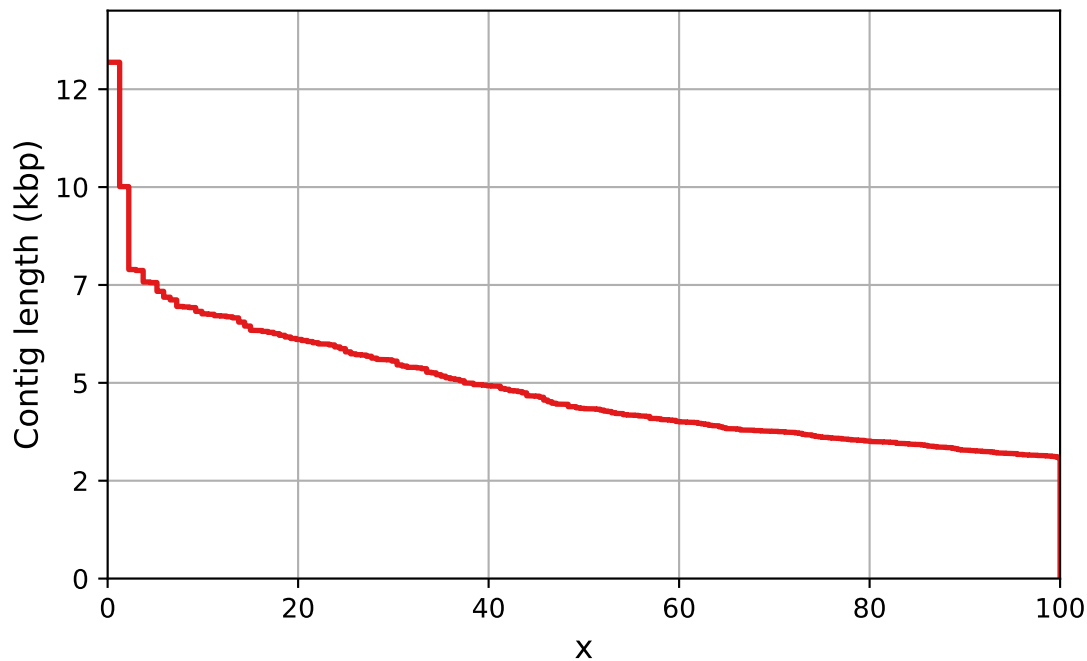


# Report

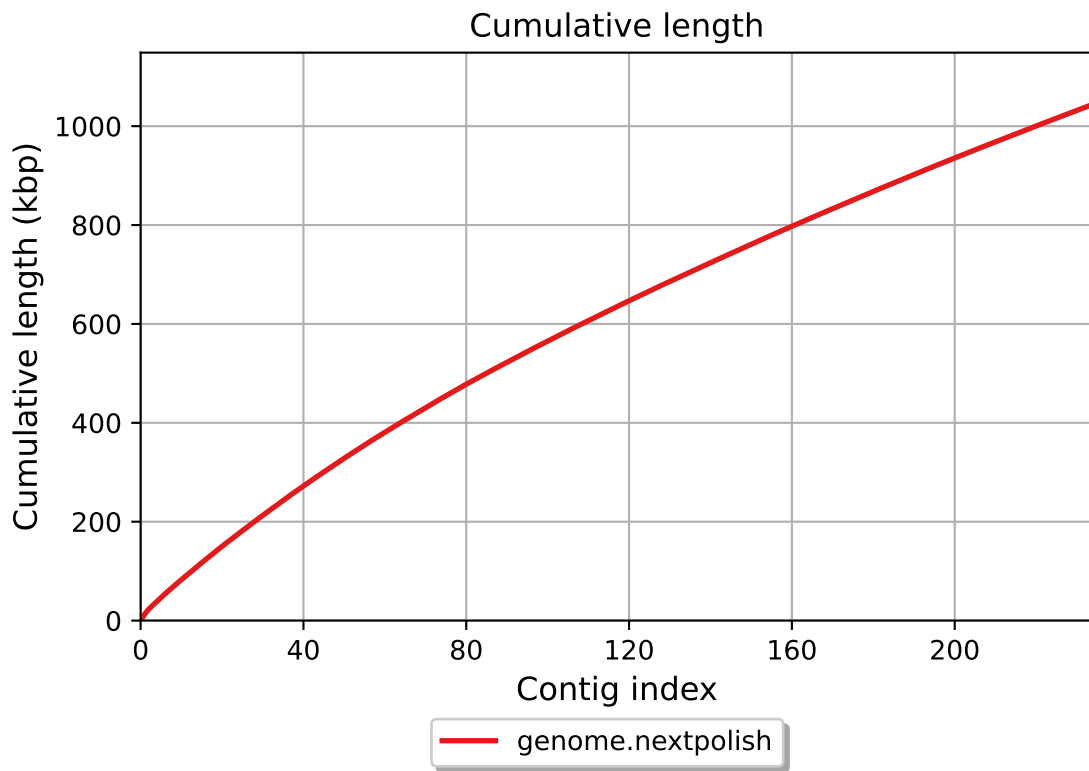
	genome.nextpolish
# contigs (>= 0 bp)	234
# contigs (>= 1000 bp)	234
# contigs (>= 5000 bp)	62
# contigs (>= 10000 bp)	2
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1044278
Total length (>= 1000 bp)	1044278
Total length (>= 5000 bp)	391021
Total length (>= 10000 bp)	23200
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	234
Largest contig	13188
Total length	1044278
GC (%)	39.95
N50	4348
N90	3278
auN	4872.2
L50	90
L90	202
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 2 part

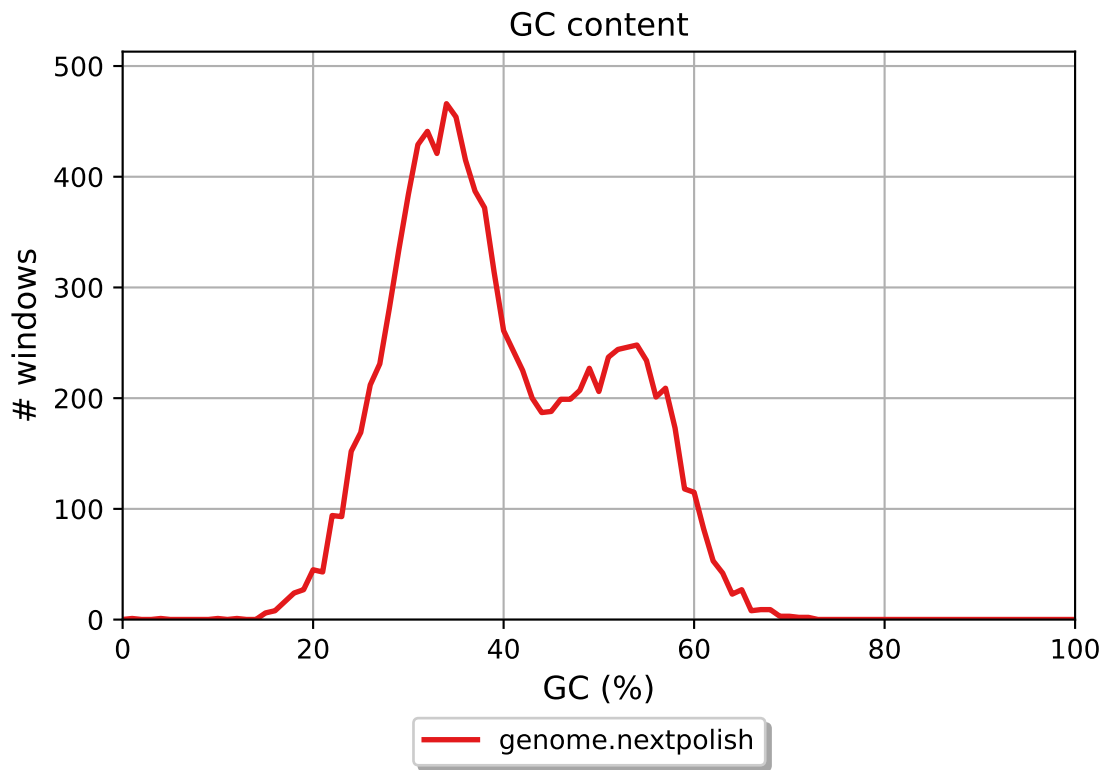
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

