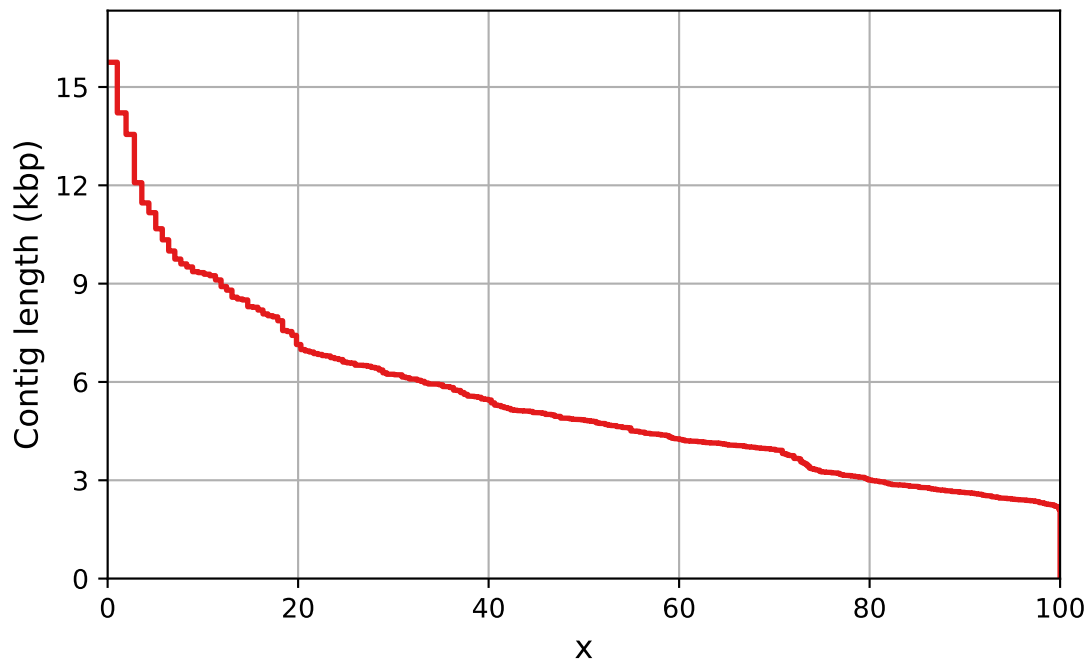


# Report

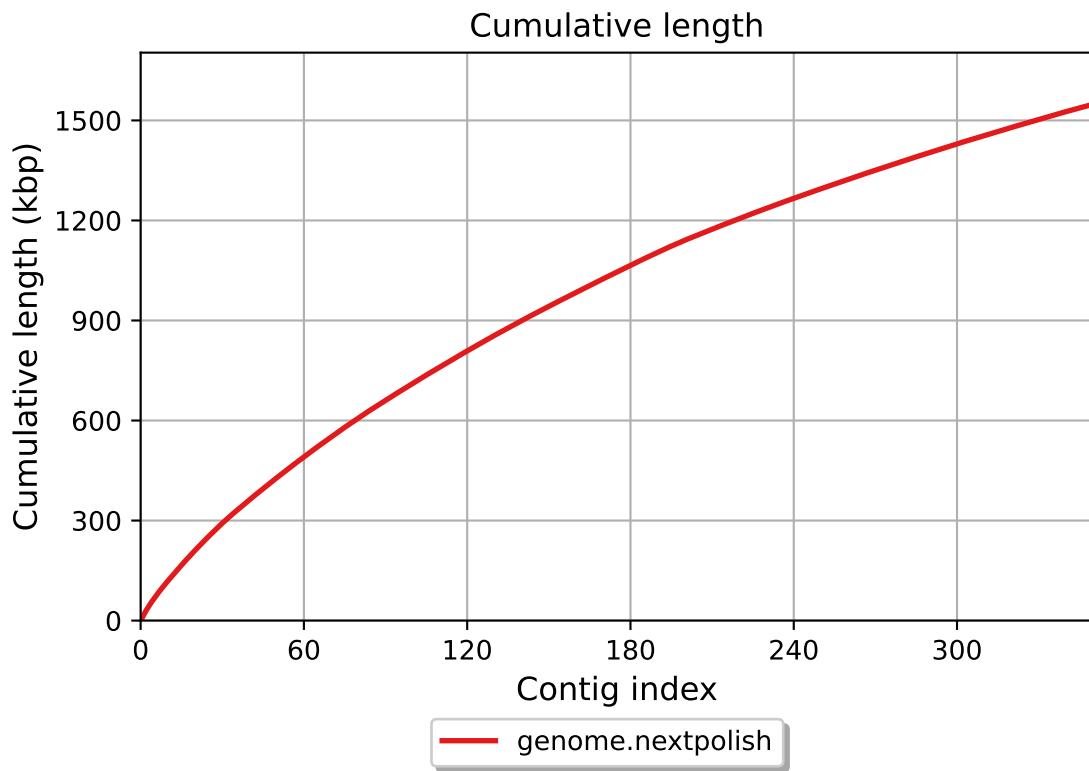
	genome.nextpolish
# contigs (>= 0 bp)	350
# contigs (>= 1000 bp)	350
# contigs (>= 5000 bp)	102
# contigs (>= 10000 bp)	8
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1548440
Total length (>= 1000 bp)	1548440
Total length (>= 5000 bp)	721472
Total length (>= 10000 bp)	99237
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	350
Largest contig	15754
Total length	1548440
GC (%)	40.22
N50	4845
N90	2623
auN	5476.7
L50	113
L90	287
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 0 part

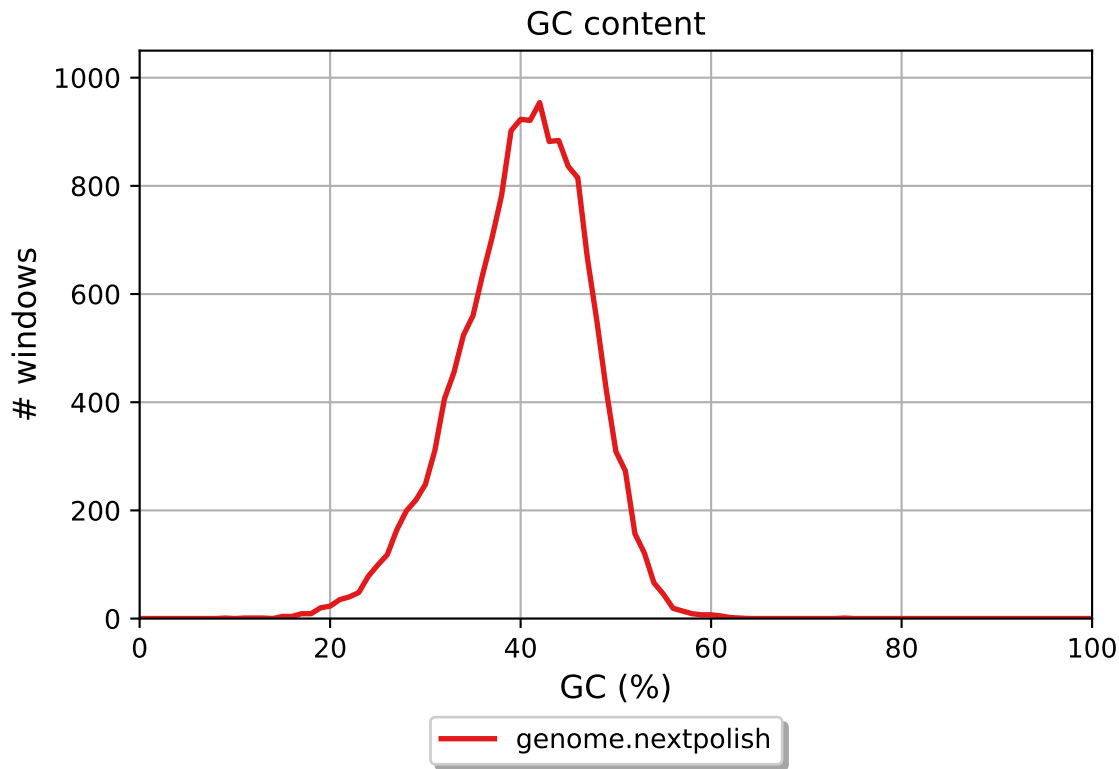
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

