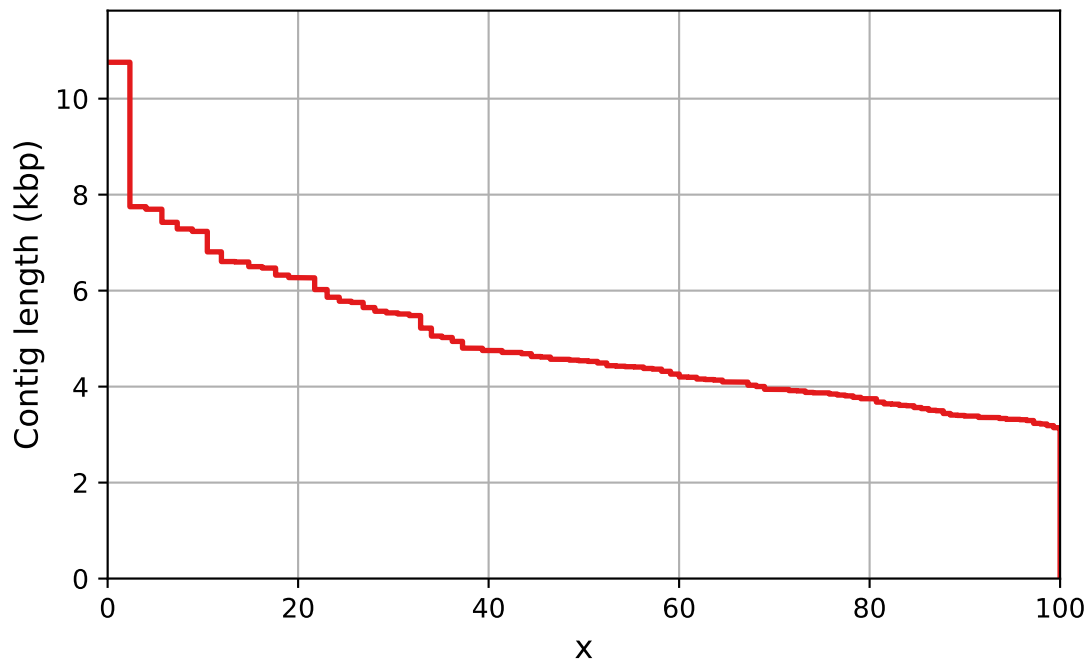


Report

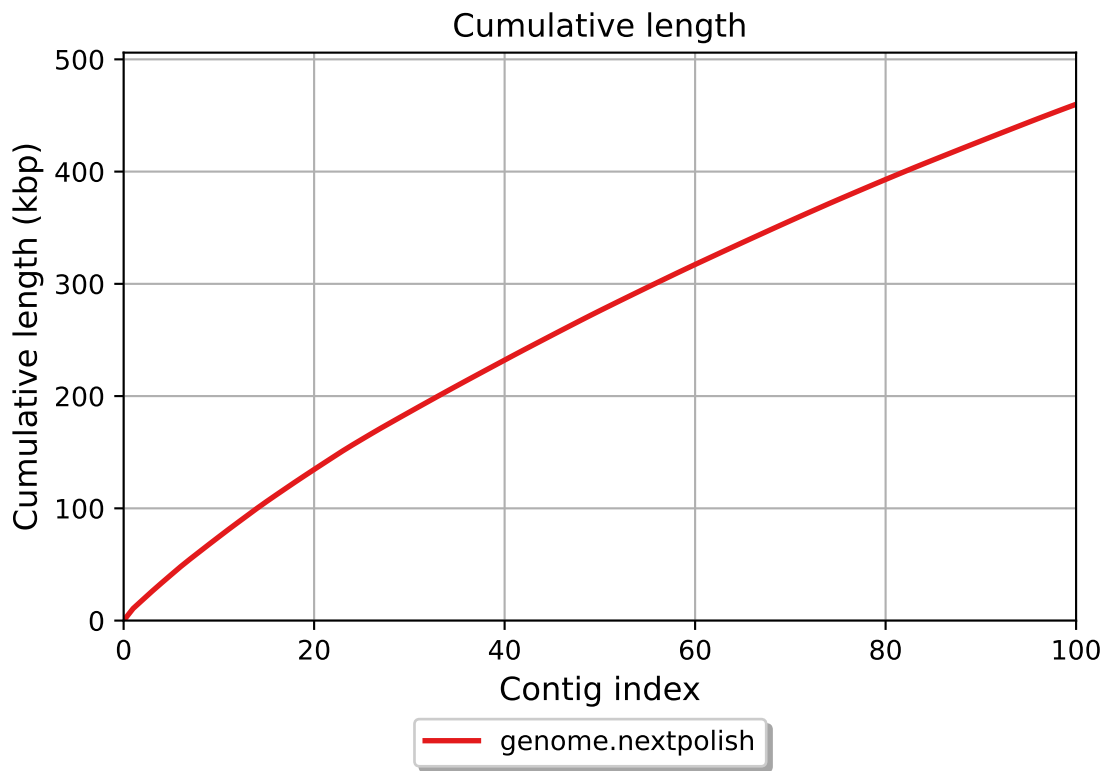
	genome.nextpolish
# contigs (>= 0 bp)	100
# contigs (>= 1000 bp)	100
# contigs (>= 5000 bp)	26
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	459982
Total length (>= 1000 bp)	459982
Total length (>= 5000 bp)	166444
Total length (>= 10000 bp)	10758
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	100
Largest contig	10758
Total length	459982
GC (%)	49.29
N50	4539
N90	3386
auN	4964.6
L50	40
L90	87
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 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).

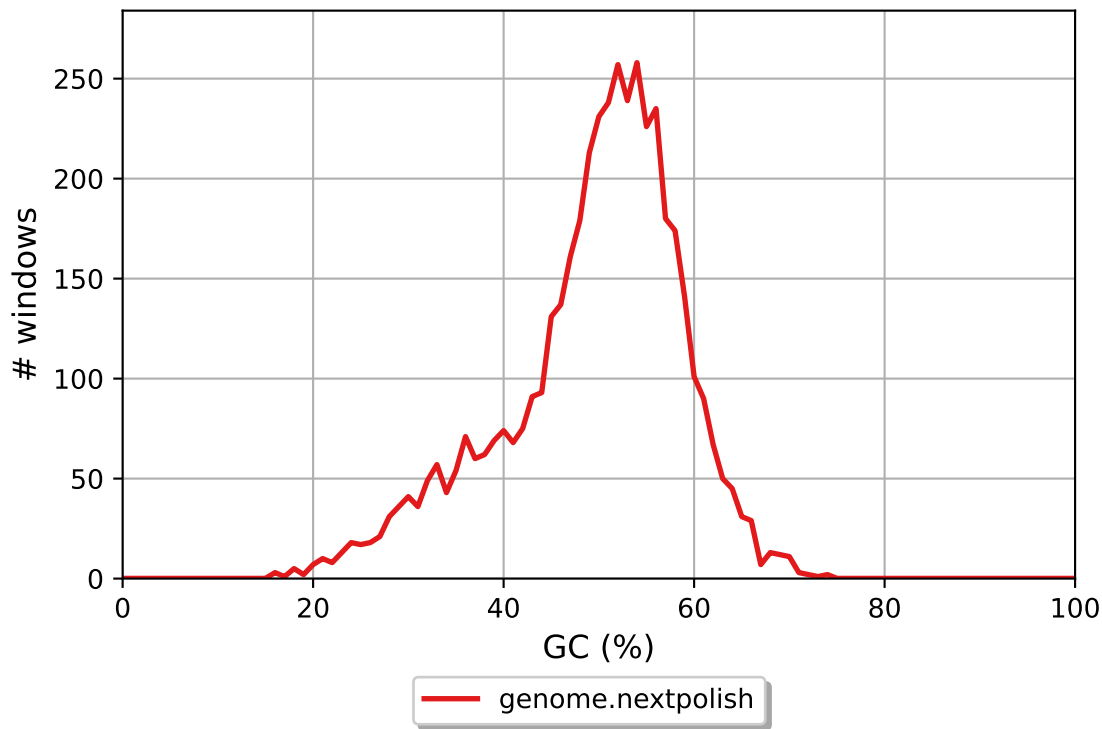
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

