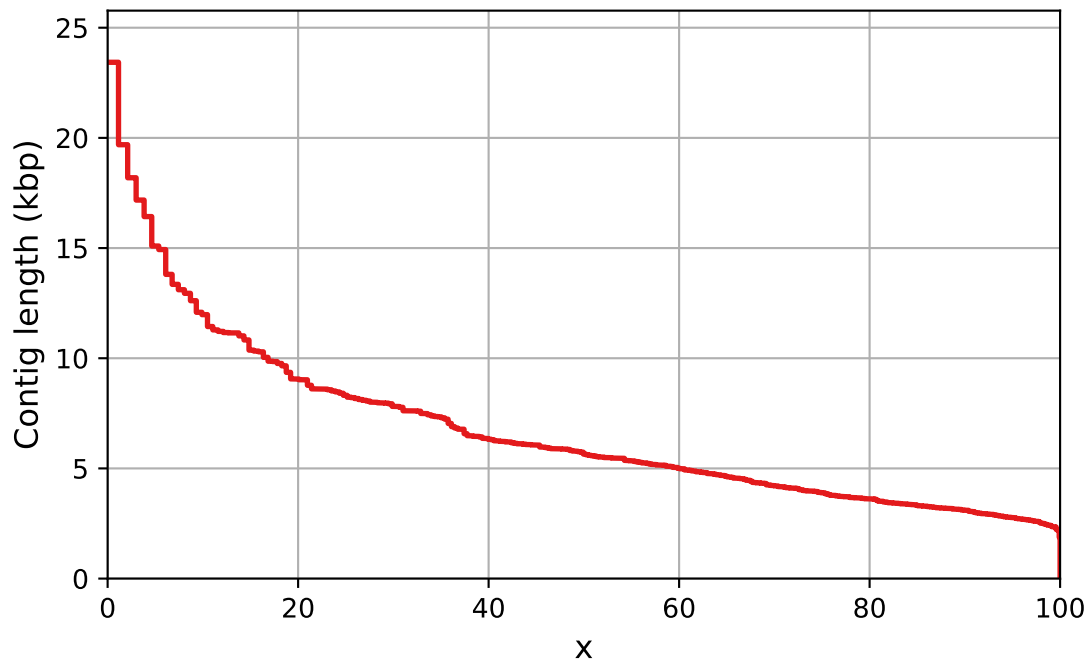


# Report

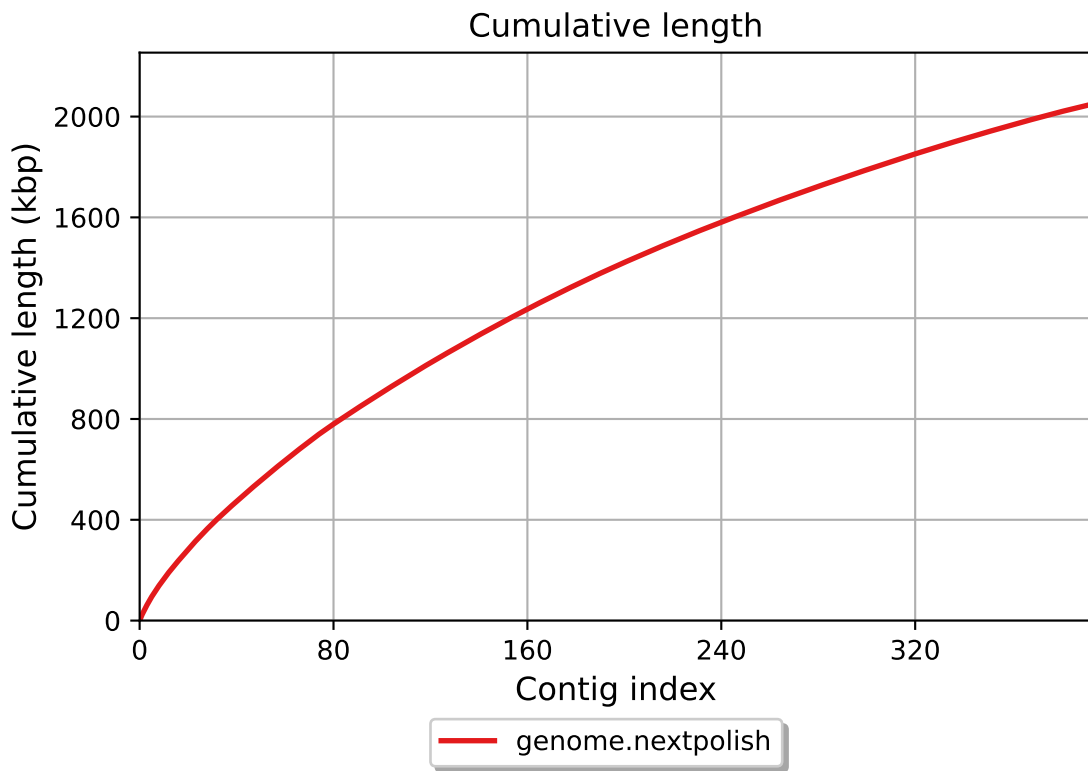
	genome.nextpolish
# contigs (>= 0 bp)	393
# contigs (>= 1000 bp)	393
# contigs (>= 5000 bp)	159
# contigs (>= 10000 bp)	26
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2048844
Total length (>= 1000 bp)	2048844
Total length (>= 5000 bp)	1231079
Total length (>= 10000 bp)	345111
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	393
Largest contig	23431
Total length	2048844
GC (%)	33.77
N50	5652
N90	3101
auN	6815.6
L50	121
L90	318
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 2 part

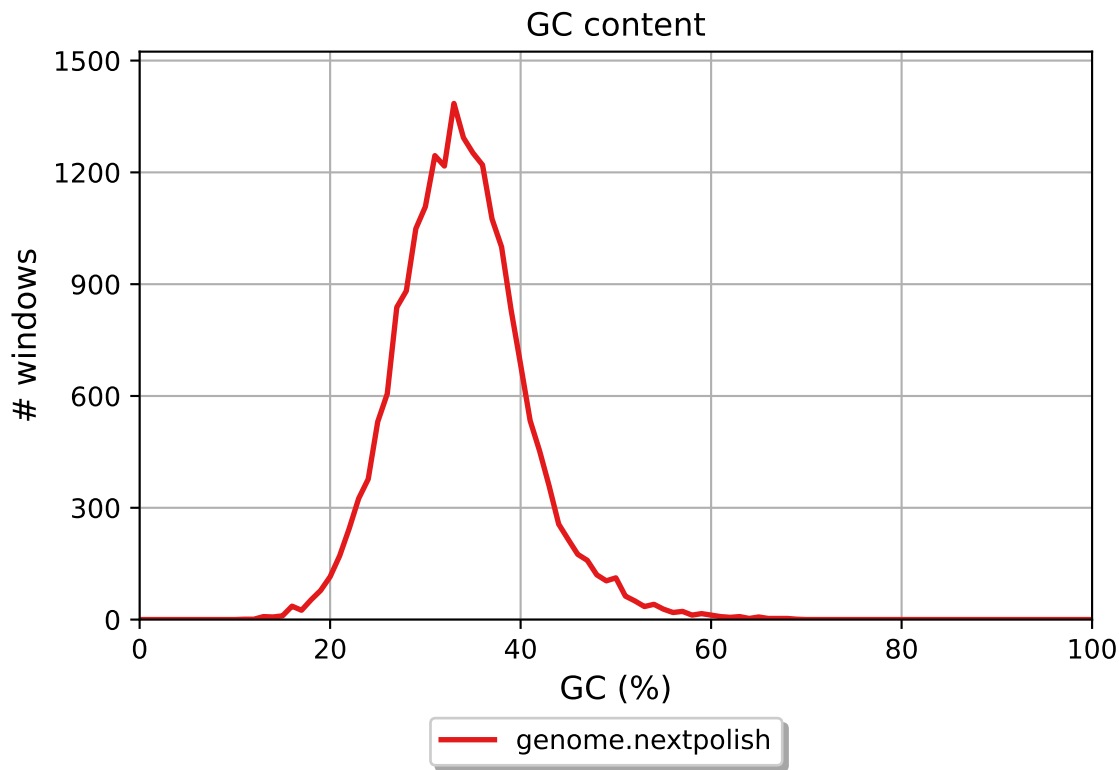
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

