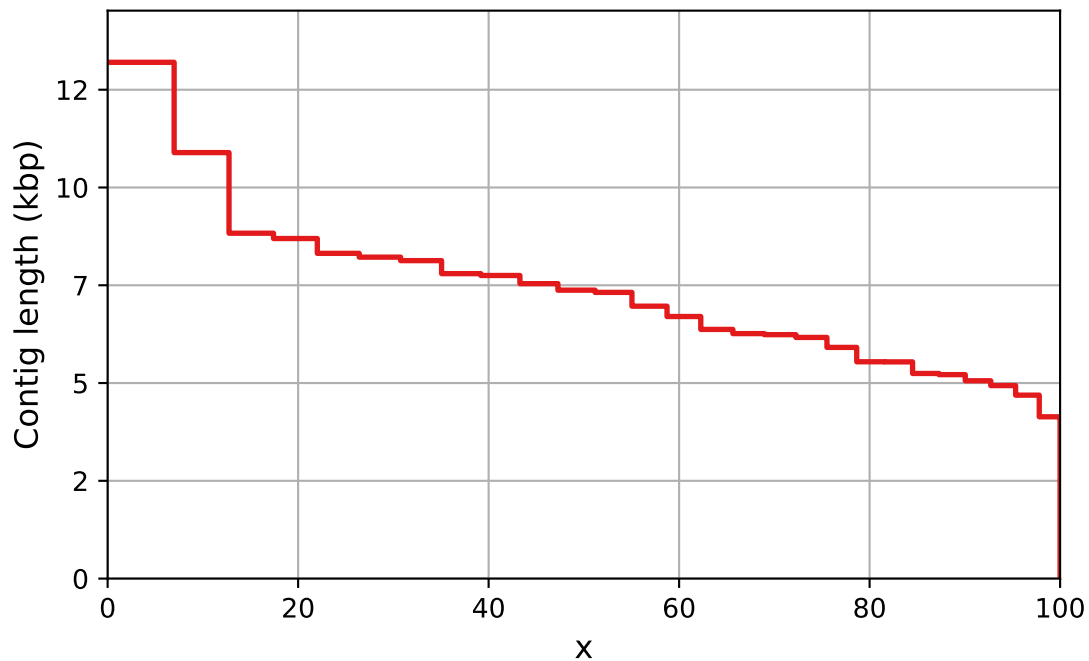


Report

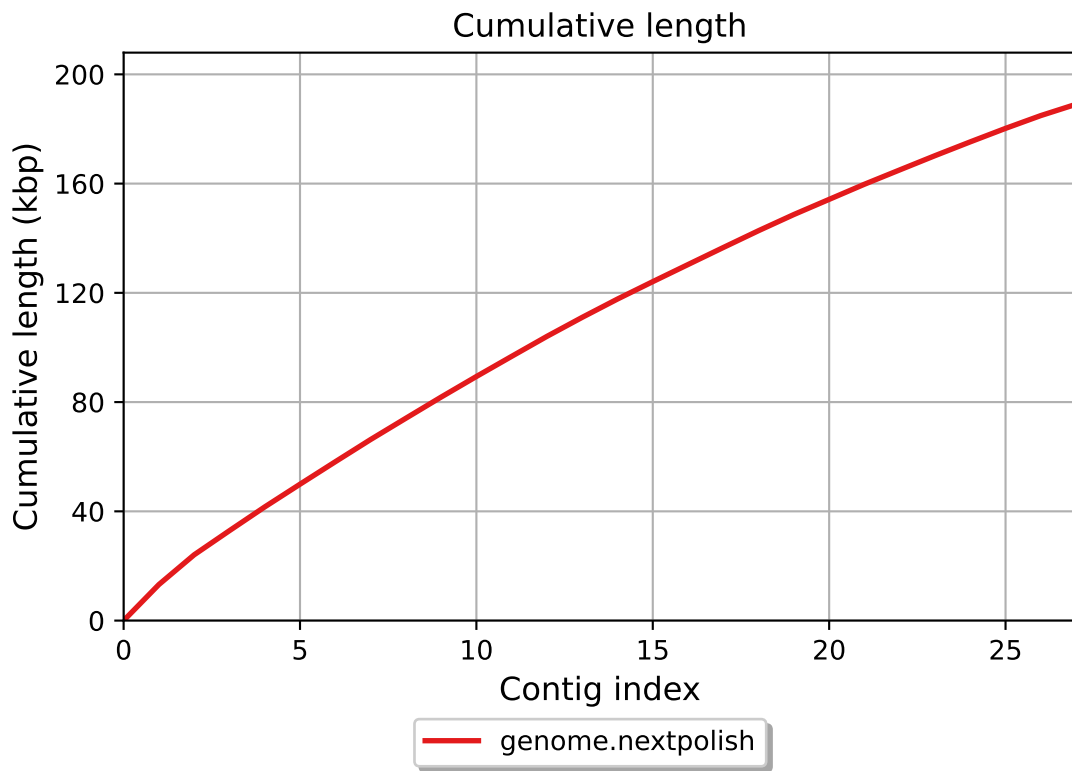
	genome.nextpolish
# contigs (>= 0 bp)	27
# contigs (>= 1000 bp)	27
# contigs (>= 5000 bp)	24
# contigs (>= 10000 bp)	2
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	189045
Total length (>= 1000 bp)	189045
Total length (>= 5000 bp)	175283
Total length (>= 10000 bp)	24093
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	27
Largest contig	13201
Total length	189045
GC (%)	46.96
N50	7375
N90	5215
auN	7539.2
L50	11
L90	23
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

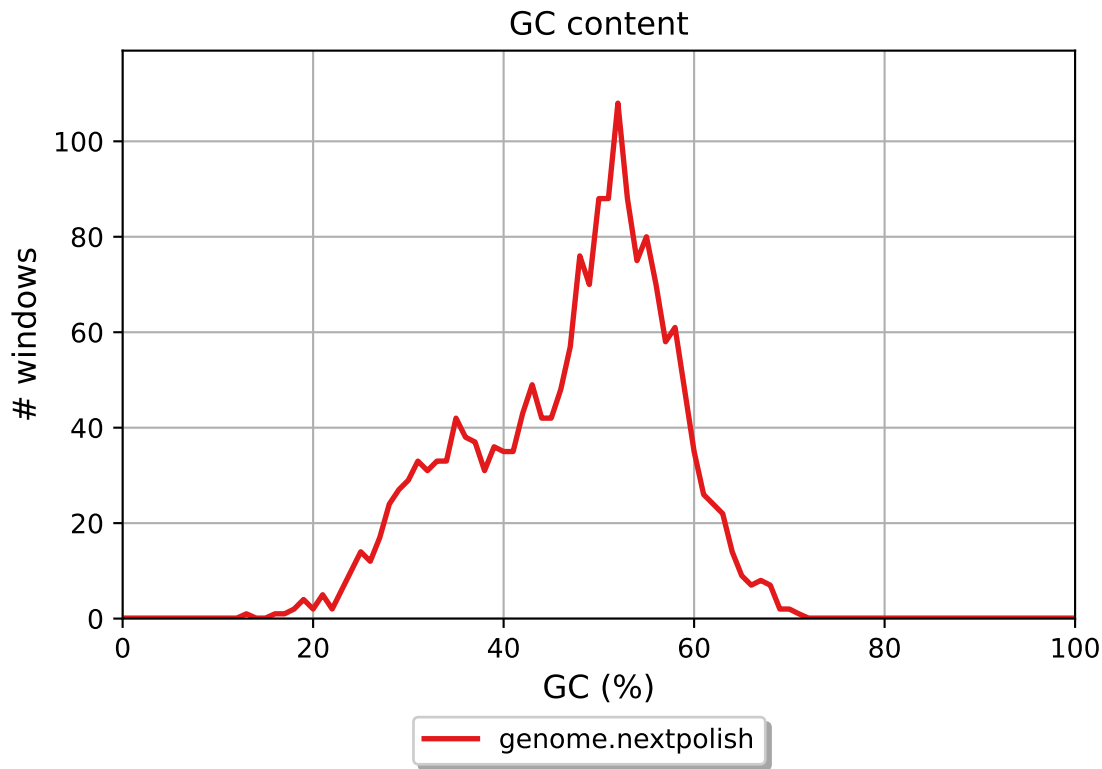
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

