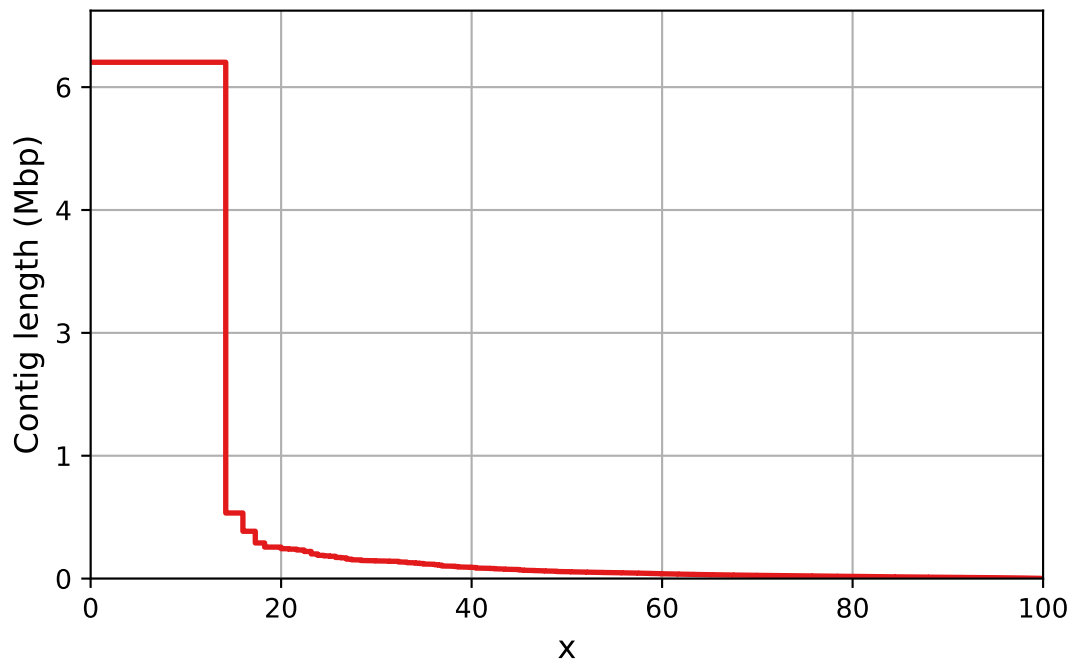


Report

	genome.nextpolish
# contigs (>= 0 bp)	1155
# contigs (>= 1000 bp)	1149
# contigs (>= 5000 bp)	1003
# contigs (>= 10000 bp)	795
# contigs (>= 25000 bp)	407
# contigs (>= 50000 bp)	181
Total length (>= 0 bp)	44435856
Total length (>= 1000 bp)	44431702
Total length (>= 5000 bp)	43941008
Total length (>= 10000 bp)	42400268
Total length (>= 25000 bp)	36203603
Total length (>= 50000 bp)	28251331
# contigs	1155
Largest contig	6303702
Total length	44435856
GC (%)	53.92
N50	84600
N90	15437
auN	1002582.1
L50	90
L90	602
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	95 + 1 part

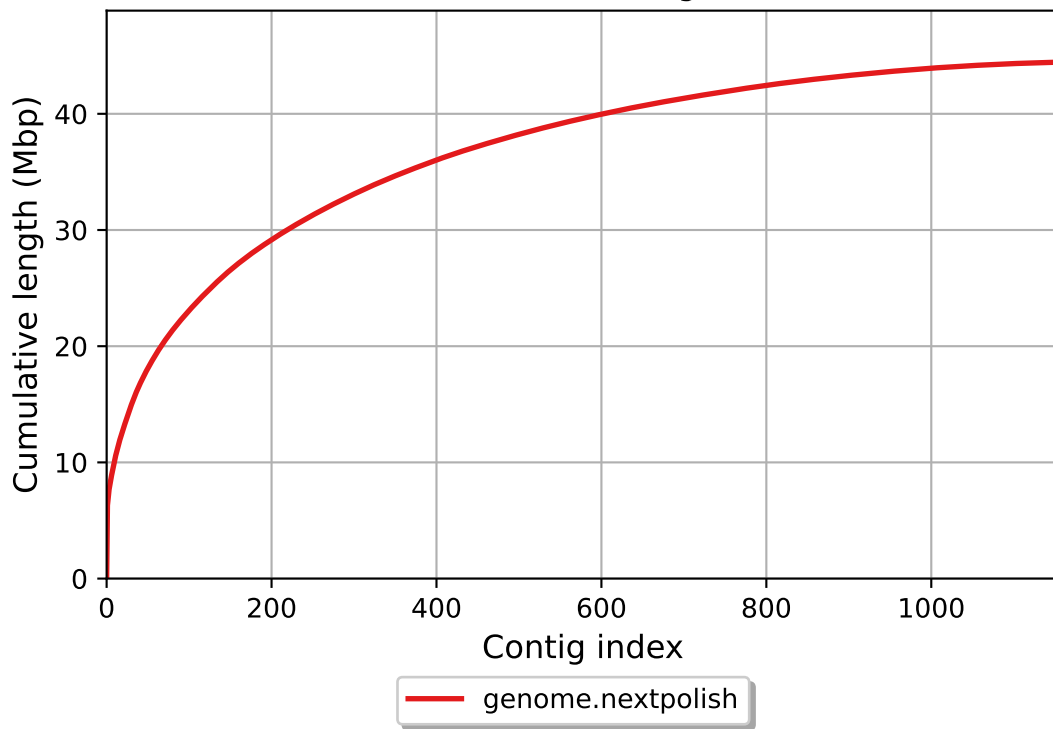
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

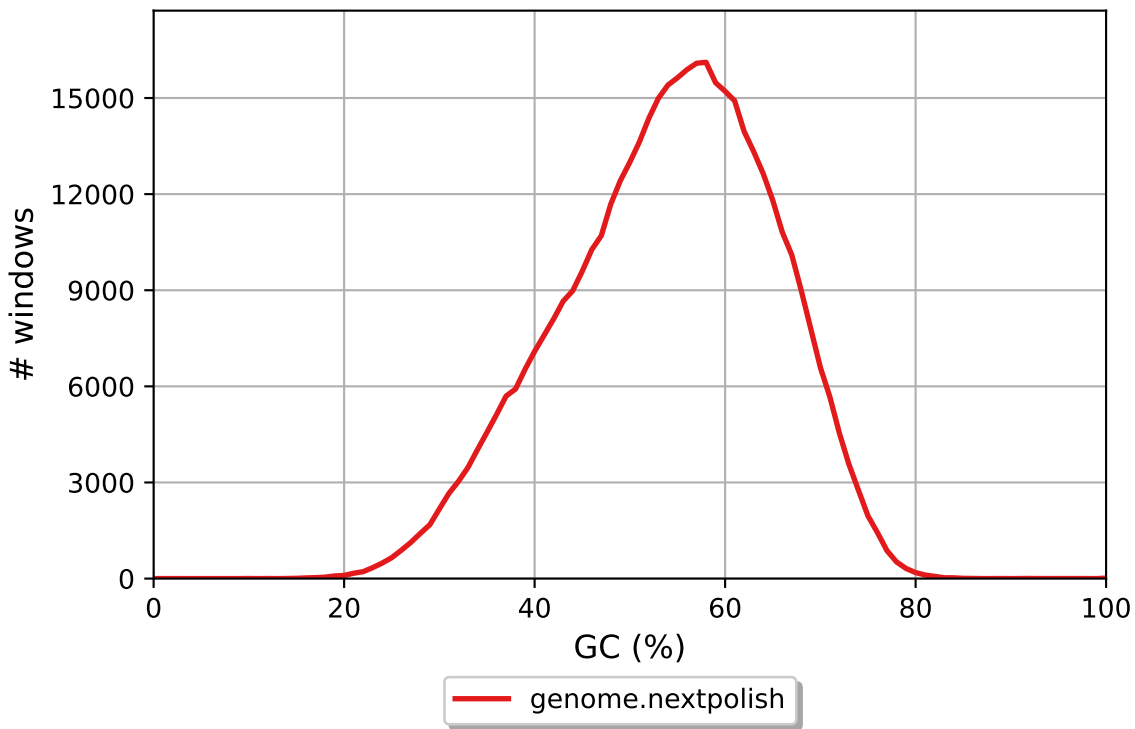


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

