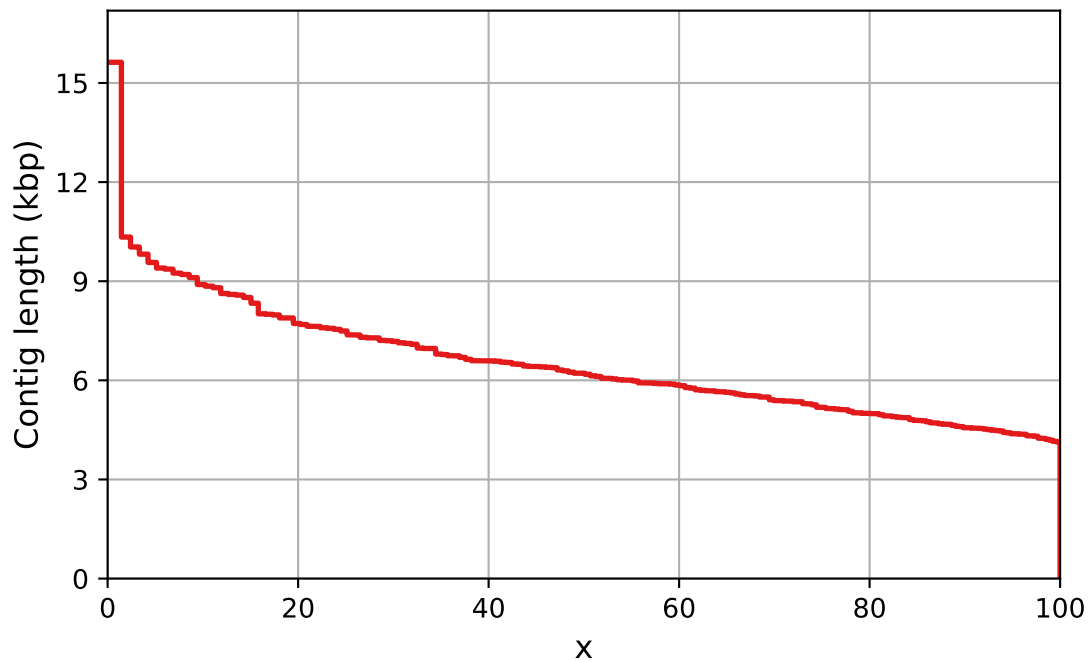


Report

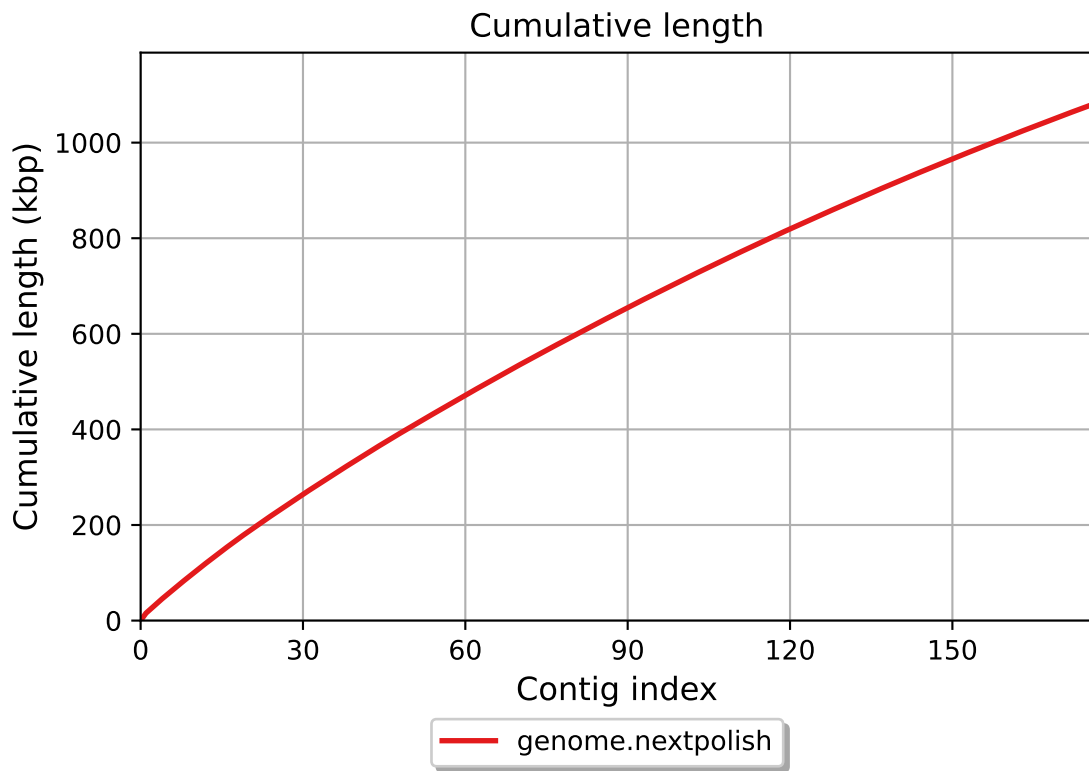
	genome.nextpolish
# contigs (>= 0 bp)	176
# contigs (>= 1000 bp)	176
# contigs (>= 5000 bp)	129
# contigs (>= 10000 bp)	3
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1080268
Total length (>= 1000 bp)	1080268
Total length (>= 5000 bp)	865035
Total length (>= 10000 bp)	35999
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	176
Largest contig	15626
Total length	1080268
GC (%)	48.10
N50	6216
N90	4566
auN	6547.8
L50	71
L90	152
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 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).

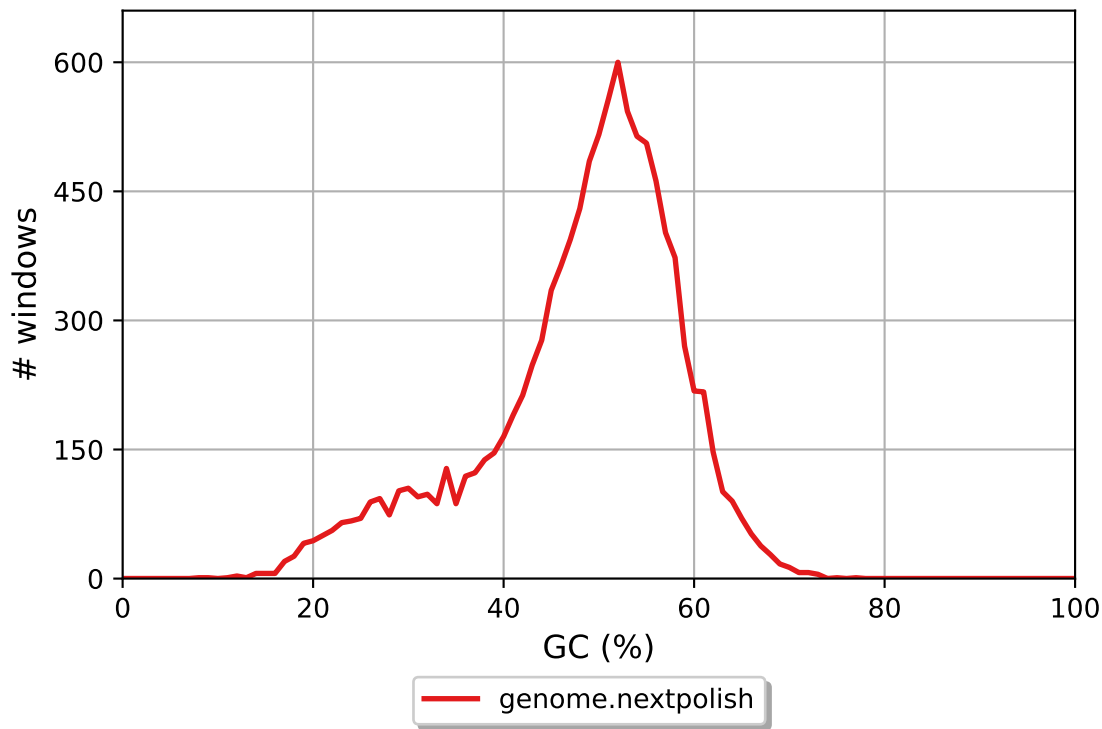
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

