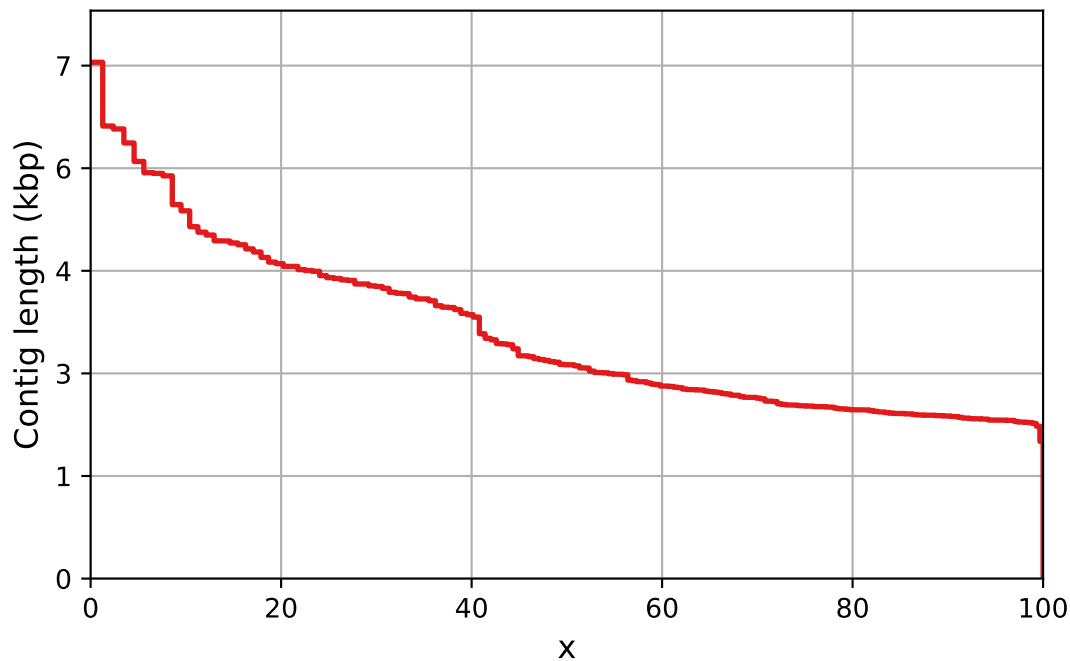


Report

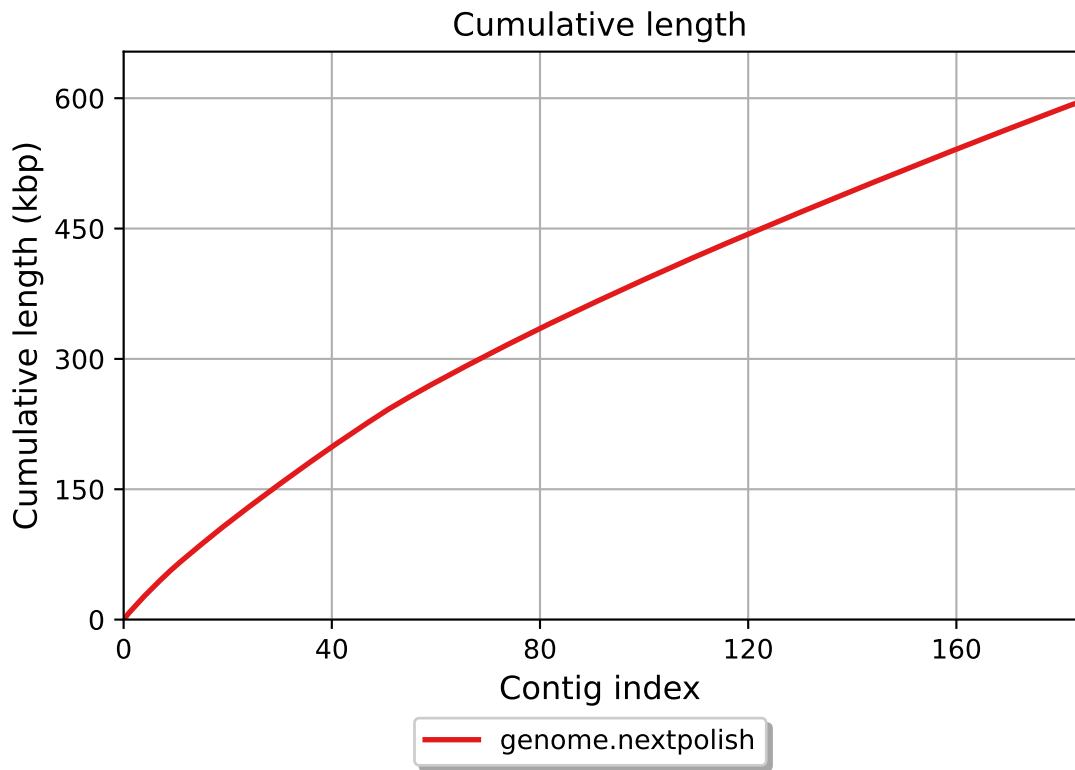
	genome.nextpolish
# contigs (>= 0 bp)	183
# contigs (>= 1000 bp)	183
# contigs (>= 5000 bp)	13
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	594181
Total length (>= 1000 bp)	594181
Total length (>= 5000 bp)	77032
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	183
Largest contig	7549
Total length	594181
GC (%)	40.34
N50	3126
N90	2377
auN	3600.1
L50	68
L90	158
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	8 + 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).

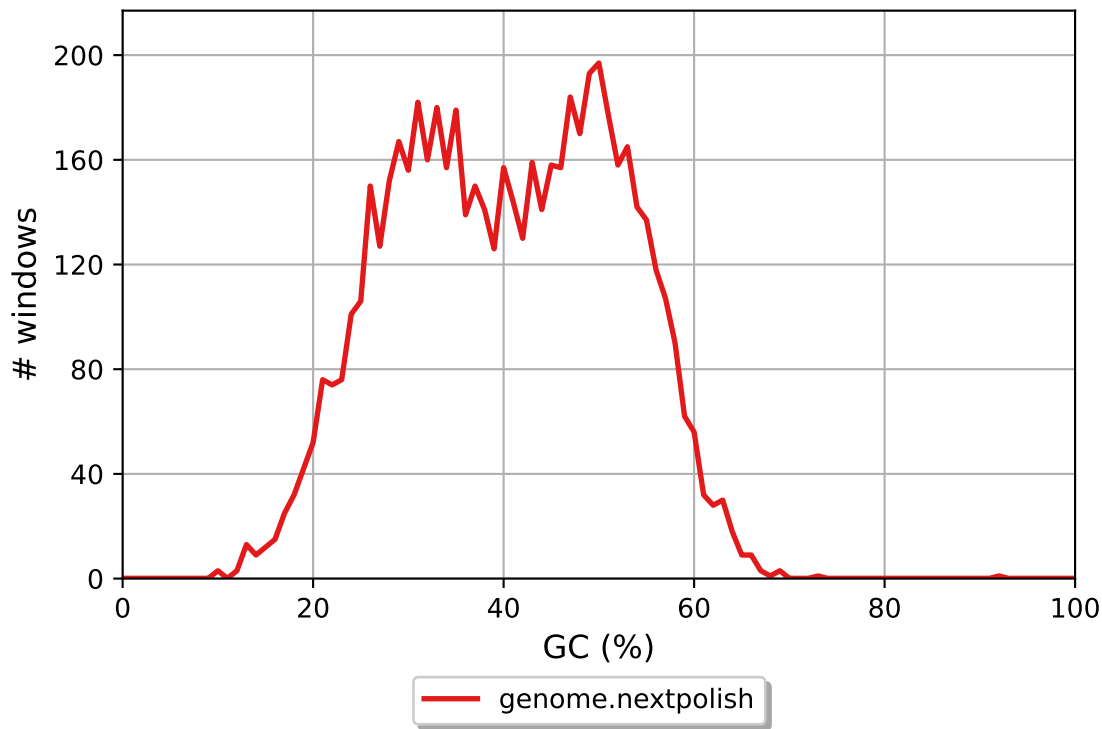
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

