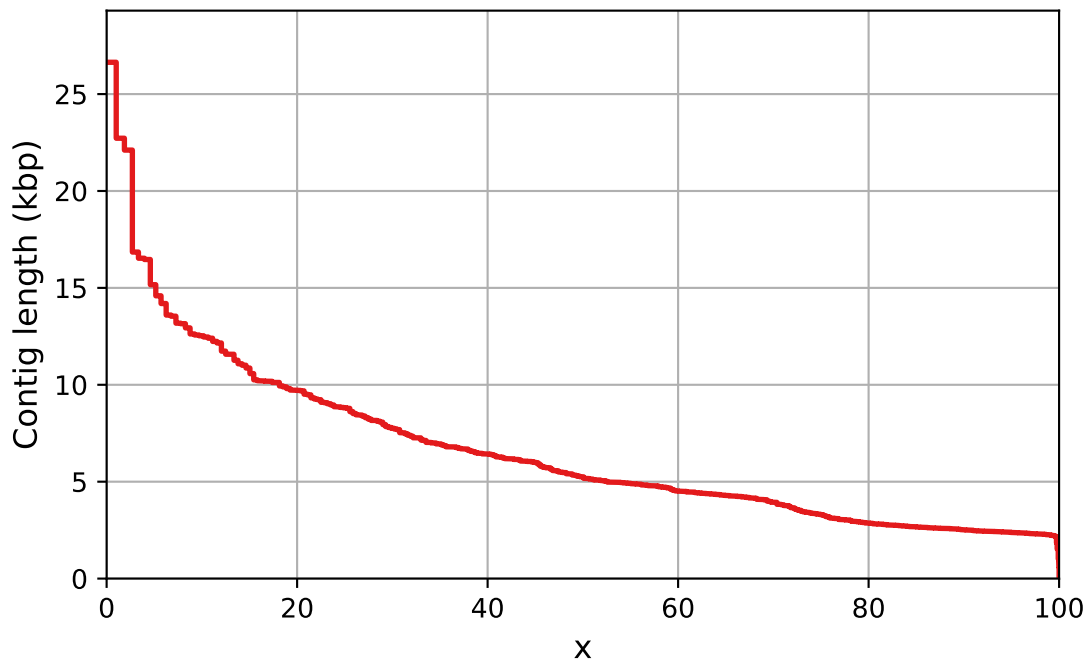


# Report

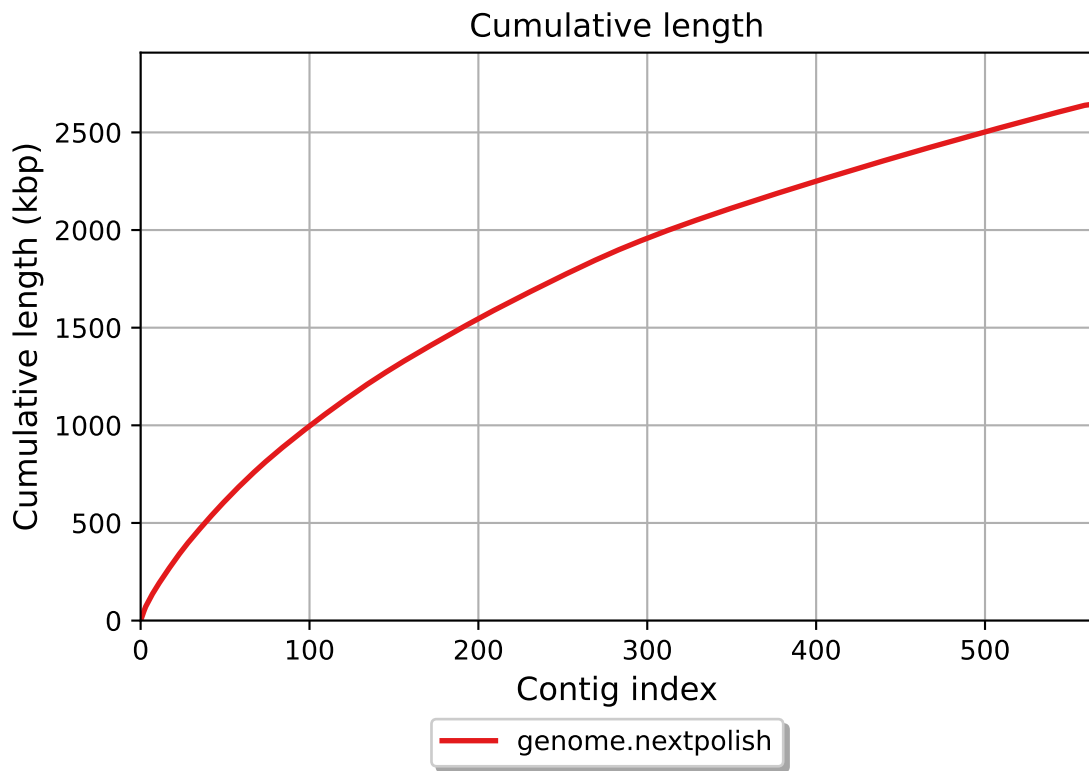
	genome.nextpolish
# contigs (>= 0 bp)	564
# contigs (>= 1000 bp)	562
# contigs (>= 5000 bp)	168
# contigs (>= 10000 bp)	36
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2644119
Total length (>= 1000 bp)	2642753
Total length (>= 5000 bp)	1390471
Total length (>= 10000 bp)	479608
Total length (>= 25000 bp)	26641
Total length (>= 50000 bp)	0
# contigs	564
Largest contig	26641
Total length	2644119
GC (%)	58.91
N50	5249
N90	2513
auN	6697.2
L50	155
L90	450
# 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).

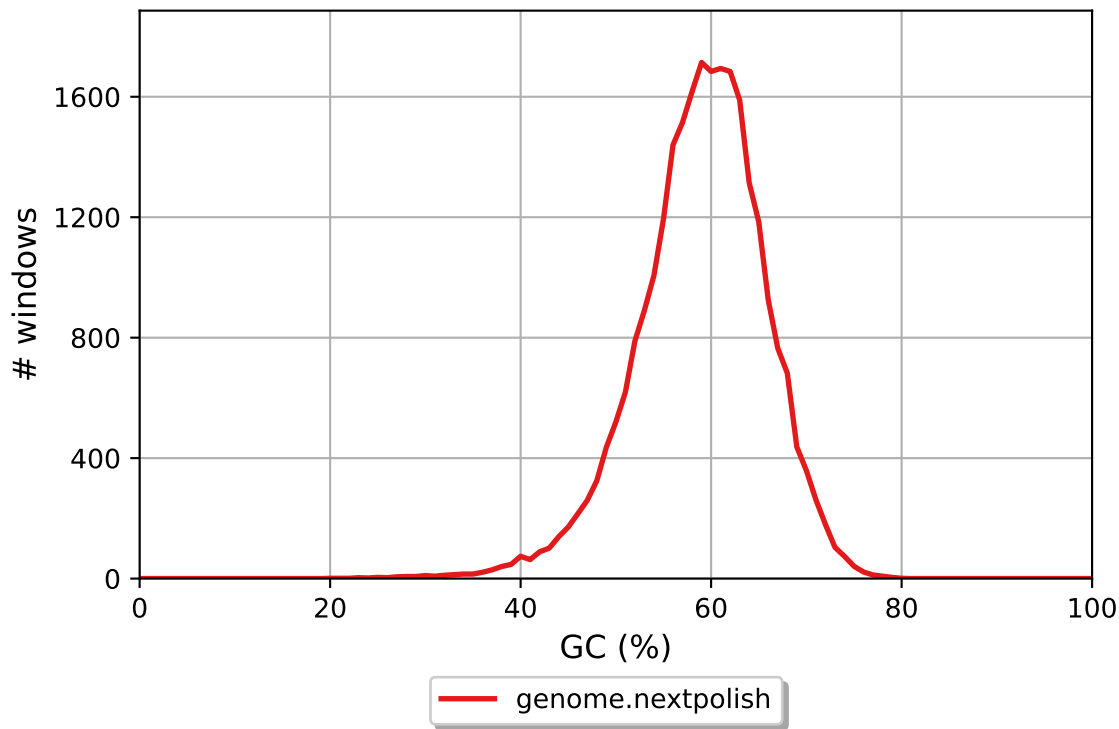
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

