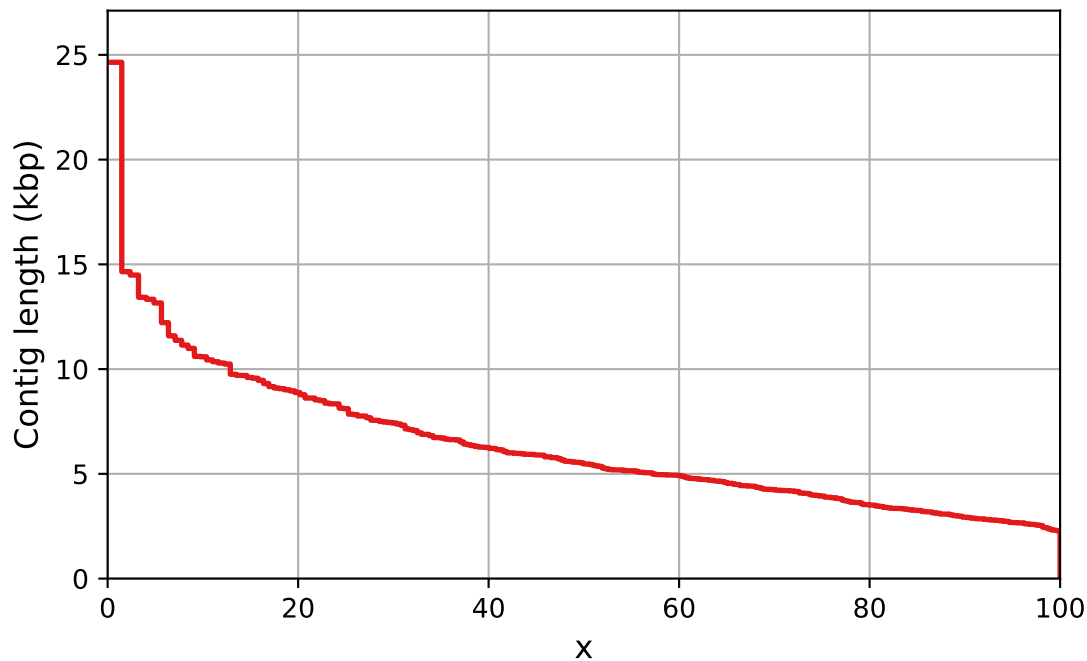


Report

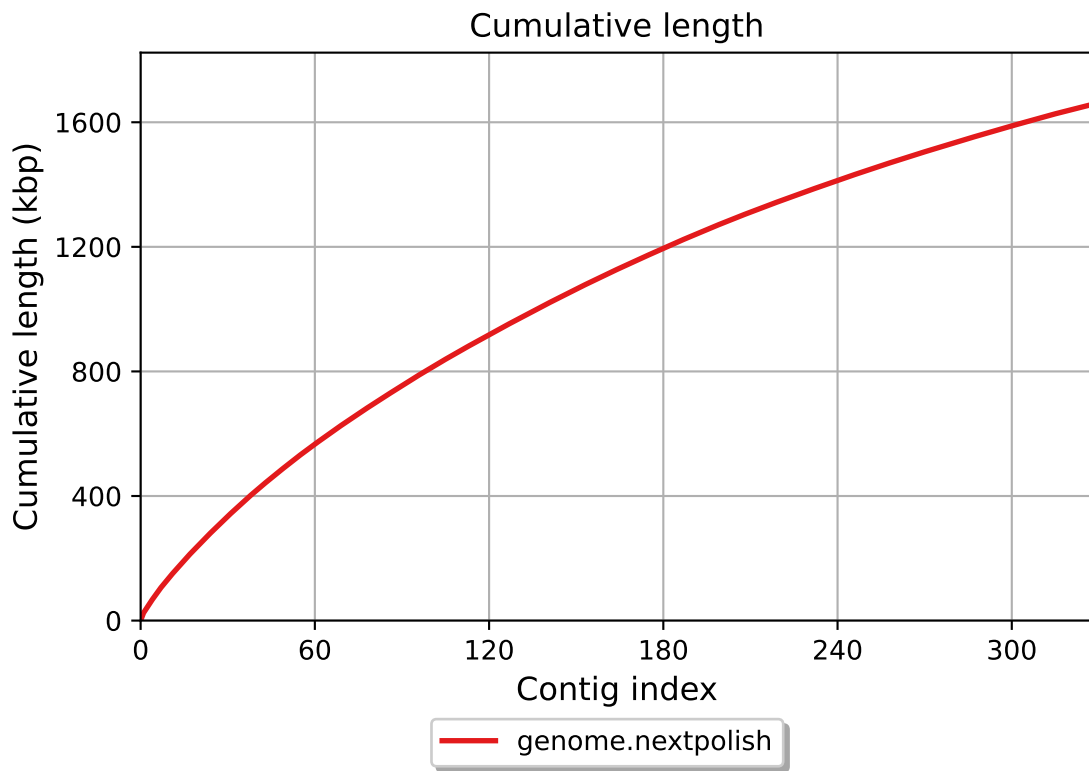
	genome.nextpolish
# contigs (>= 0 bp)	328
# contigs (>= 1000 bp)	328
# contigs (>= 5000 bp)	126
# contigs (>= 10000 bp)	17
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1657769
Total length (>= 1000 bp)	1657769
Total length (>= 5000 bp)	948210
Total length (>= 10000 bp)	213527
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	328
Largest contig	24646
Total length	1657769
GC (%)	42.97
N50	5477
N90	2935
auN	6421.4
L50	104
L90	266
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 5 part

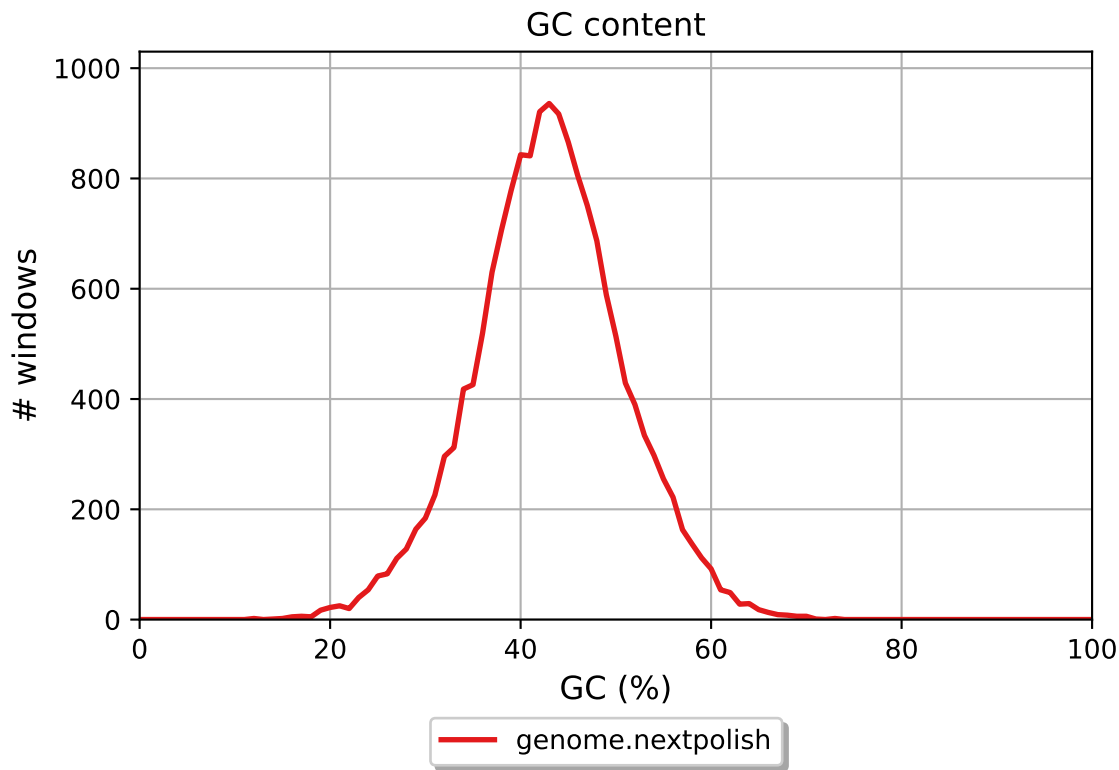
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

