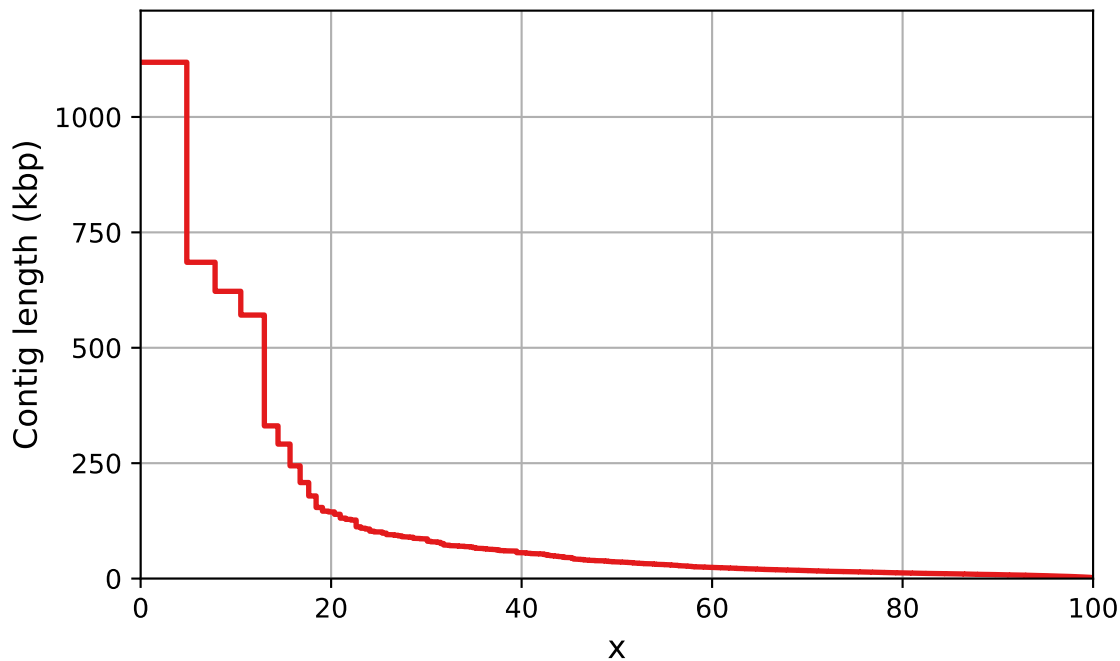


Report

	genome.nextpolish
# contigs (>= 0 bp)	1134
# contigs (>= 1000 bp)	1133
# contigs (>= 5000 bp)	931
# contigs (>= 10000 bp)	600
# contigs (>= 25000 bp)	190
# contigs (>= 50000 bp)	81
Total length (>= 0 bp)	23069836
Total length (>= 1000 bp)	23068986
Total length (>= 5000 bp)	22352319
Total length (>= 10000 bp)	19890820
Total length (>= 25000 bp)	13650403
Total length (>= 50000 bp)	9919147
# contigs	1134
Largest contig	1118480
Total length	23069836
GC (%)	46.97
N50	35976
N90	8498
auN	152219.7
L50	120
L90	696
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	36 + 6 part

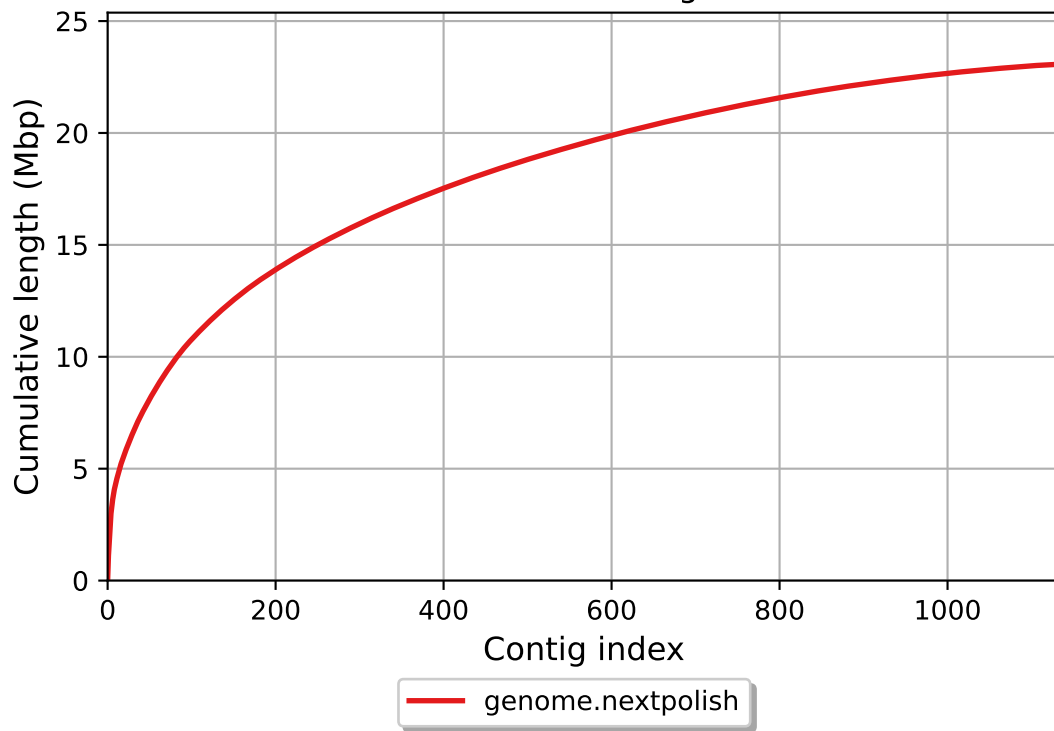
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

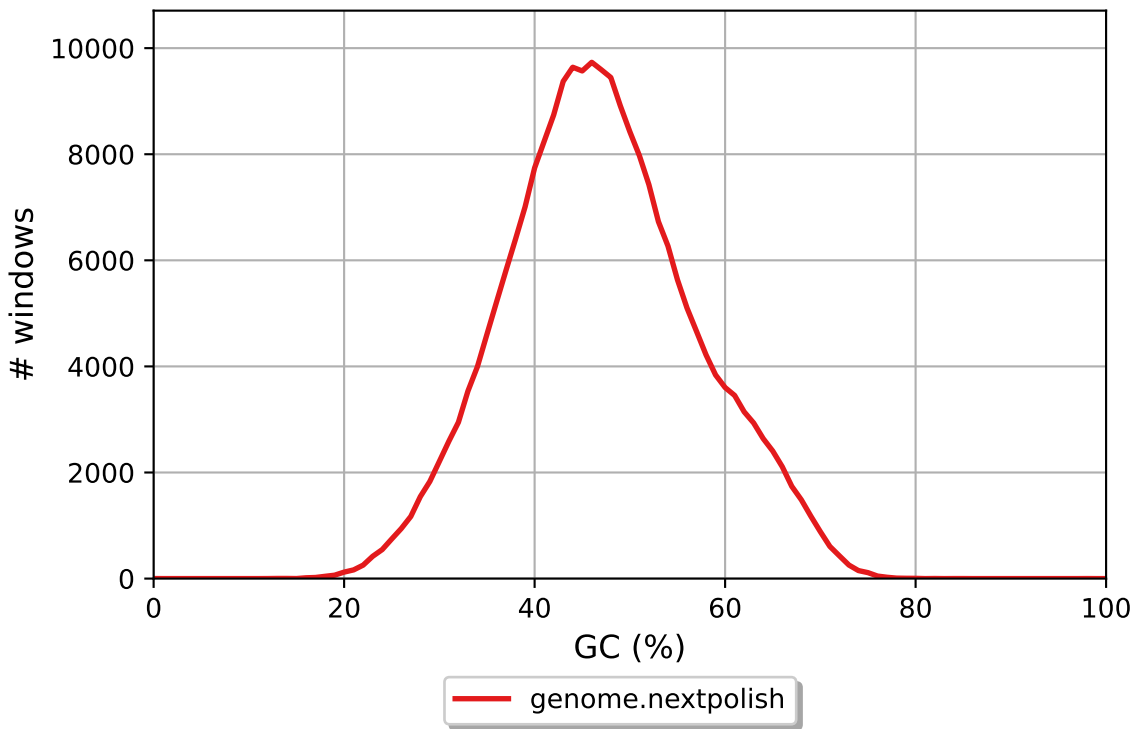


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

