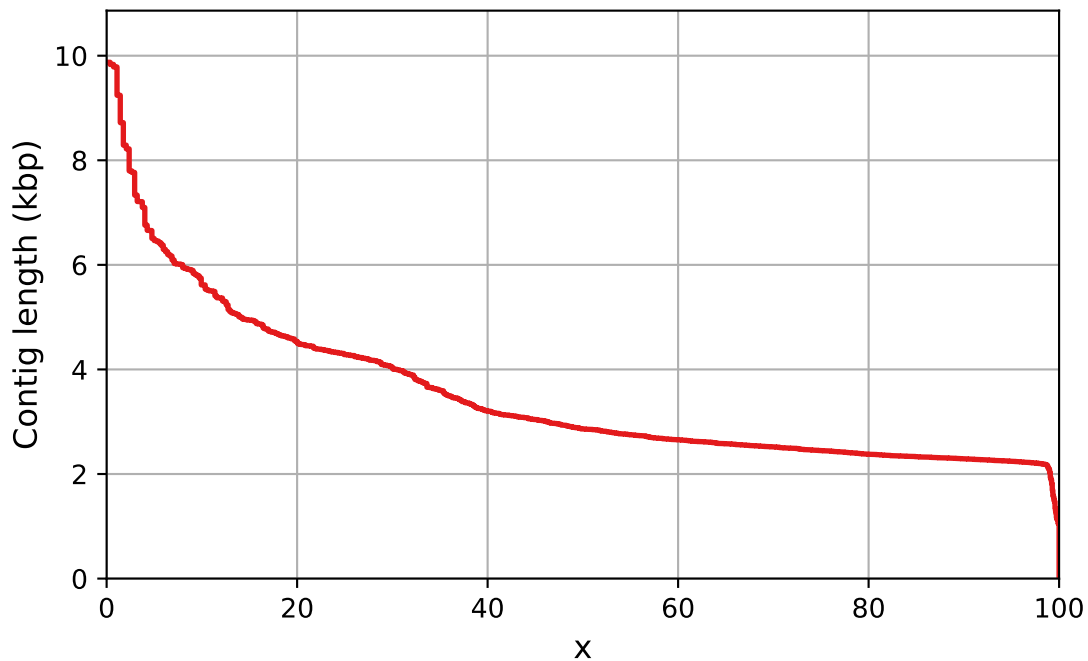


# Report

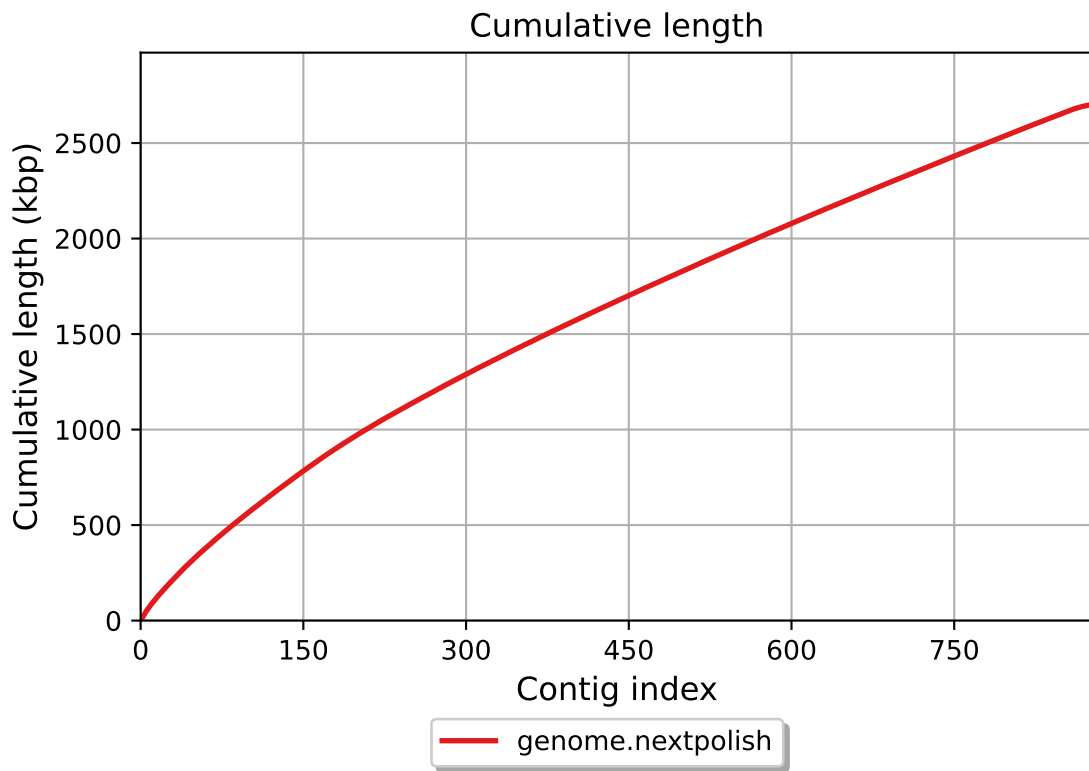
	genome.nextpolish
# contigs (>= 0 bp)	878
# contigs (>= 1000 bp)	878
# contigs (>= 5000 bp)	60
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2702967
Total length (>= 1000 bp)	2702967
Total length (>= 5000 bp)	380168
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	878
Largest contig	9874
Total length	2702967
GC (%)	37.83
N50	2862
N90	2292
auN	3528.9
L50	322
L90	751
# 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).

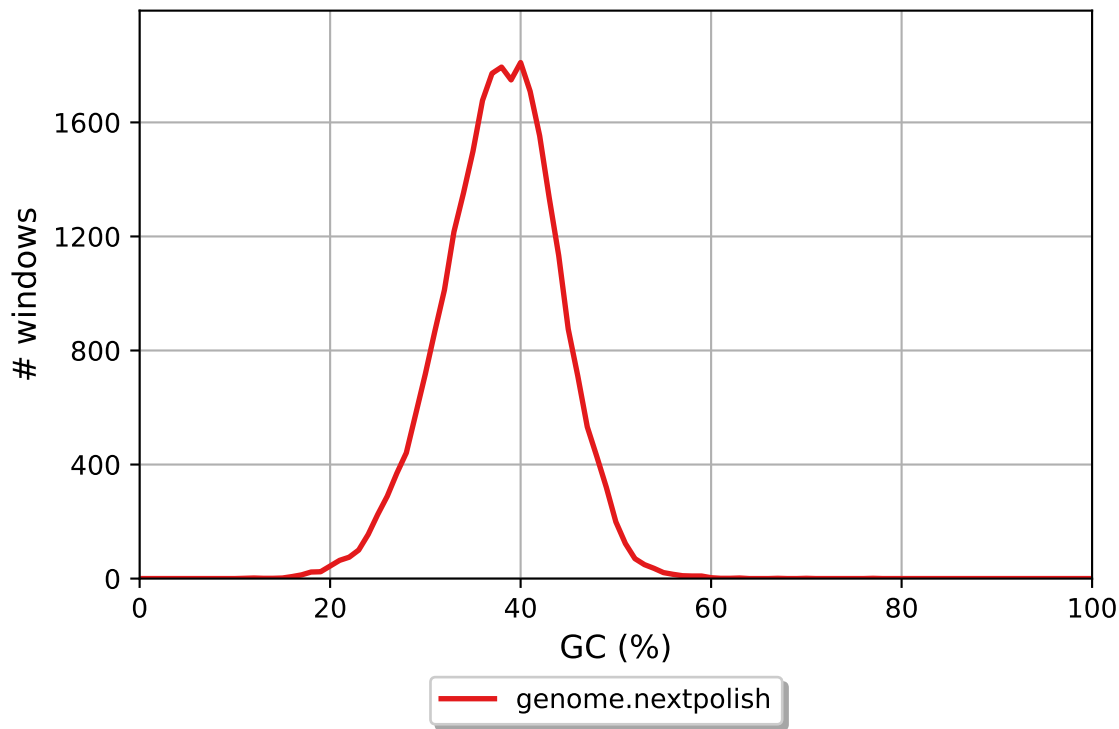
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

