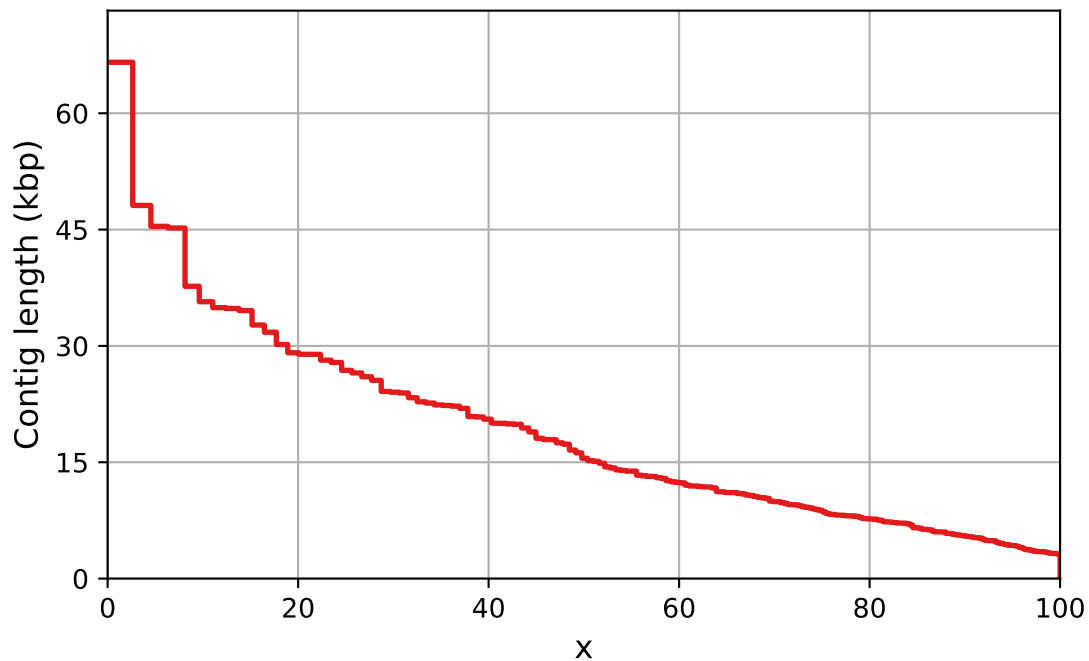


# Report

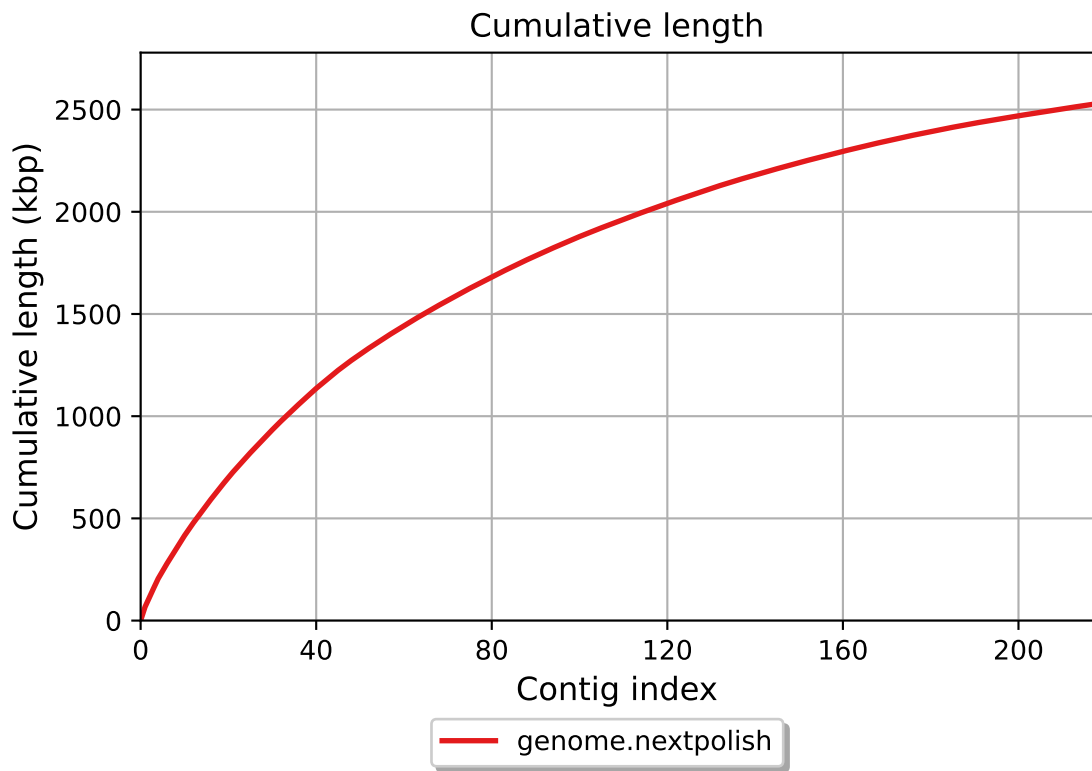
	genome.nextpolish
# contigs (>= 0 bp)	217
# contigs (>= 1000 bp)	217
# contigs (>= 5000 bp)	166
# contigs (>= 10000 bp)	87
# contigs (>= 25000 bp)	21
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	2526019
Total length (>= 1000 bp)	2526019
Total length (>= 5000 bp)	2326824
Total length (>= 10000 bp)	1755051
Total length (>= 25000 bp)	725593
Total length (>= 50000 bp)	66573
# contigs	217
Largest contig	66573
Total length	2526019
GC (%)	59.48
N50	15515
N90	5533
auN	19557.4
L50	48
L90	156
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 1 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).

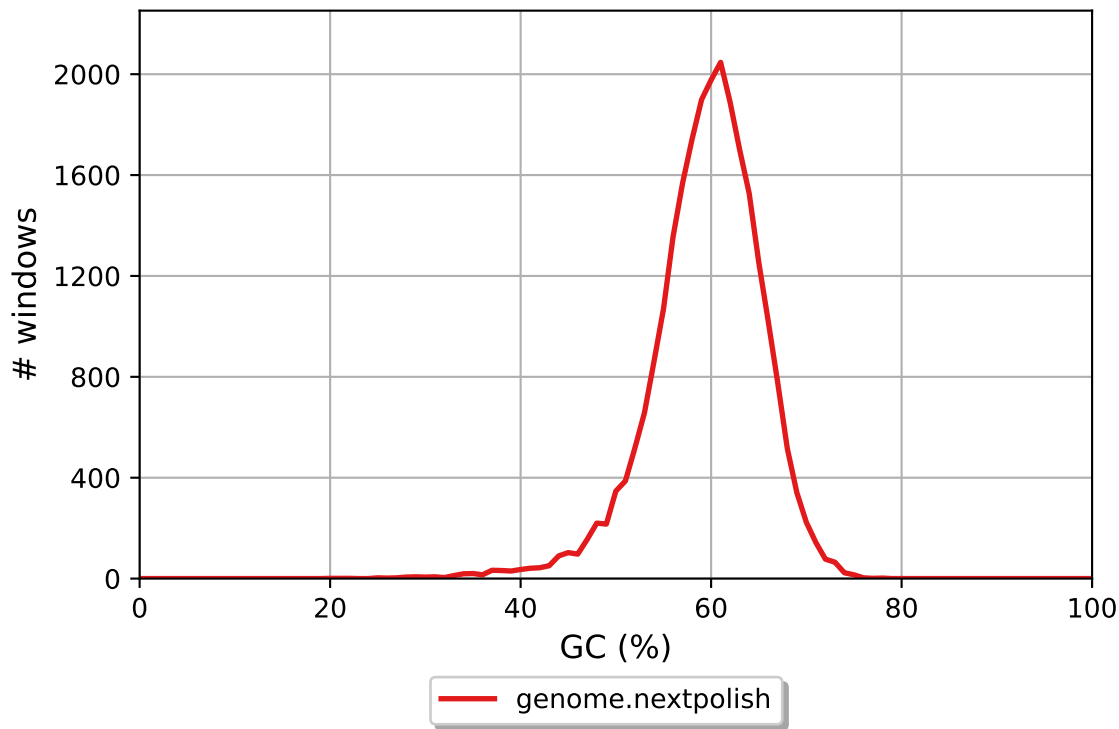
Nx



— genome.nextpolish



## GC content



genome.nextpolish GC content

