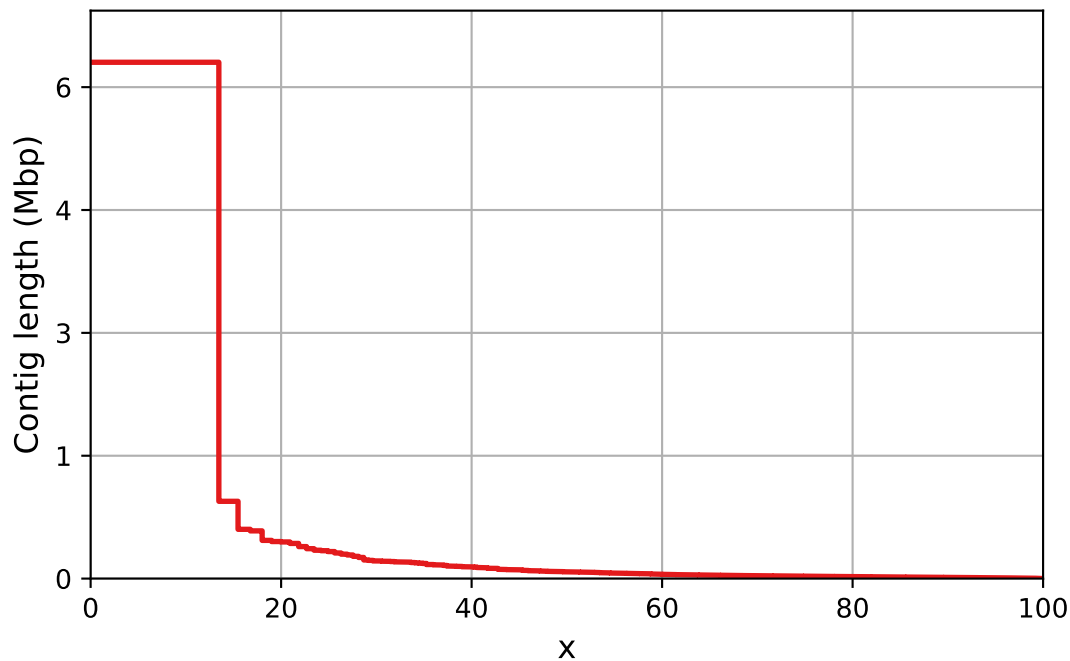


Report

	genome.nextpolish
# contigs (>= 0 bp)	1296
# contigs (>= 1000 bp)	1289
# contigs (>= 5000 bp)	1125
# contigs (>= 10000 bp)	856
# contigs (>= 25000 bp)	421
# contigs (>= 50000 bp)	171
Total length (>= 0 bp)	46823303
Total length (>= 1000 bp)	46818189
Total length (>= 5000 bp)	46264326
Total length (>= 10000 bp)	44278810
Total length (>= 25000 bp)	37155176
Total length (>= 50000 bp)	28551667
# contigs	1296
Largest contig	6303729
Total length	46823303
GC (%)	53.64
N50	82631
N90	14448
auN	970702.1
L50	93
L90	681
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	91 + 1 part

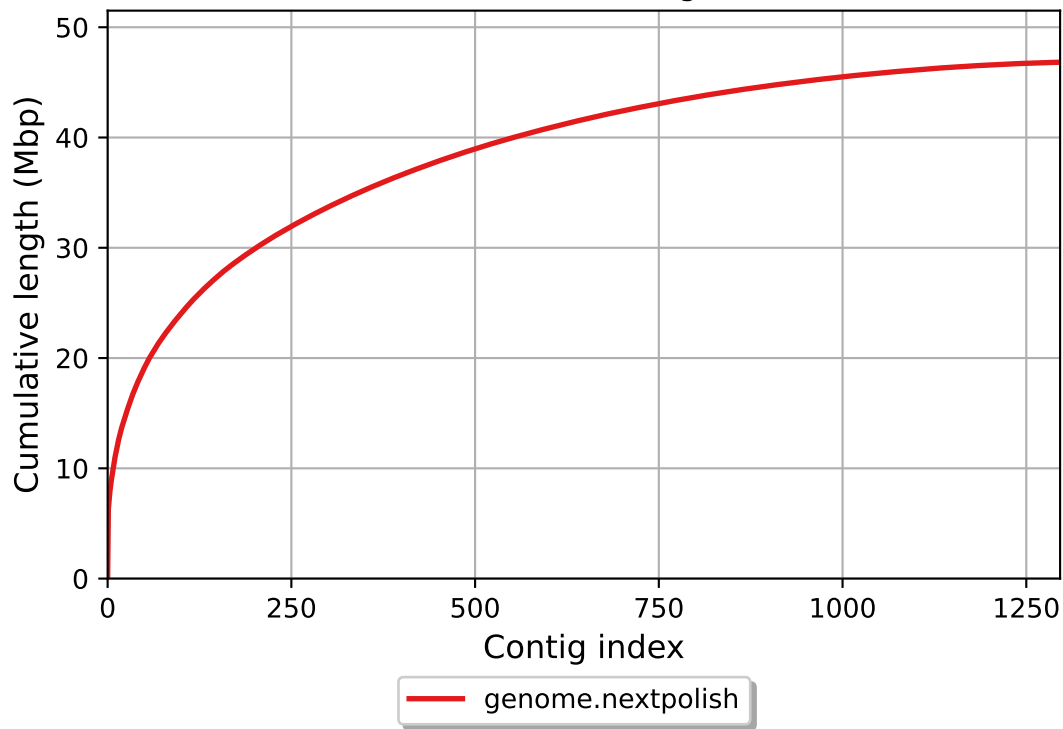
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

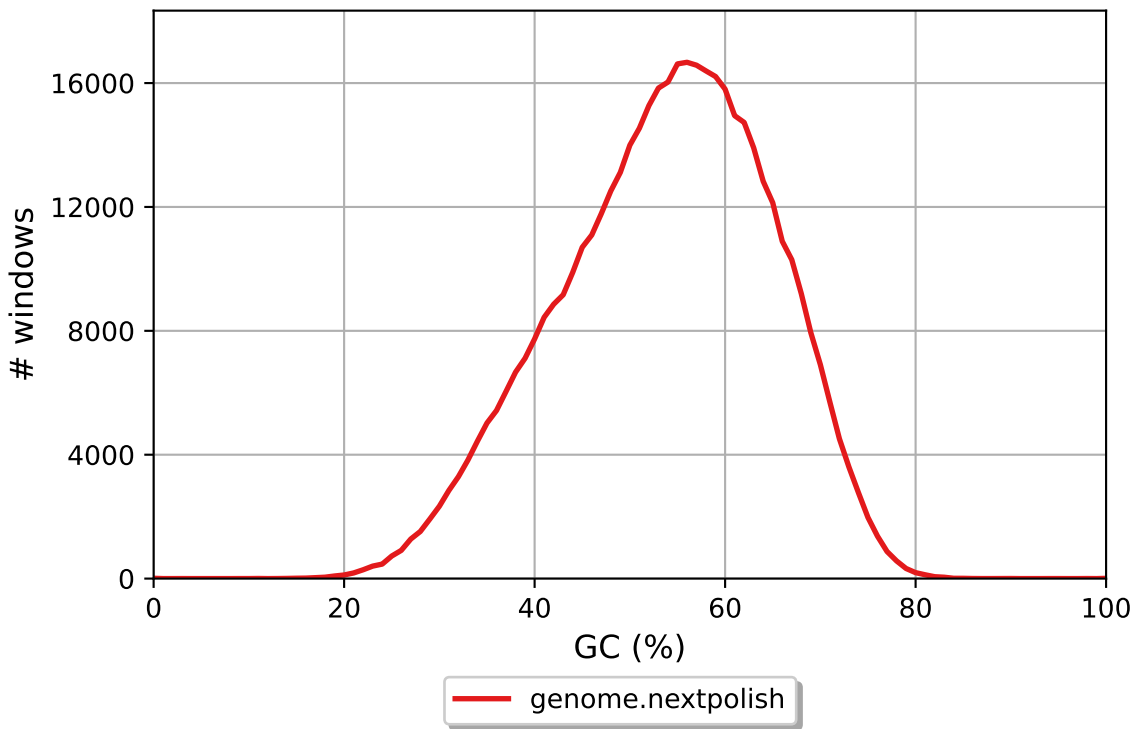


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

