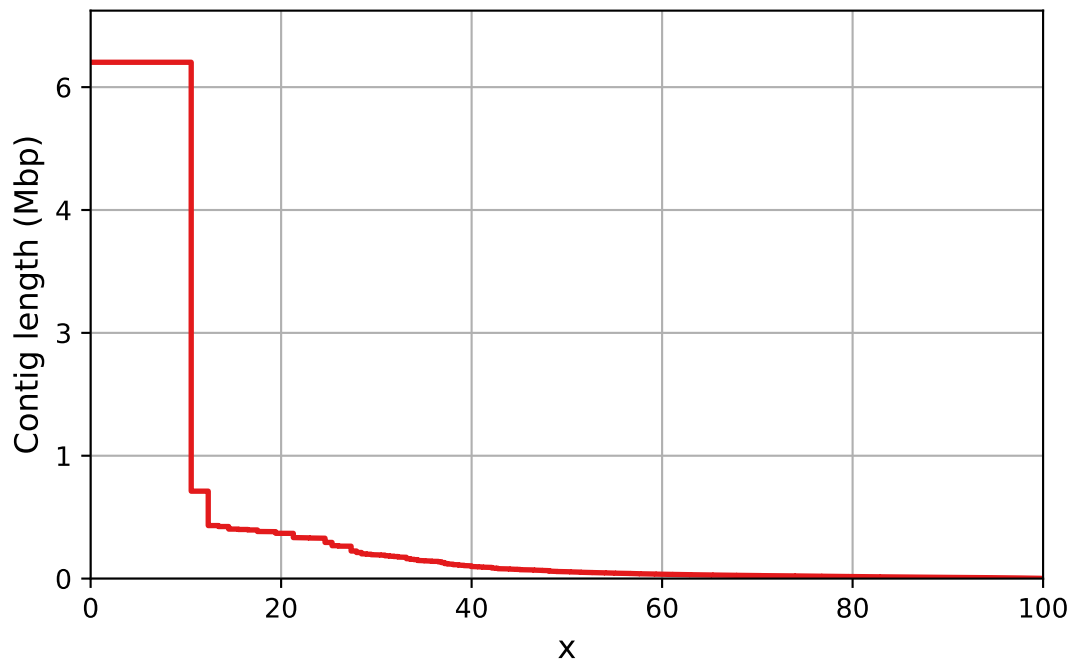


Report

	genome.nextpolish
# contigs (>= 0 bp)	1576
# contigs (>= 1000 bp)	1569
# contigs (>= 5000 bp)	1413
# contigs (>= 10000 bp)	1099
# contigs (>= 25000 bp)	521
# contigs (>= 50000 bp)	217
Total length (>= 0 bp)	59691422
Total length (>= 1000 bp)	59686194
Total length (>= 5000 bp)	59146960
Total length (>= 10000 bp)	56854461
Total length (>= 25000 bp)	47545420
Total length (>= 50000 bp)	36857670
# contigs	1576
Largest contig	6303671
Total length	59691422
GC (%)	53.13
N50	83630
N90	14598
auN	824450.2
L50	107
L90	845
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	145 + 4 part

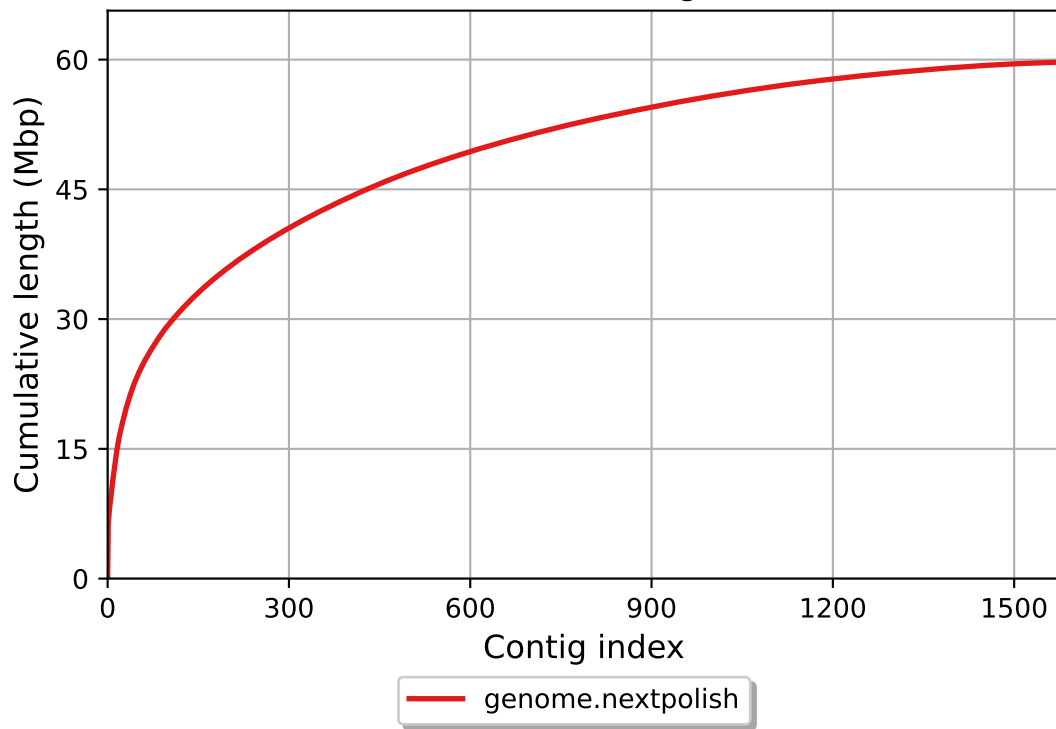
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

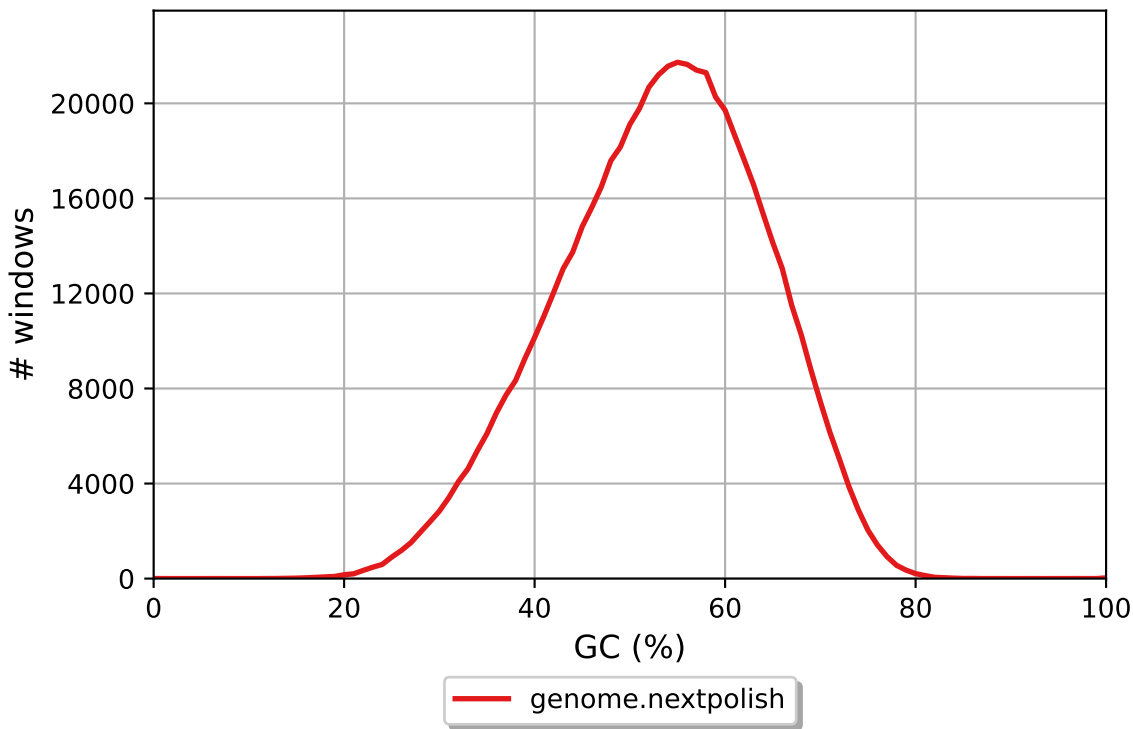


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

