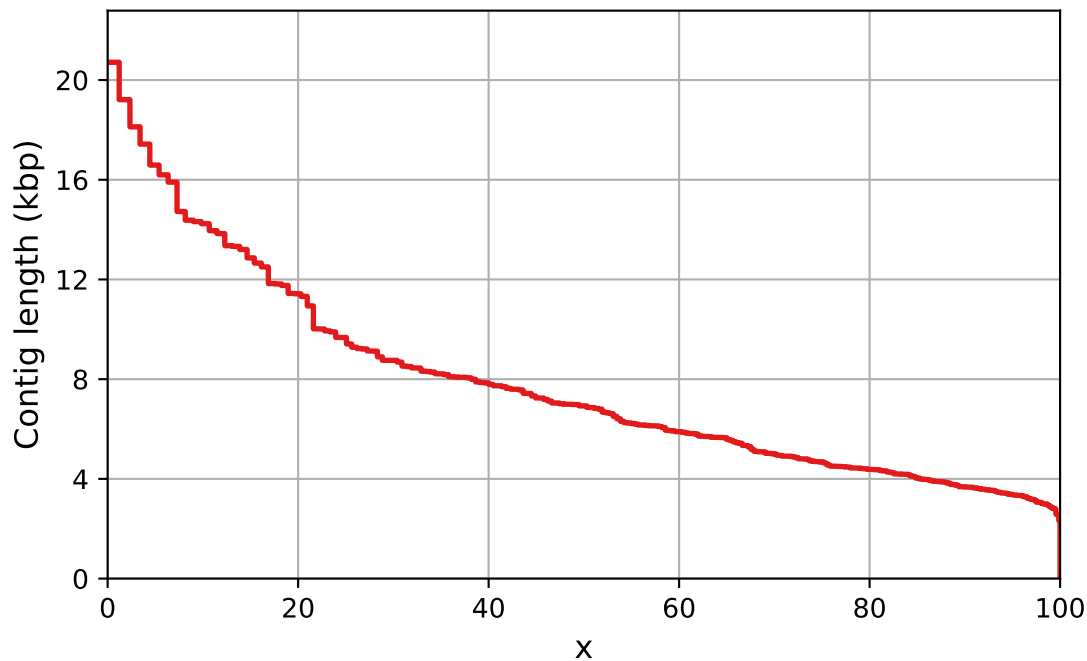


# Report

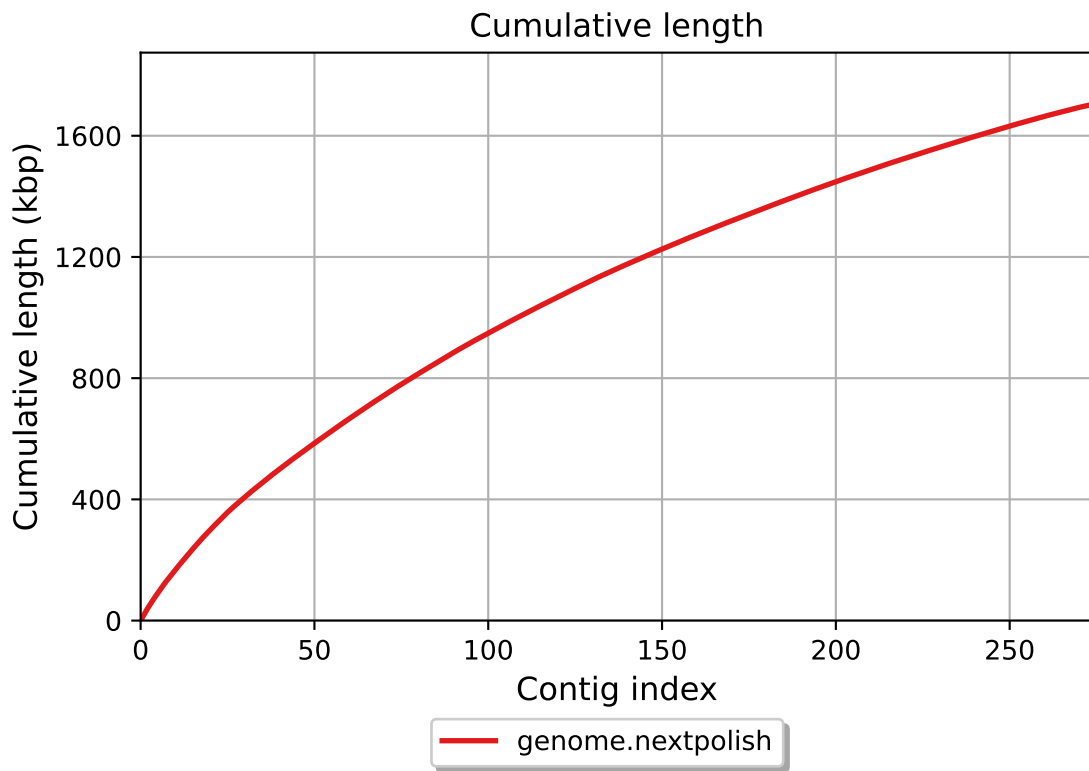
	genome.nextpolish
# contigs (>= 0 bp)	274
# contigs (>= 1000 bp)	274
# contigs (>= 5000 bp)	143
# contigs (>= 10000 bp)	28
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1703870
Total length (>= 1000 bp)	1703870
Total length (>= 5000 bp)	1191589
Total length (>= 10000 bp)	388100
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	274
Largest contig	20712
Total length	1703870
GC (%)	56.78
N50	6926
N90	3676
auN	7889.3
L50	86
L90	223
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	7 + 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).

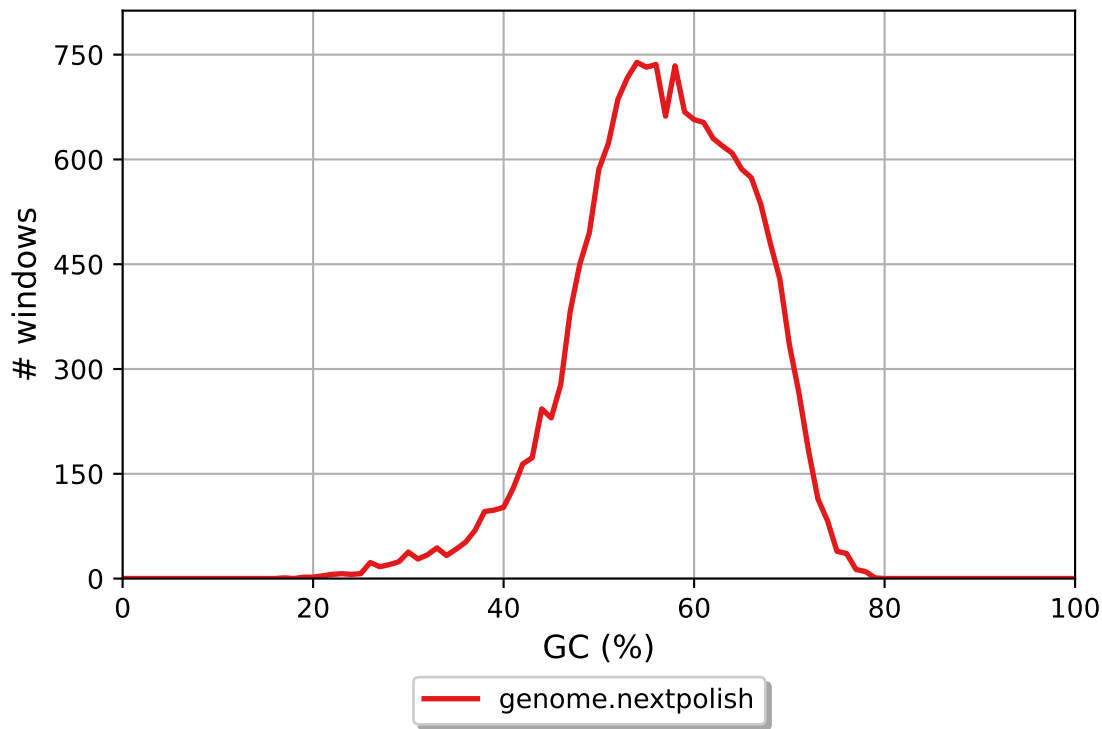
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

