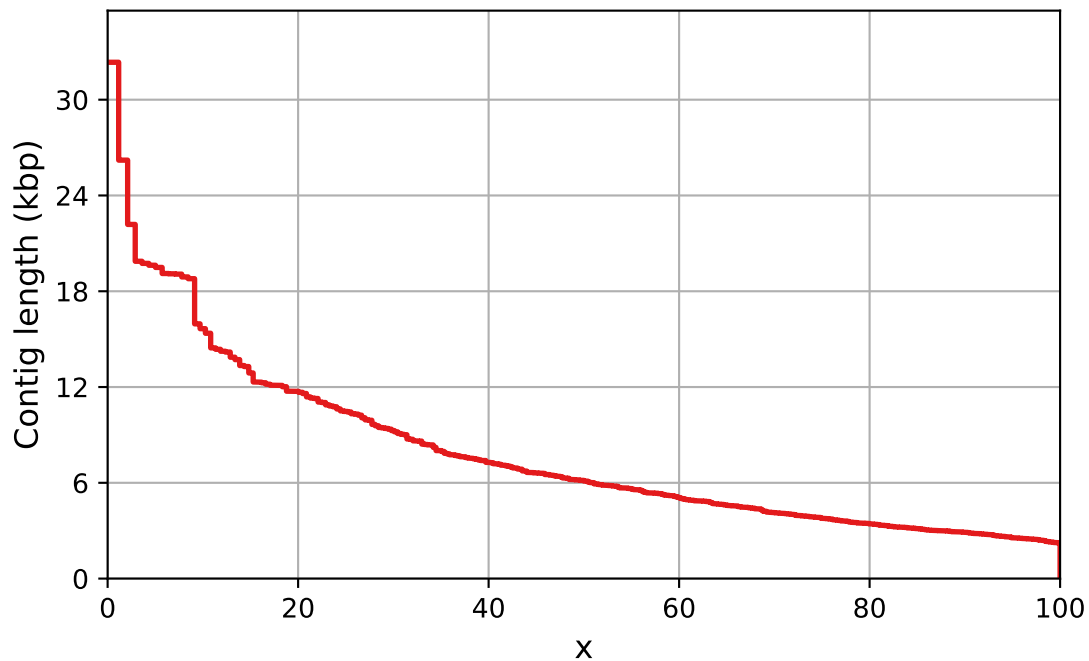


Report

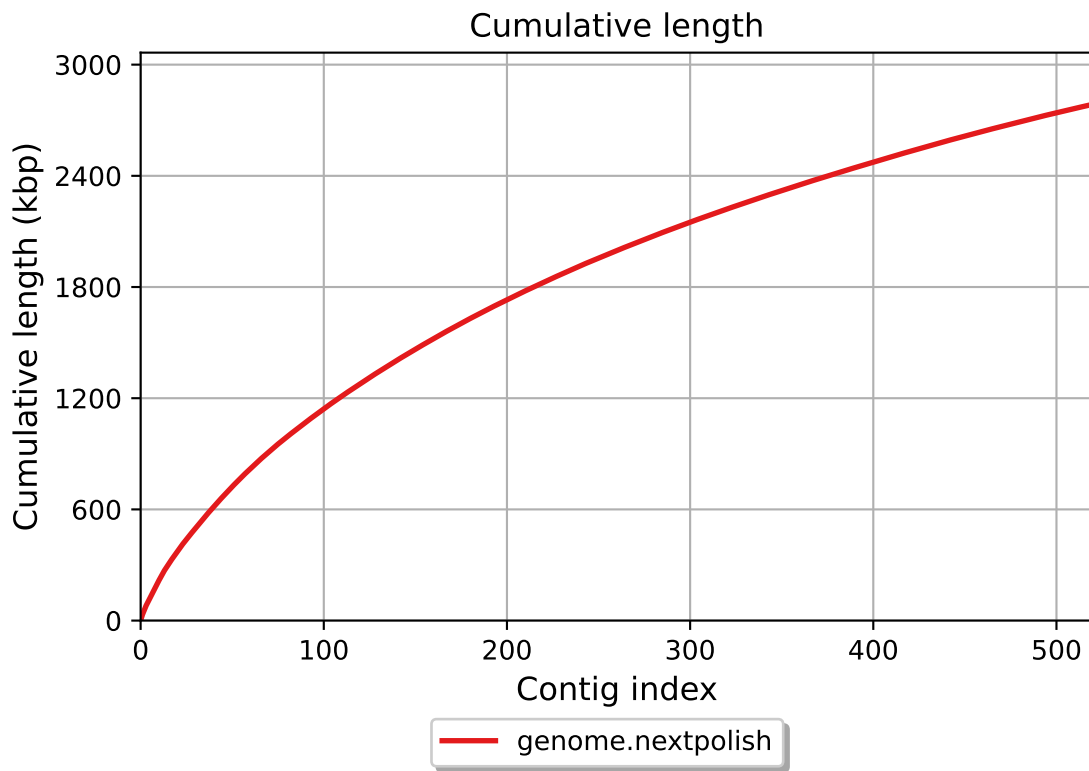
	genome.nextpolish
# contigs (>= 0 bp)	520
# contigs (>= 1000 bp)	520
# contigs (>= 5000 bp)	190
# contigs (>= 10000 bp)	53
# contigs (>= 25000 bp)	2
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2786155
Total length (>= 1000 bp)	2786155
Total length (>= 5000 bp)	1681950
Total length (>= 10000 bp)	753146
Total length (>= 25000 bp)	58557
Total length (>= 50000 bp)	0
# contigs	520
Largest contig	32344
Total length	2786155
GC (%)	35.41
N50	6113
N90	2899
auN	7949.6
L50	139
L90	412
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 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).

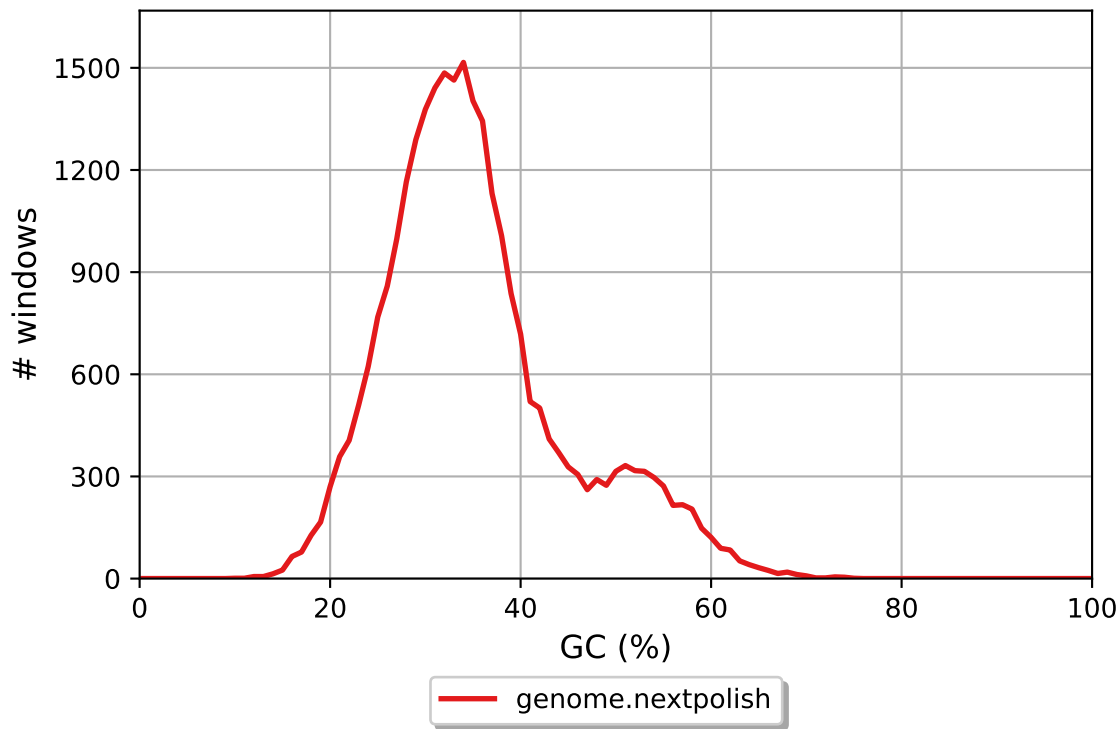
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

