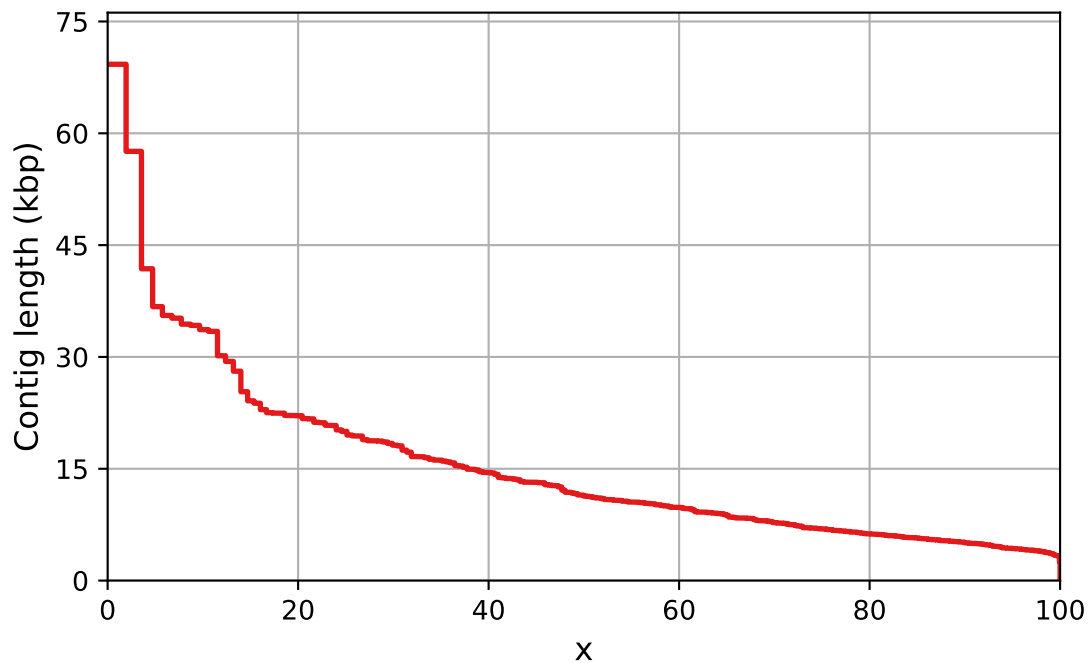


# Report

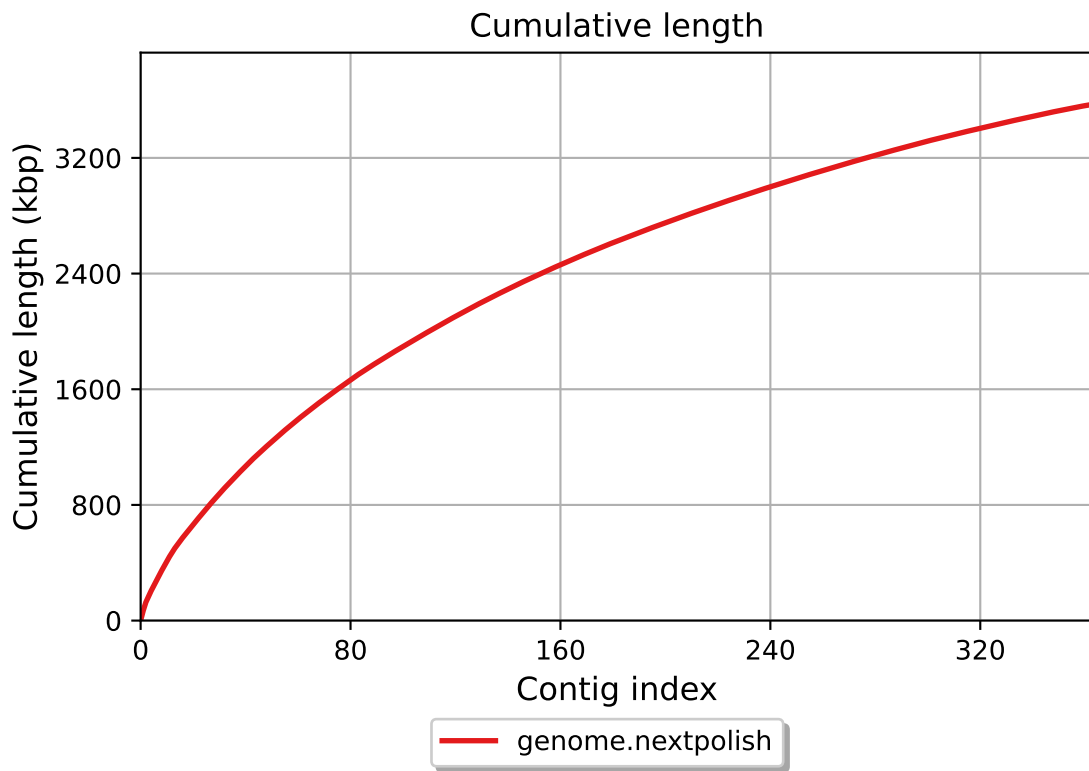
	genome.nextpolish
# contigs (>= 0 bp)	363
# contigs (>= 1000 bp)	363
# contigs (>= 5000 bp)	284
# contigs (>= 10000 bp)	120
# contigs (>= 25000 bp)	14
# contigs (>= 50000 bp)	2
Total length (>= 0 bp)	3570910
Total length (>= 1000 bp)	3570910
Total length (>= 5000 bp)	3238229
Total length (>= 10000 bp)	2103873
Total length (>= 25000 bp)	524890
Total length (>= 50000 bp)	126840
# contigs	363
Largest contig	69260
Total length	3570910
GC (%)	39.84
N50	11338
N90	5068
auN	15781.4
L50	91
L90	280
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	7 + 3 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).

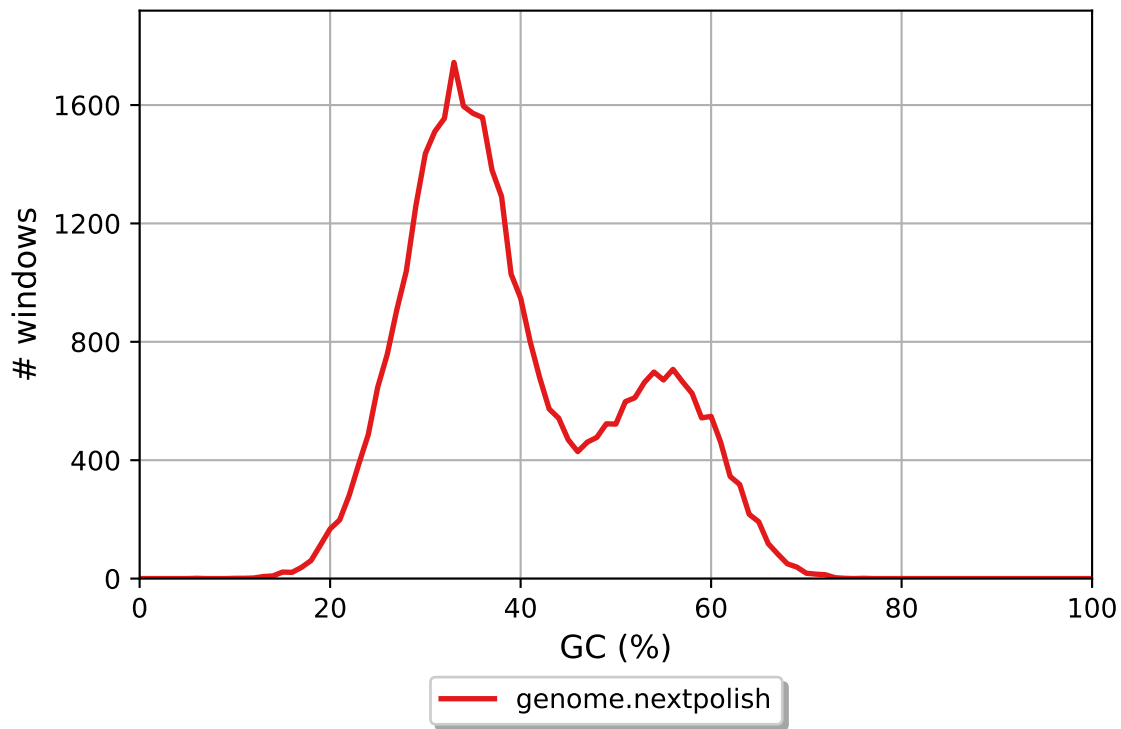
Nx



— genome.nextpolish



GC content



genome.nextpolish GC content

