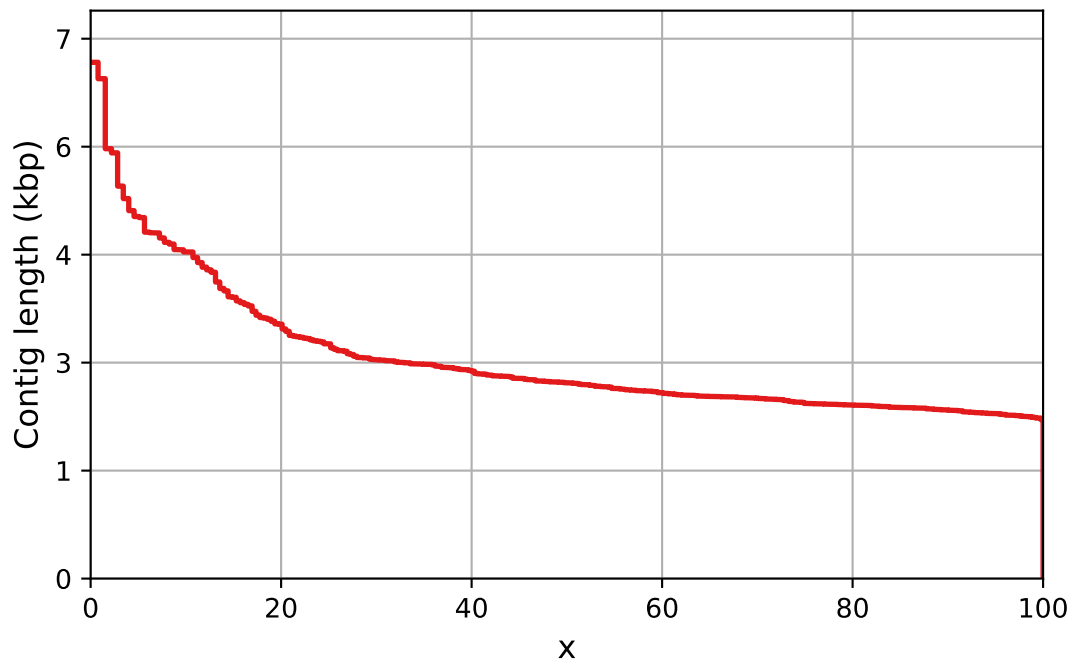


# Report

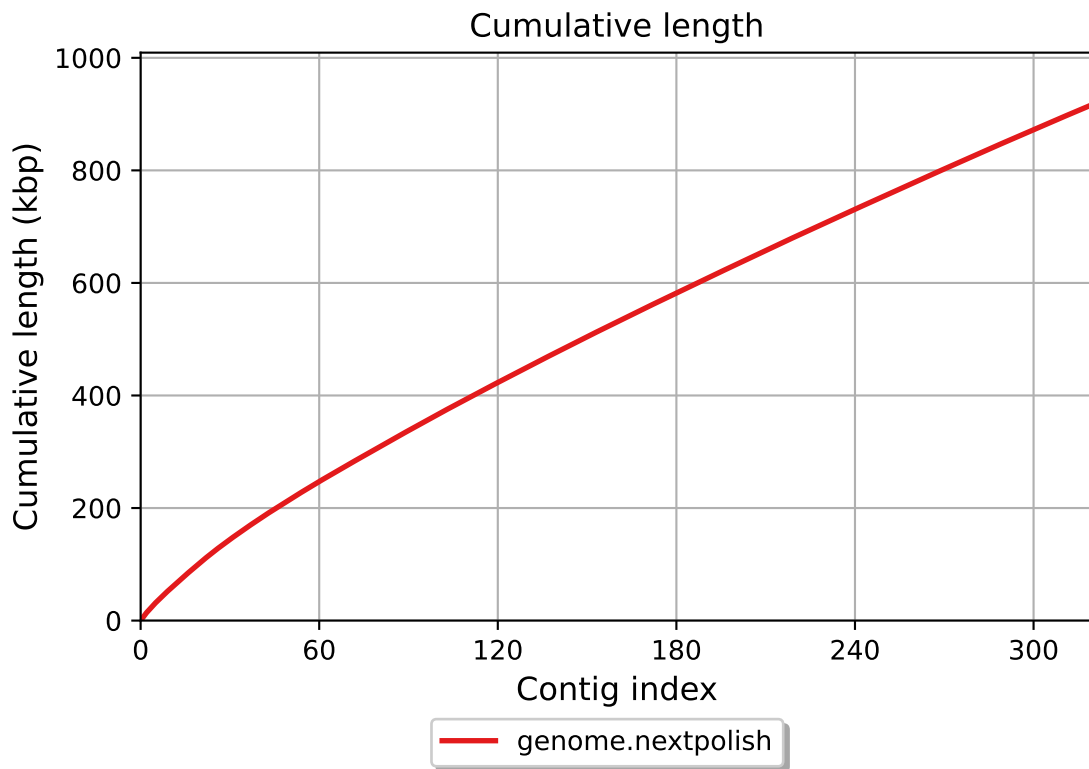
	genome.nextpolish
# contigs (>= 0 bp)	320
# contigs (>= 1000 bp)	320
# contigs (>= 5000 bp)	9
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	917476
Total length (>= 1000 bp)	917476
Total length (>= 5000 bp)	51891
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	320
Largest contig	7172
Total length	917476
GC (%)	50.68
N50	2719
N90	2343
auN	3061.9
L50	134
L90	280
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	3 + 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).

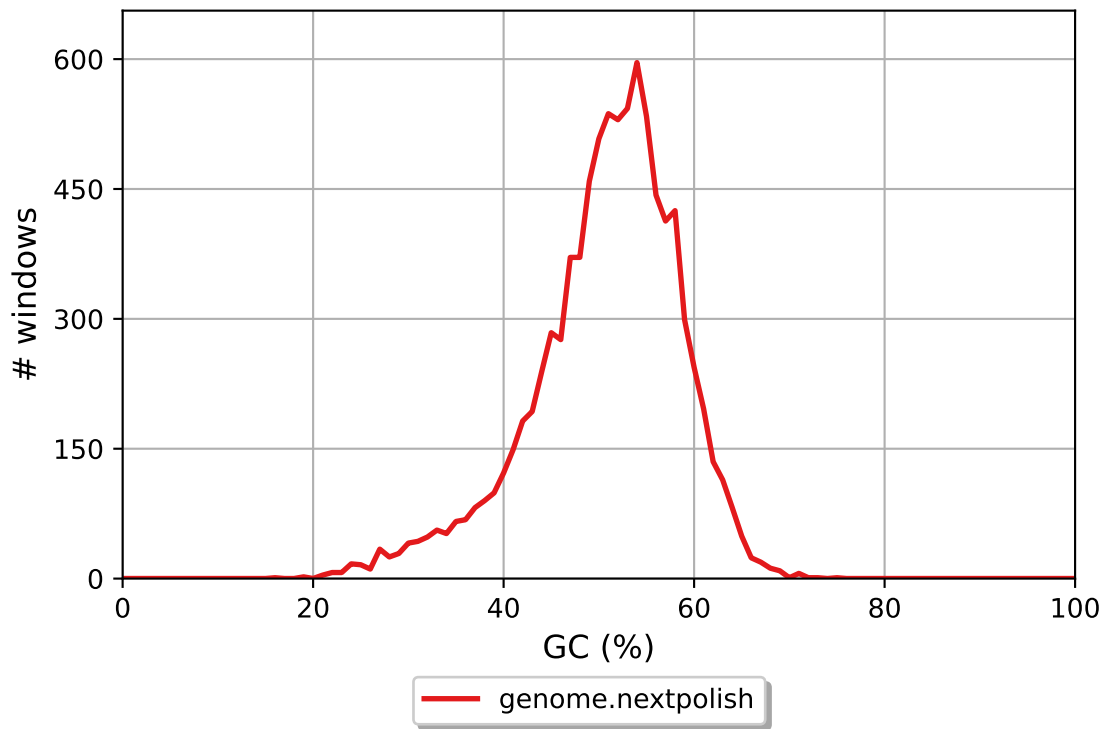
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

