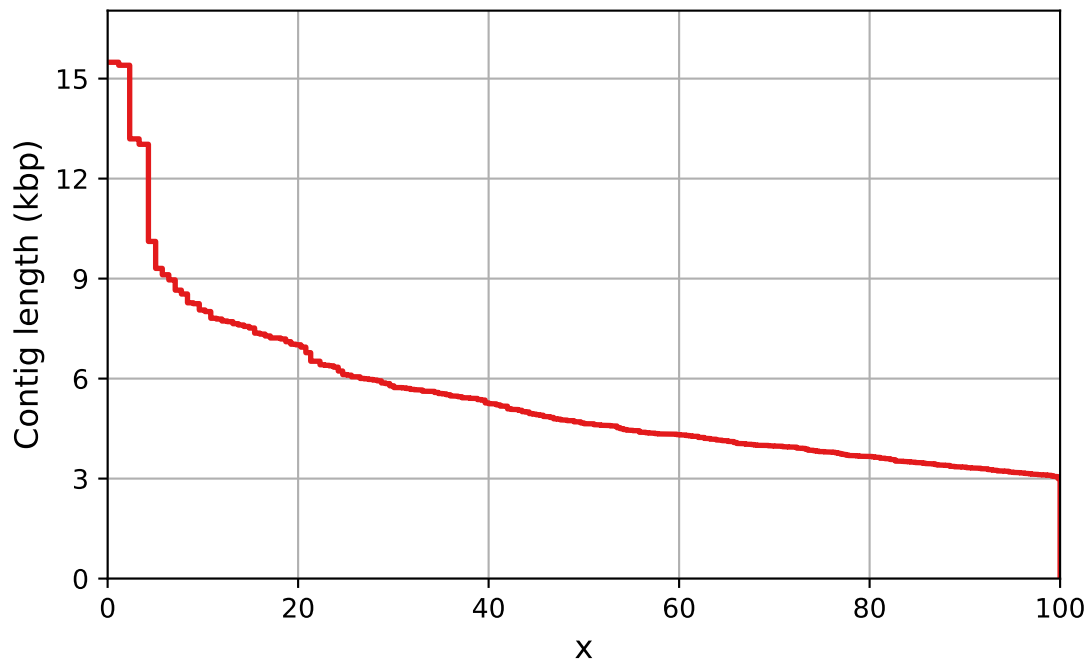


Report

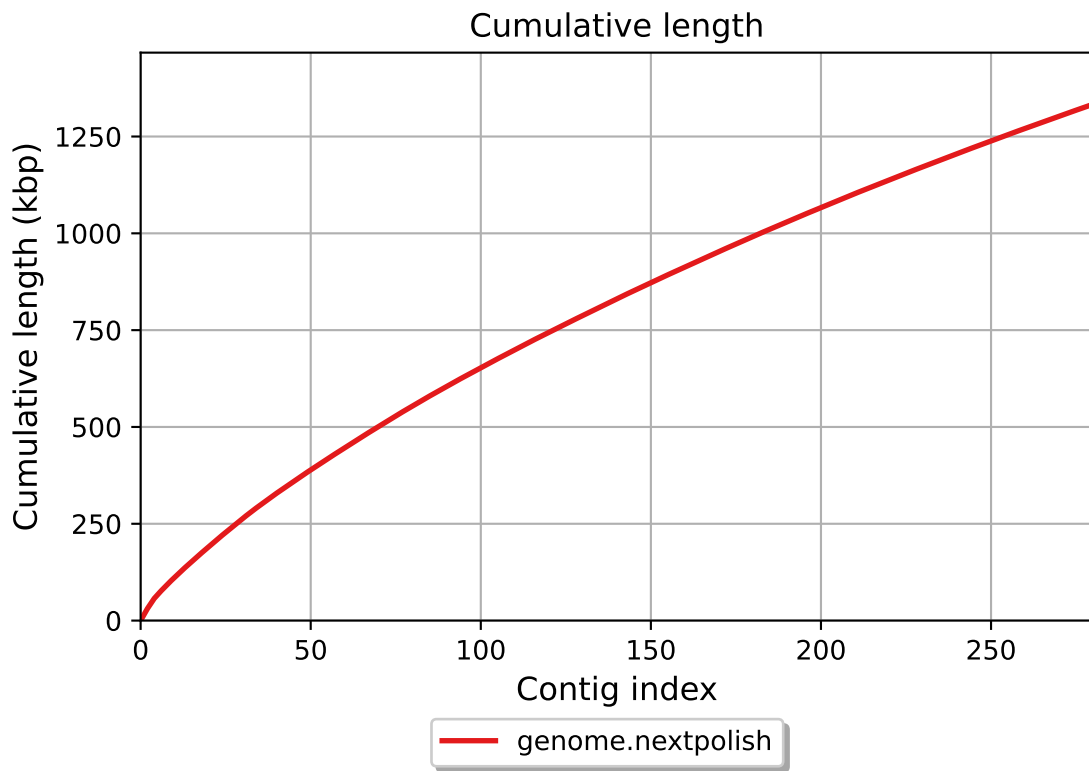
	genome.nextpolish
# contigs (>= 0 bp)	280
# contigs (>= 1000 bp)	280
# contigs (>= 5000 bp)	86
# contigs (>= 10000 bp)	5
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1333262
Total length (>= 1000 bp)	1333262
Total length (>= 5000 bp)	585079
Total length (>= 10000 bp)	67226
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	280
Largest contig	15490
Total length	1333262
GC (%)	49.73
N50	4669
N90	3344
auN	5455.9
L50	103
L90	239
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	7 + 2 part

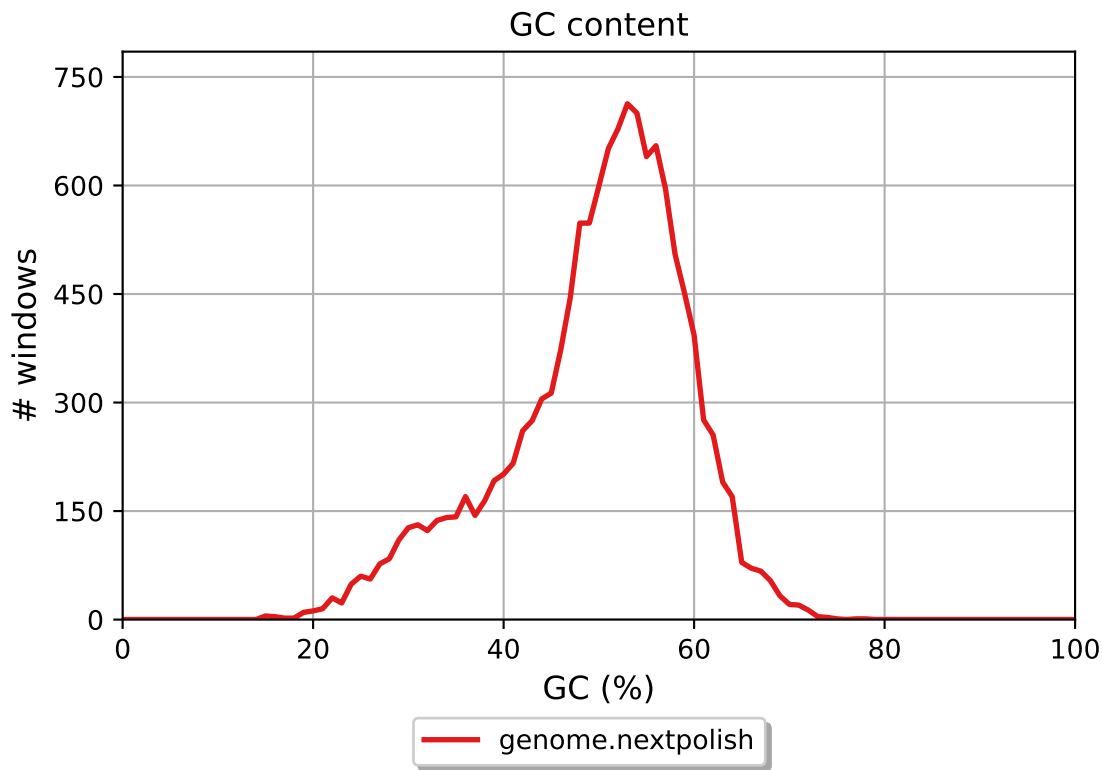
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

