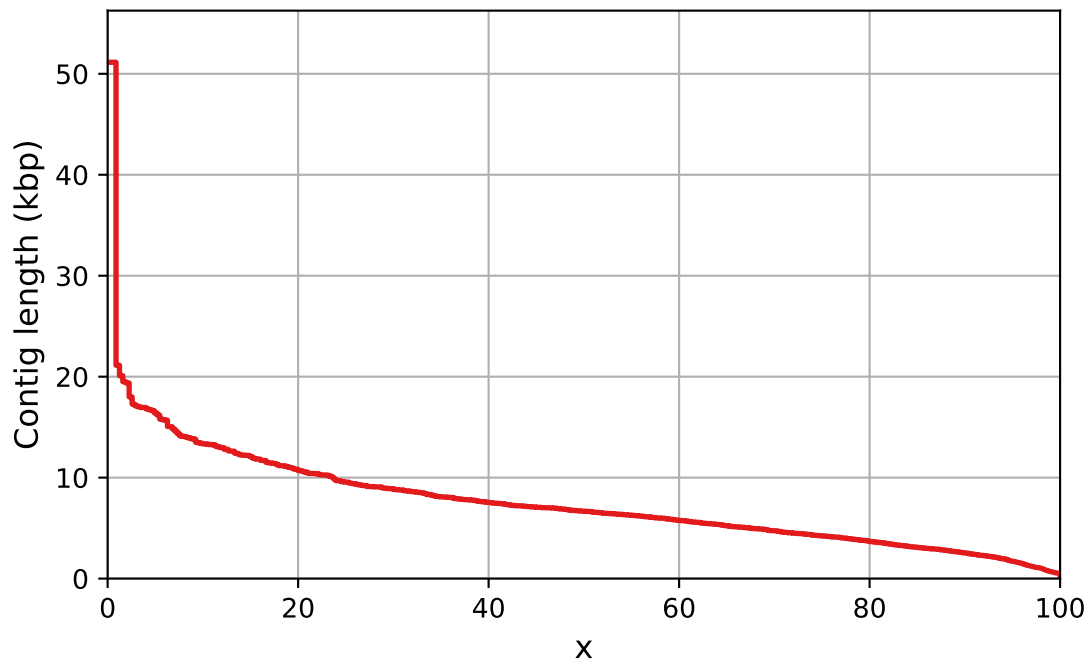


Report

	genome.nextpolish
# contigs (>= 0 bp)	1267
# contigs (>= 1000 bp)	1095
# contigs (>= 5000 bp)	469
# contigs (>= 10000 bp)	103
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	5803509
Total length (>= 1000 bp)	5689158
Total length (>= 5000 bp)	3912886
Total length (>= 10000 bp)	1373213
Total length (>= 25000 bp)	51142
Total length (>= 50000 bp)	51142
# contigs	1259
Largest contig	51142
Total length	5800637
GC (%)	53.68
N50	6667
N90	2545
auN	7677.5
L50	296
L90	826
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	13 + 2 part

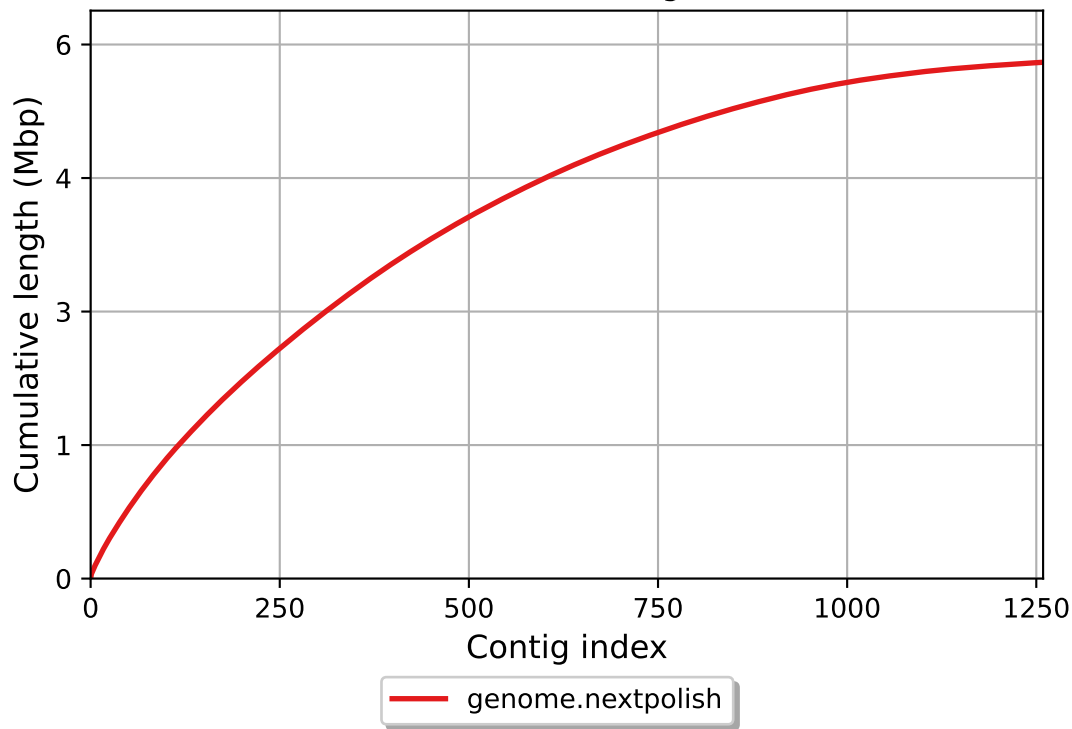
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

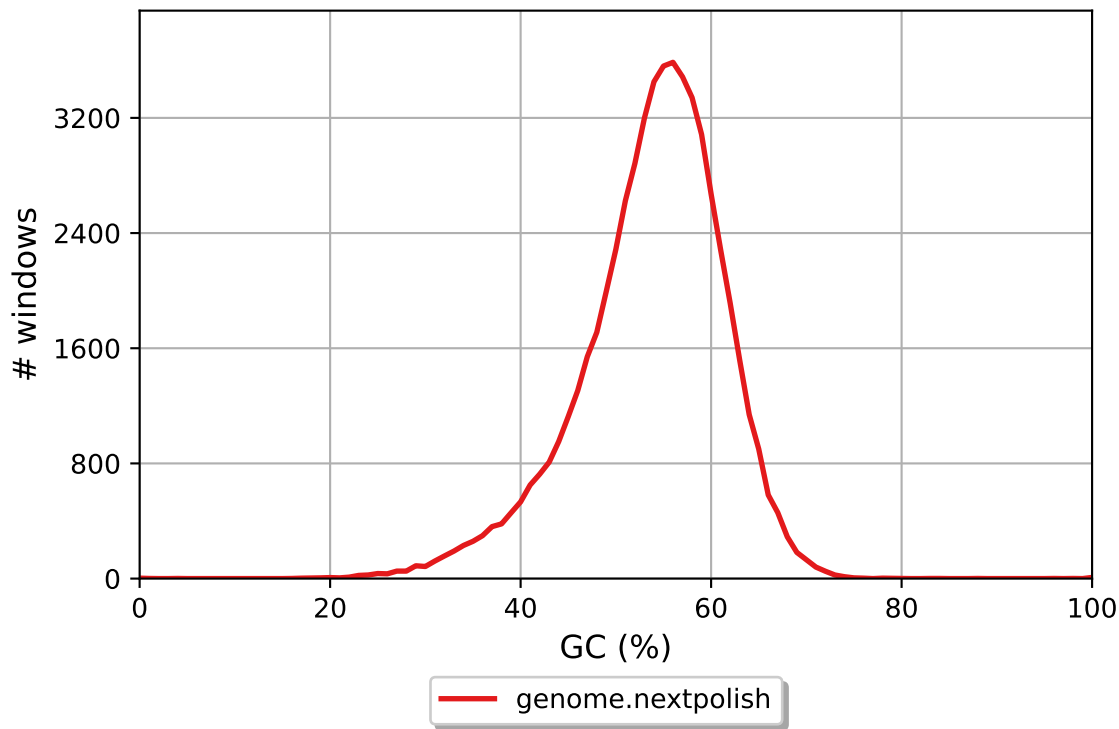


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

