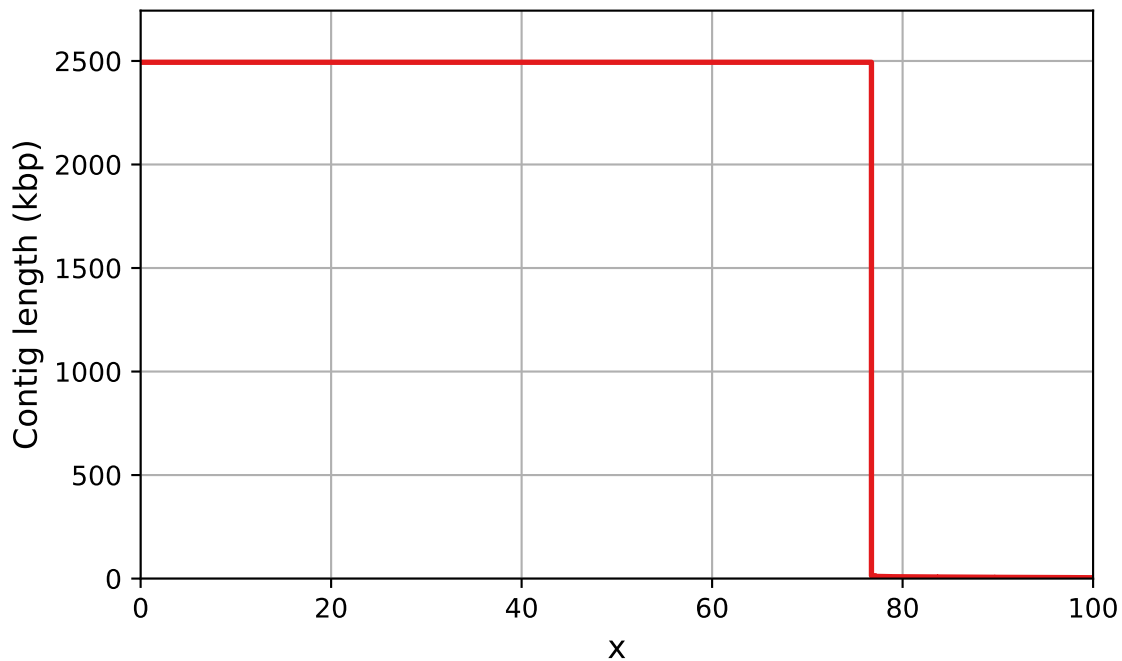


Report

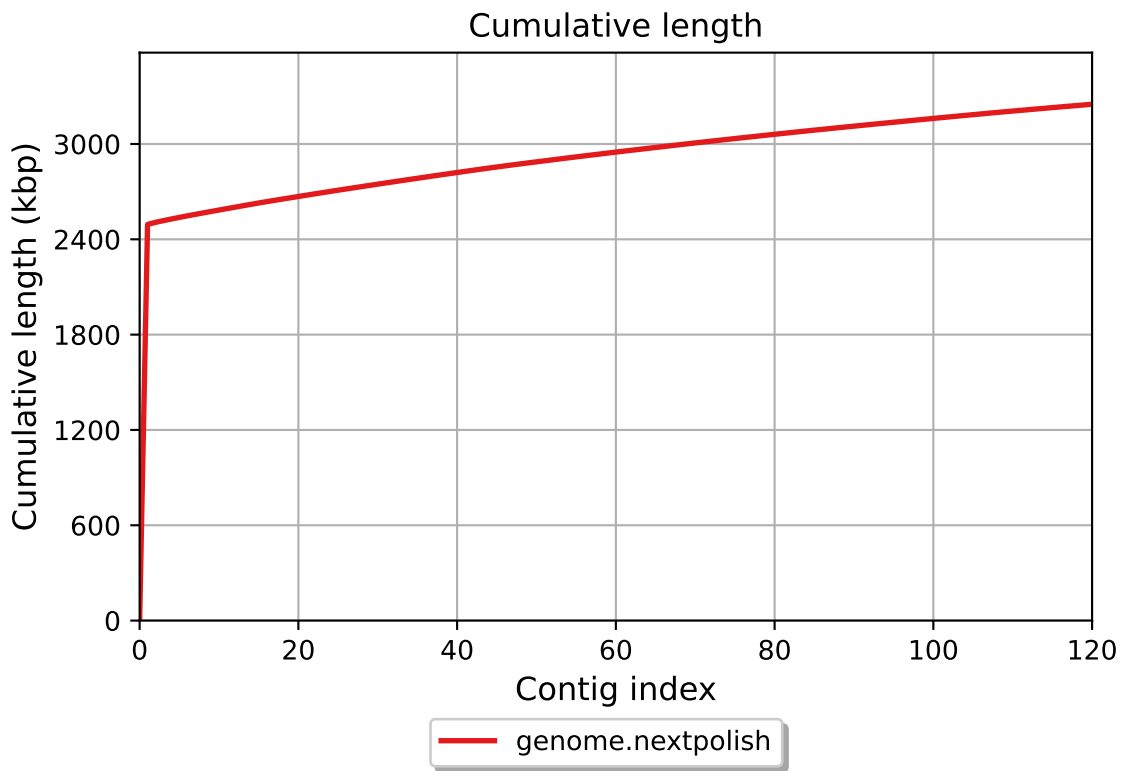
	genome.nextpolish
# contigs (>= 0 bp)	120
# contigs (>= 1000 bp)	120
# contigs (>= 5000 bp)	88
# contigs (>= 10000 bp)	5
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	3250022
Total length (>= 1000 bp)	3250022
Total length (>= 5000 bp)	3102295
Total length (>= 10000 bp)	2537921
Total length (>= 25000 bp)	2493726
Total length (>= 50000 bp)	2493726
# contigs	120
Largest contig	2493726
Total length	3250022
GC (%)	57.95
N50	2493726
N90	6038
auN	1915007.5
L50	1
L90	57
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	10 + 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).

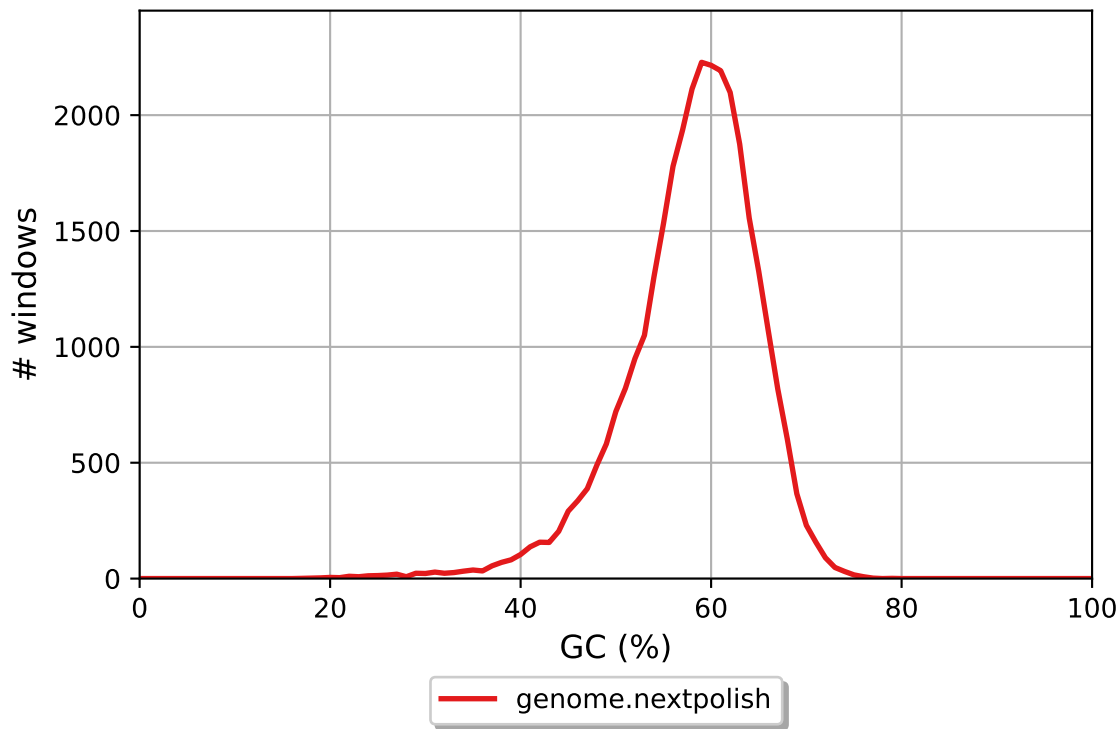
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

