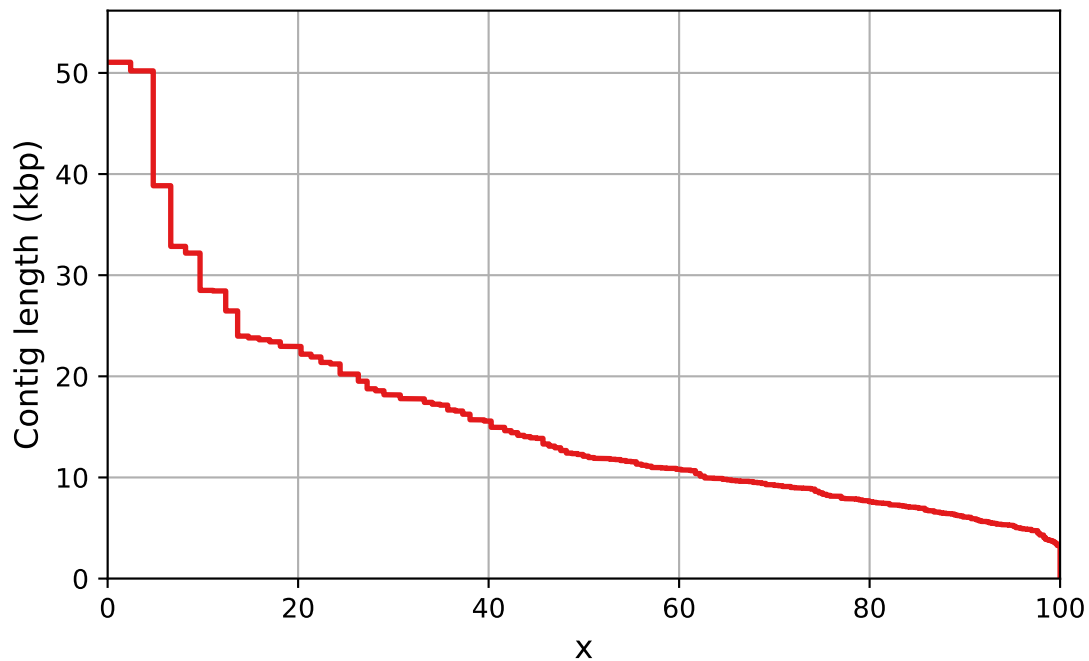


Report

	genome.nextpolish
# contigs (>= 0 bp)	191
# contigs (>= 1000 bp)	191
# contigs (>= 5000 bp)	170
# contigs (>= 10000 bp)	76
# contigs (>= 25000 bp)	8
# contigs (>= 50000 bp)	2
Total length (>= 0 bp)	2113859
Total length (>= 1000 bp)	2113859
Total length (>= 5000 bp)	2025453
Total length (>= 10000 bp)	1325082
Total length (>= 25000 bp)	288536
Total length (>= 50000 bp)	101254
# contigs	191
Largest contig	51053
Total length	2113859
GC (%)	58.47
N50	12099
N90	6088
auN	16008.7
L50	53
L90	148
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 0 part

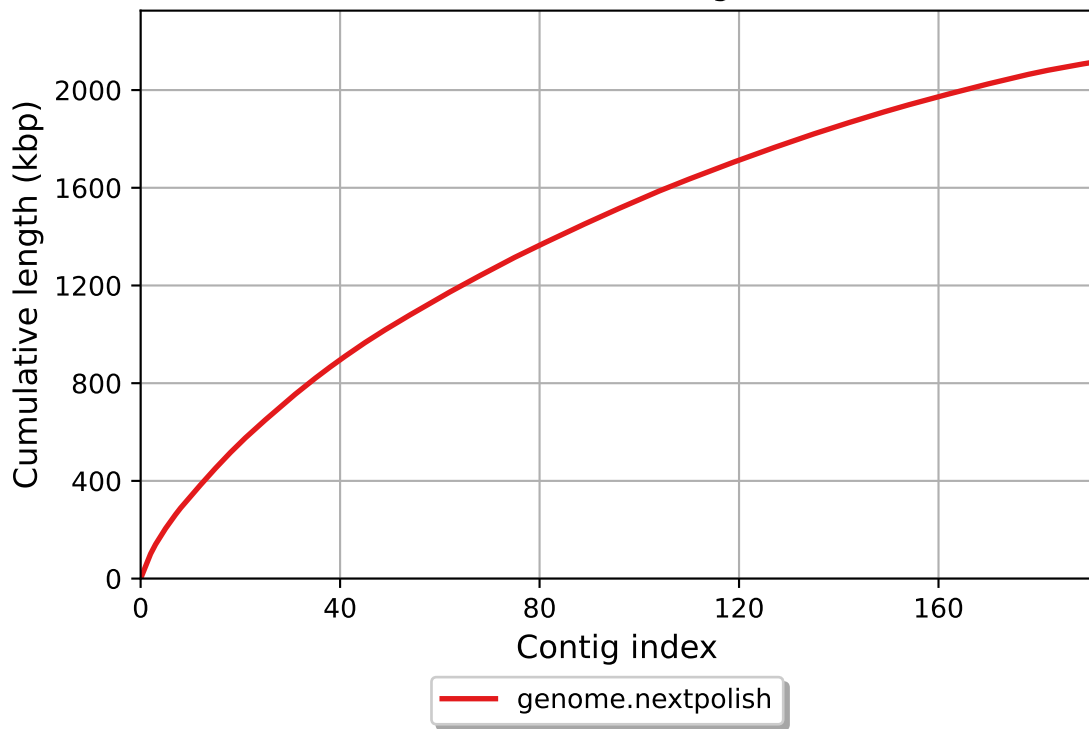
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

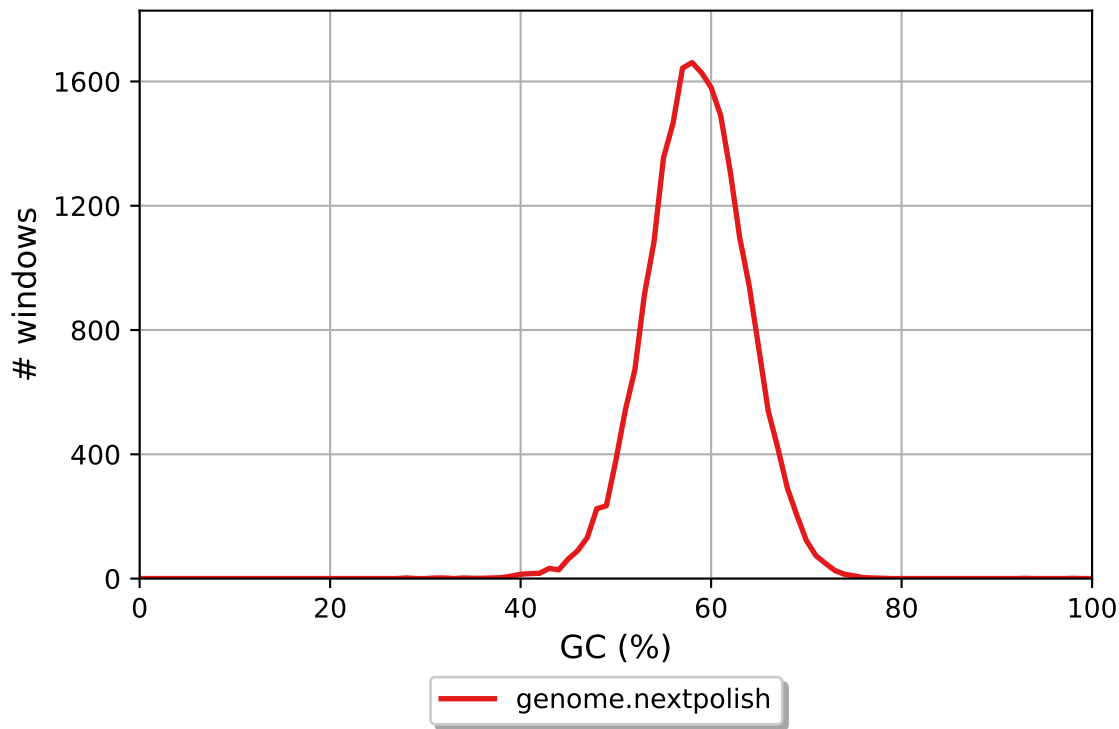


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

