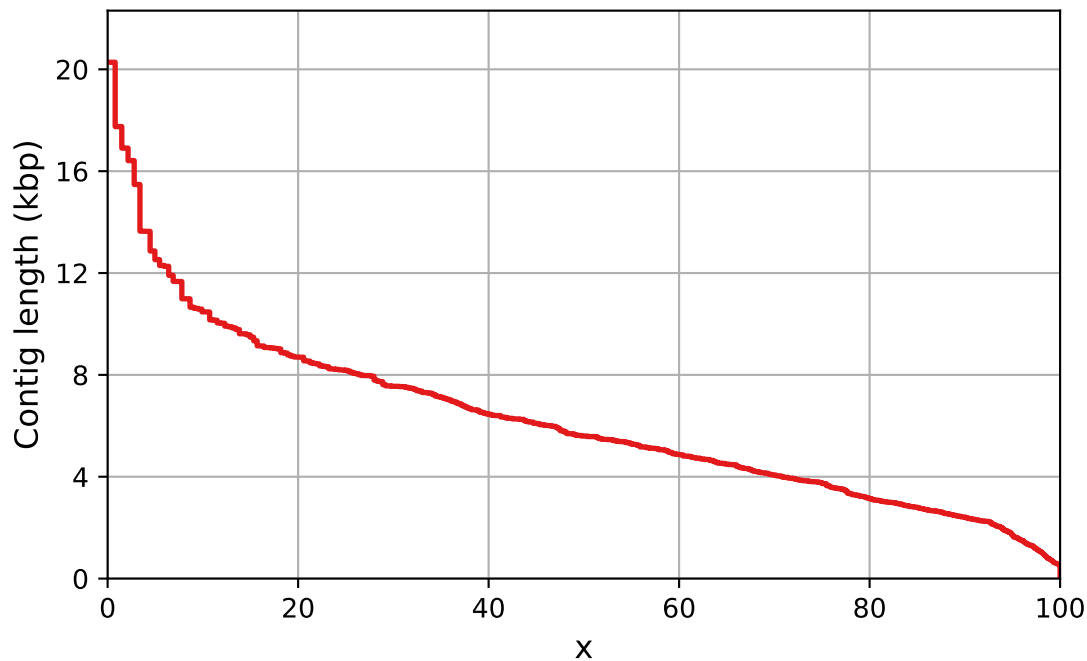


Report

	genome.nextpolish
# contigs (>= 0 bp)	616
# contigs (>= 1000 bp)	546
# contigs (>= 5000 bp)	199
# contigs (>= 10000 bp)	25
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2559133
Total length (>= 1000 bp)	2511571
Total length (>= 5000 bp)	1506290
Total length (>= 10000 bp)	314453
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	612
Largest contig	20277
Total length	2557756
GC (%)	39.88
N50	5599
N90	2414
auN	6245.5
L50	157
L90	425
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	2 + 0 part

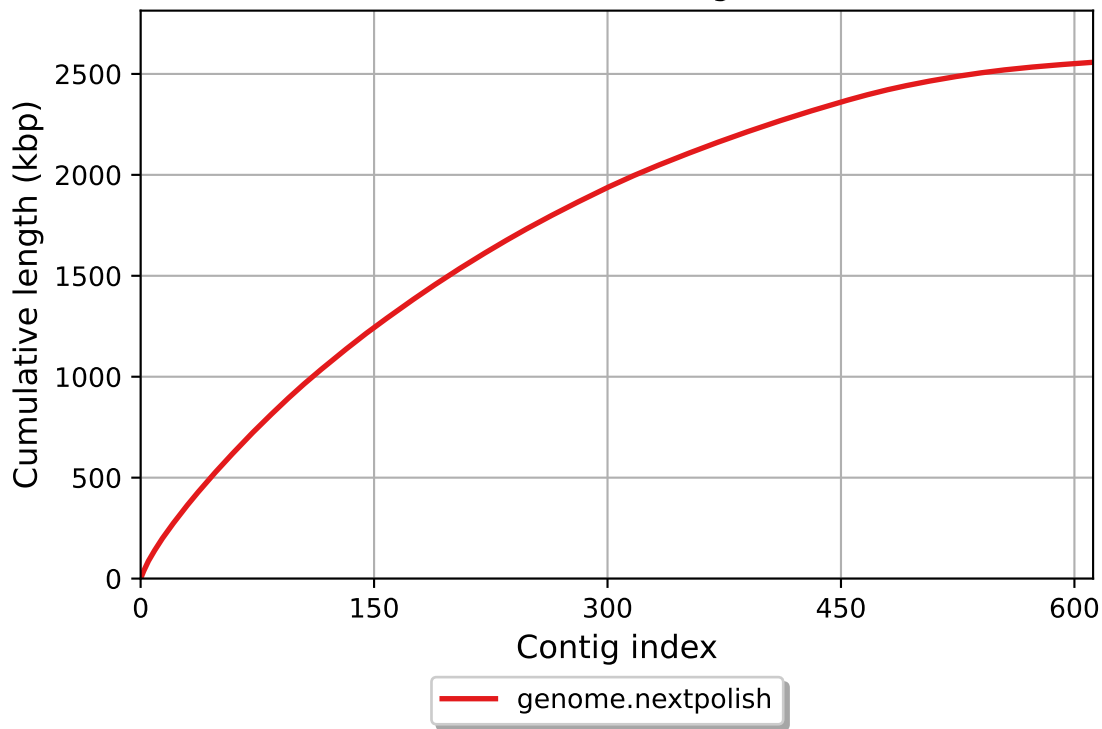
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

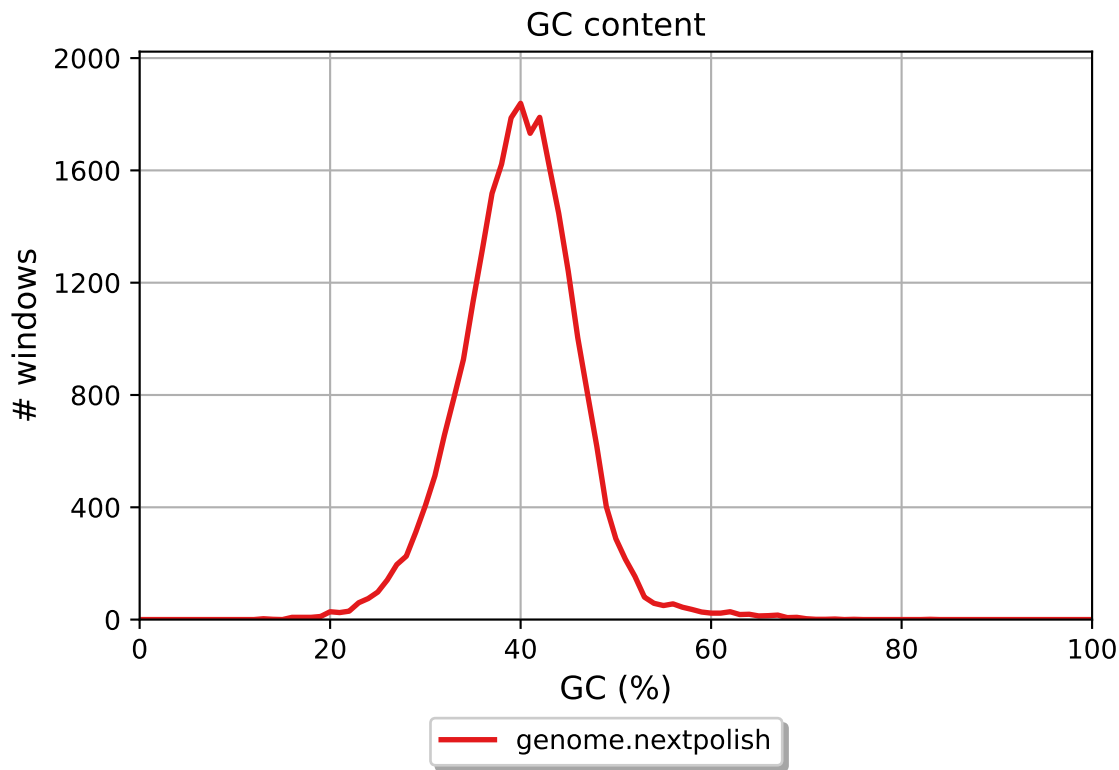
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

