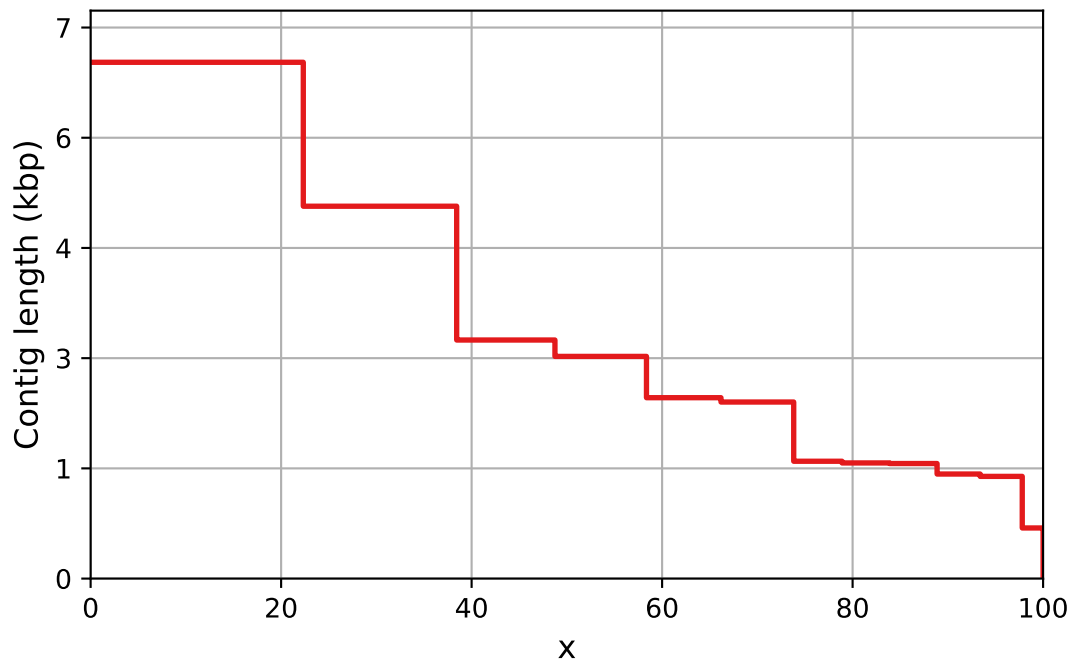


Report

	genome.nextpolish
# contigs (>= 0 bp)	12
# contigs (>= 1000 bp)	11
# contigs (>= 5000 bp)	2
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	31471
Total length (>= 1000 bp)	30782
Total length (>= 5000 bp)	12095
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	12
Largest contig	7027
Total length	31471
GC (%)	51.24
N50	3024
N90	1423
auN	3765.4
L50	4
L90	10
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 0 part

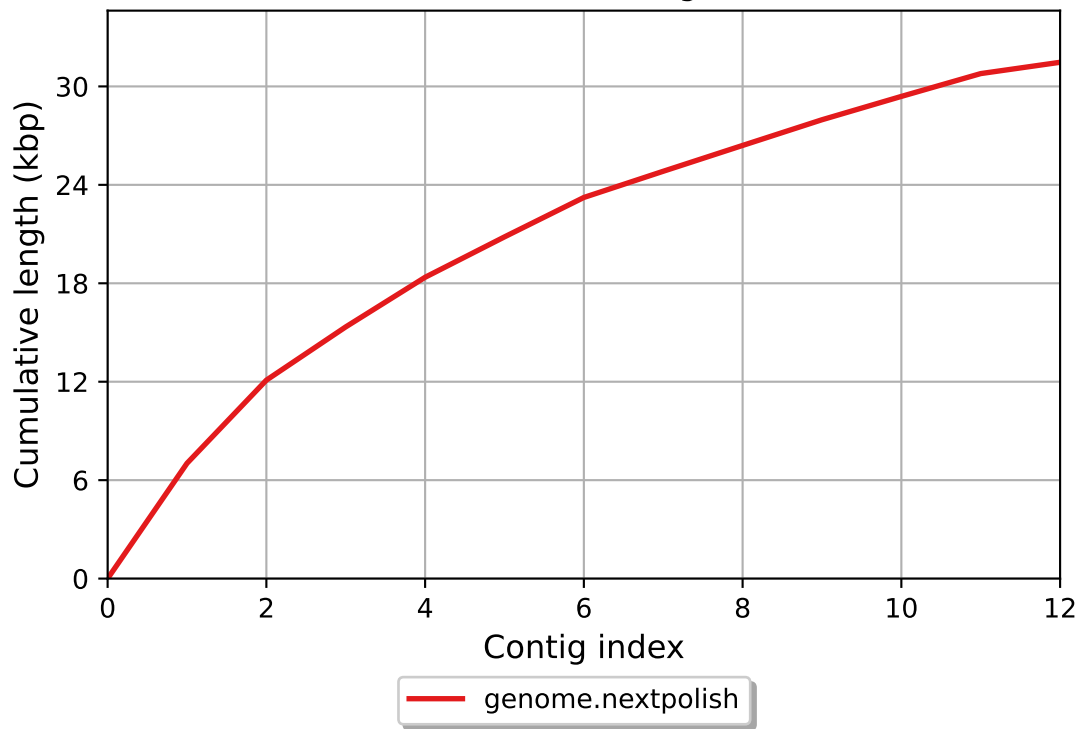
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

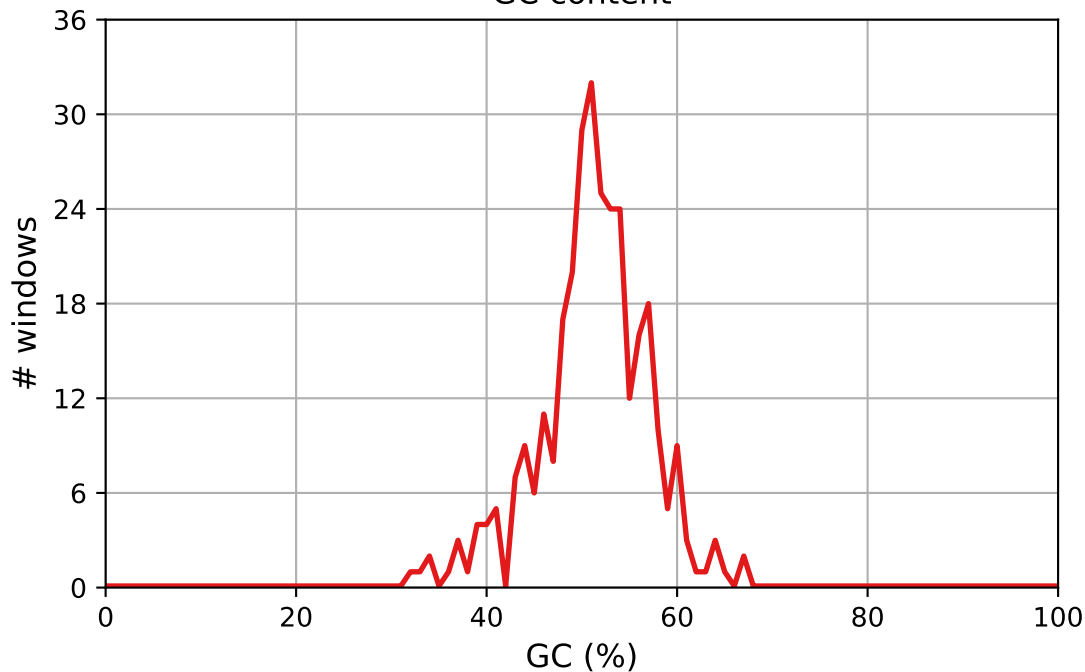


— genome.nextpolish

Cumulative length



GC content



— genome.nextpolish

genome.nextpolish GC content

