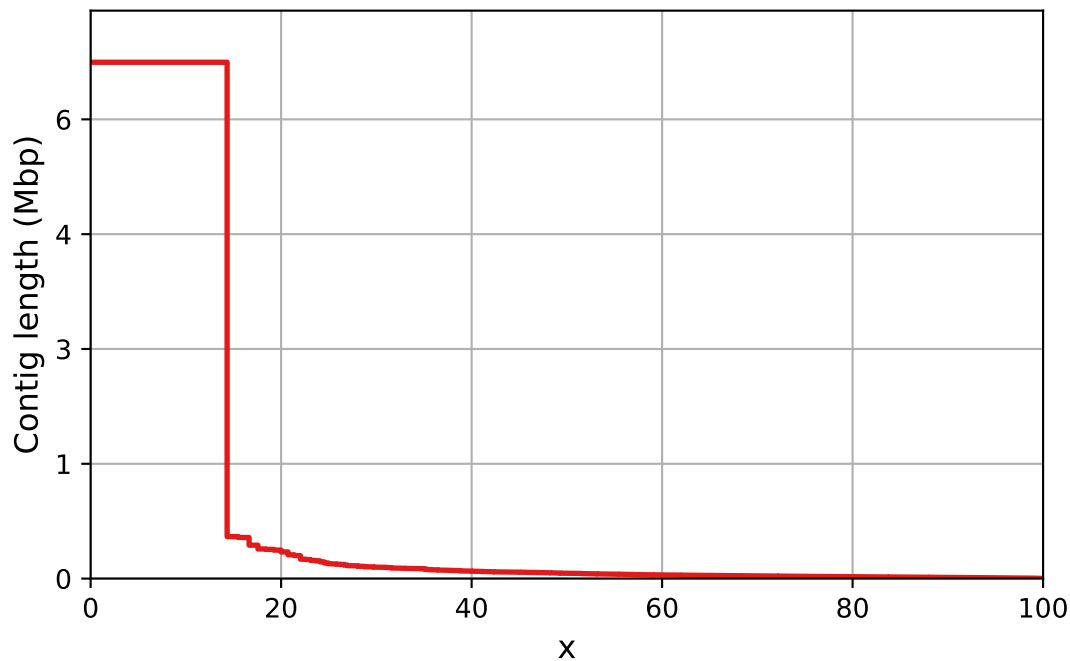


Report

	genome.nextpolish
# contigs (>= 0 bp)	1281
# contigs (>= 1000 bp)	1278
# contigs (>= 5000 bp)	1176
# contigs (>= 10000 bp)	919
# contigs (>= 25000 bp)	478
# contigs (>= 50000 bp)	191
Total length (>= 0 bp)	47073784
Total length (>= 1000 bp)	47072052
Total length (>= 5000 bp)	46719962
Total length (>= 10000 bp)	44810847
Total length (>= 25000 bp)	37480228
Total length (>= 50000 bp)	27393240
# contigs	1281
Largest contig	6745846
Total length	47073784
GC (%)	51.57
N50	69648
N90	15358
auN	1050955.8
L50	127
L90	725
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	75 + 5 part

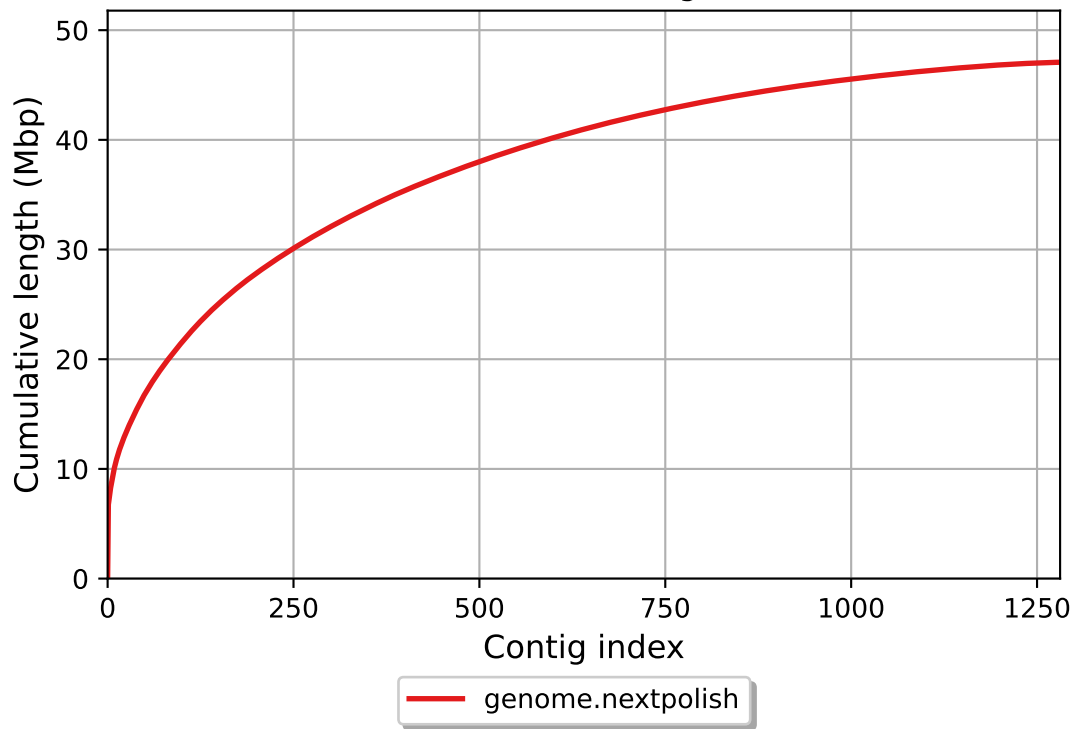
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

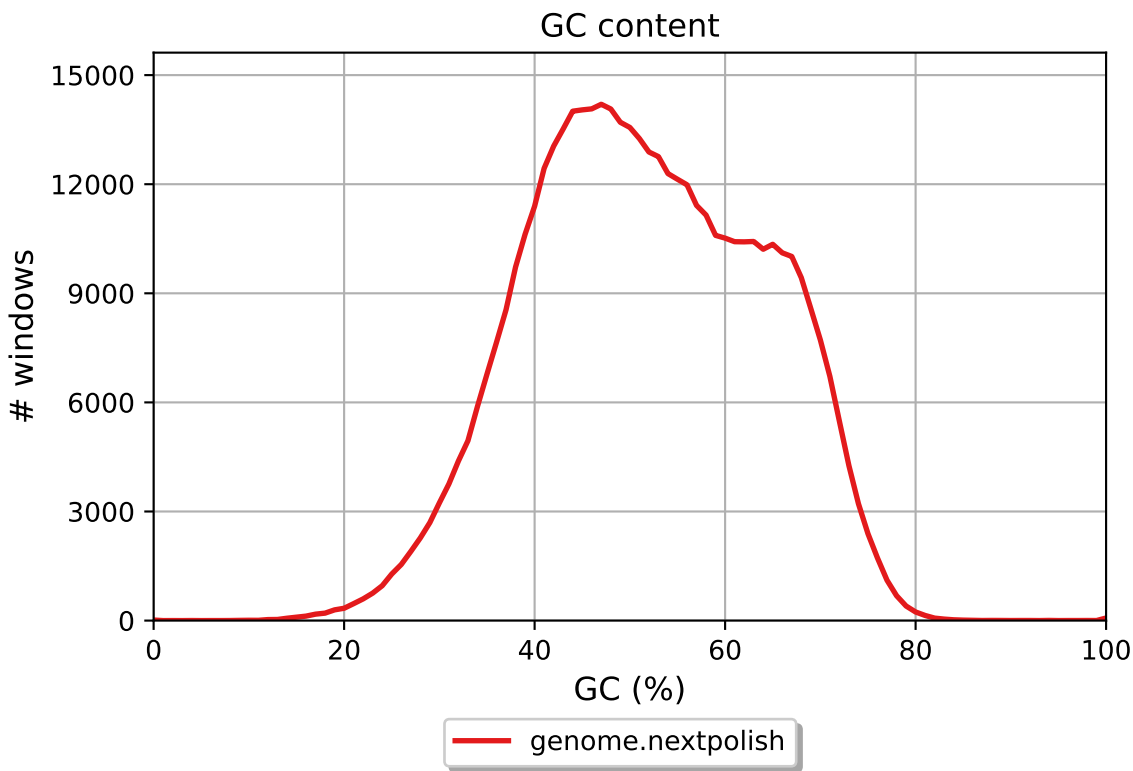
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

