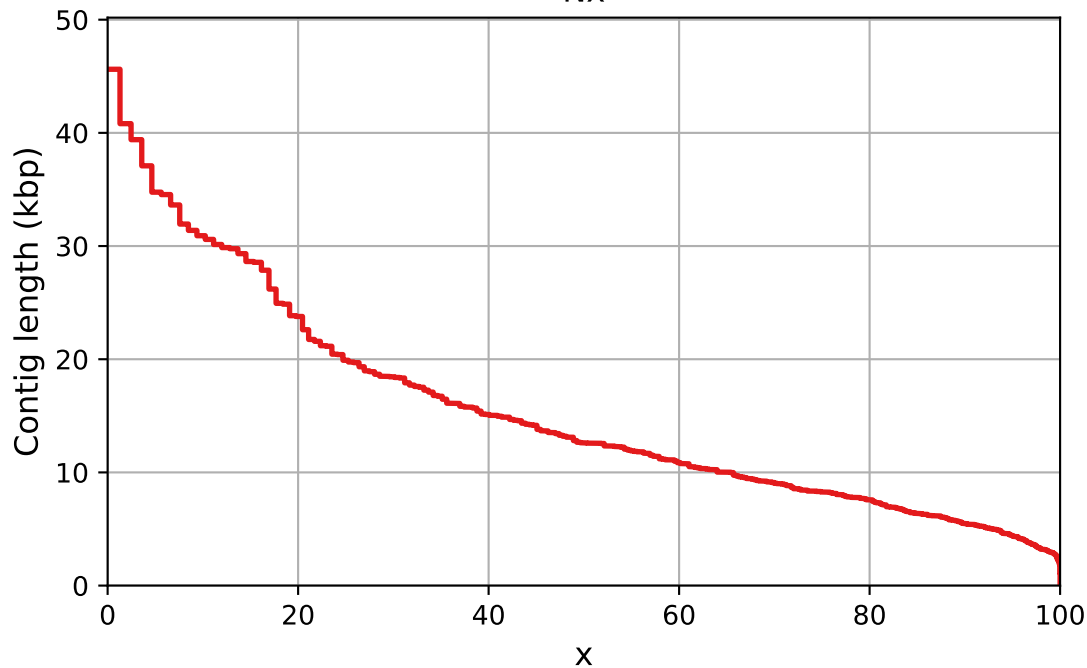


# Report

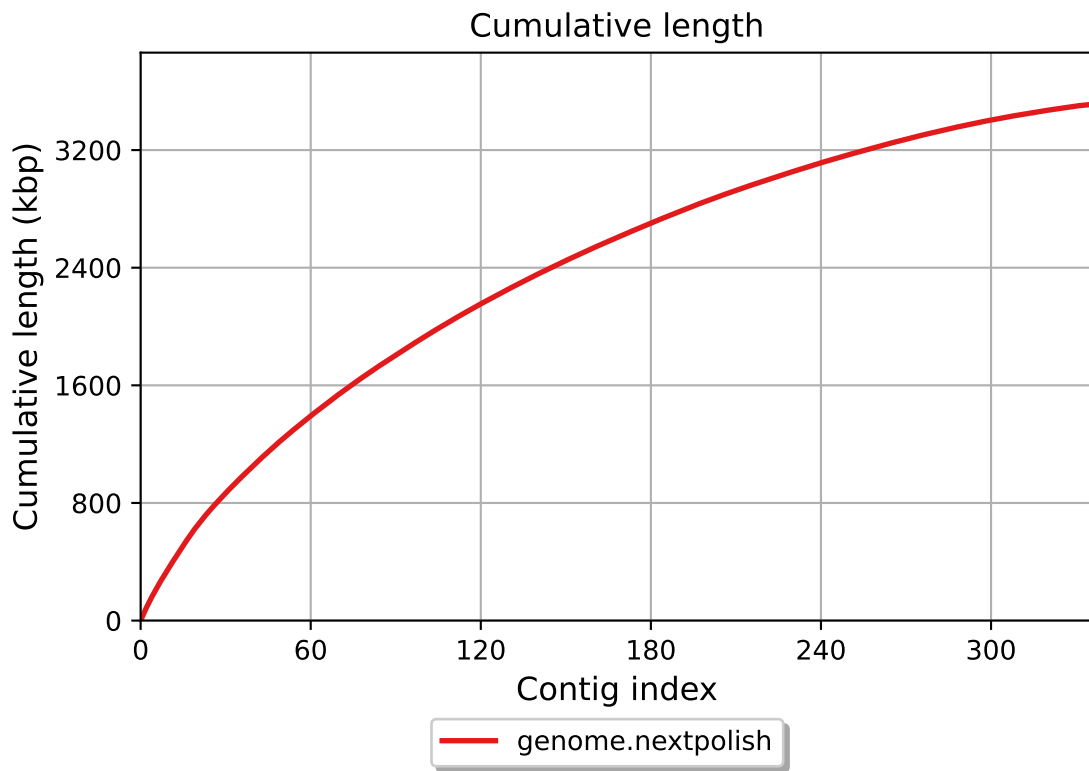
	genome.nextpolish
# contigs (>= 0 bp)	336
# contigs (>= 1000 bp)	336
# contigs (>= 5000 bp)	268
# contigs (>= 10000 bp)	134
# contigs (>= 25000 bp)	19
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	3511529
Total length (>= 1000 bp)	3511529
Total length (>= 5000 bp)	3264036
Total length (>= 10000 bp)	2297132
Total length (>= 25000 bp)	621036
Total length (>= 50000 bp)	0
# contigs	336
Largest contig	45619
Total length	3511529
GC (%)	56.03
N50	12599
N90	5477
auN	15652.6
L50	87
L90	249
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	5 + 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).

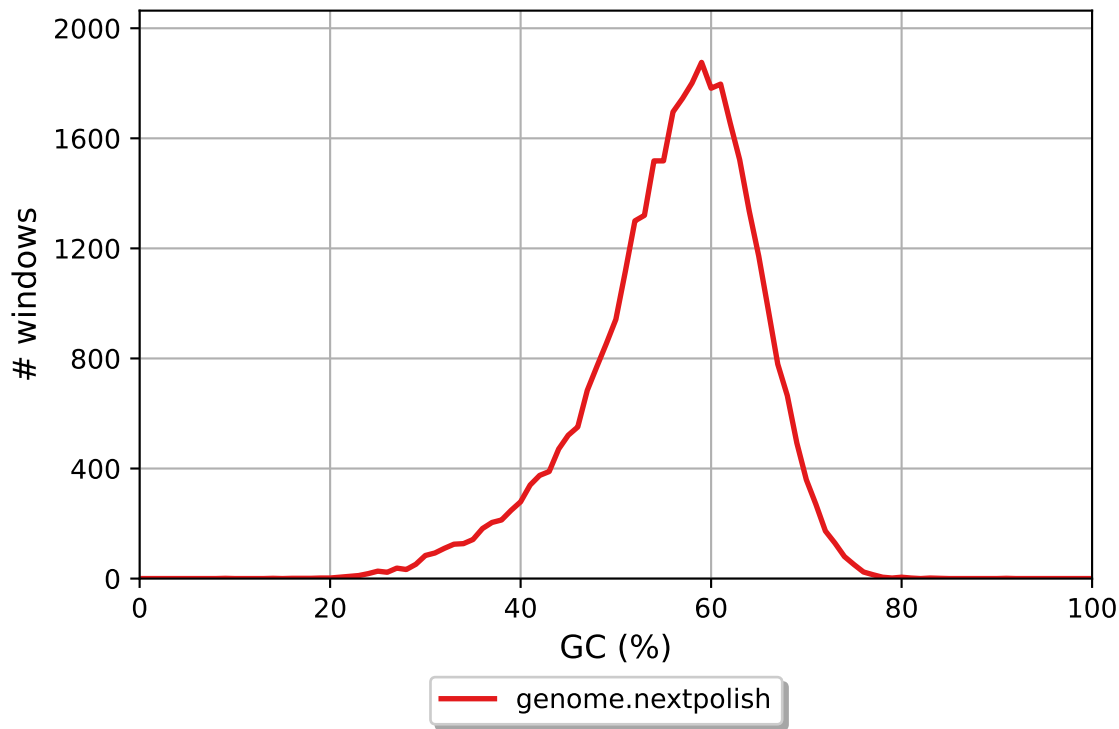
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

