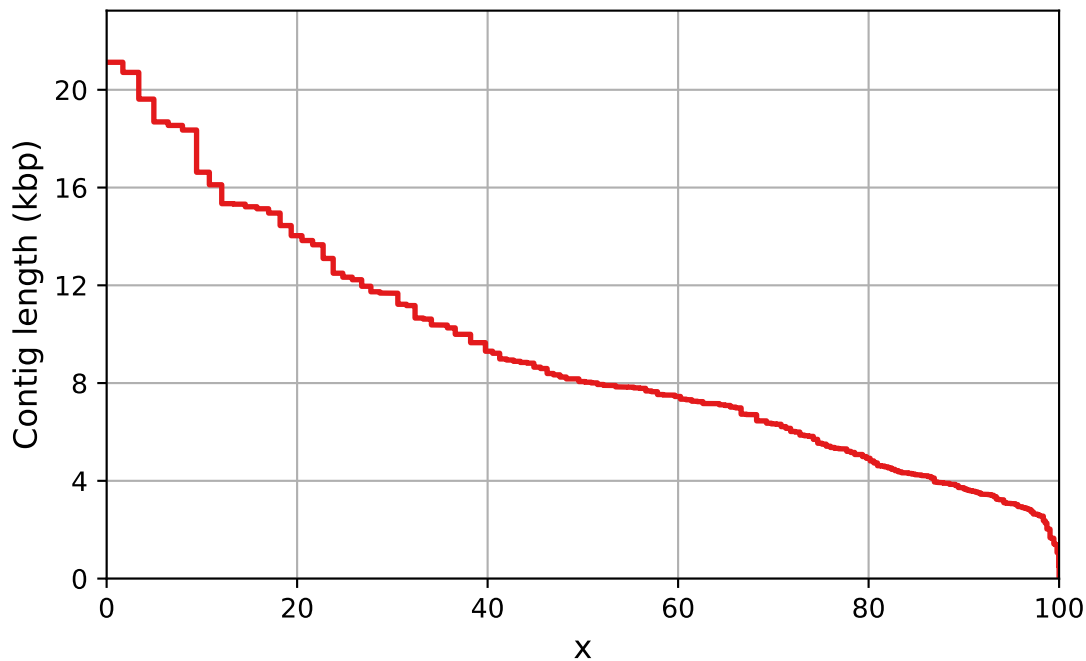


# Report

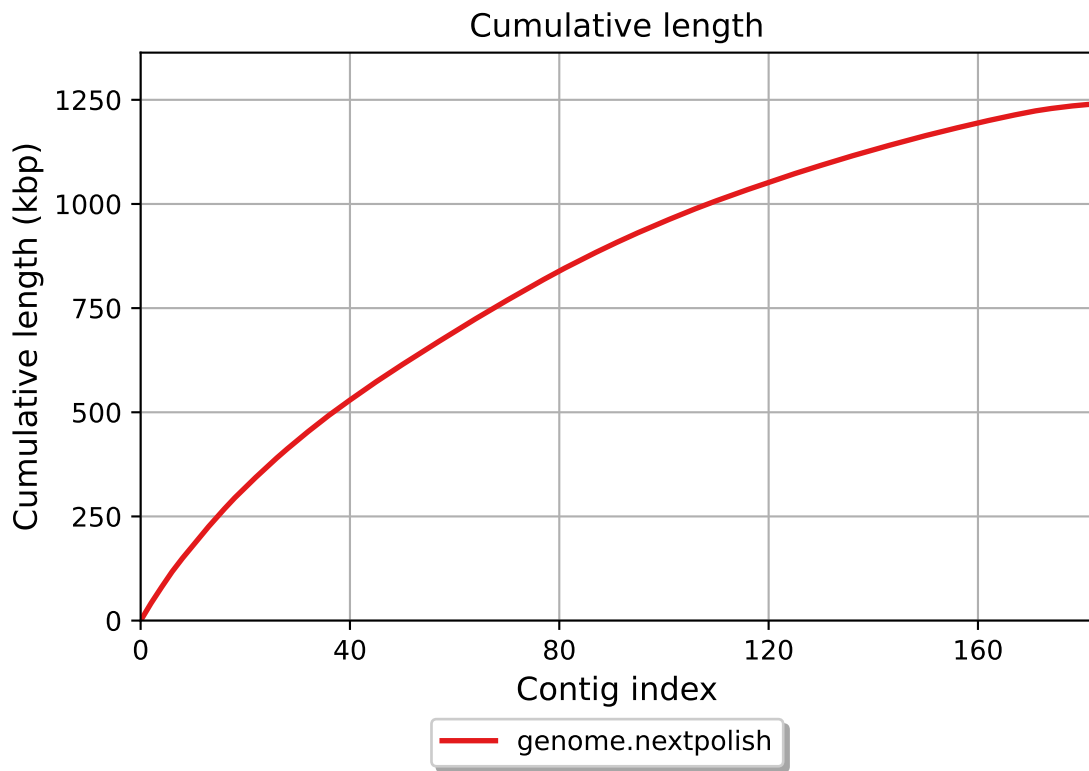
	genome.nextpolish
# contigs (>= 0 bp)	182
# contigs (>= 1000 bp)	181
# contigs (>= 5000 bp)	105
# contigs (>= 10000 bp)	32
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1239317
Total length (>= 1000 bp)	1238807
Total length (>= 5000 bp)	983522
Total length (>= 10000 bp)	453602
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	182
Largest contig	21126
Total length	1239317
GC (%)	37.99
N50	8062
N90	3718
auN	9394.7
L50	51
L90	136
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 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).

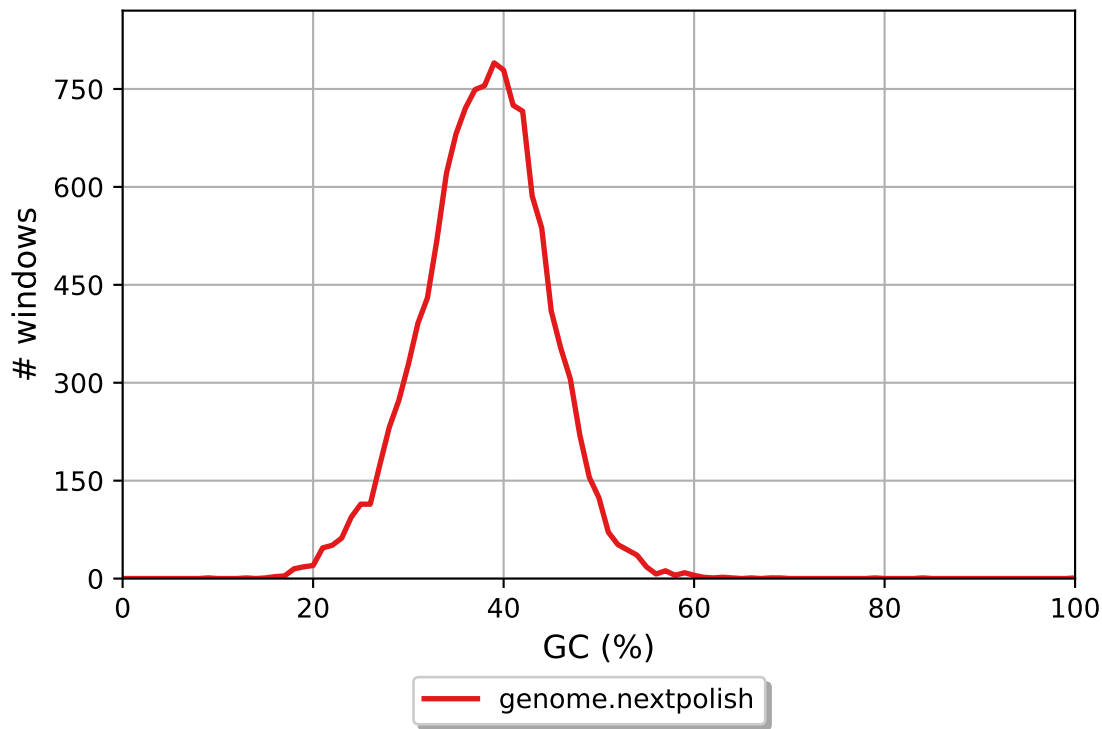
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

