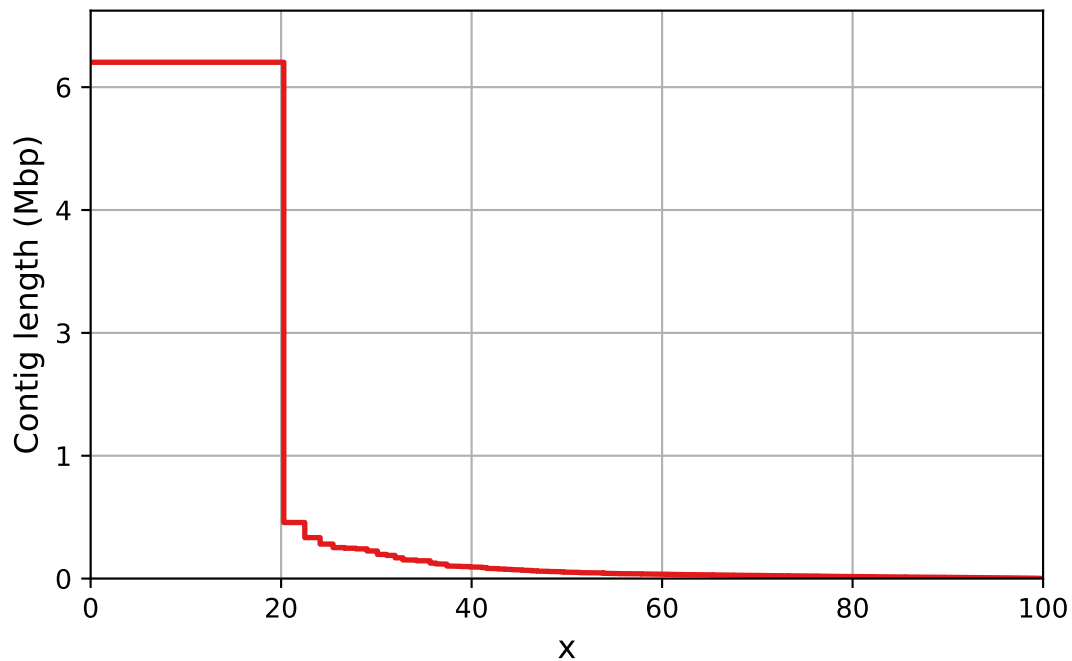


# Report

	genome.nextpolish
# contigs (>= 0 bp)	817
# contigs (>= 1000 bp)	815
# contigs (>= 5000 bp)	732
# contigs (>= 10000 bp)	554
# contigs (>= 25000 bp)	268
# contigs (>= 50000 bp)	111
Total length (>= 0 bp)	31081158
Total length (>= 1000 bp)	31079770
Total length (>= 5000 bp)	30778657
Total length (>= 10000 bp)	29449914
Total length (>= 25000 bp)	24774788
Total length (>= 50000 bp)	19093914
# contigs	817
Largest contig	6304013
Total length	31081158
GC (%)	55.53
N50	78561
N90	14769
auN	1373008.6
L50	54
L90	433
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	68 + 6 part

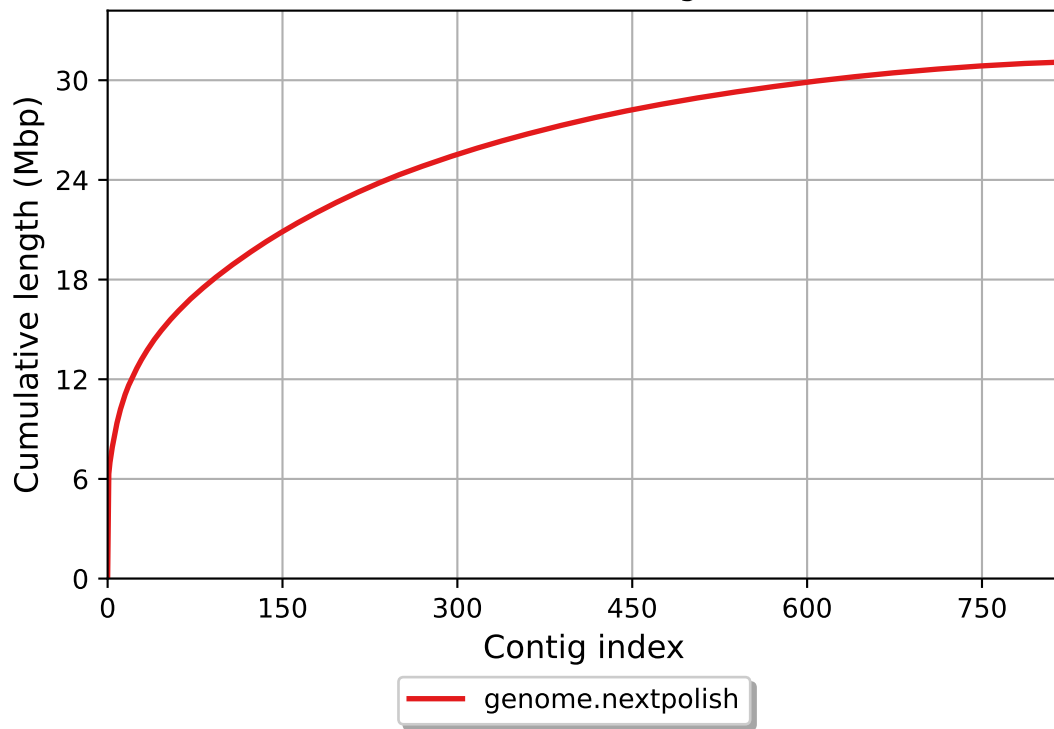
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

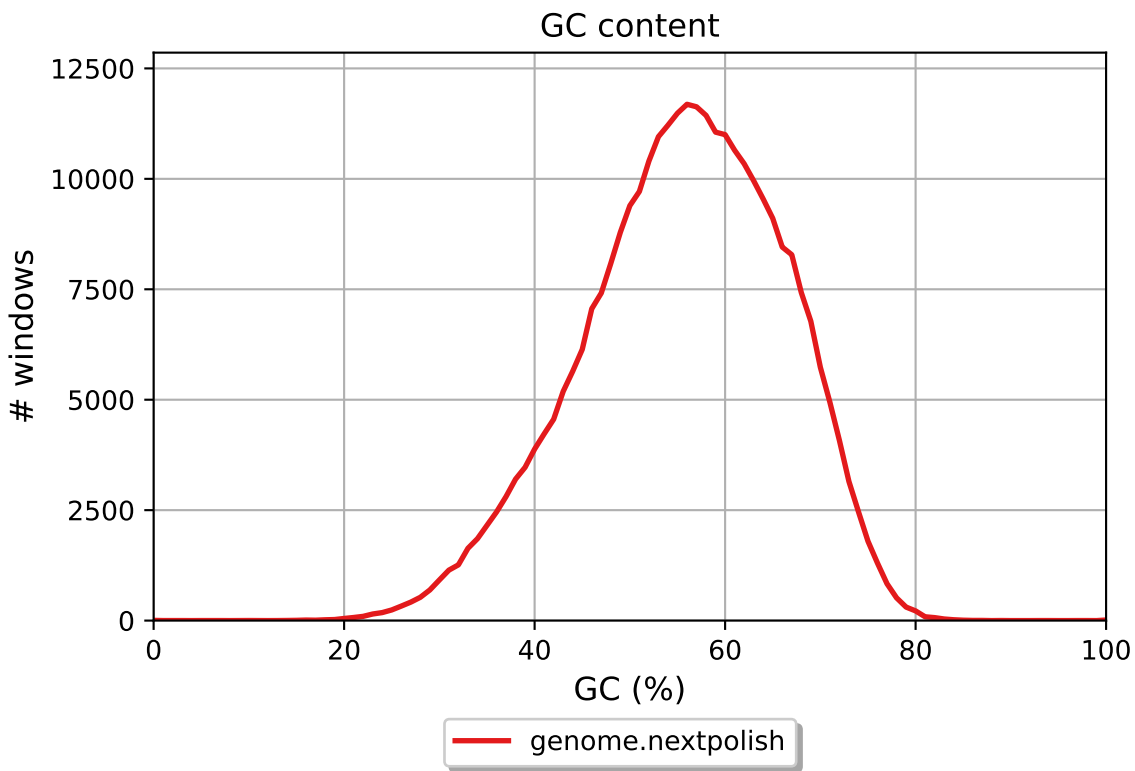
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

