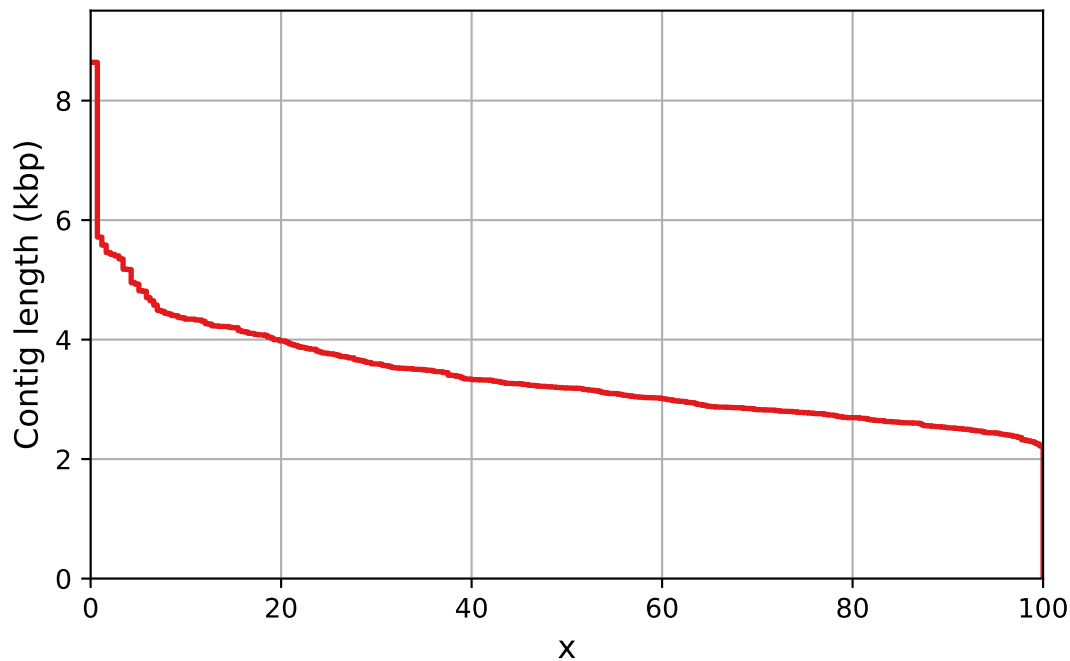


Report

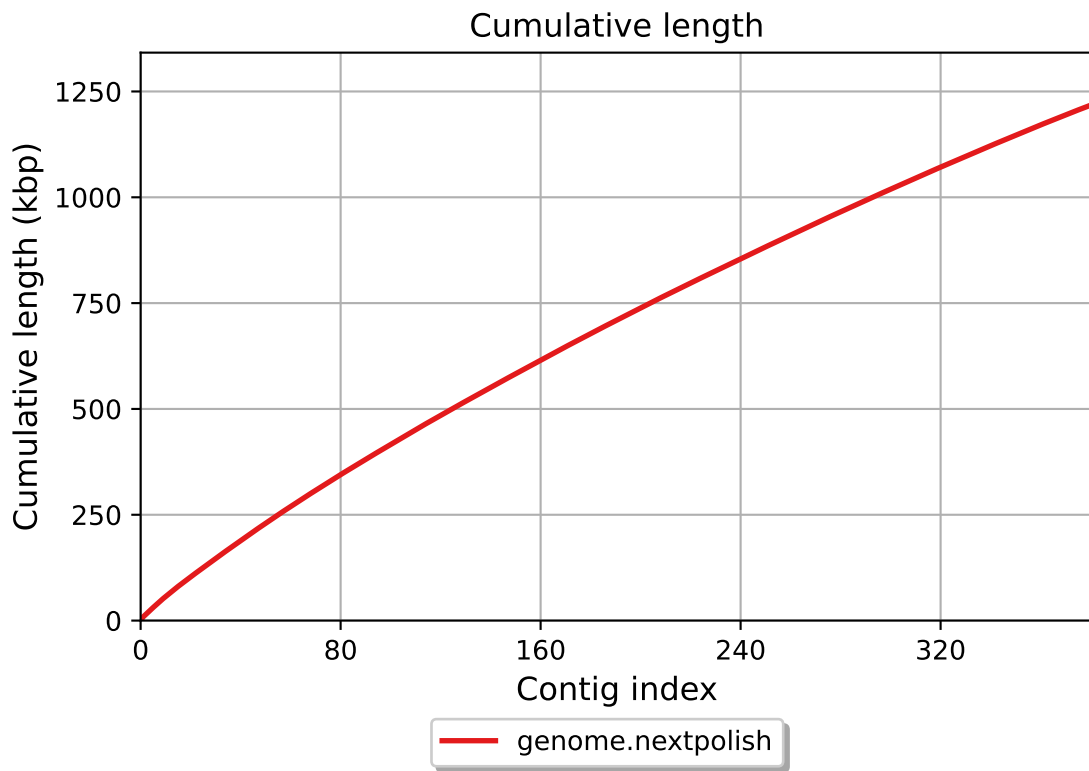
	genome.nextpolish
# contigs (>= 0 bp)	381
# contigs (>= 1000 bp)	381
# contigs (>= 5000 bp)	9
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1219761
Total length (>= 1000 bp)	1219761
Total length (>= 5000 bp)	51924
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	381
Largest contig	8641
Total length	1219761
GC (%)	41.49
N50	3190
N90	2524
auN	3366.6
L50	159
L90	331
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	1 + 1 part

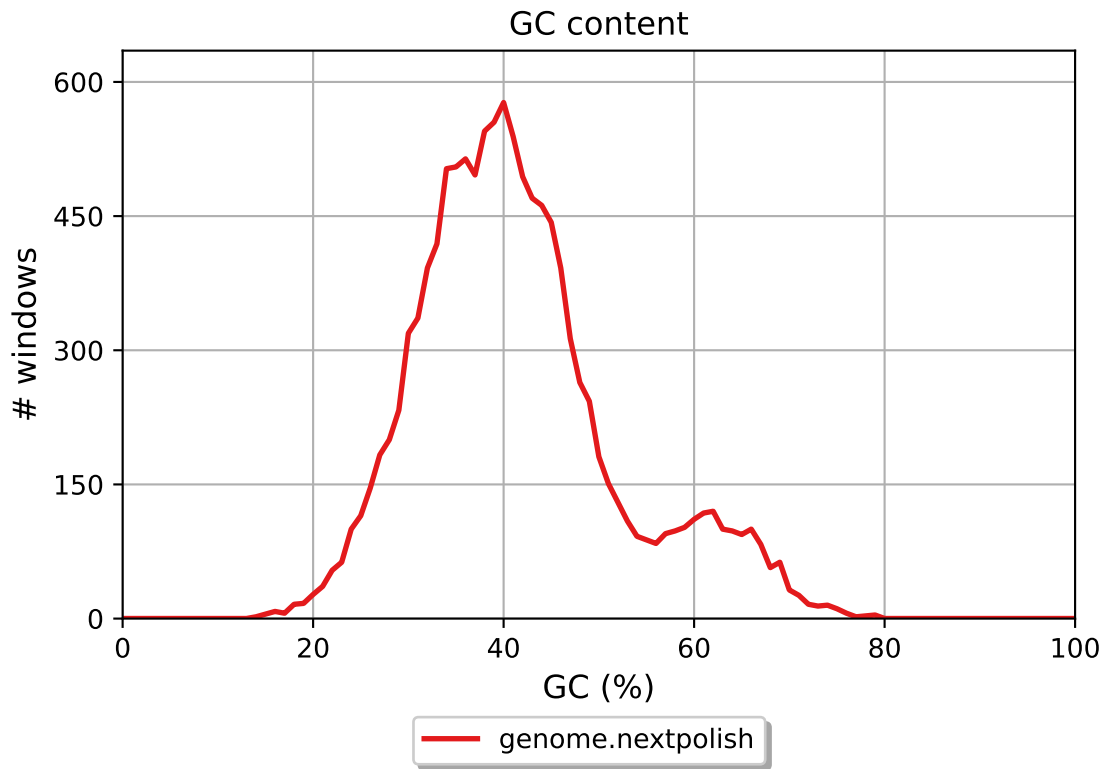
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

