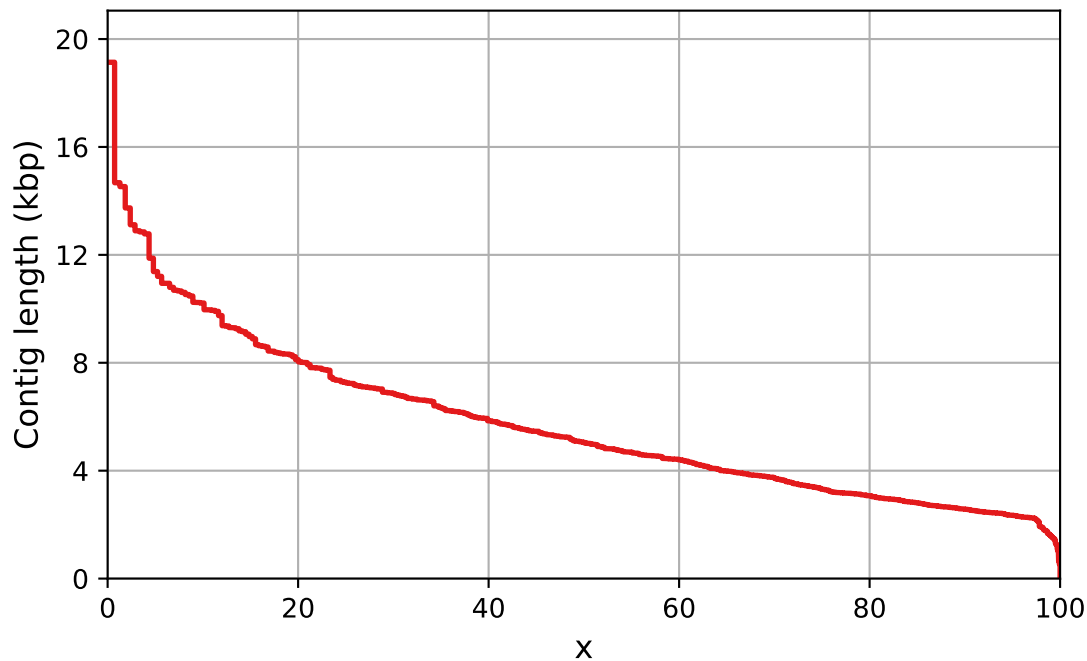


Report

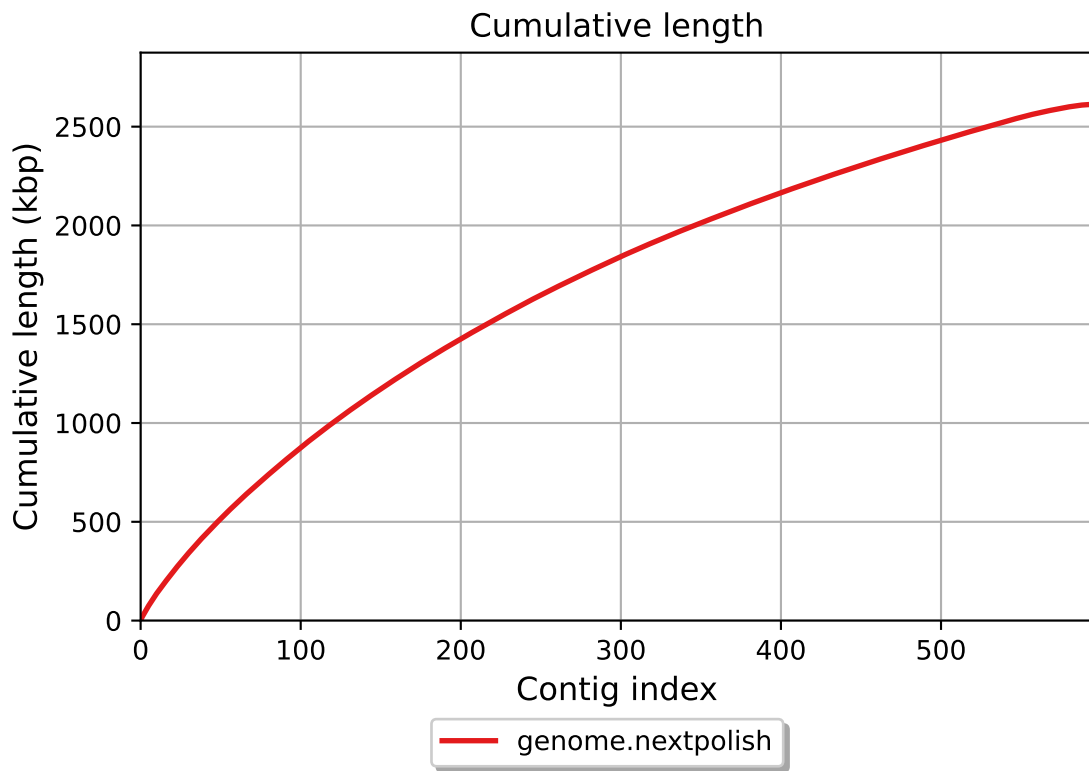
	genome.nextpolish
# contigs (>= 0 bp)	595
# contigs (>= 1000 bp)	587
# contigs (>= 5000 bp)	179
# contigs (>= 10000 bp)	22
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2613308
Total length (>= 1000 bp)	2608552
Total length (>= 5000 bp)	1324115
Total length (>= 10000 bp)	264517
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	595
Largest contig	19139
Total length	2613308
GC (%)	43.42
N50	5047
N90	2574
auN	5766.6
L50	176
L90	469
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	6 + 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).

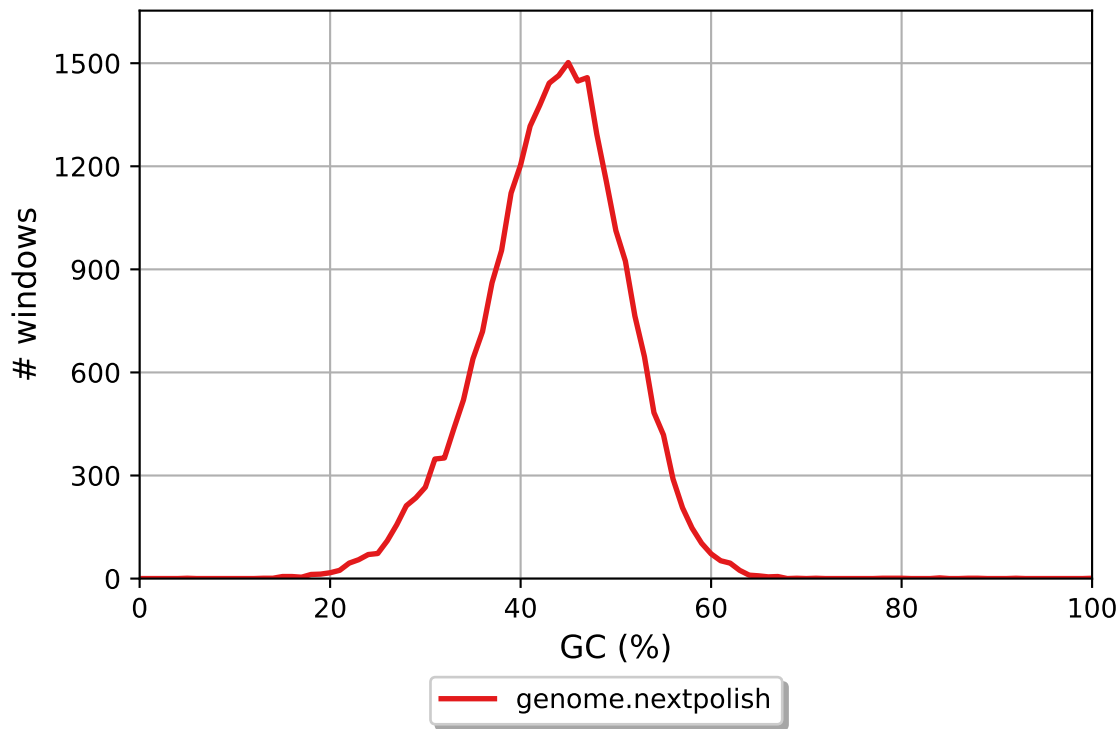
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

