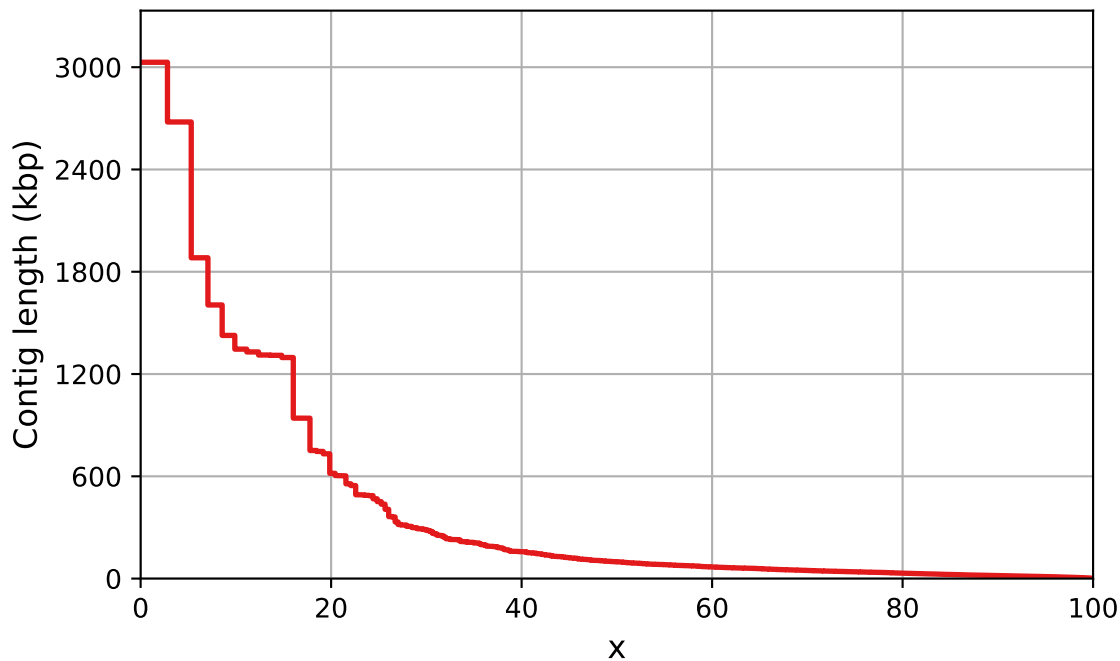


Report

	genome.nextpolish
# contigs (>= 0 bp)	2435
# contigs (>= 1000 bp)	2424
# contigs (>= 5000 bp)	2133
# contigs (>= 10000 bp)	1734
# contigs (>= 25000 bp)	943
# contigs (>= 50000 bp)	476
Total length (>= 0 bp)	107409973
Total length (>= 1000 bp)	107402099
Total length (>= 5000 bp)	106404093
Total length (>= 10000 bp)	103498085
Total length (>= 25000 bp)	90649932
Total length (>= 50000 bp)	73901875
# contigs	2434
Largest contig	3028910
Total length	107409474
GC (%)	51.73
N50	98324
N90	17692
auN	440626.3
L50	183
L90	1231
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	209 + 17 part

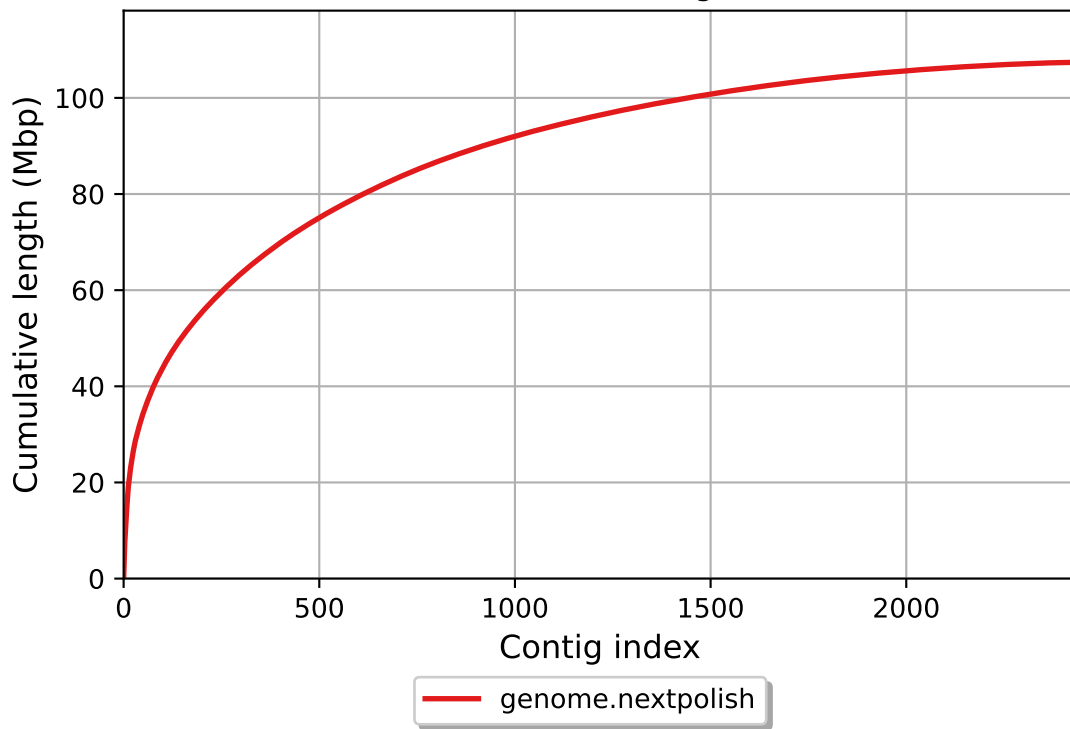
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

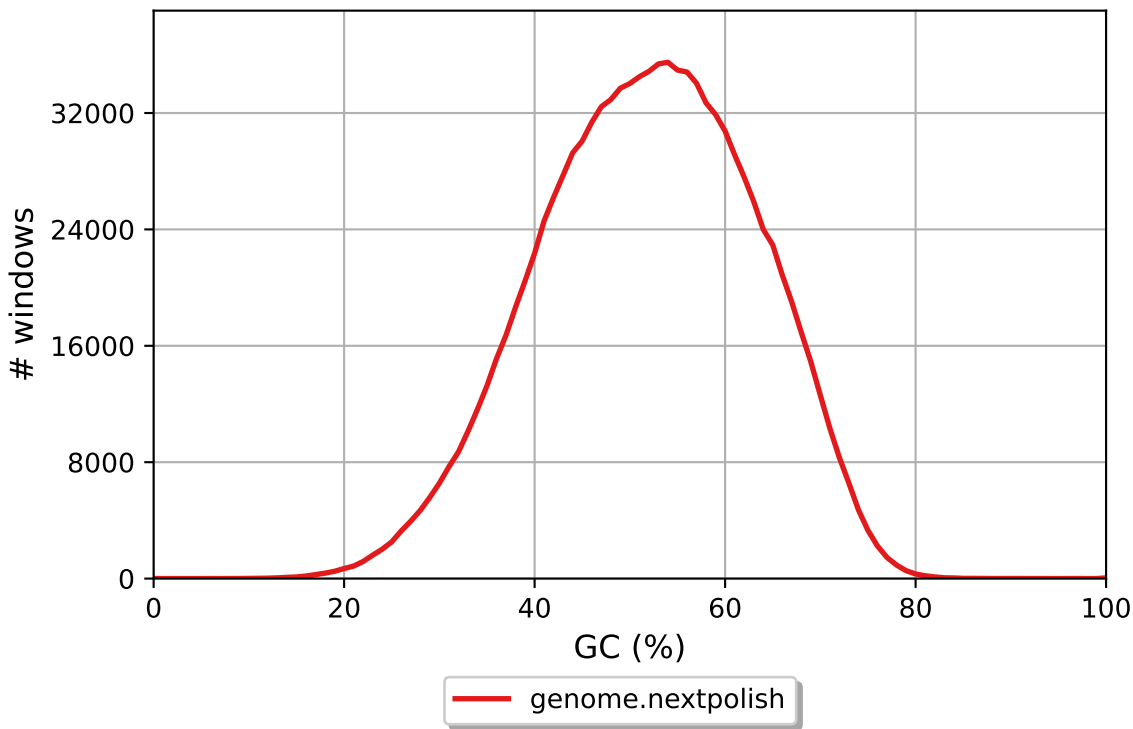


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

