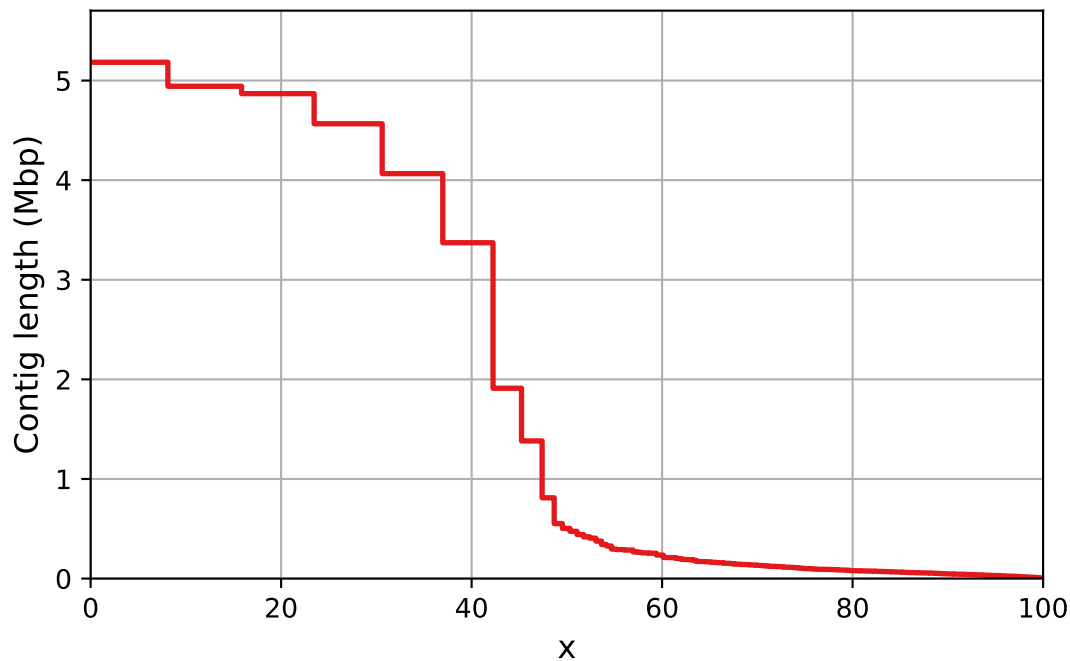


# Report

	genome.nextpolish
# contigs (>= 0 bp)	512
# contigs (>= 1000 bp)	512
# contigs (>= 5000 bp)	499
# contigs (>= 10000 bp)	460
# contigs (>= 25000 bp)	345
# contigs (>= 50000 bp)	224
Total length (>= 0 bp)	63903418
Total length (>= 1000 bp)	63903418
Total length (>= 5000 bp)	63852113
Total length (>= 10000 bp)	63549147
Total length (>= 25000 bp)	61584368
Total length (>= 50000 bp)	57132106
# contigs	512
Largest contig	5183495
Total length	63903418
GC (%)	43.66
N50	503695
N90	47728
auN	2111520.8
L50	11
L90	232
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	178 + 7 part

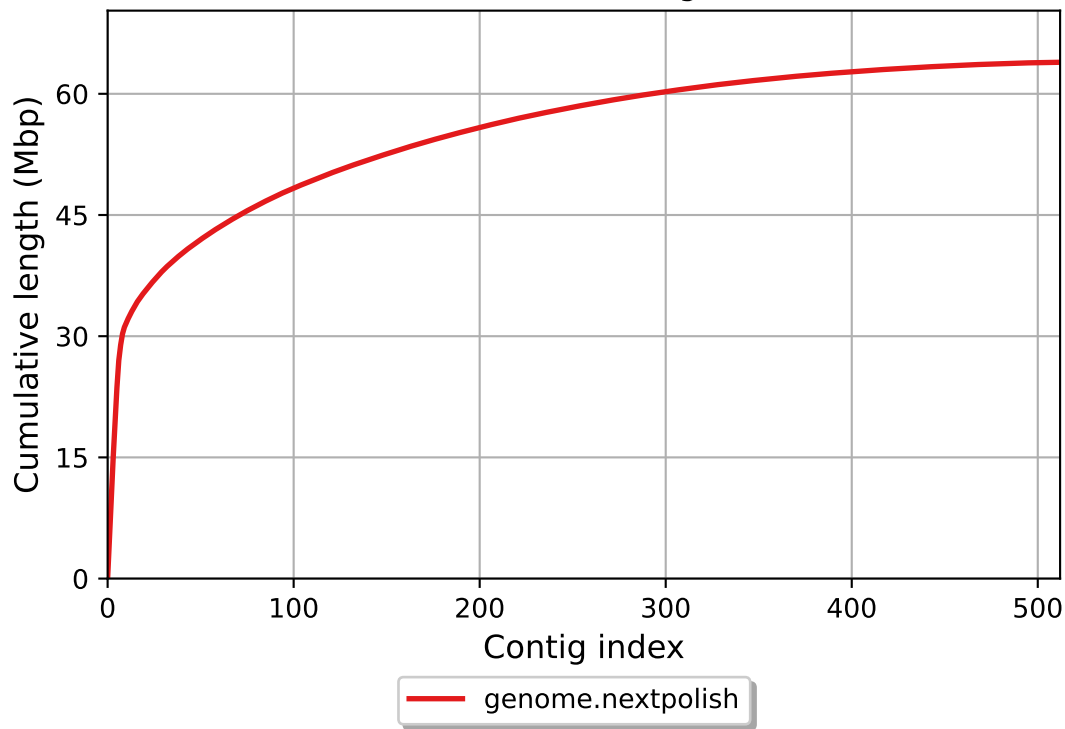
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

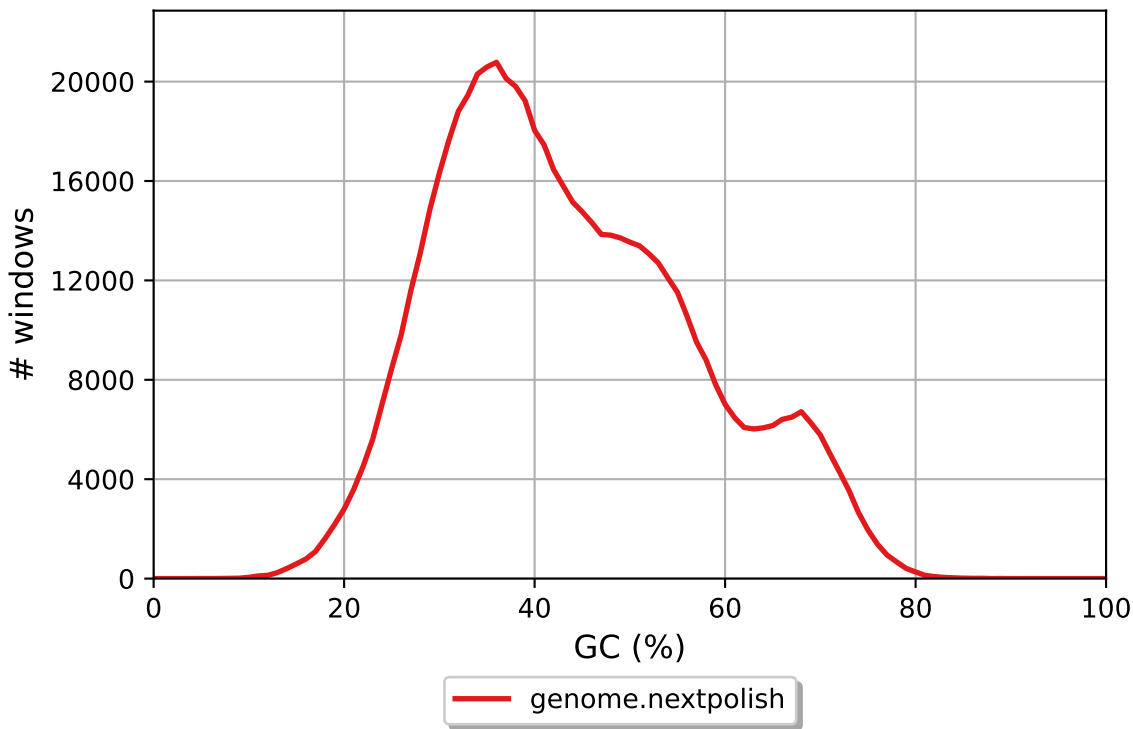


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

