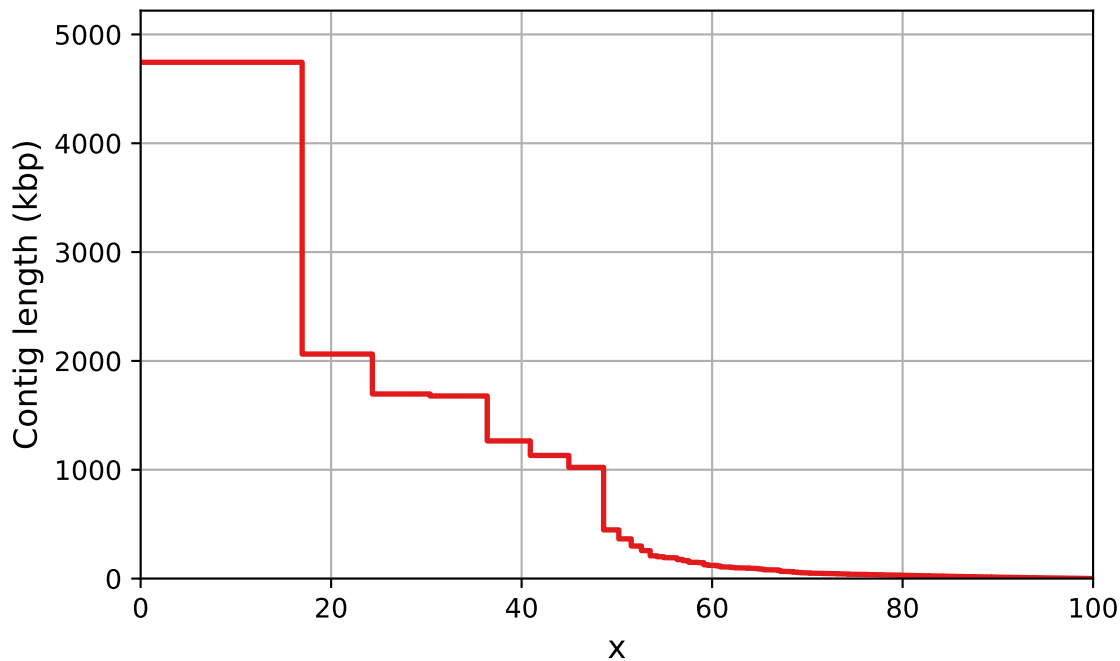


Report

	genome.nextpolish
# contigs (>= 0 bp)	973
# contigs (>= 1000 bp)	849
# contigs (>= 5000 bp)	475
# contigs (>= 10000 bp)	326
# contigs (>= 25000 bp)	158
# contigs (>= 50000 bp)	58
Total length (>= 0 bp)	27986471
Total length (>= 1000 bp)	27911070
Total length (>= 5000 bp)	26912543
Total length (>= 10000 bp)	25857347
Total length (>= 25000 bp)	23226108
Total length (>= 50000 bp)	19666715
# contigs	946
Largest contig	4744226
Total length	27978269
GC (%)	42.44
N50	447281
N90	13347
auN	1344360.3
L50	8
L90	266
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	126 + 2 part

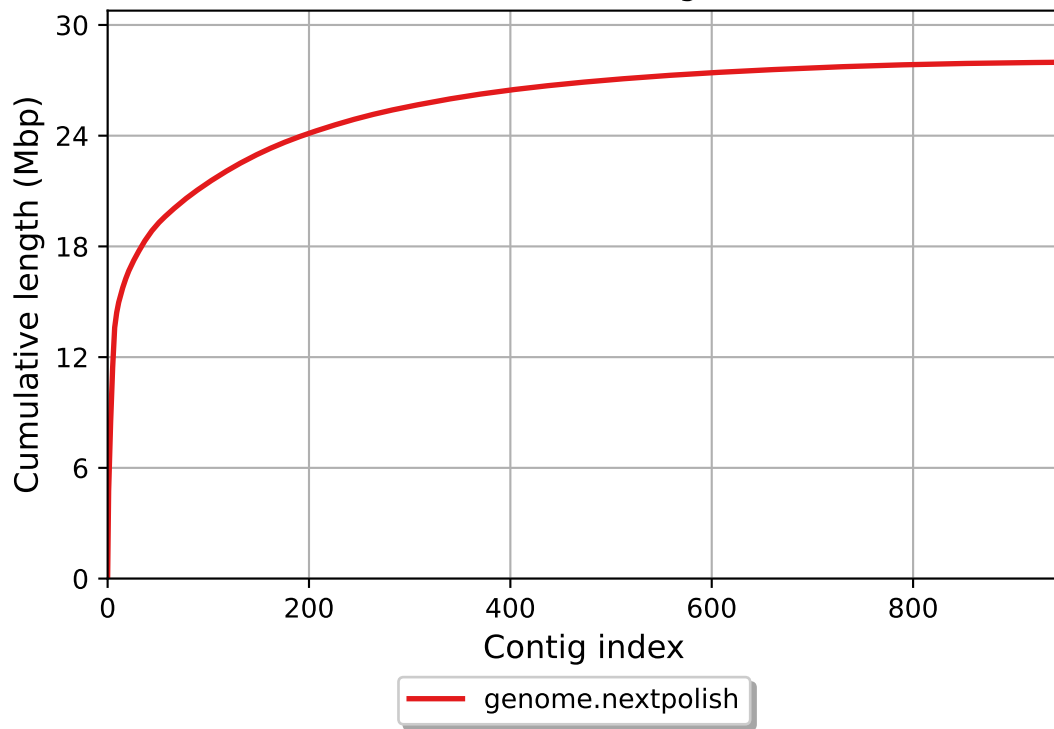
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

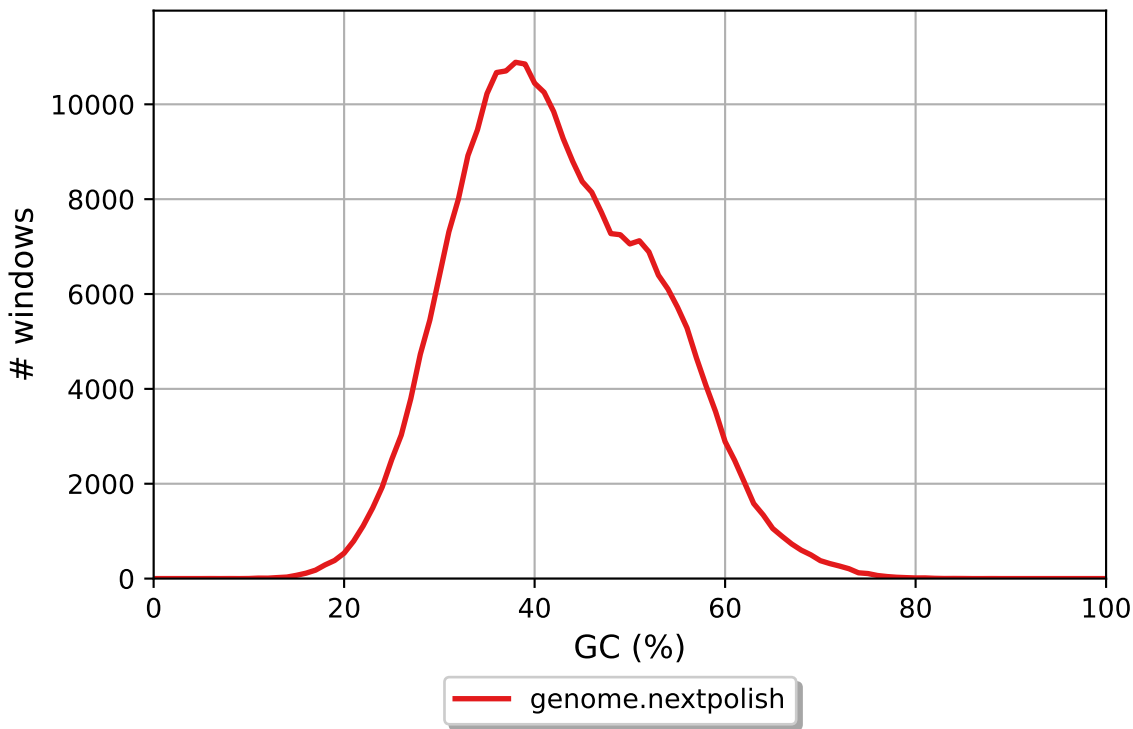


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

