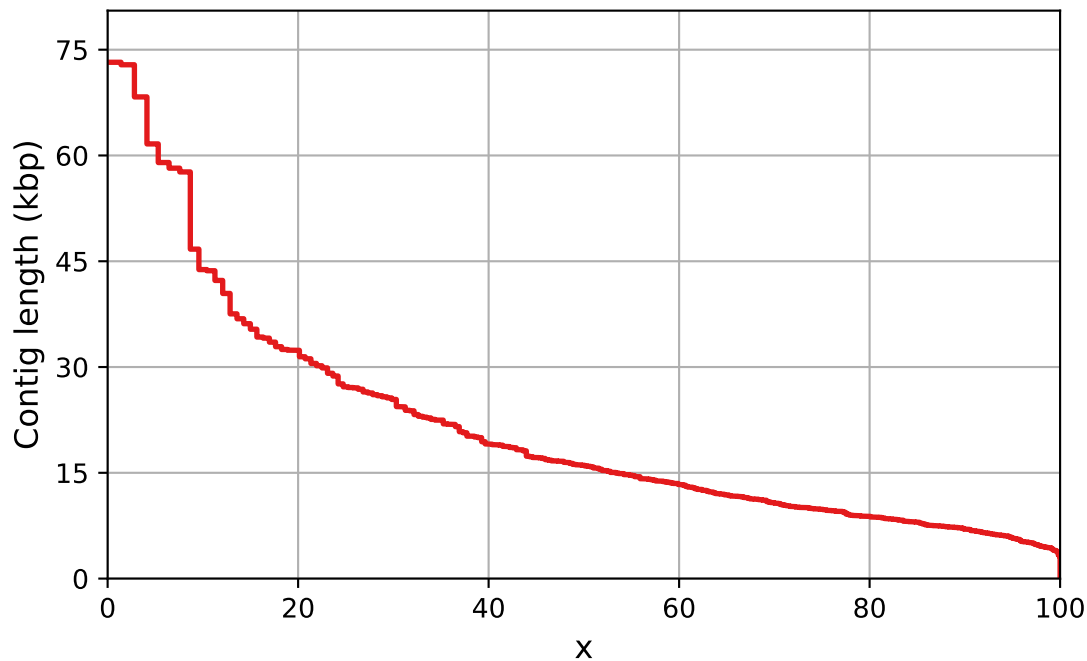


Report

	genome.nextpolish
# contigs (>= 0 bp)	389
# contigs (>= 1000 bp)	389
# contigs (>= 5000 bp)	355
# contigs (>= 10000 bp)	192
# contigs (>= 25000 bp)	42
# contigs (>= 50000 bp)	7
Total length (>= 0 bp)	5192955
Total length (>= 1000 bp)	5192955
Total length (>= 5000 bp)	5046597
Total length (>= 10000 bp)	3820765
Total length (>= 25000 bp)	1574100
Total length (>= 50000 bp)	450884
# contigs	389
Largest contig	73216
Total length	5192955
GC (%)	57.80
N50	16068
N90	7010
auN	21679.5
L50	95
L90	294
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	9 + 2 part

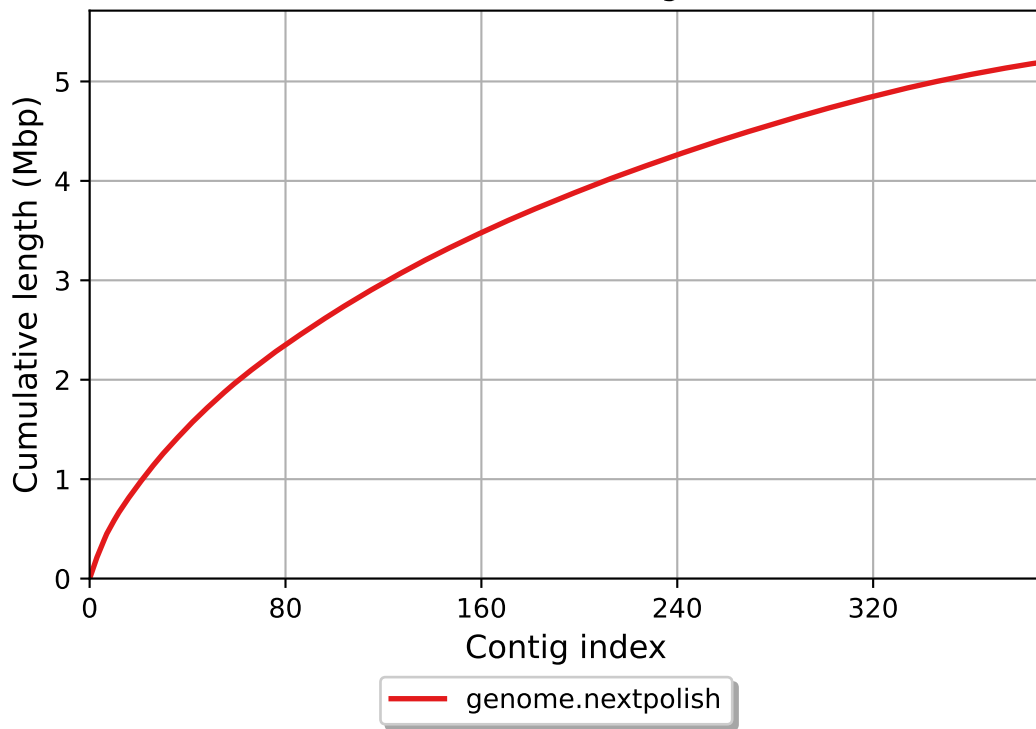
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

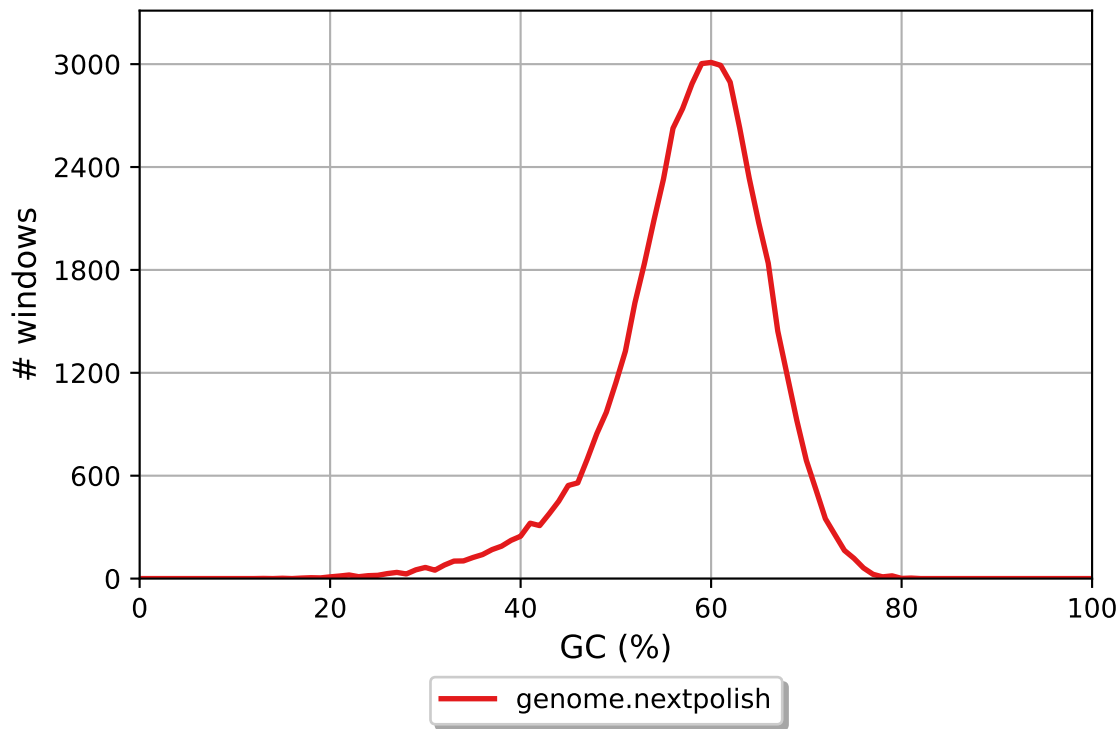


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

