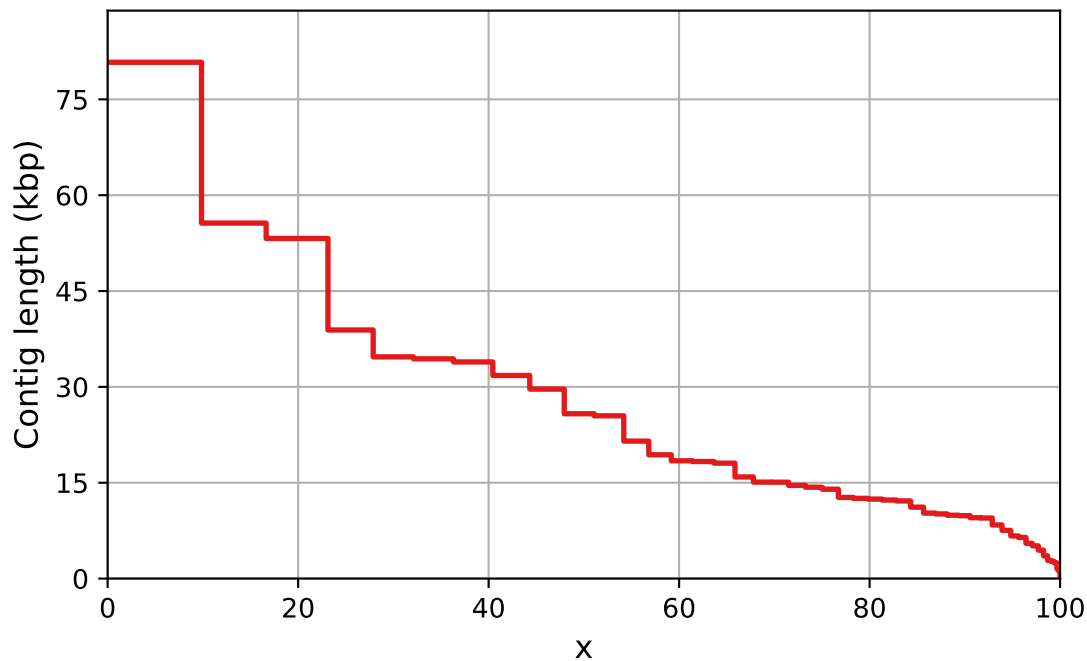


Report

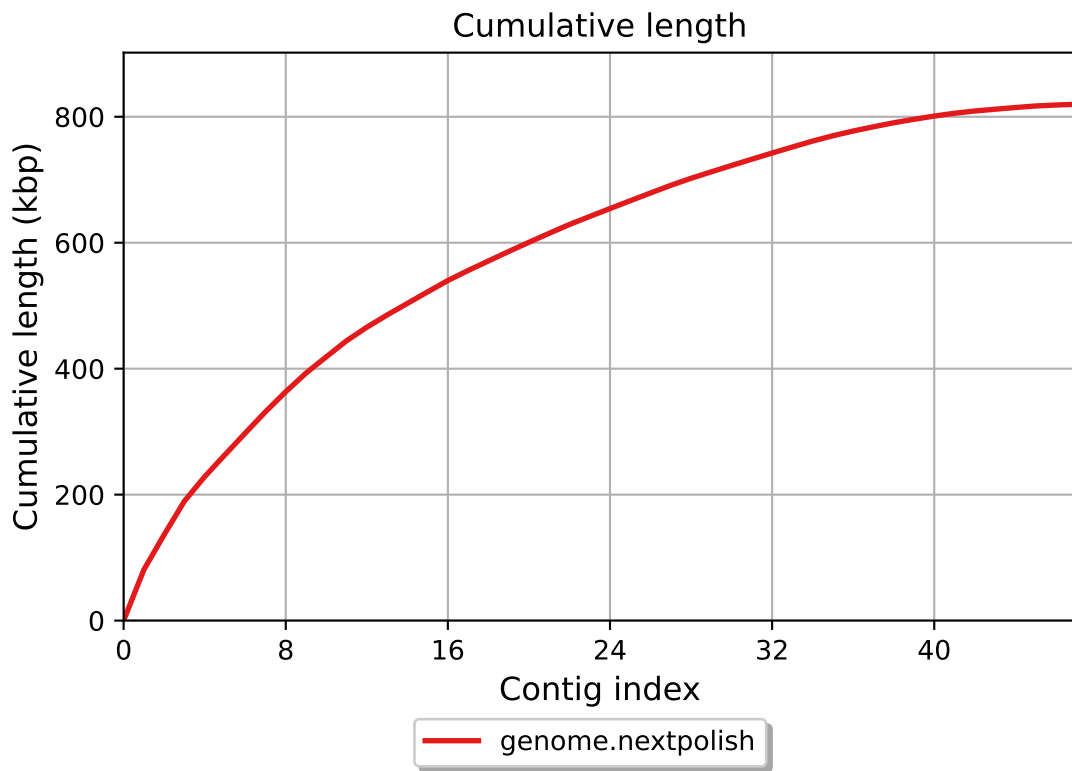
	genome.nextpolish
# contigs (>= 0 bp)	47
# contigs (>= 1000 bp)	47
# contigs (>= 5000 bp)	40
# contigs (>= 10000 bp)	30
# contigs (>= 25000 bp)	11
# contigs (>= 50000 bp)	3
Total length (>= 0 bp)	819826
Total length (>= 1000 bp)	819826
Total length (>= 5000 bp)	801097
Total length (>= 10000 bp)	722656
Total length (>= 25000 bp)	444280
Total length (>= 50000 bp)	189670
# contigs	47
Largest contig	80803
Total length	819826
GC (%)	40.17
N50	25796
N90	9846
auN	31326.5
L50	10
L90	32
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 0 part

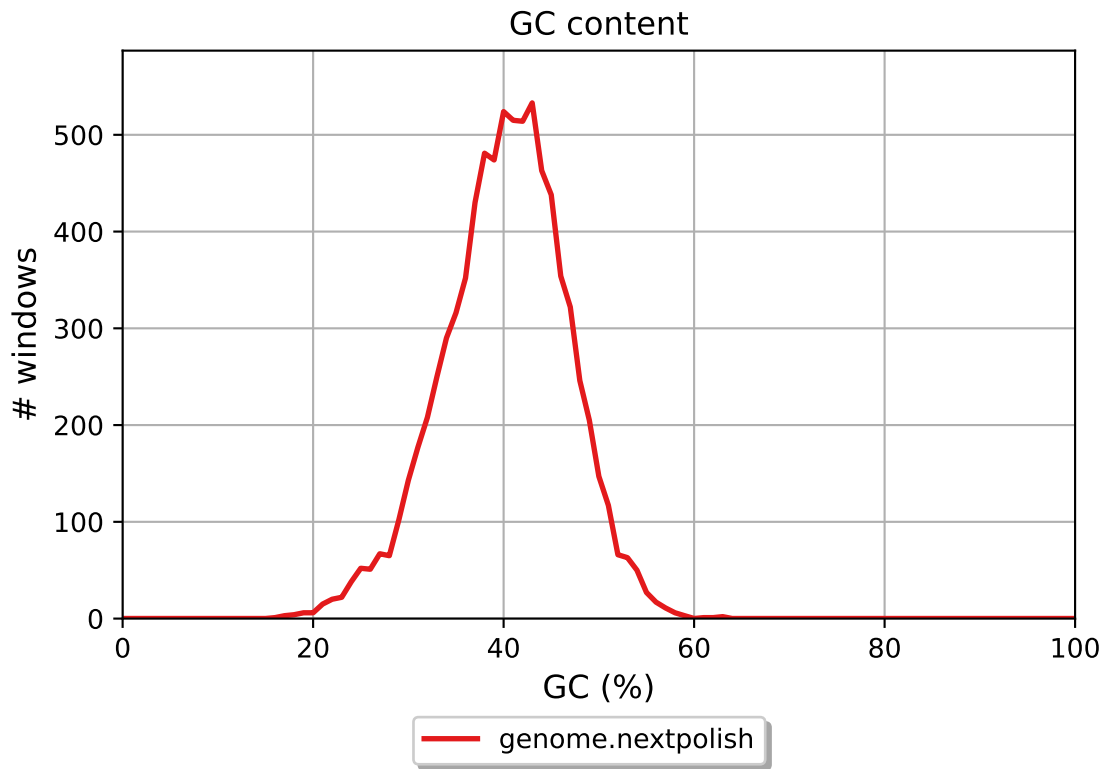
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

