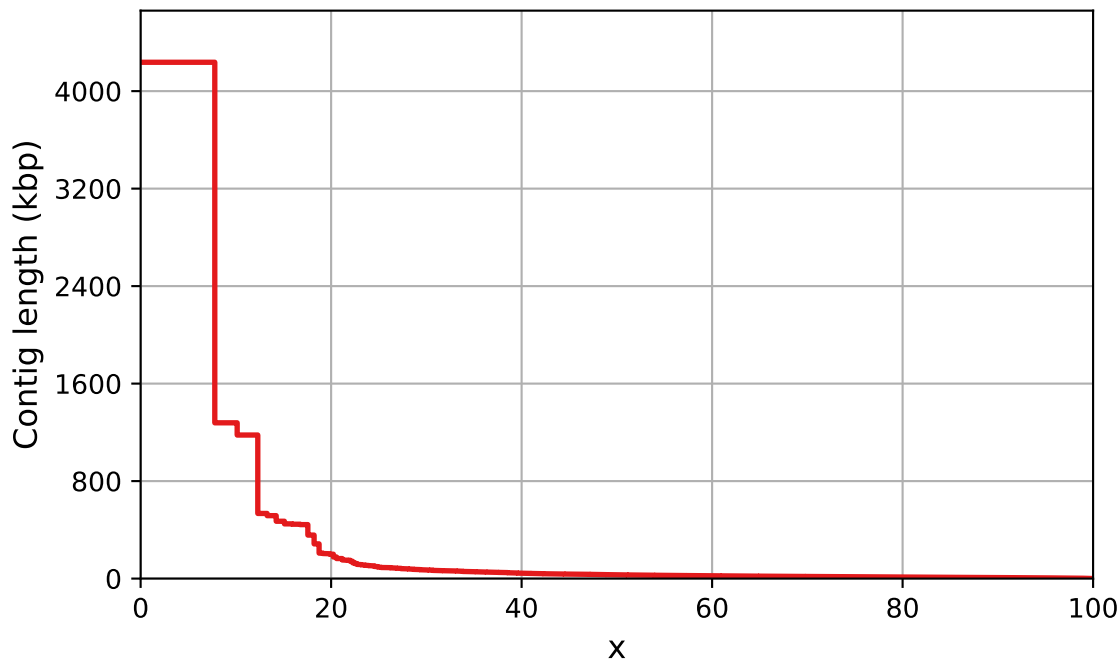


Report

	genome.nextpolish
# contigs (>= 0 bp)	2851
# contigs (>= 1000 bp)	2829
# contigs (>= 5000 bp)	2201
# contigs (>= 10000 bp)	1438
# contigs (>= 25000 bp)	455
# contigs (>= 50000 bp)	146
Total length (>= 0 bp)	54409014
Total length (>= 1000 bp)	54393881
Total length (>= 5000 bp)	52368481
Total length (>= 10000 bp)	46712255
Total length (>= 25000 bp)	31085146
Total length (>= 50000 bp)	20824976
# contigs	2848
Largest contig	4236804
Total length	54407640
GC (%)	54.91
N50	31086
N90	8319
auN	445059.7
L50	315
L90	1685
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	61 + 6 part

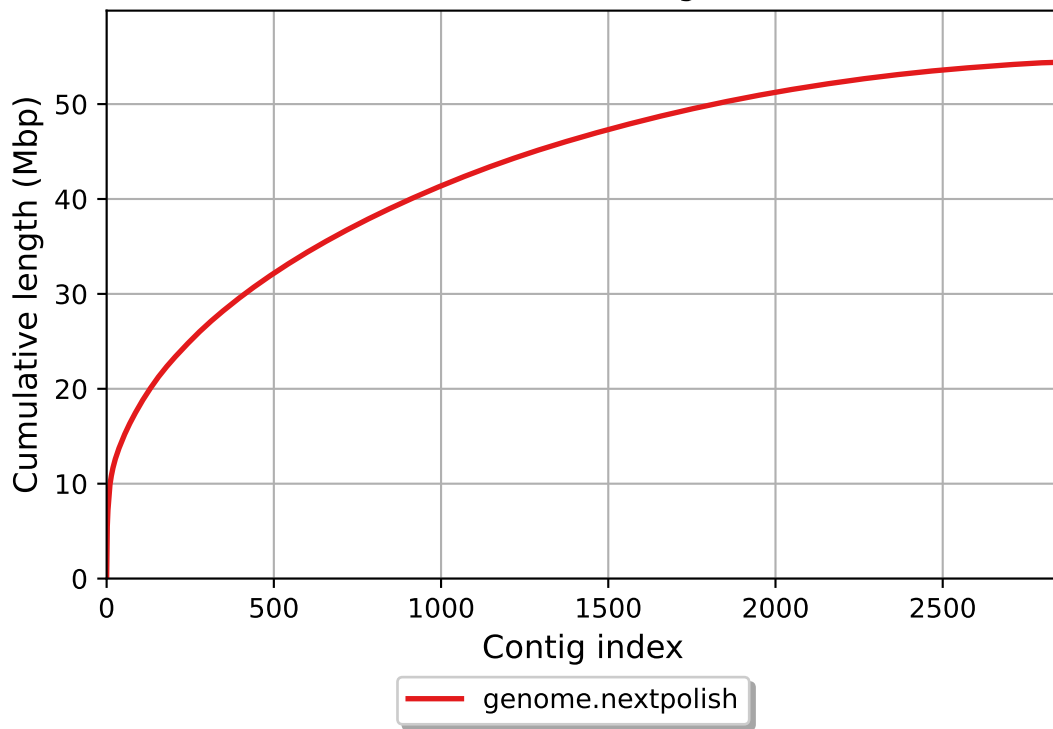
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

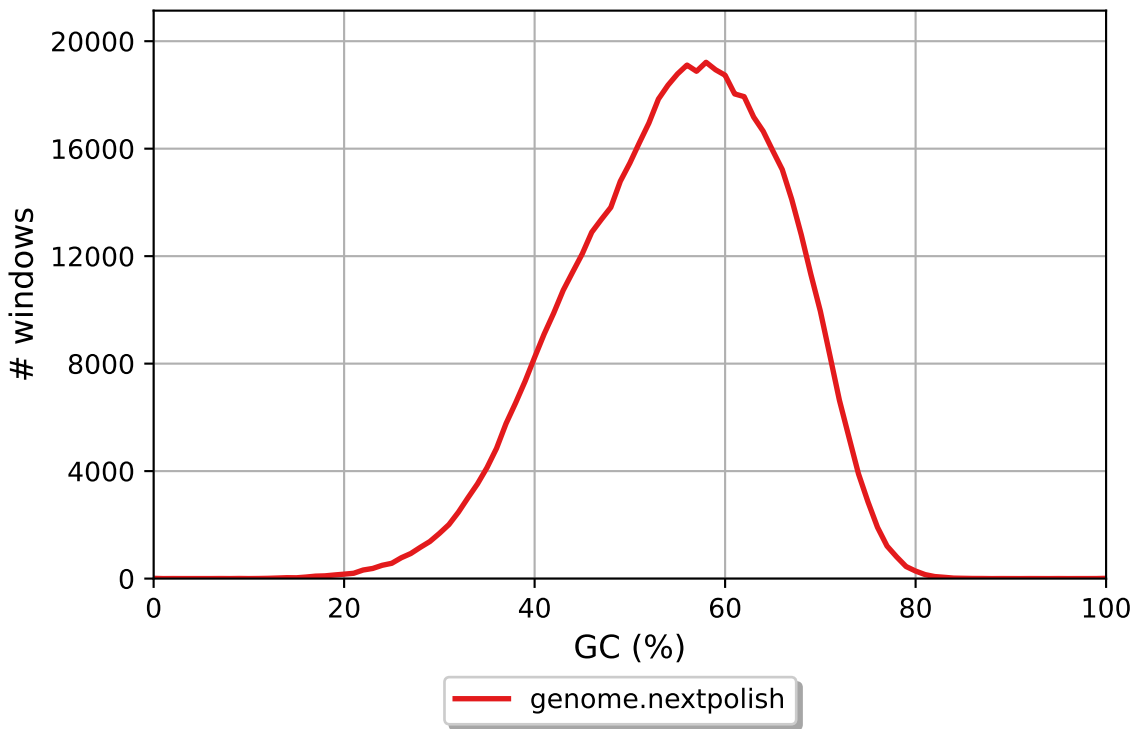


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

