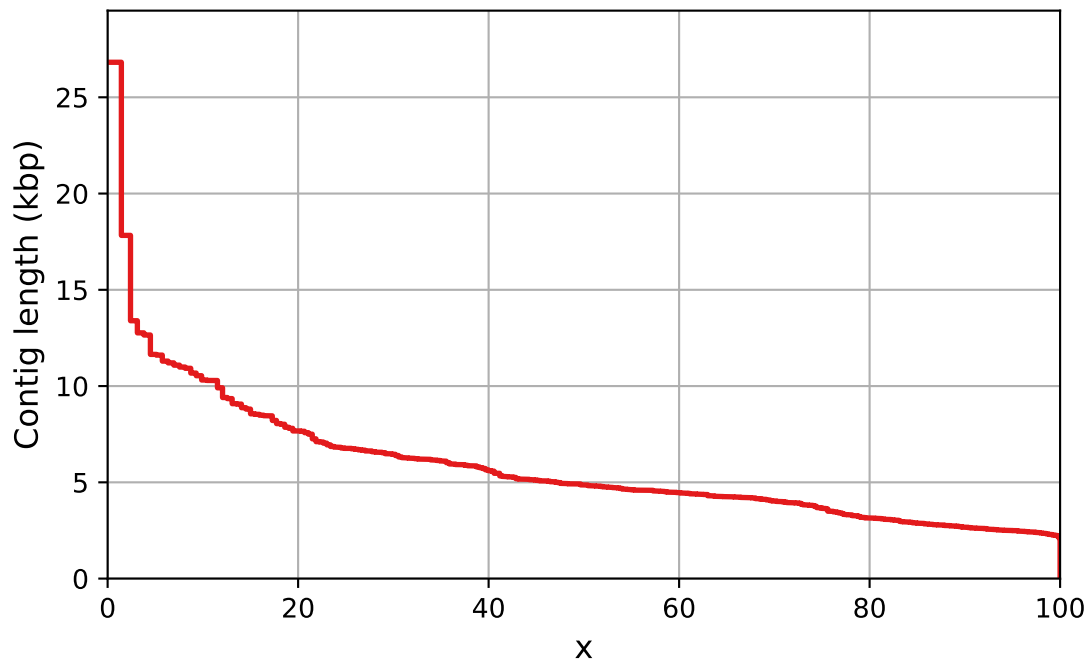


# Report

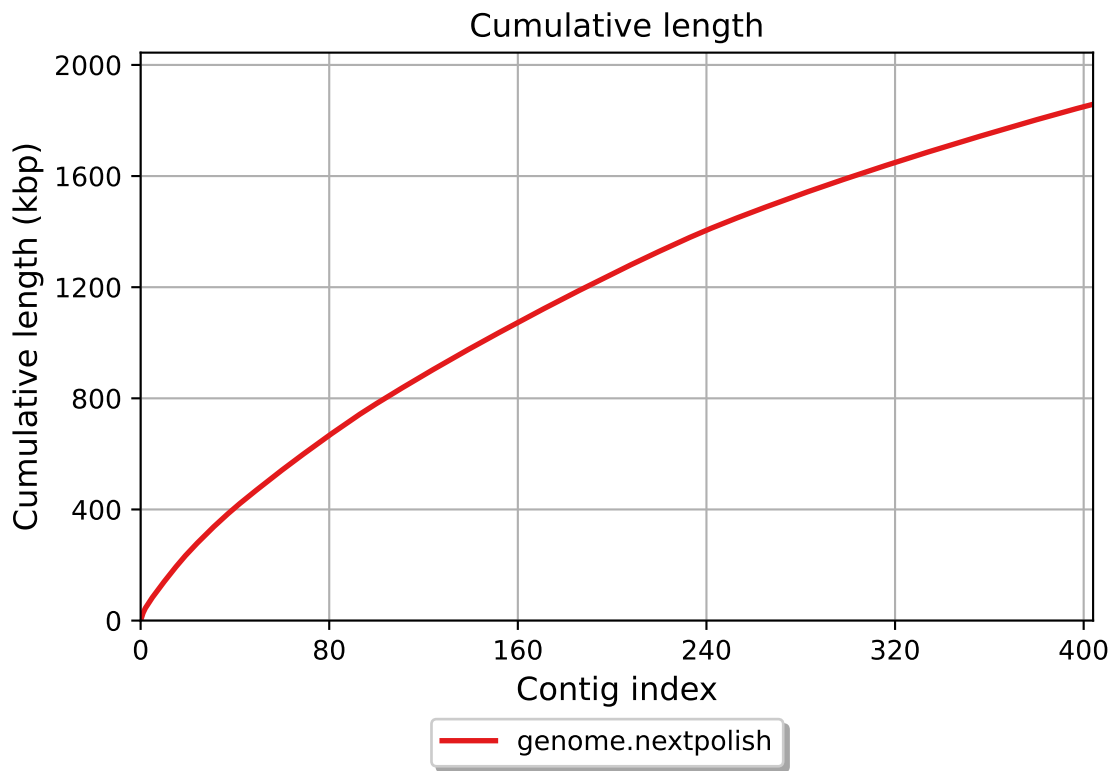
	genome.nextpolish
# contigs (>= 0 bp)	404
# contigs (>= 1000 bp)	404
# contigs (>= 5000 bp)	119
# contigs (>= 10000 bp)	17
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1858419
Total length (>= 1000 bp)	1858419
Total length (>= 5000 bp)	878410
Total length (>= 10000 bp)	214315
Total length (>= 25000 bp)	26819
Total length (>= 50000 bp)	0
# contigs	404
Largest contig	26819
Total length	1858419
GC (%)	33.59
N50	4867
N90	2669
auN	5927.3
L50	130
L90	329
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 2 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).

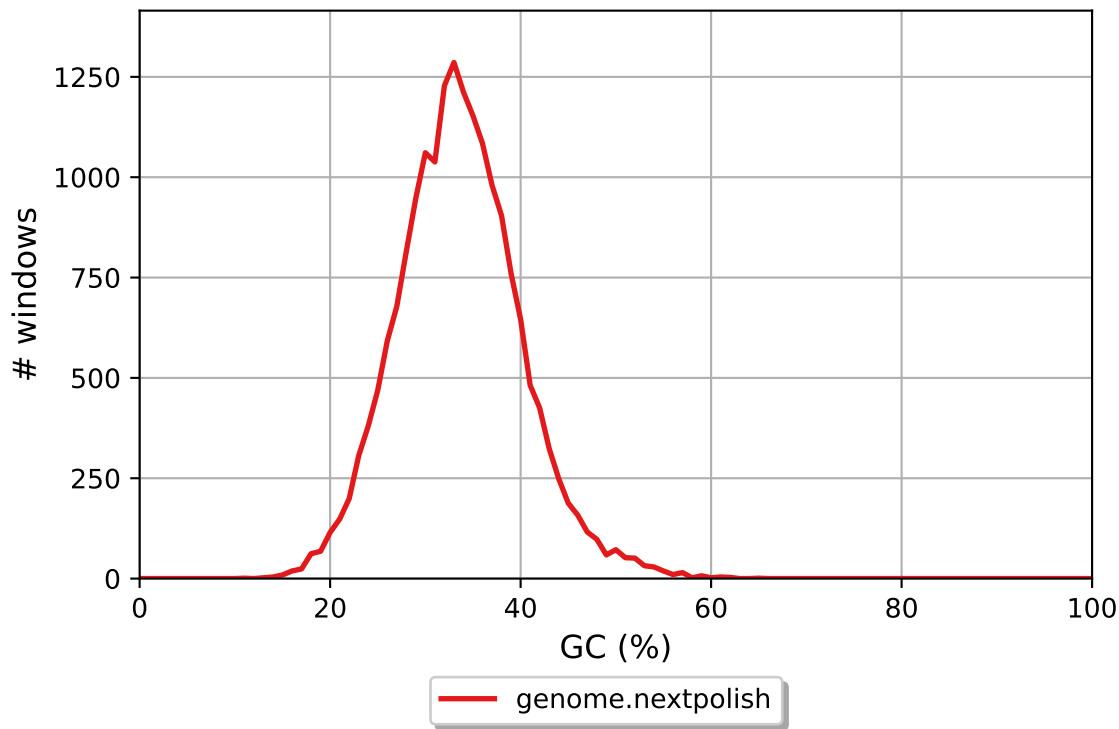
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

