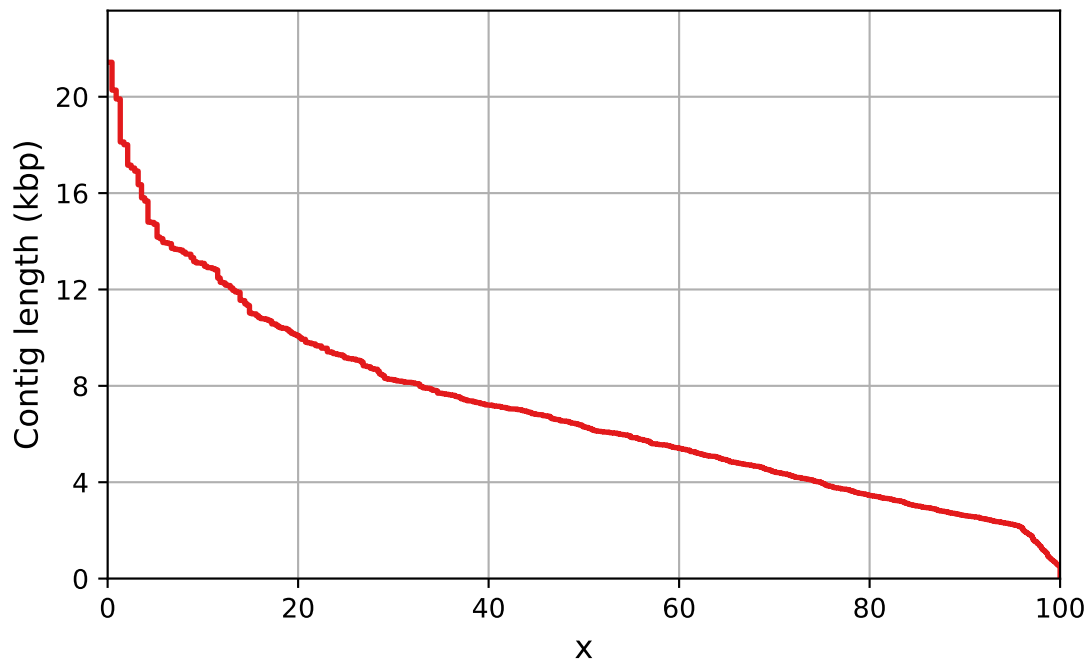


# Report

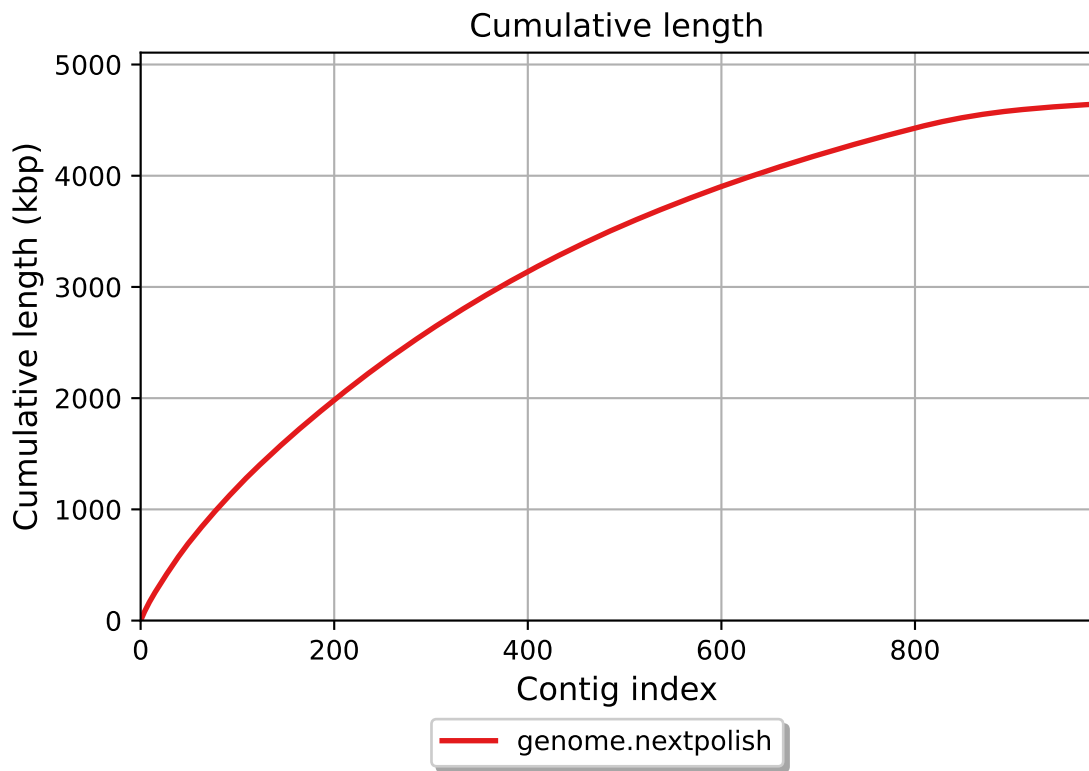
	genome.nextpolish
# contigs (>= 0 bp)	989
# contigs (>= 1000 bp)	895
# contigs (>= 5000 bp)	368
# contigs (>= 10000 bp)	73
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	4643724
Total length (>= 1000 bp)	4580681
Total length (>= 5000 bp)	2983096
Total length (>= 10000 bp)	944344
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	984
Largest contig	21430
Total length	4642567
GC (%)	52.04
N50	6298
N90	2616
auN	7045.3
L50	251
L90	697
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	6 + 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).

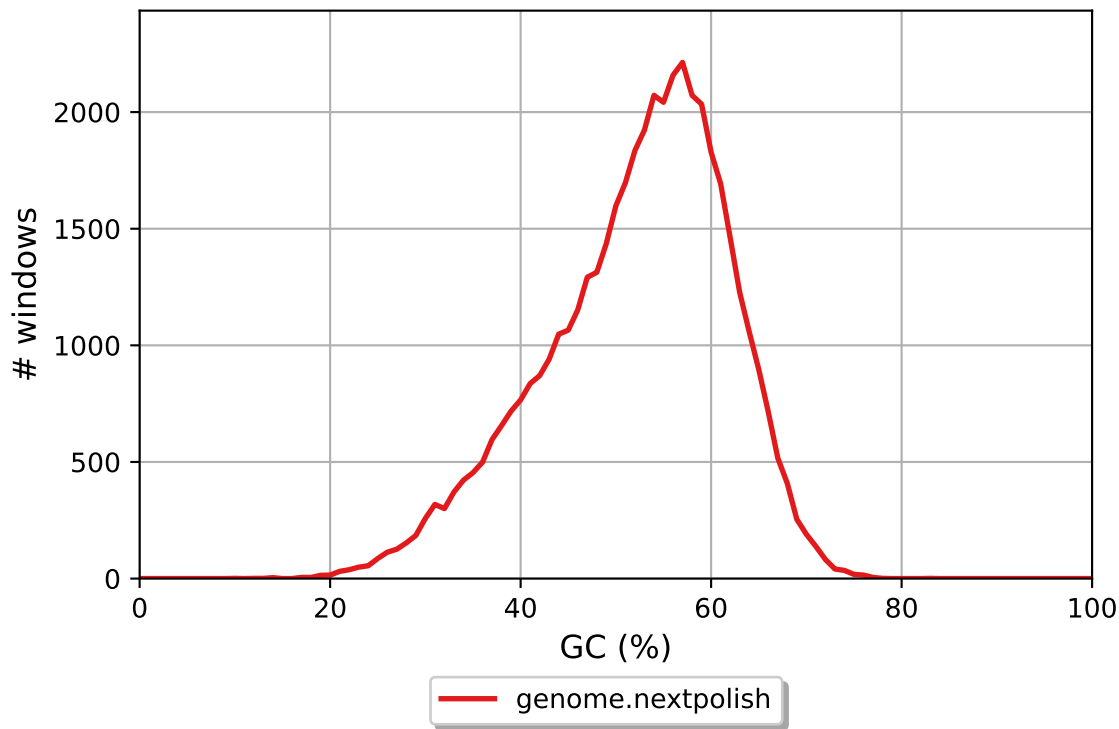
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

