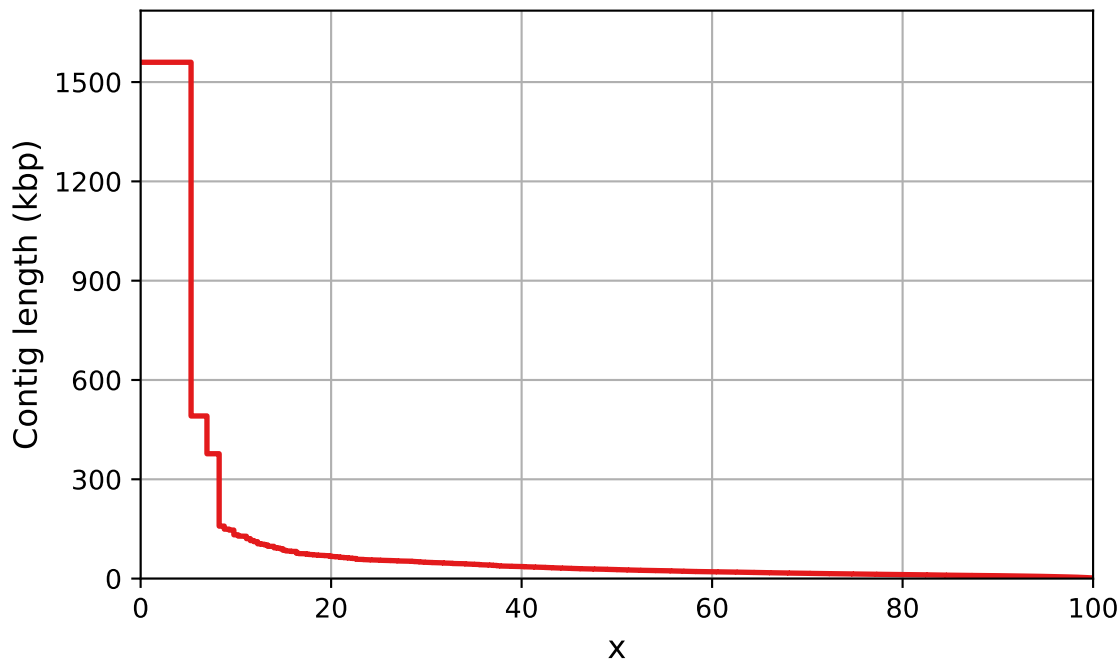


Report

	genome.nextpolish
# contigs (>= 0 bp)	1607
# contigs (>= 1000 bp)	1596
# contigs (>= 5000 bp)	1358
# contigs (>= 10000 bp)	903
# contigs (>= 25000 bp)	286
# contigs (>= 50000 bp)	89
Total length (>= 0 bp)	29452175
Total length (>= 1000 bp)	29444704
Total length (>= 5000 bp)	28635163
Total length (>= 10000 bp)	25169946
Total length (>= 25000 bp)	15497863
Total length (>= 50000 bp)	8679755
# contigs	1606
Largest contig	1559855
Total length	29451676
GC (%)	43.64
N50	27103
N90	8604
auN	127767.9
L50	257
L90	1047
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	71 + 9 part

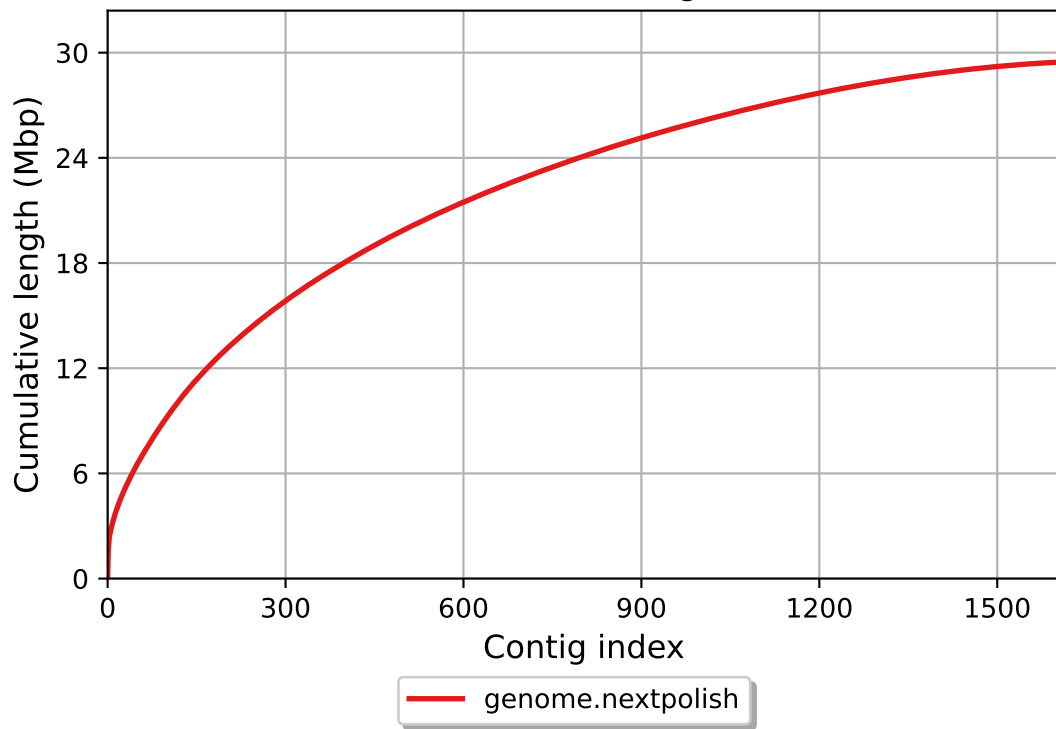
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

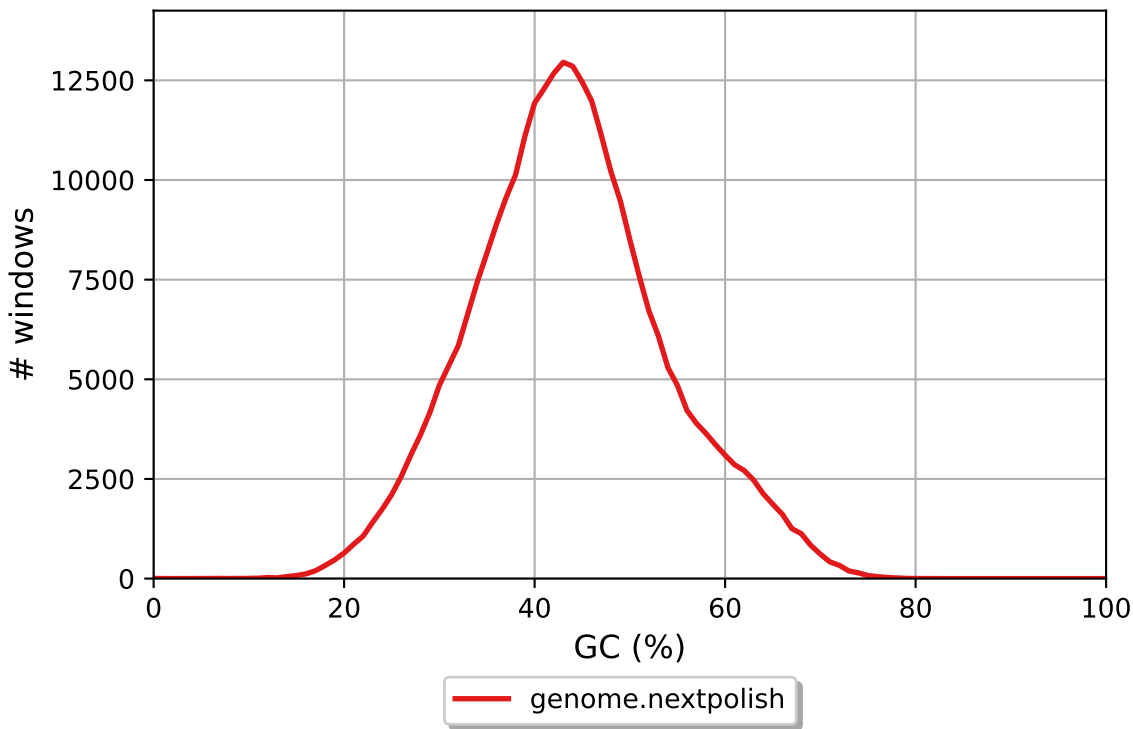


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

