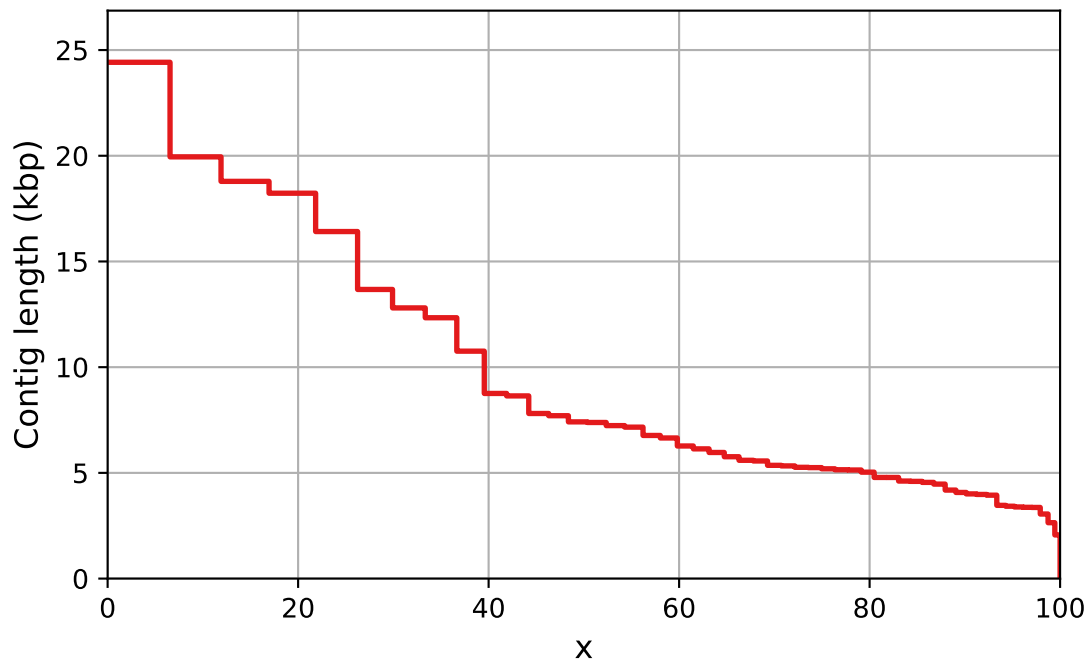


Report

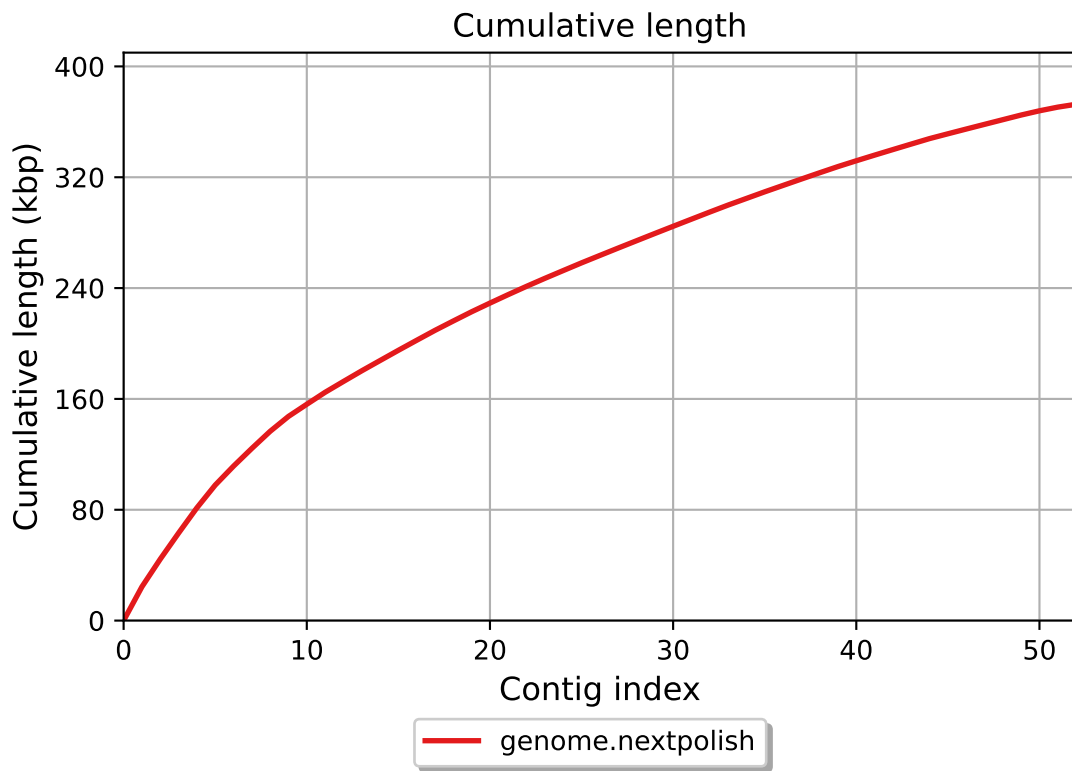
	genome.nextpolish
# contigs (>= 0 bp)	52
# contigs (>= 1000 bp)	52
# contigs (>= 5000 bp)	33
# contigs (>= 10000 bp)	9
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	372670
Total length (>= 1000 bp)	372670
Total length (>= 5000 bp)	299908
Total length (>= 10000 bp)	147375
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	52
Largest contig	24421
Total length	372670
GC (%)	60.74
N50	7413
N90	4074
auN	10350.5
L50	14
L90	41
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 1 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).

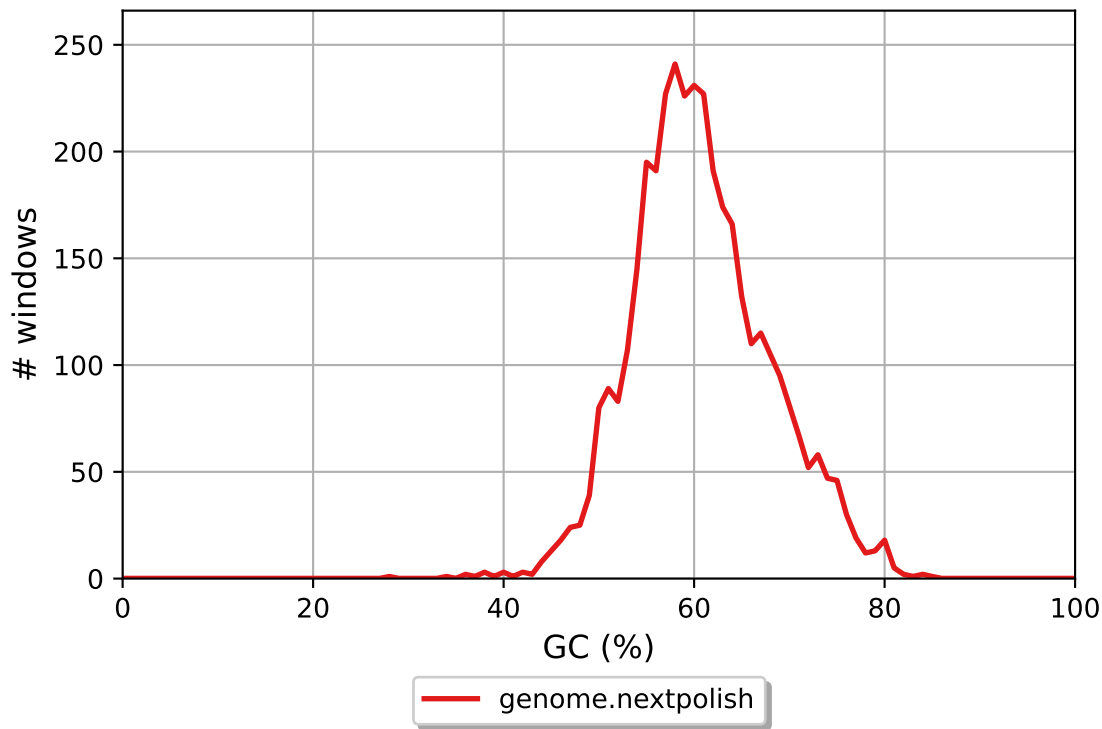
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

