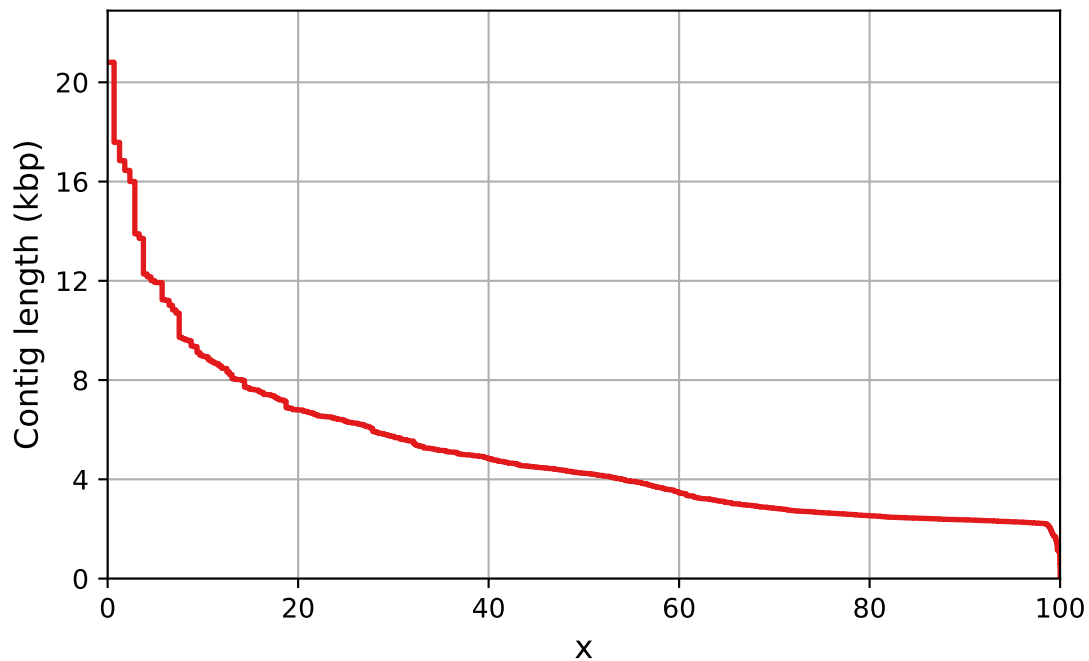


# Report

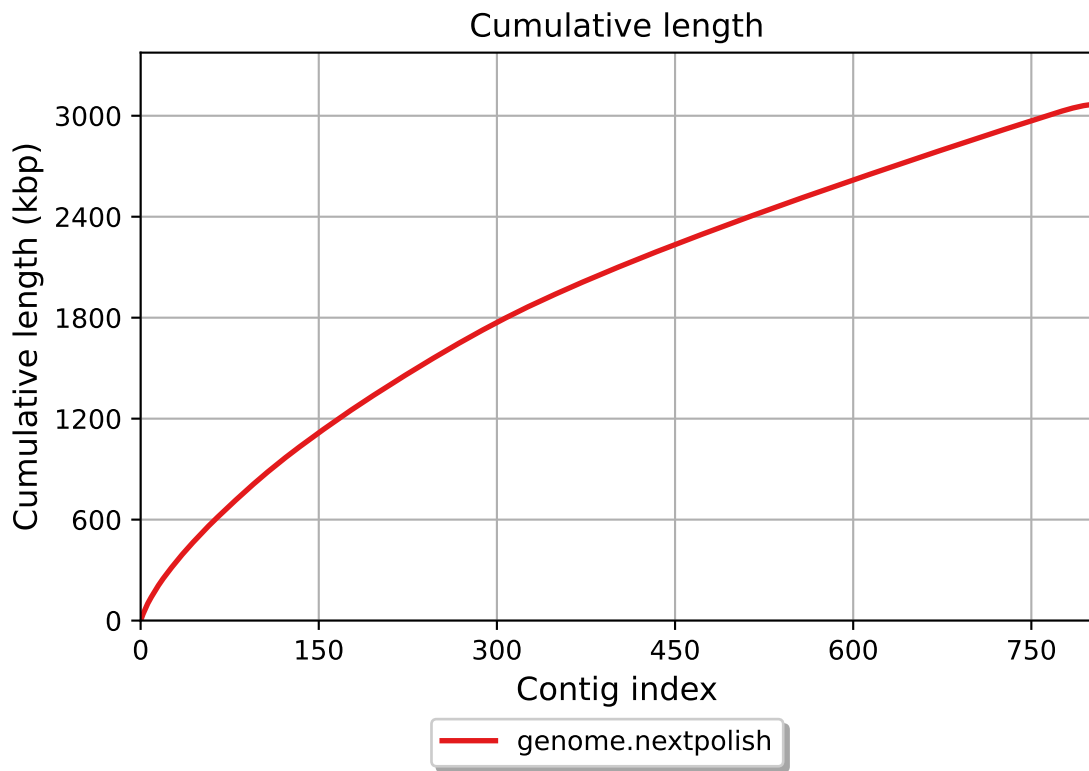
	genome.nextpolish
# contigs (>= 0 bp)	802
# contigs (>= 1000 bp)	800
# contigs (>= 5000 bp)	156
# contigs (>= 10000 bp)	17
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	3068235
Total length (>= 1000 bp)	3066753
Total length (>= 5000 bp)	1146402
Total length (>= 10000 bp)	230569
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	802
Largest contig	20807
Total length	3068235
GC (%)	37.86
N50	4244
N90	2367
auN	5114.9
L50	241
L90	660
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 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).

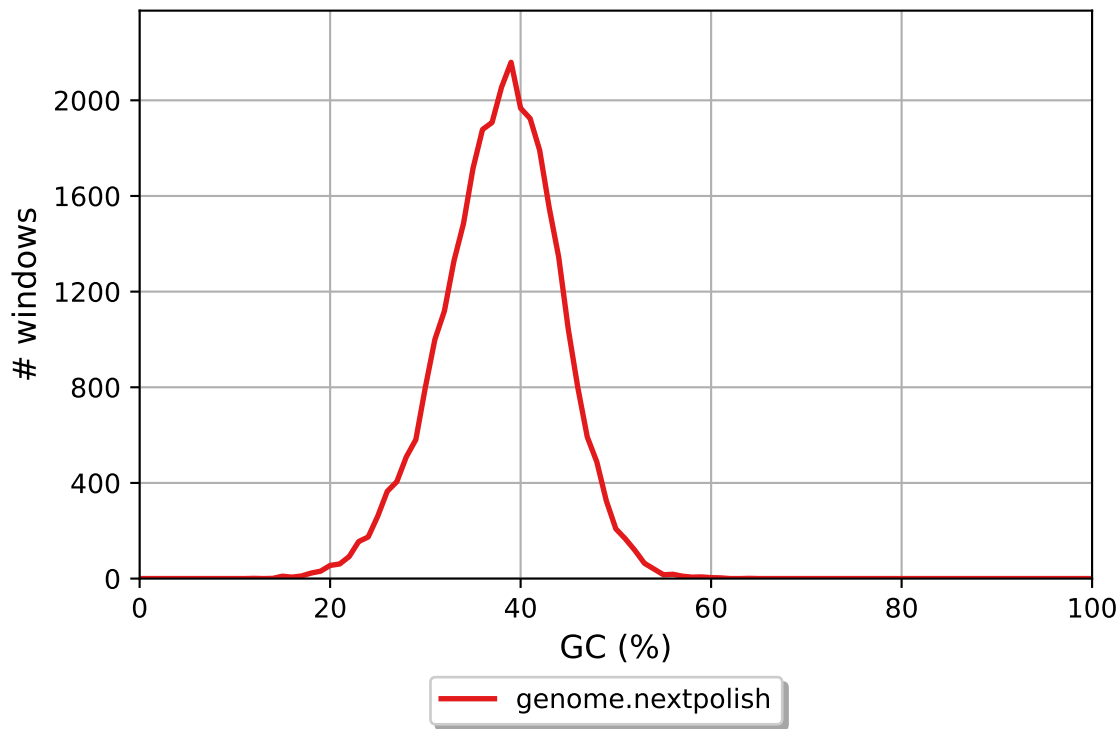
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

