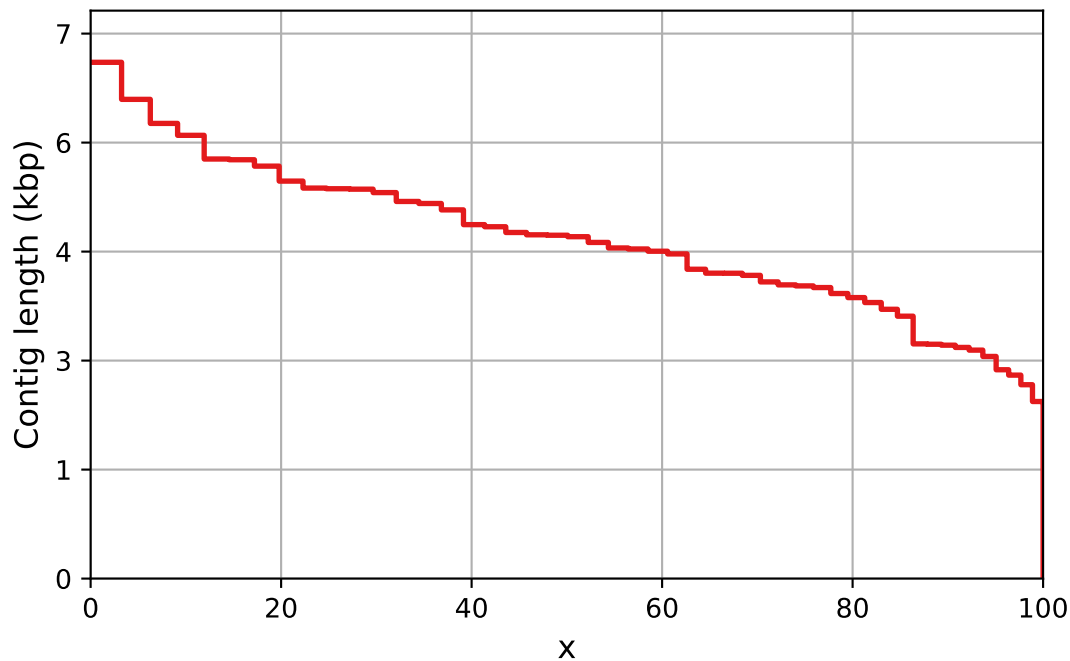


Report

	genome.nextpolish
# contigs (>= 0 bp)	49
# contigs (>= 1000 bp)	49
# contigs (>= 5000 bp)	15
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	218633
Total length (>= 1000 bp)	218633
Total length (>= 5000 bp)	85584
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	49
Largest contig	7105
Total length	218633
GC (%)	49.48
N50	4725
N90	3212
auN	4711.2
L50	20
L90	42
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

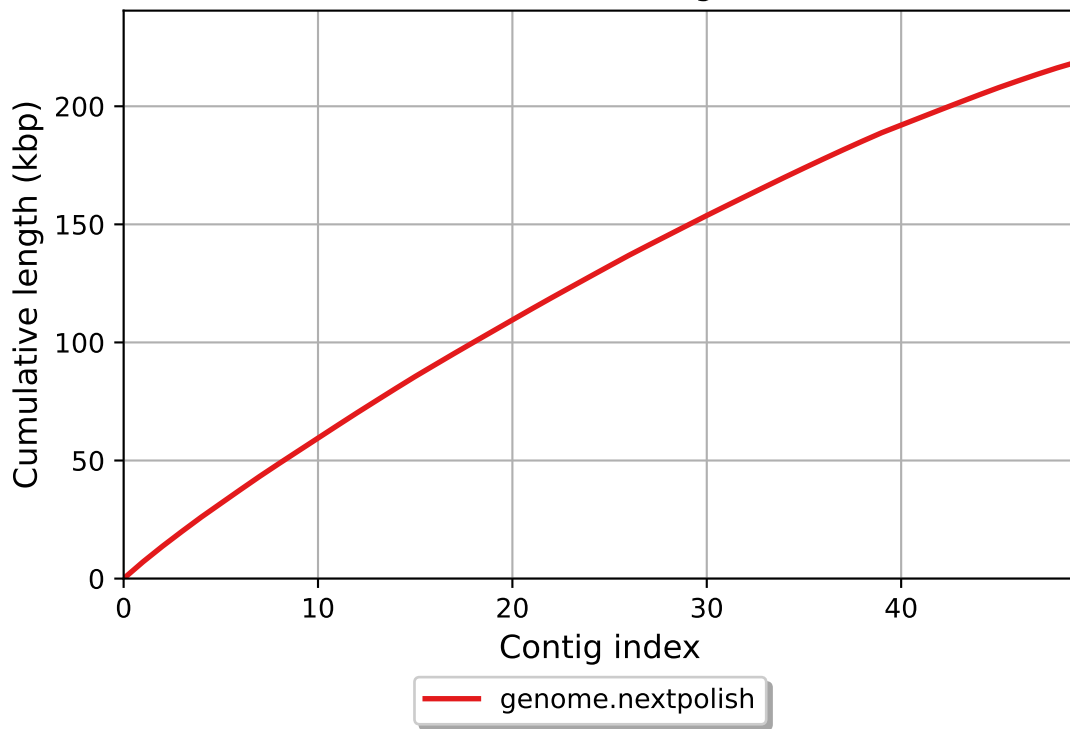
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

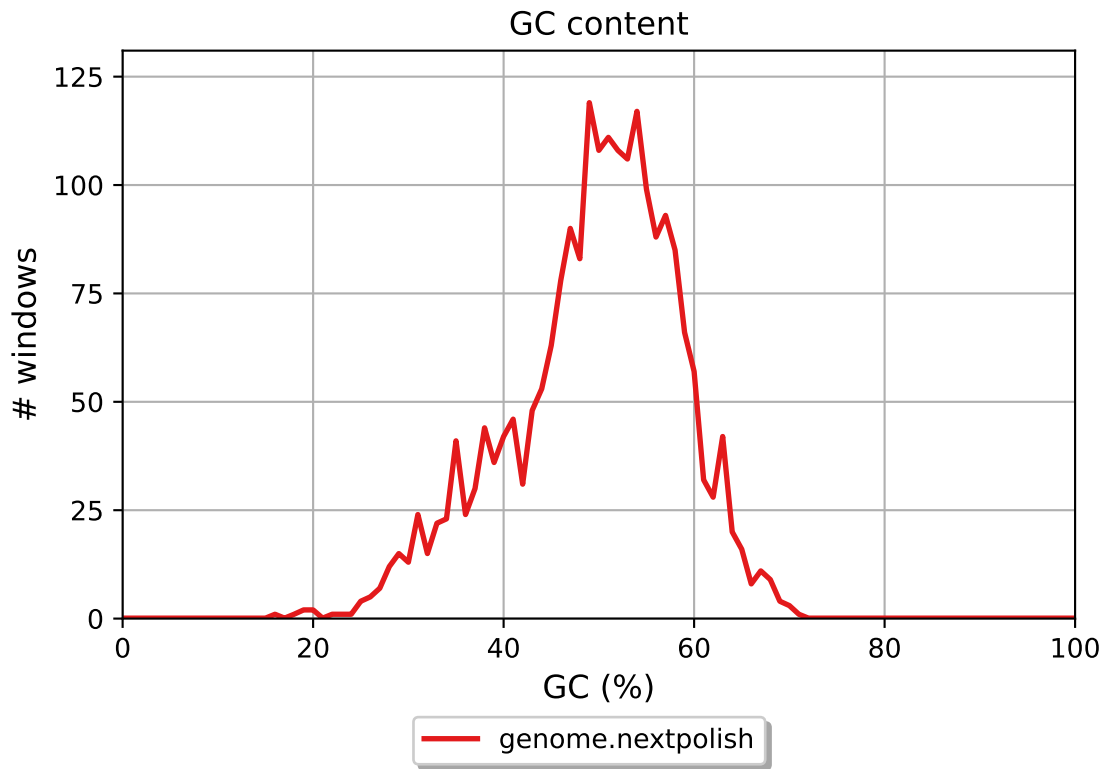
Nx



genome.nextpolish

Cumulative length





genome.nextpolish GC content

