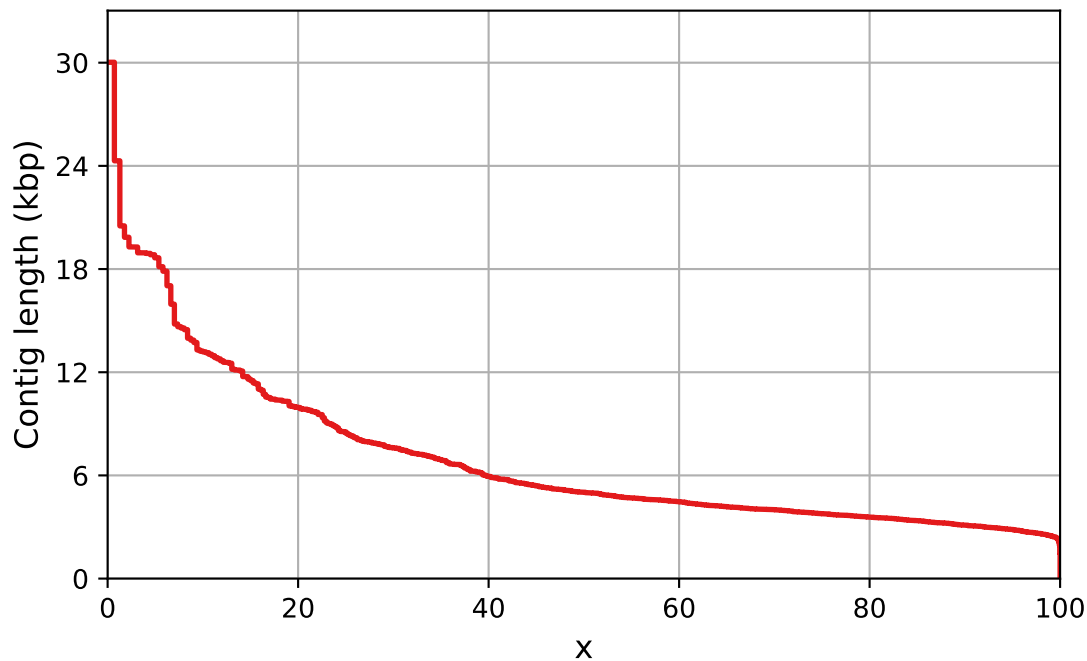


Report

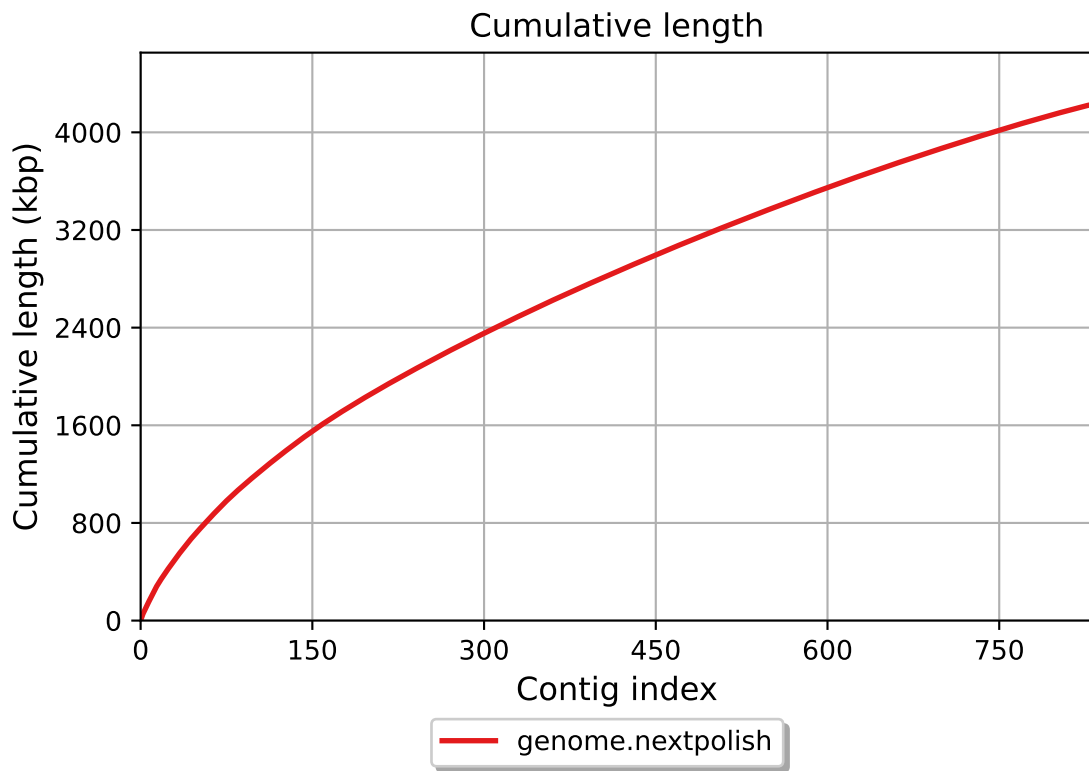
	genome.nextpolish
# contigs (>= 0 bp)	832
# contigs (>= 1000 bp)	832
# contigs (>= 5000 bp)	252
# contigs (>= 10000 bp)	59
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	4229959
Total length (>= 1000 bp)	4229959
Total length (>= 5000 bp)	2122069
Total length (>= 10000 bp)	825955
Total length (>= 25000 bp)	30024
Total length (>= 50000 bp)	0
# contigs	832
Largest contig	30024
Total length	4229959
GC (%)	39.19
N50	5019
N90	3104
auN	6973.6
L50	251
L90	680
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	8 + 1 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).

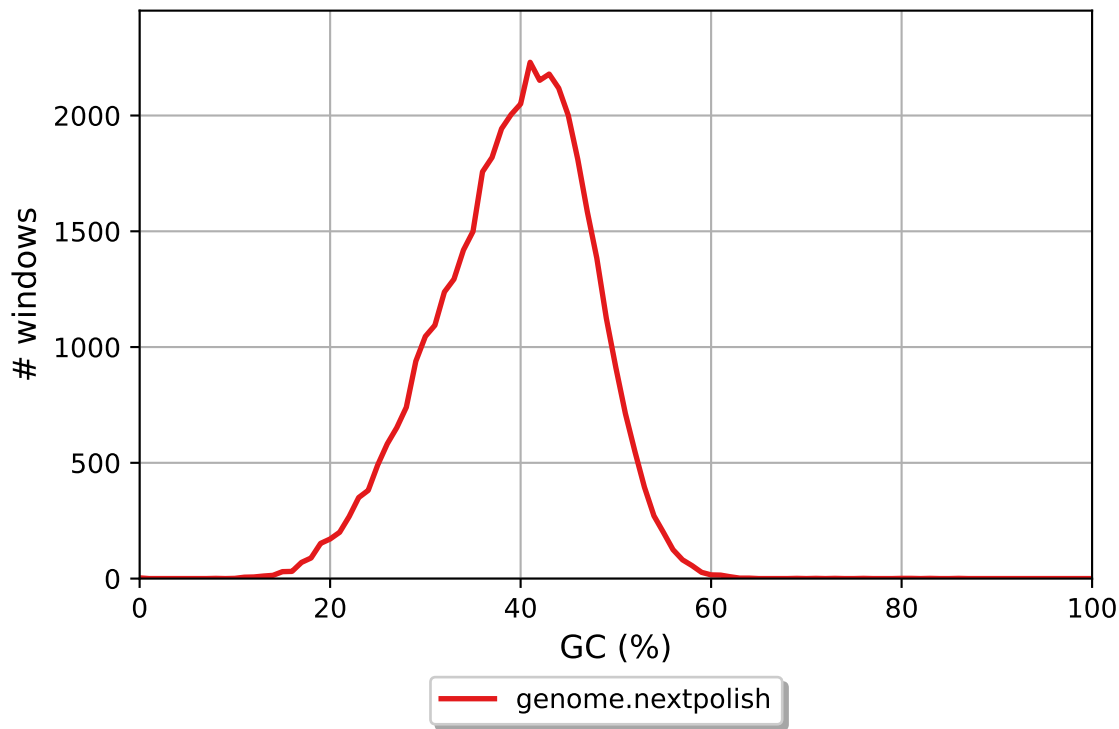
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

