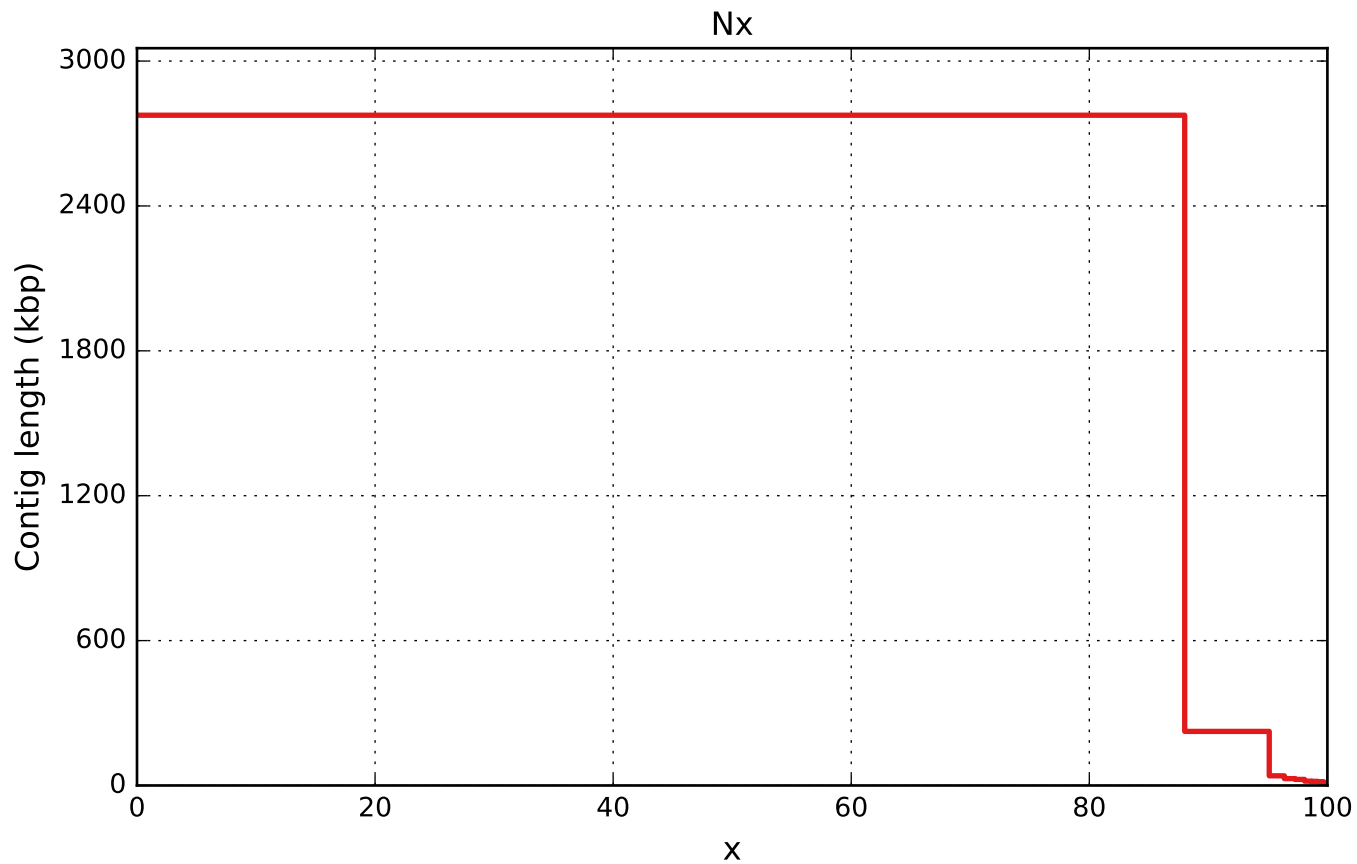


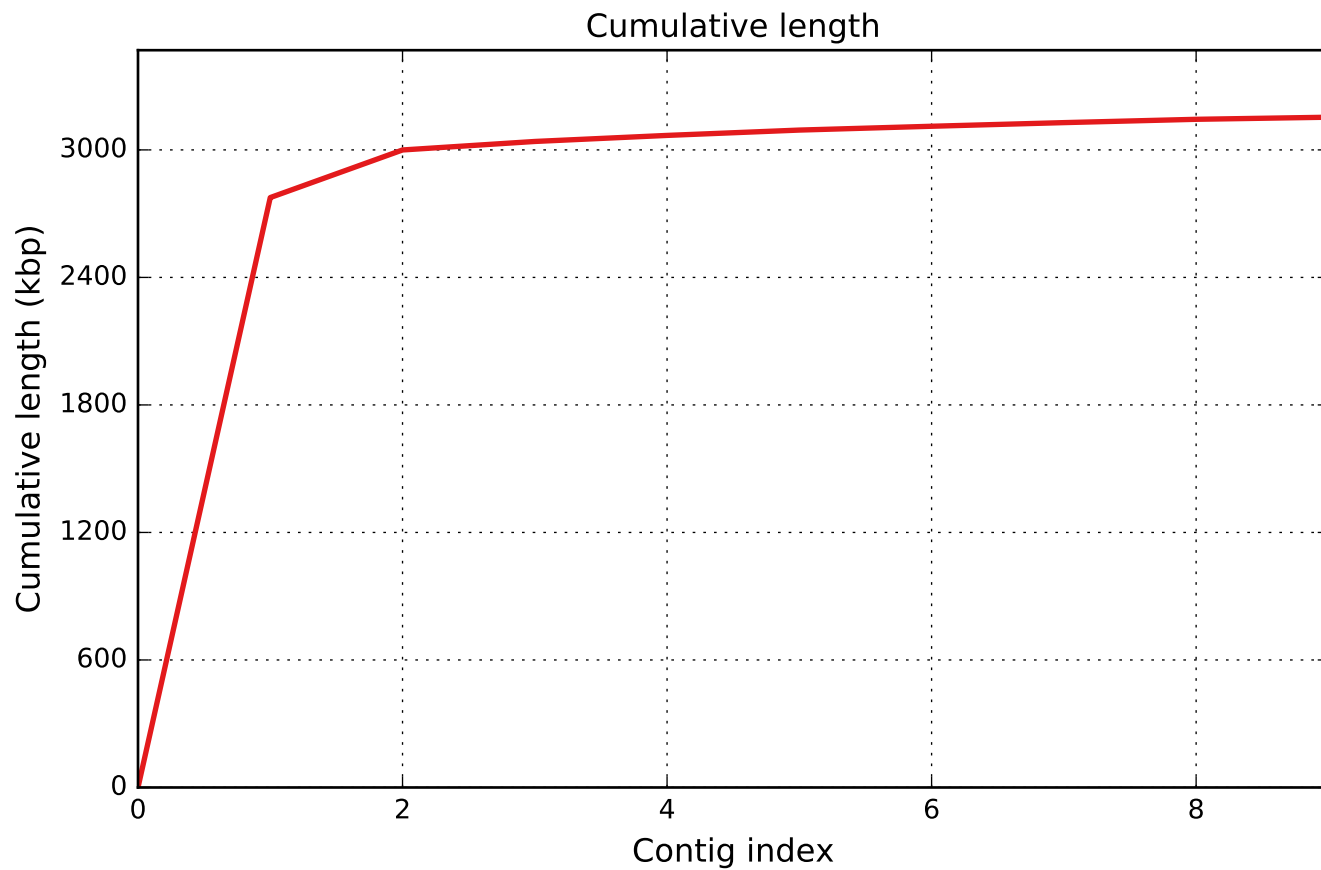
# Report

|                            | pilon_combined_genome |
|----------------------------|-----------------------|
| # contigs (>= 0 bp)        | 9                     |
| # contigs (>= 1000 bp)     | 9                     |
| # contigs (>= 5000 bp)     | 9                     |
| # contigs (>= 10000 bp)    | 8                     |
| # contigs (>= 25000 bp)    | 5                     |
| # contigs (>= 50000 bp)    | 2                     |
| Total length (>= 0 bp)     | 3153758               |
| Total length (>= 1000 bp)  | 3153758               |
| Total length (>= 5000 bp)  | 3153758               |
| Total length (>= 10000 bp) | 3143962               |
| Total length (>= 25000 bp) | 3093281               |
| Total length (>= 50000 bp) | 2999691               |
| # contigs                  | 9                     |
| Largest contig             | 2775723               |
| Total length               | 3153758               |
| GC (%)                     | 37.77                 |
| N50                        | 2775723               |
| N75                        | 2775723               |
| L50                        | 1                     |
| L75                        | 1                     |
| # N's per 100 kbp          | 0.00                  |

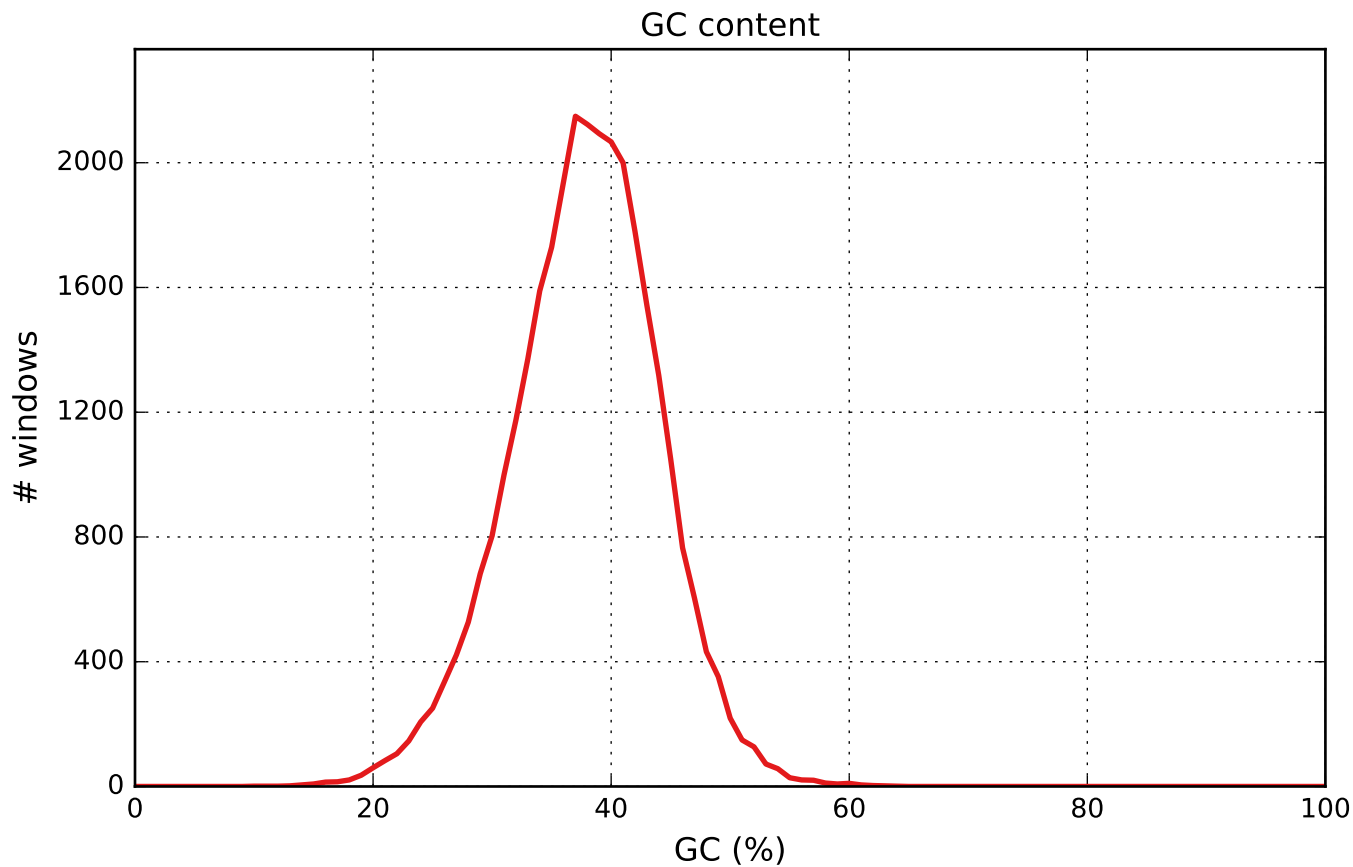
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— pilon\_combined\_genome

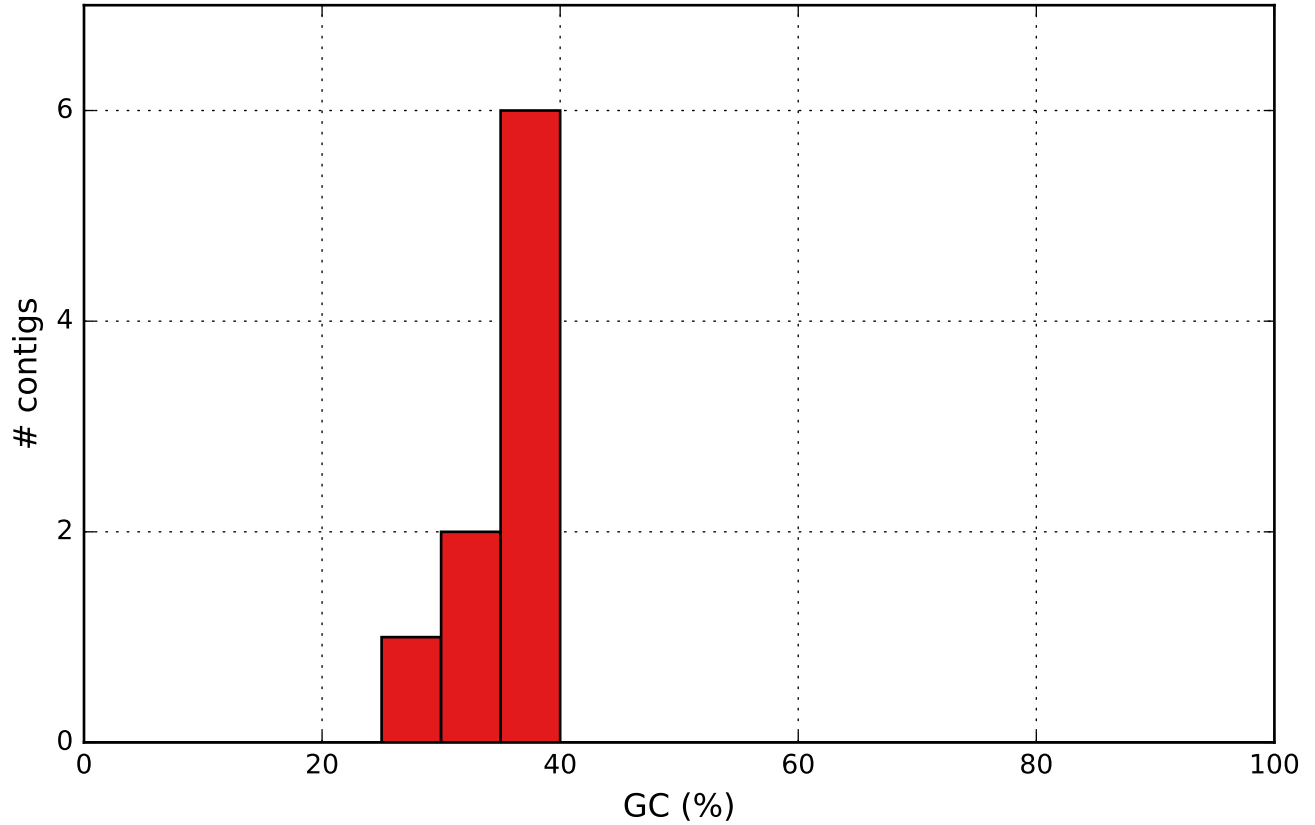



— pilon\_combined\_genome



— pilon\_combined\_genome

pilon\_combined\_genome GC content



 pilon\_combined\_genome