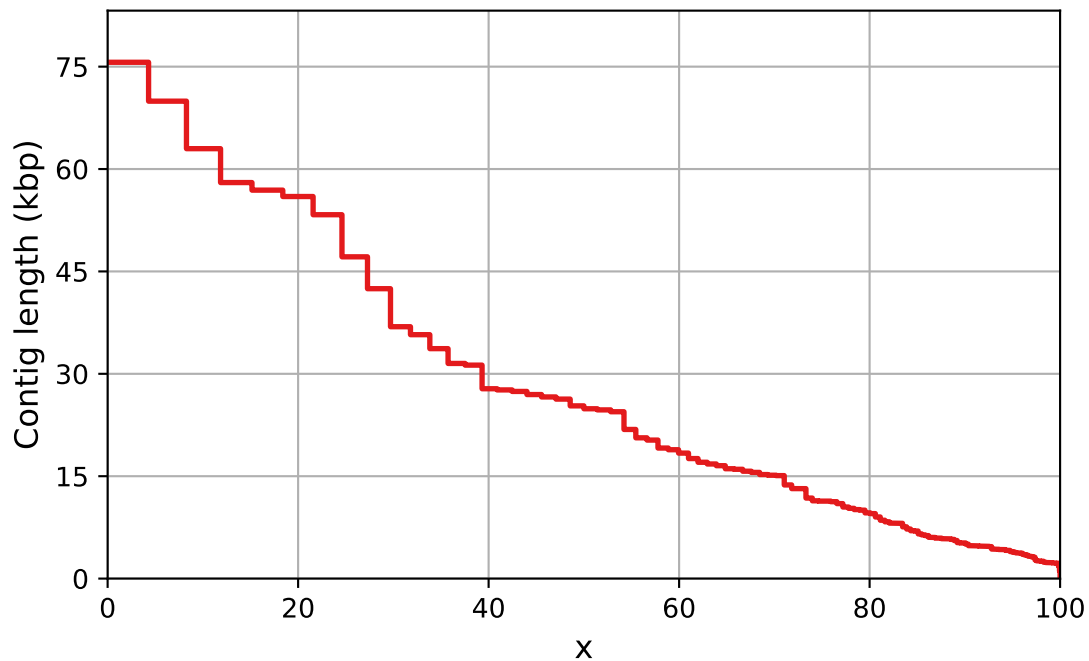


# Report

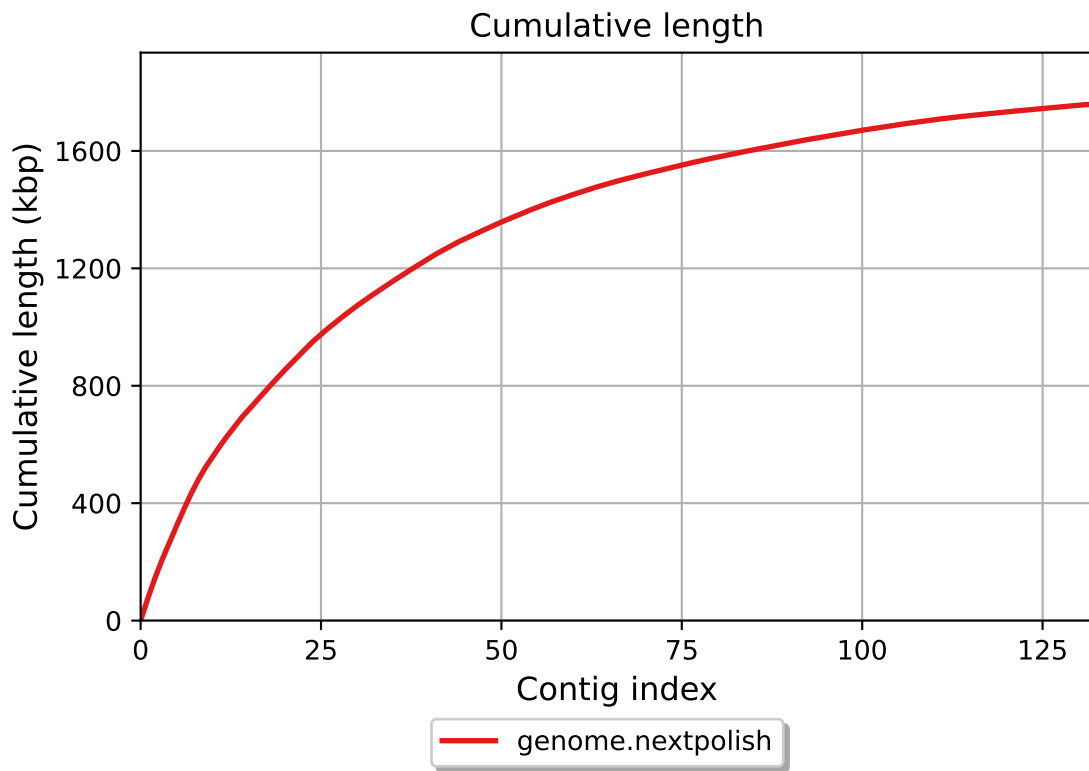
	genome.nextpolish
# contigs (>= 0 bp)	132
# contigs (>= 1000 bp)	132
# contigs (>= 5000 bp)	82
# contigs (>= 10000 bp)	54
# contigs (>= 25000 bp)	21
# contigs (>= 50000 bp)	7
Total length (>= 0 bp)	1758947
Total length (>= 1000 bp)	1758947
Total length (>= 5000 bp)	1589606
Total length (>= 10000 bp)	1398665
Total length (>= 25000 bp)	879536
Total length (>= 50000 bp)	432774
# contigs	132
Largest contig	75641
Total length	1758947
GC (%)	32.38
N50	25304
N90	5219
auN	29984.6
L50	21
L90	81
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	5 + 0 part

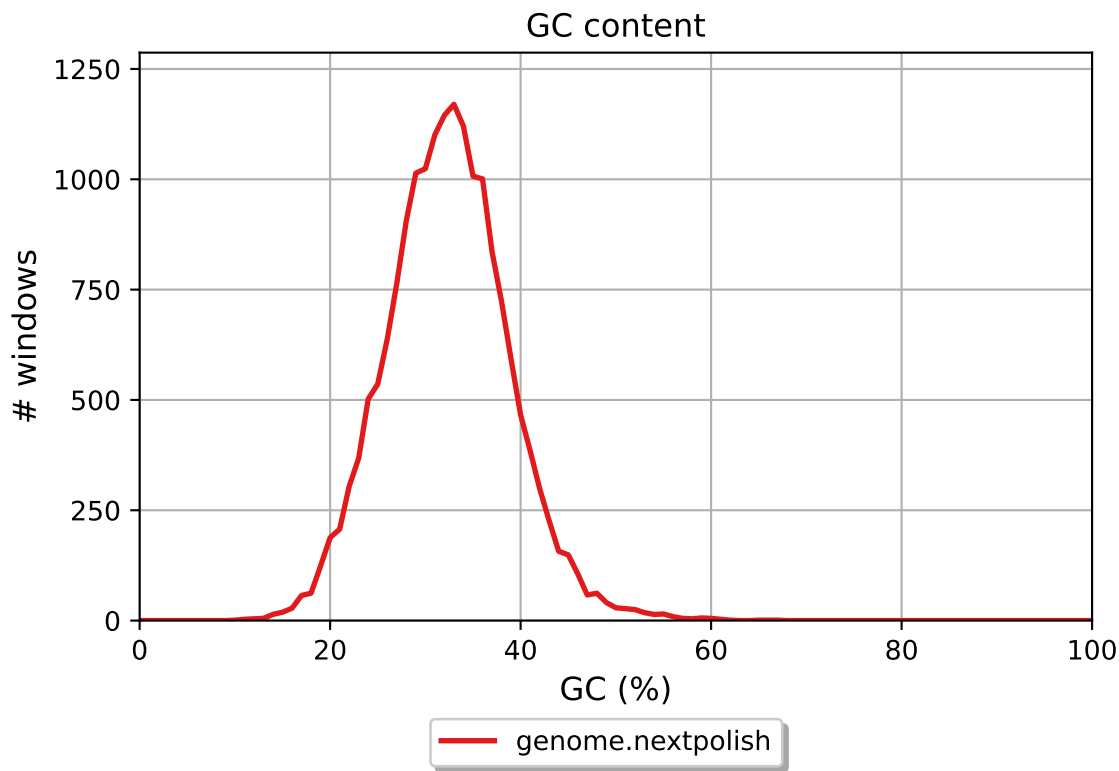
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



— genome.nextpolish





genome.nextpolish GC content

