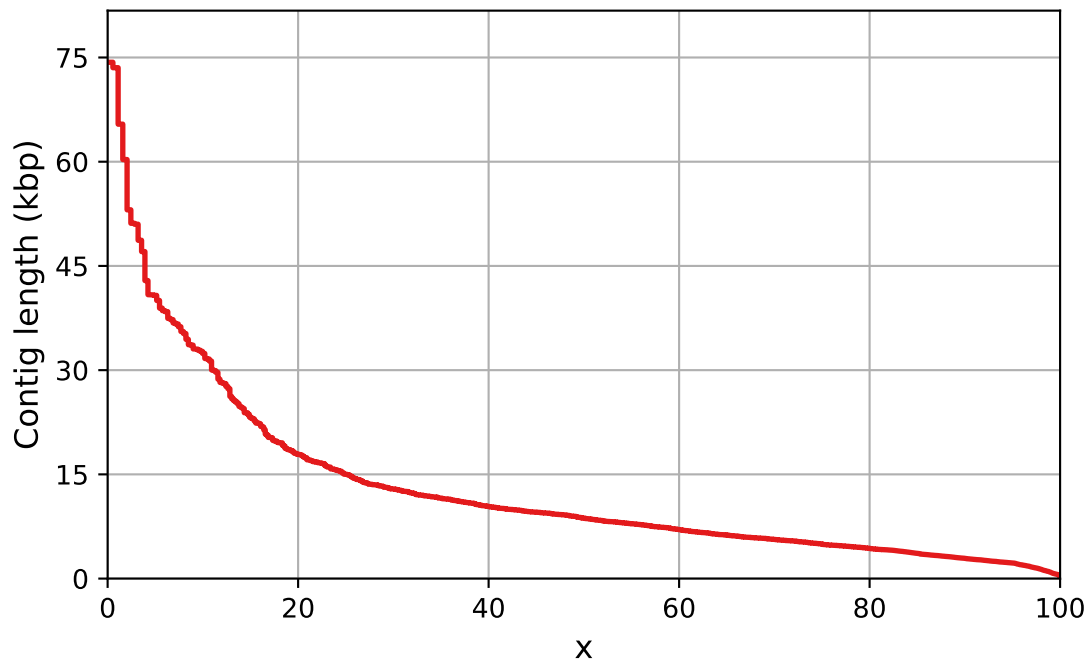


Report

	genome.nextpolish
# contigs (>= 0 bp)	2290
# contigs (>= 1000 bp)	2049
# contigs (>= 5000 bp)	944
# contigs (>= 10000 bp)	319
# contigs (>= 25000 bp)	49
# contigs (>= 50000 bp)	7
Total length (>= 0 bp)	13399425
Total length (>= 1000 bp)	13238647
Total length (>= 5000 bp)	10038772
Total length (>= 10000 bp)	5607827
Total length (>= 25000 bp)	1847446
Total length (>= 50000 bp)	428774
# contigs	2276
Largest contig	74312
Total length	13394430
GC (%)	44.54
N50	8699
N90	2916
auN	13098.6
L50	435
L90	1463
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	23 + 5 part

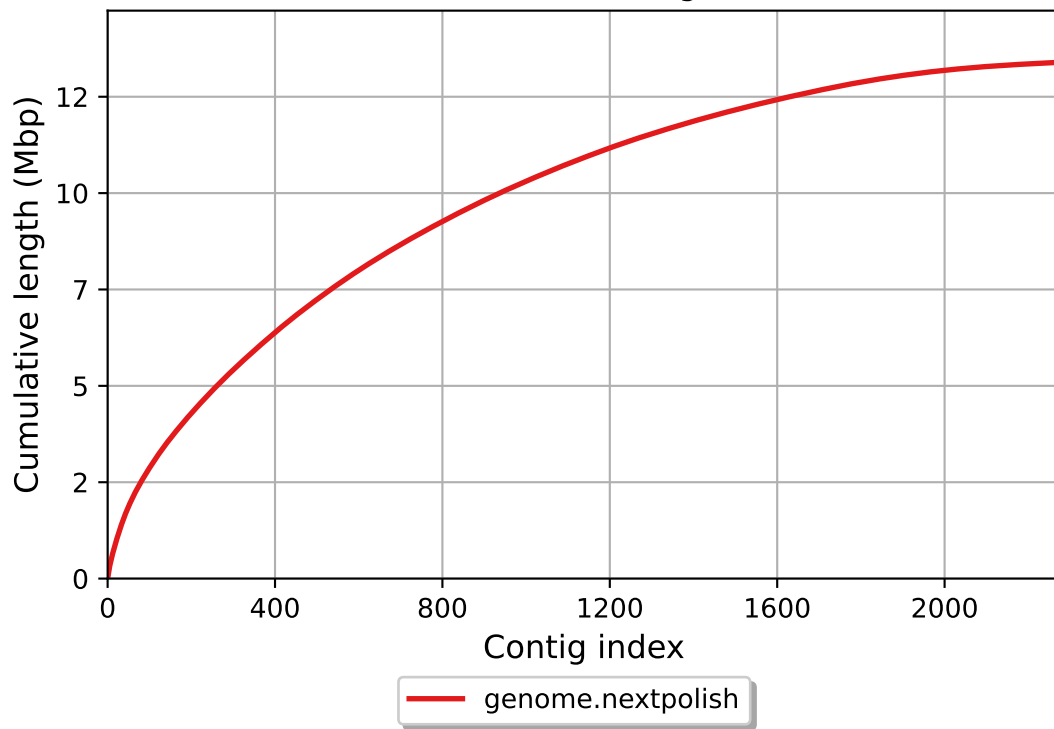
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

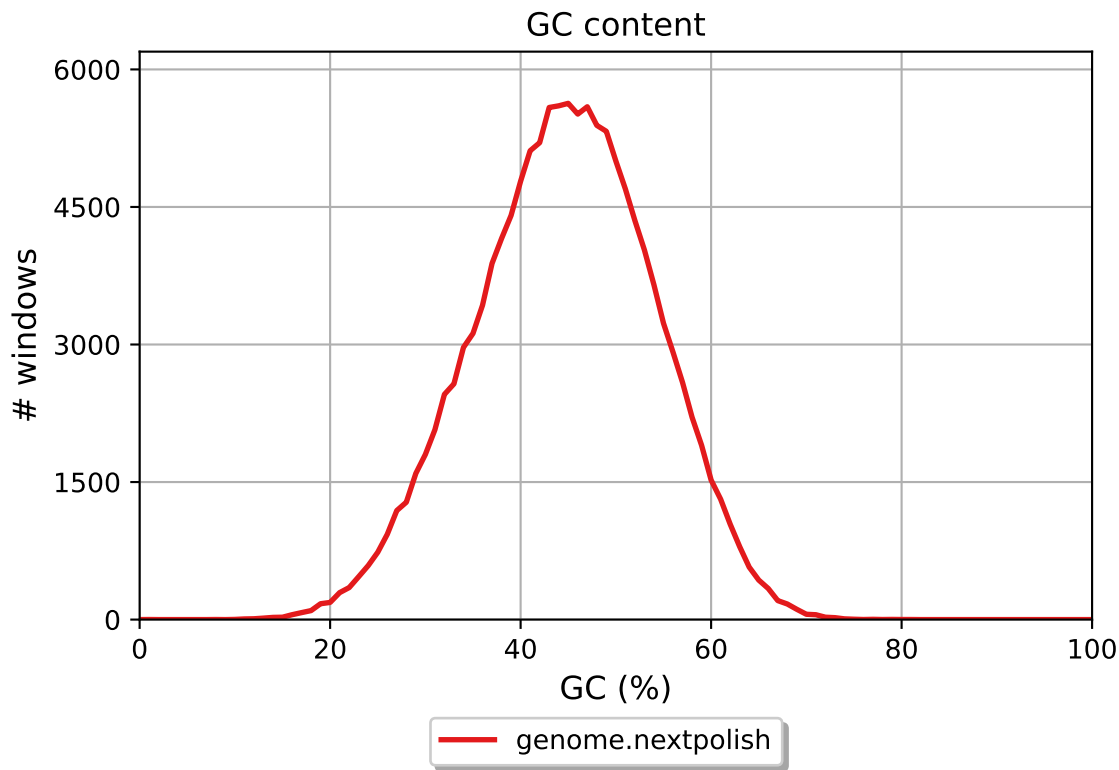
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

