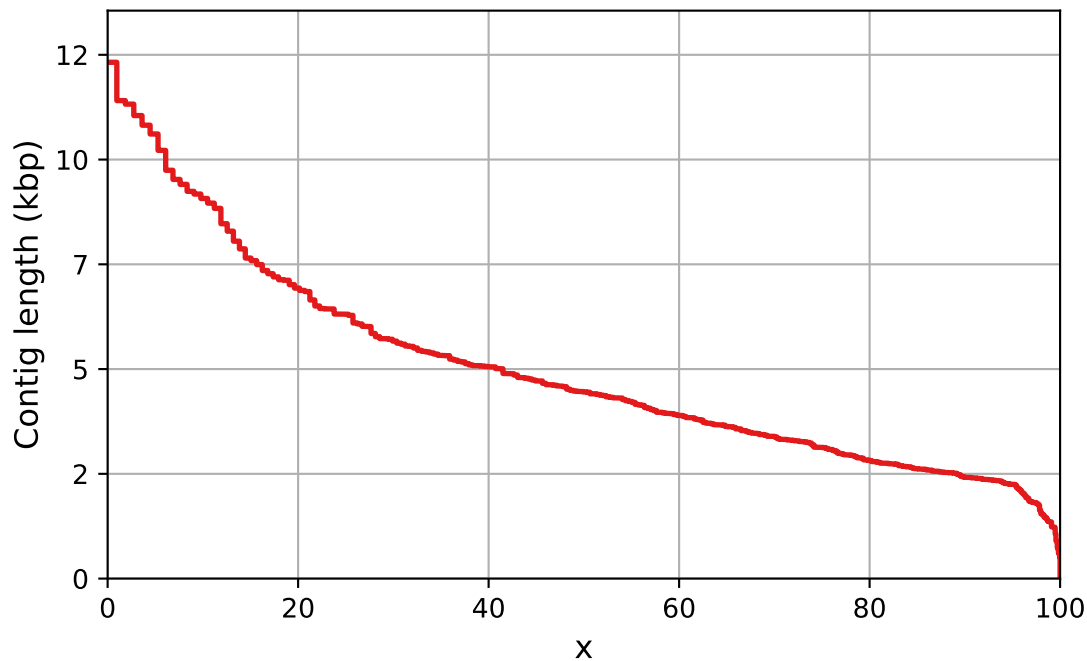


Report

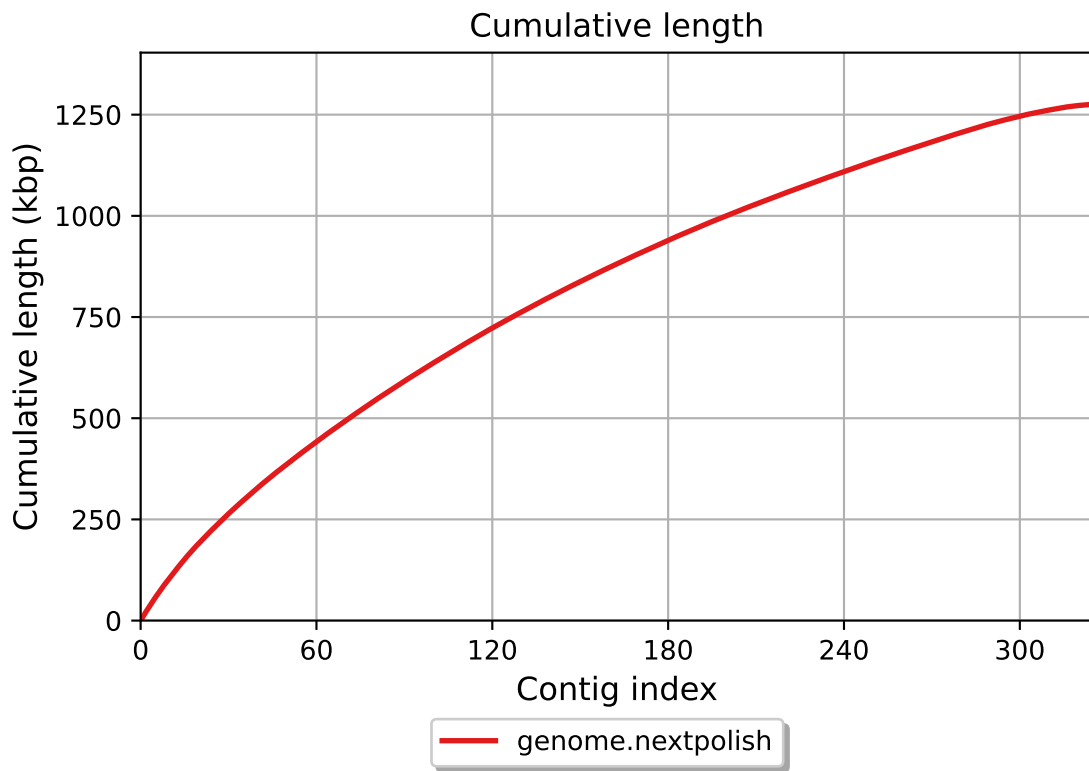
	genome.nextpolish
# contigs (>= 0 bp)	326
# contigs (>= 1000 bp)	317
# contigs (>= 5000 bp)	77
# contigs (>= 10000 bp)	7
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1276114
Total length (>= 1000 bp)	1270121
Total length (>= 5000 bp)	529491
Total length (>= 10000 bp)	77750
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	325
Largest contig	12321
Total length	1275667
GC (%)	35.68
N50	4455
N90	2417
auN	5018.0
L50	101
L90	256
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	1 + 2 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).

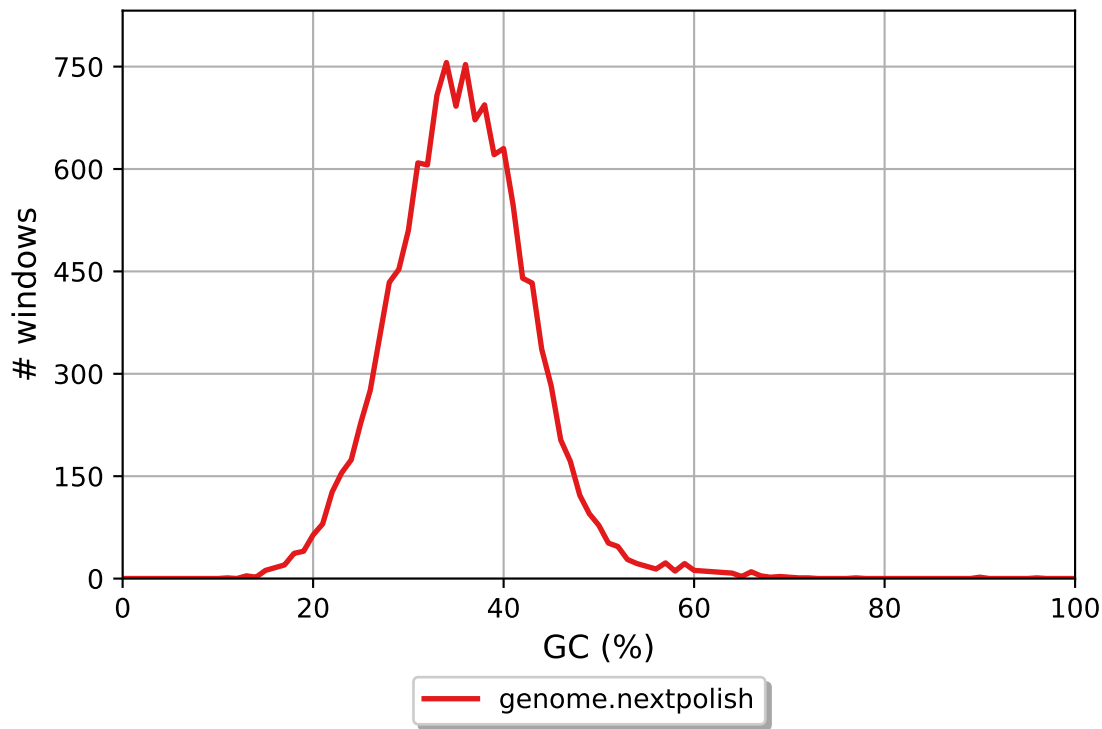
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

