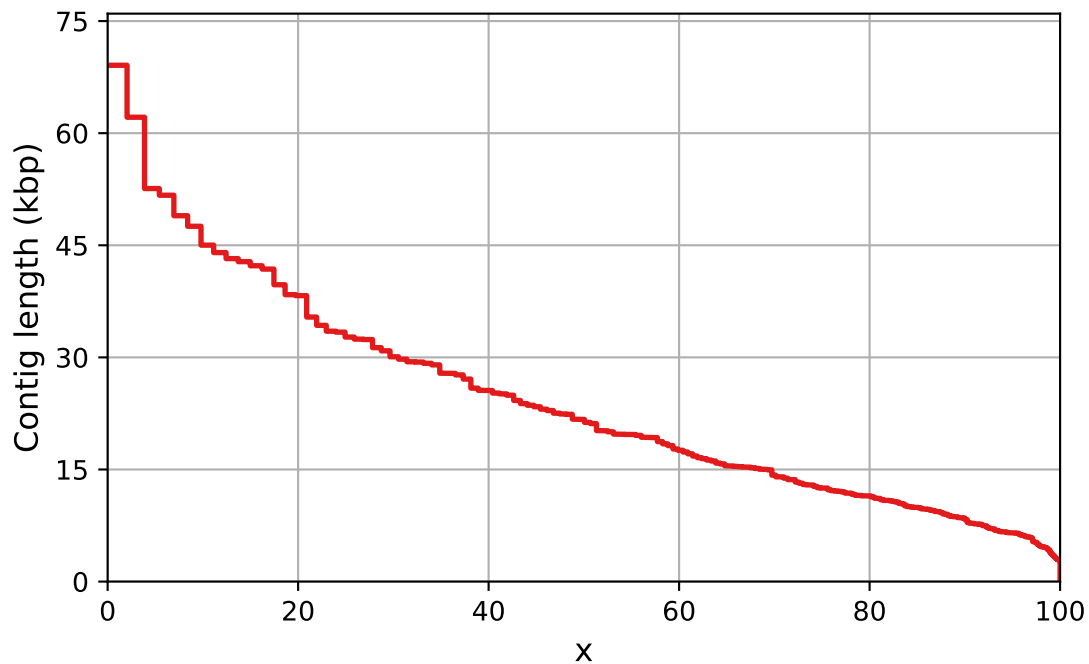


# Report

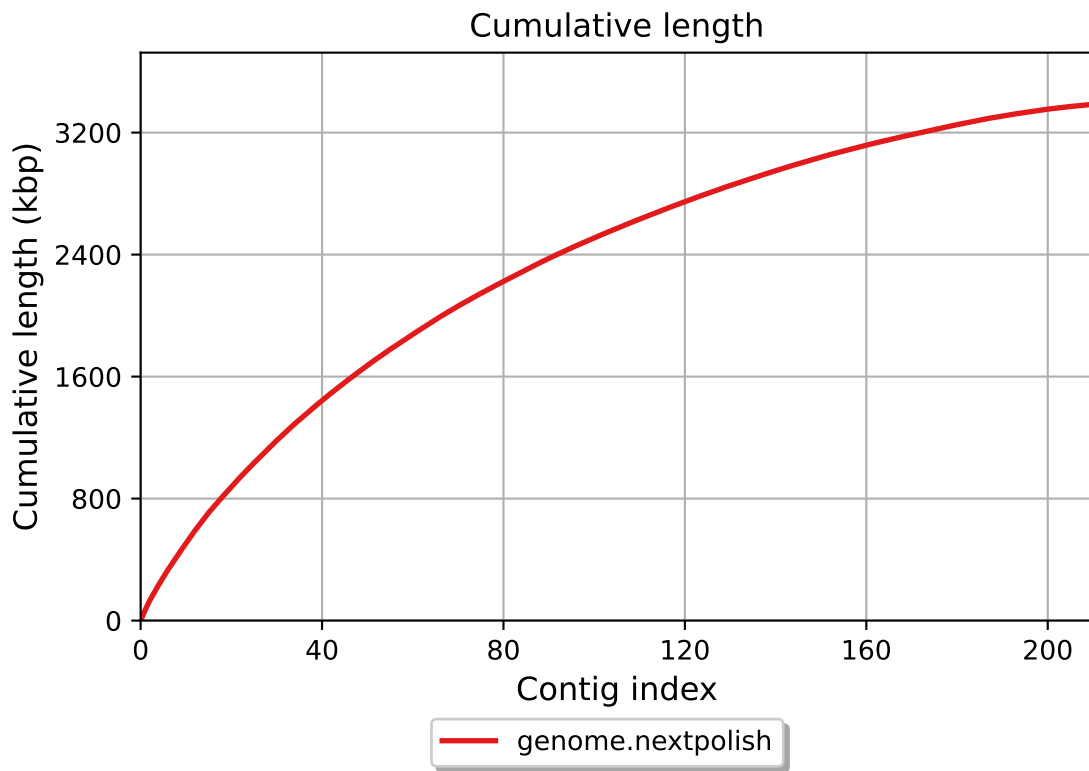
	genome.nextpolish
# contigs (>= 0 bp)	210
# contigs (>= 1000 bp)	210
# contigs (>= 5000 bp)	189
# contigs (>= 10000 bp)	130
# contigs (>= 25000 bp)	39
# contigs (>= 50000 bp)	4
Total length (>= 0 bp)	3385720
Total length (>= 1000 bp)	3385720
Total length (>= 5000 bp)	3303817
Total length (>= 10000 bp)	2852980
Total length (>= 25000 bp)	1418504
Total length (>= 50000 bp)	235508
# contigs	210
Largest contig	69086
Total length	3385720
GC (%)	57.81
N50	21697
N90	8516
auN	24744.4
L50	51
L90	151
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	5 + 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).

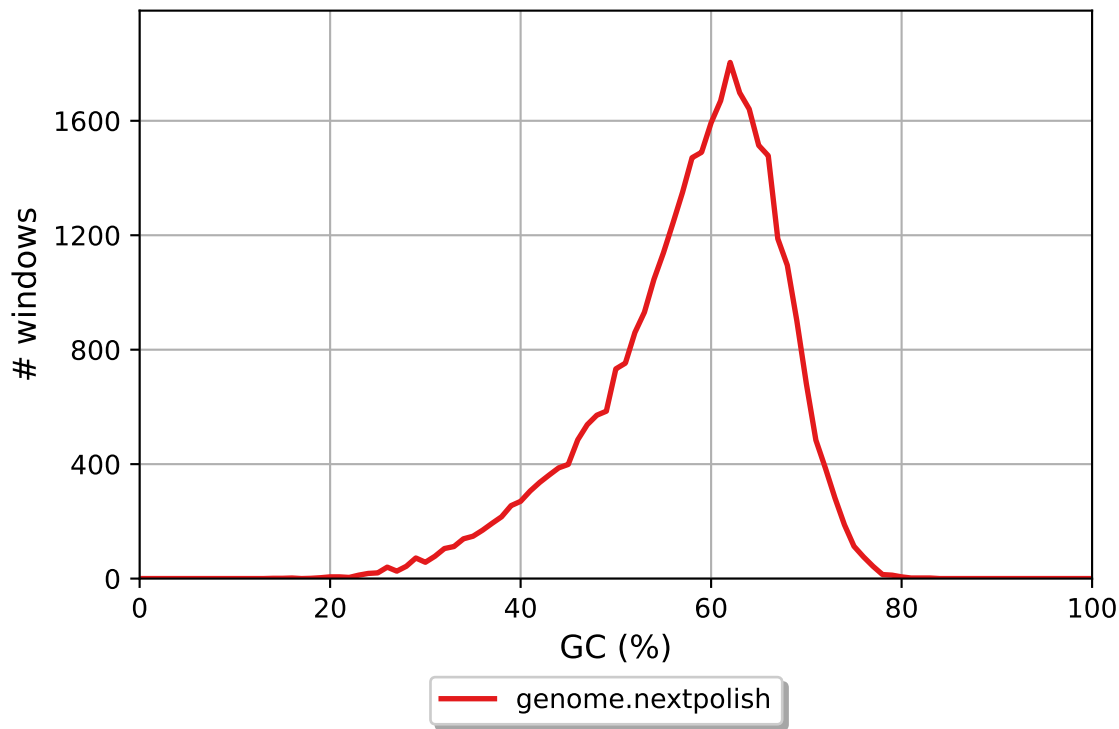
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

