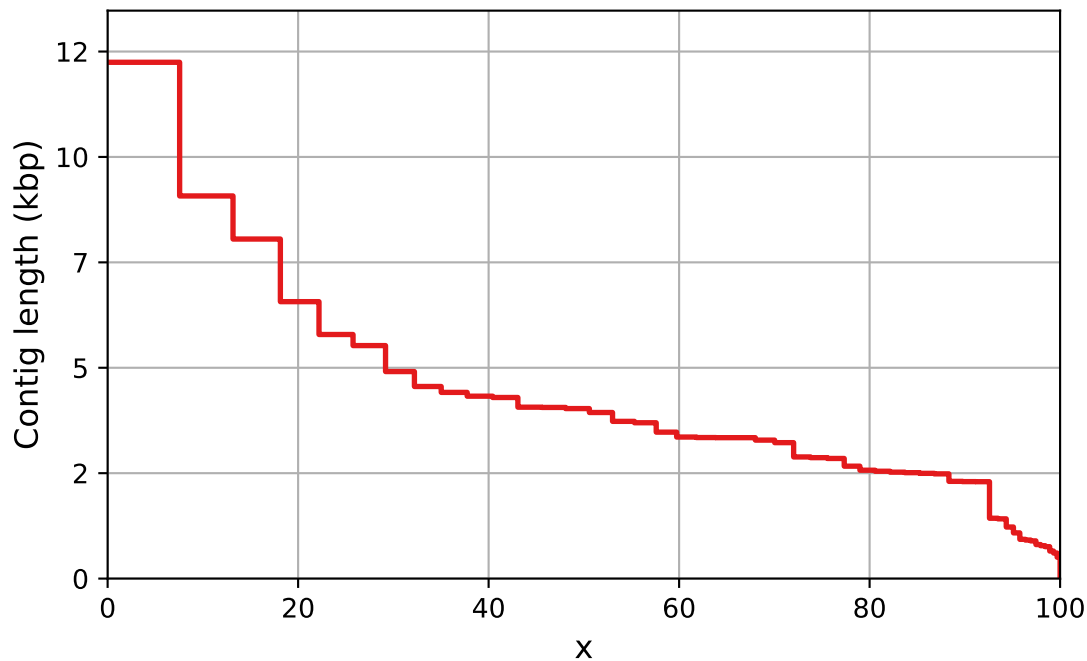


Report

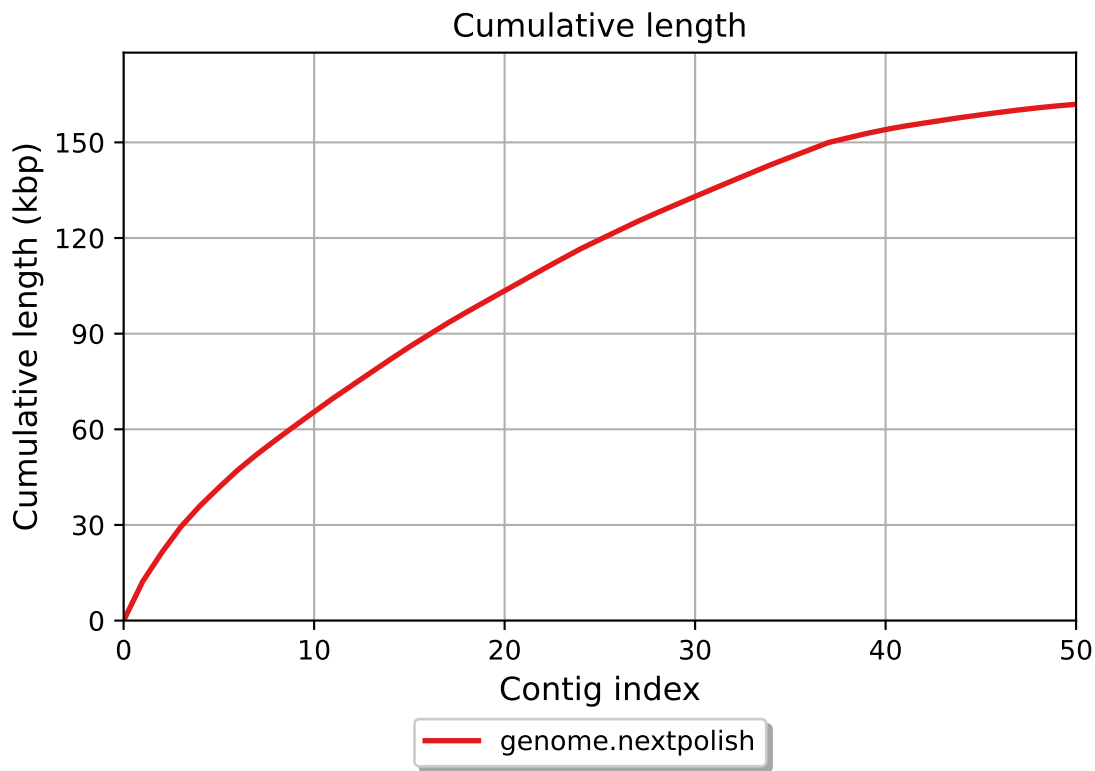
	genome.nextpolish
# contigs (>= 0 bp)	50
# contigs (>= 1000 bp)	41
# contigs (>= 5000 bp)	6
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	161954
Total length (>= 1000 bp)	155105
Total length (>= 5000 bp)	47256
Total length (>= 10000 bp)	12245
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	50
Largest contig	12245
Total length	161954
GC (%)	56.95
N50	4032
N90	2300
auN	4780.2
L50	14
L90	36
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	1 + 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).

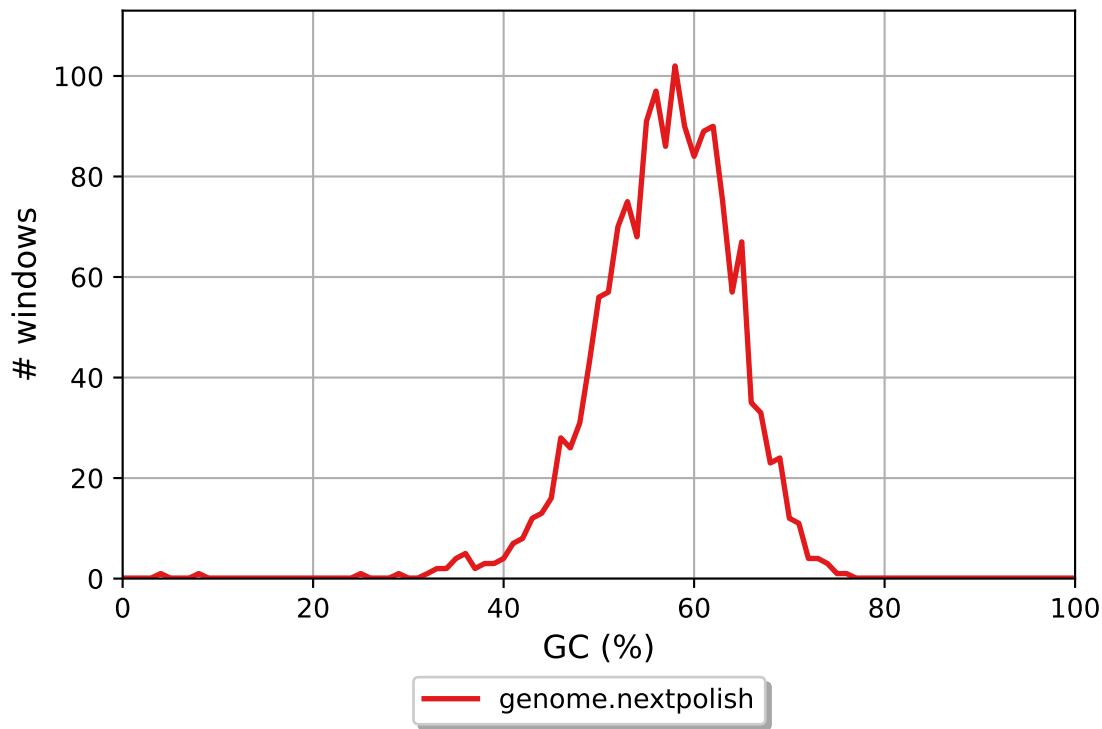
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

