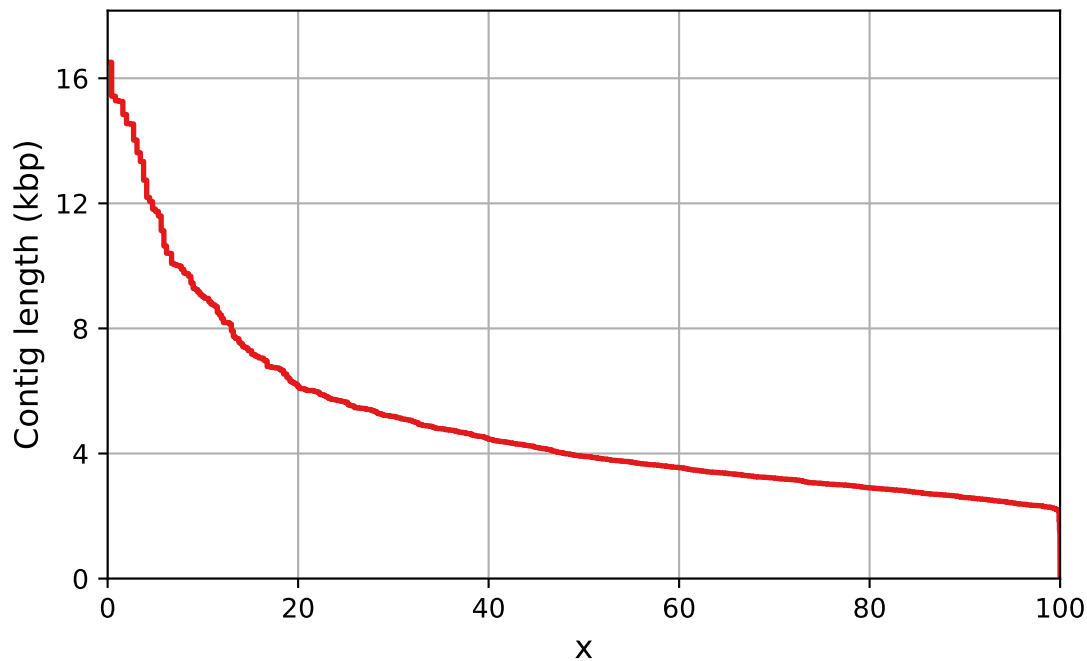


# Report

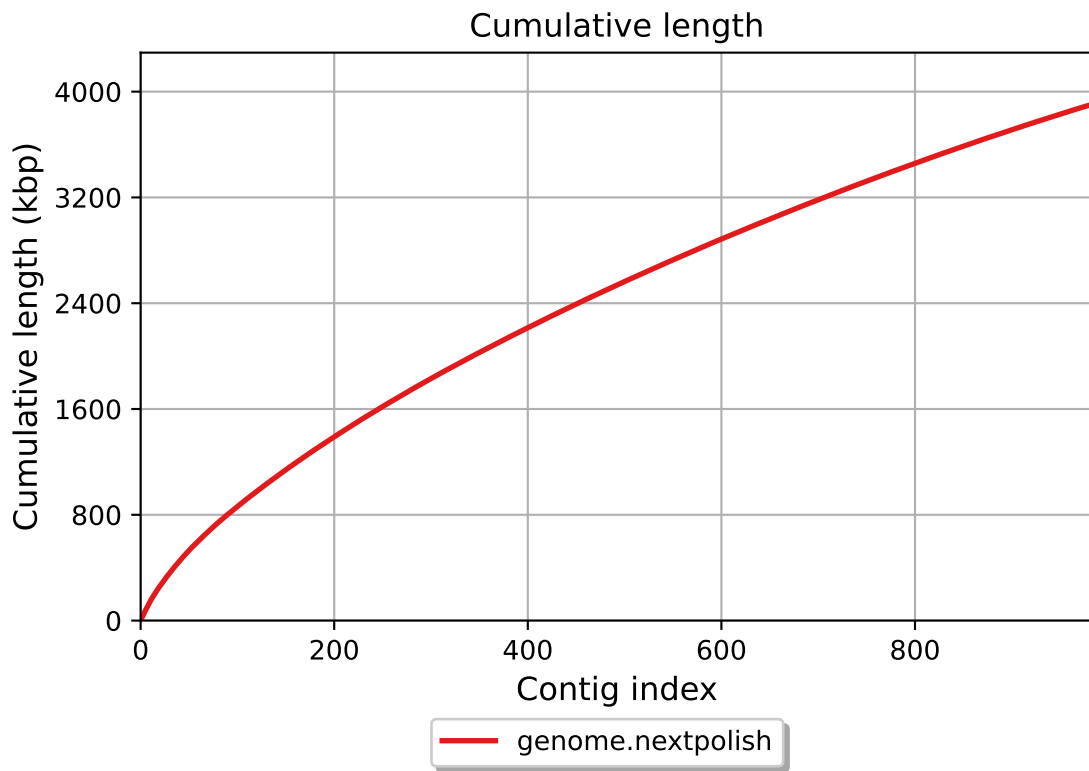
	genome.nextpolish
# contigs (>= 0 bp)	984
# contigs (>= 1000 bp)	983
# contigs (>= 5000 bp)	174
# contigs (>= 10000 bp)	23
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	3904469
Total length (>= 1000 bp)	3903526
Total length (>= 5000 bp)	1263505
Total length (>= 10000 bp)	292217
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	984
Largest contig	16511
Total length	3904469
GC (%)	37.24
N50	3911
N90	2594
auN	4953.6
L50	331
L90	822
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	6 + 5 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).

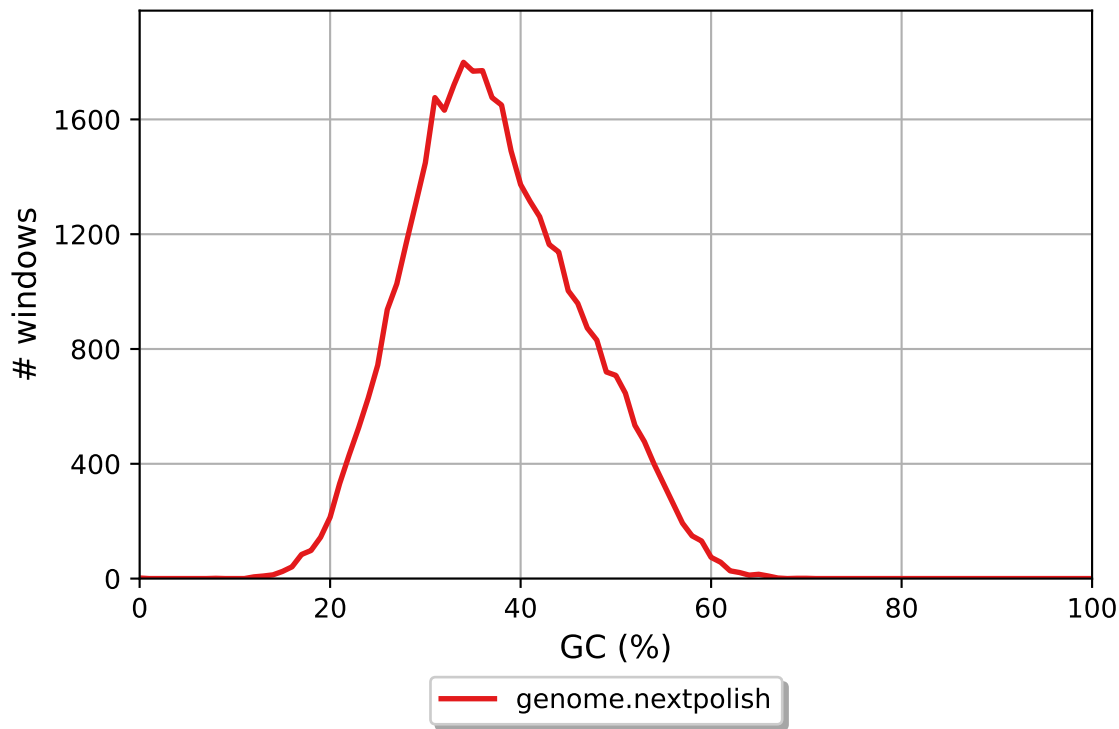
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

