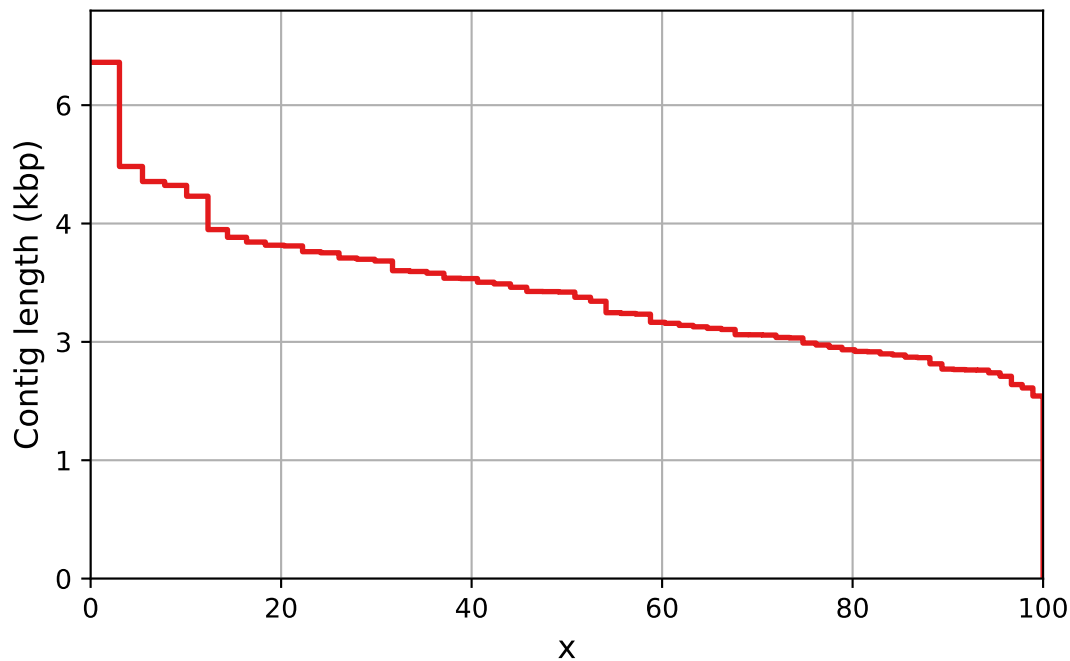


Report

	genome.nextpolish
# contigs (>= 0 bp)	62
# contigs (>= 1000 bp)	62
# contigs (>= 5000 bp)	3
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	216086
Total length (>= 1000 bp)	216086
Total length (>= 5000 bp)	16798
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	62
Largest contig	6543
Total length	216086
GC (%)	50.76
N50	3631
N90	2655
auN	3663.4
L50	26
L90	54
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 1 part

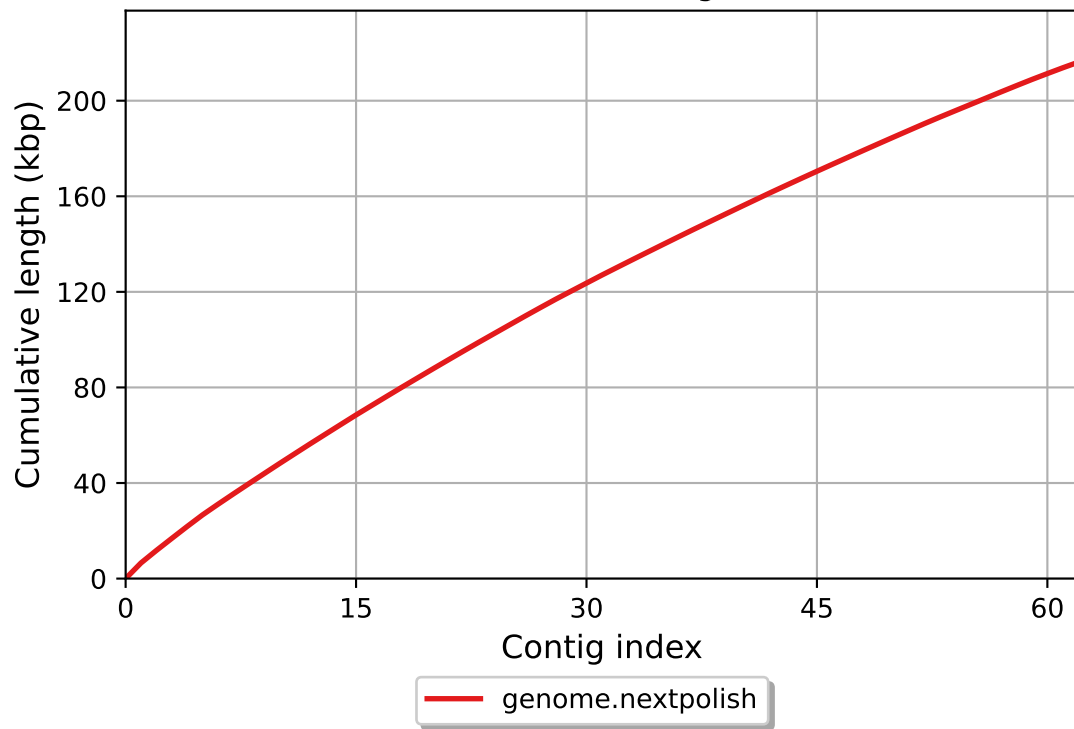
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

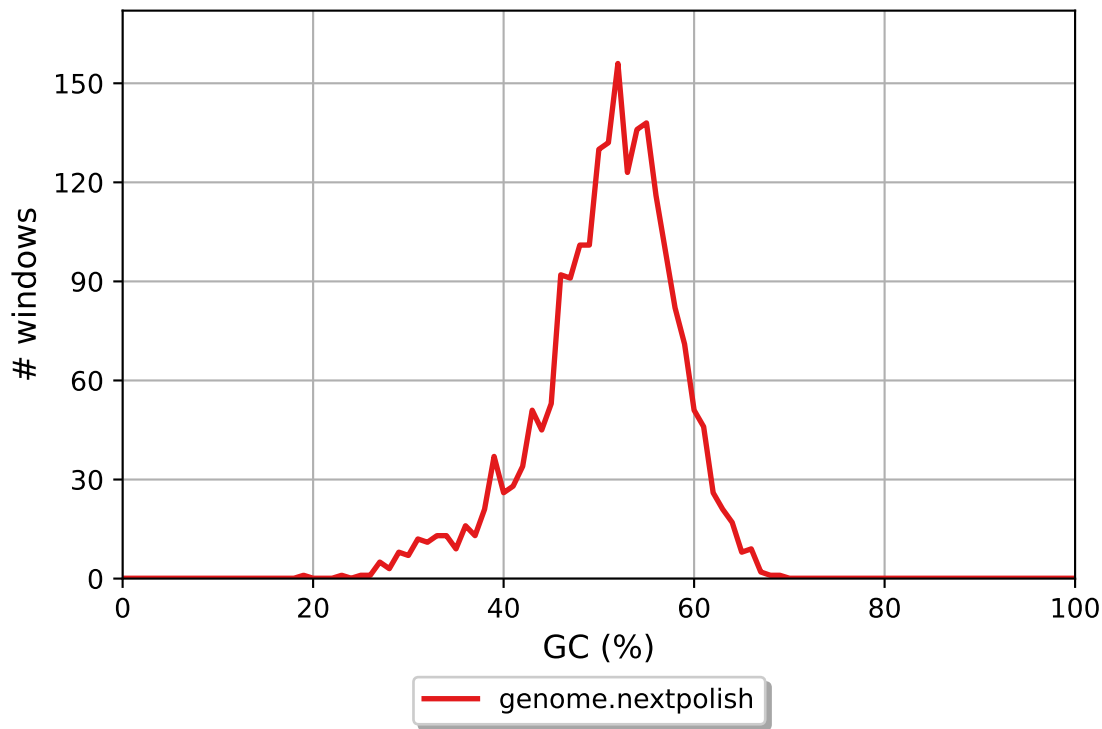


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

