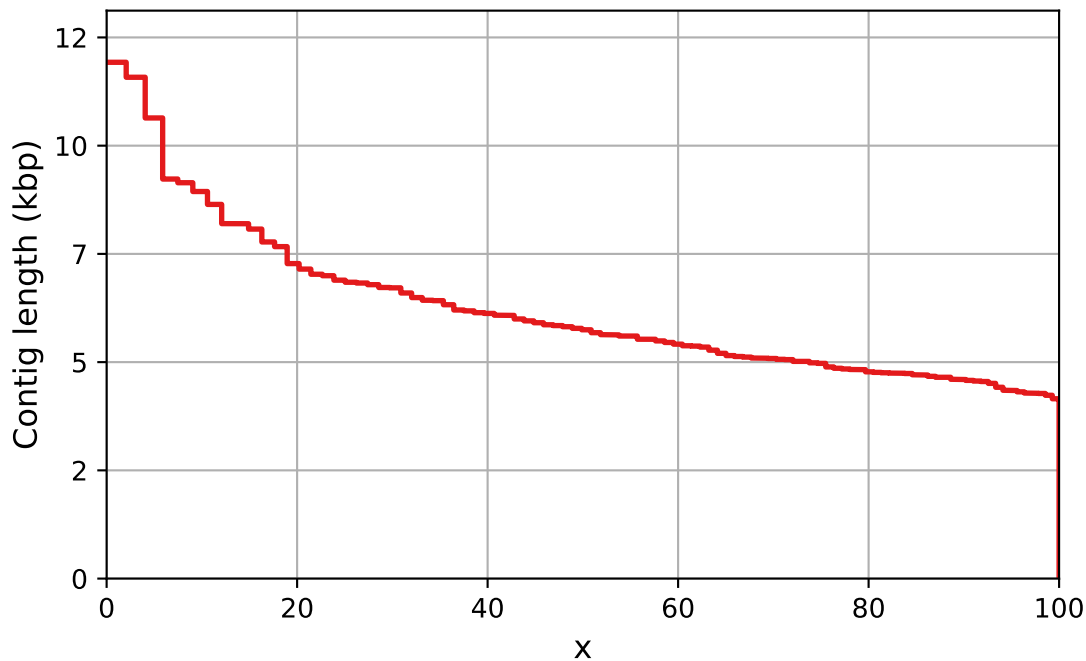


# Report

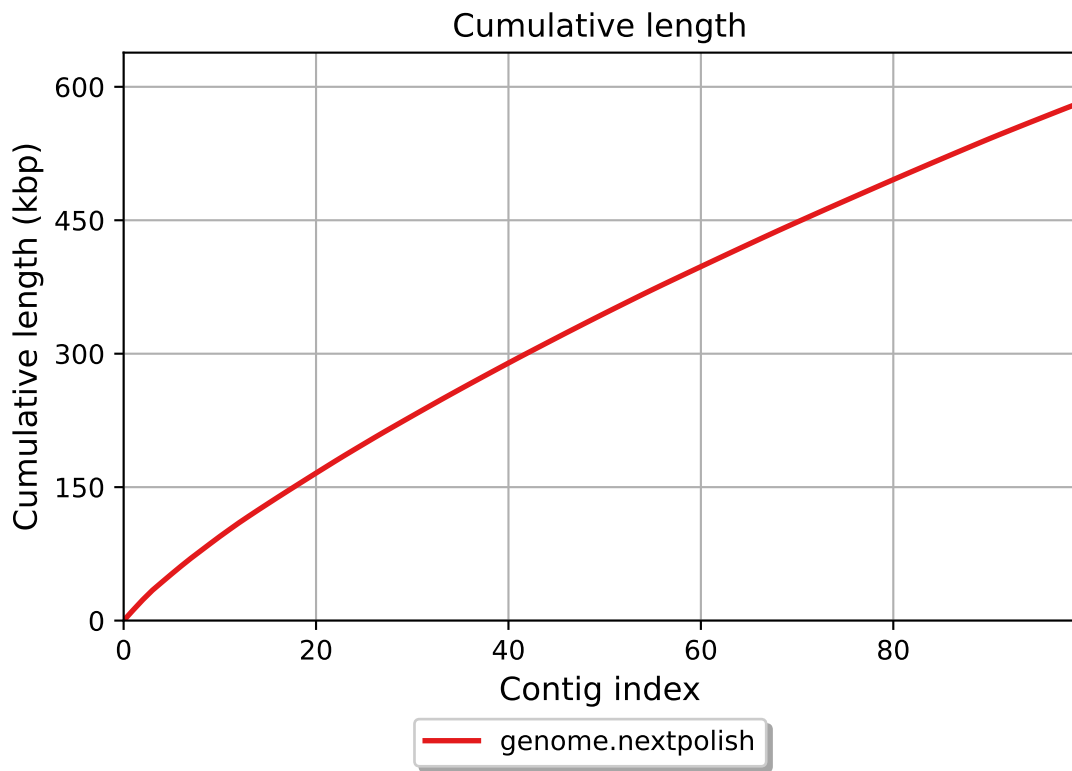
	genome.nextpolish
# contigs (>= 0 bp)	99
# contigs (>= 1000 bp)	99
# contigs (>= 5000 bp)	66
# contigs (>= 10000 bp)	3
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	580379
Total length (>= 1000 bp)	580379
Total length (>= 5000 bp)	428216
Total length (>= 10000 bp)	34146
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	99
Largest contig	11926
Total length	580379
GC (%)	46.72
N50	5747
N90	4598
auN	6259.0
L50	41
L90	86
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 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).

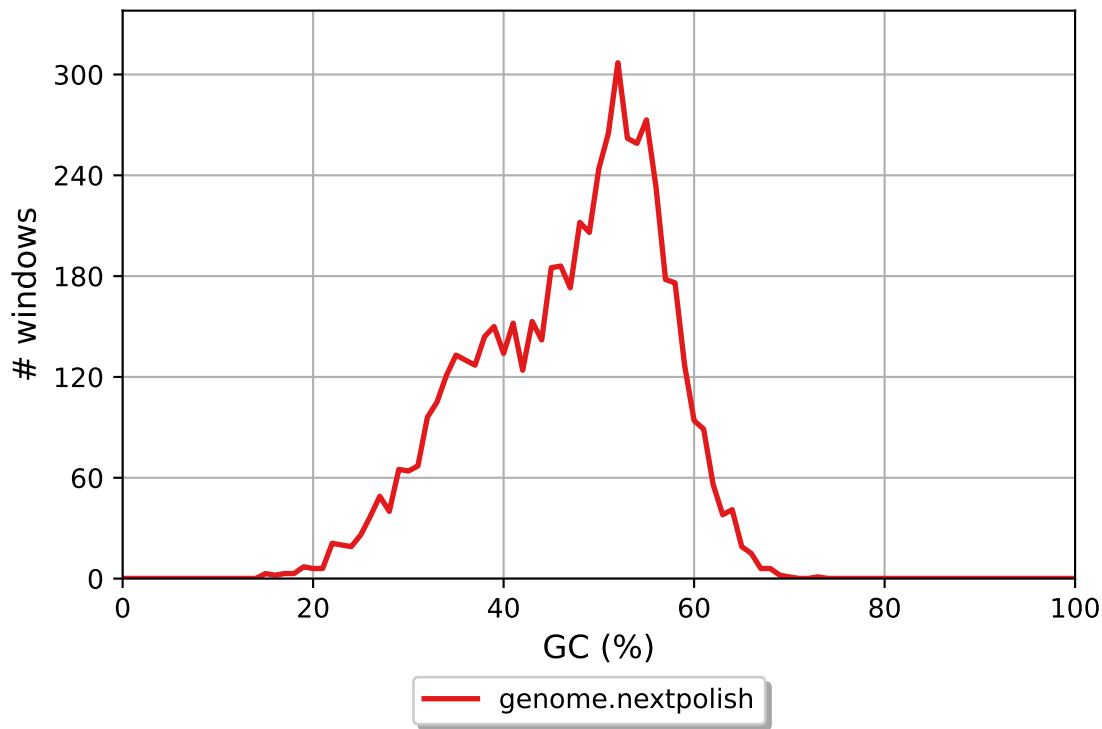
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

