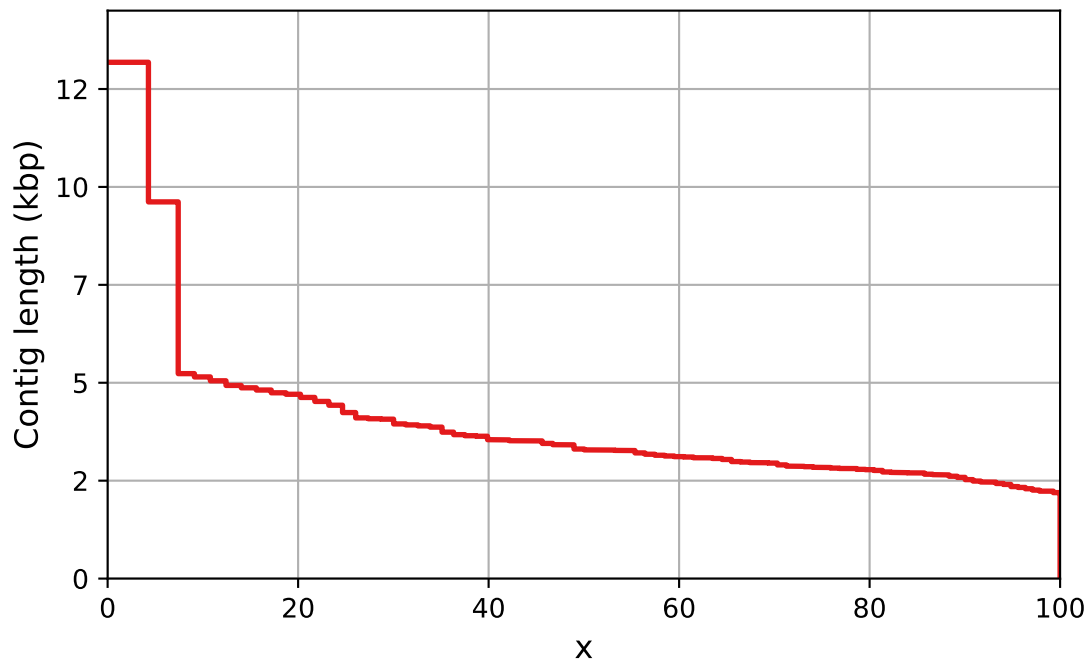


# Report

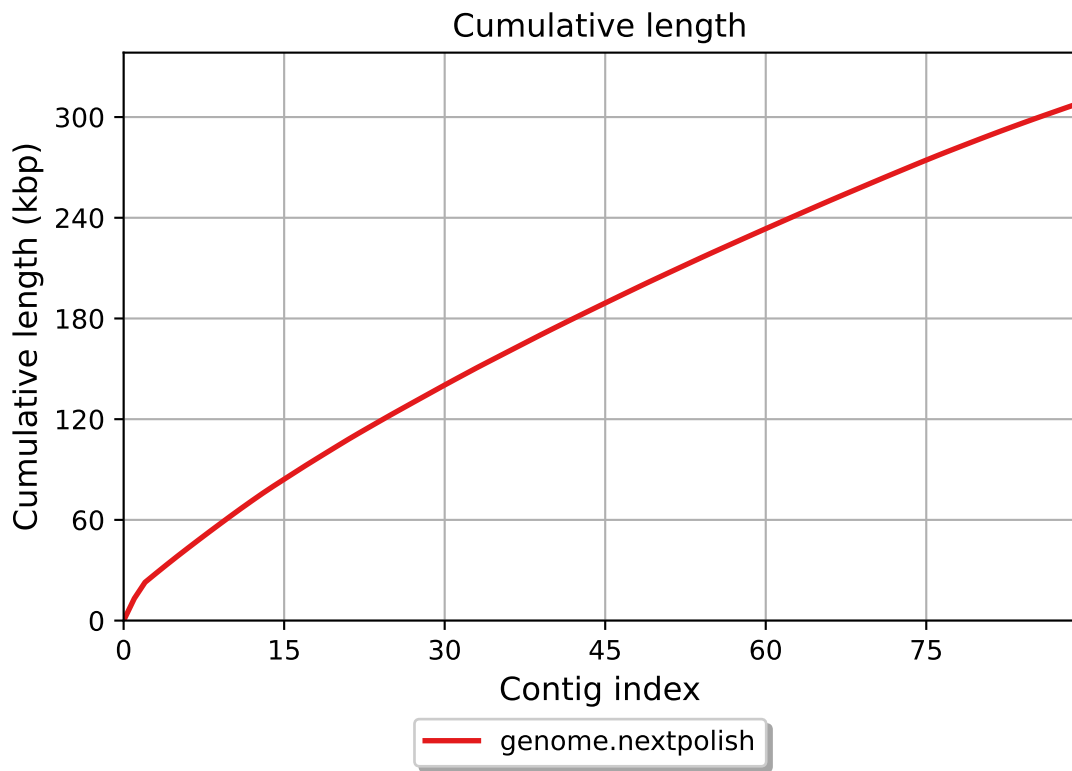
	genome.nextpolish
# contigs (>= 0 bp)	89
# contigs (>= 1000 bp)	89
# contigs (>= 5000 bp)	5
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	307623
Total length (>= 1000 bp)	307623
Total length (>= 5000 bp)	38232
Total length (>= 10000 bp)	13182
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	89
Largest contig	13182
Total length	307623
GC (%)	53.46
N50	3308
N90	2584
auN	4061.0
L50	34
L90	76
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

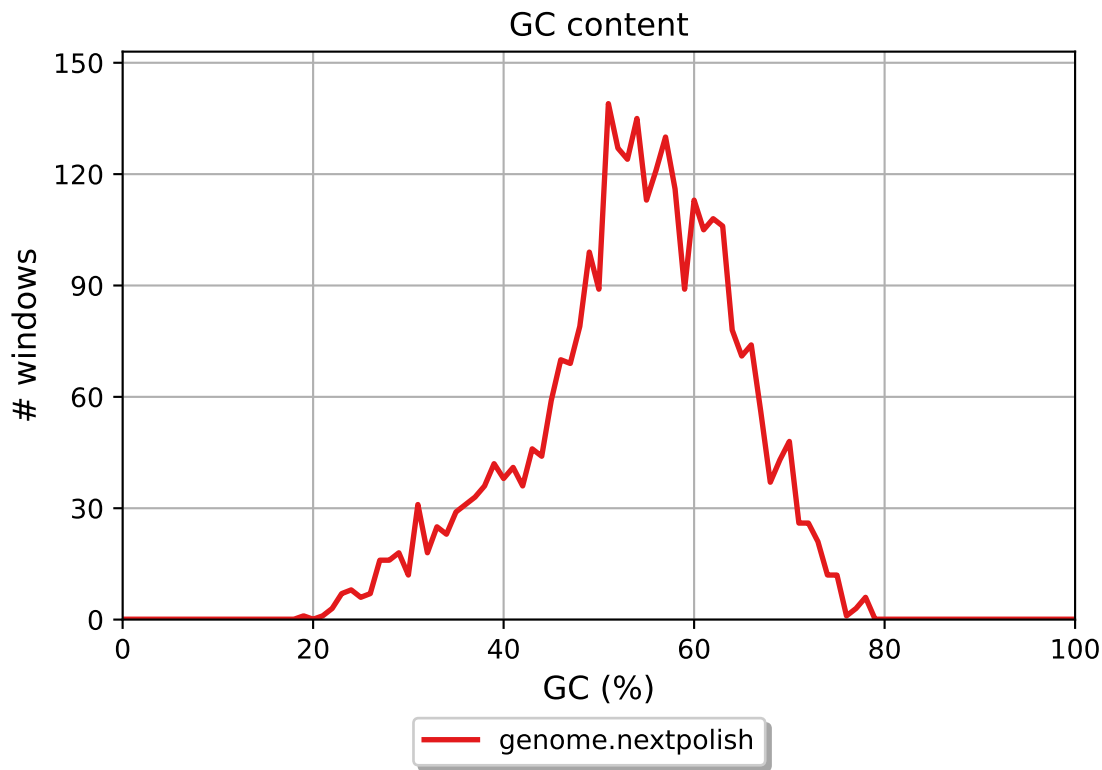
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

