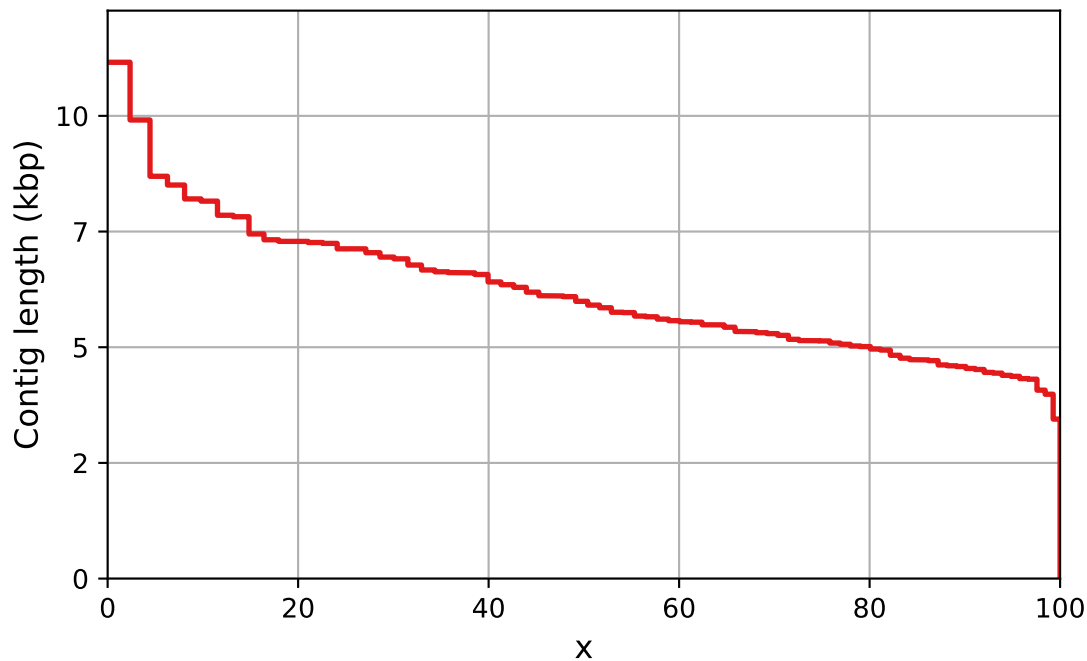


Report

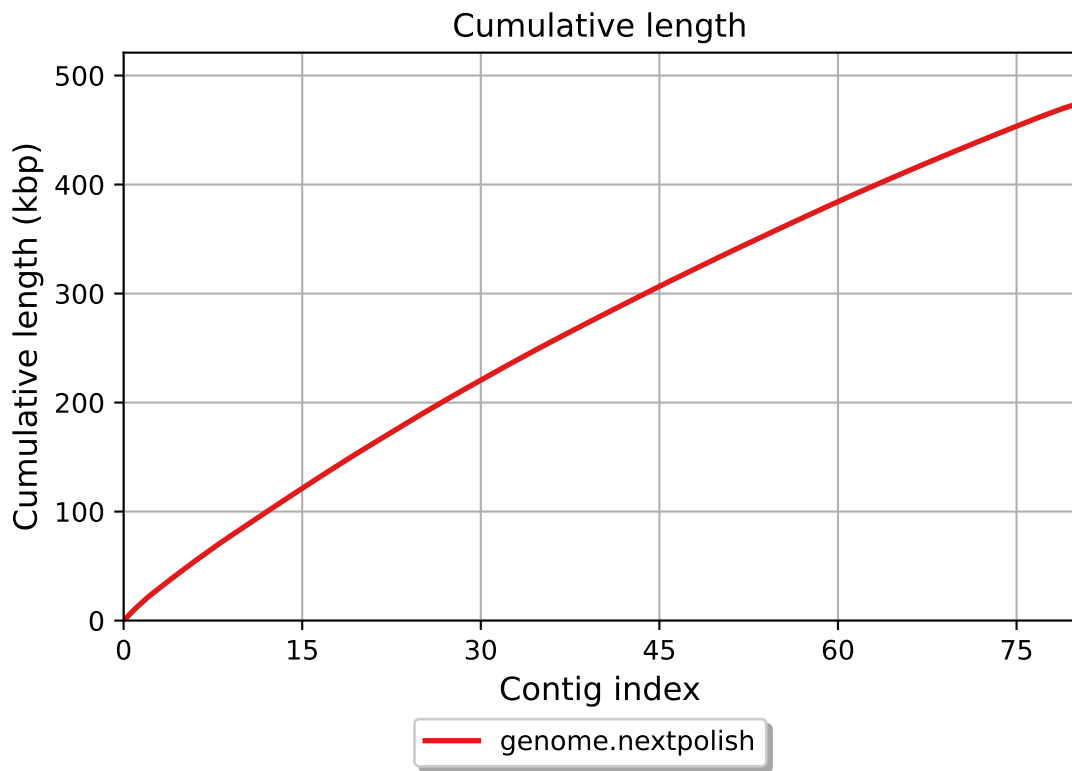
	genome.nextpolish
# contigs (>= 0 bp)	80
# contigs (>= 1000 bp)	80
# contigs (>= 5000 bp)	59
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	473632
Total length (>= 1000 bp)	473632
Total length (>= 5000 bp)	379324
Total length (>= 10000 bp)	11158
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	80
Largest contig	11158
Total length	473632
GC (%)	62.54
N50	5992
N90	4583
auN	6239.1
L50	33
L90	69
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 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).

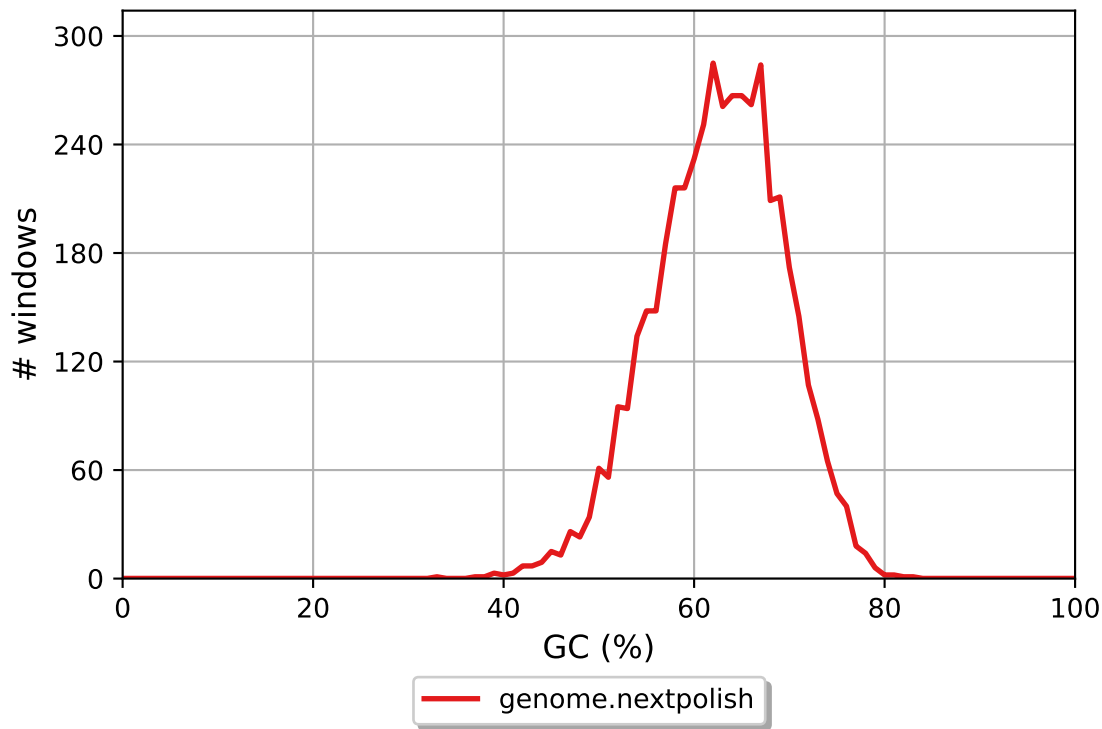
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

