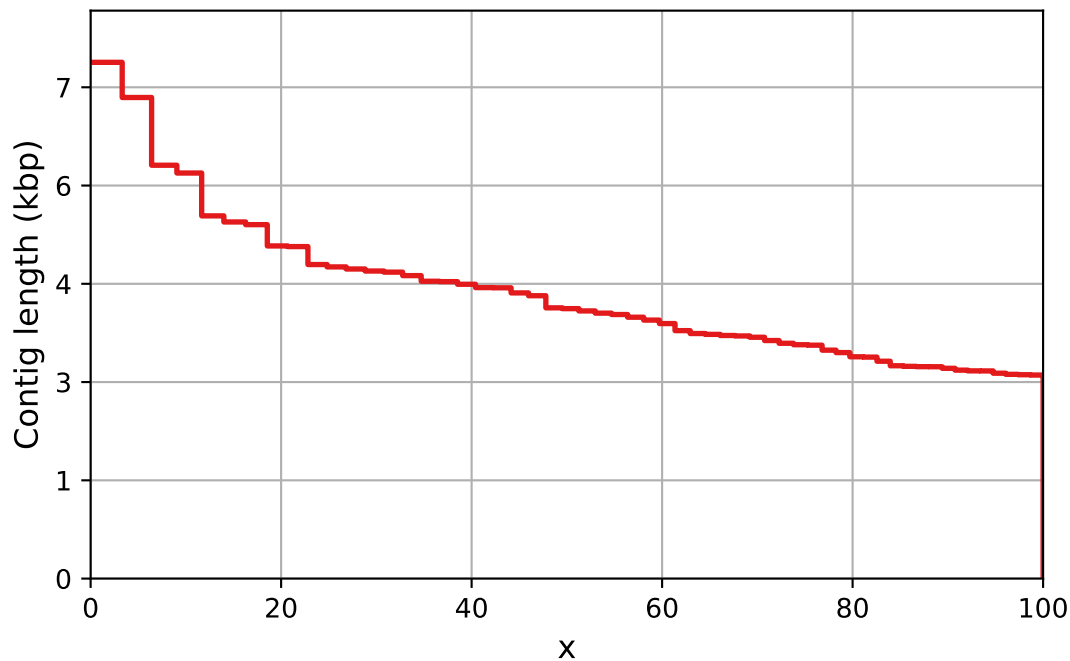


Report

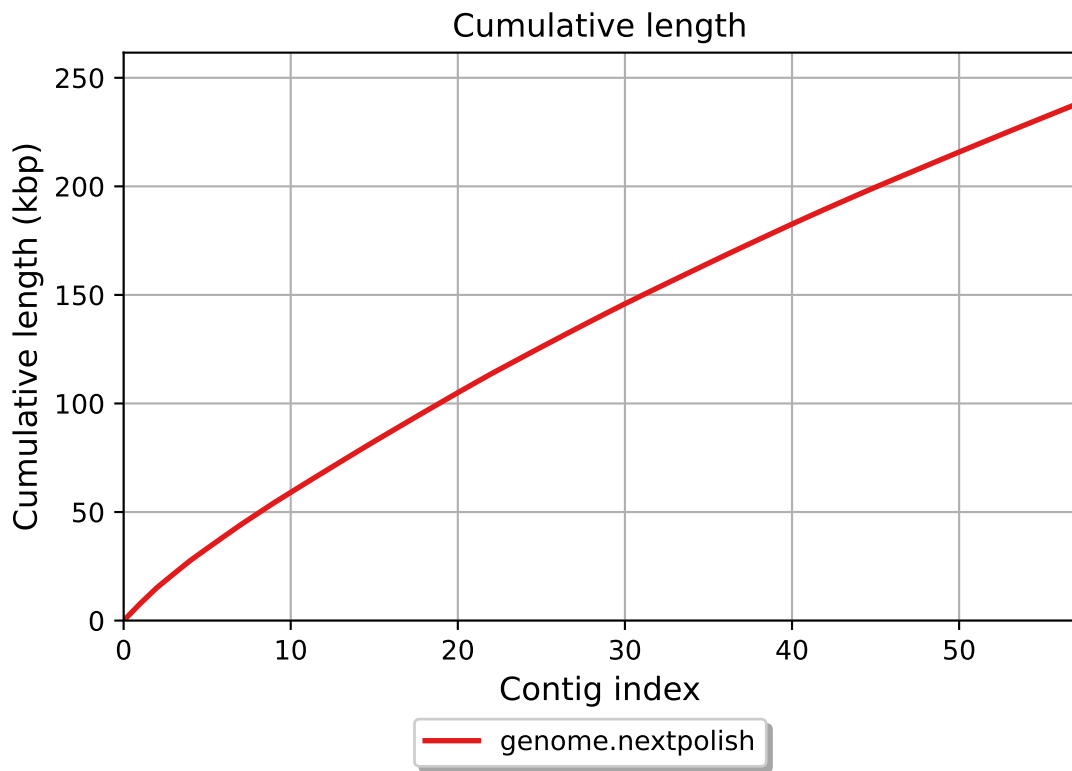
	genome.nextpolish
# contigs (>= 0 bp)	57
# contigs (>= 1000 bp)	57
# contigs (>= 5000 bp)	9
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	237806
Total length (>= 1000 bp)	237806
Total length (>= 5000 bp)	54256
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	57
Largest contig	7881
Total length	237806
GC (%)	55.76
N50	4121
N90	3210
auN	4420.1
L50	24
L90	50
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	0 + 1 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).

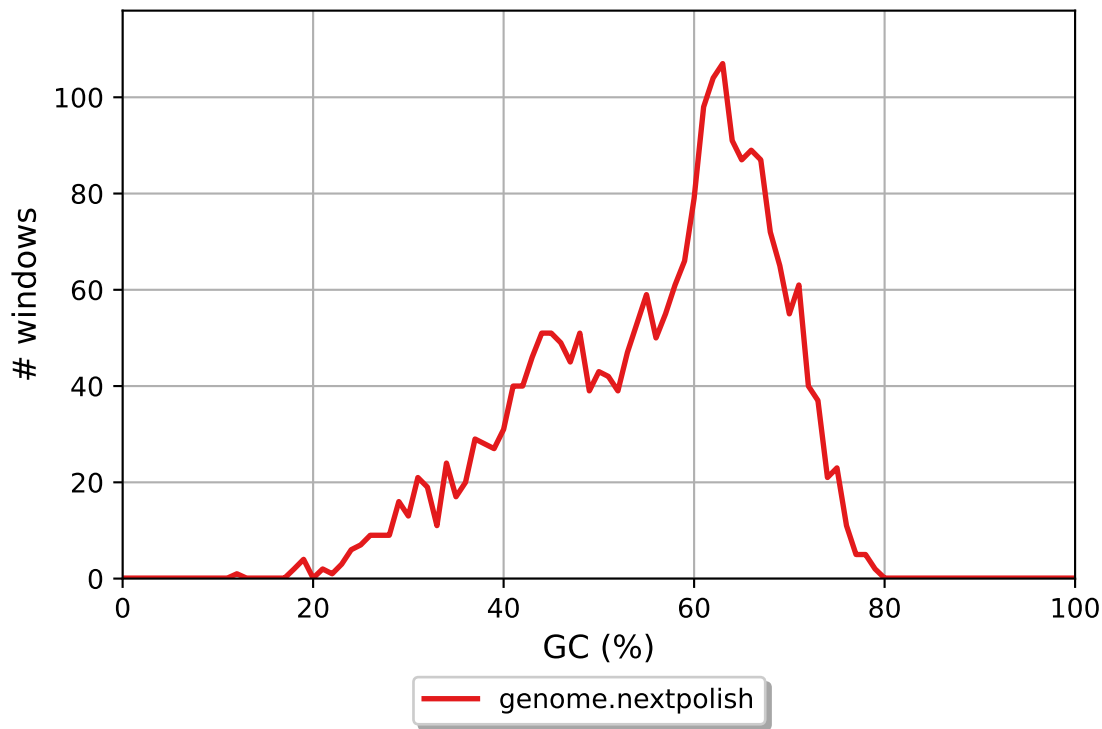
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

