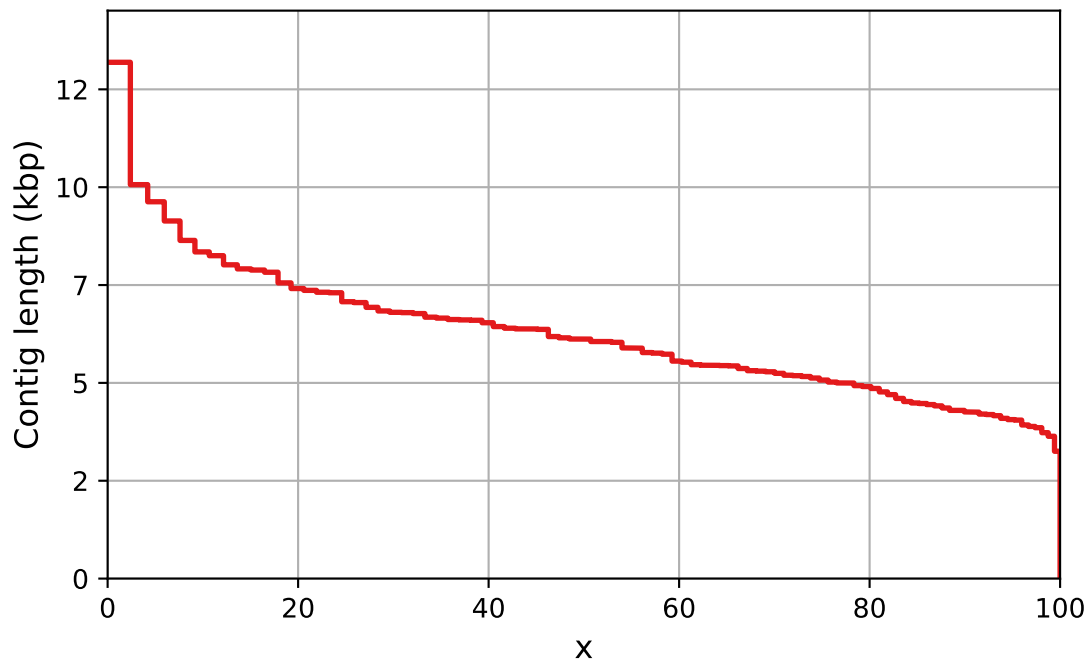


# Report

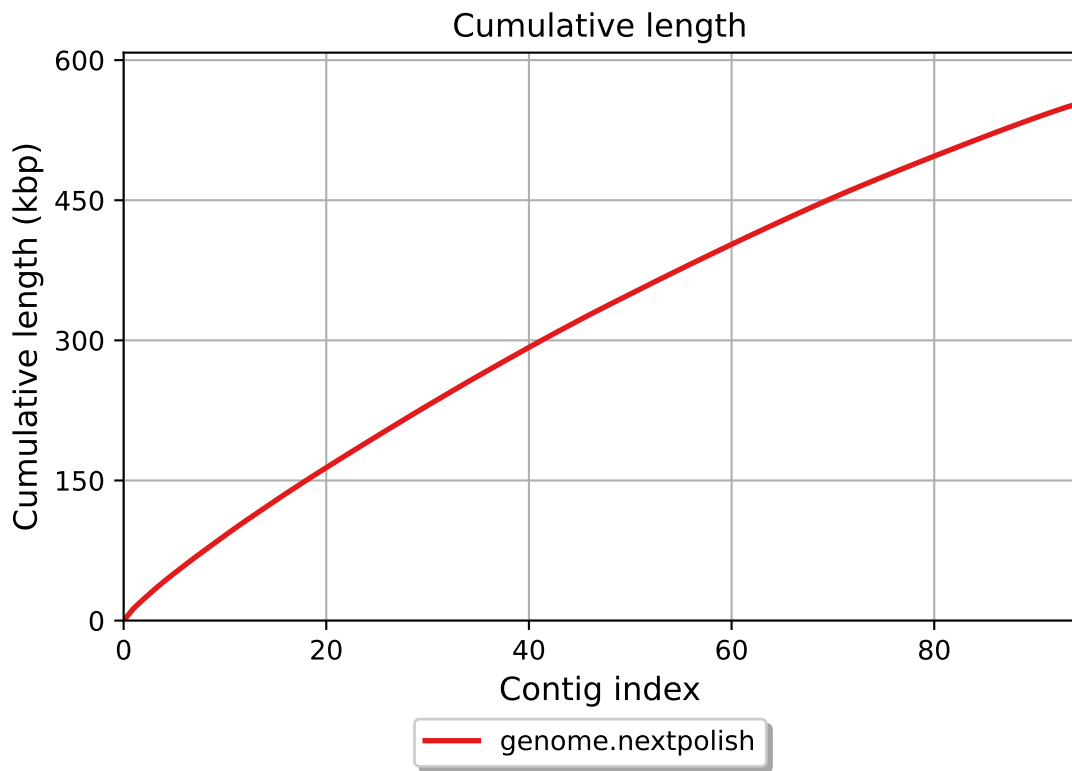
	genome.nextpolish
# contigs (>= 0 bp)	94
# contigs (>= 1000 bp)	94
# contigs (>= 5000 bp)	64
# contigs (>= 10000 bp)	2
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	552750
Total length (>= 1000 bp)	552750
Total length (>= 5000 bp)	423136
Total length (>= 10000 bp)	23256
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	94
Largest contig	13191
Total length	552750
GC (%)	51.21
N50	6119
N90	4259
auN	6313.9
L50	38
L90	81
# 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).

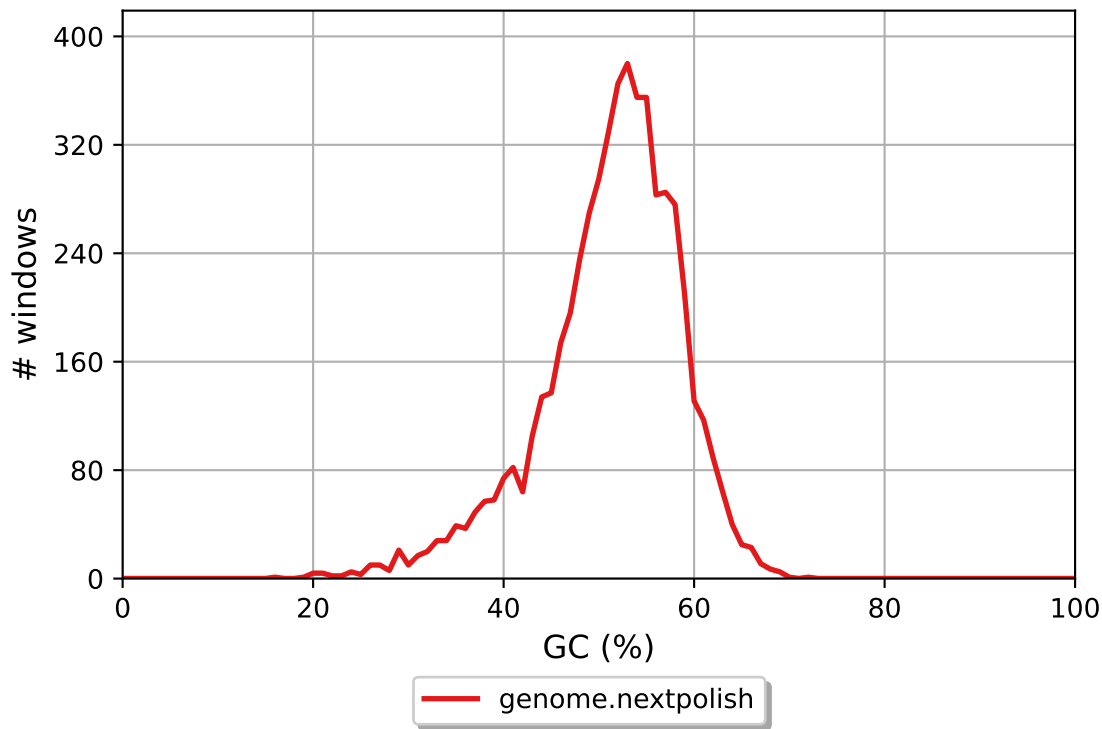
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

