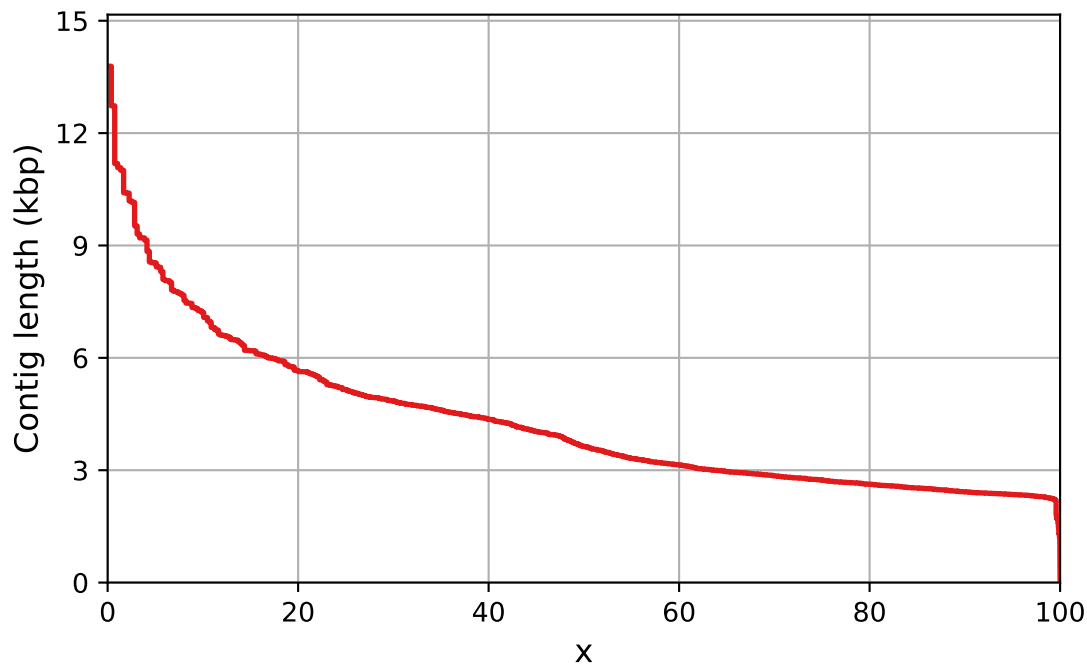


Report

	genome.nextpolish
# contigs (>= 0 bp)	986
# contigs (>= 1000 bp)	986
# contigs (>= 5000 bp)	144
# contigs (>= 10000 bp)	9
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	3563758
Total length (>= 1000 bp)	3563758
Total length (>= 5000 bp)	959266
Total length (>= 10000 bp)	100926
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	986
Largest contig	13784
Total length	3563758
GC (%)	51.14
N50	3635
N90	2424
auN	4305.5
L50	333
L90	831
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	4 + 1 part

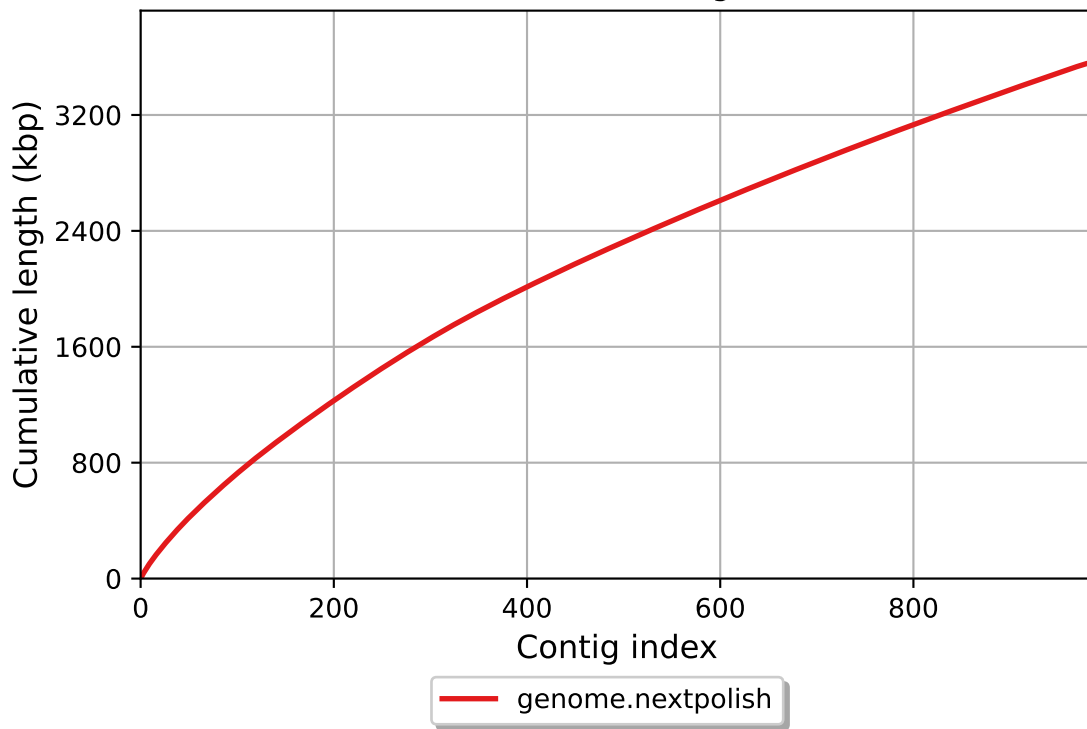
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

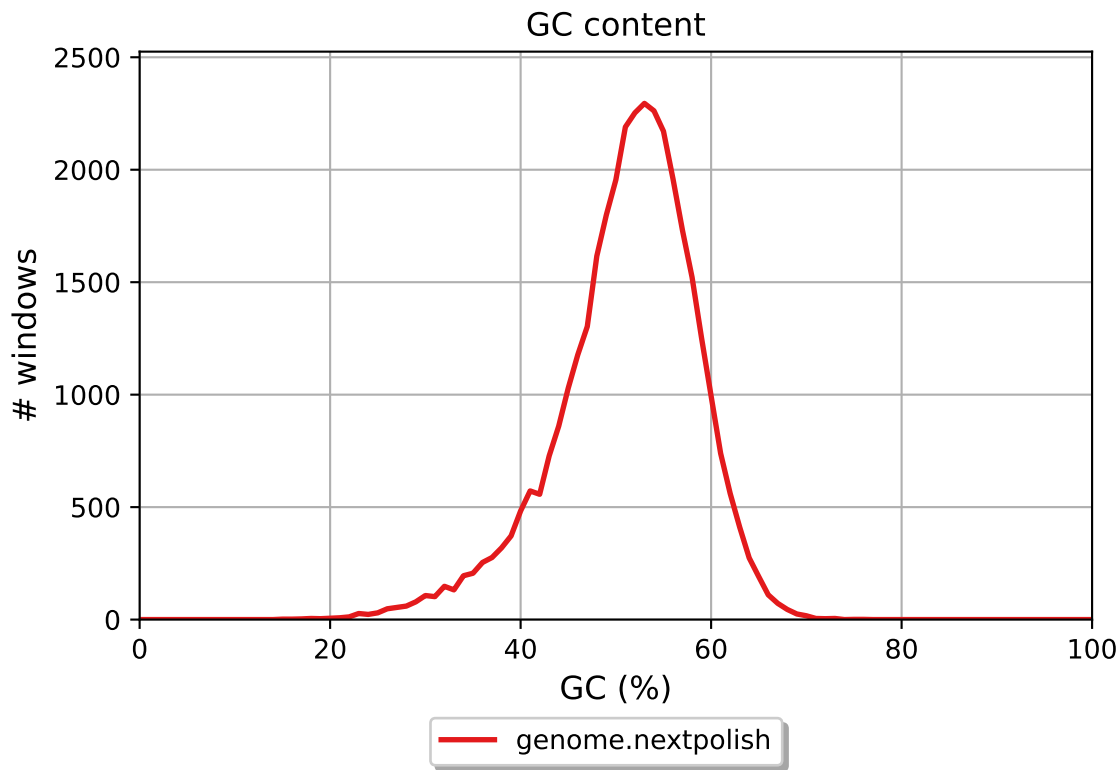
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

