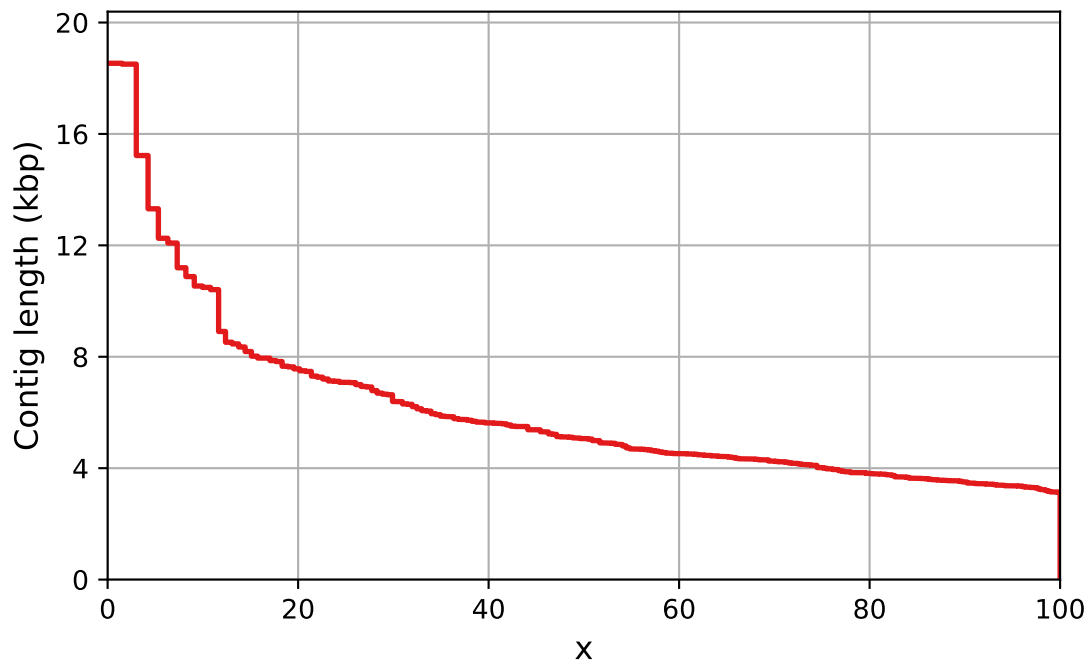


# Report

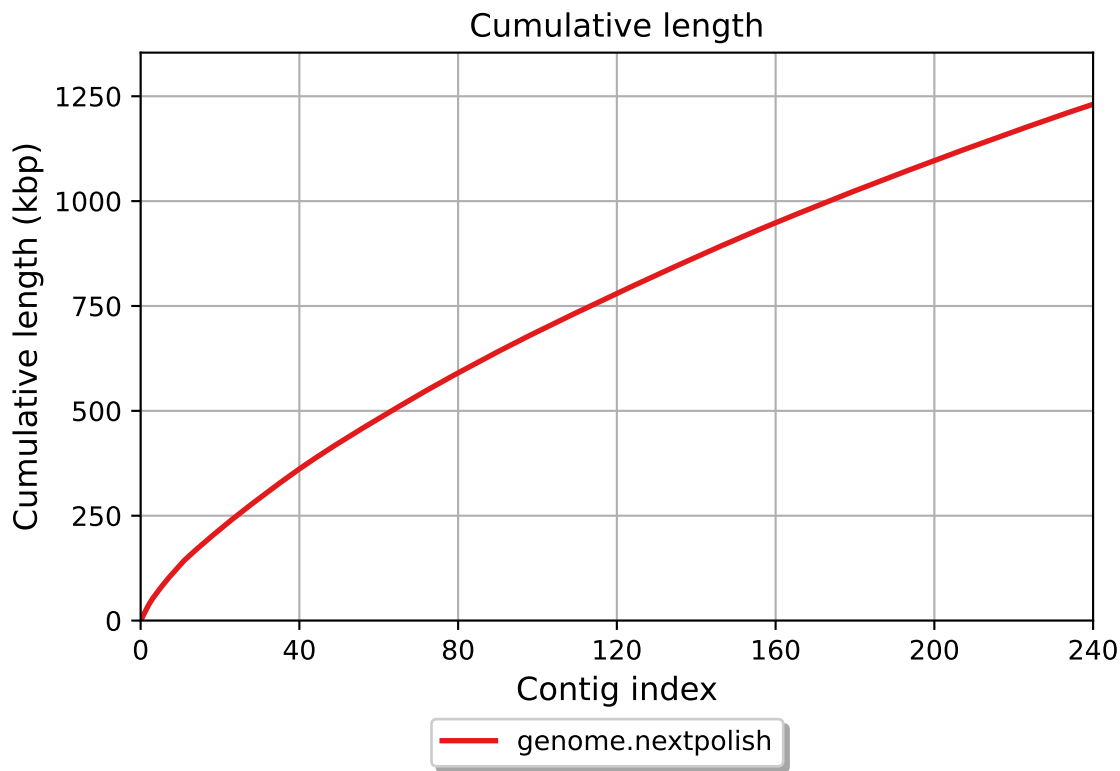
	genome.nextpolish
# contigs (>= 0 bp)	240
# contigs (>= 1000 bp)	240
# contigs (>= 5000 bp)	87
# contigs (>= 10000 bp)	11
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1230894
Total length (>= 1000 bp)	1230894
Total length (>= 5000 bp)	626081
Total length (>= 10000 bp)	143433
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	240
Largest contig	18537
Total length	1230894
GC (%)	47.64
N50	5063
N90	3503
auN	6126.8
L50	85
L90	204
# 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).

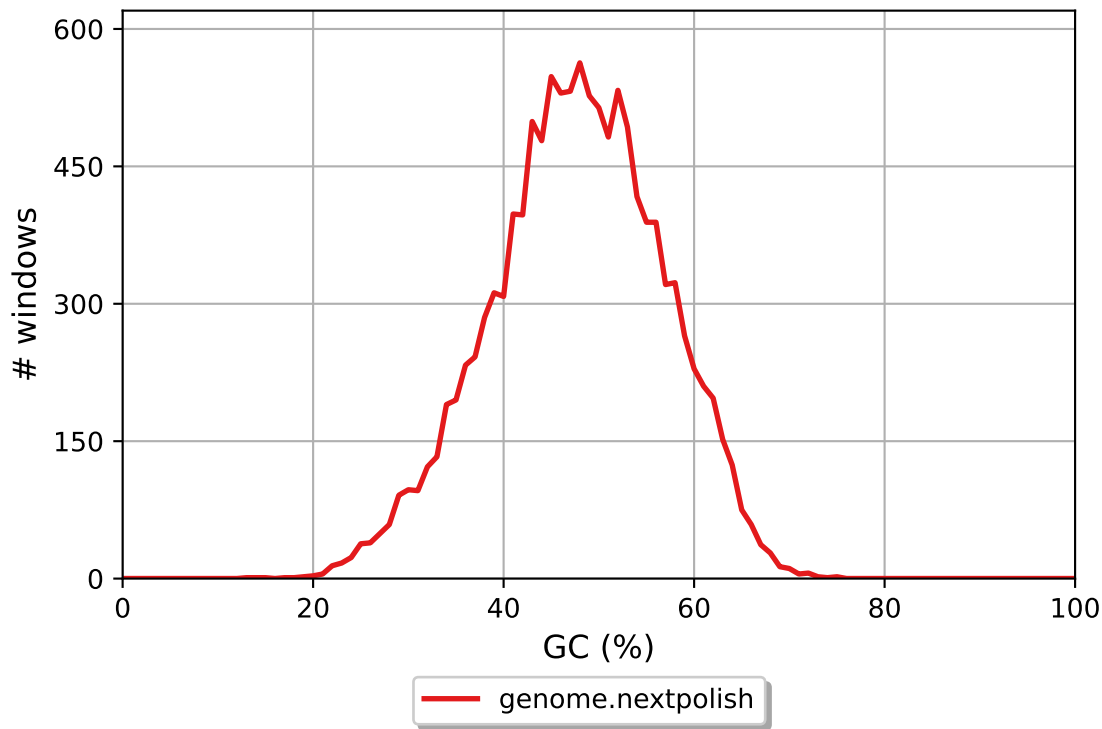
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

