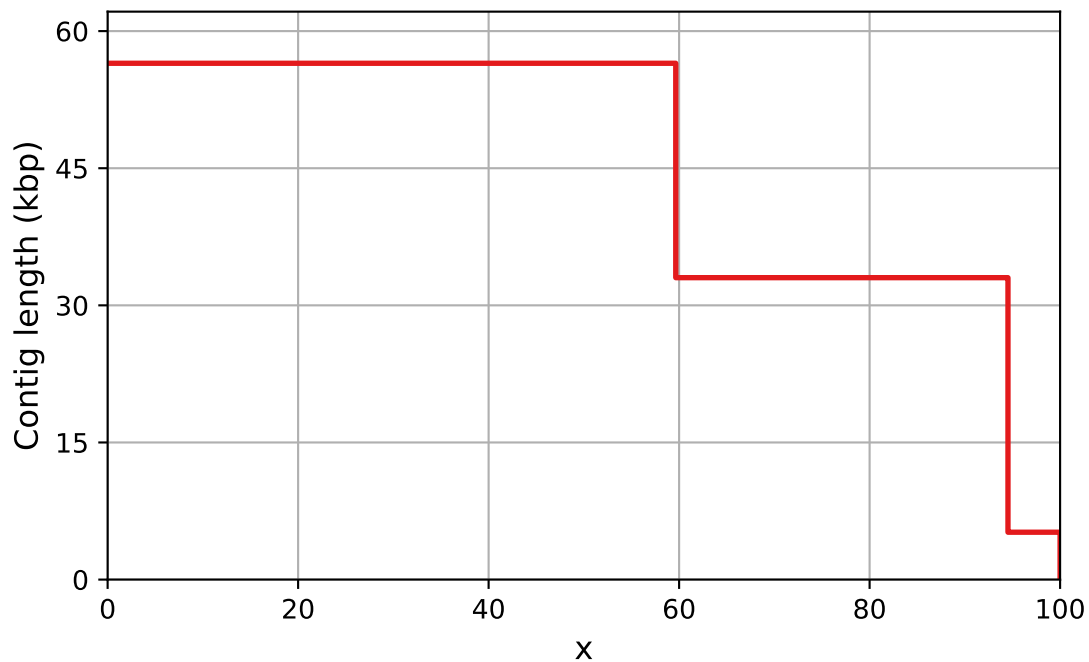


Report

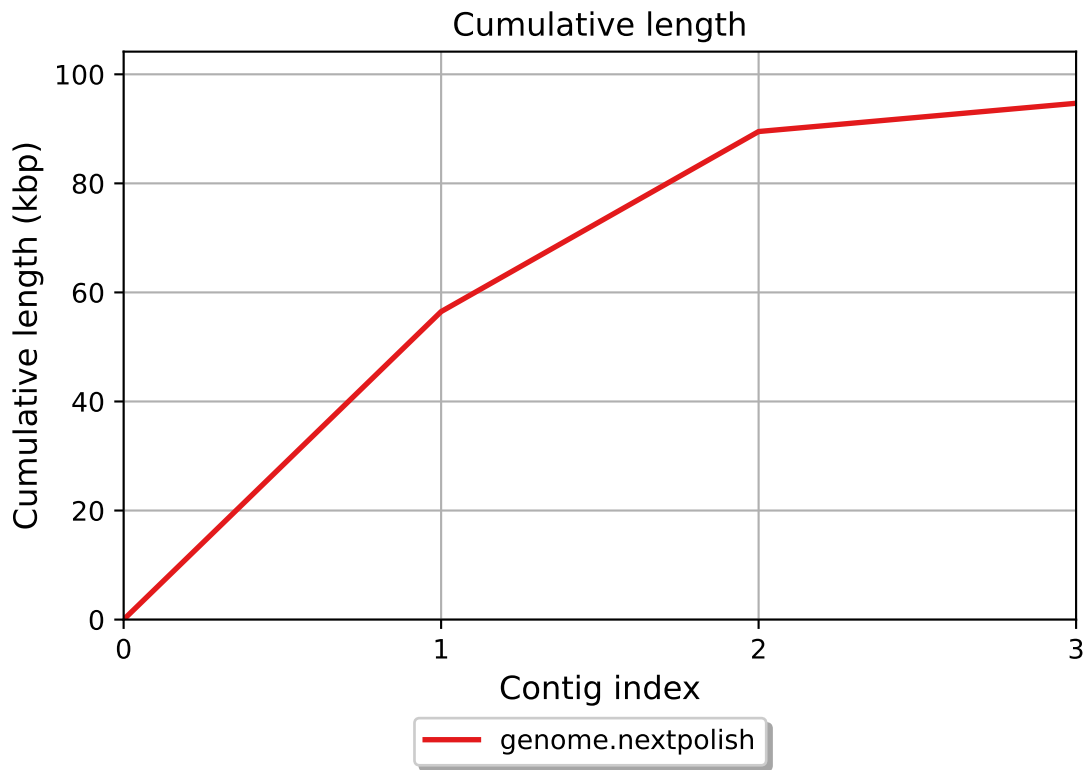
	genome.nextpolish
# contigs (>= 0 bp)	3
# contigs (>= 1000 bp)	3
# contigs (>= 5000 bp)	3
# contigs (>= 10000 bp)	2
# contigs (>= 25000 bp)	2
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	94689
Total length (>= 1000 bp)	94689
Total length (>= 5000 bp)	94689
Total length (>= 10000 bp)	89507
Total length (>= 25000 bp)	89507
Total length (>= 50000 bp)	56480
# contigs	3
Largest contig	56480
Total length	94689
GC (%)	59.82
N50	56480
N90	33027
auN	45492.4
L50	1
L90	2
# 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).

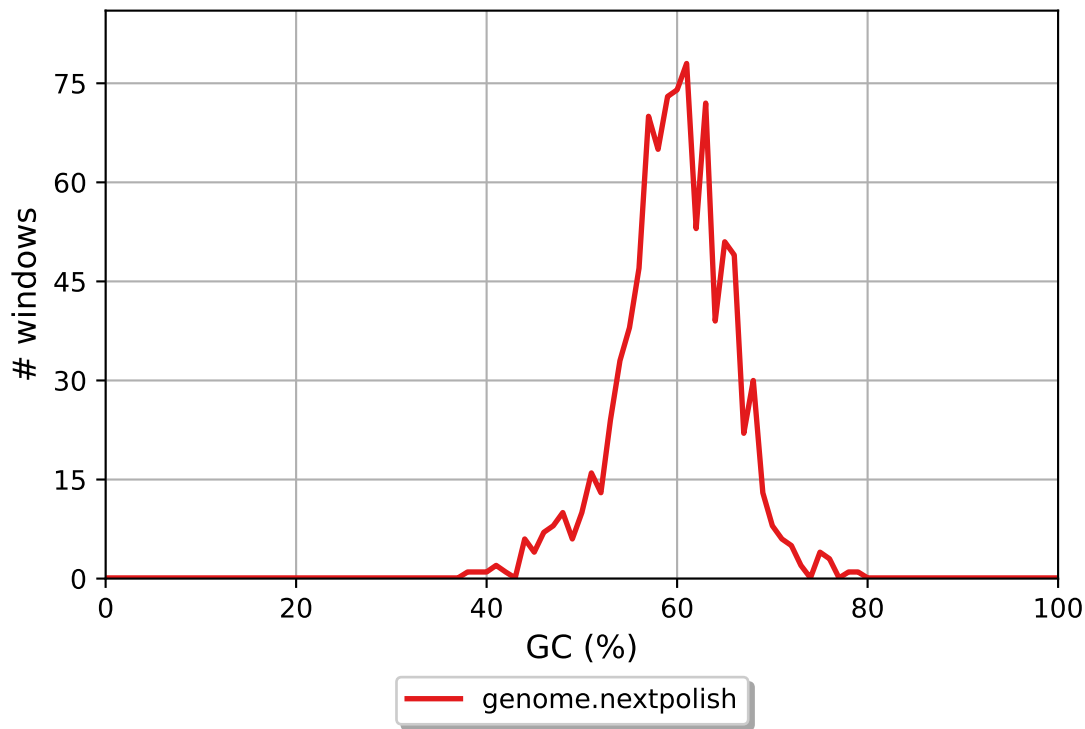
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

