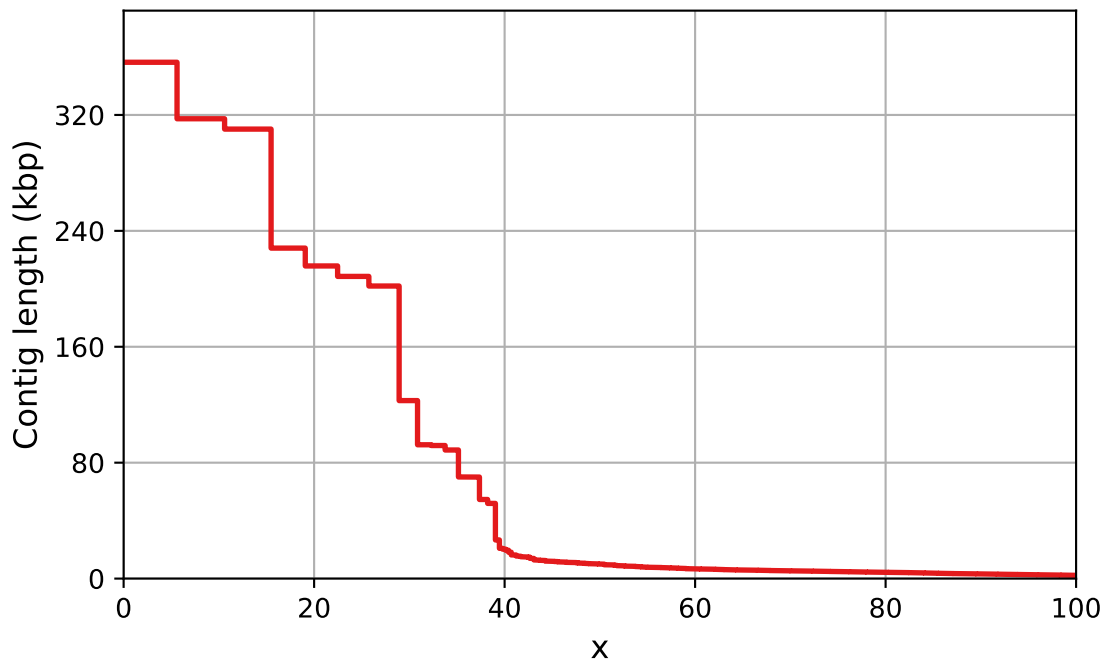


# Report

	genome.nextpolish
# contigs (>= 0 bp)	793
# contigs (>= 1000 bp)	793
# contigs (>= 5000 bp)	298
# contigs (>= 10000 bp)	69
# contigs (>= 25000 bp)	16
# contigs (>= 50000 bp)	15
Total length (>= 0 bp)	6355487
Total length (>= 1000 bp)	6355487
Total length (>= 5000 bp)	4668035
Total length (>= 10000 bp)	3177203
Total length (>= 25000 bp)	2507464
Total length (>= 50000 bp)	2480800
# contigs	793
Largest contig	356358
Total length	6355487
GC (%)	54.26
N50	9997
N90	3200
auN	92508.2
L50	70
L90	558
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	6 + 1 part

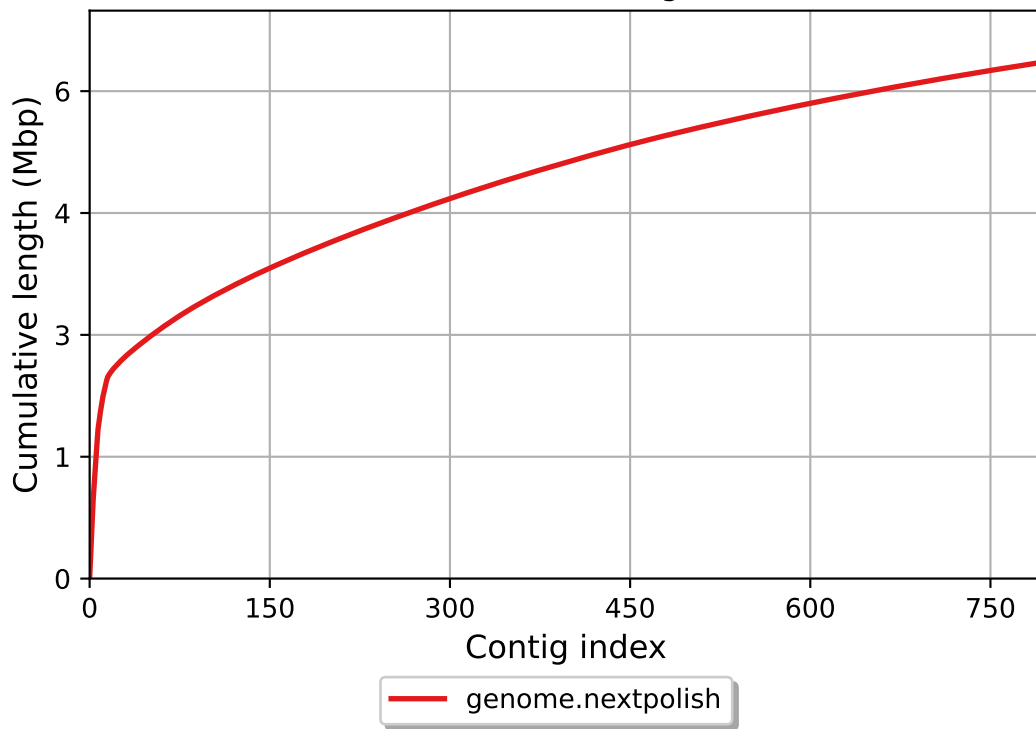
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

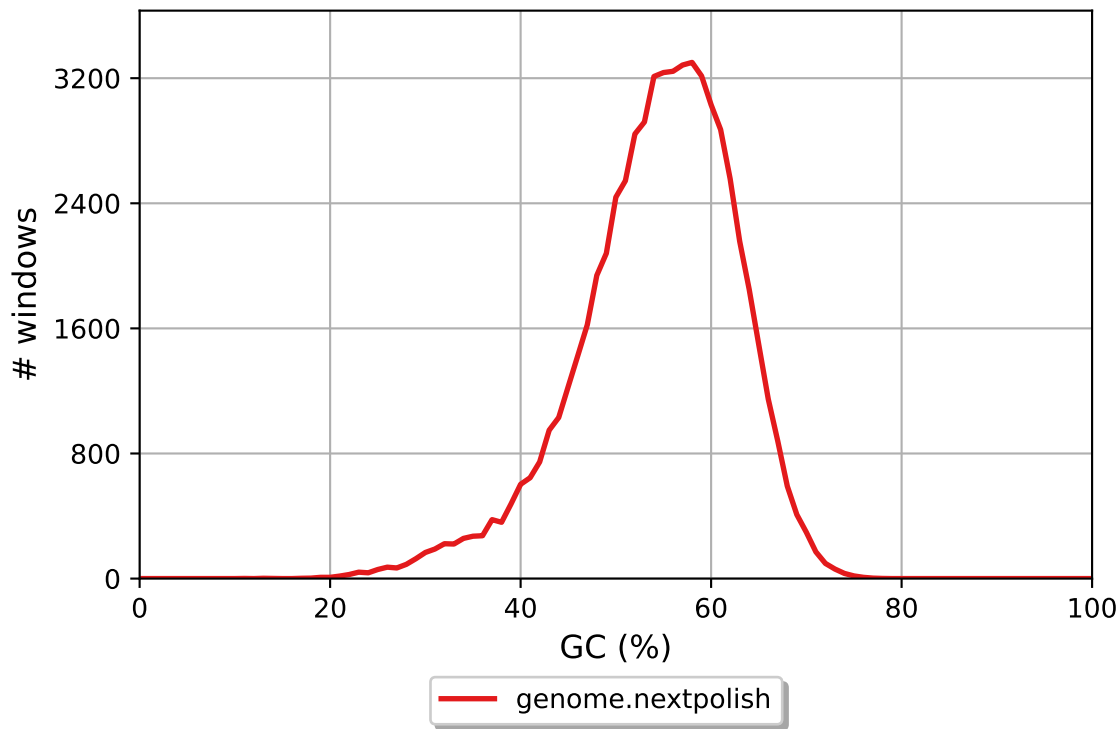


— genome.nextpolish

Cumulative length



## GC content



genome.nextpolish GC content

