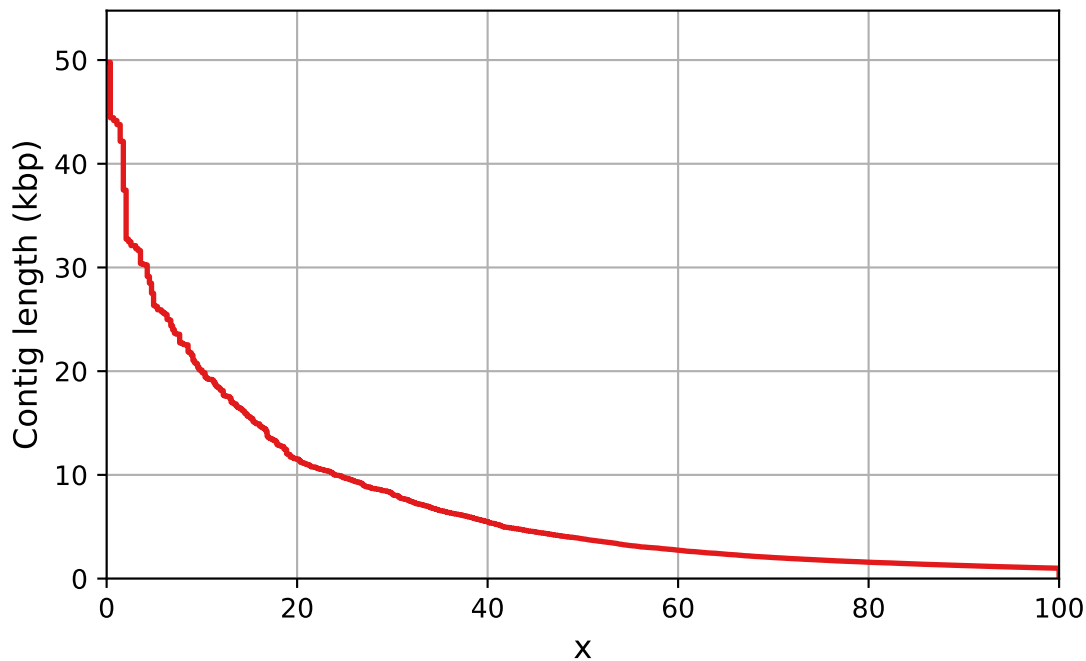


Report

| | MEGAHIT-MaxBin2-3108596.006.fa |
|----------------------------|--------------------------------|
| # contigs (>= 0 bp) | 4462 |
| # contigs (>= 1000 bp) | 4462 |
| # contigs (>= 5000 bp) | 497 |
| # contigs (>= 10000 bp) | 177 |
| # contigs (>= 25000 bp) | 25 |
| # contigs (>= 50000 bp) | 0 |
| Total length (>= 0 bp) | 12794923 |
| Total length (>= 1000 bp) | 12794923 |
| Total length (>= 5000 bp) | 5329929 |
| Total length (>= 10000 bp) | 3062183 |
| Total length (>= 25000 bp) | 811896 |
| Total length (>= 50000 bp) | 0 |
| # contigs | 4462 |
| Largest contig | 49786 |
| Total length | 12794923 |
| GC (%) | 44.13 |
| N50 | 3832 |
| N75 | 1787 |
| L50 | 742 |
| L75 | 2024 |
| # N's per 100 kbp | 0.00 |
| # predicted rRNA genes | 3 + 3 part |

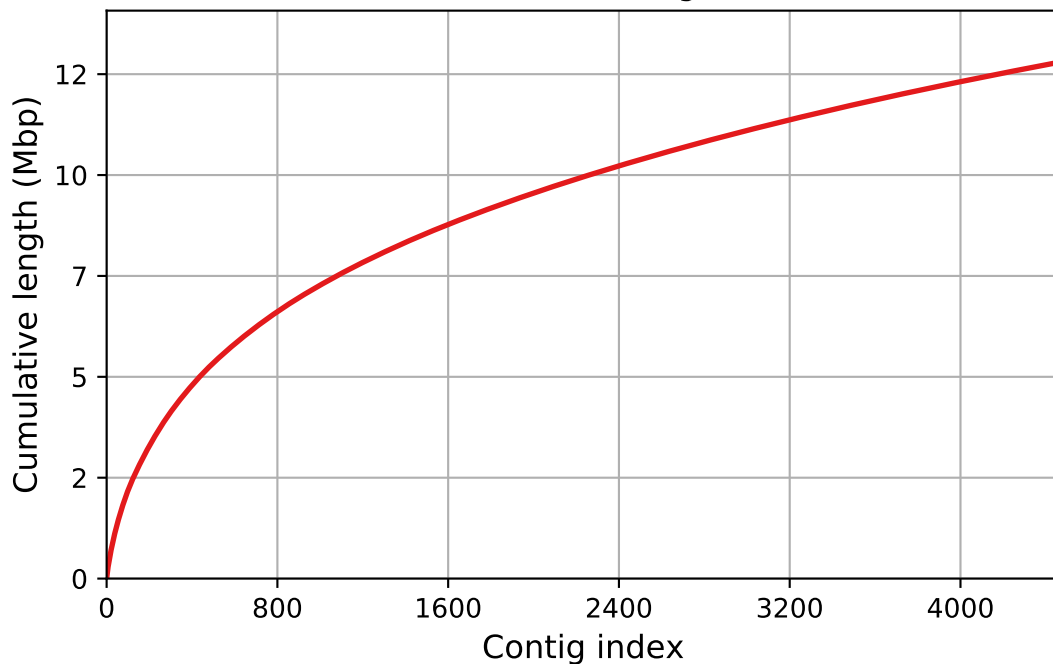
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Nx



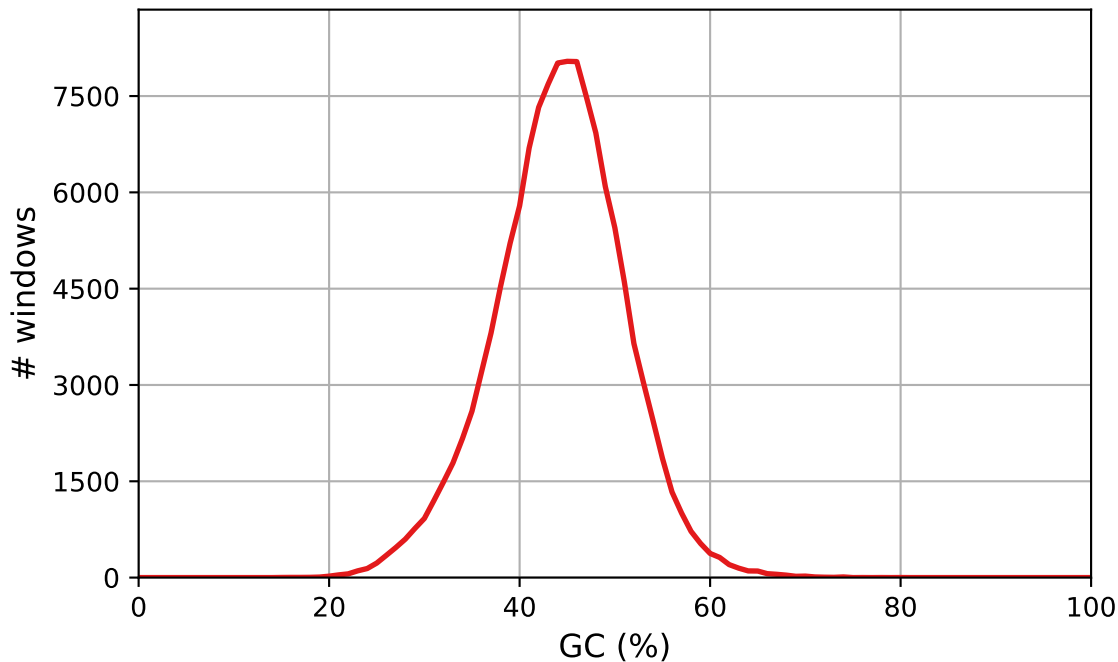
MEGAHIT-MaxBin2-3108596.006.fa

Cumulative length



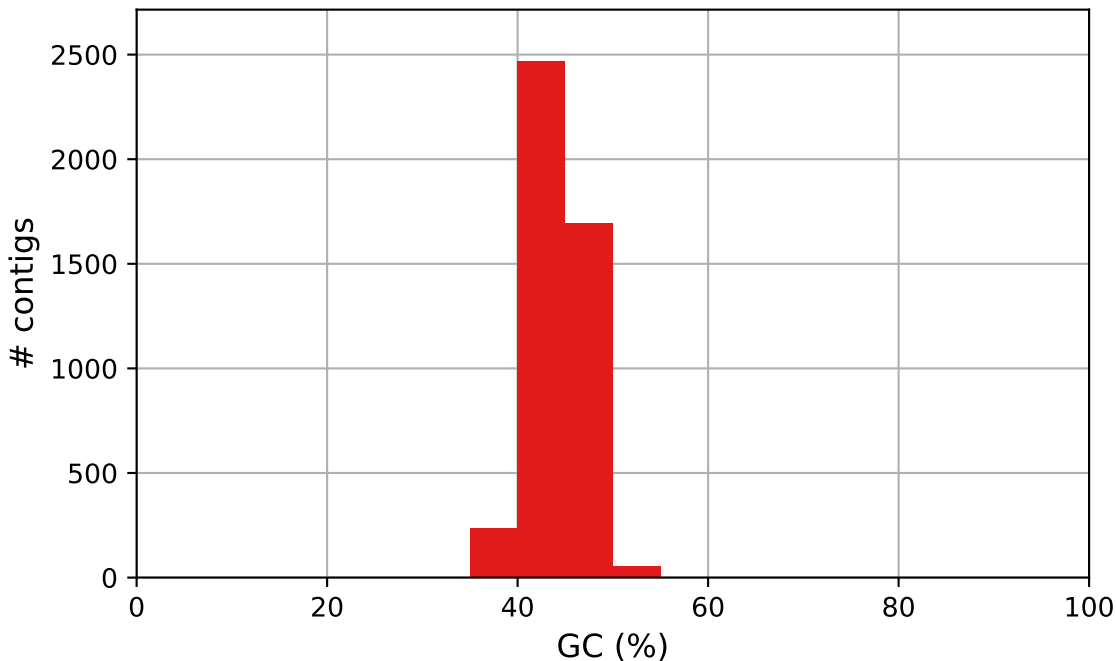
— MEGAHIT-MaxBin2-3108596.006.fa

GC content



MEGAHIT-MaxBin2-3108596.006.fa

MEGAHIT-MaxBin2-3108596.006.fa GC content



MEGAHit-MaxBin2-3108596.006.fa