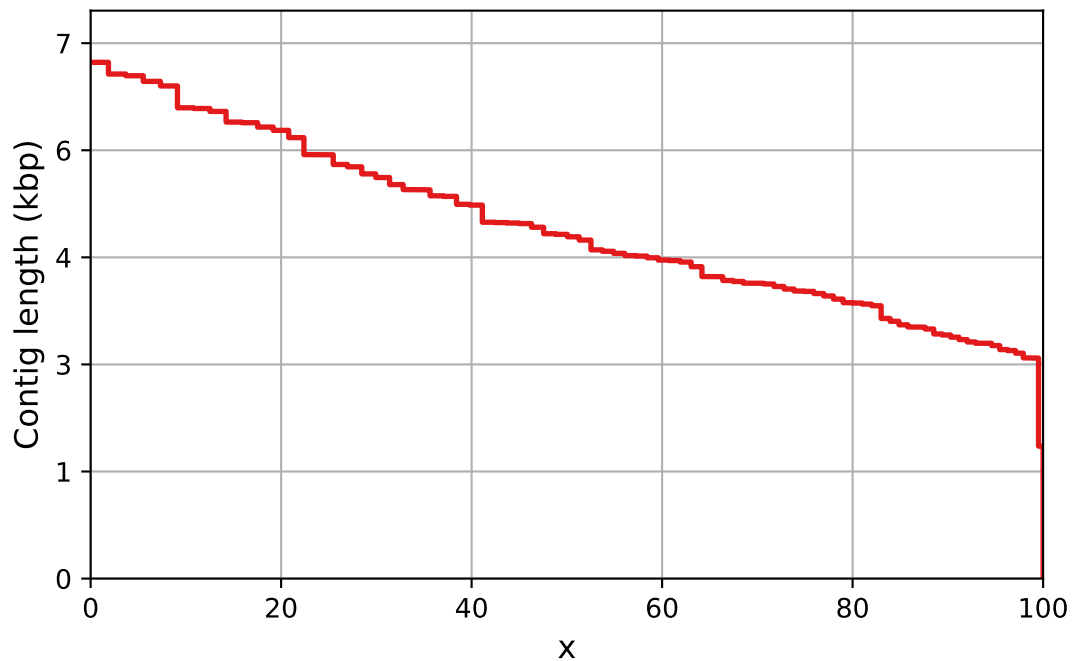


Report

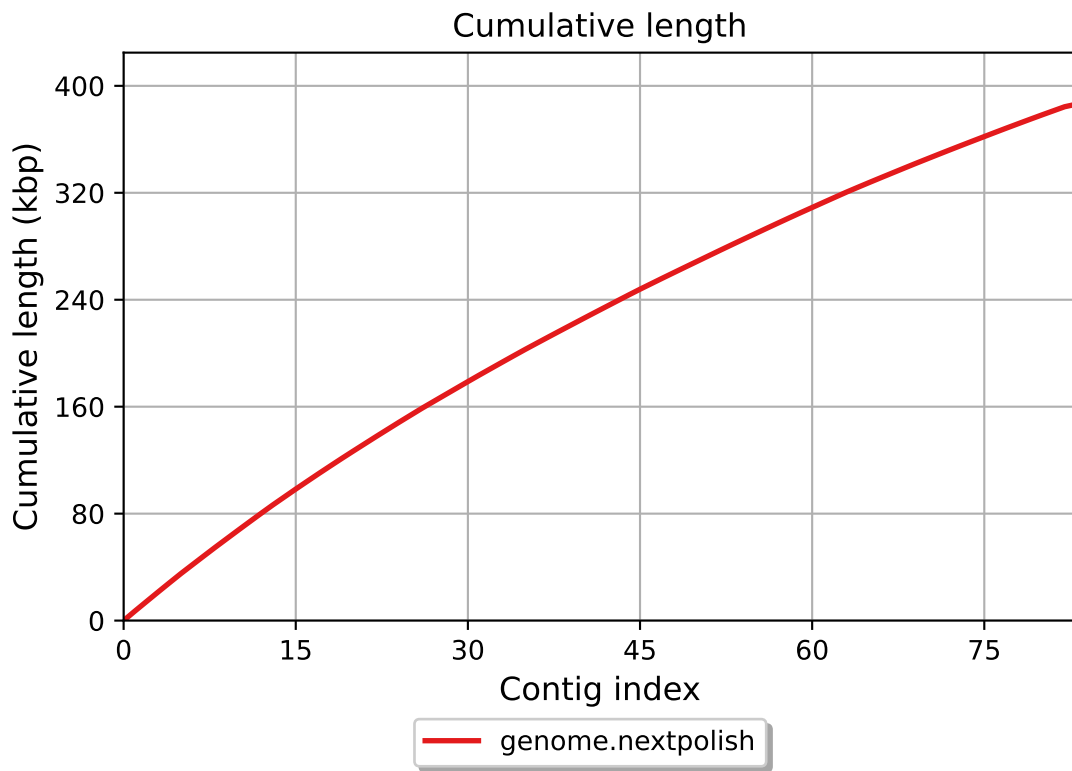
	genome.nextpolish
# contigs (>= 0 bp)	83
# contigs (>= 1000 bp)	83
# contigs (>= 5000 bp)	26
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	386215
Total length (>= 1000 bp)	386215
Total length (>= 5000 bp)	158833
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	83
Largest contig	7232
Total length	386215
GC (%)	51.74
N50	4822
N90	3412
auN	4946.2
L50	33
L90	71
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 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).

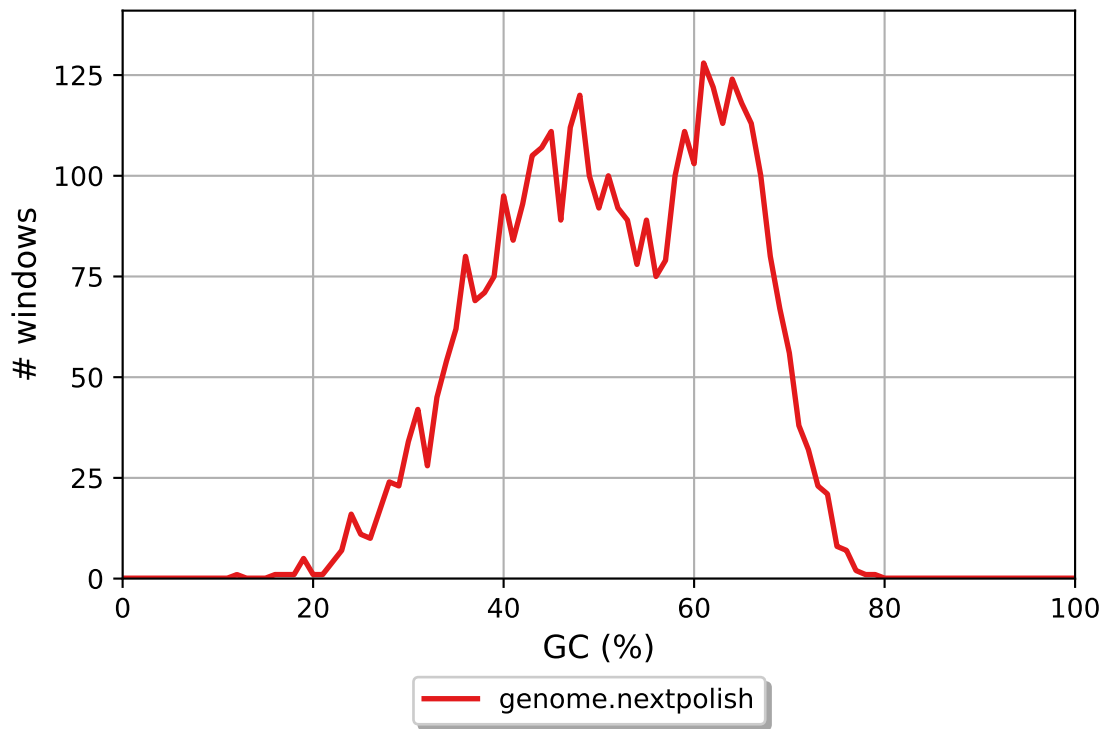
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

