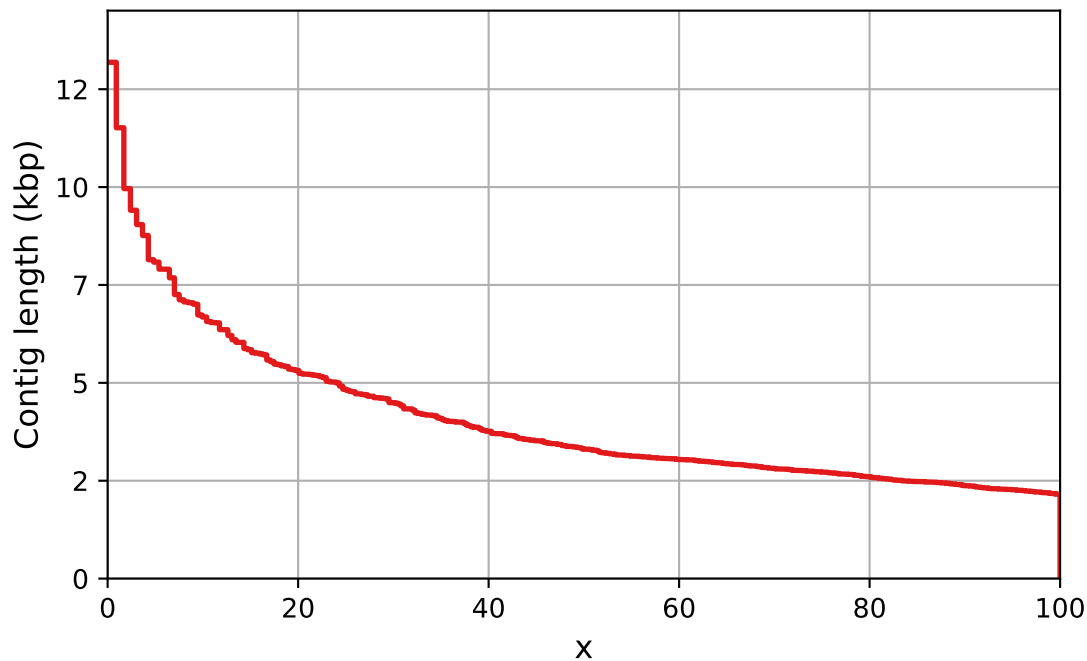


# Report

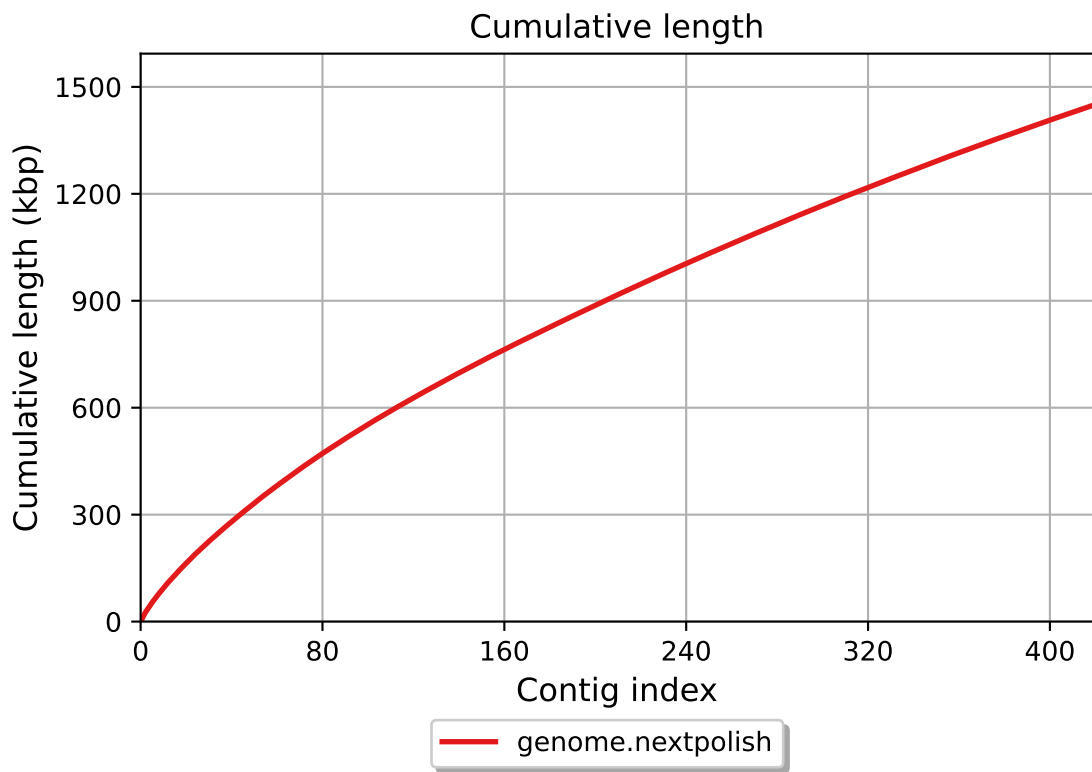
	genome.nextpolish
# contigs (>= 0 bp)	419
# contigs (>= 1000 bp)	419
# contigs (>= 5000 bp)	53
# contigs (>= 10000 bp)	2
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1448443
Total length (>= 1000 bp)	1448443
Total length (>= 5000 bp)	347523
Total length (>= 10000 bp)	24704
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	419
Largest contig	13188
Total length	1448443
GC (%)	42.08
N50	3314
N90	2374
auN	4076.9
L50	148
L90	356
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 1 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).

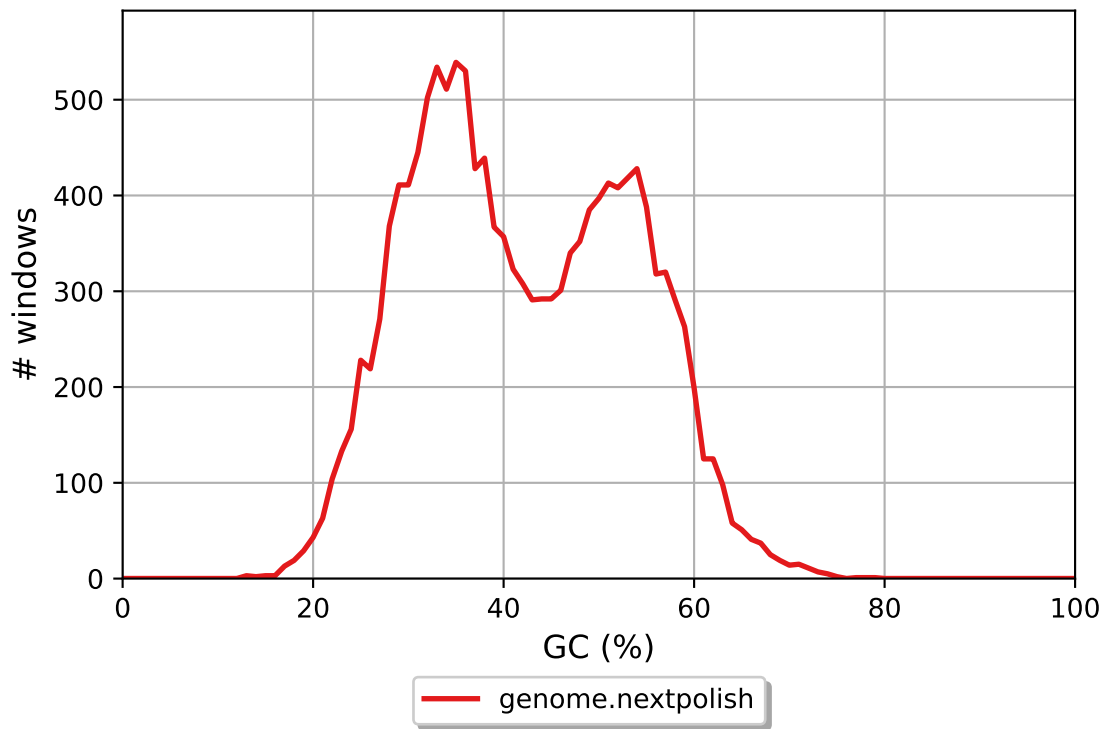
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

