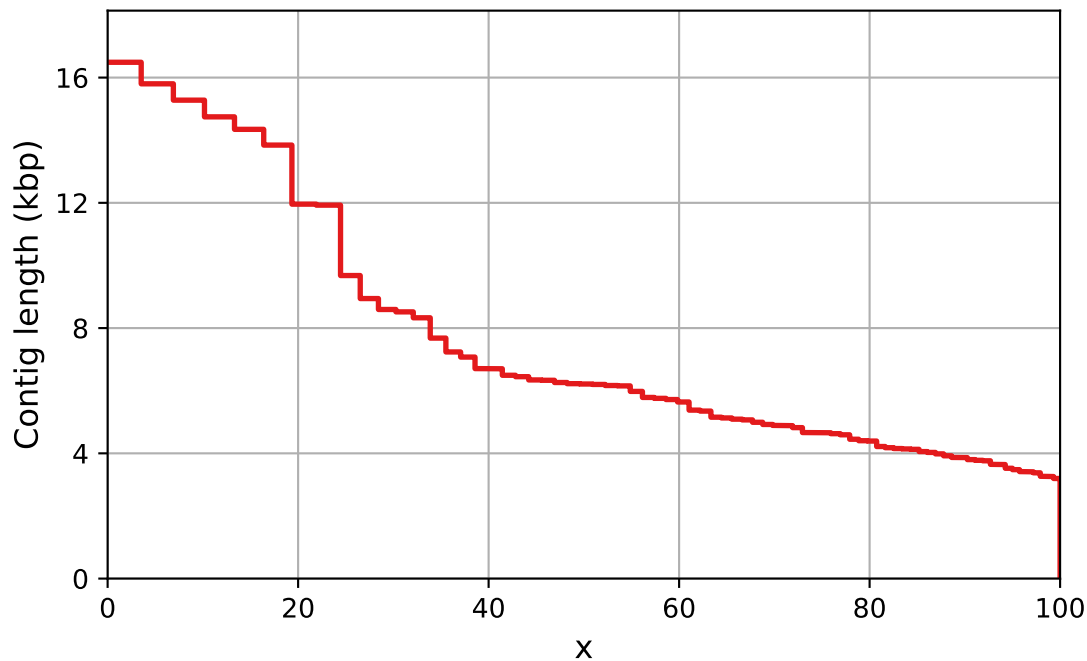


# Report

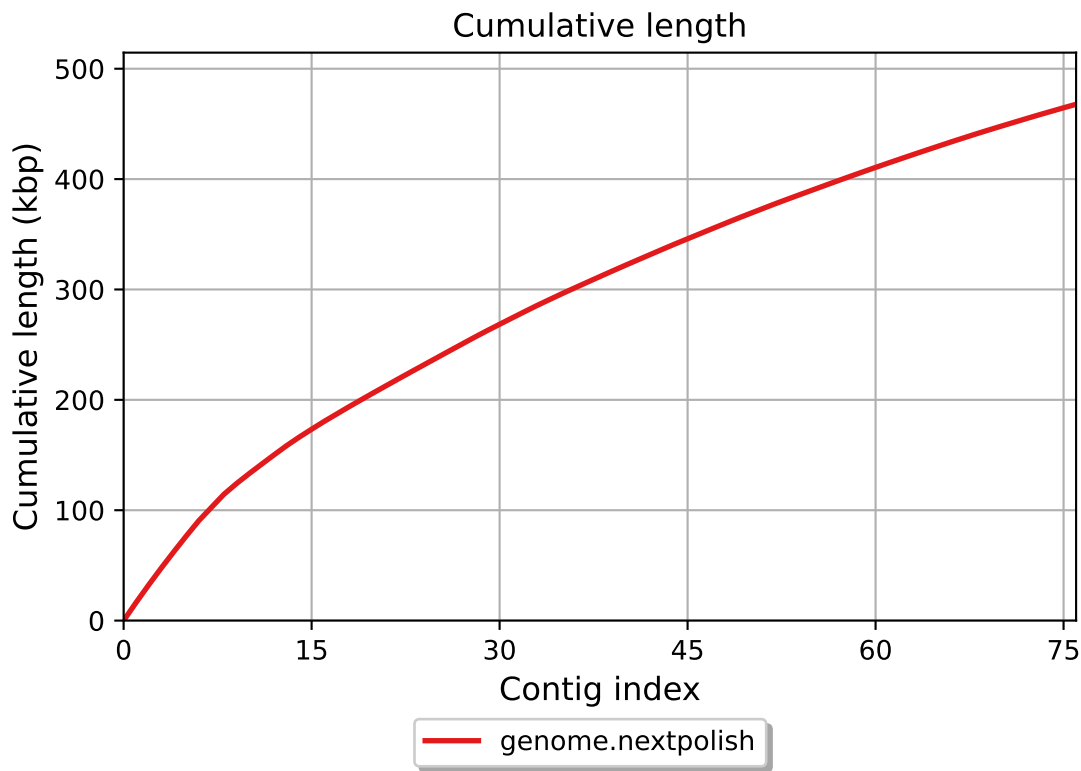
	genome.nextpolish
# contigs (>= 0 bp)	76
# contigs (>= 1000 bp)	76
# contigs (>= 5000 bp)	39
# contigs (>= 10000 bp)	8
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	467796
Total length (>= 1000 bp)	467796
Total length (>= 5000 bp)	316739
Total length (>= 10000 bp)	114390
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	76
Largest contig	16489
Total length	467796
GC (%)	33.02
N50	6214
N90	3864
auN	7795.6
L50	25
L90	63
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 1 part

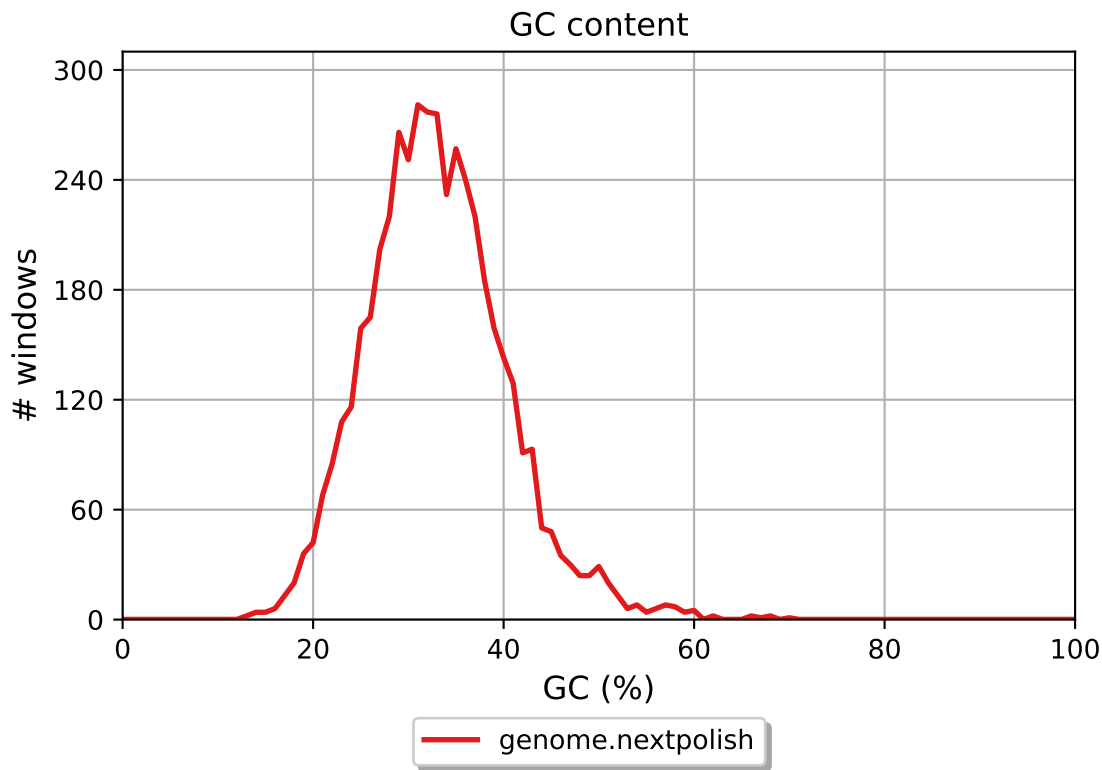
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

