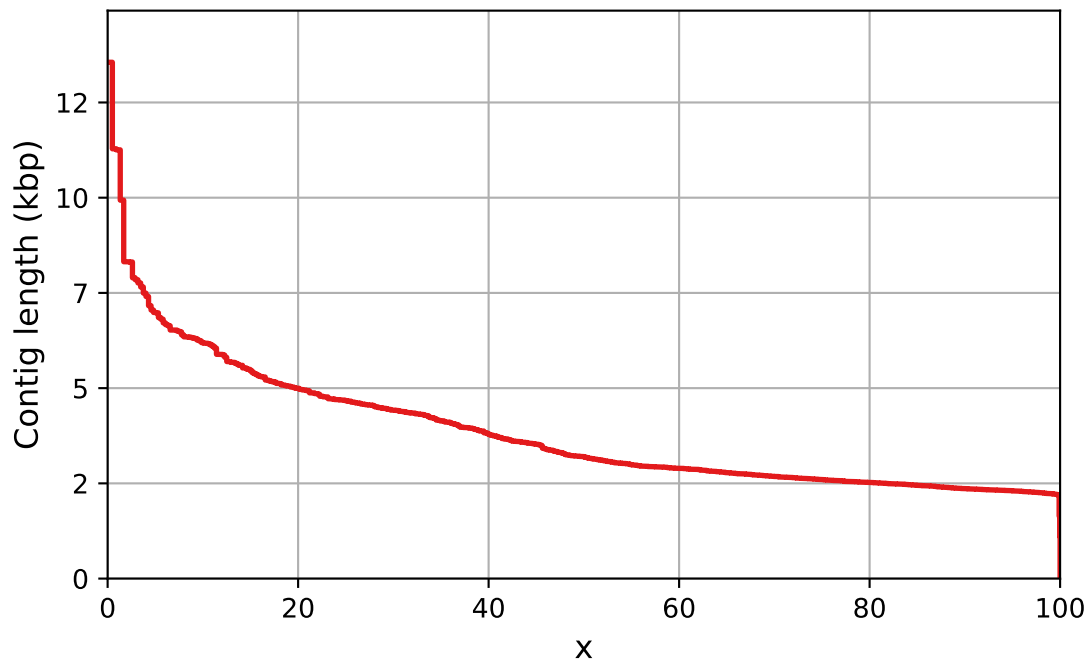


# Report

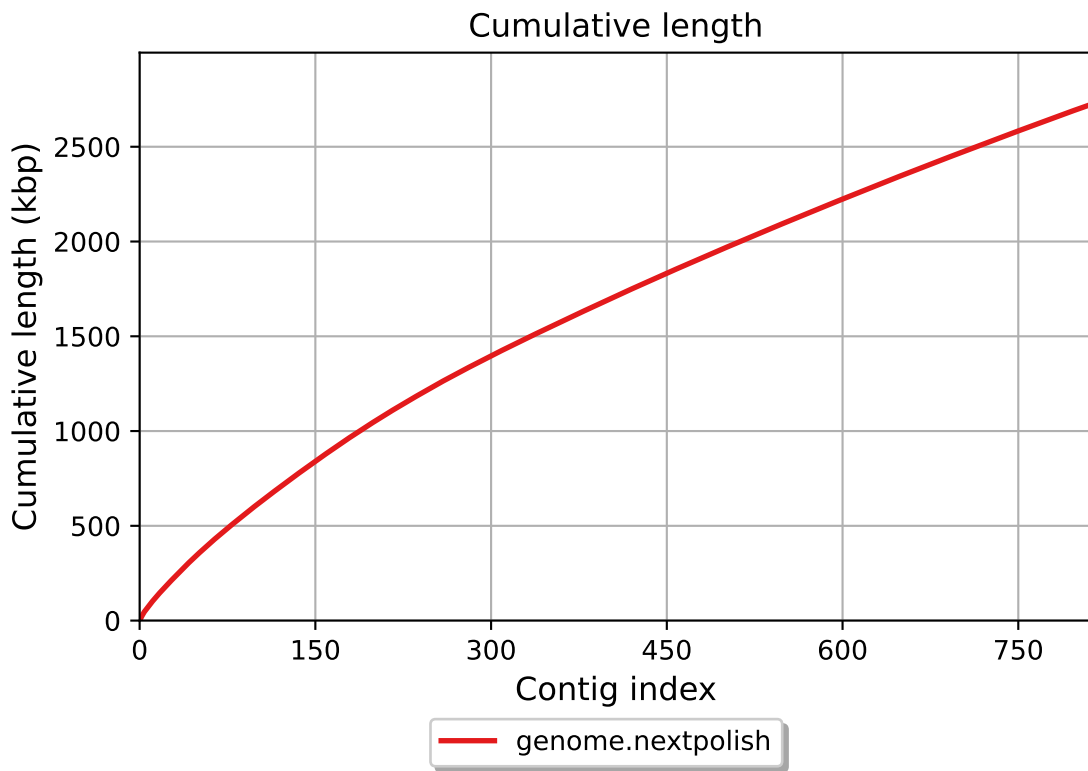
	genome.nextpolish
# contigs (>= 0 bp)	813
# contigs (>= 1000 bp)	813
# contigs (>= 5000 bp)	86
# contigs (>= 10000 bp)	3
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2724133
Total length (>= 1000 bp)	2724133
Total length (>= 5000 bp)	542636
Total length (>= 10000 bp)	36095
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	813
Largest contig	13555
Total length	2724133
GC (%)	50.76
N50	3199
N90	2363
auN	3871.7
L50	290
L90	694
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 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).

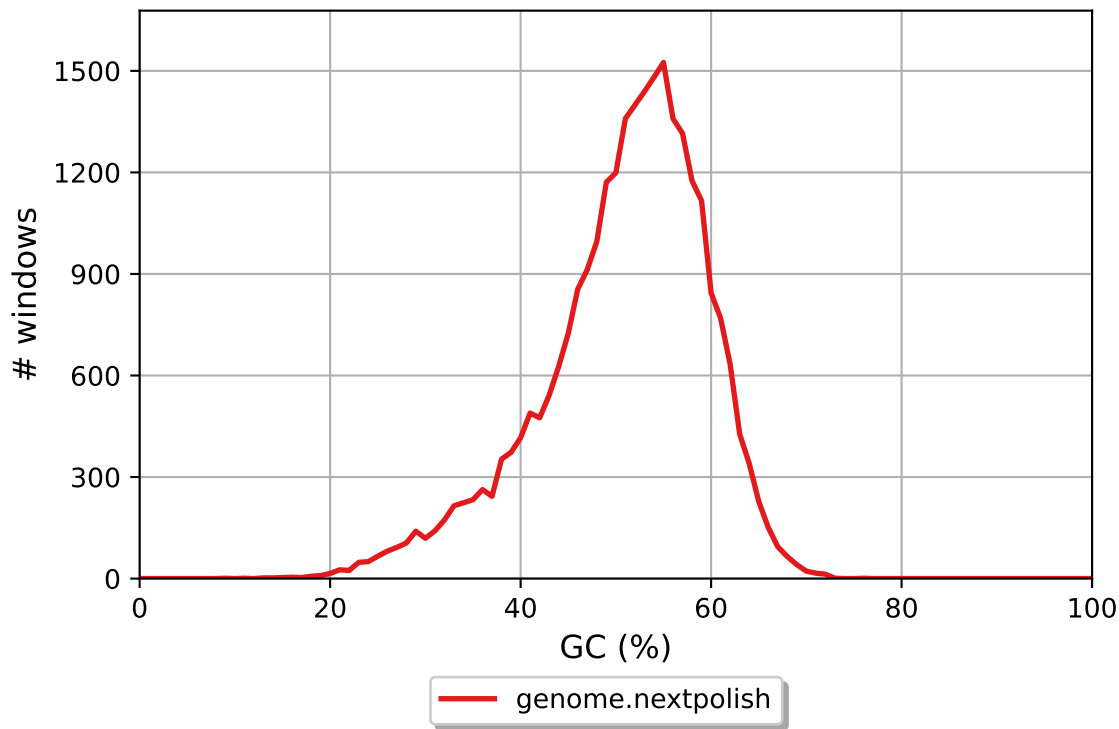
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

