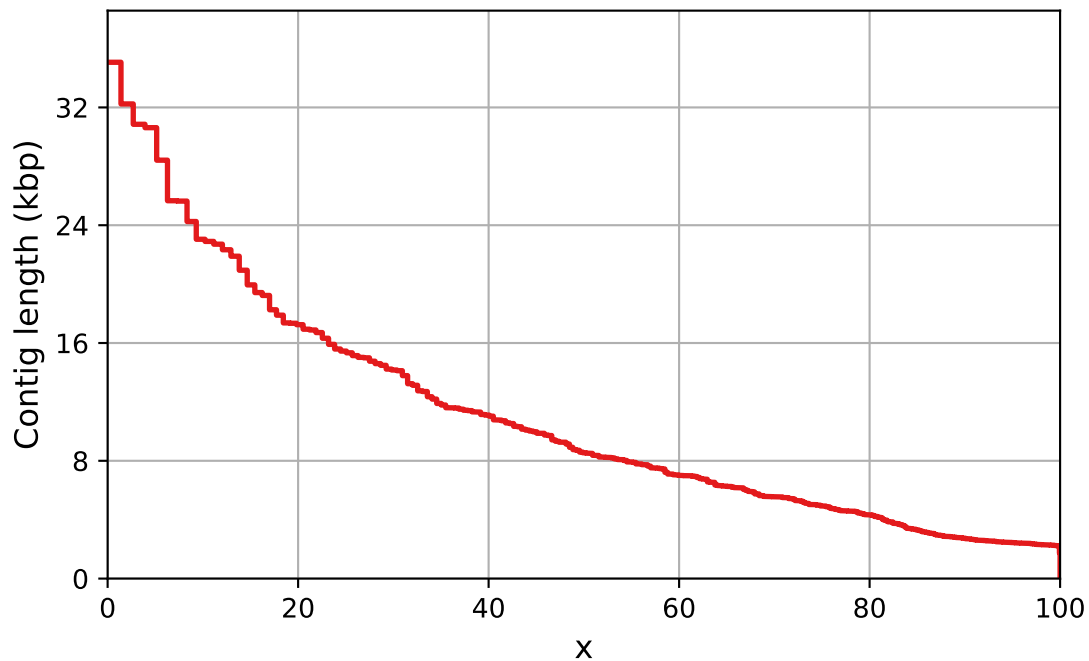


Report

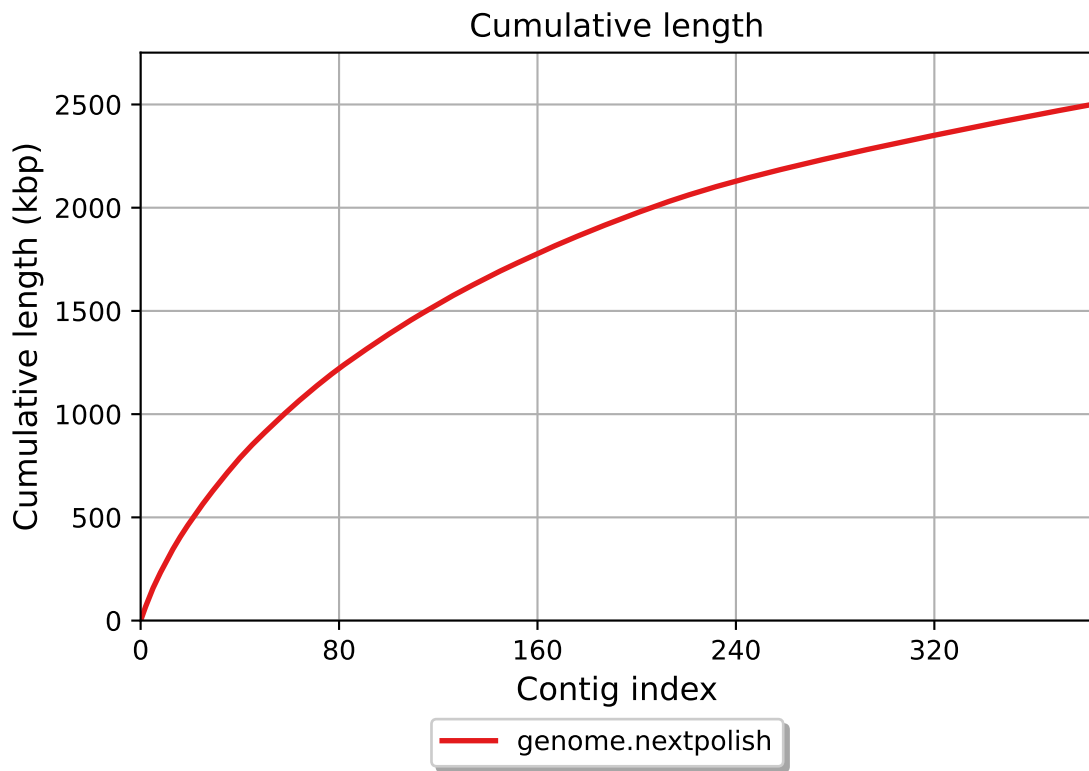
	genome.nextpolish
# contigs (>= 0 bp)	384
# contigs (>= 1000 bp)	384
# contigs (>= 5000 bp)	176
# contigs (>= 10000 bp)	69
# contigs (>= 25000 bp)	7
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2501033
Total length (>= 1000 bp)	2501033
Total length (>= 5000 bp)	1861396
Total length (>= 10000 bp)	1117103
Total length (>= 25000 bp)	208502
Total length (>= 50000 bp)	0
# contigs	384
Largest contig	35067
Total length	2501033
GC (%)	38.57
N50	8546
N90	2740
auN	11197.1
L50	84
L90	282
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	9 + 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).

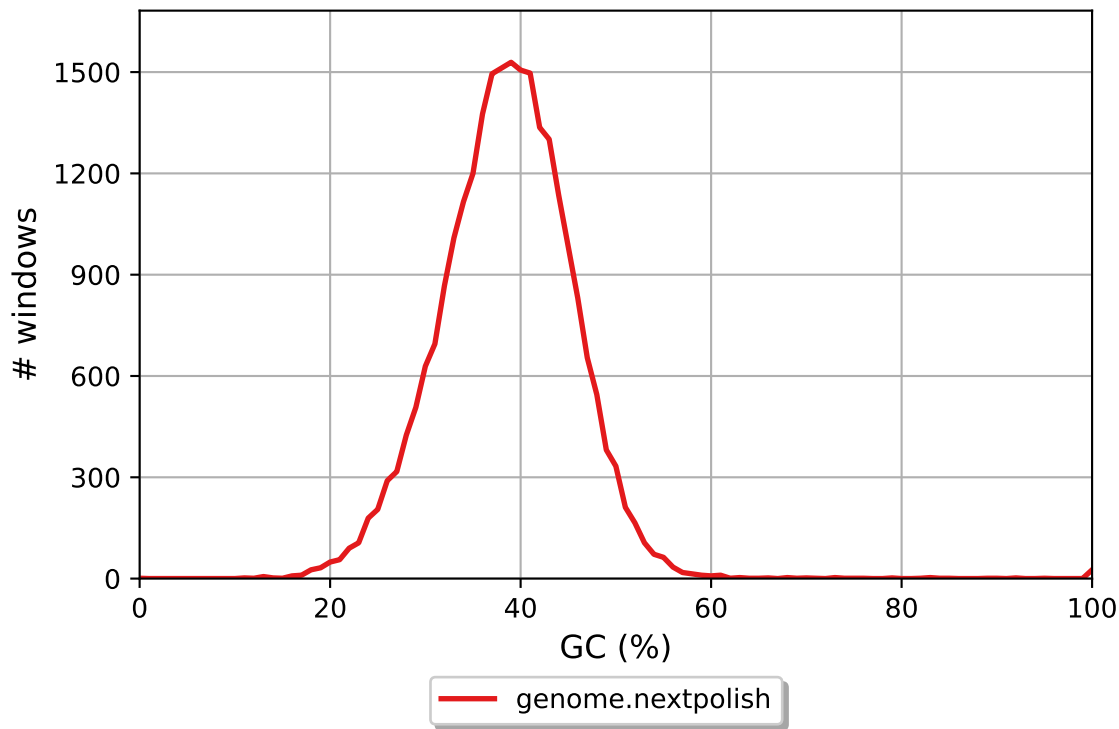
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

