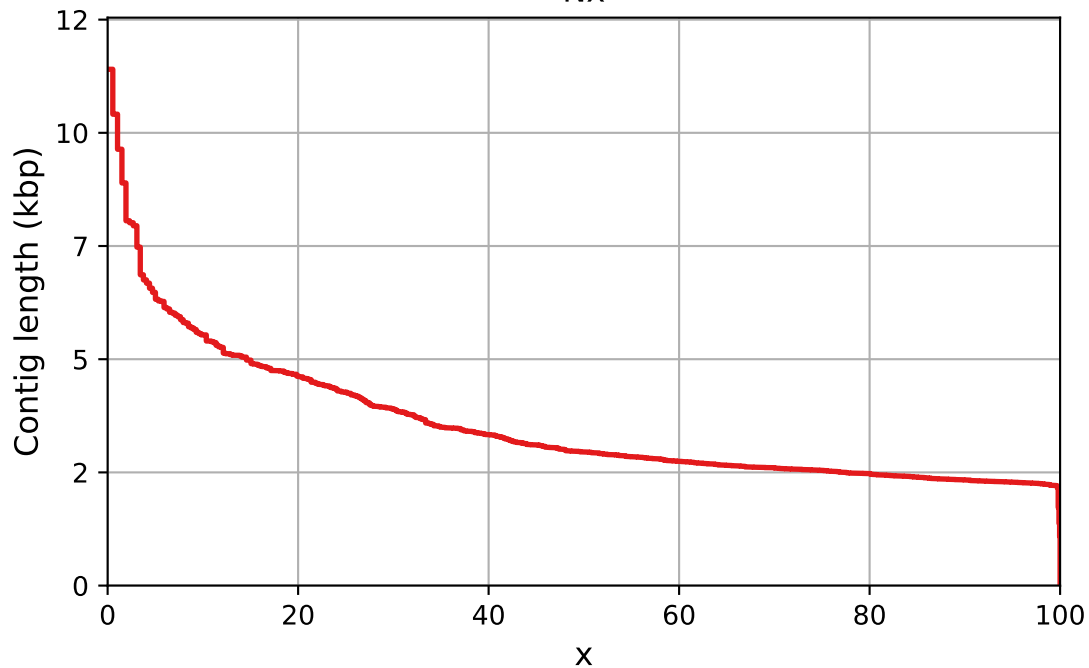


# Report

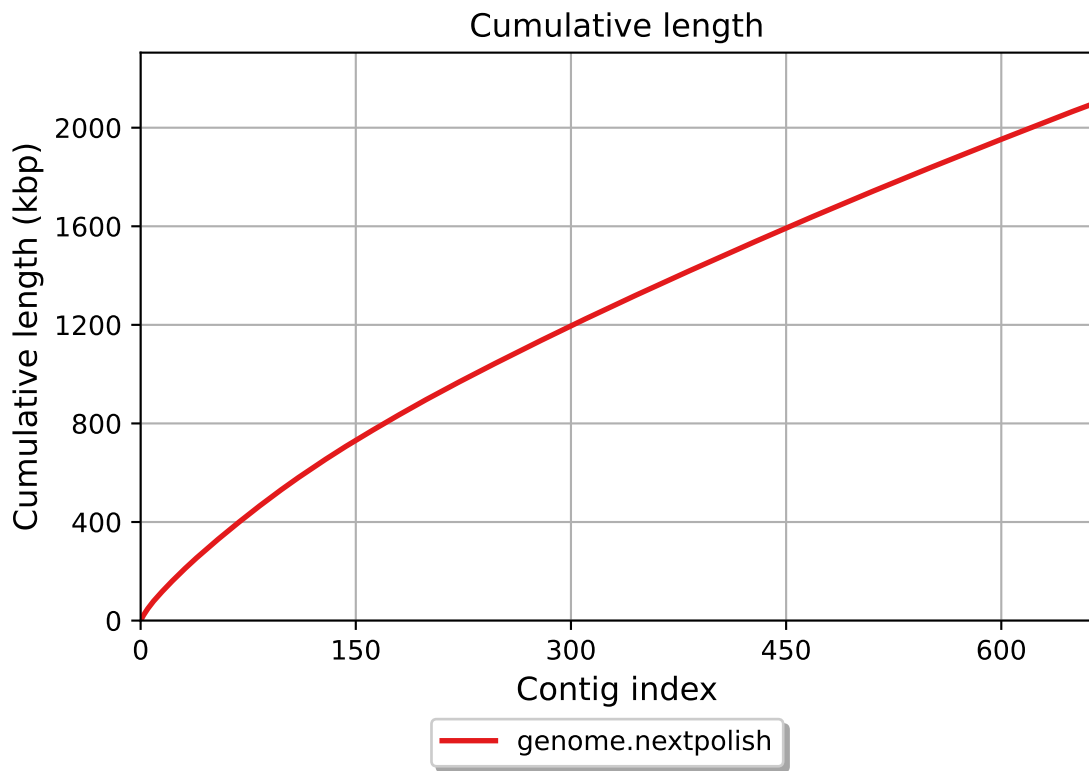
	genome.nextpolish
# contigs (>= 0 bp)	664
# contigs (>= 1000 bp)	664
# contigs (>= 5000 bp)	49
# contigs (>= 10000 bp)	2
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2095086
Total length (>= 1000 bp)	2095086
Total length (>= 5000 bp)	305349
Total length (>= 10000 bp)	21814
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	664
Largest contig	11403
Total length	2095086
GC (%)	50.64
N50	2950
N90	2334
auN	3577.5
L50	249
L90	571
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 2 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).

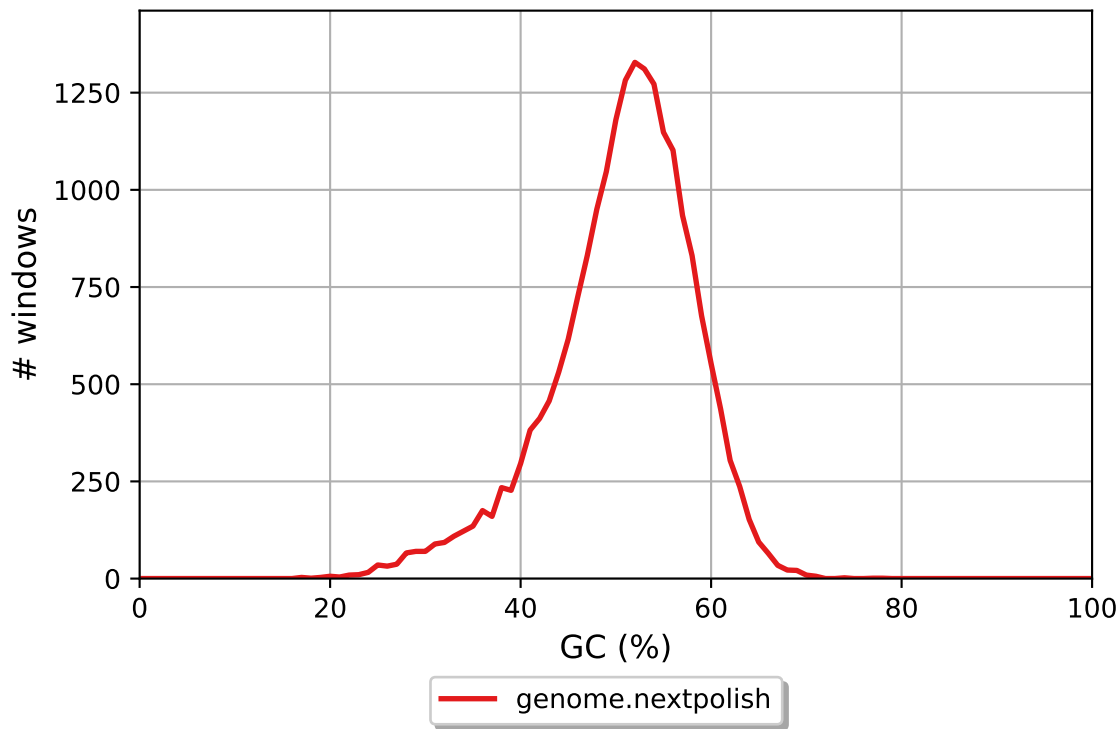
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

