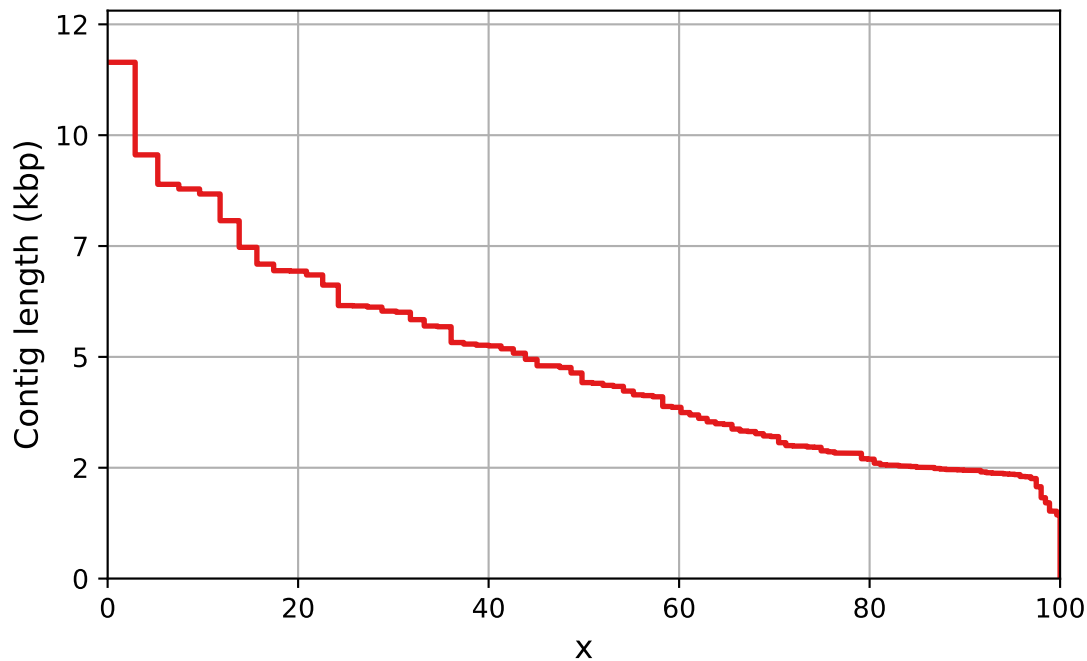


Report

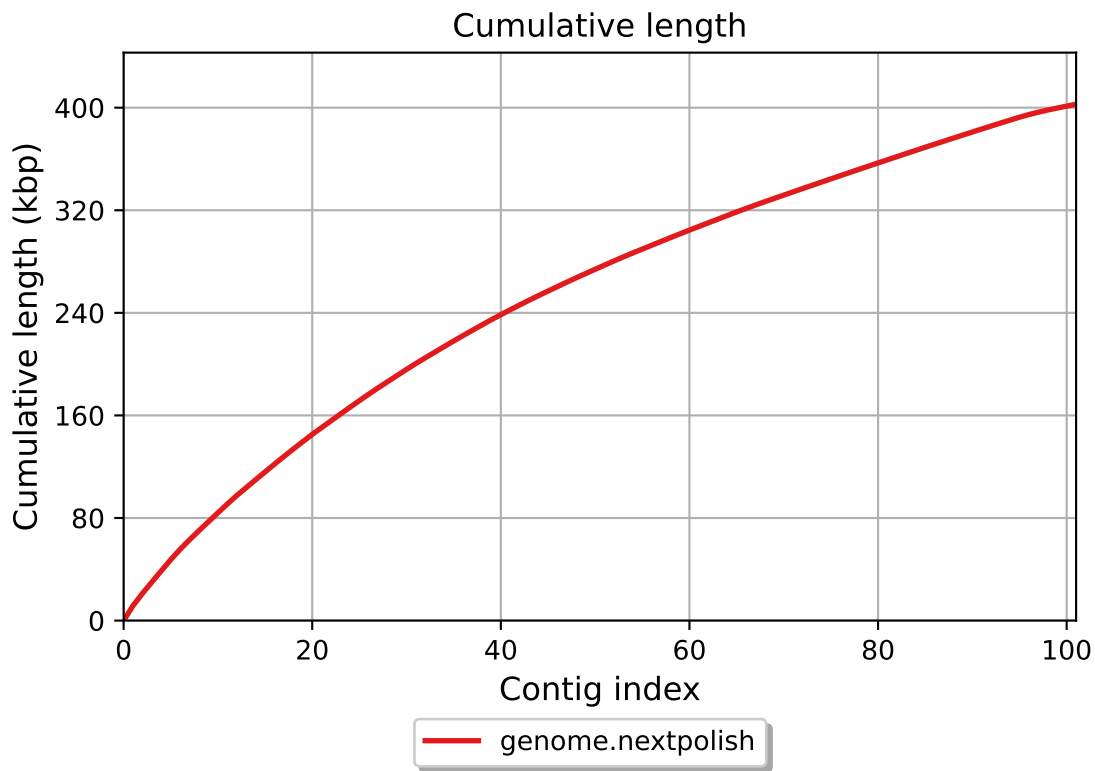
	genome.nextpolish
# contigs (>= 0 bp)	101
# contigs (>= 1000 bp)	101
# contigs (>= 5000 bp)	26
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	402659
Total length (>= 1000 bp)	402659
Total length (>= 5000 bp)	176606
Total length (>= 10000 bp)	11644
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	101
Largest contig	11644
Total length	402659
GC (%)	58.07
N50	4420
N90	2443
auN	4966.3
L50	32
L90	83
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 2 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).

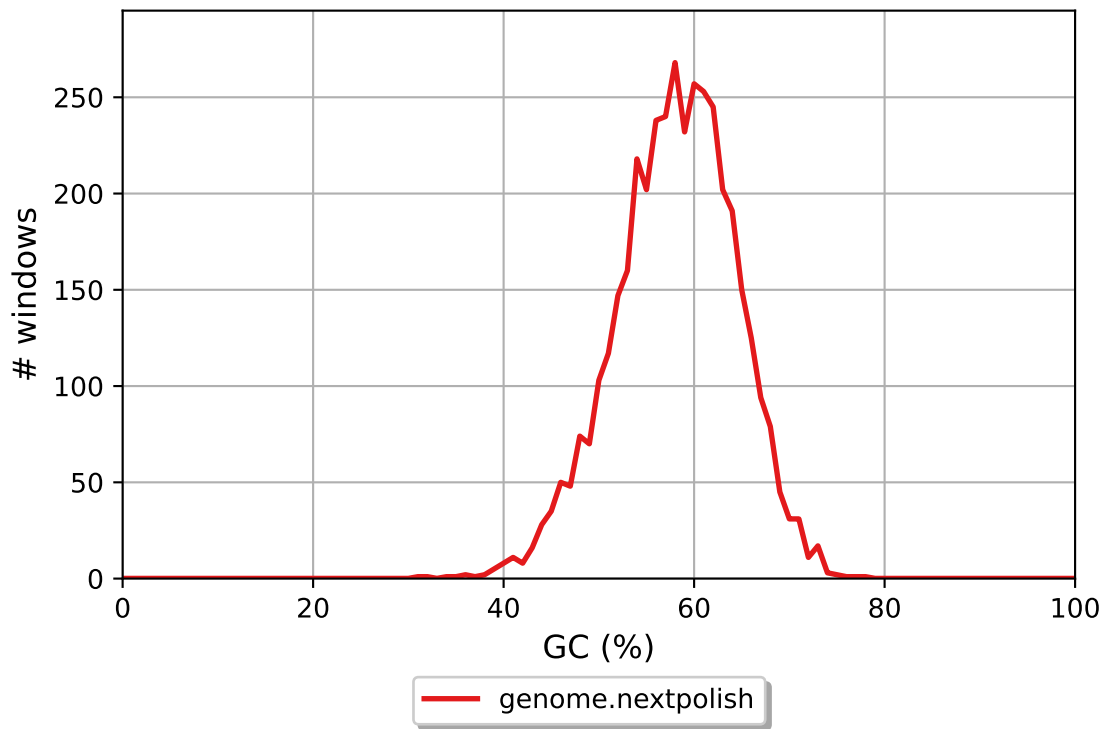
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

