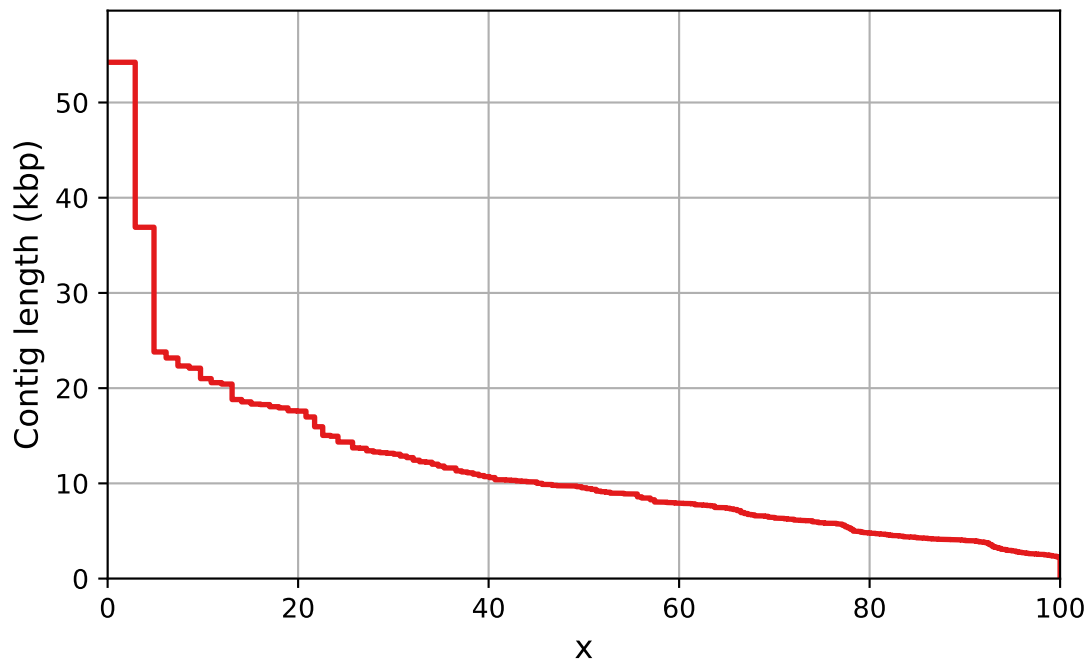


Report

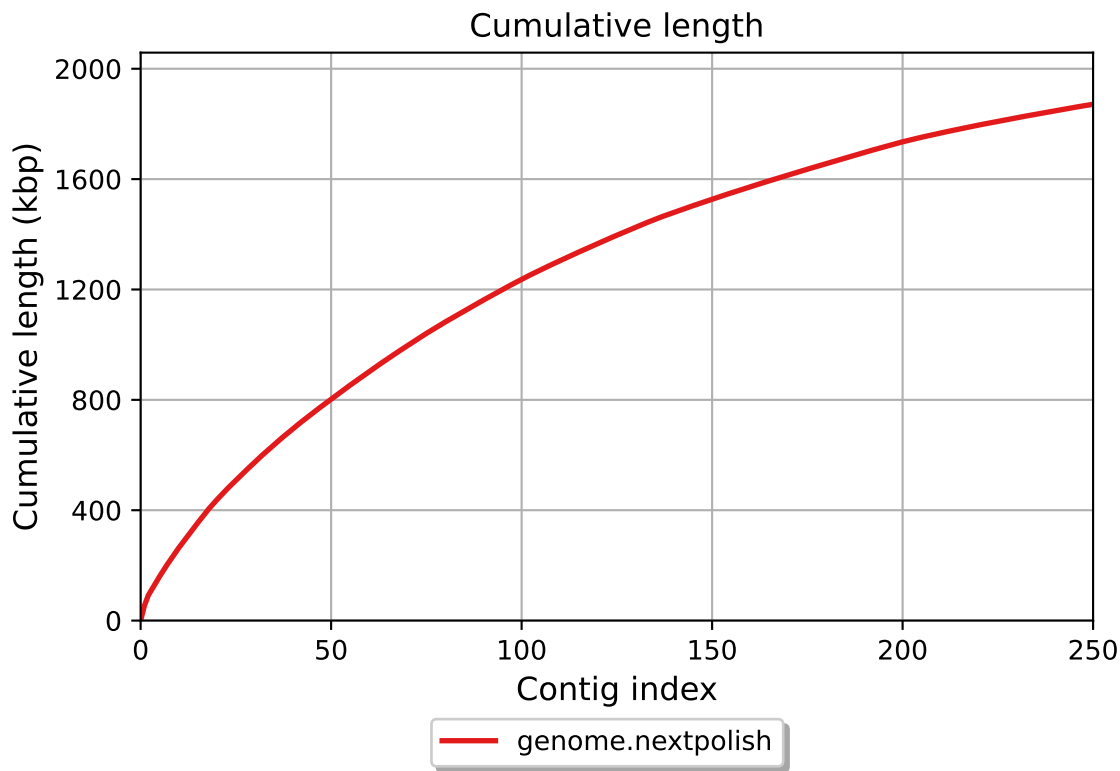
	genome.nextpolish
# contigs (>= 0 bp)	250
# contigs (>= 1000 bp)	250
# contigs (>= 5000 bp)	137
# contigs (>= 10000 bp)	55
# contigs (>= 25000 bp)	2
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	1871275
Total length (>= 1000 bp)	1871275
Total length (>= 5000 bp)	1464685
Total length (>= 10000 bp)	853312
Total length (>= 25000 bp)	91121
Total length (>= 50000 bp)	54225
# contigs	250
Largest contig	54225
Total length	1871275
GC (%)	38.26
N50	9535
N90	4029
auN	11937.8
L50	64
L90	187
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	8 + 0 part

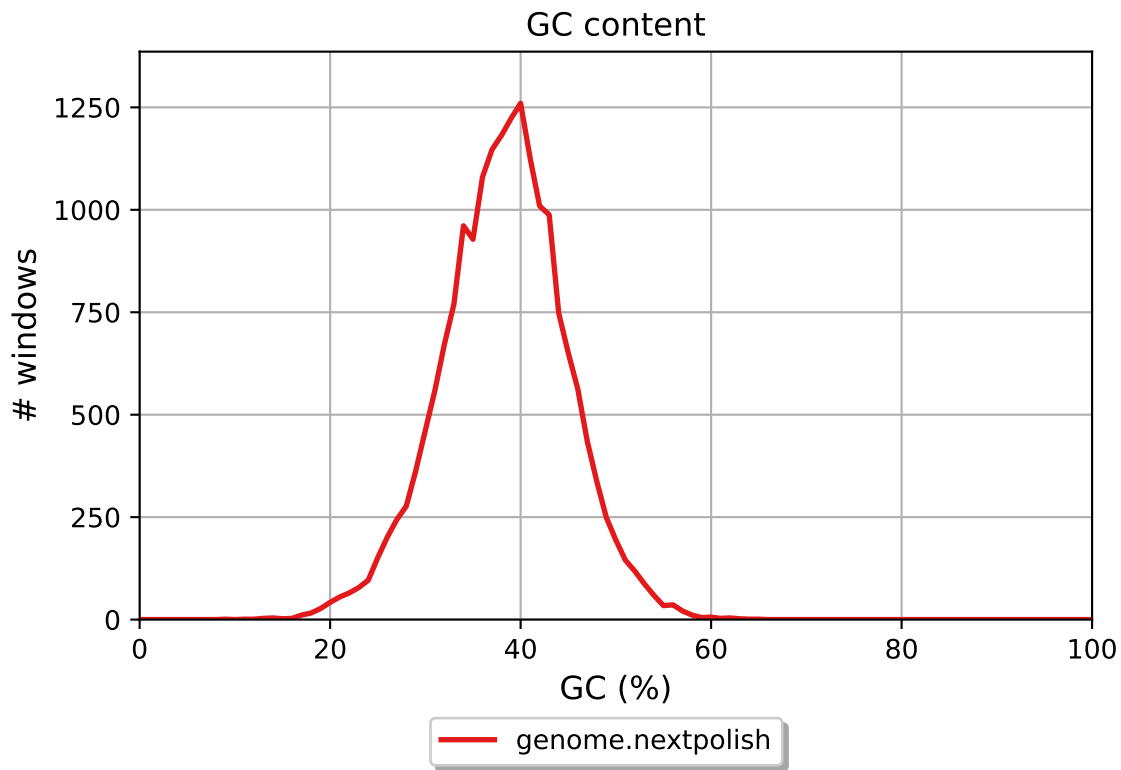
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

