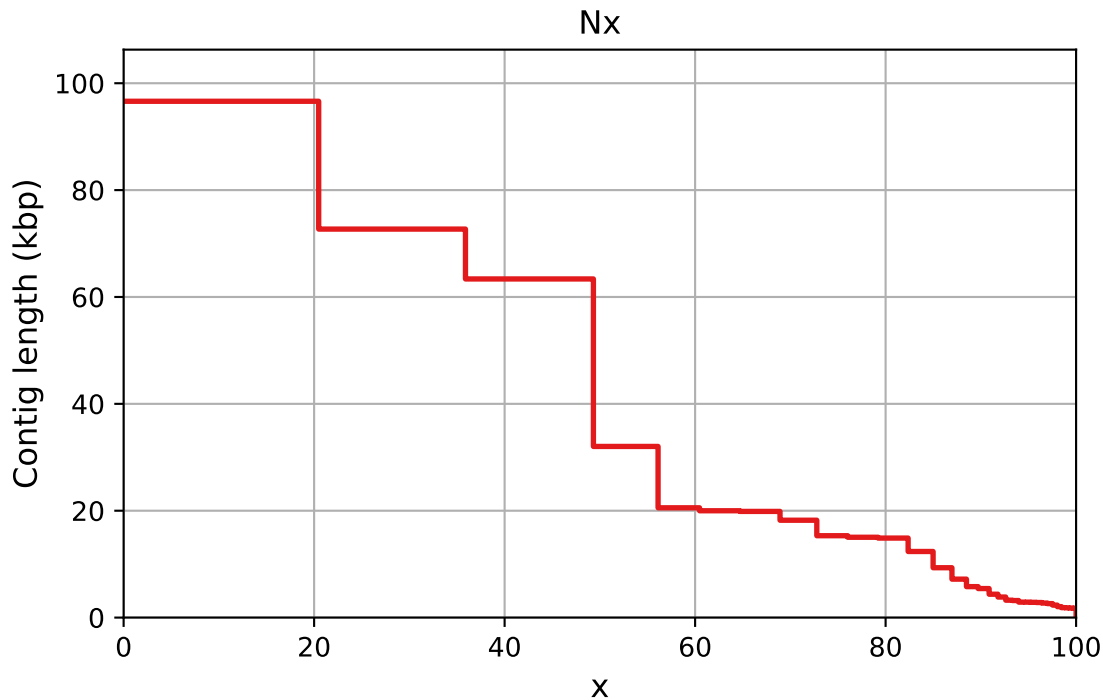


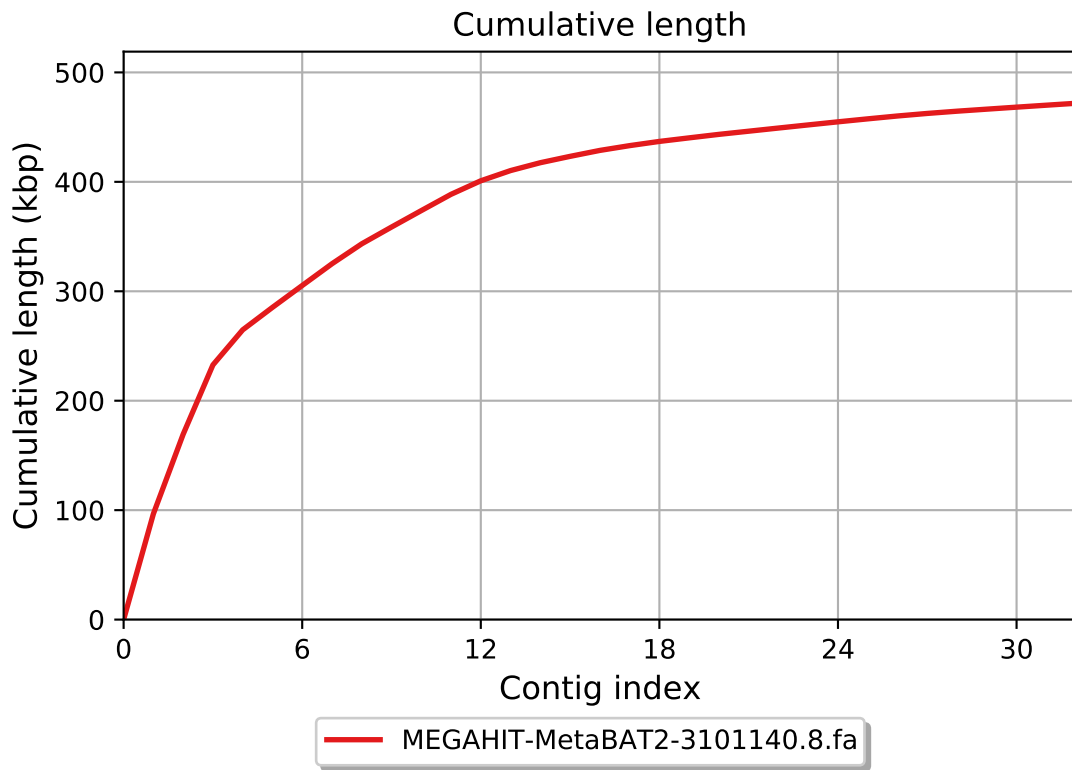
Report

| | MEGAHIT-MetaBAT2-3101140.8.fa |
|----------------------------|-------------------------------|
| # contigs (>= 0 bp) | 32 |
| # contigs (>= 1000 bp) | 32 |
| # contigs (>= 5000 bp) | 16 |
| # contigs (>= 10000 bp) | 12 |
| # contigs (>= 25000 bp) | 4 |
| # contigs (>= 50000 bp) | 3 |
| Total length (>= 0 bp) | 471795 |
| Total length (>= 1000 bp) | 471795 |
| Total length (>= 5000 bp) | 428715 |
| Total length (>= 10000 bp) | 400963 |
| Total length (>= 25000 bp) | 264730 |
| Total length (>= 50000 bp) | 232701 |
| # contigs | 32 |
| Largest contig | 96624 |
| Total length | 471795 |
| GC (%) | 39.86 |
| N50 | 32029 |
| N75 | 15321 |
| L50 | 4 |
| L75 | 9 |
| # N's per 100 kbp | 0.00 |
| # predicted rRNA genes | 0 + 0 part |

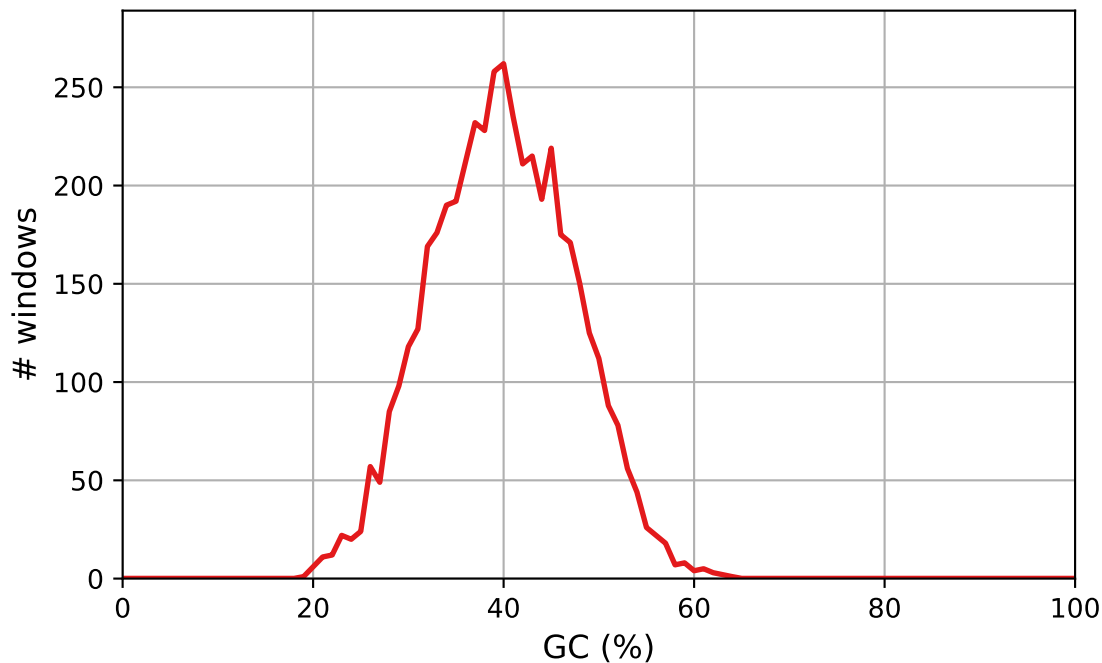
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— MEGAHIT-MetaBAT2-3101140.8.fa

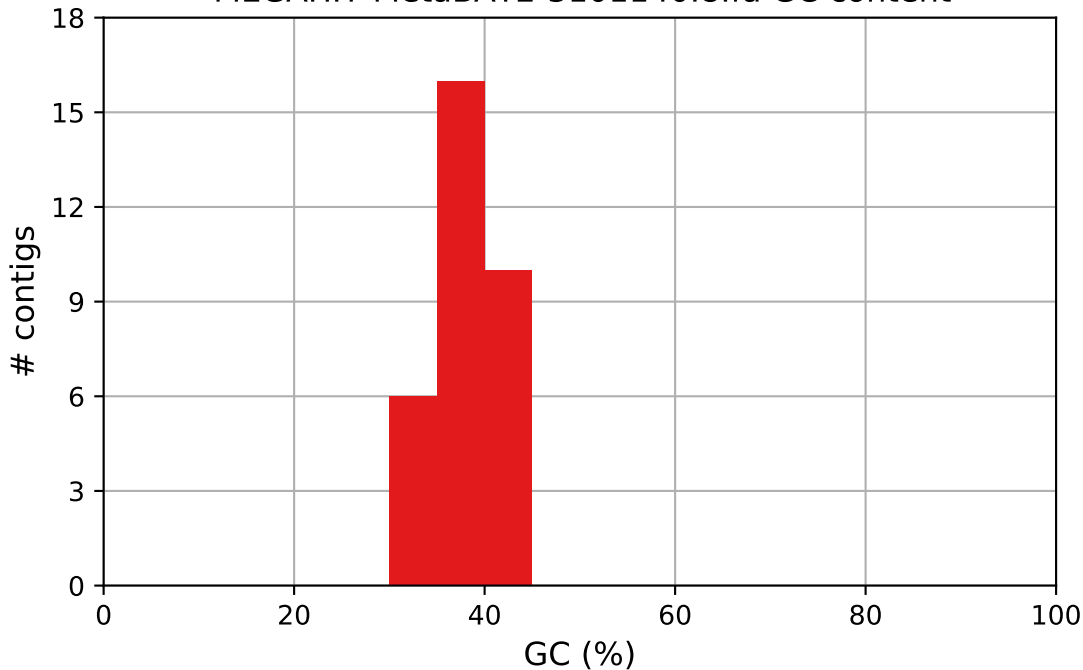


GC content



— MEGAHIT-MetaBAT2-3101140.8.fa

MEGAHIT-MetaBAT2-3101140.8.fa GC content



MEGAHit-MetaBAT2-3101140.8.fa