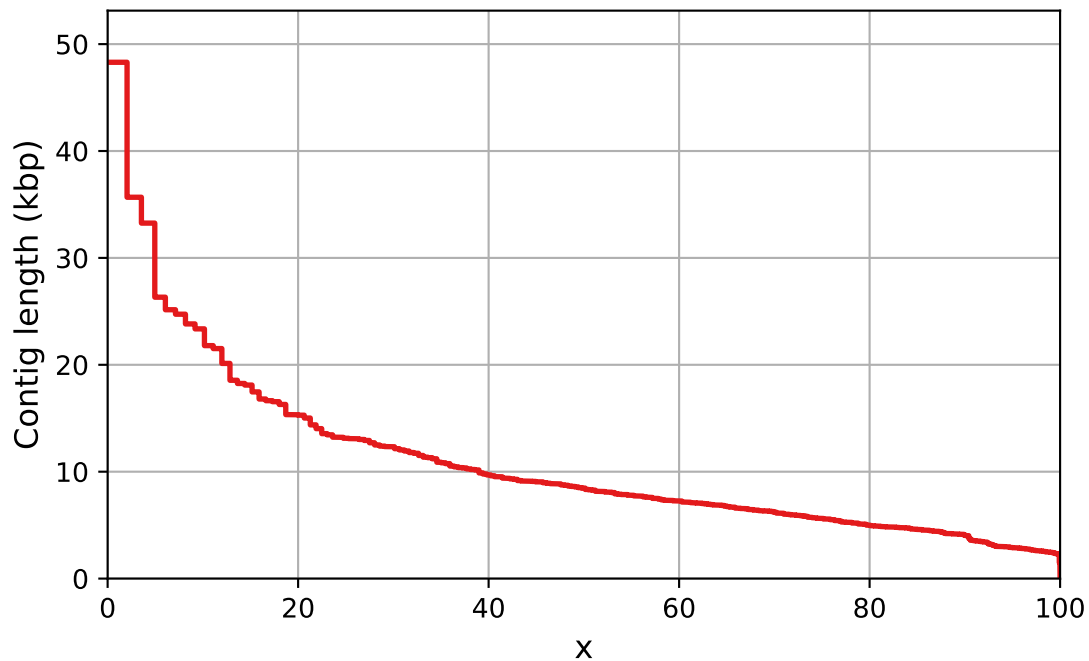


Report

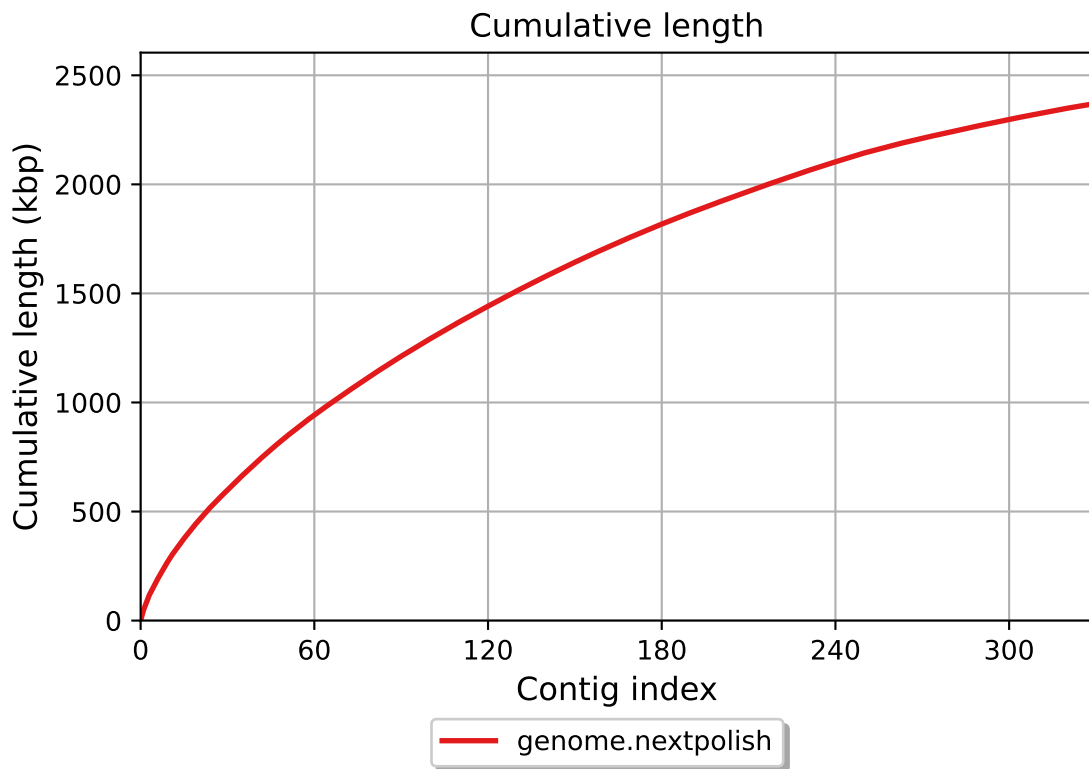
	genome.nextpolish
# contigs (>= 0 bp)	329
# contigs (>= 1000 bp)	328
# contigs (>= 5000 bp)	194
# contigs (>= 10000 bp)	58
# contigs (>= 25000 bp)	5
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2367212
Total length (>= 1000 bp)	2366708
Total length (>= 5000 bp)	1890578
Total length (>= 10000 bp)	922923
Total length (>= 25000 bp)	168705
Total length (>= 50000 bp)	0
# contigs	329
Largest contig	48301
Total length	2367212
GC (%)	33.22
N50	8466
N90	4063
auN	11171.2
L50	87
L90	247
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	1 + 4 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).

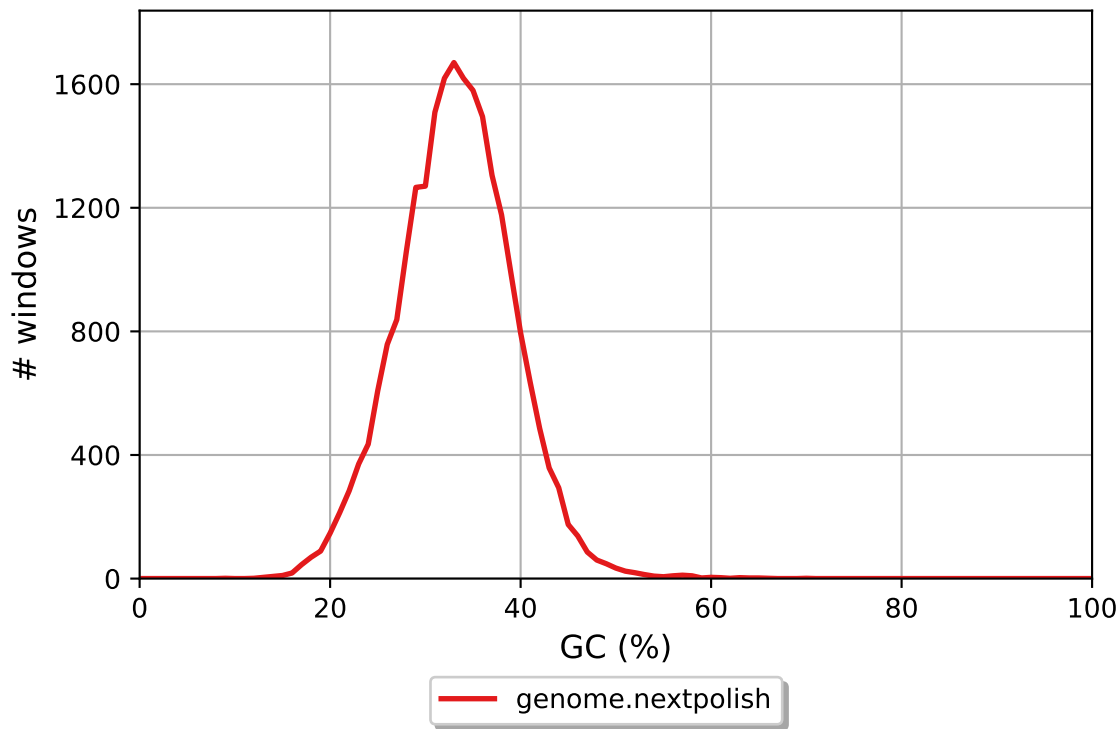
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

