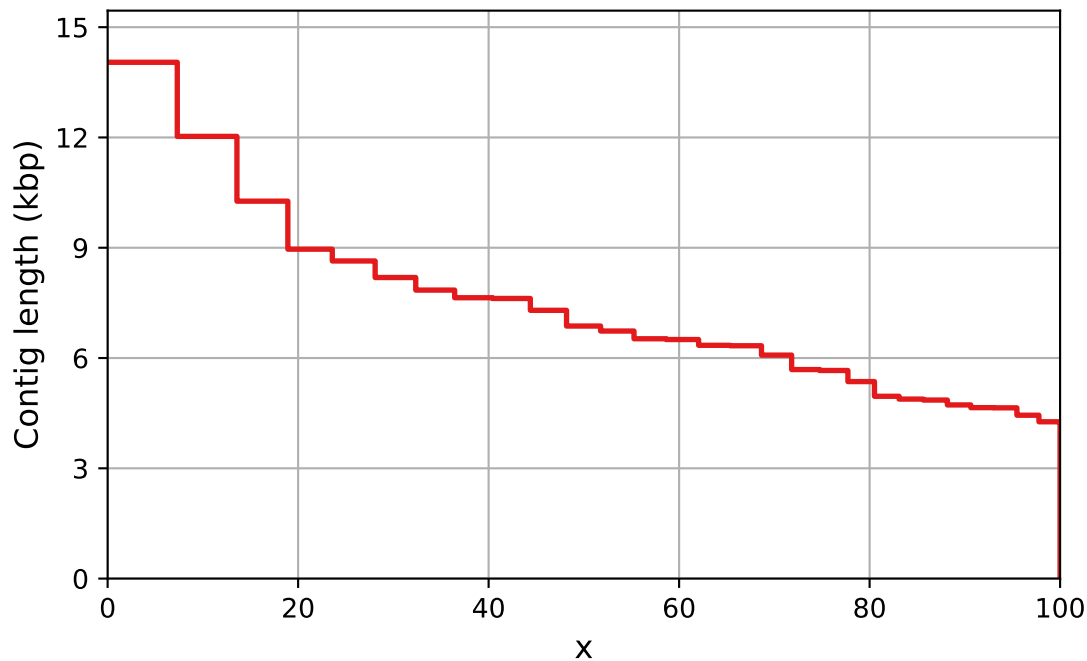


# Report

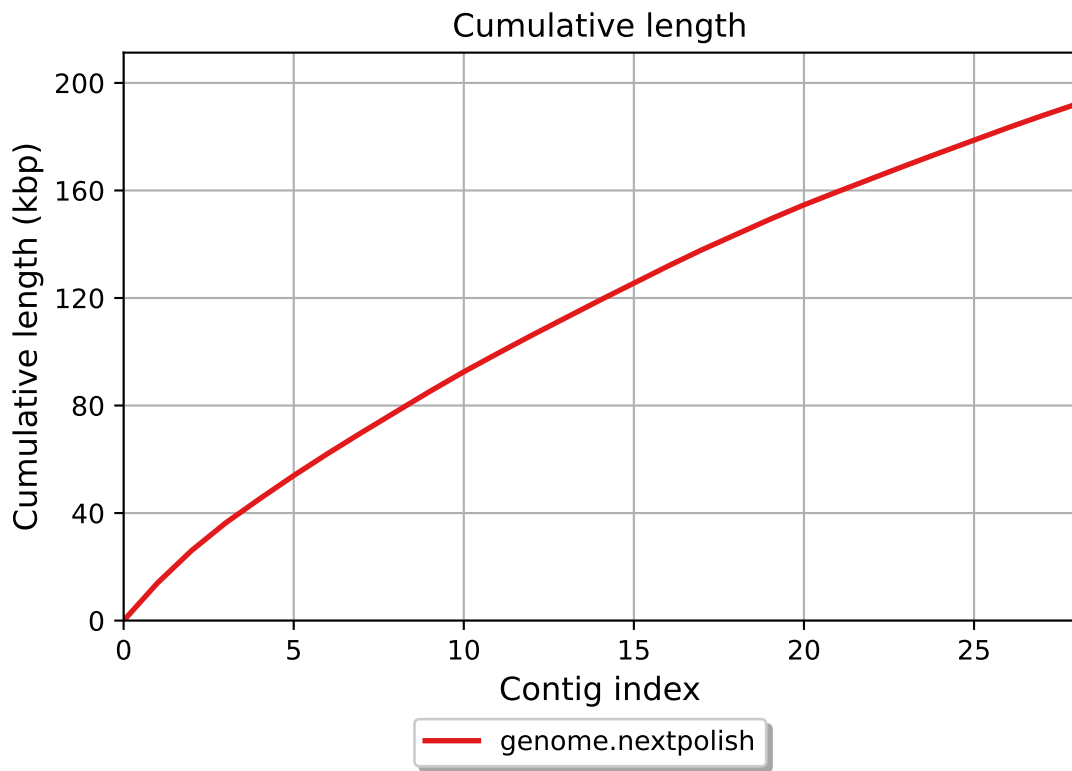
	genome.nextpolish
# contigs (>= 0 bp)	28
# contigs (>= 1000 bp)	28
# contigs (>= 5000 bp)	20
# contigs (>= 10000 bp)	3
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	192062
Total length (>= 1000 bp)	192062
Total length (>= 5000 bp)	154634
Total length (>= 10000 bp)	36341
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	28
Largest contig	14045
Total length	192062
GC (%)	27.71
N50	6870
N90	4724
auN	7622.6
L50	11
L90	24
# 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).

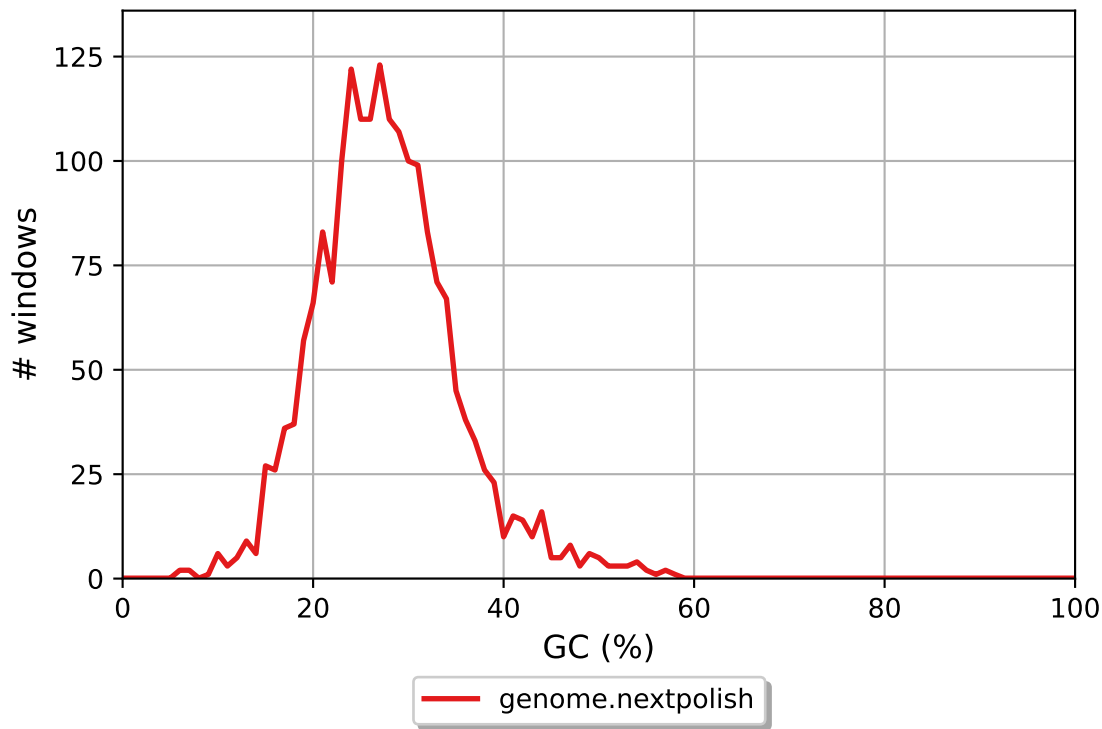
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

