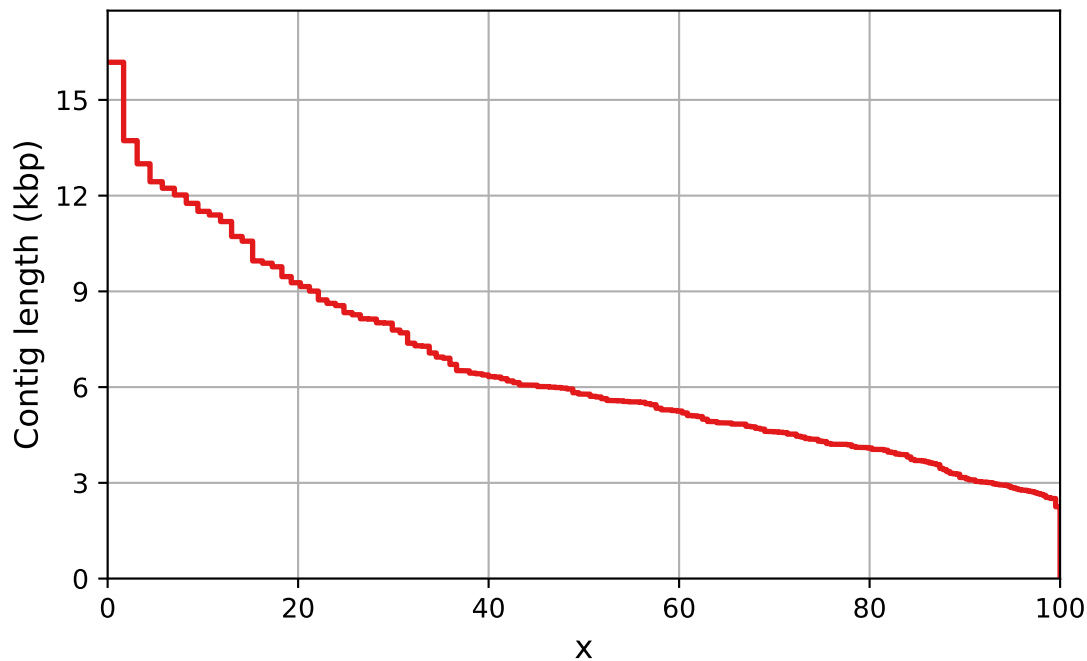


# Report

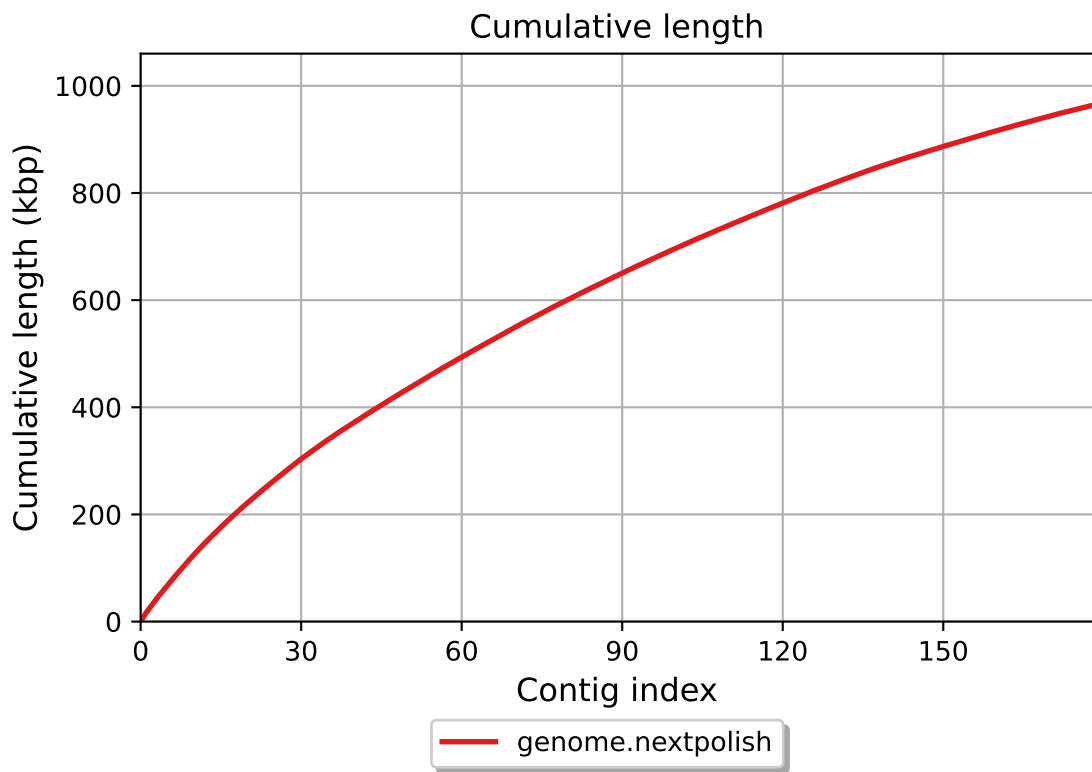
	genome.nextpolish
# contigs (>= 0 bp)	178
# contigs (>= 1000 bp)	178
# contigs (>= 5000 bp)	80
# contigs (>= 10000 bp)	12
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	963773
Total length (>= 1000 bp)	963773
Total length (>= 5000 bp)	601862
Total length (>= 10000 bp)	146728
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	178
Largest contig	16179
Total length	963773
GC (%)	42.18
N50	5783
N90	3167
auN	6629.0
L50	58
L90	144
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 1 part

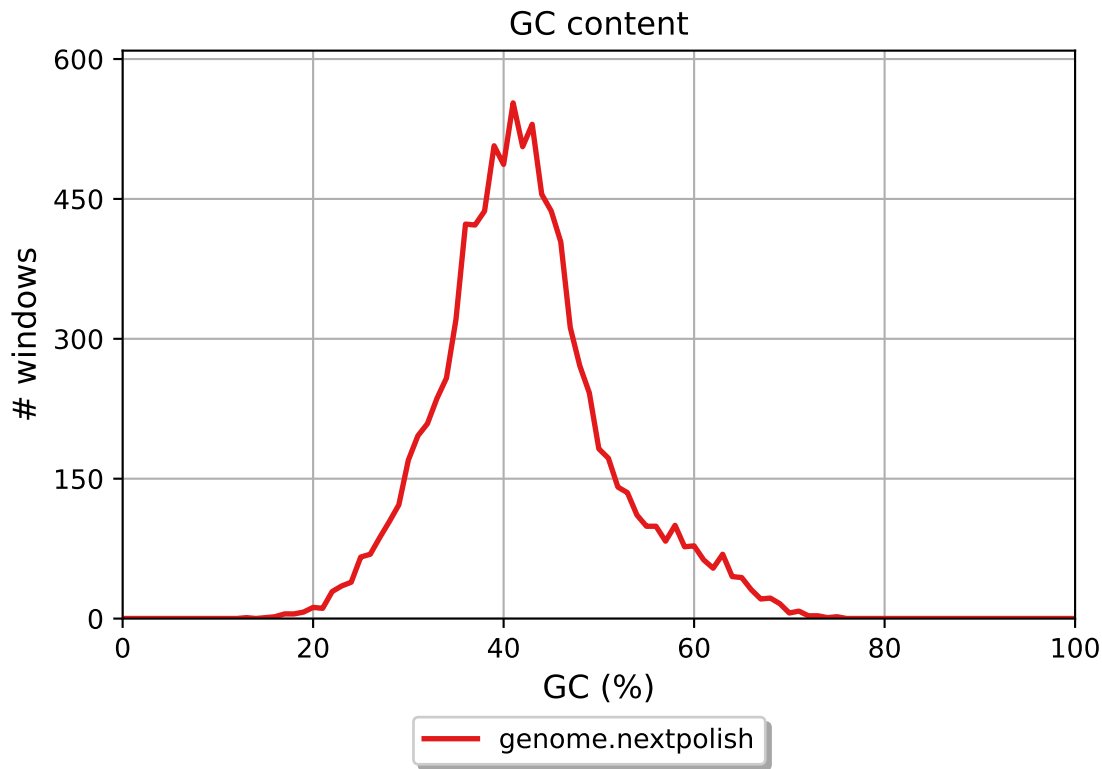
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

