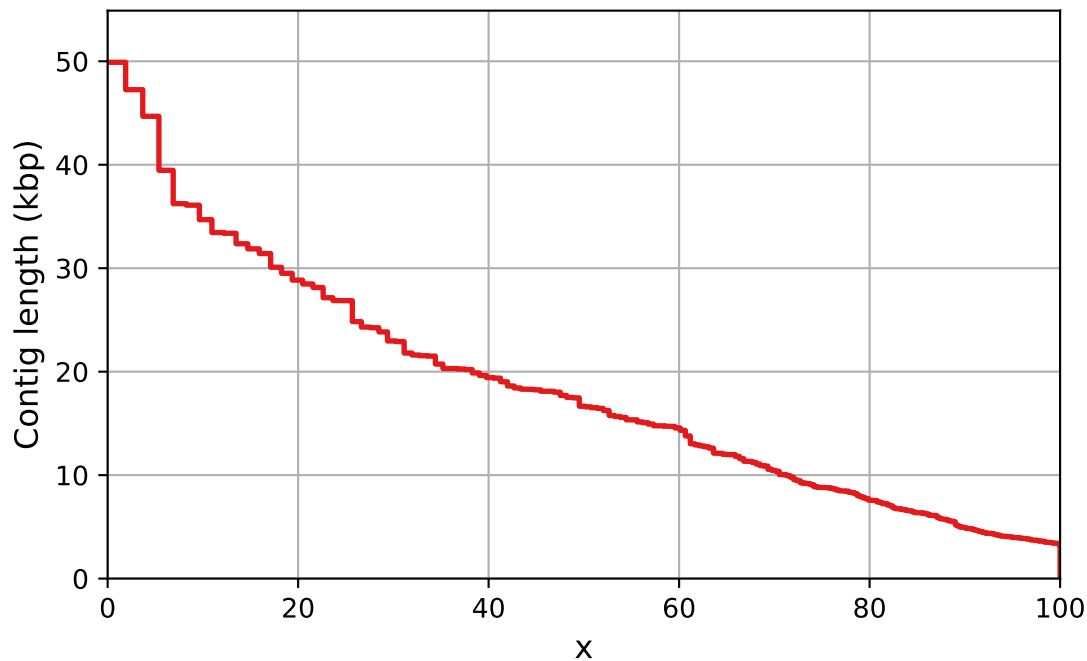


# Report

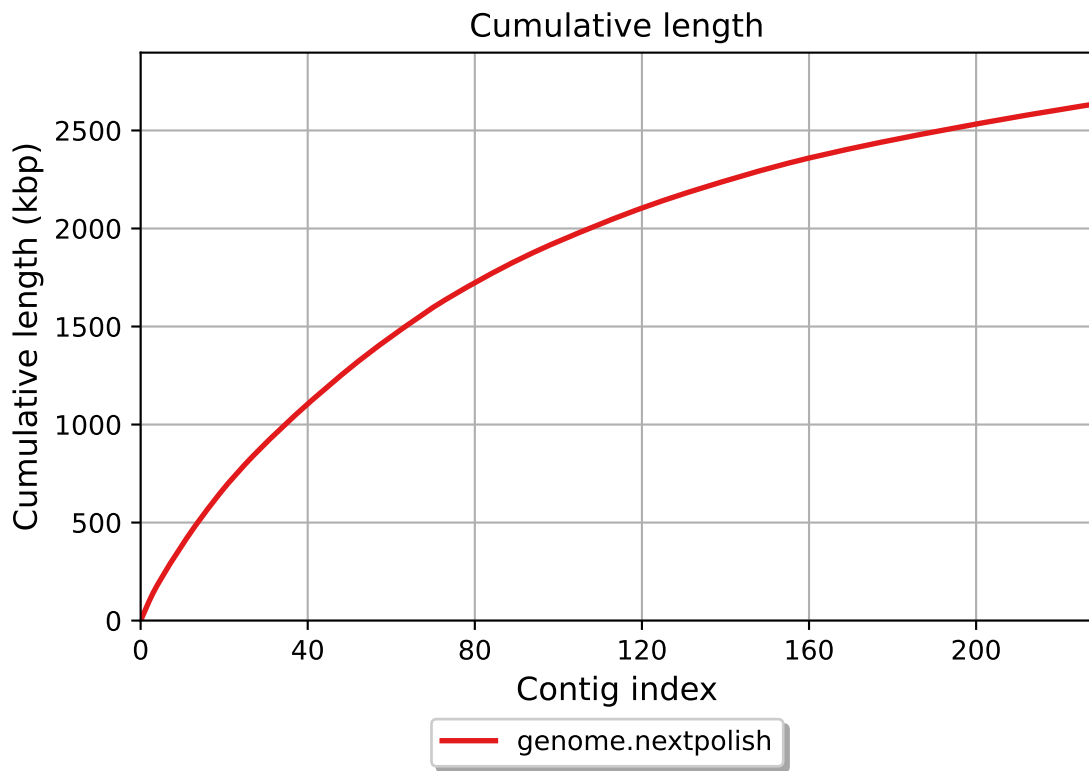
	genome.nextpolish
# contigs (>= 0 bp)	228
# contigs (>= 1000 bp)	228
# contigs (>= 5000 bp)	160
# contigs (>= 10000 bp)	94
# contigs (>= 25000 bp)	20
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2633626
Total length (>= 1000 bp)	2633626
Total length (>= 5000 bp)	2359376
Total length (>= 10000 bp)	1877170
Total length (>= 25000 bp)	676734
Total length (>= 50000 bp)	0
# contigs	228
Largest contig	49904
Total length	2633626
GC (%)	57.76
N50	16657
N90	4909
auN	18679.8
L50	52
L90	163
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 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).

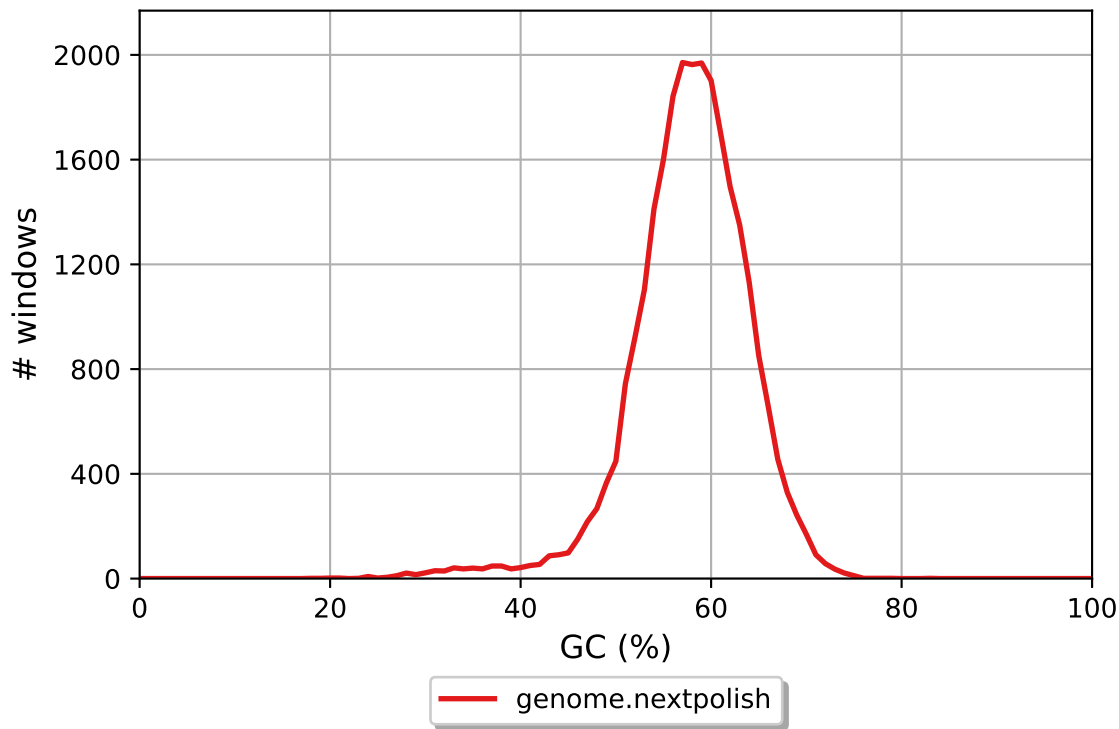
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

