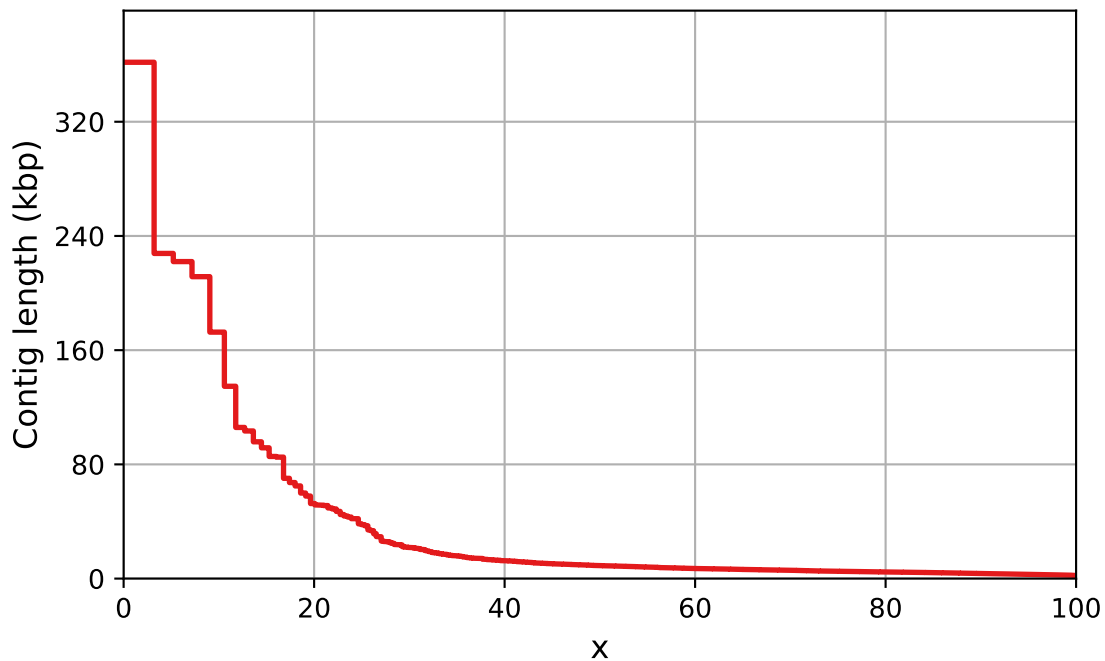


Report

	genome.nextpolish
# contigs (>= 0 bp)	1432
# contigs (>= 1000 bp)	1431
# contigs (>= 5000 bp)	674
# contigs (>= 10000 bp)	187
# contigs (>= 25000 bp)	42
# contigs (>= 50000 bp)	21
Total length (>= 0 bp)	11302780
Total length (>= 1000 bp)	11302215
Total length (>= 5000 bp)	8594650
Total length (>= 10000 bp)	5241856
Total length (>= 25000 bp)	3185548
Total length (>= 50000 bp)	2424436
# contigs	1432
Largest contig	361661
Total length	11302780
GC (%)	45.71
N50	9086
N90	3558
auN	44621.4
L50	230
L90	1042
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	30 + 2 part

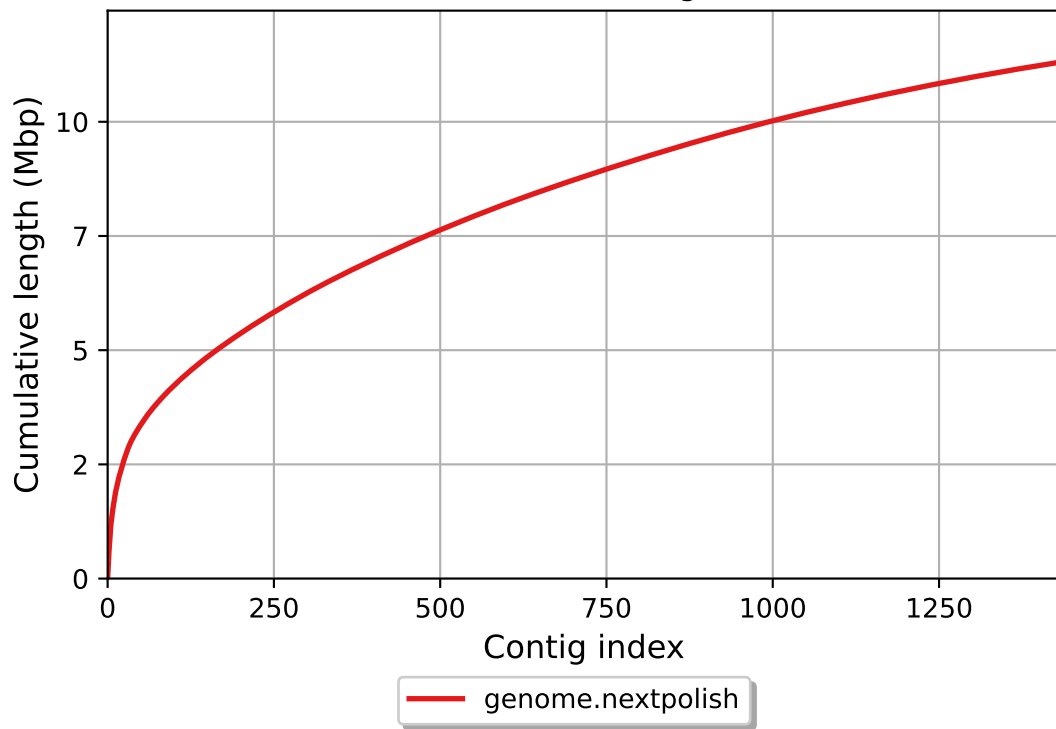
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

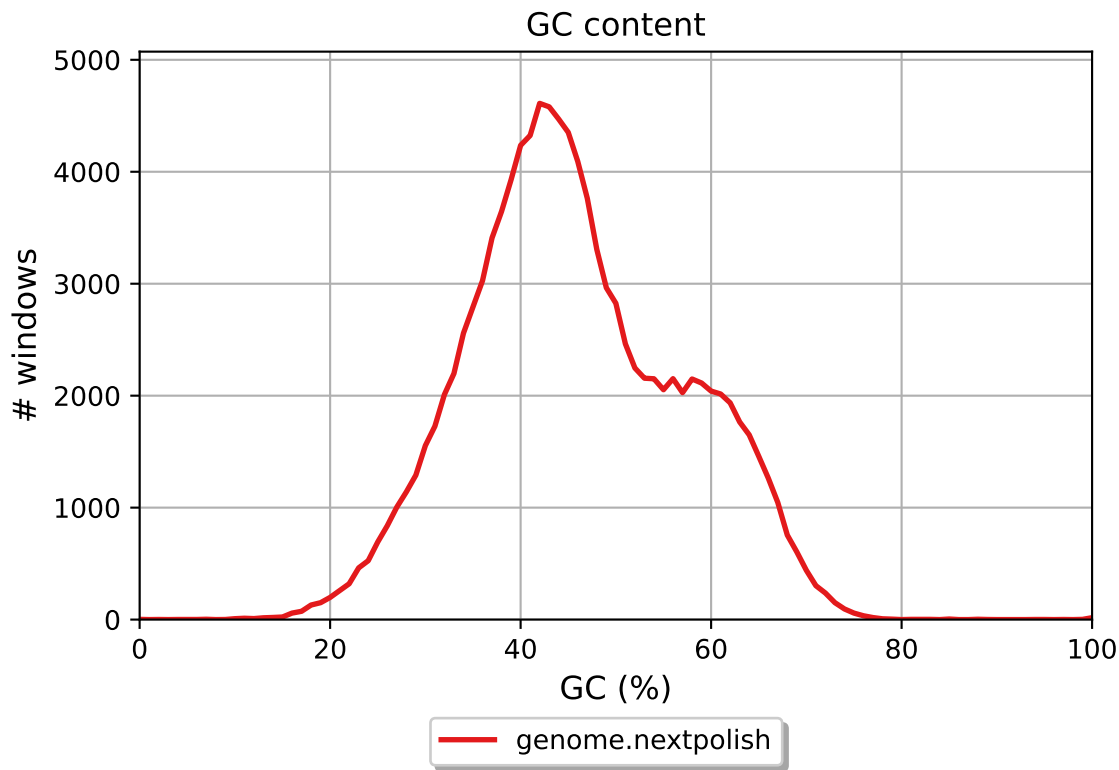
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

