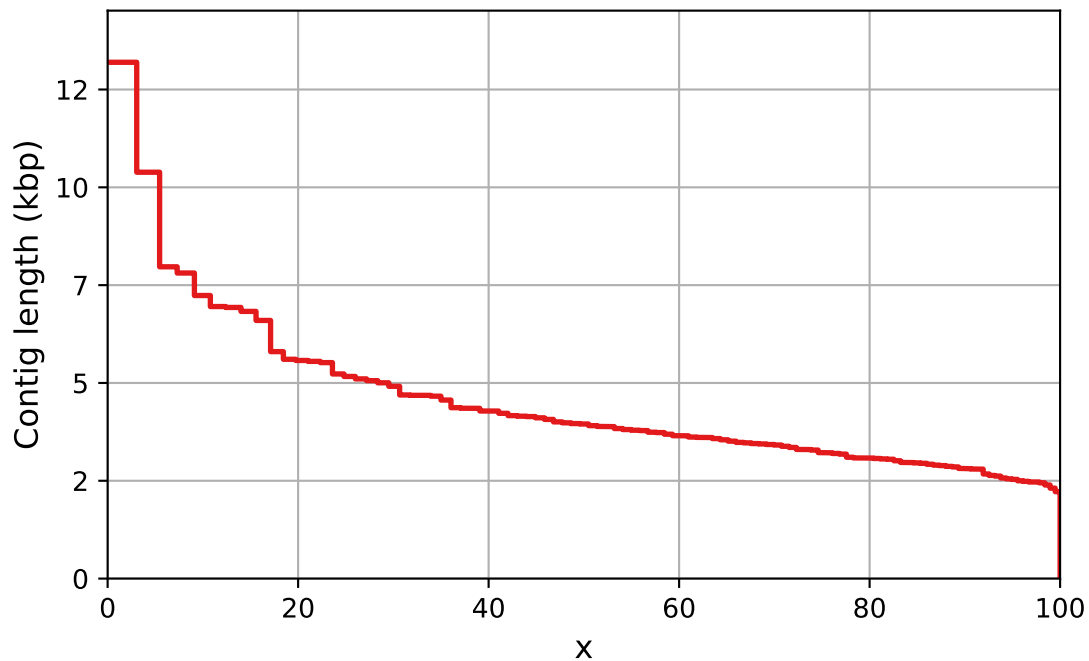


Report

	genome.nextpolish
# contigs (>= 0 bp)	108
# contigs (>= 1000 bp)	108
# contigs (>= 5000 bp)	19
# contigs (>= 10000 bp)	2
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	431974
Total length (>= 1000 bp)	431974
Total length (>= 5000 bp)	127522
Total length (>= 10000 bp)	23582
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	108
Largest contig	13196
Total length	431974
GC (%)	50.86
N50	3951
N90	2809
auN	4659.0
L50	40
L90	92
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 0 part

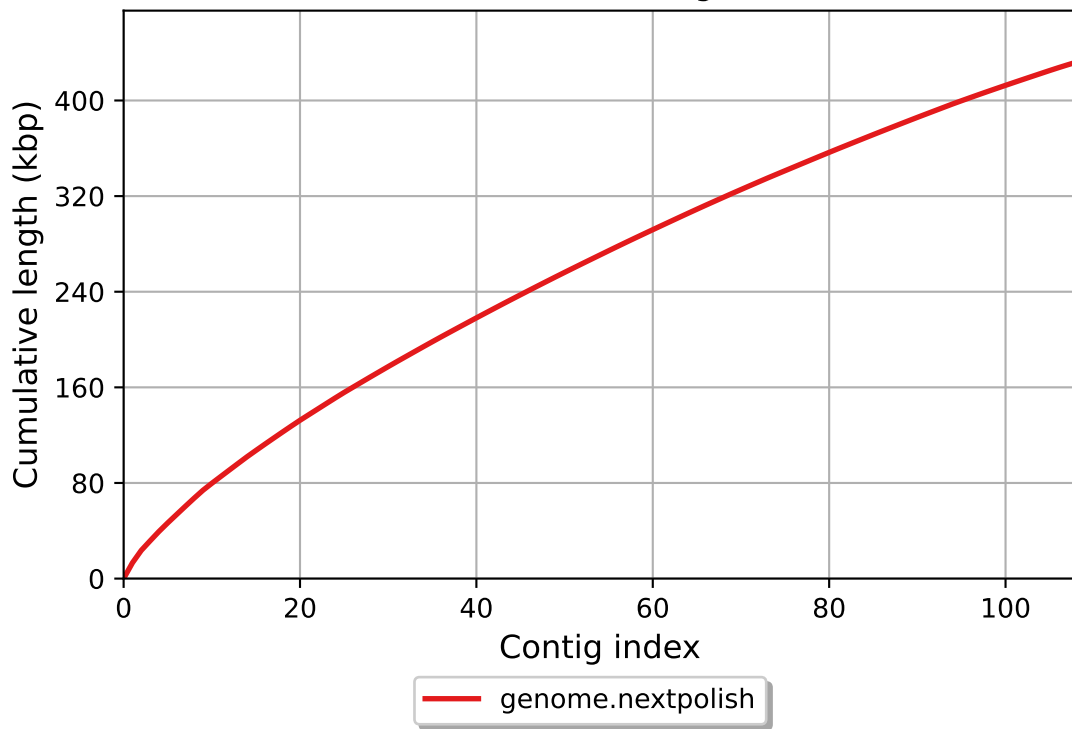
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

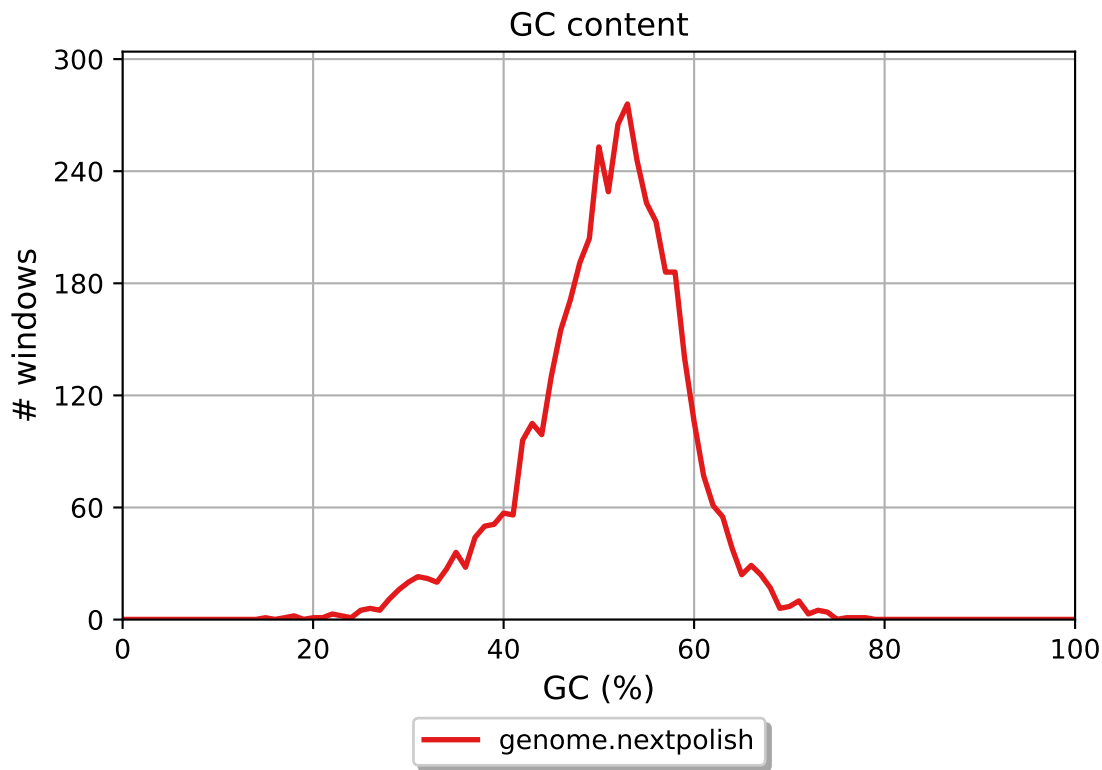
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

