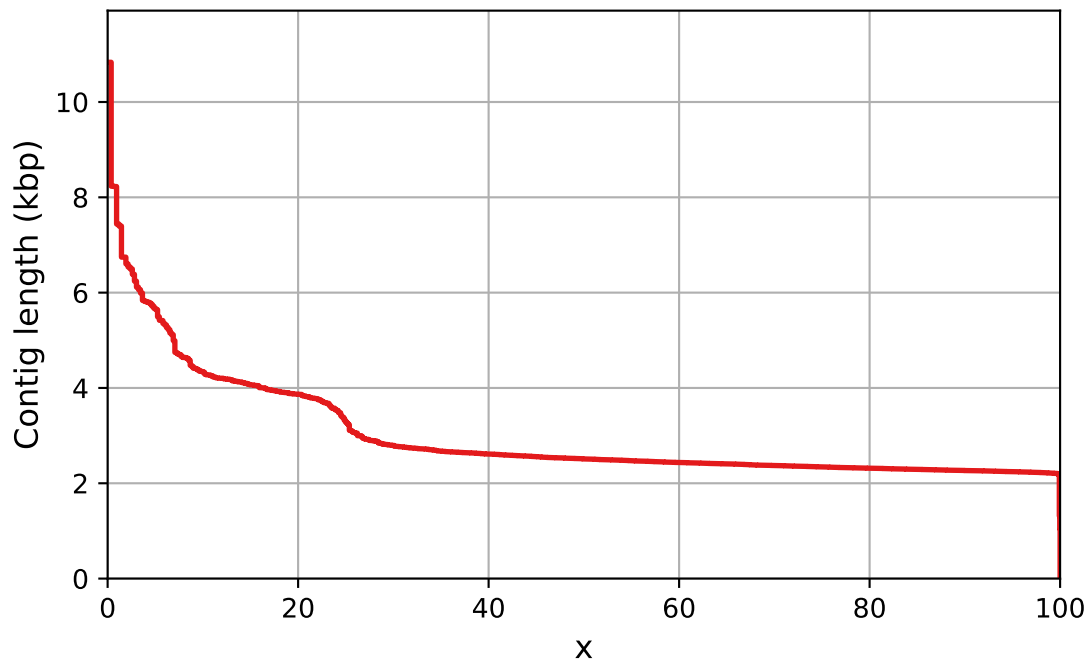


# Report

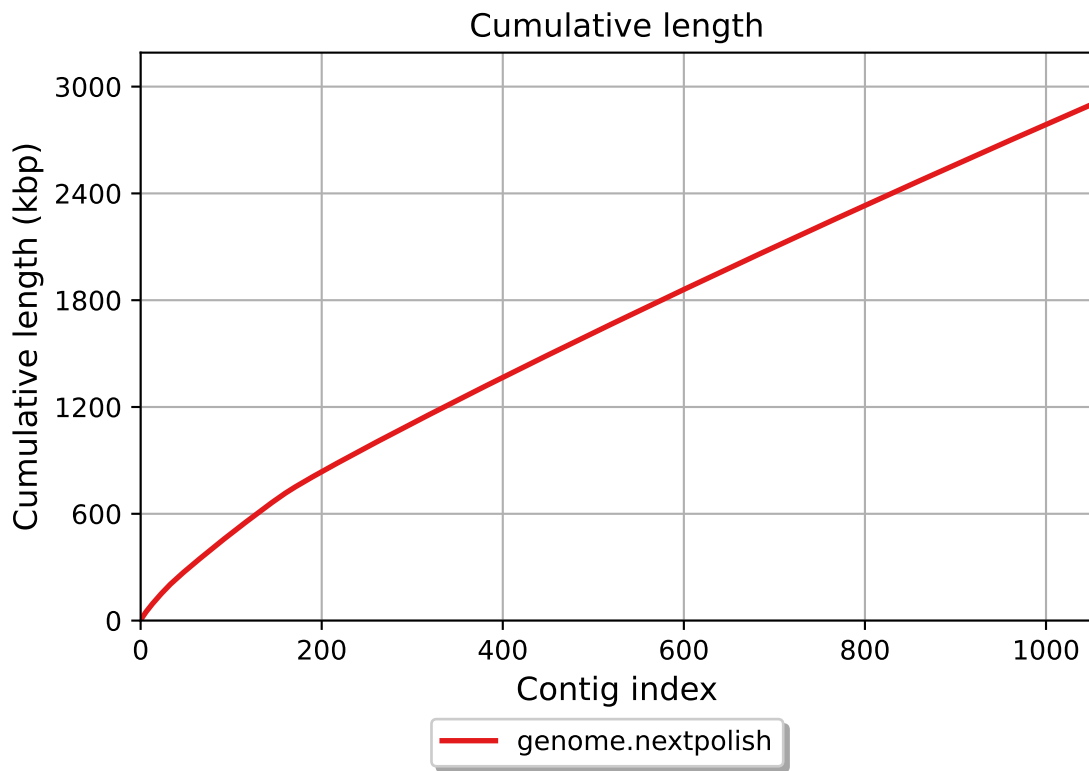
	genome.nextpolish
# contigs (>= 0 bp)	1052
# contigs (>= 1000 bp)	1052
# contigs (>= 5000 bp)	32
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	2900802
Total length (>= 1000 bp)	2900802
Total length (>= 5000 bp)	199857
Total length (>= 10000 bp)	10833
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	1052
Largest contig	10833
Total length	2900802
GC (%)	40.89
N50	2510
N90	2267
auN	3024.4
L50	434
L90	922
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	5 + 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).

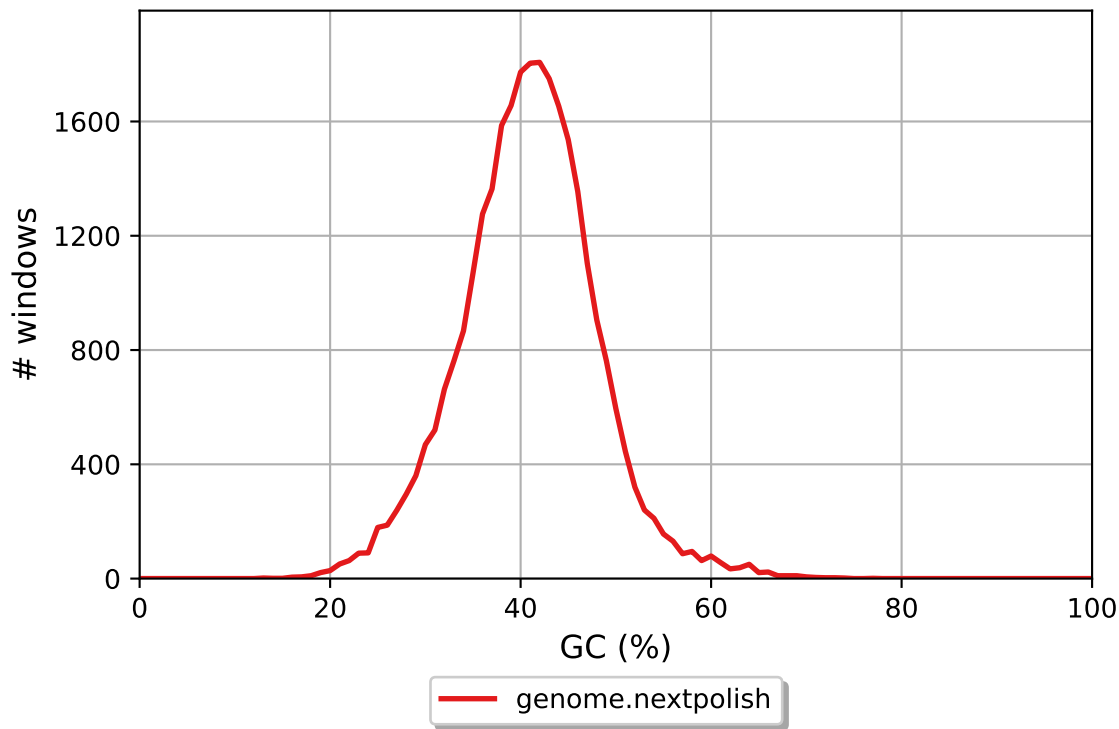
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

