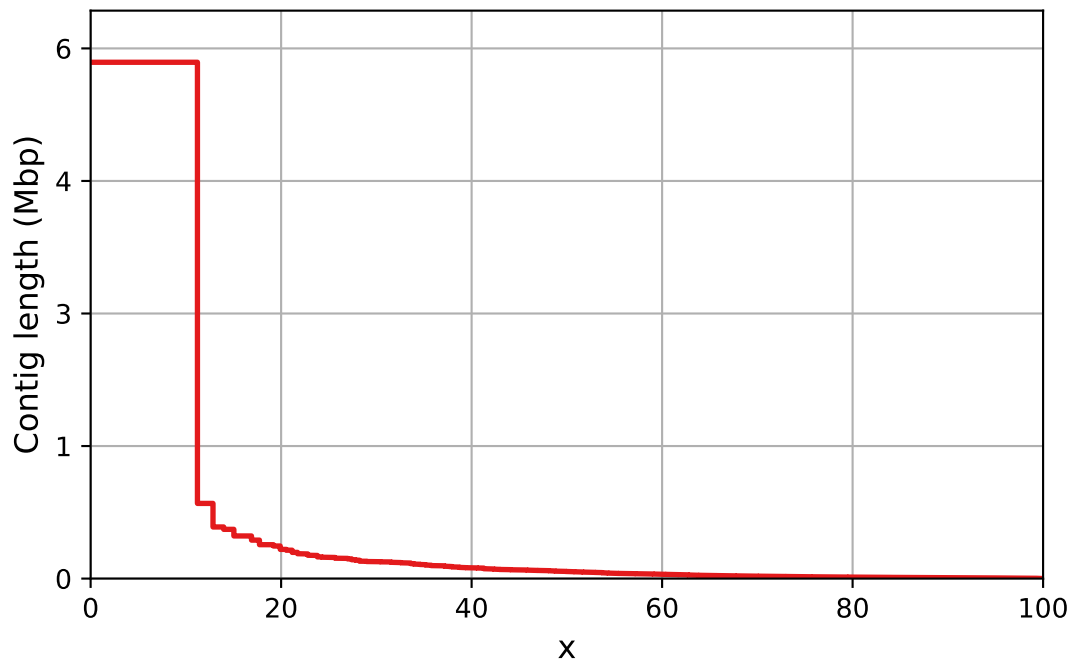


Report

	genome.nextpolish
# contigs (>= 0 bp)	1875
# contigs (>= 1000 bp)	1866
# contigs (>= 5000 bp)	1534
# contigs (>= 10000 bp)	1001
# contigs (>= 25000 bp)	387
# contigs (>= 50000 bp)	203
Total length (>= 0 bp)	52101236
Total length (>= 1000 bp)	52095702
Total length (>= 5000 bp)	50864766
Total length (>= 10000 bp)	46892912
Total length (>= 25000 bp)	37425822
Total length (>= 50000 bp)	31237722
# contigs	1875
Largest contig	5843176
Total length	52101236
GC (%)	56.93
N50	80559
N90	10003
auN	767407.7
L50	119
L90	1001
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	86 + 14 part

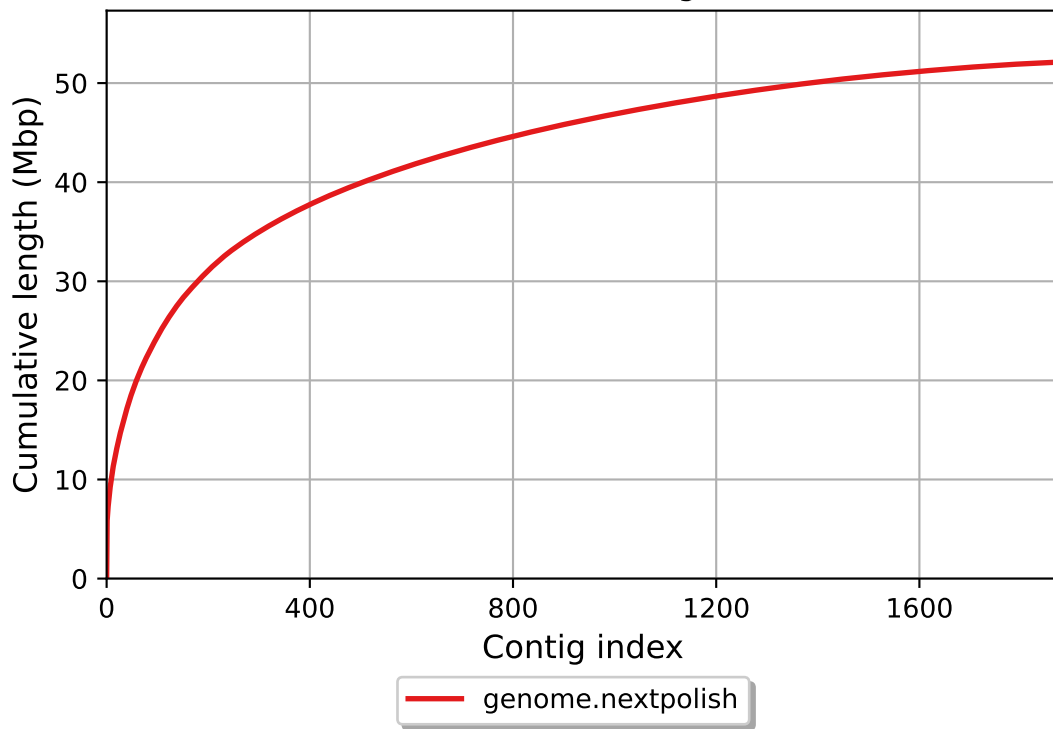
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx

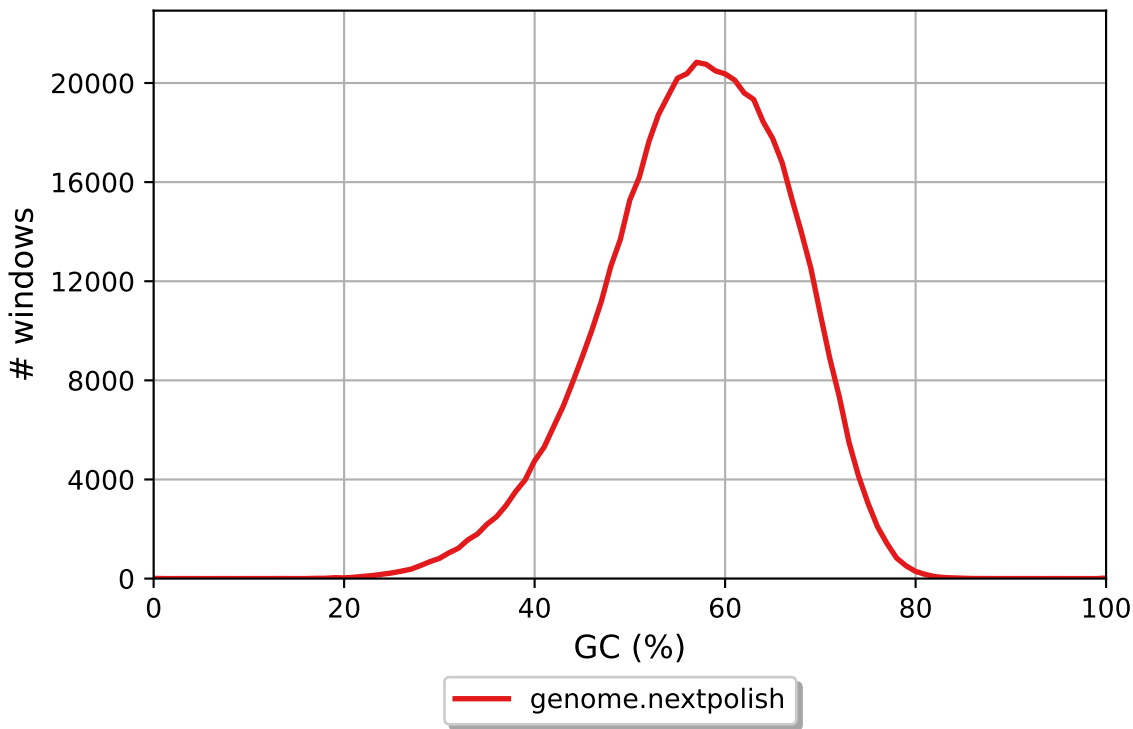


— genome.nextpolish

Cumulative length



GC content



genome.nextpolish GC content

