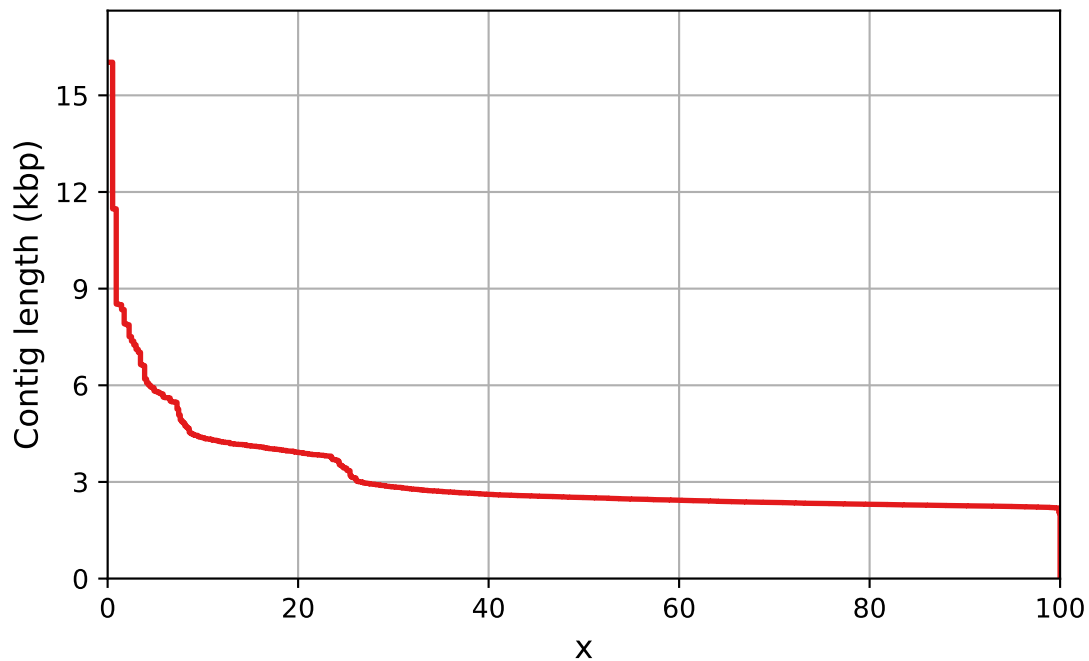


# Report

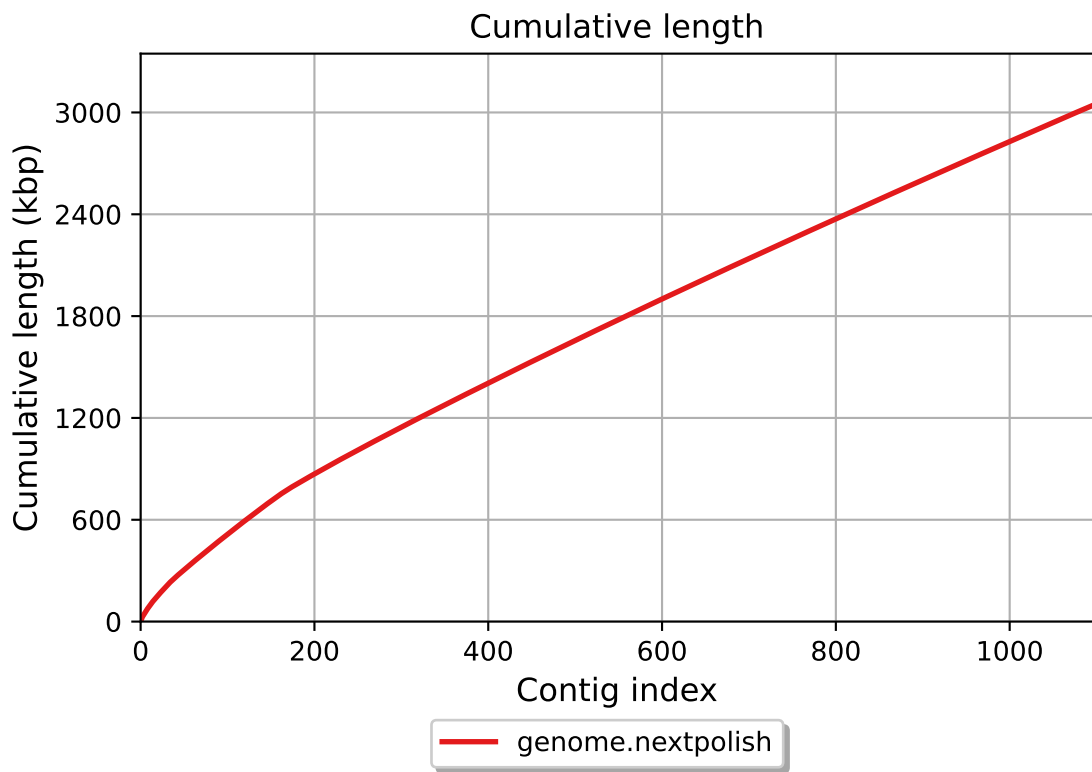
	genome.nextpolish
# contigs (>= 0 bp)	1096
# contigs (>= 1000 bp)	1096
# contigs (>= 5000 bp)	34
# contigs (>= 10000 bp)	2
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	3042344
Total length (>= 1000 bp)	3042344
Total length (>= 5000 bp)	231970
Total length (>= 10000 bp)	27503
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	1096
Largest contig	16026
Total length	3042344
GC (%)	40.98
N50	2516
N90	2259
auN	3126.7
L50	446
L90	960
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	5 + 0 part

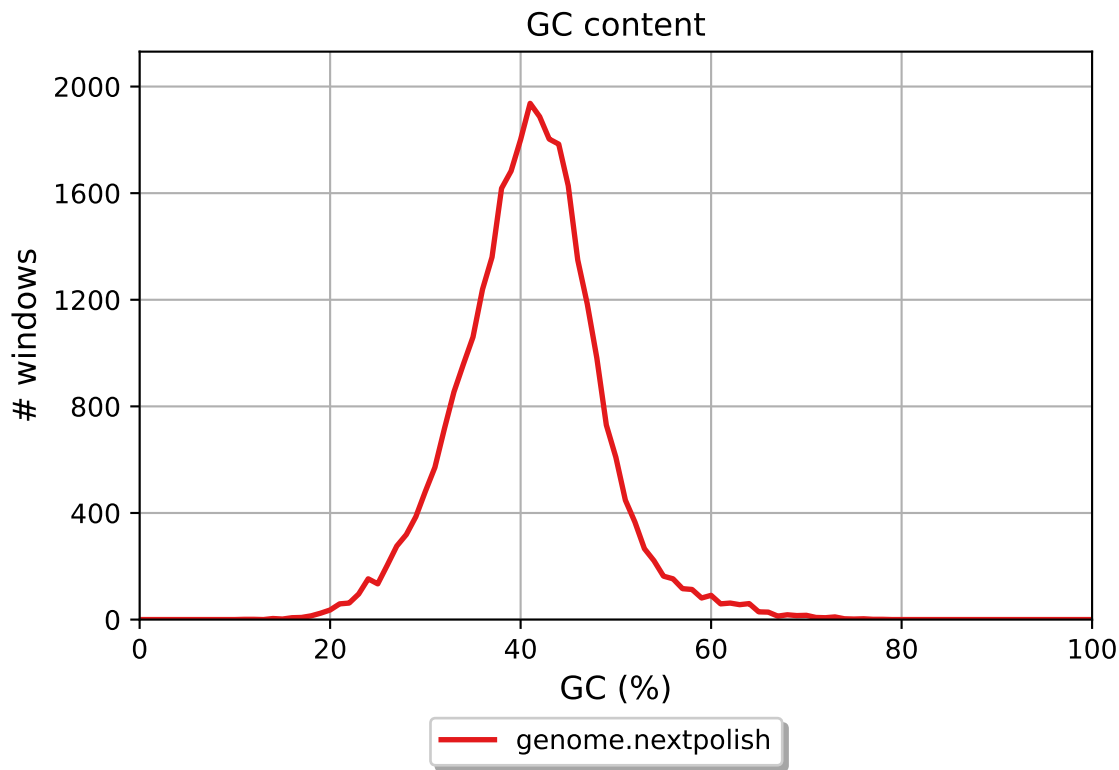
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

