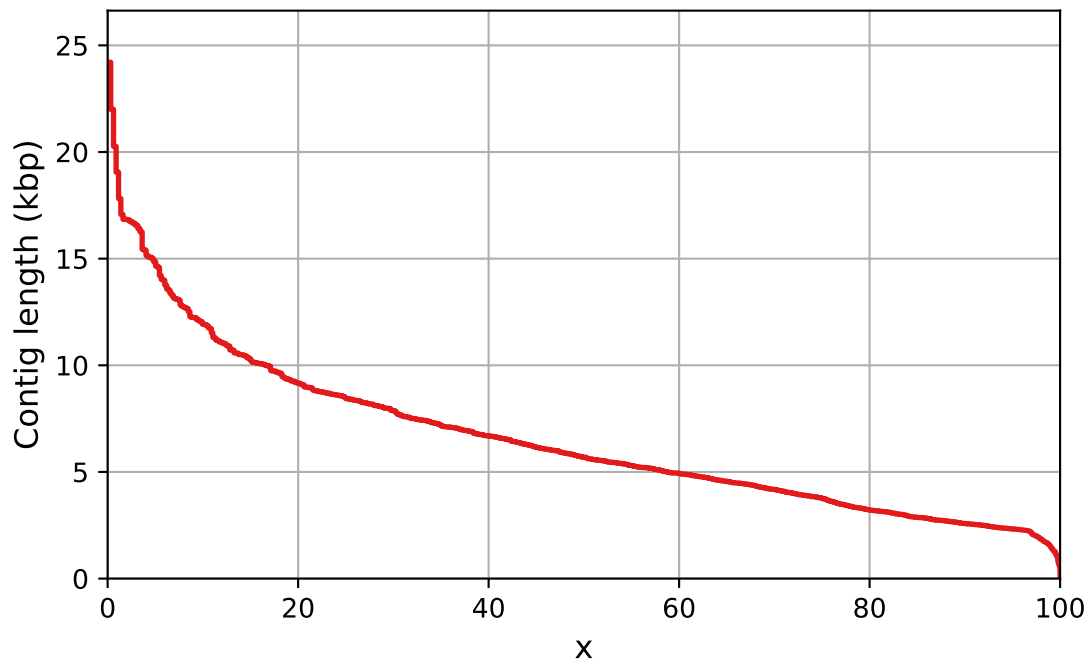


Report

	genome.nextpolish
# contigs (>= 0 bp)	1575
# contigs (>= 1000 bp)	1543
# contigs (>= 5000 bp)	557
# contigs (>= 10000 bp)	96
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	7455255
Total length (>= 1000 bp)	7435230
Total length (>= 5000 bp)	4379401
Total length (>= 10000 bp)	1235909
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	1573
Largest contig	24207
Total length	7455051
GC (%)	51.14
N50	5694
N90	2582
auN	6613.7
L50	435
L90	1201
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	14 + 5 part

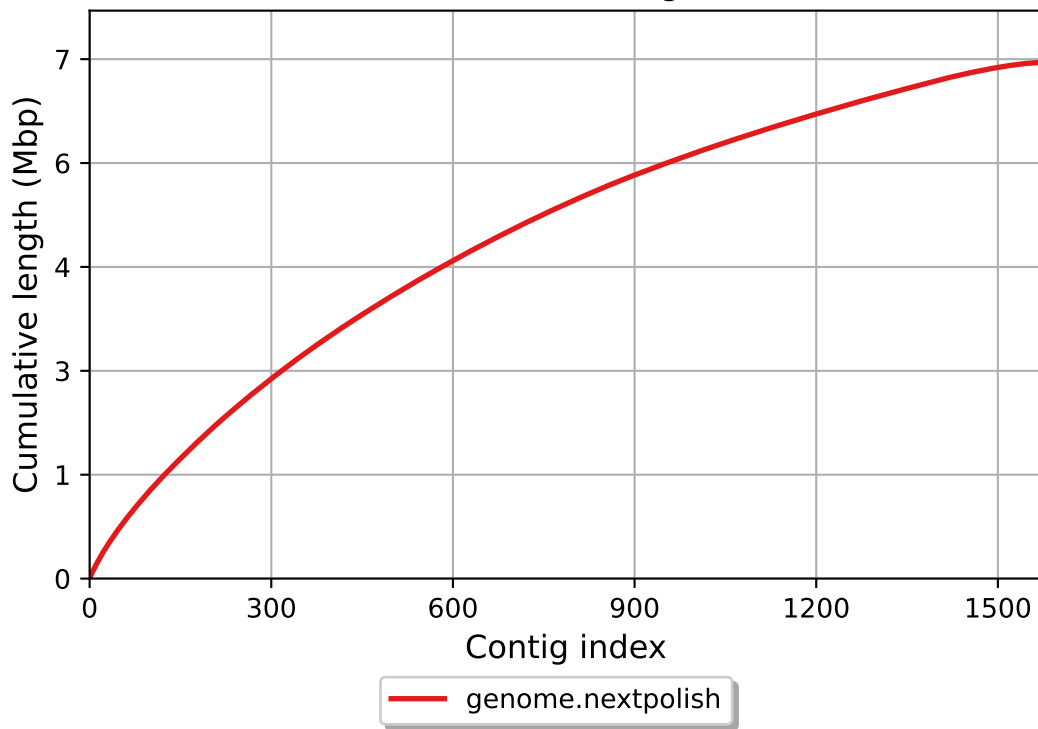
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

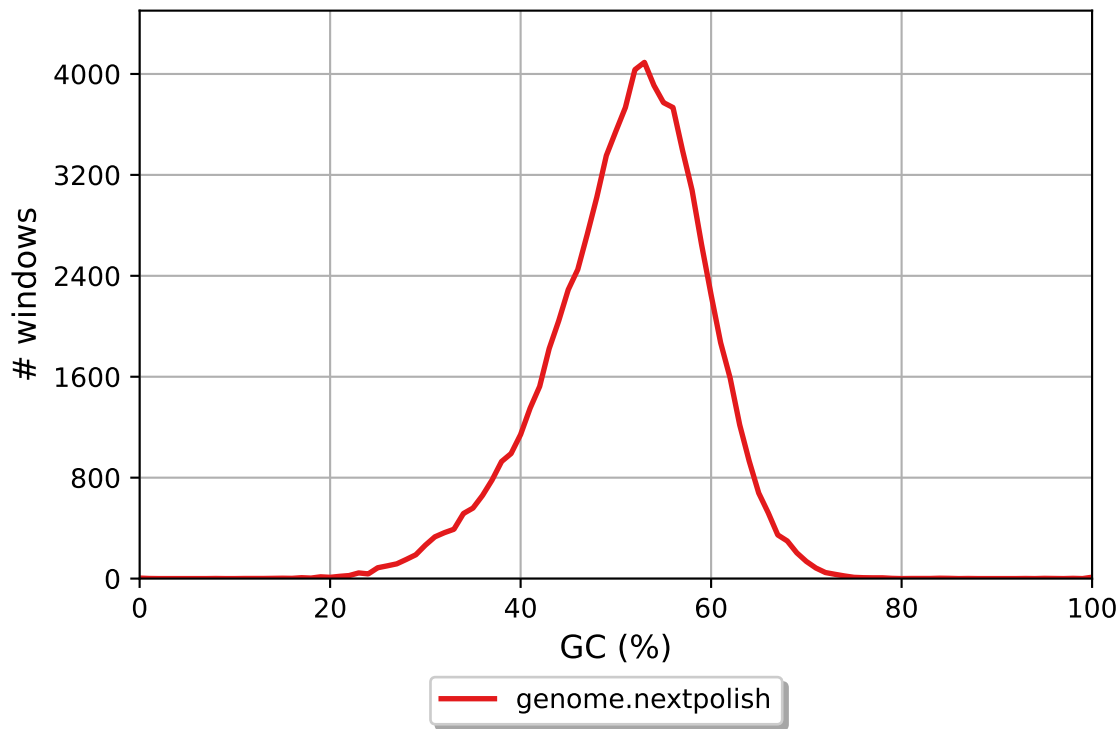


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

