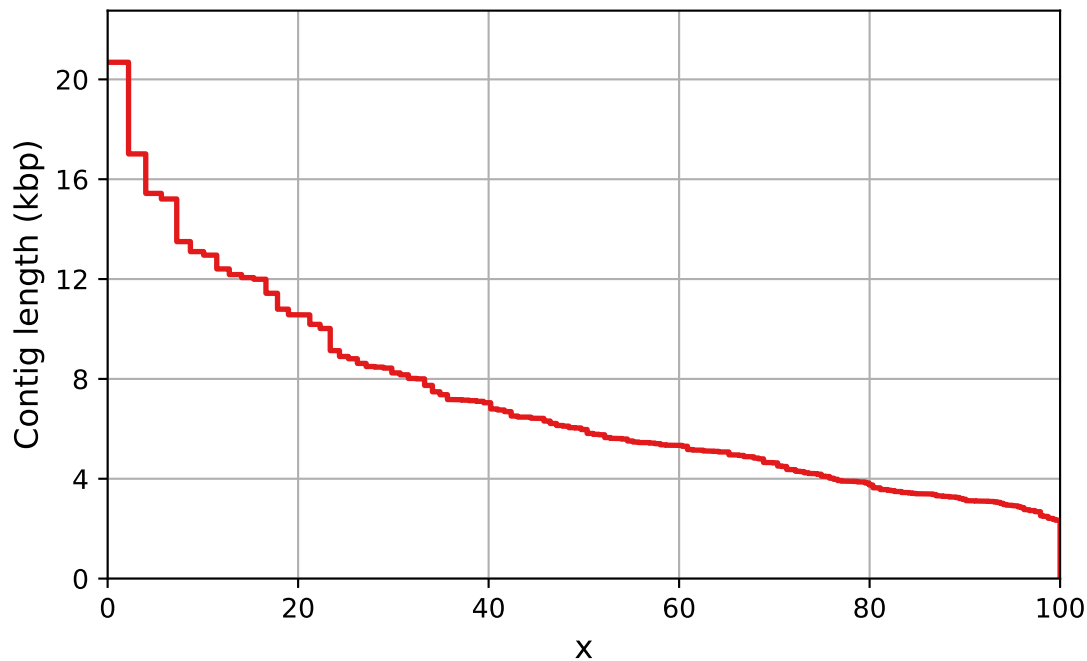


Report

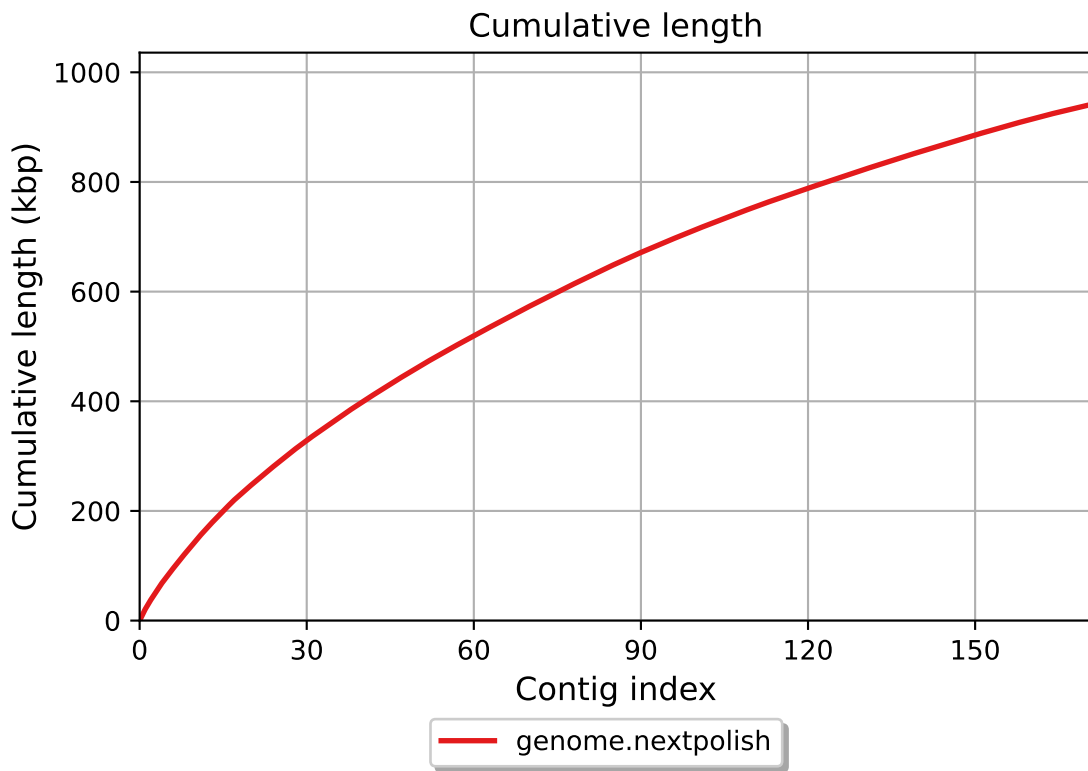
	genome.nextpolish
# contigs (>= 0 bp)	171
# contigs (>= 1000 bp)	171
# contigs (>= 5000 bp)	78
# contigs (>= 10000 bp)	17
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	941700
Total length (>= 1000 bp)	941700
Total length (>= 5000 bp)	614242
Total length (>= 10000 bp)	220097
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	171
Largest contig	20685
Total length	941700
GC (%)	41.12
N50	5972
N90	3178
auN	7225.0
L50	52
L90	138
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	6 + 0 part

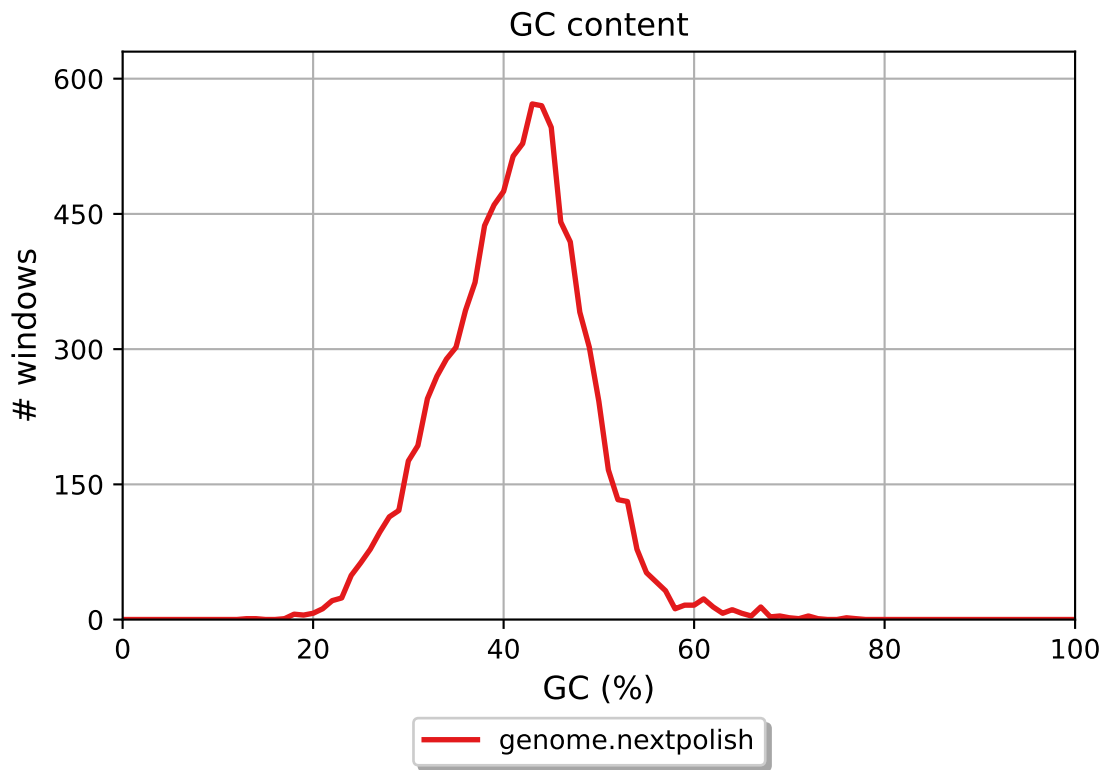
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

