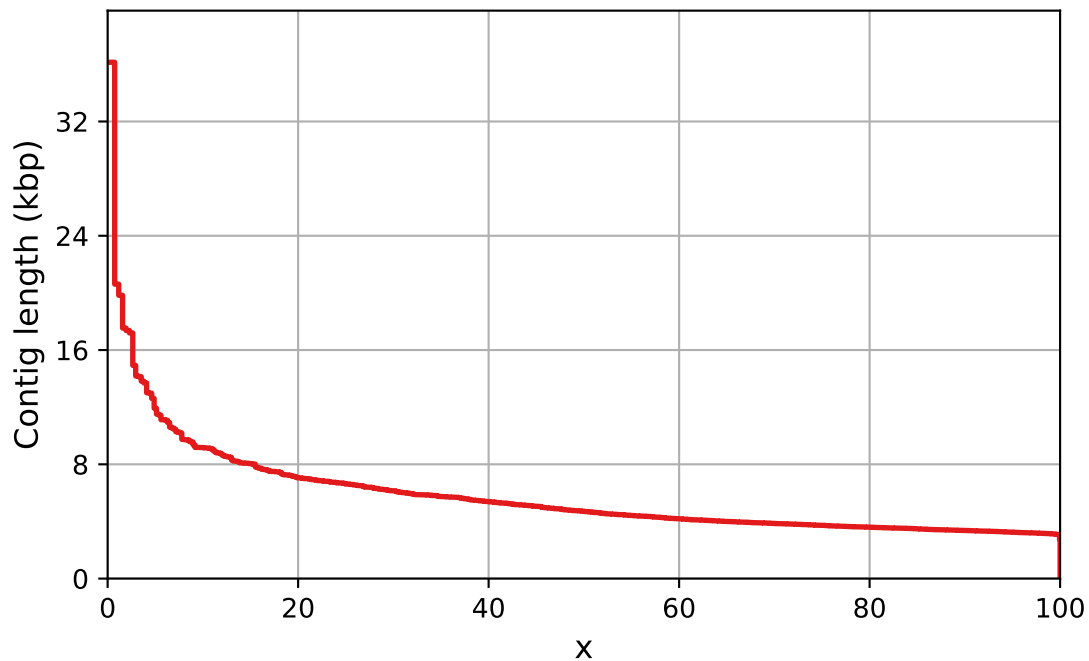


# Report

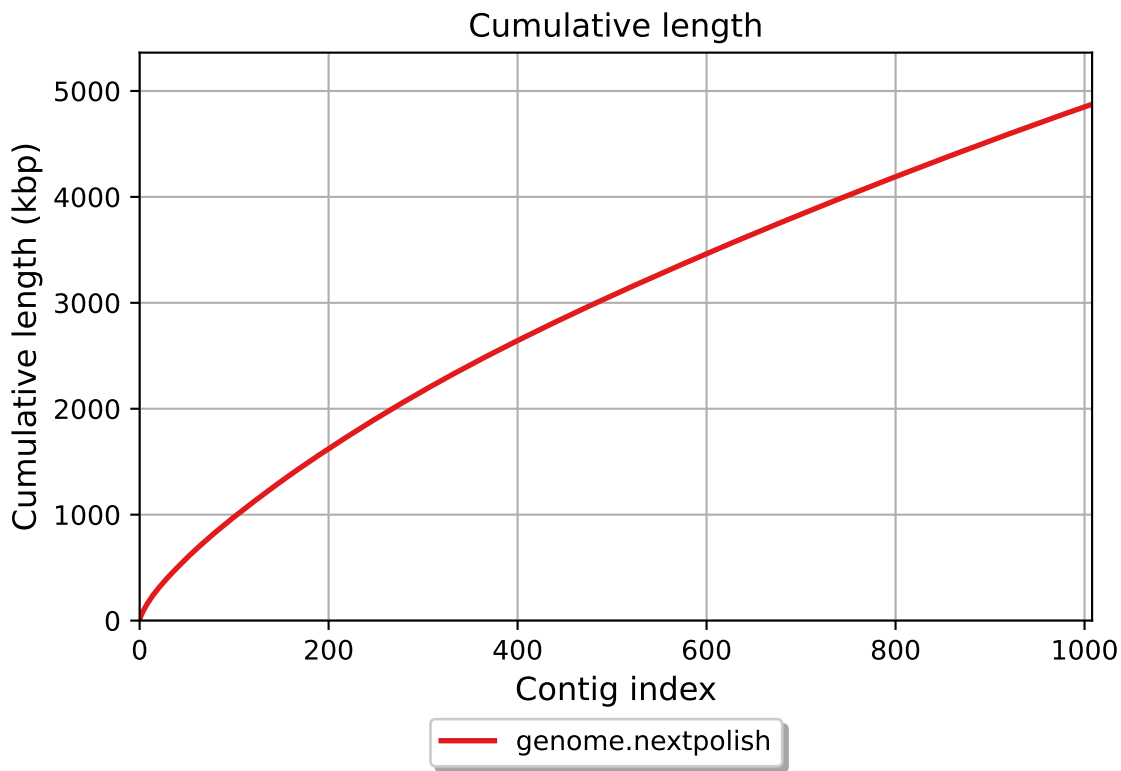
	genome.nextpolish
# contigs (>= 0 bp)	1008
# contigs (>= 1000 bp)	1008
# contigs (>= 5000 bp)	310
# contigs (>= 10000 bp)	27
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	4874548
Total length (>= 1000 bp)	4874548
Total length (>= 5000 bp)	2219967
Total length (>= 10000 bp)	379514
Total length (>= 25000 bp)	36146
Total length (>= 50000 bp)	0
# contigs	1008
Largest contig	36146
Total length	4874548
GC (%)	36.87
N50	4710
N90	3371
auN	5889.0
L50	355
L90	858
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	10 + 4 part

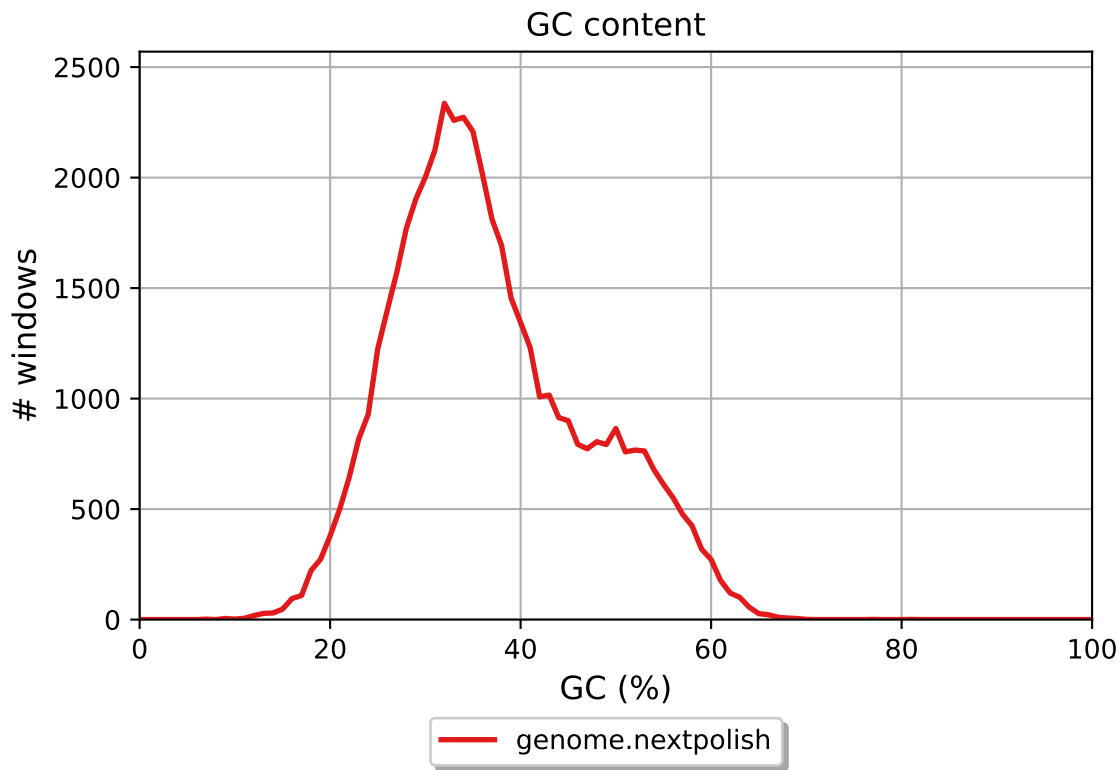
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

