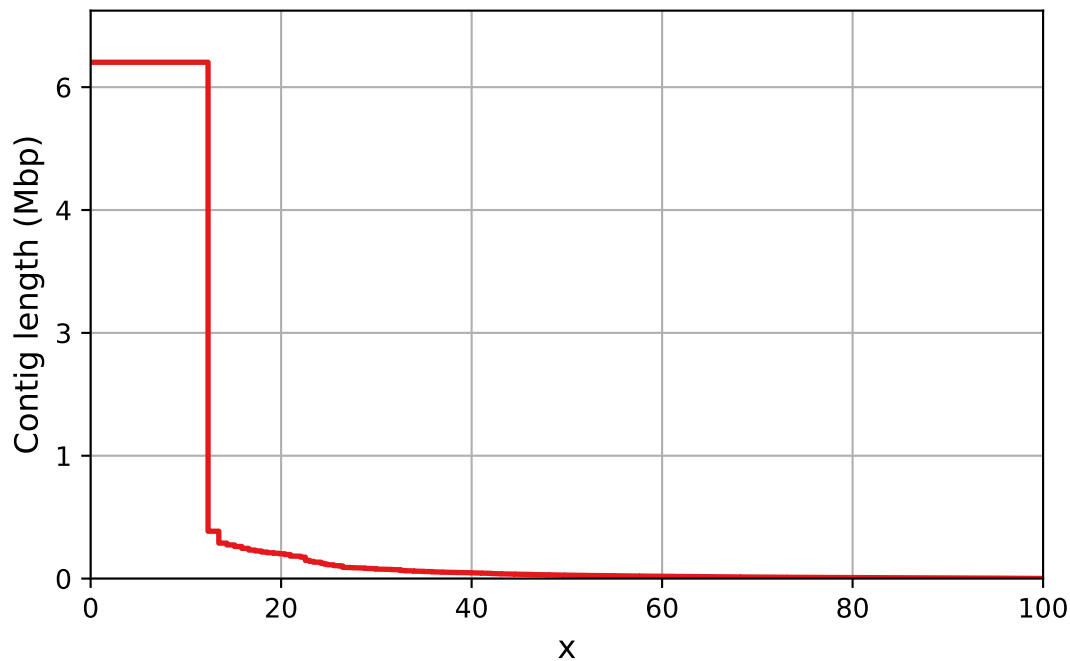


Report

	genome.nextpolish
# contigs (>= 0 bp)	2623
# contigs (>= 1000 bp)	2602
# contigs (>= 5000 bp)	1899
# contigs (>= 10000 bp)	1123
# contigs (>= 25000 bp)	402
# contigs (>= 50000 bp)	153
Total length (>= 0 bp)	51136116
Total length (>= 1000 bp)	51120869
Total length (>= 5000 bp)	48823755
Total length (>= 10000 bp)	43181073
Total length (>= 25000 bp)	32157018
Total length (>= 50000 bp)	23614183
# contigs	2623
Largest contig	6303950
Total length	51136116
GC (%)	55.01
N50	41806
N90	7504
auN	849501.7
L50	196
L90	1450
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	91 + 6 part

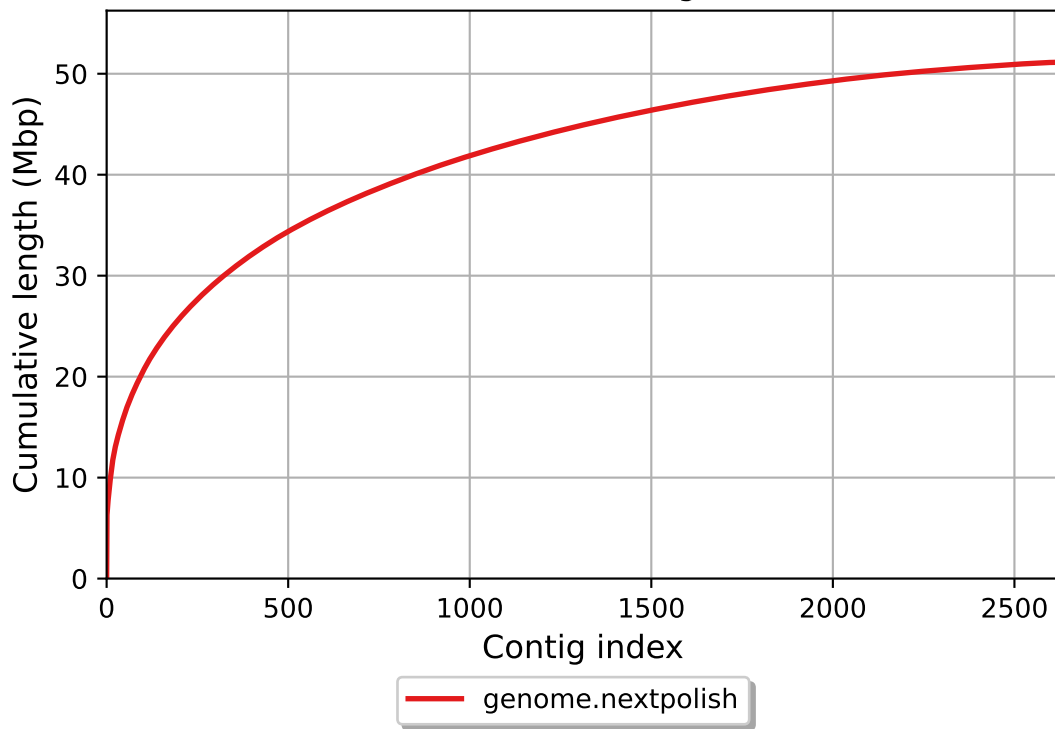
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

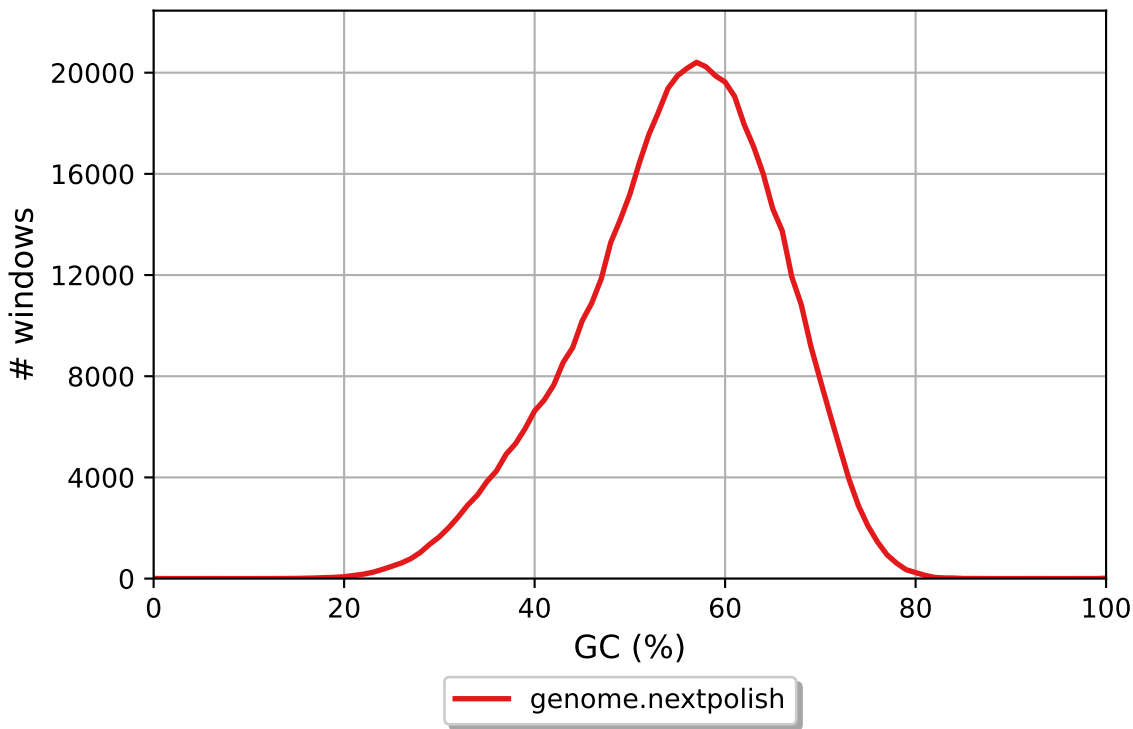


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

