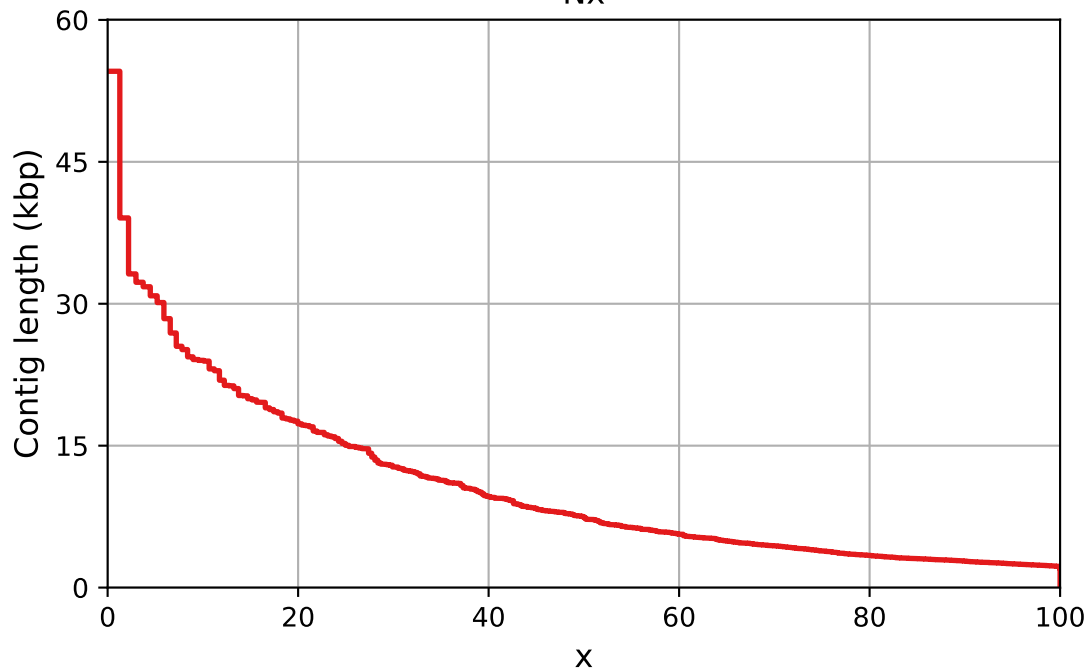


# Report

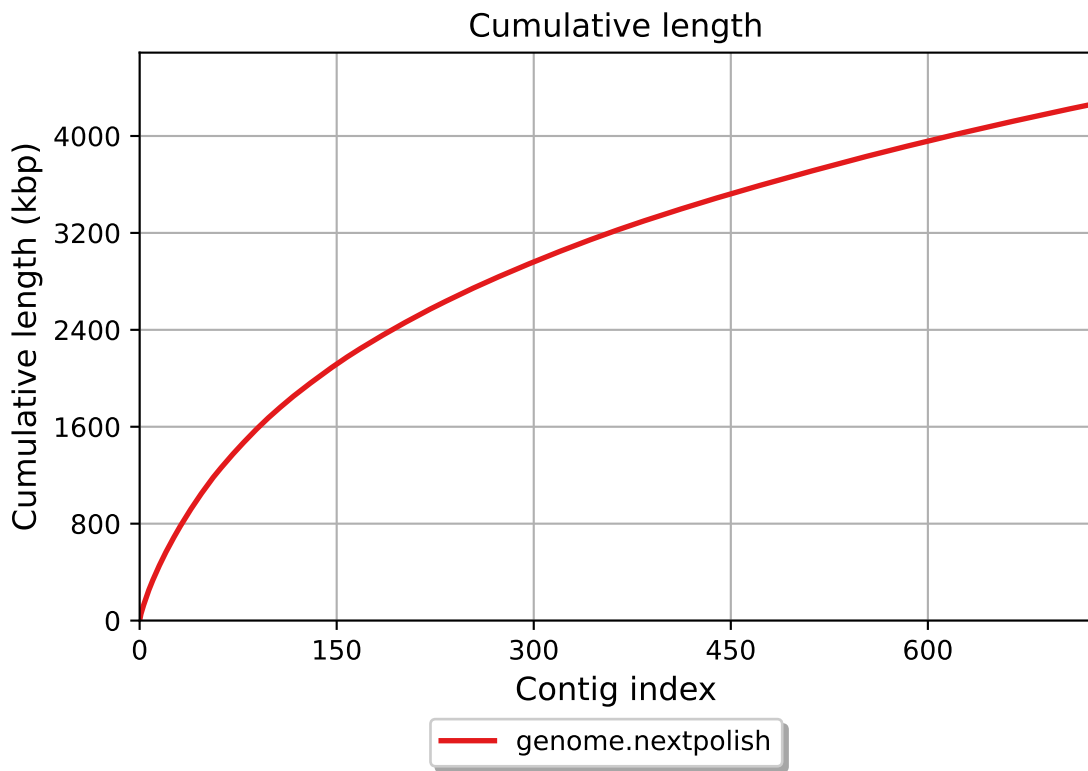
	genome.nextpolish
# contigs (>= 0 bp)	725
# contigs (>= 1000 bp)	725
# contigs (>= 5000 bp)	256
# contigs (>= 10000 bp)	98
# contigs (>= 25000 bp)	11
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	4261880
Total length (>= 1000 bp)	4261880
Total length (>= 5000 bp)	2754566
Total length (>= 10000 bp)	1673848
Total length (>= 25000 bp)	357819
Total length (>= 50000 bp)	54571
# contigs	725
Largest contig	54571
Total length	4261880
GC (%)	35.08
N50	7482
N90	2801
auN	10888.5
L50	152
L90	555
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	8 + 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).

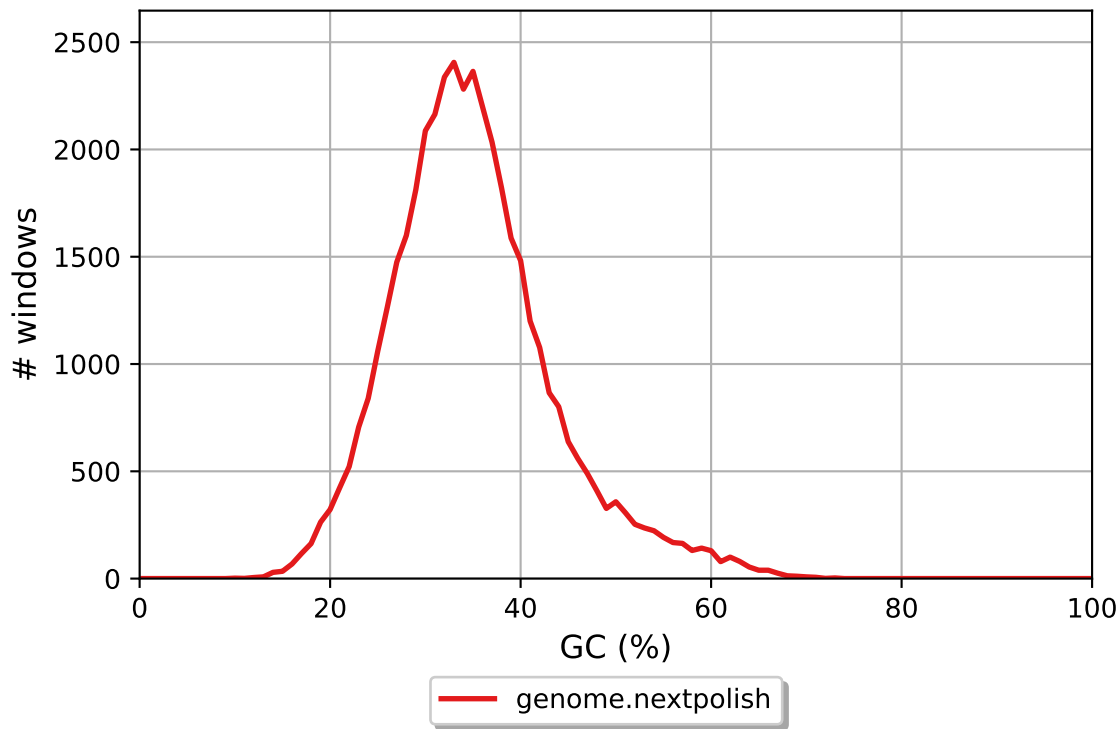
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

