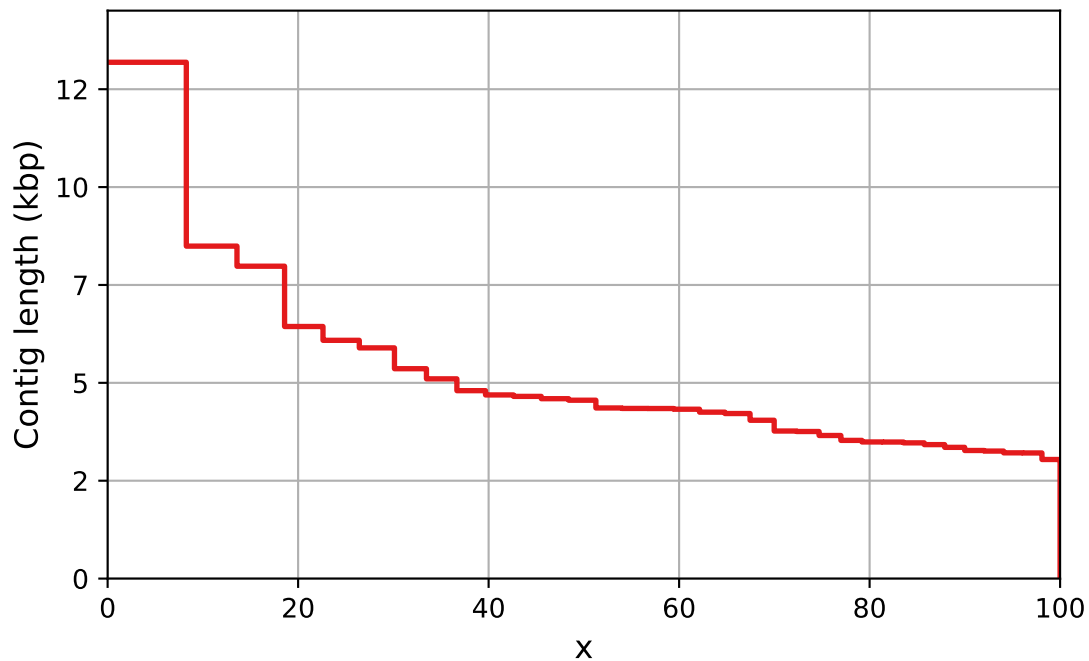


Report

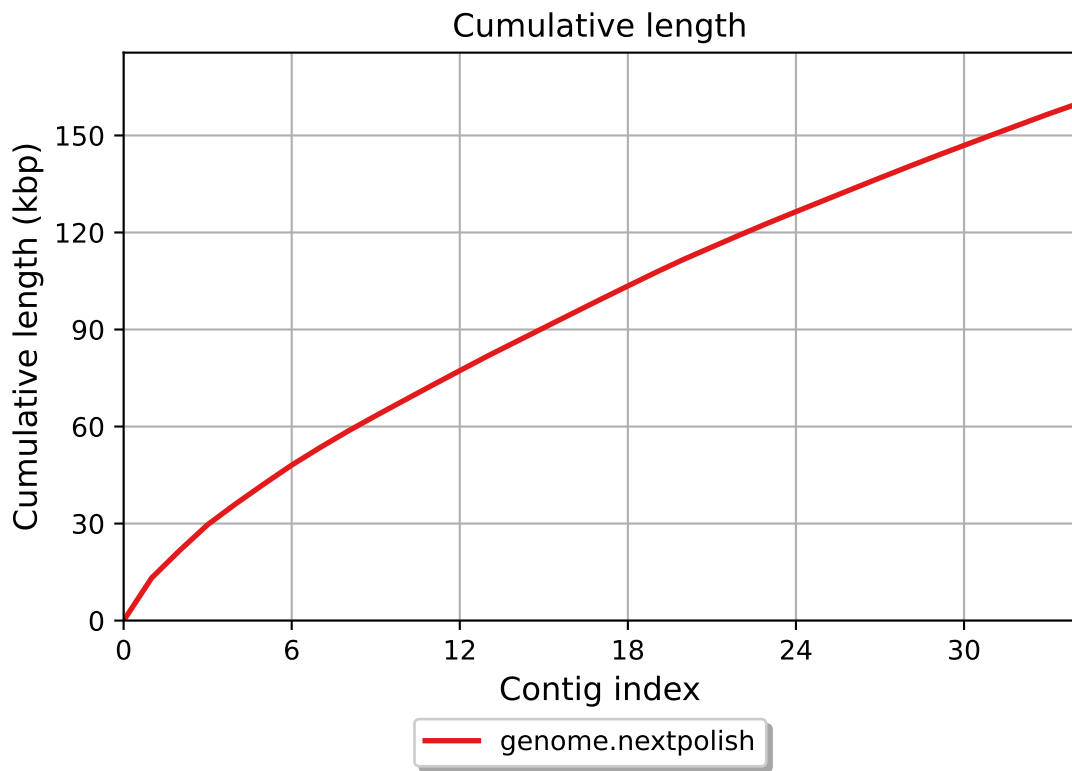
	genome.nextpolish
# contigs (>= 0 bp)	34
# contigs (>= 1000 bp)	34
# contigs (>= 5000 bp)	8
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	159643
Total length (>= 1000 bp)	159643
Total length (>= 5000 bp)	58540
Total length (>= 10000 bp)	13188
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	34
Largest contig	13188
Total length	159643
GC (%)	46.39
N50	4556
N90	3272
auN	5503.4
L50	13
L90	30
# 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).

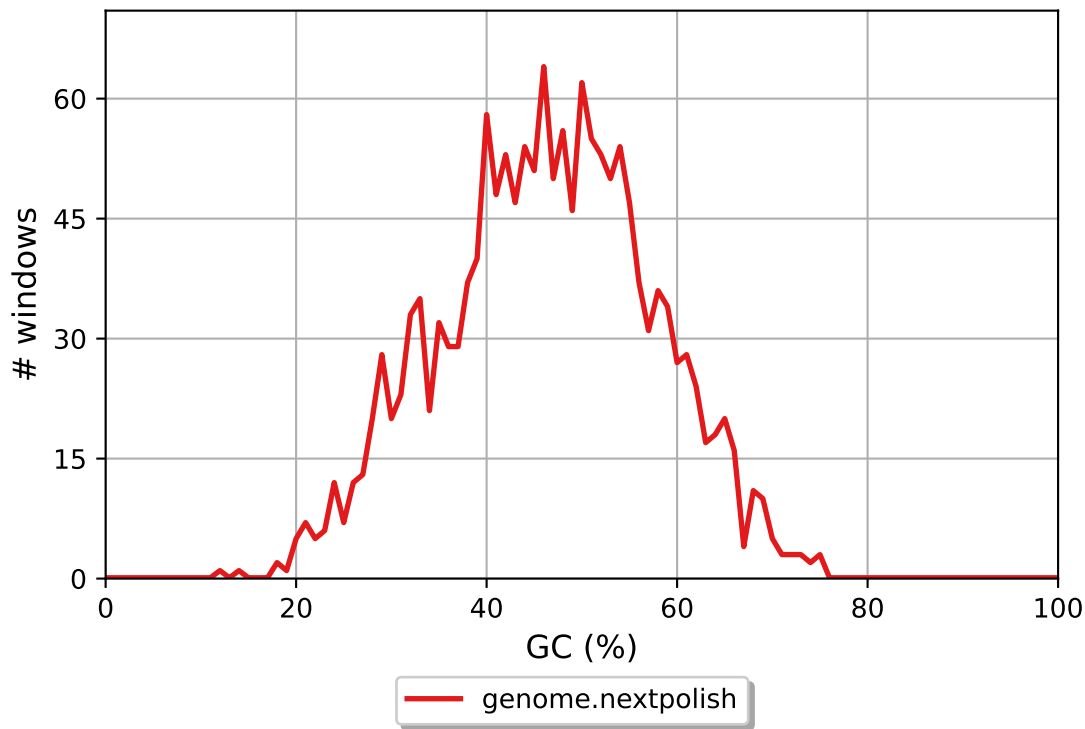
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

