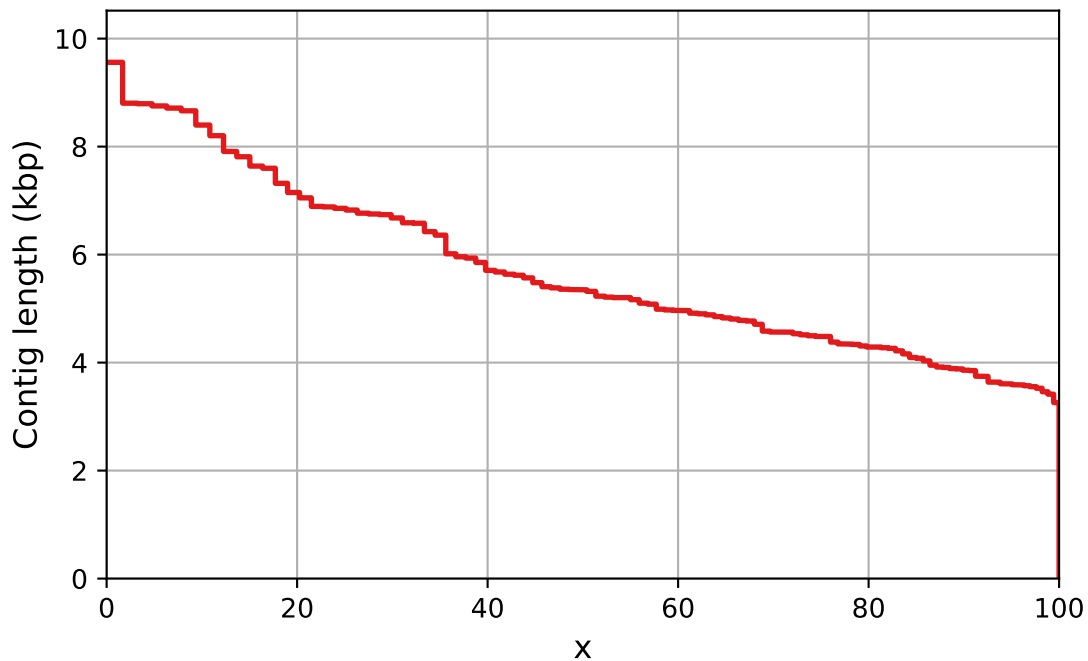


# Report

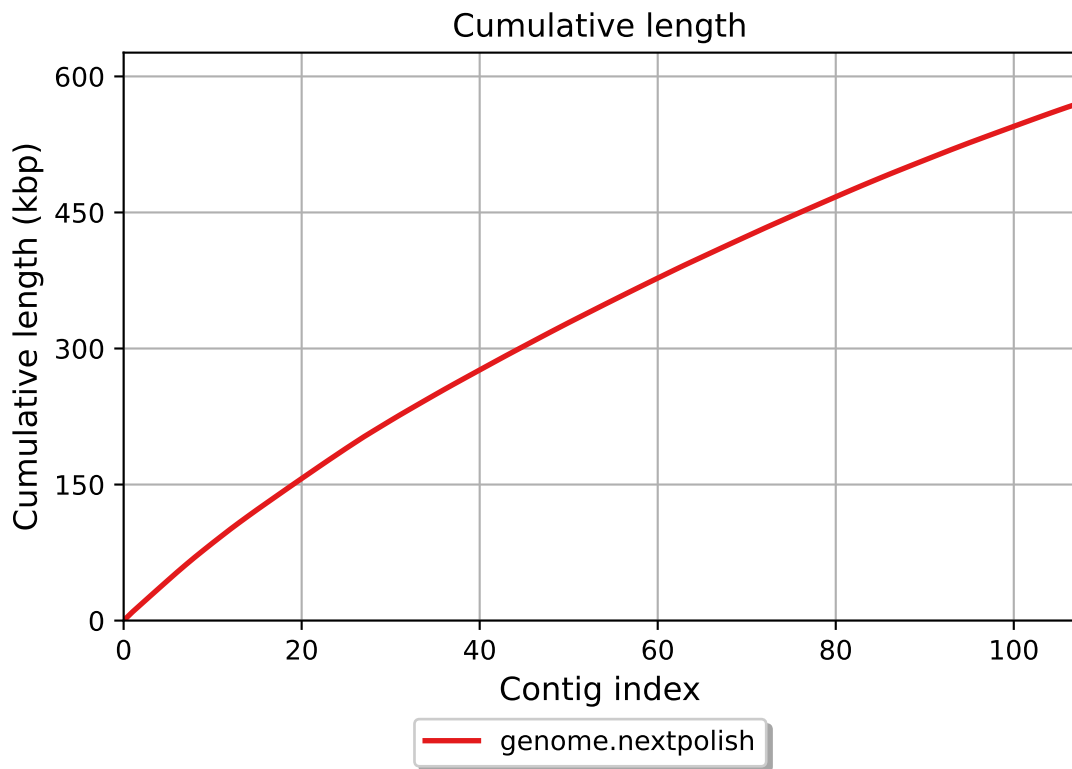
	genome.nextpolish
# contigs (>= 0 bp)	107
# contigs (>= 1000 bp)	107
# contigs (>= 5000 bp)	50
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	569290
Total length (>= 1000 bp)	569290
Total length (>= 5000 bp)	328539
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	107
Largest contig	9561
Total length	569290
GC (%)	50.50
N50	5348
N90	3859
auN	5734.6
L50	42
L90	92
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

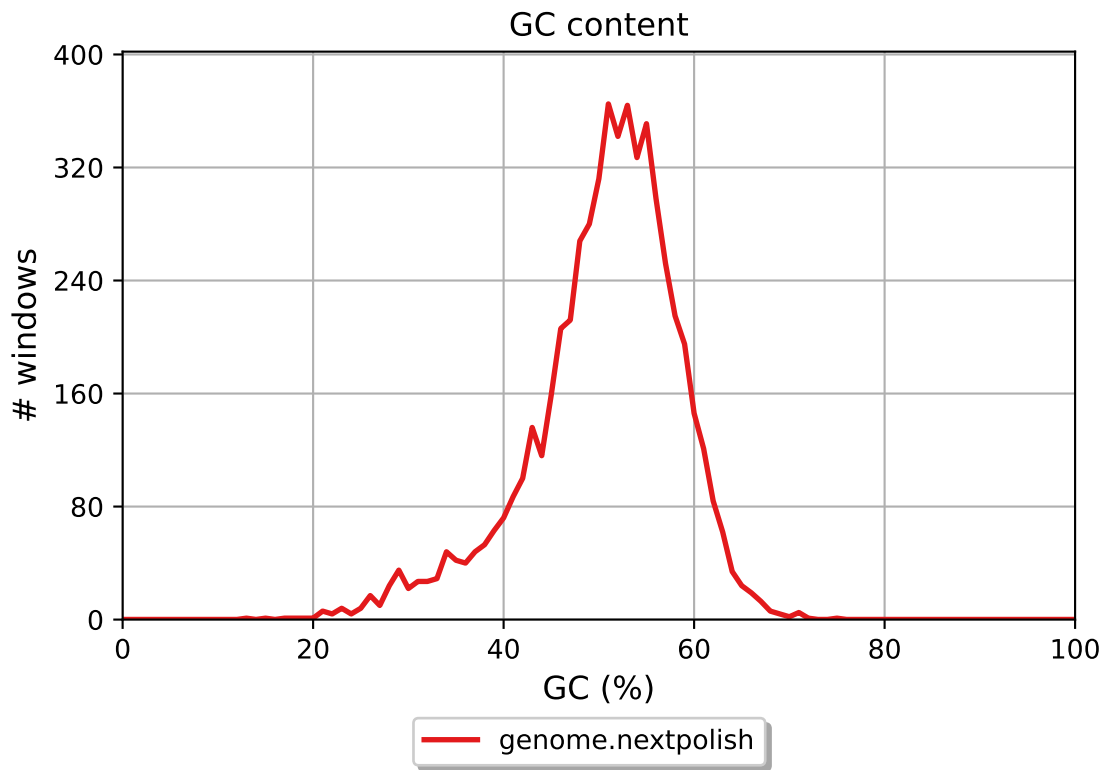
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

