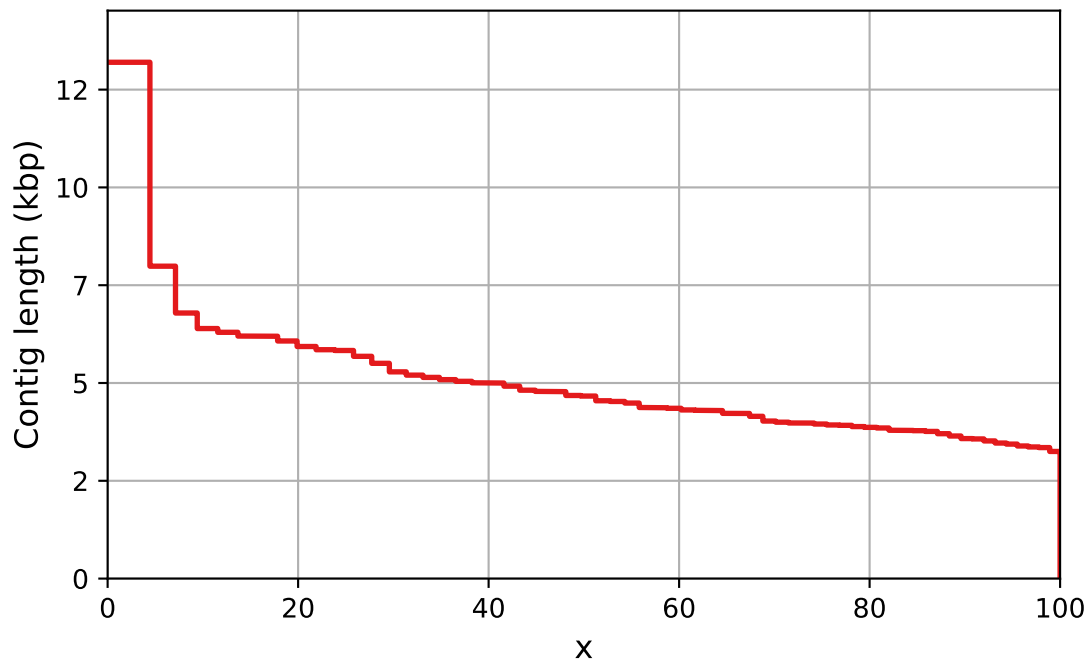


Report

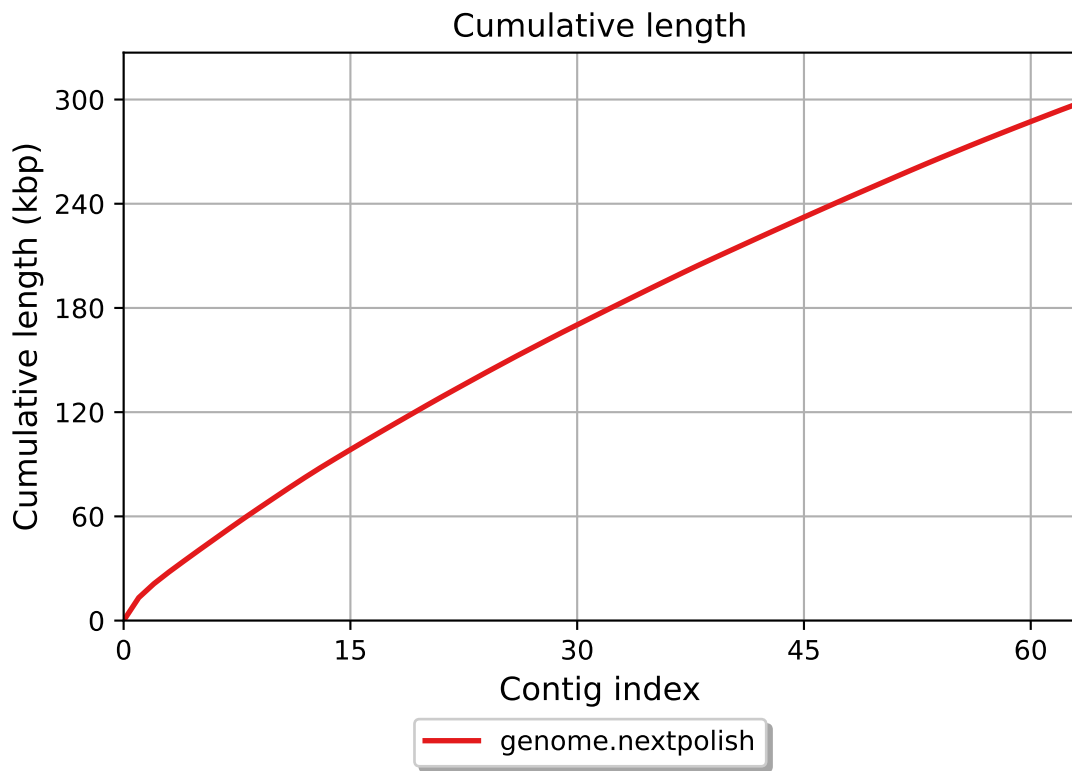
	genome.nextpolish
# contigs (>= 0 bp)	63
# contigs (>= 1000 bp)	63
# contigs (>= 5000 bp)	19
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	297269
Total length (>= 1000 bp)	297269
Total length (>= 5000 bp)	118694
Total length (>= 10000 bp)	13199
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	63
Largest contig	13199
Total length	297269
GC (%)	51.01
N50	4666
N90	3579
auN	5168.2
L50	26
L90	55
# 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).

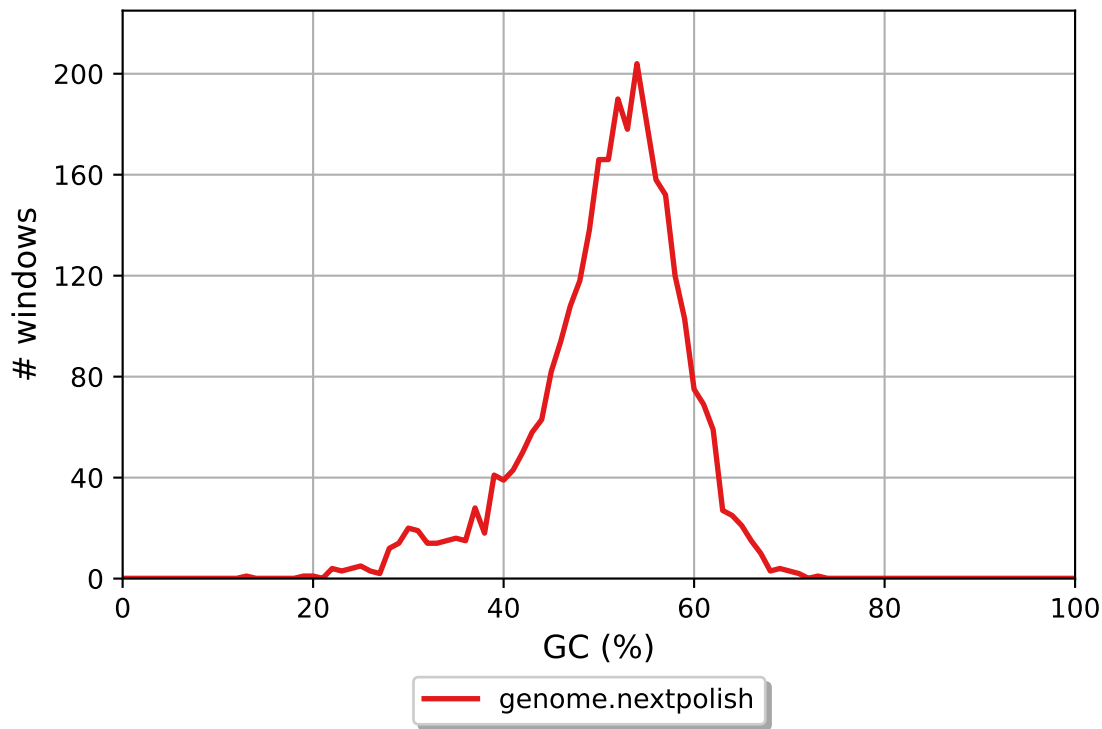
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

