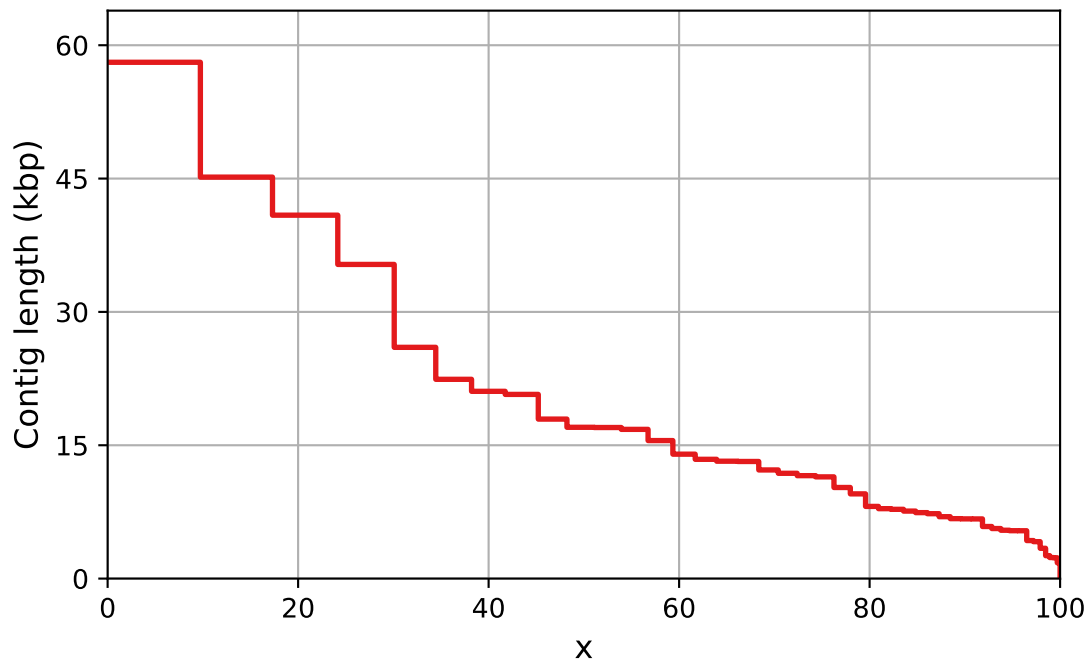


Report

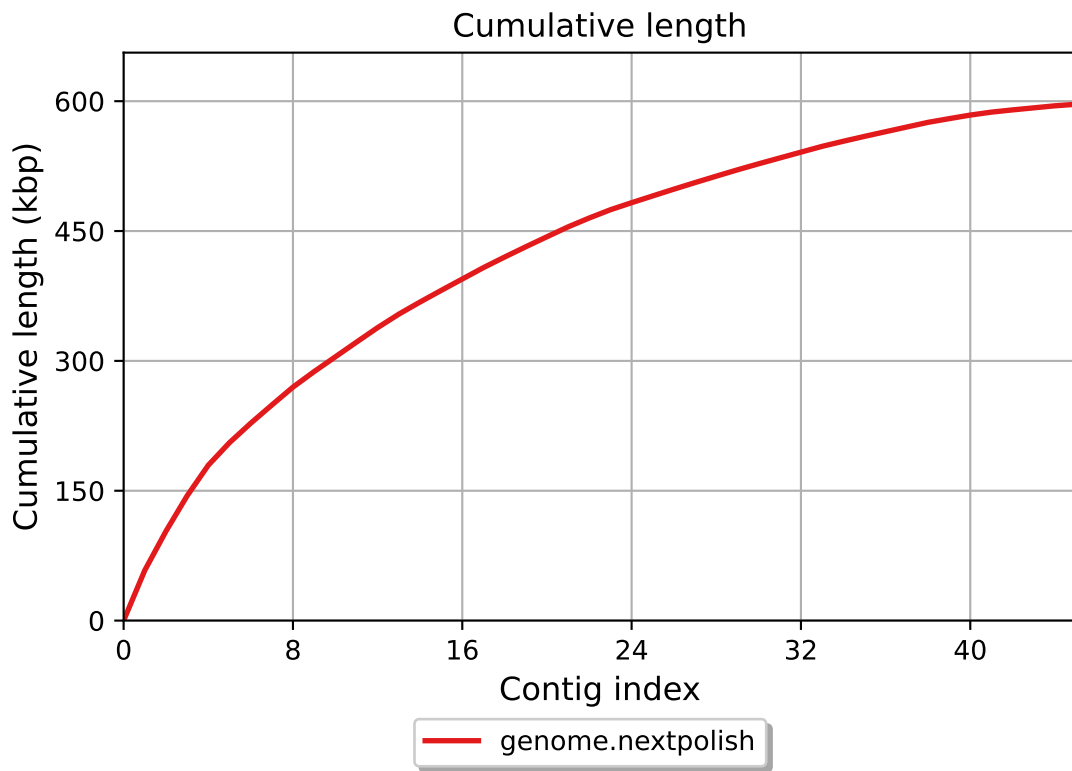
	genome.nextpolish
# contigs (>= 0 bp)	45
# contigs (>= 1000 bp)	45
# contigs (>= 5000 bp)	38
# contigs (>= 10000 bp)	22
# contigs (>= 25000 bp)	5
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	596441
Total length (>= 1000 bp)	596441
Total length (>= 5000 bp)	575532
Total length (>= 10000 bp)	465081
Total length (>= 25000 bp)	205480
Total length (>= 50000 bp)	58082
# contigs	45
Largest contig	58082
Total length	596441
GC (%)	52.16
N50	17020
N90	6718
auN	23545.8
L50	10
L90	32
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 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).

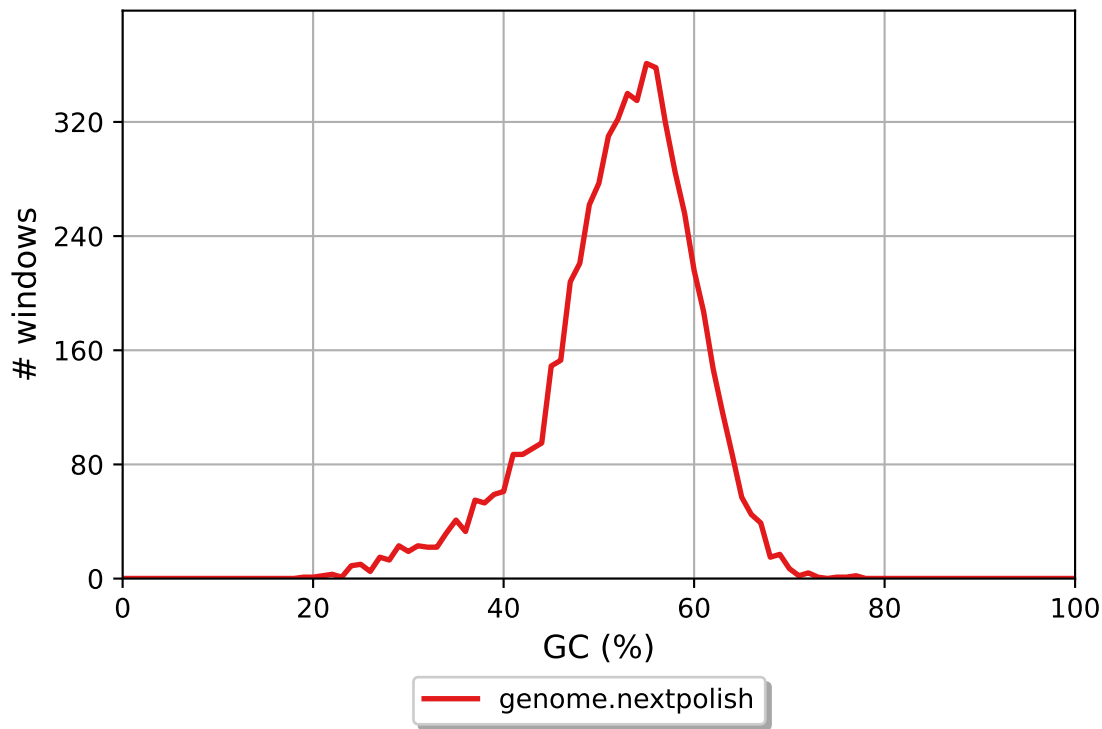
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

