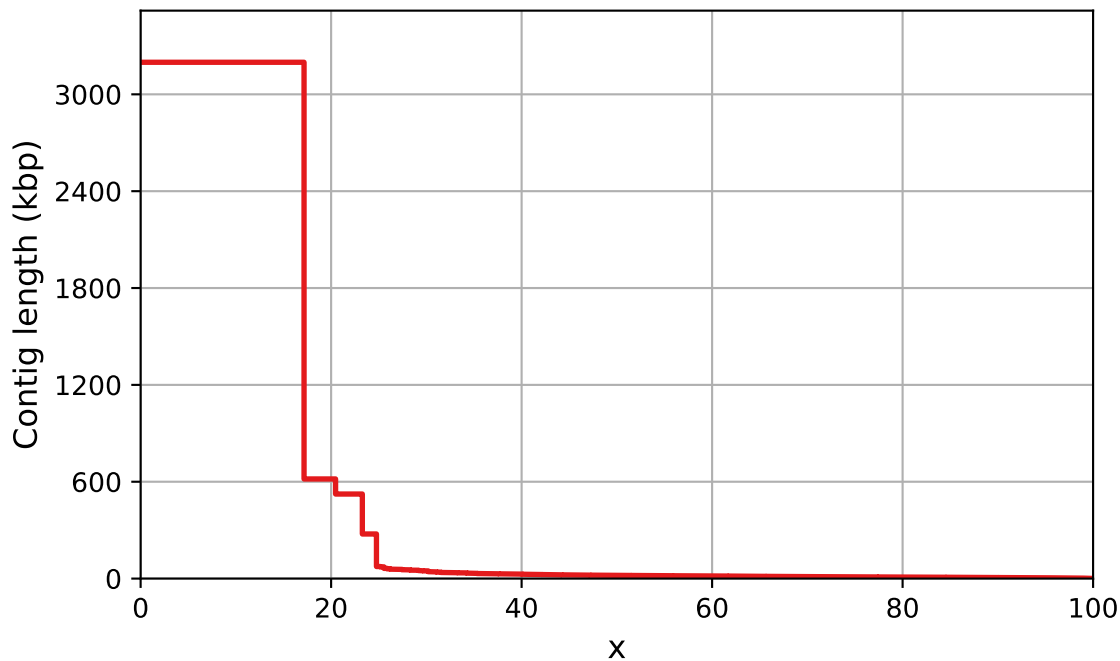


# Report

	genome.nextpolish
# contigs (>= 0 bp)	1252
# contigs (>= 1000 bp)	1250
# contigs (>= 5000 bp)	961
# contigs (>= 10000 bp)	522
# contigs (>= 25000 bp)	93
# contigs (>= 50000 bp)	18
Total length (>= 0 bp)	18642710
Total length (>= 1000 bp)	18641225
Total length (>= 5000 bp)	17663225
Total length (>= 10000 bp)	14446860
Total length (>= 25000 bp)	7860346
Total length (>= 50000 bp)	5436515
# contigs	1252
Largest contig	3198608
Total length	18642710
GC (%)	47.55
N50	20001
N90	6459
auN	602295.8
L50	160
L90	806
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	51 + 10 part

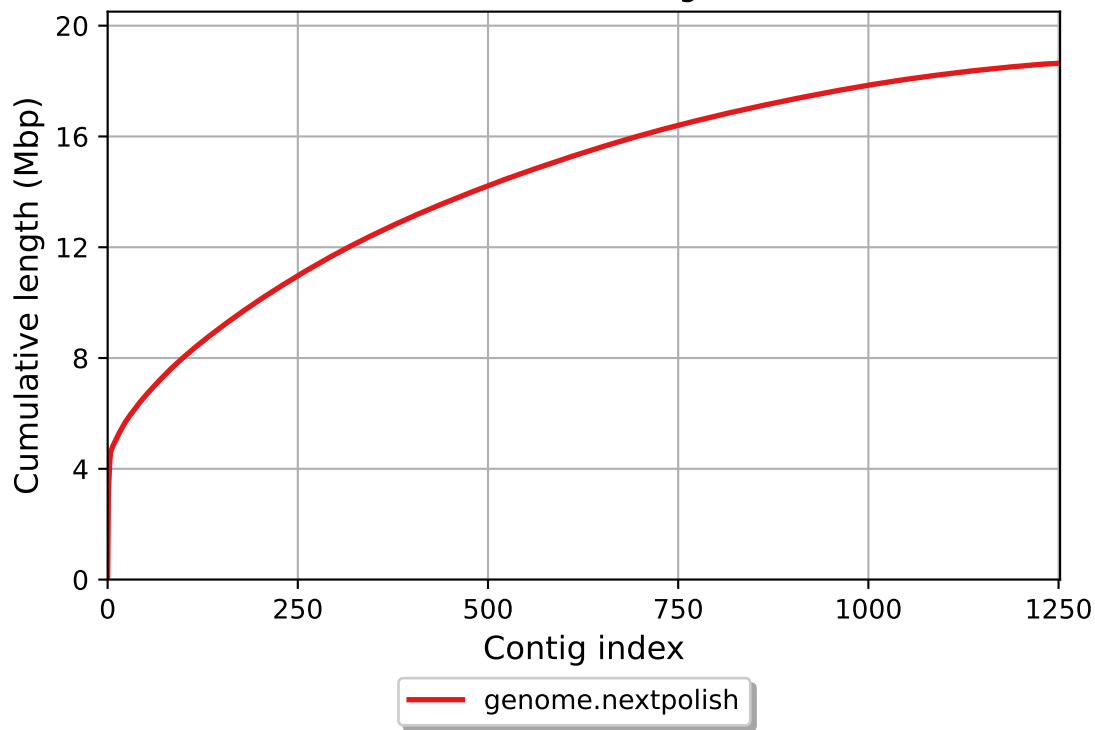
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

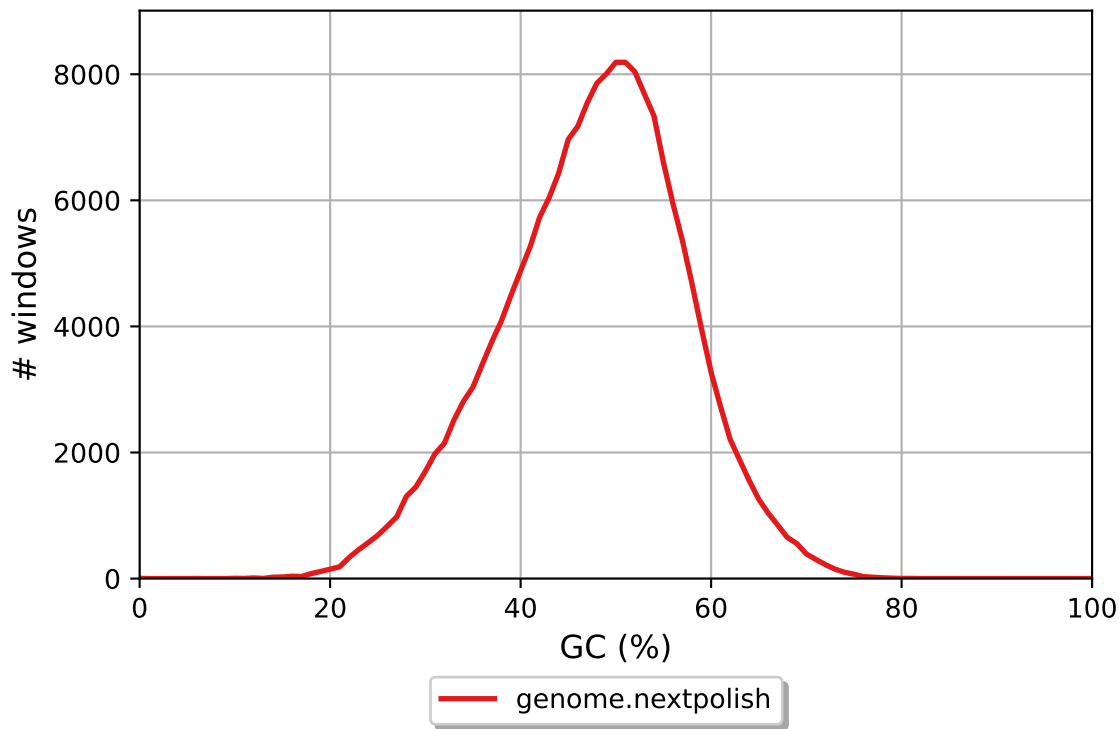


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

