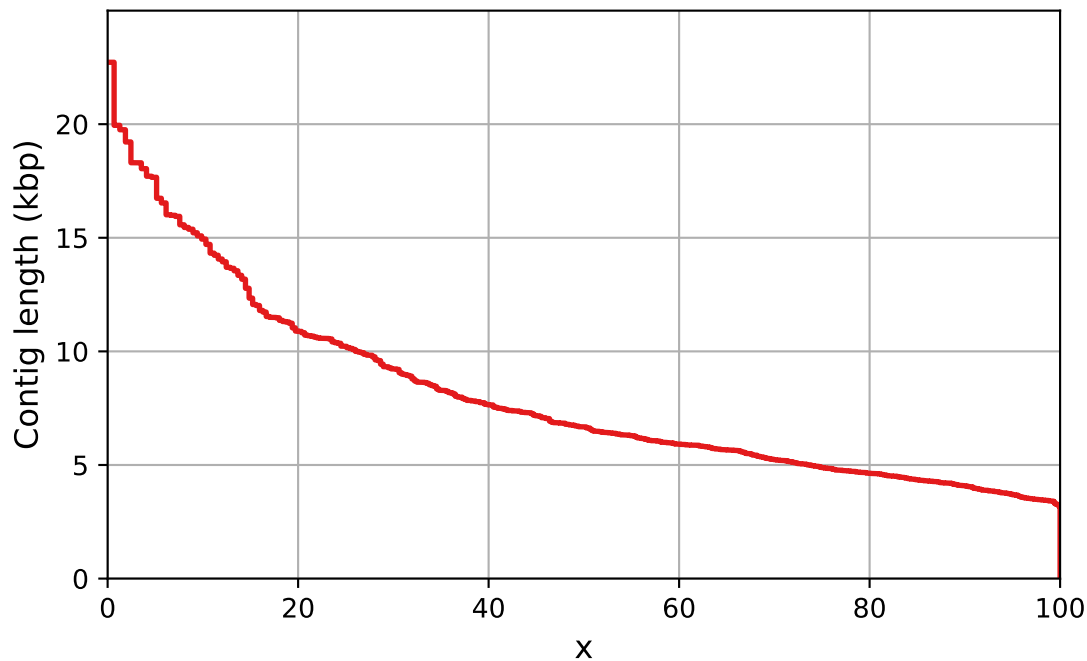


Report

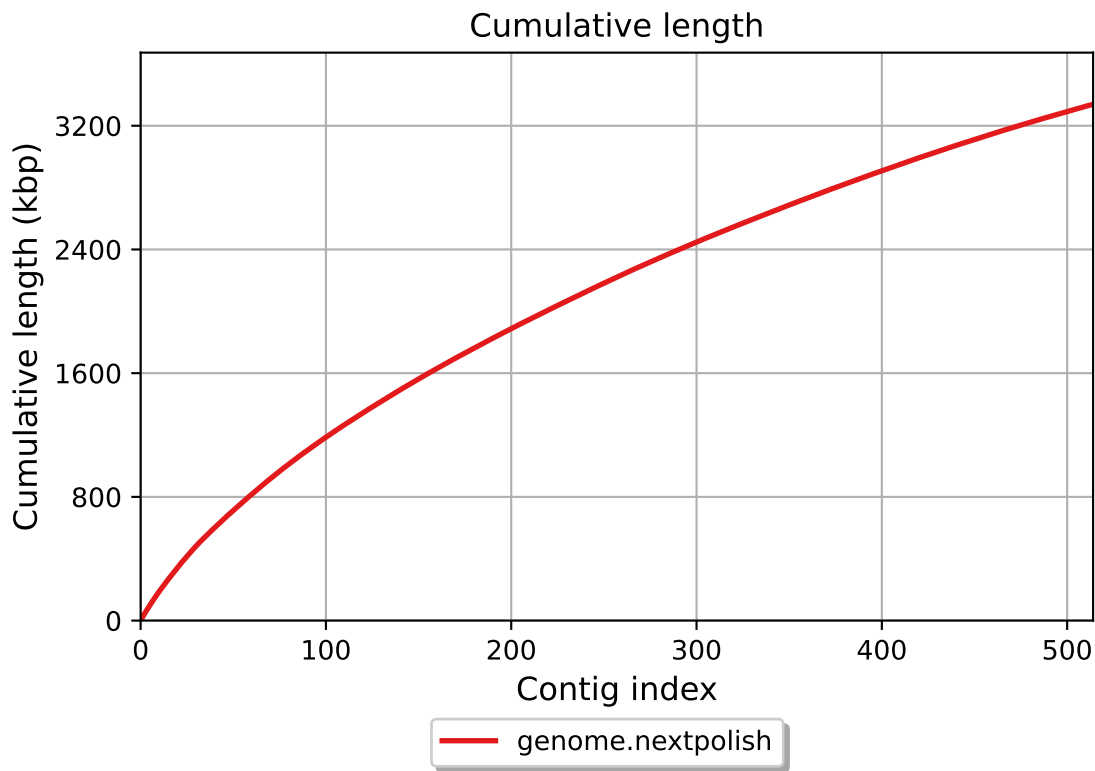
	genome.nextpolish
# contigs (>= 0 bp)	514
# contigs (>= 1000 bp)	514
# contigs (>= 5000 bp)	302
# contigs (>= 10000 bp)	66
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	3338879
Total length (>= 1000 bp)	3338879
Total length (>= 5000 bp)	2457946
Total length (>= 10000 bp)	878339
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	514
Largest contig	22724
Total length	3338879
GC (%)	38.13
N50	6677
N90	4082
auN	8073.5
L50	166
L90	424
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 1 part

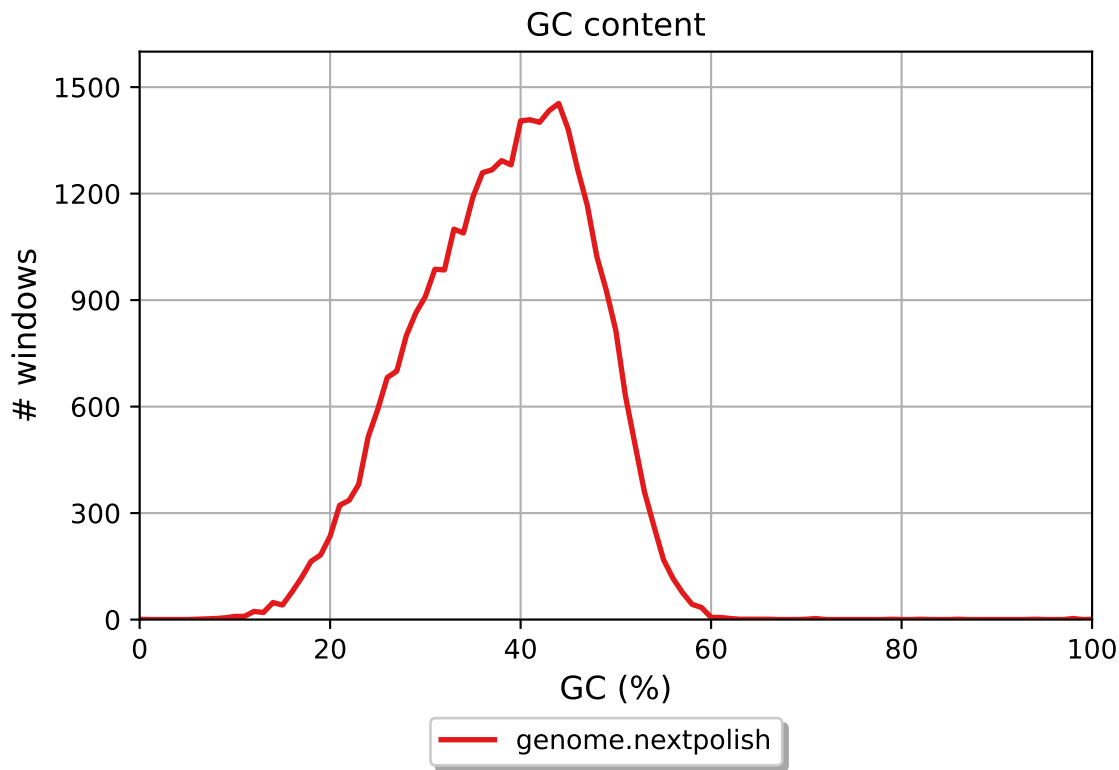
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

