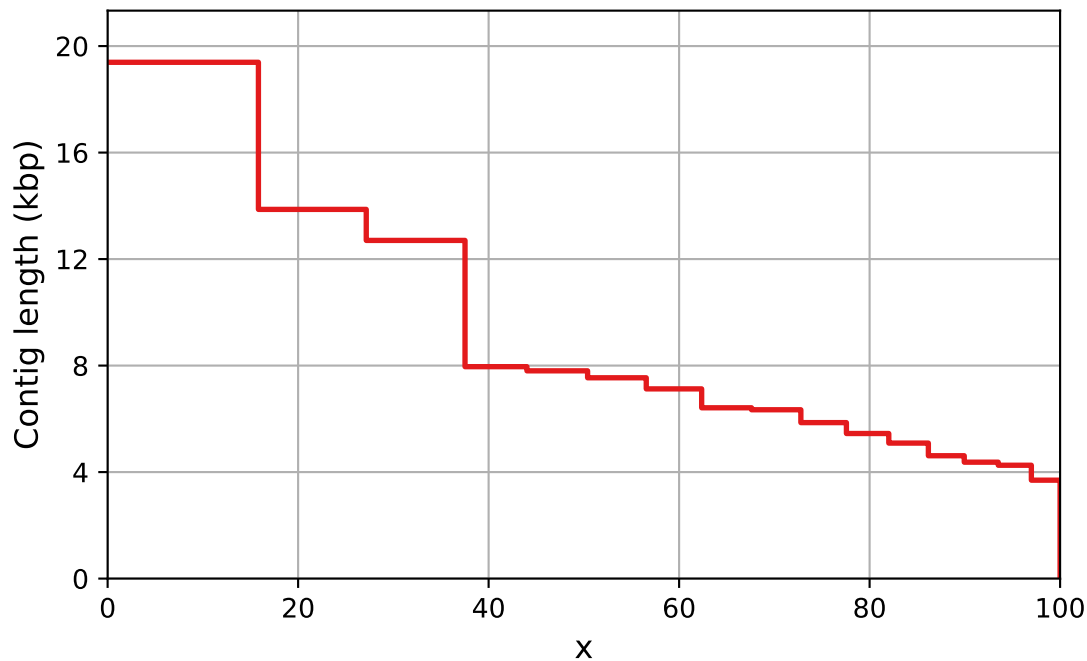


Report

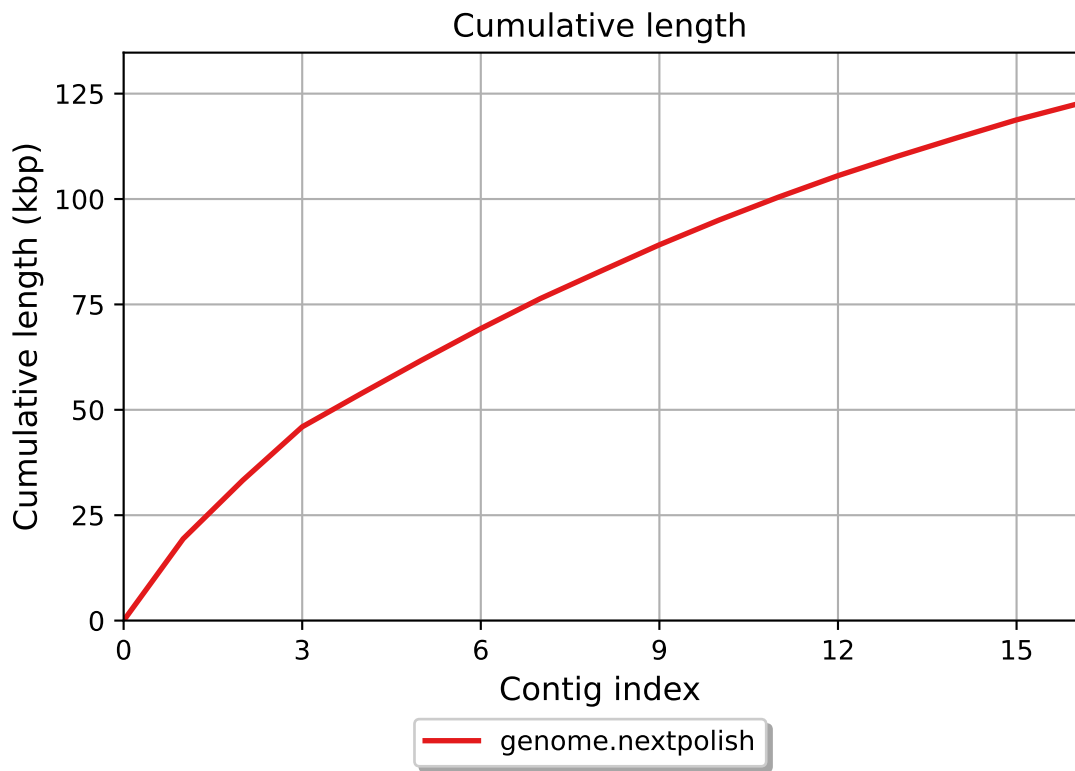
	genome.nextpolish
# contigs (>= 0 bp)	16
# contigs (>= 1000 bp)	16
# contigs (>= 5000 bp)	12
# contigs (>= 10000 bp)	3
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	122478
Total length (>= 1000 bp)	122478
Total length (>= 5000 bp)	105539
Total length (>= 10000 bp)	45958
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	16
Largest contig	19391
Total length	122478
GC (%)	57.93
N50	7803
N90	4372
auN	9837.9
L50	5
L90	14
# 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).

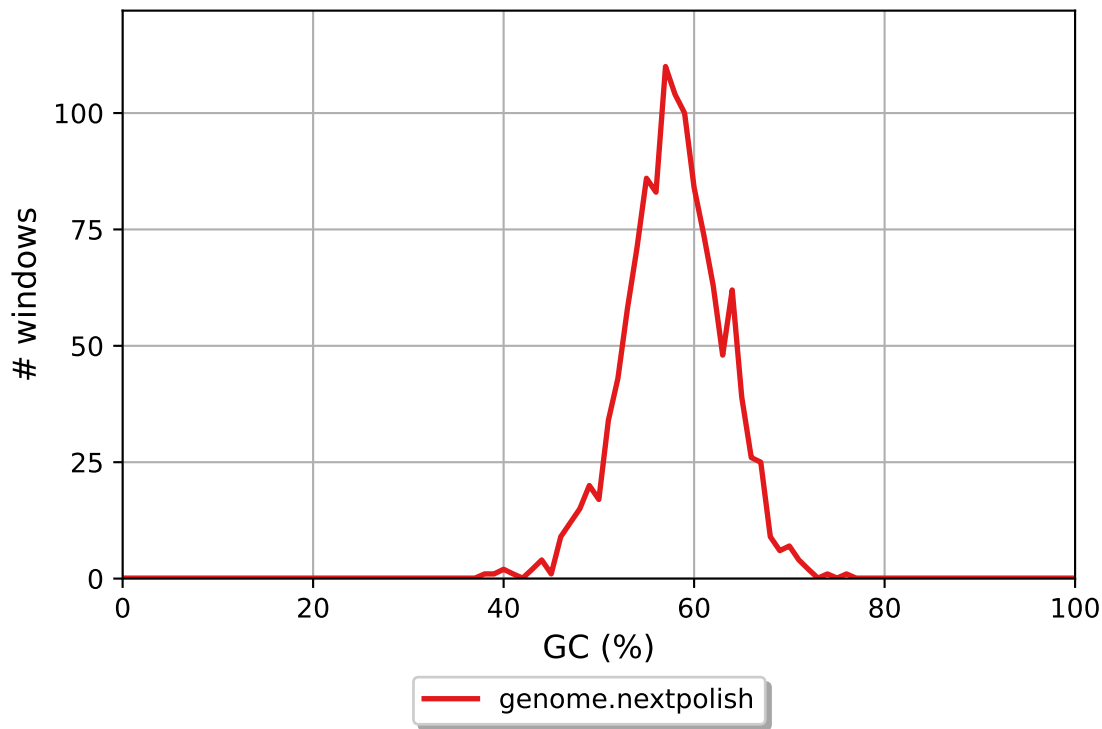
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

