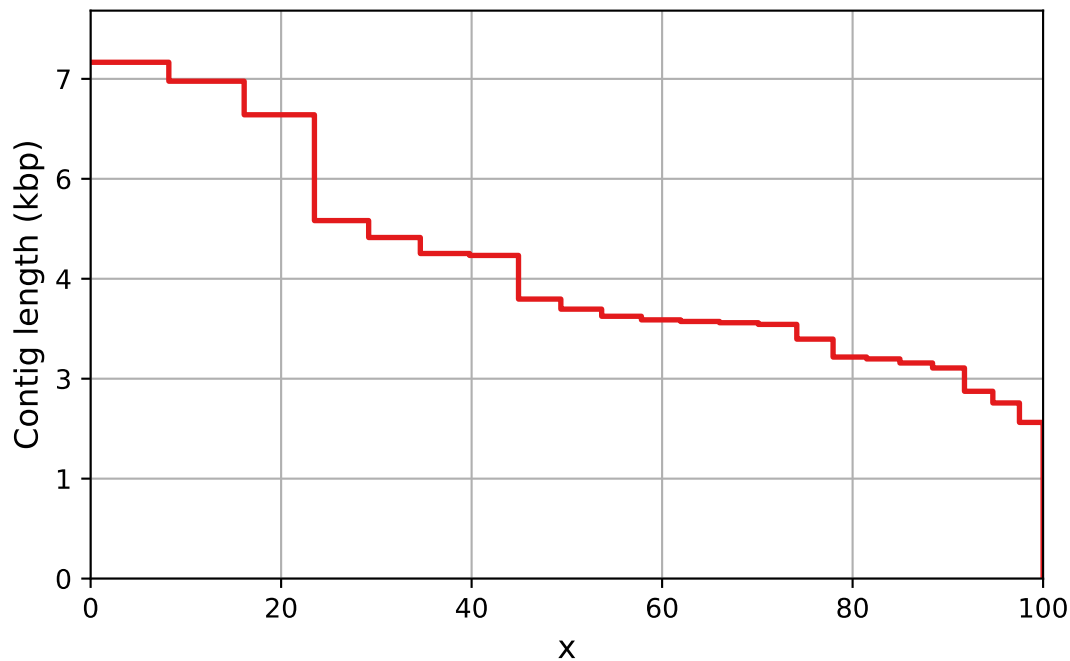


Report

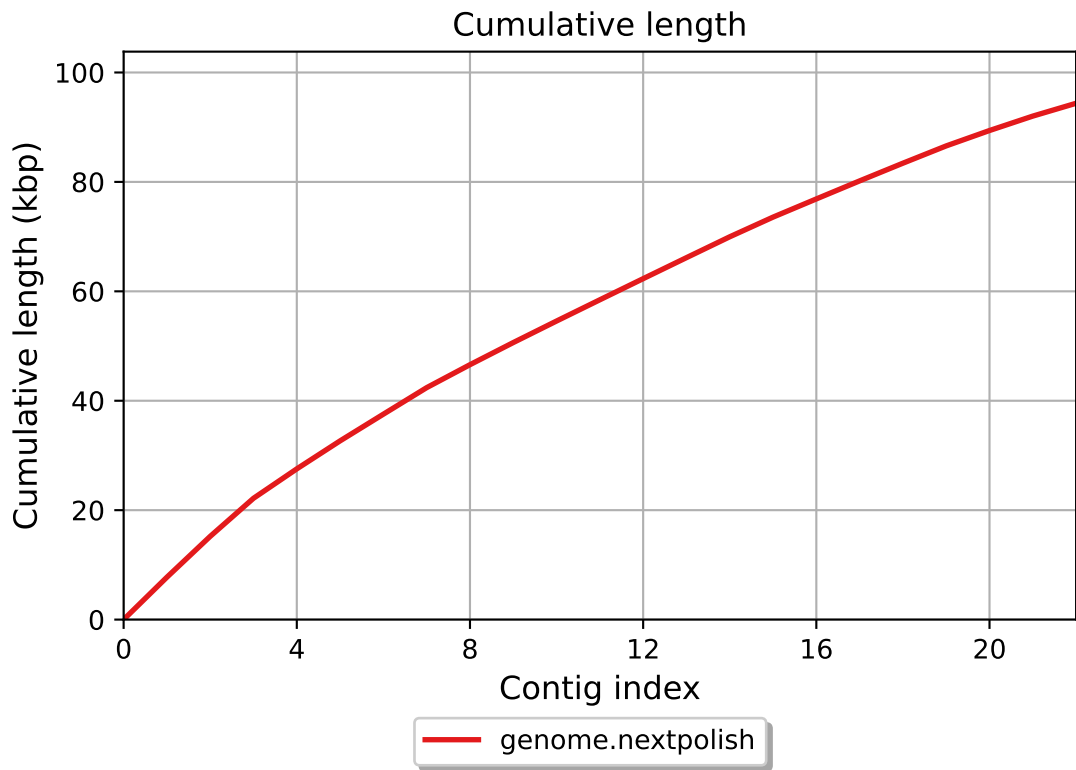
	genome.nextpolish
# contigs (>= 0 bp)	22
# contigs (>= 1000 bp)	22
# contigs (>= 5000 bp)	5
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	94373
Total length (>= 1000 bp)	94373
Total length (>= 5000 bp)	32667
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	22
Largest contig	7749
Total length	94373
GC (%)	55.67
N50	4044
N90	3161
auN	4778.9
L50	9
L90	19
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 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).

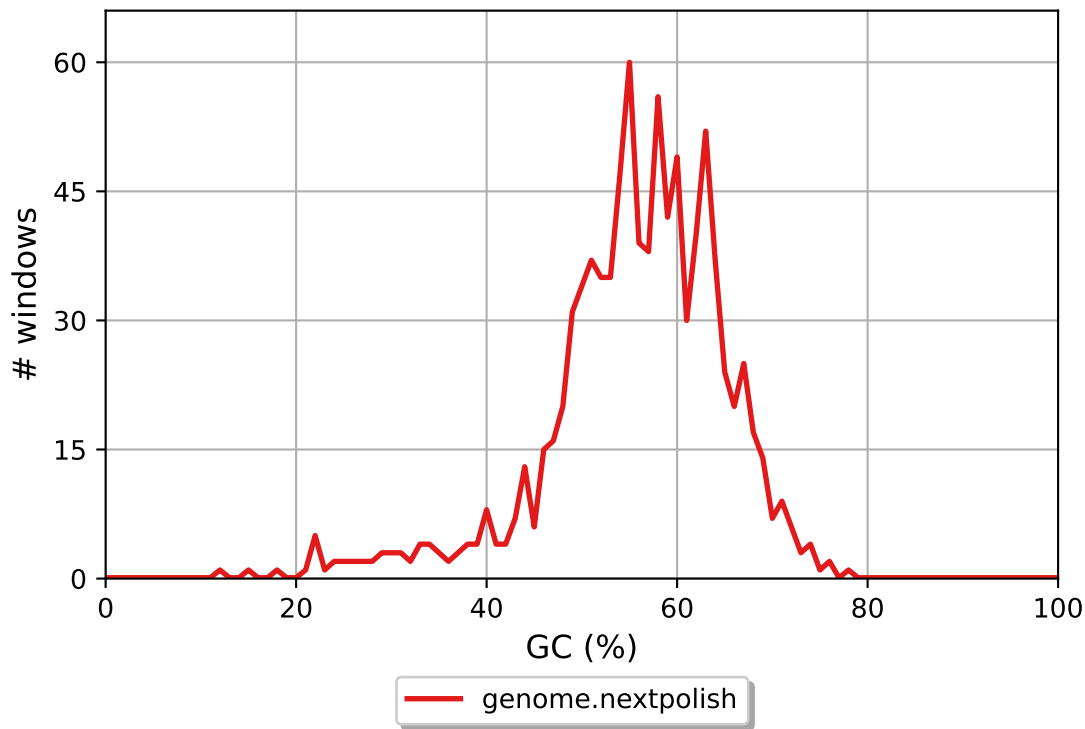
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

