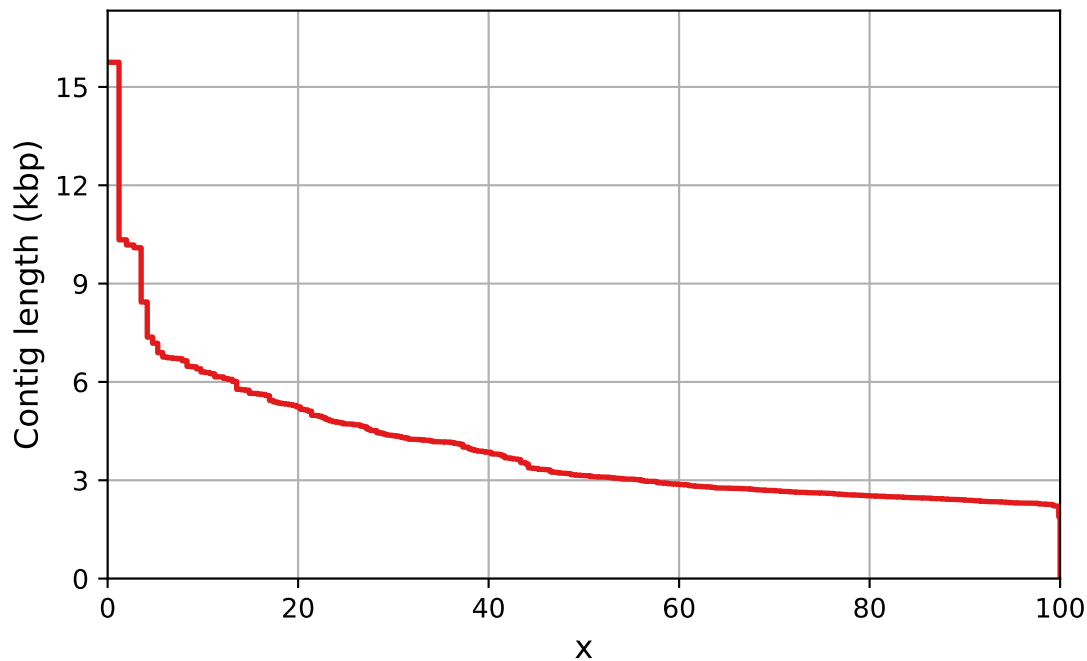


Report

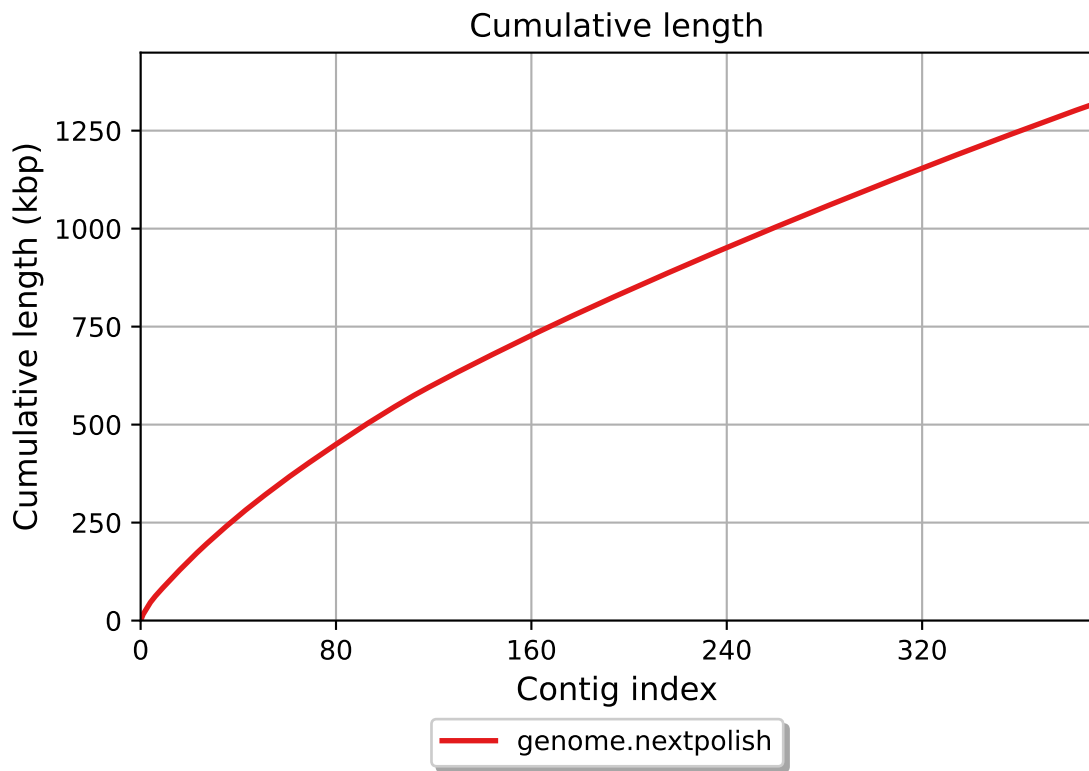
	genome.nextpolish
# contigs (>= 0 bp)	390
# contigs (>= 1000 bp)	390
# contigs (>= 5000 bp)	43
# contigs (>= 10000 bp)	4
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	1317292
Total length (>= 1000 bp)	1317292
Total length (>= 5000 bp)	281989
Total length (>= 10000 bp)	46360
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	390
Largest contig	15752
Total length	1317292
GC (%)	42.16
N50	3146
N90	2401
auN	3997.7
L50	138
L90	333
# 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).

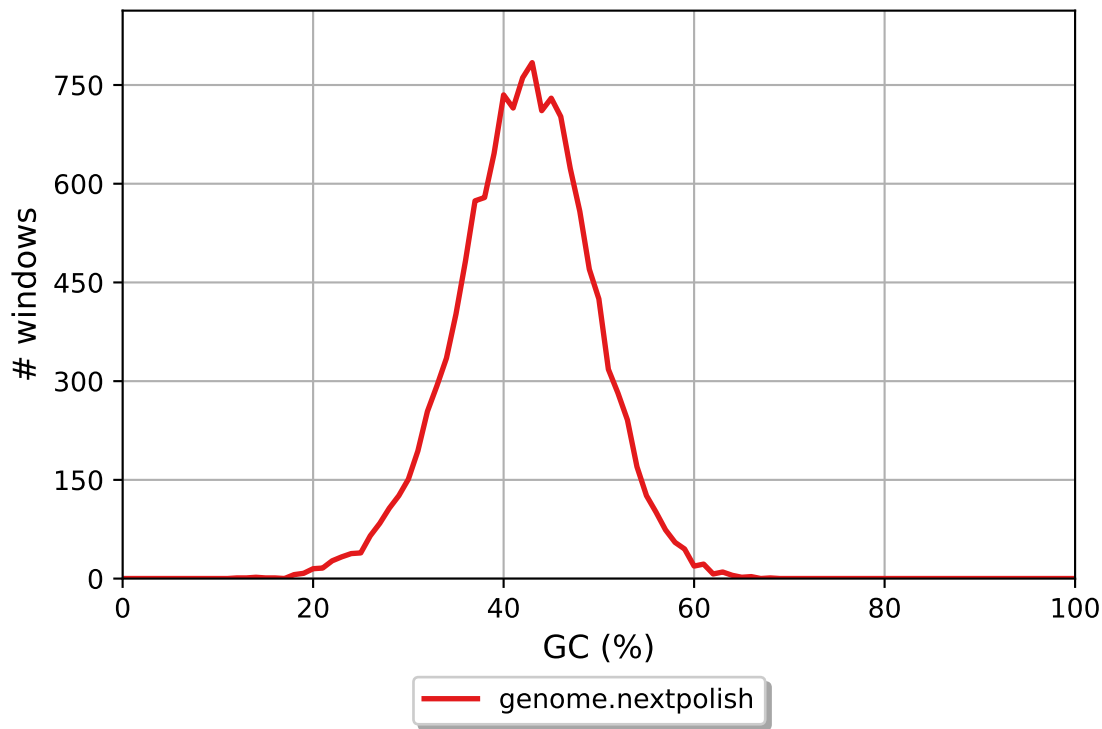
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

