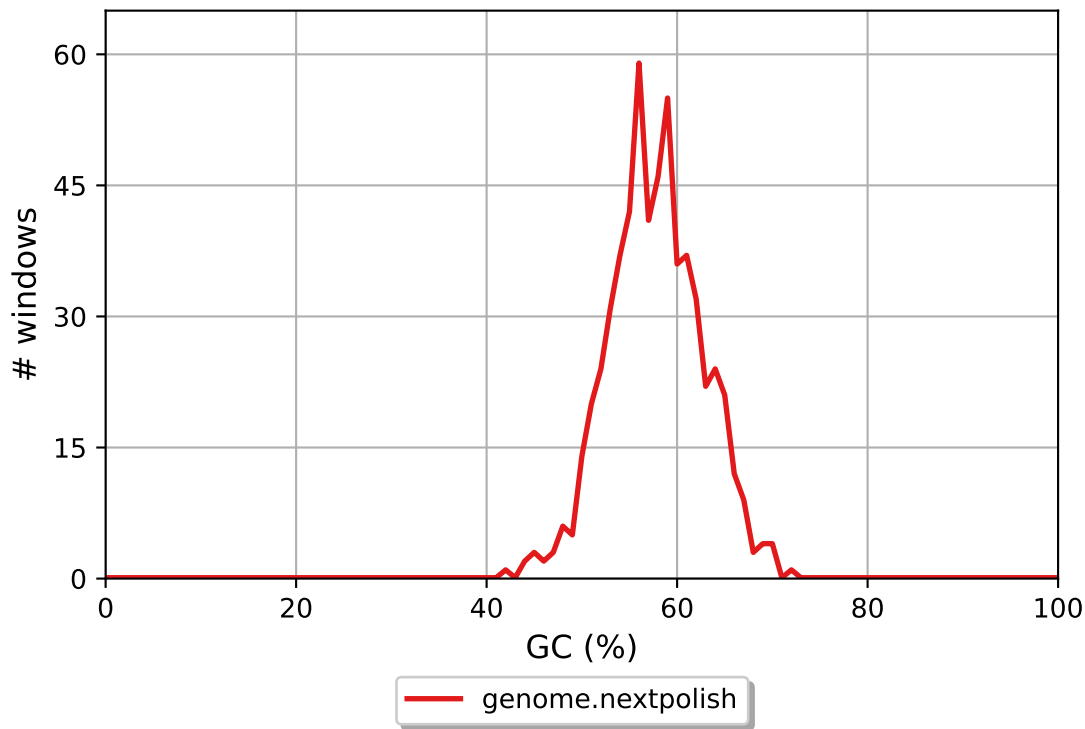


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

