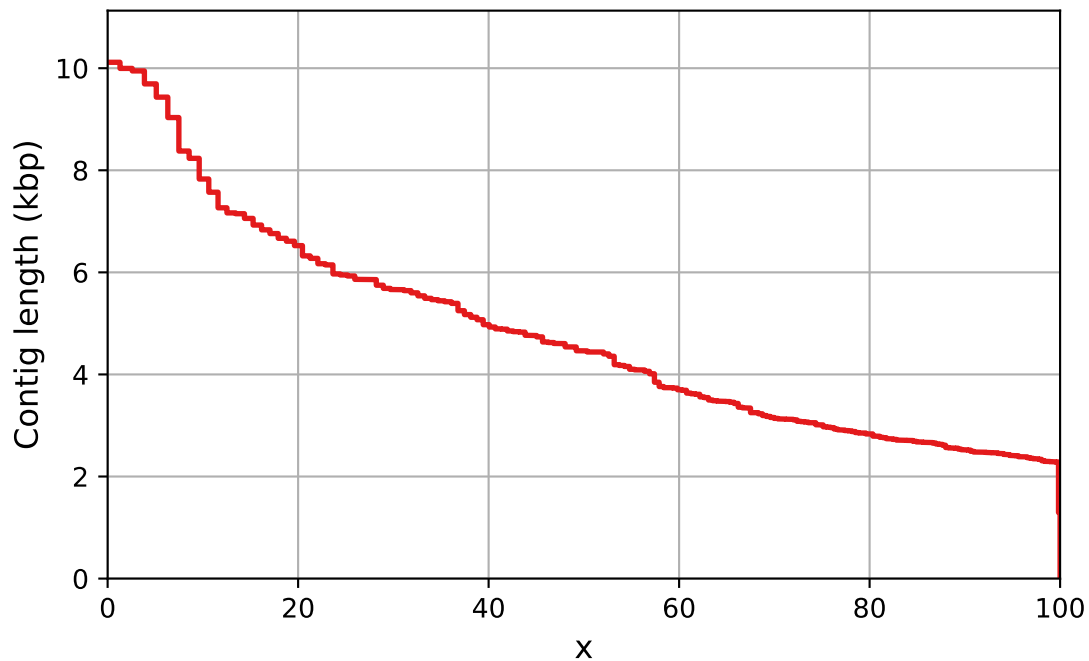


Report

	genome.nextpolish
# contigs (>= 0 bp)	193
# contigs (>= 1000 bp)	193
# contigs (>= 5000 bp)	46
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	778250
Total length (>= 1000 bp)	778250
Total length (>= 5000 bp)	306891
Total length (>= 10000 bp)	10116
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	193
Largest contig	10116
Total length	778250
GC (%)	50.67
N50	4462
N90	2521
auN	4812.0
L50	64
L90	161
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 1 part

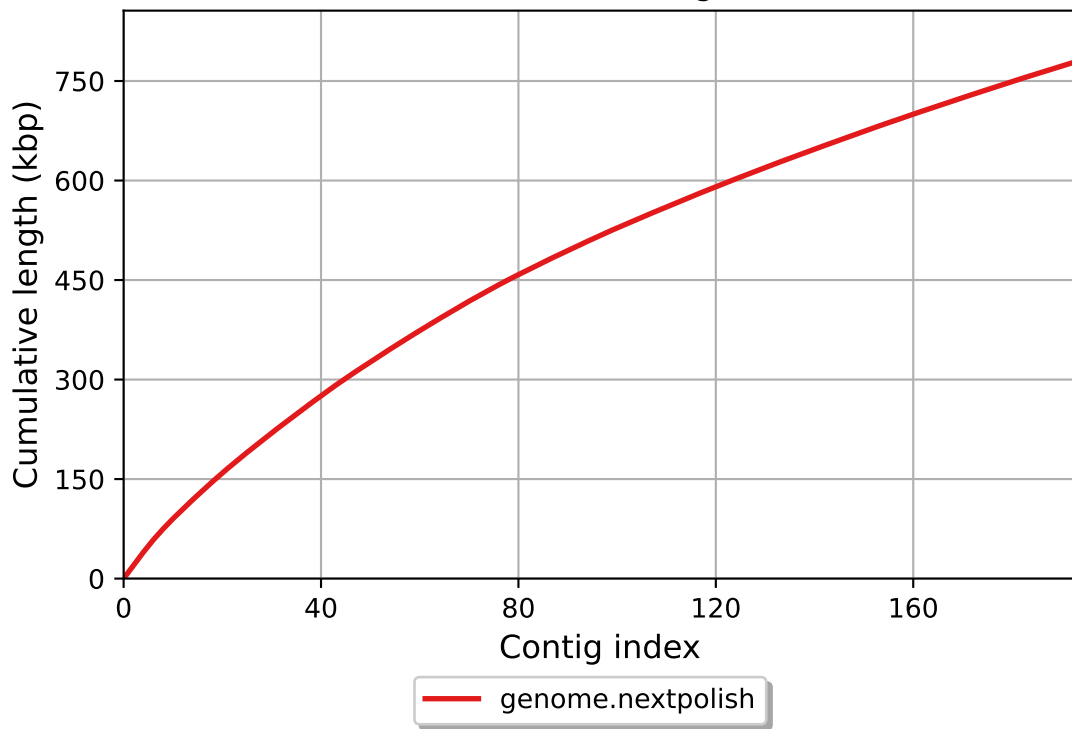
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

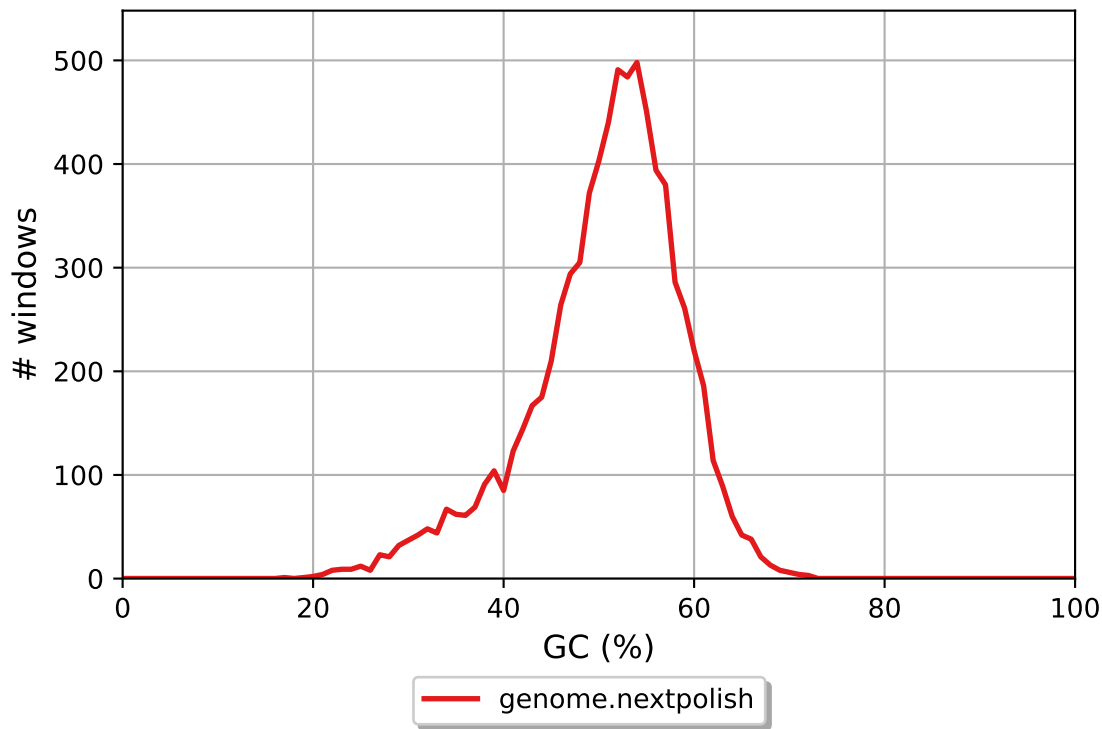


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

