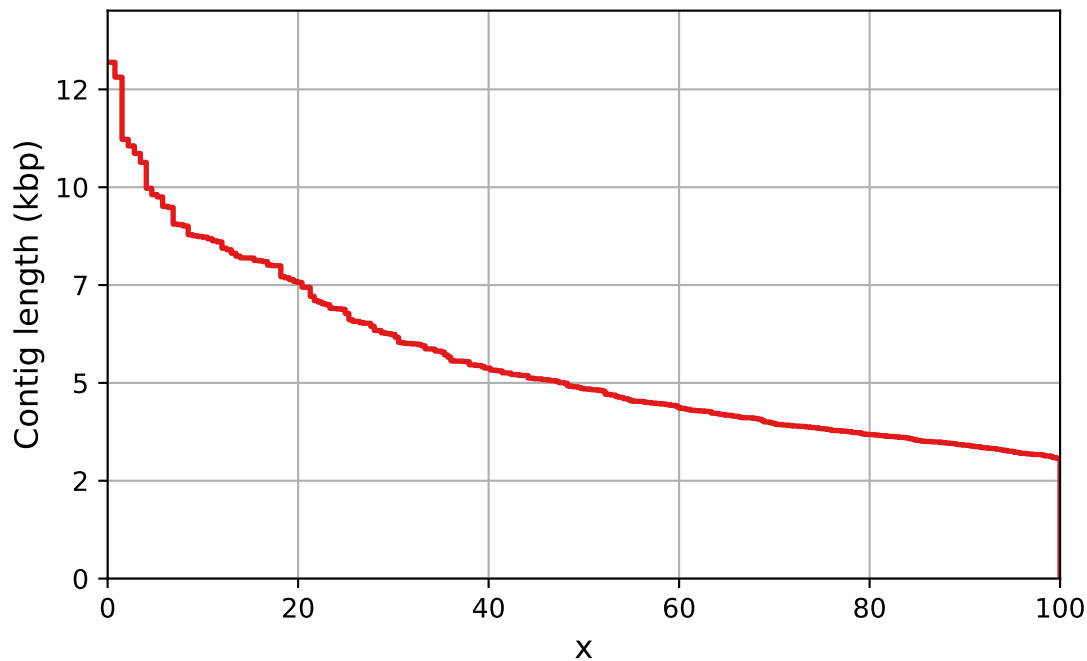


Report

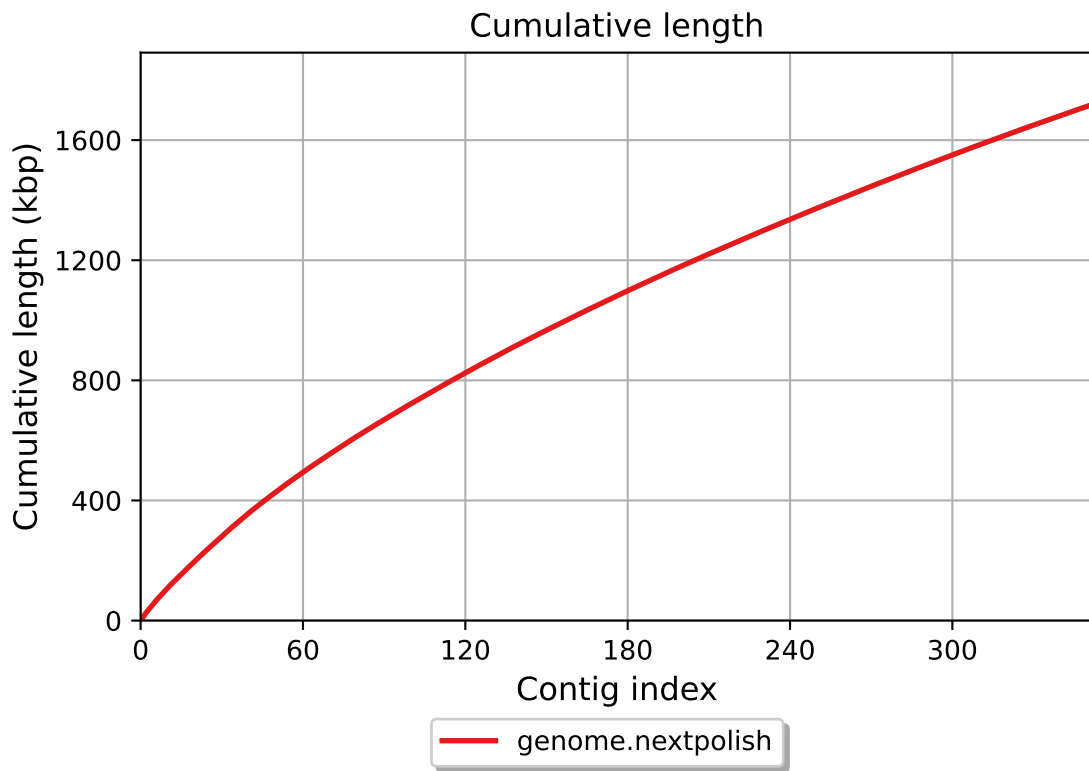
	genome.nextpolish
# contigs (>= 0 bp)	352
# contigs (>= 1000 bp)	352
# contigs (>= 5000 bp)	120
# contigs (>= 10000 bp)	6
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1719394
Total length (>= 1000 bp)	1719394
Total length (>= 5000 bp)	825065
Total length (>= 10000 bp)	69783
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	352
Largest contig	13192
Total length	1719394
GC (%)	34.60
N50	4847
N90	3410
auN	5551.8
L50	128
L90	300
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 3 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).

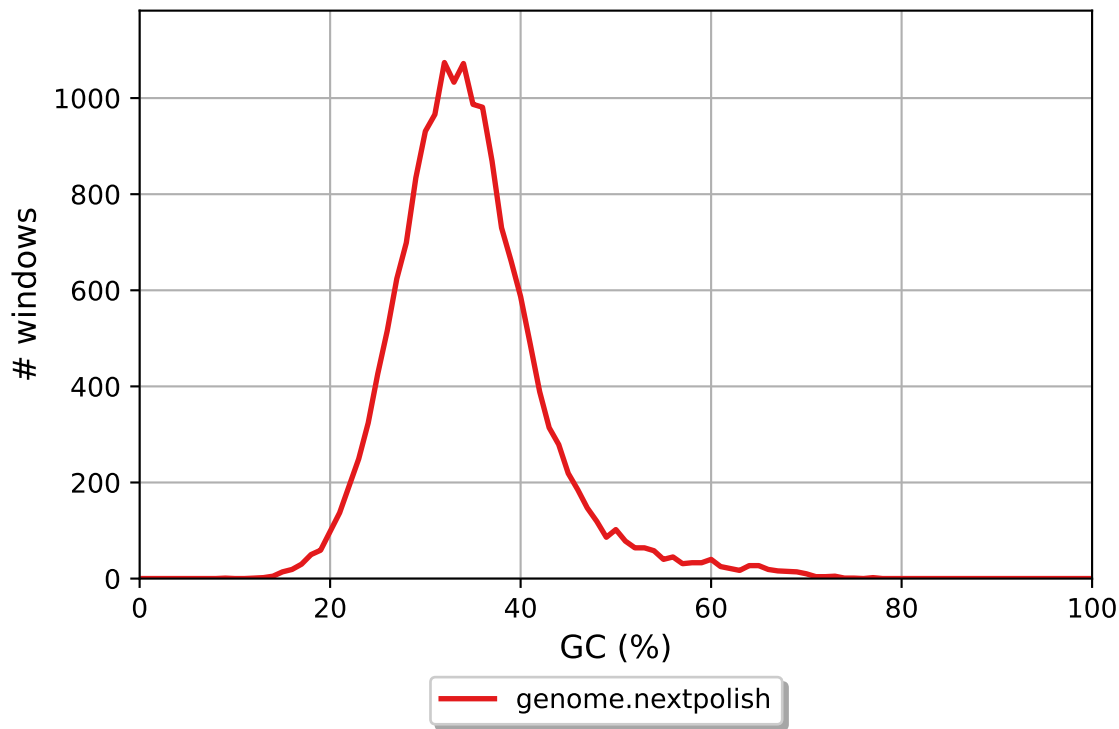
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

