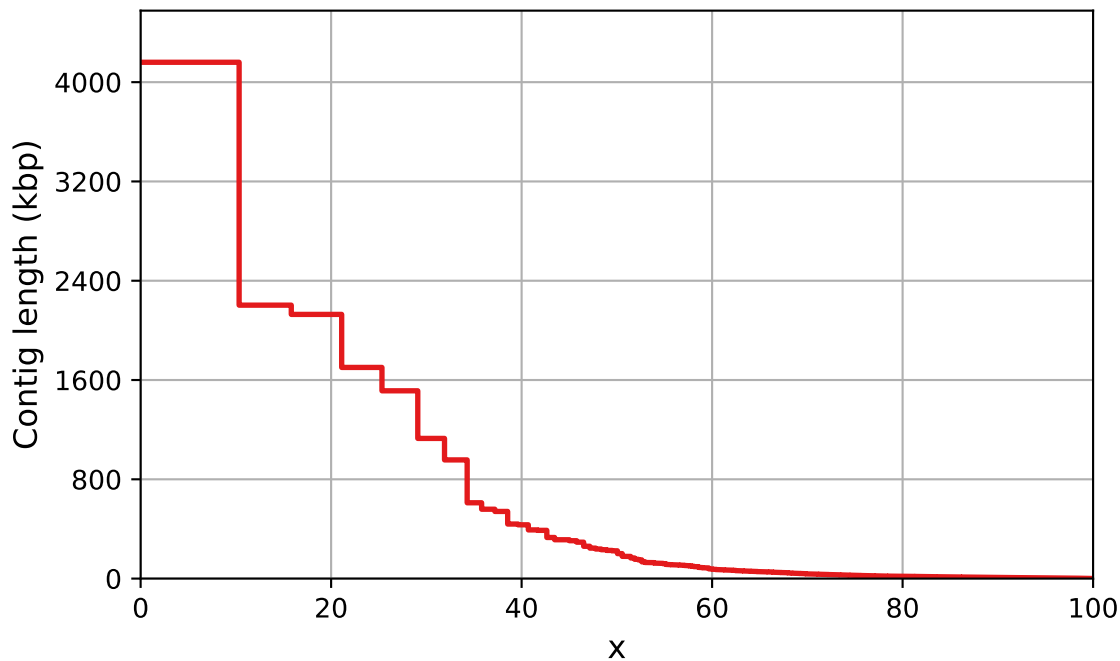


Report

	genome.nextpolish
# contigs (>= 0 bp)	1451
# contigs (>= 1000 bp)	1447
# contigs (>= 5000 bp)	942
# contigs (>= 10000 bp)	559
# contigs (>= 25000 bp)	205
# contigs (>= 50000 bp)	104
Total length (>= 0 bp)	40231023
Total length (>= 1000 bp)	40228595
Total length (>= 5000 bp)	38589561
Total length (>= 10000 bp)	35876086
Total length (>= 25000 bp)	30407525
Total length (>= 50000 bp)	26905780
# contigs	1451
Largest contig	4160311
Total length	40231023
GC (%)	45.88
N50	223813
N90	9184
auN	931475.4
L50	25
L90	594
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	89 + 1 part

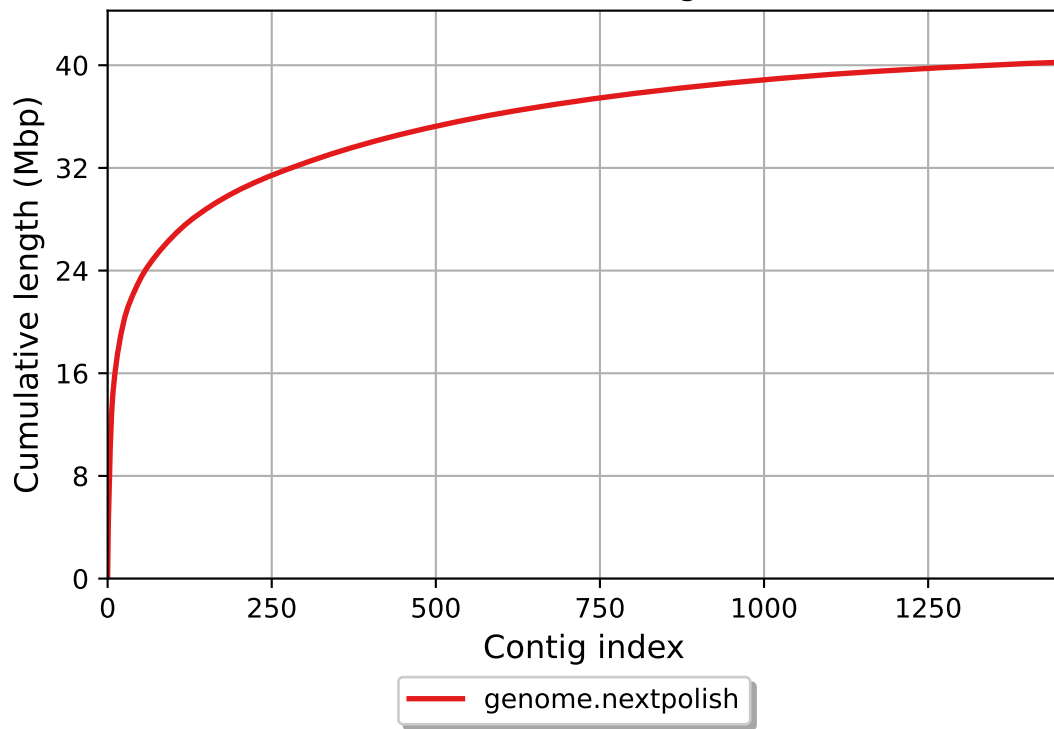
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

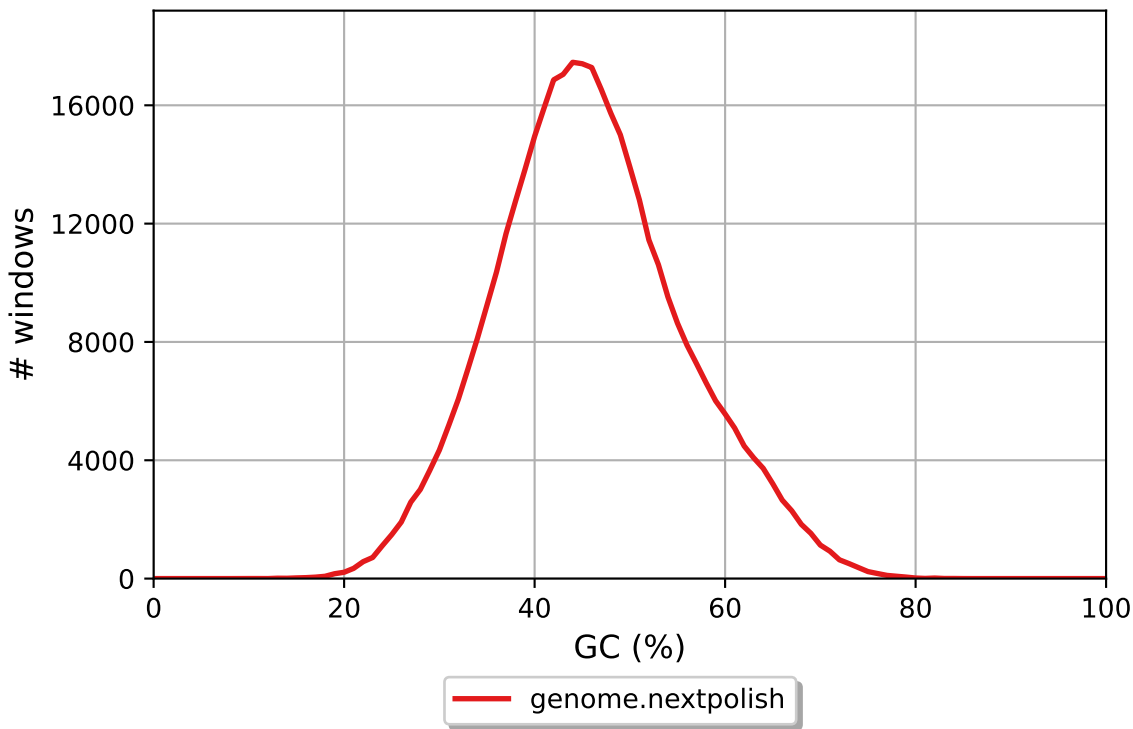


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

