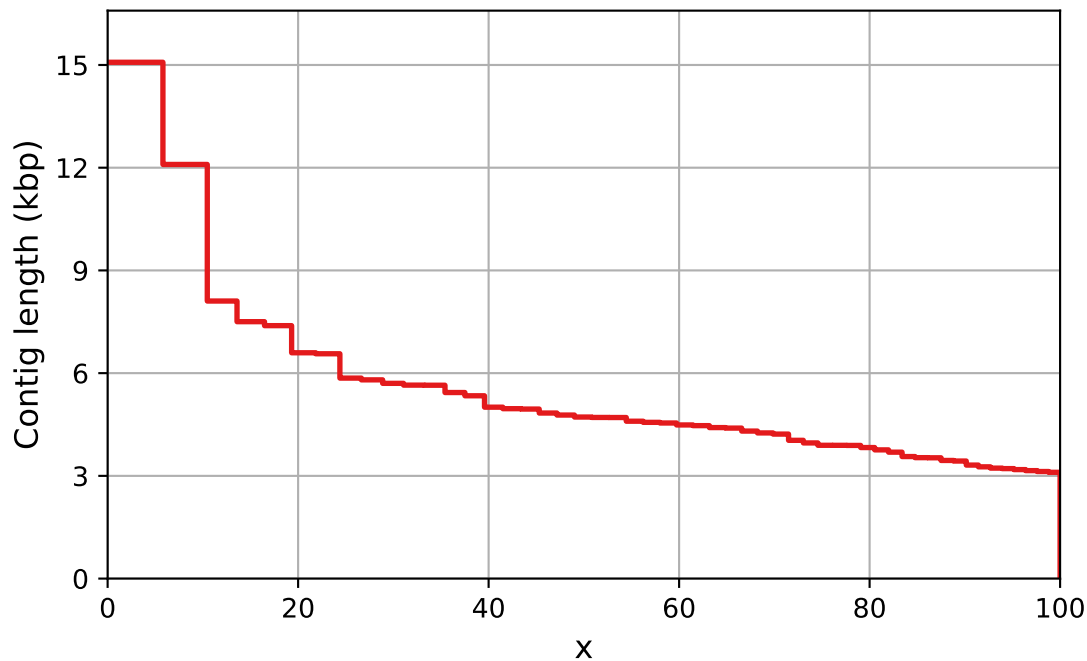


Report

	genome.nextpolish
# contigs (>= 0 bp)	53
# contigs (>= 1000 bp)	53
# contigs (>= 5000 bp)	15
# contigs (>= 10000 bp)	2
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	259707
Total length (>= 1000 bp)	259707
Total length (>= 5000 bp)	107768
Total length (>= 10000 bp)	27174
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	53
Largest contig	15079
Total length	259707
GC (%)	51.71
N50	4718
N90	3433
auN	5795.0
L50	20
L90	45
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

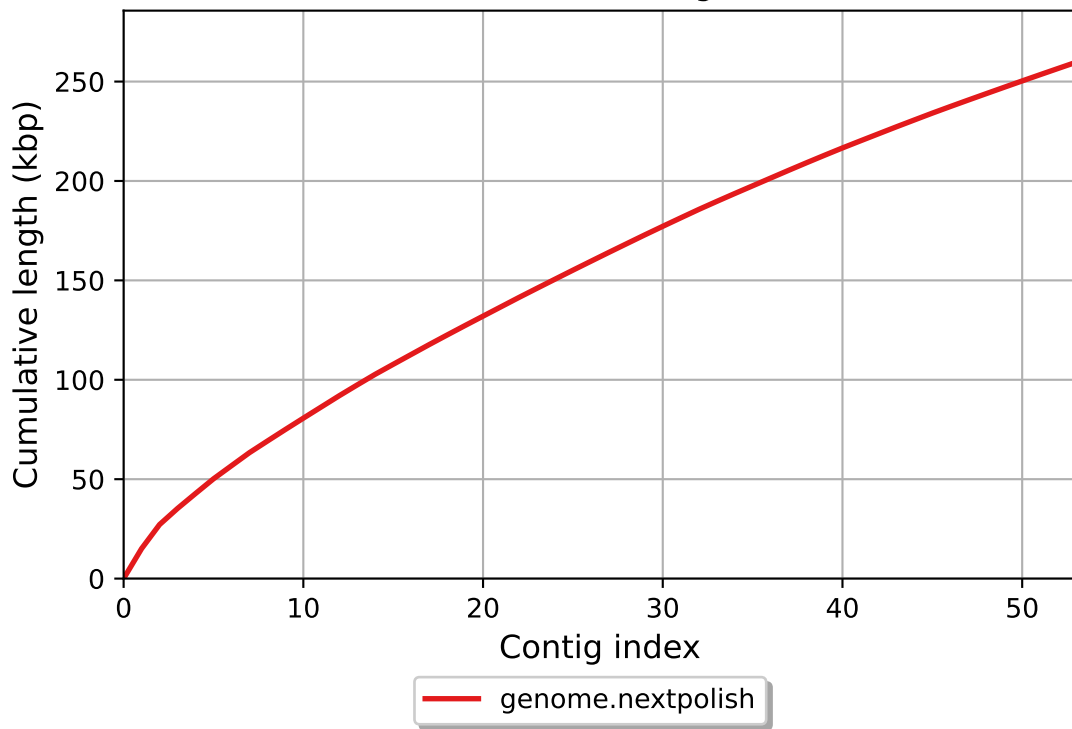
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

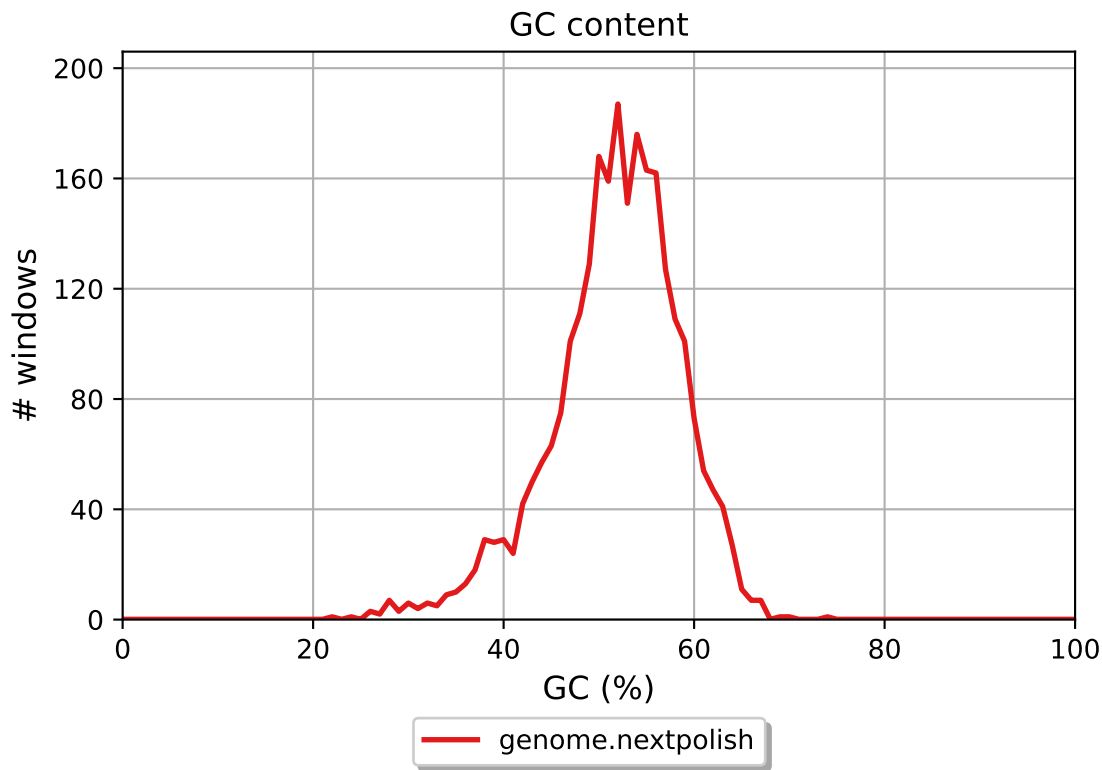
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

