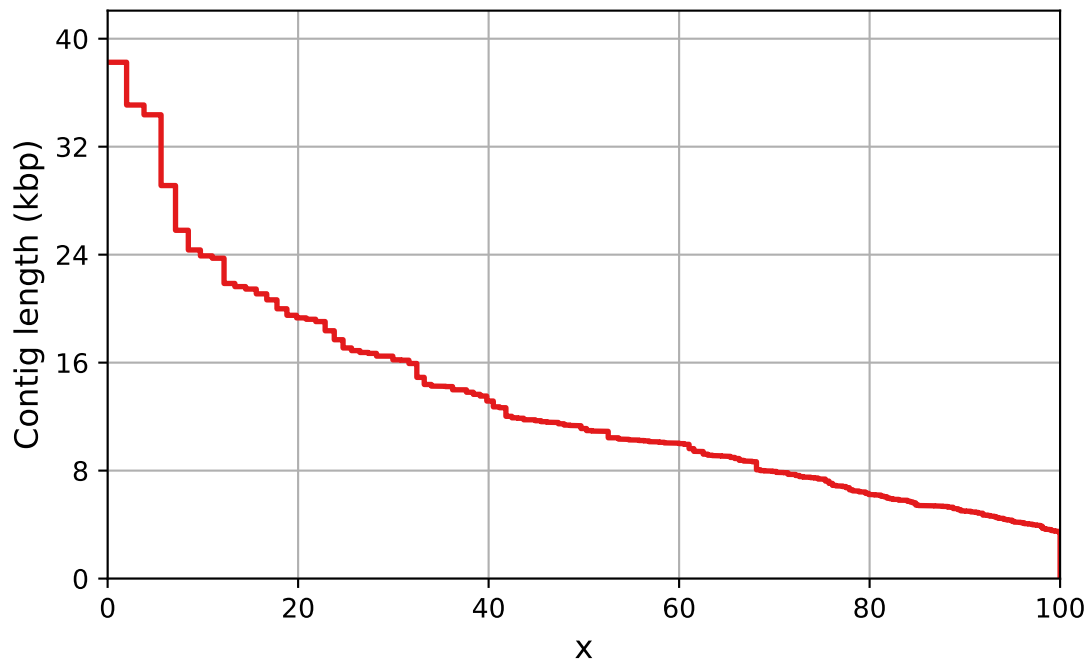


# Report

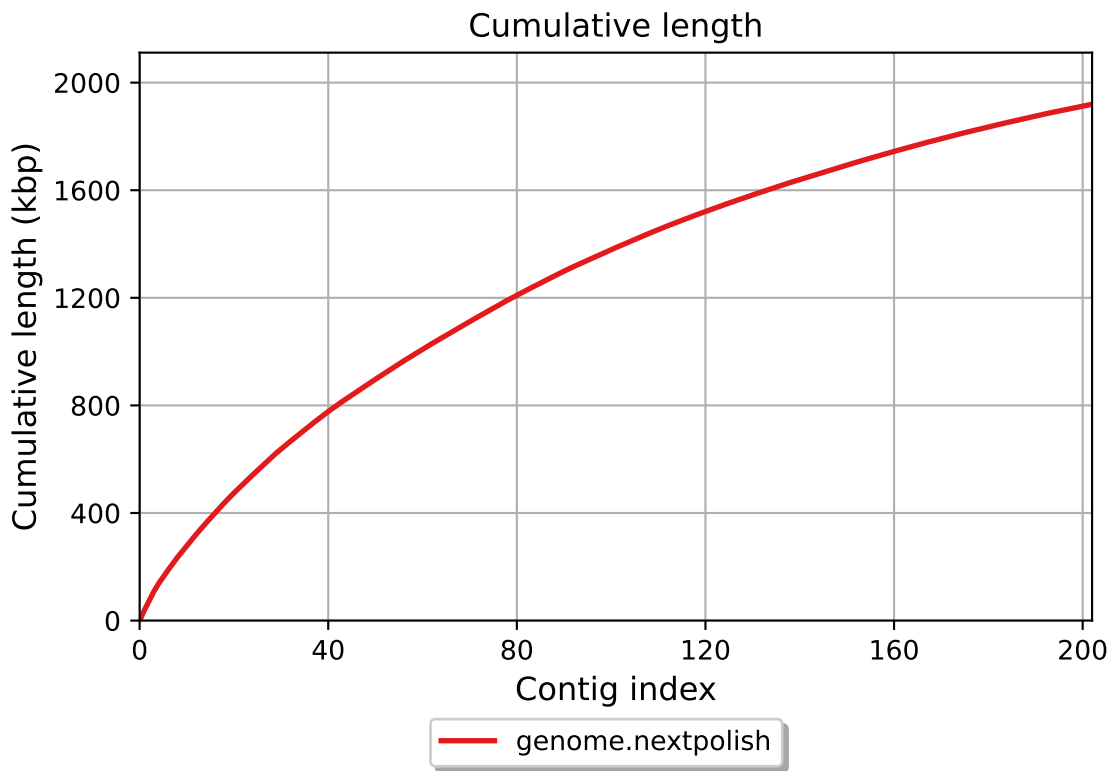
	genome.nextpolish
# contigs (>= 0 bp)	202
# contigs (>= 1000 bp)	202
# contigs (>= 5000 bp)	157
# contigs (>= 10000 bp)	75
# contigs (>= 25000 bp)	5
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1919552
Total length (>= 1000 bp)	1919552
Total length (>= 5000 bp)	1729435
Total length (>= 10000 bp)	1161848
Total length (>= 25000 bp)	162631
Total length (>= 50000 bp)	0
# contigs	202
Largest contig	38254
Total length	1919552
GC (%)	57.92
N50	11117
N90	5004
auN	13387.2
L50	56
L90	157
# 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).

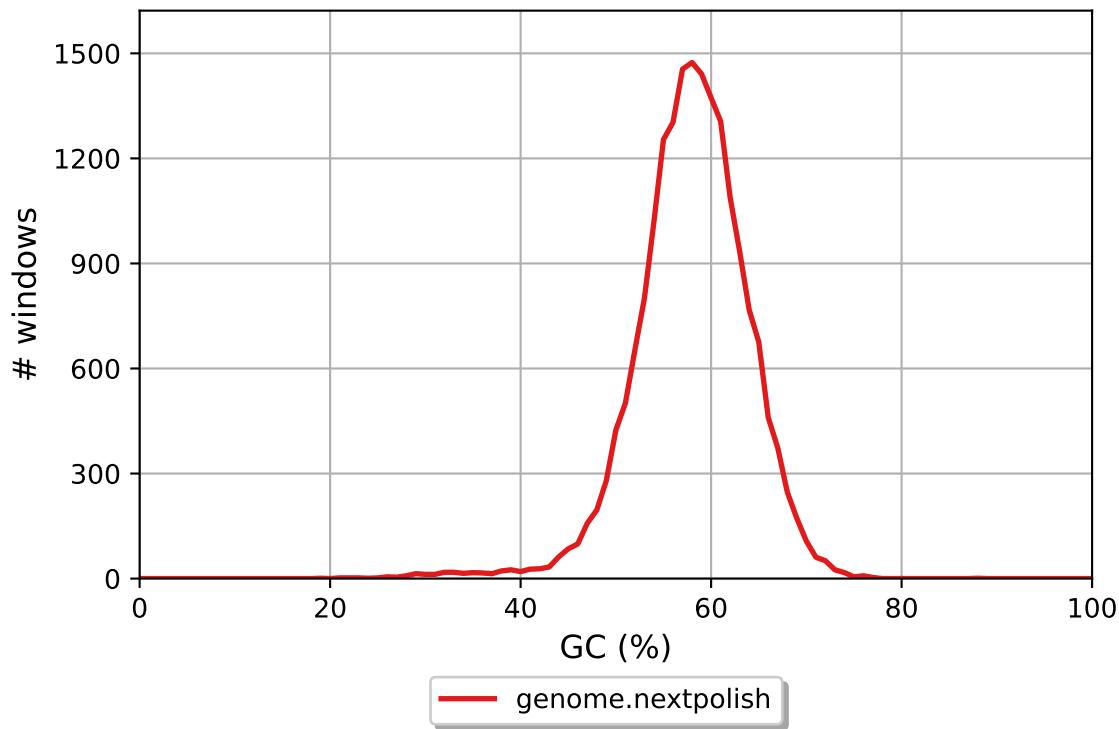
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

