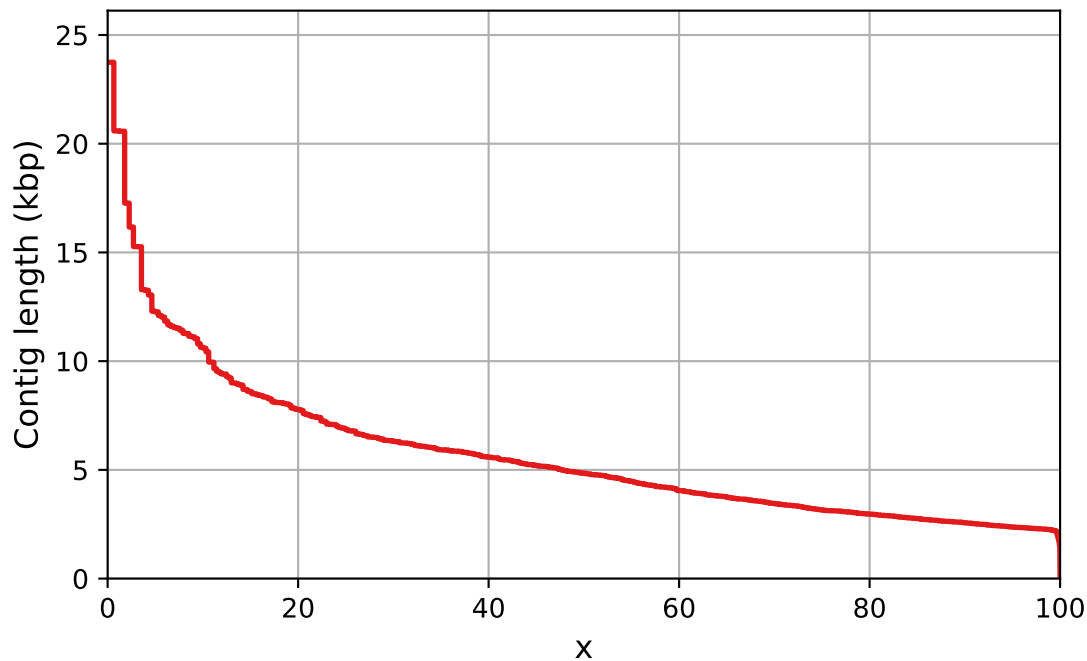


# Report

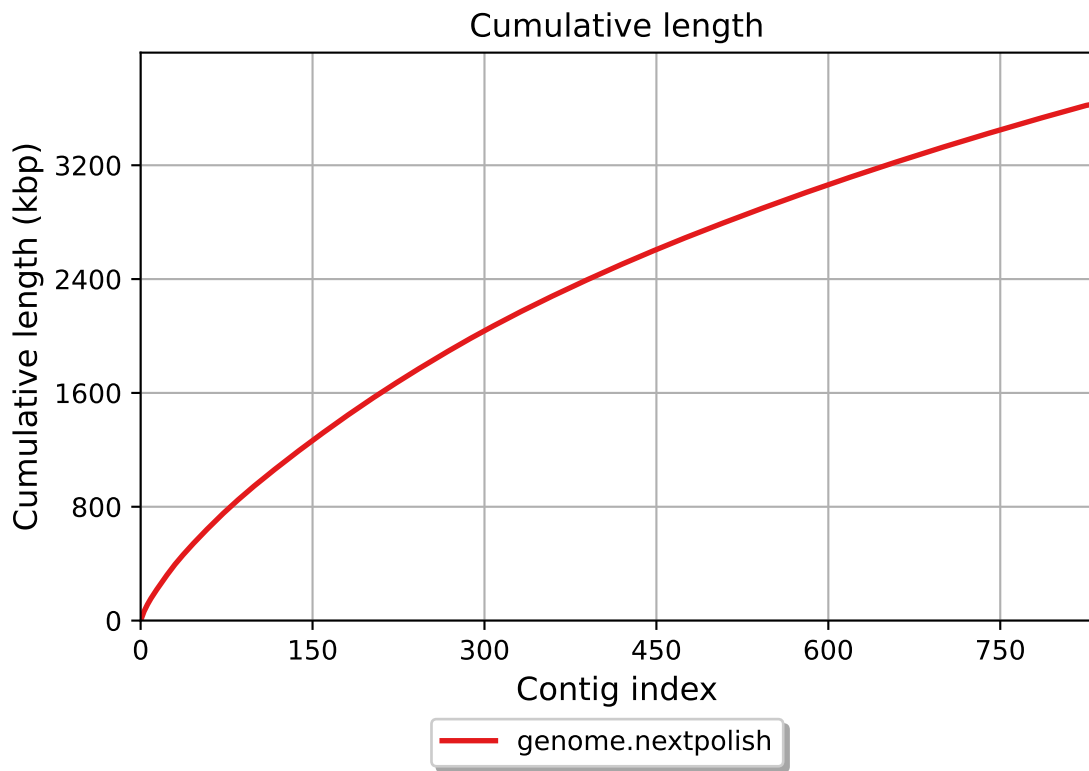
	genome.nextpolish
# contigs (>= 0 bp)	831
# contigs (>= 1000 bp)	829
# contigs (>= 5000 bp)	235
# contigs (>= 10000 bp)	29
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	3628559
Total length (>= 1000 bp)	3627466
Total length (>= 5000 bp)	1731693
Total length (>= 10000 bp)	385018
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	831
Largest contig	23745
Total length	3628559
GC (%)	44.71
N50	4849
N90	2564
auN	5774.6
L50	252
L90	676
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	9 + 0 part

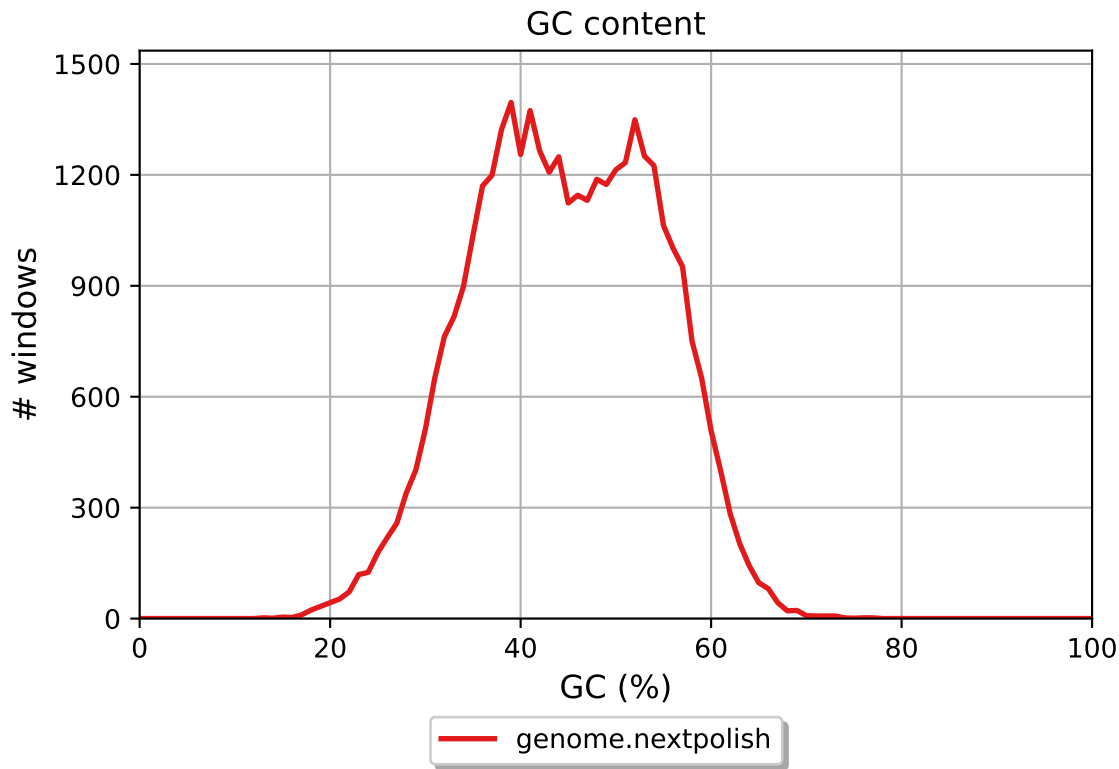
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

