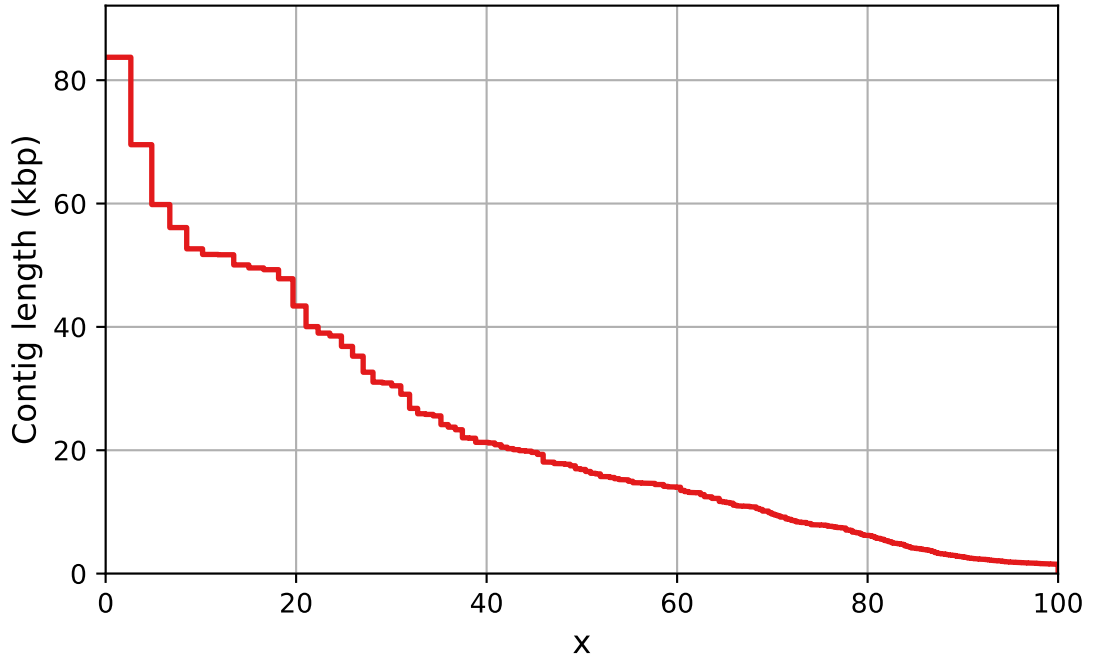


# Report

|                            | MEGAHIT-MetaBAT2-3101140.3.fa |
|----------------------------|-------------------------------|
| # contigs (>= 0 bp)        | 382                           |
| # contigs (>= 1000 bp)     | 382                           |
| # contigs (>= 5000 bp)     | 153                           |
| # contigs (>= 10000 bp)    | 96                            |
| # contigs (>= 25000 bp)    | 26                            |
| # contigs (>= 50000 bp)    | 8                             |
| Total length (>= 0 bp)     | 3162192                       |
| Total length (>= 1000 bp)  | 3162192                       |
| Total length (>= 5000 bp)  | 2614554                       |
| Total length (>= 10000 bp) | 2201042                       |
| Total length (>= 25000 bp) | 1113228                       |
| Total length (>= 50000 bp) | 475348                        |
| # contigs                  | 382                           |
| Largest contig             | 83717                         |
| Total length               | 3162192                       |
| GC (%)                     | 42.42                         |
| N50                        | 16879                         |
| N75                        | 7885                          |
| L50                        | 50                            |
| L75                        | 116                           |
| # N's per 100 kbp          | 0.00                          |
| # predicted rRNA genes     | 0 + 0 part                    |

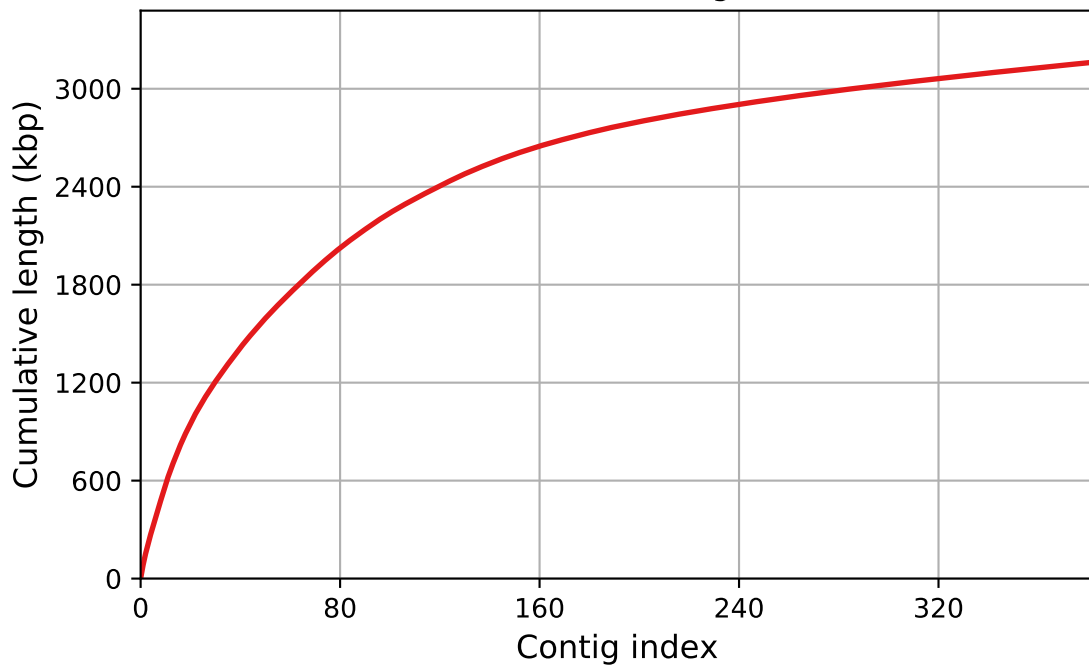
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



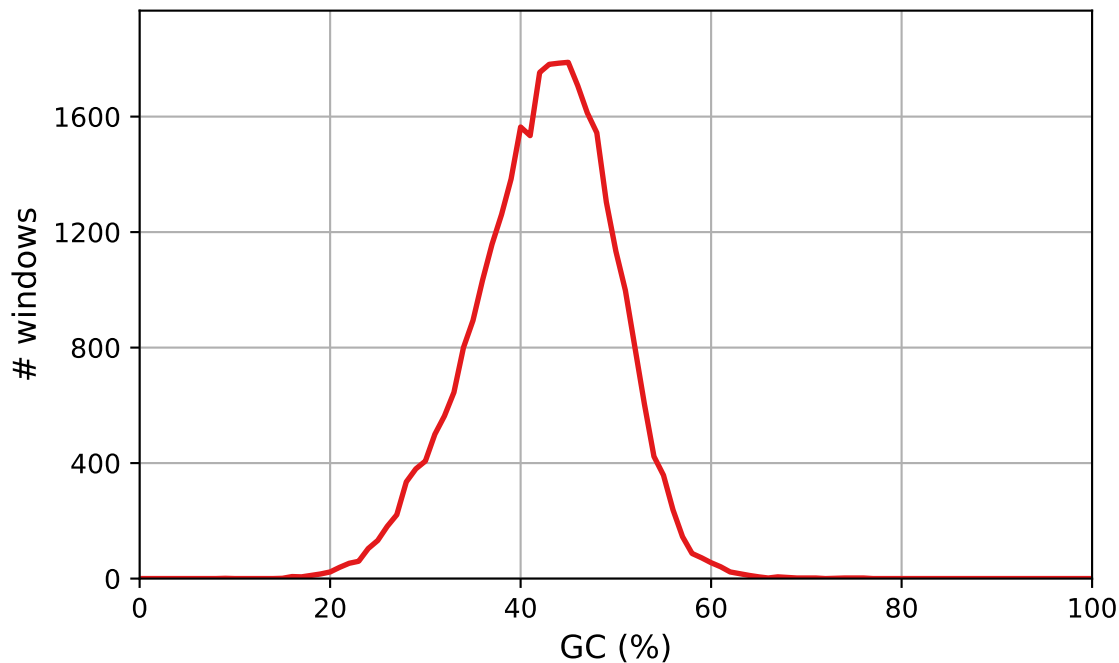
— MEGAHIT-MetaBAT2-3101140.3.fa

Cumulative length



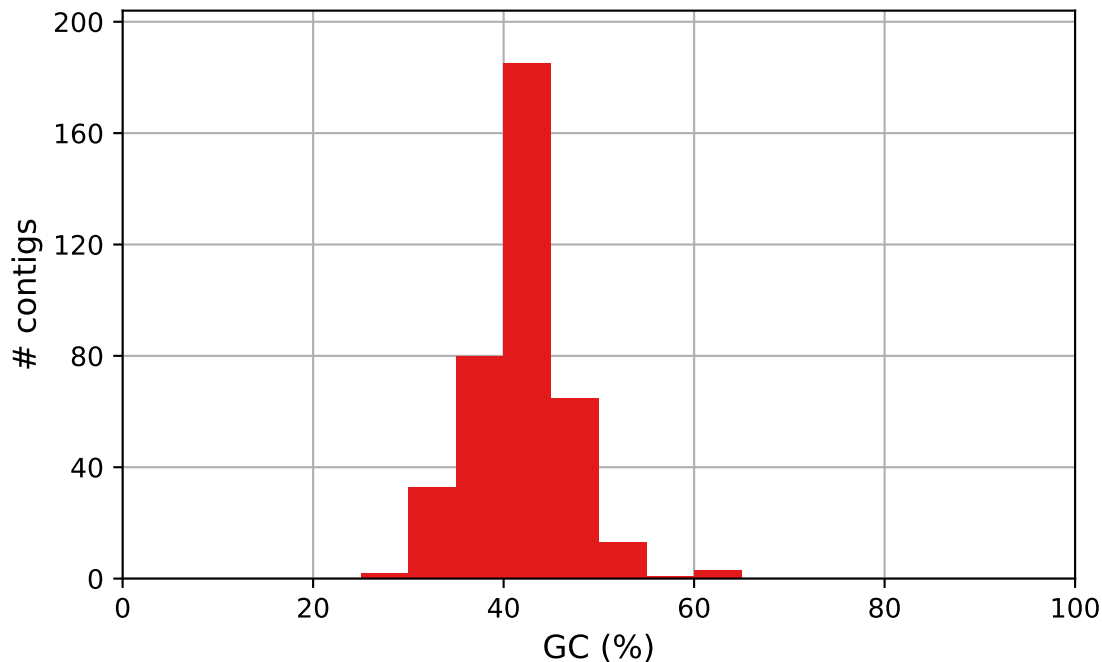
— MEGAHIT-MetaBAT2-3101140.3.fa

## GC content



— MEGAHit-MetaBAT2-3101140.3.fa

MEGAHIT-MetaBAT2-3101140.3.fa GC content



MEGAHit-MetaBAT2-3101140.3.fa