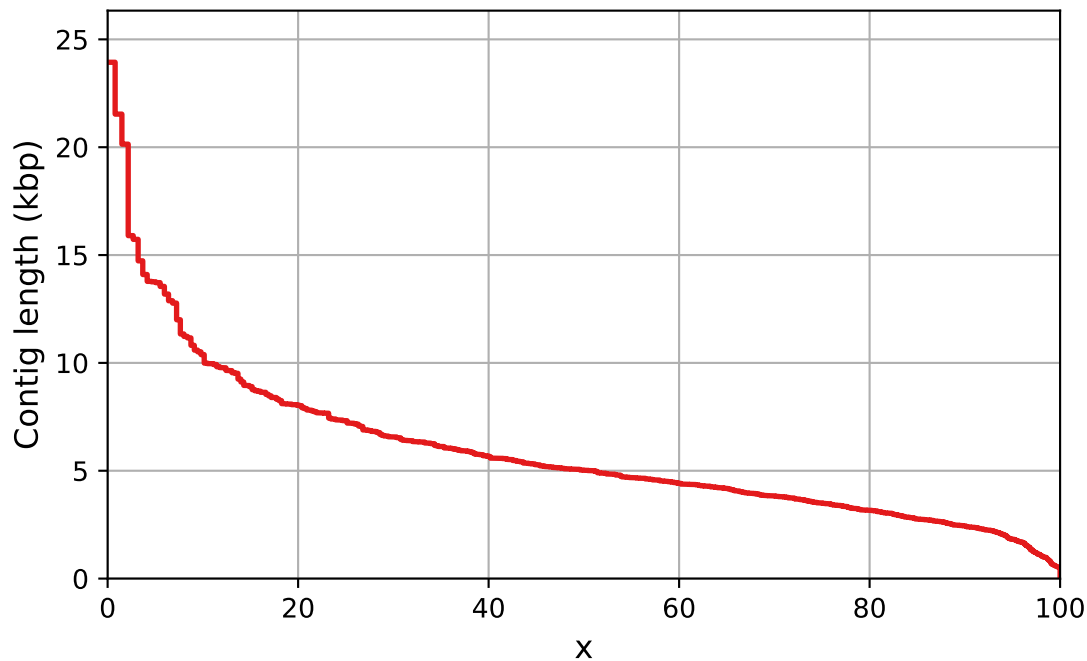


Report

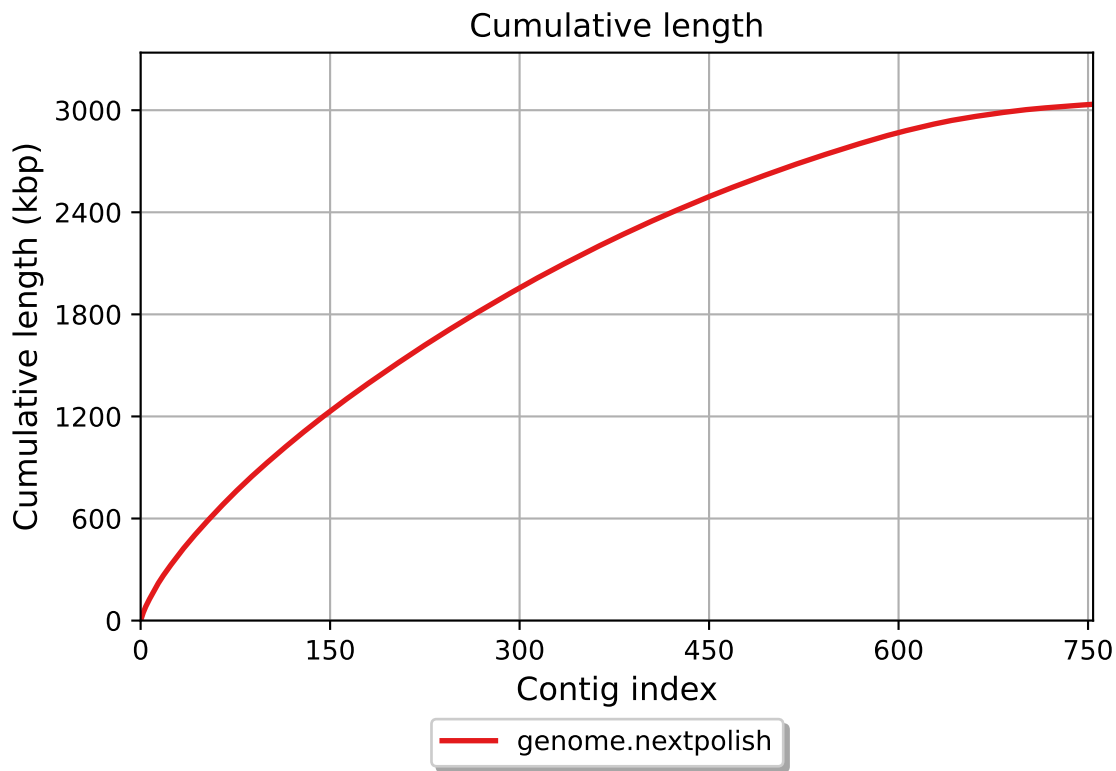
	genome.nextpolish
# contigs (>= 0 bp)	757
# contigs (>= 1000 bp)	679
# contigs (>= 5000 bp)	211
# contigs (>= 10000 bp)	22
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	3035784
Total length (>= 1000 bp)	2982983
Total length (>= 5000 bp)	1550869
Total length (>= 10000 bp)	307761
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	754
Largest contig	23938
Total length	3034908
GC (%)	44.31
N50	5023
N90	2446
auN	5950.7
L50	205
L90	539
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	5 + 2 part

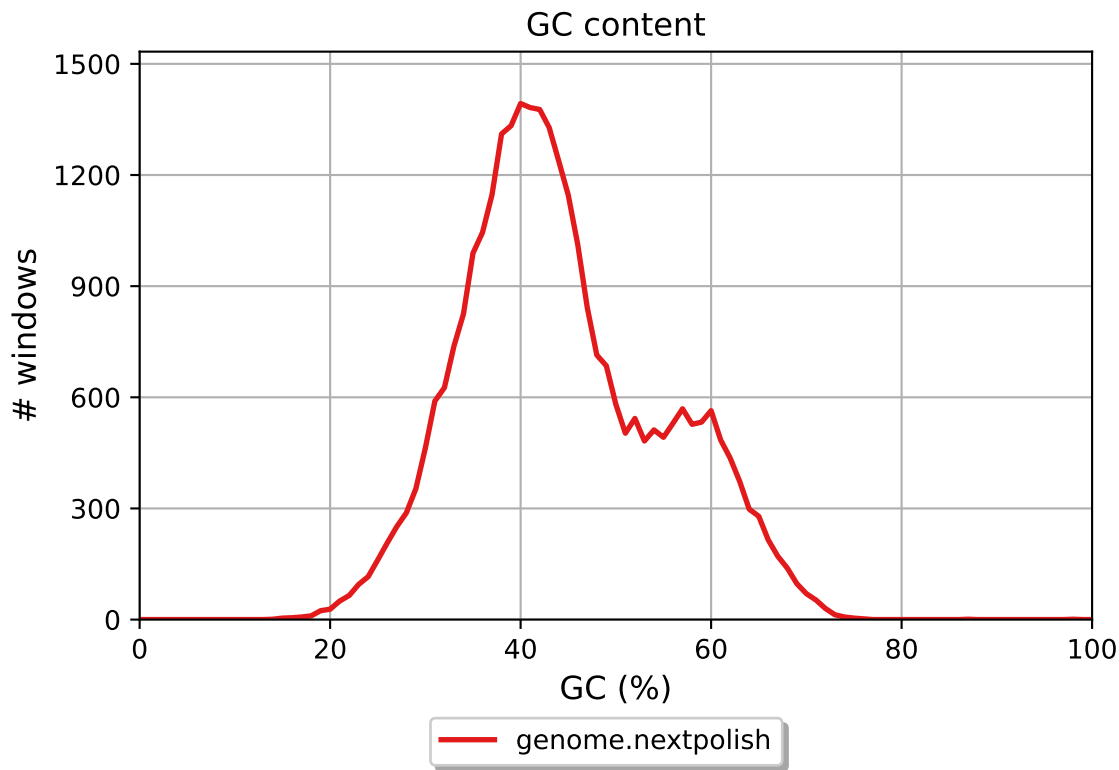
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

