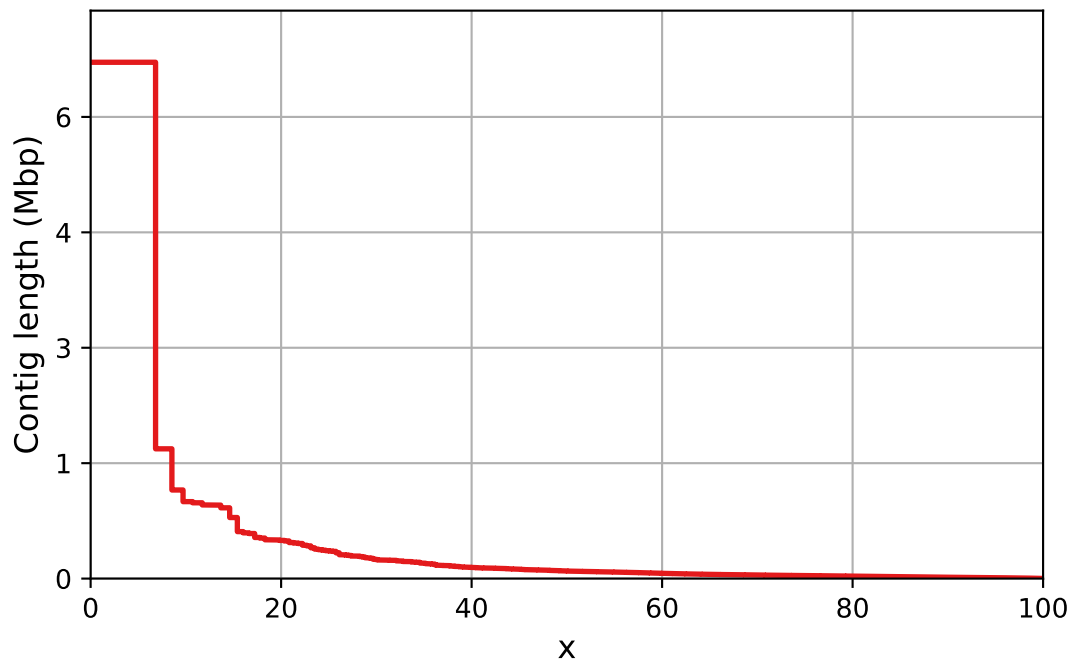


# Report

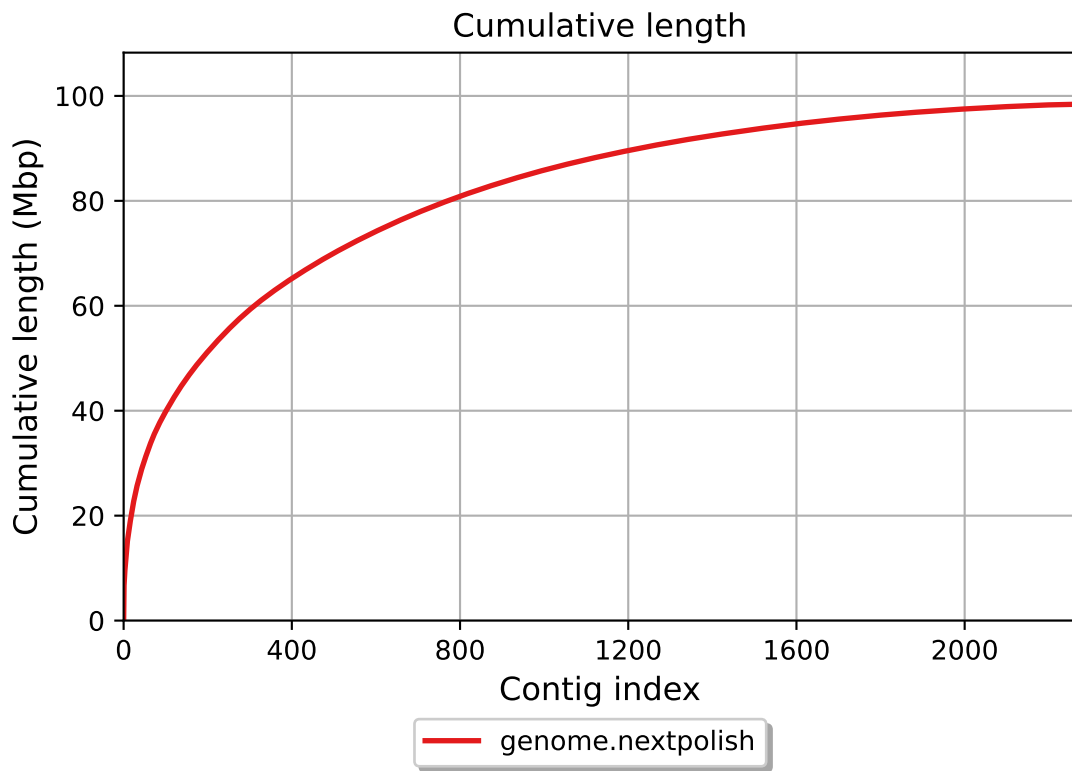
	genome.nextpolish
# contigs (>= 0 bp)	2265
# contigs (>= 1000 bp)	2255
# contigs (>= 5000 bp)	1992
# contigs (>= 10000 bp)	1585
# contigs (>= 25000 bp)	896
# contigs (>= 50000 bp)	430
Total length (>= 0 bp)	98405955
Total length (>= 1000 bp)	98398602
Total length (>= 5000 bp)	97475673
Total length (>= 10000 bp)	94526895
Total length (>= 25000 bp)	83443271
Total length (>= 50000 bp)	66725579
# contigs	2265
Largest contig	6710367
Total length	98405955
GC (%)	52.03
N50	98834
N90	17502
auN	668629.4
L50	179
L90	1140
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	184 + 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).

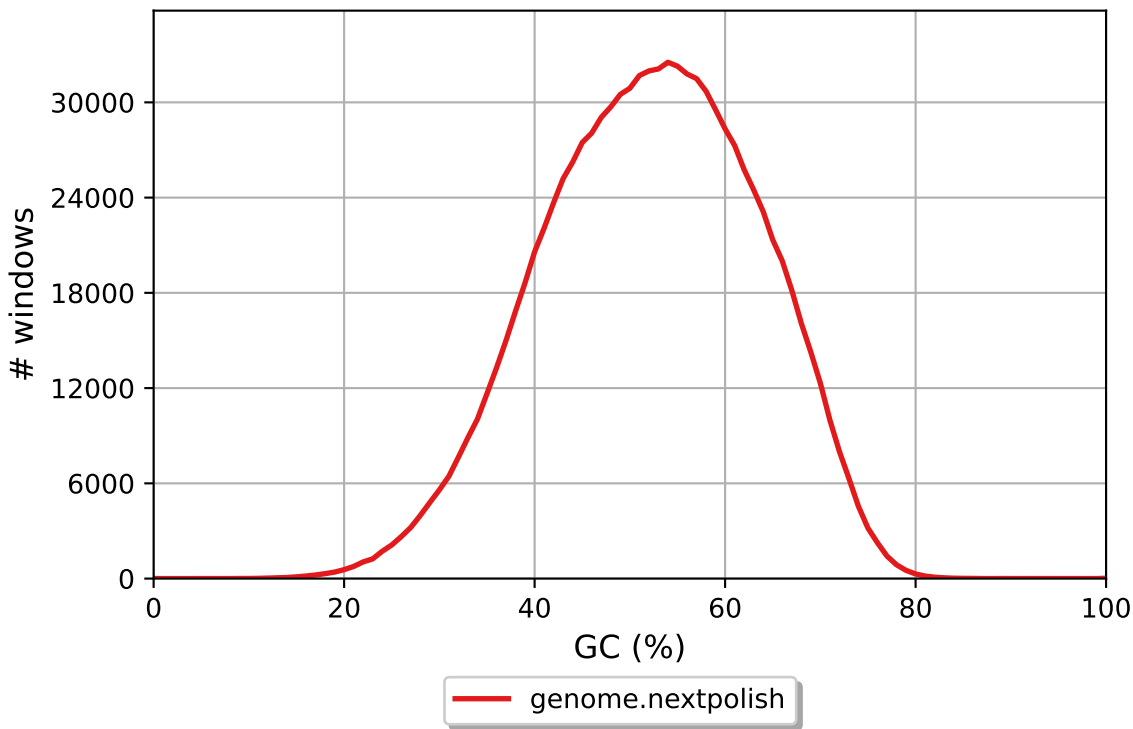
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

