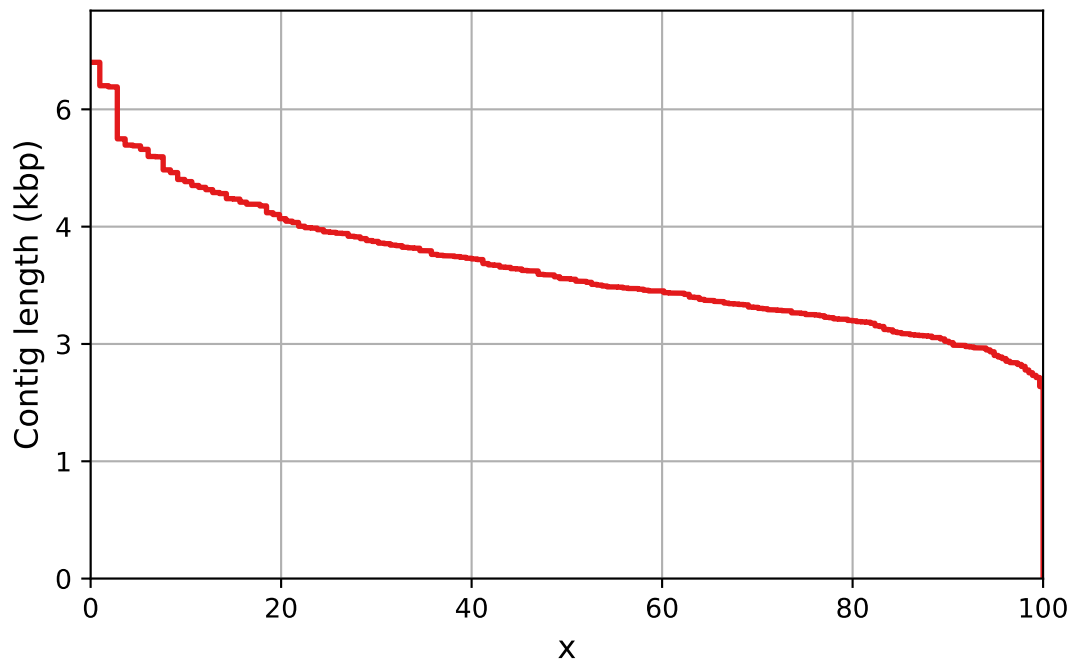


# Report

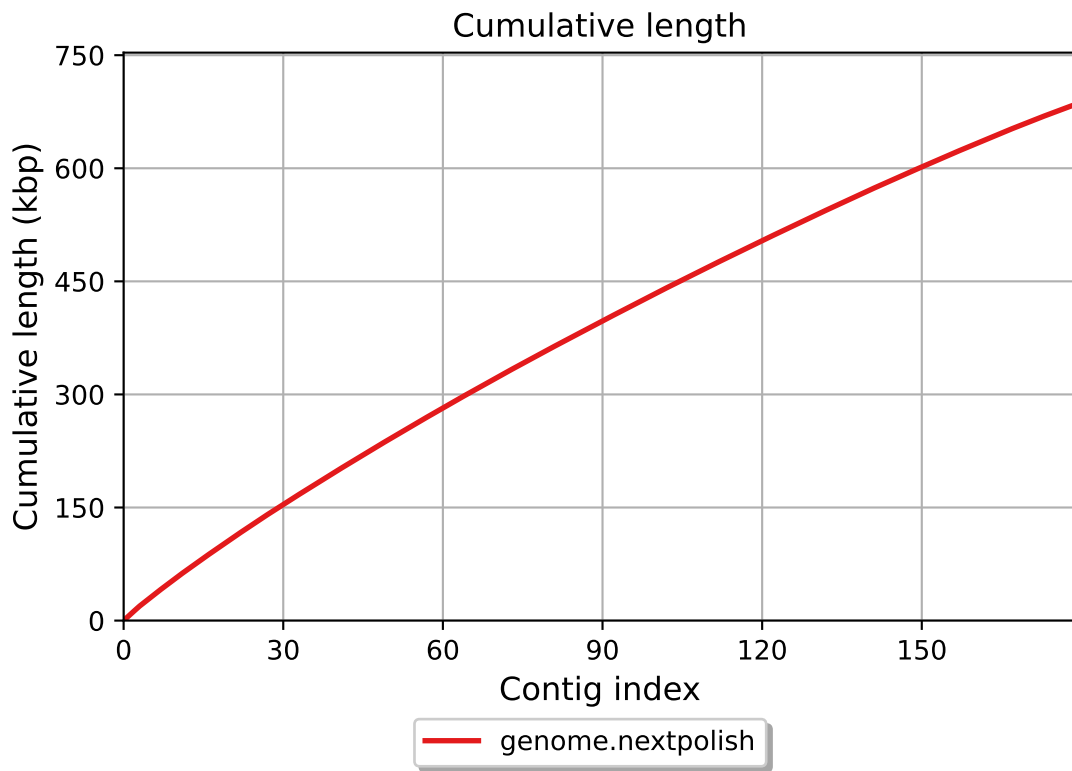
	genome.nextpolish
# contigs (>= 0 bp)	179
# contigs (>= 1000 bp)	179
# contigs (>= 5000 bp)	15
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	684896
Total length (>= 1000 bp)	684896
Total length (>= 5000 bp)	82791
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	179
Largest contig	6601
Total length	684896
GC (%)	41.29
N50	3834
N90	3034
auN	3981.4
L50	76
L90	155
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 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).

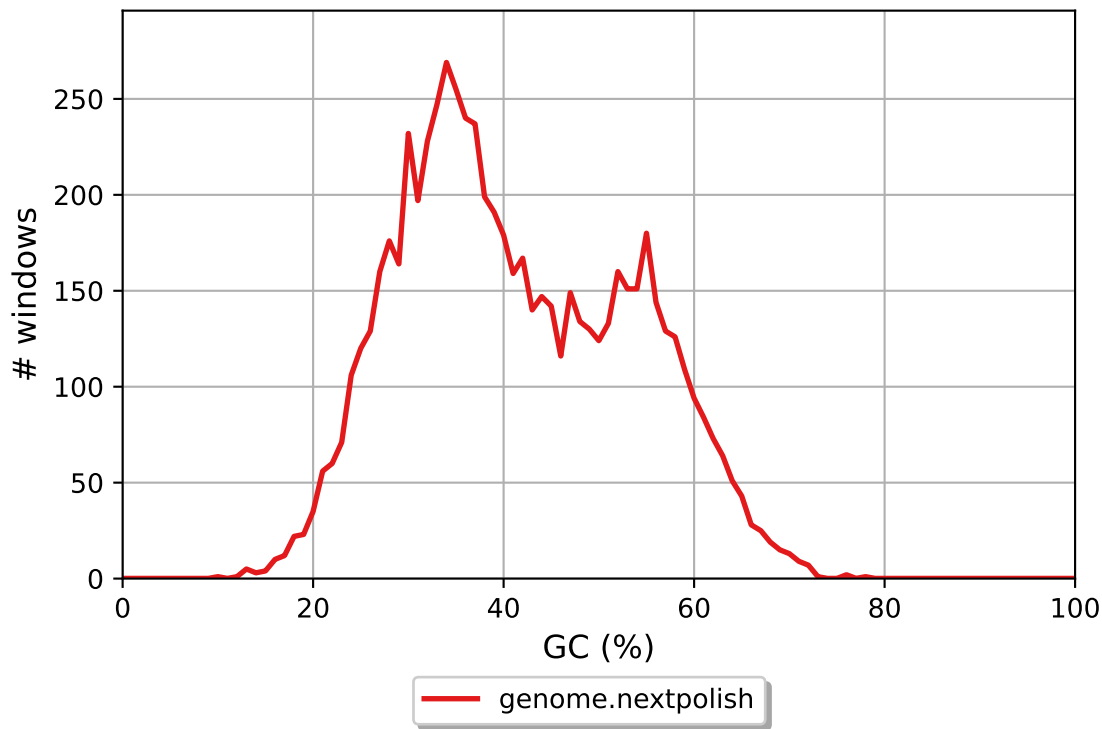
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

