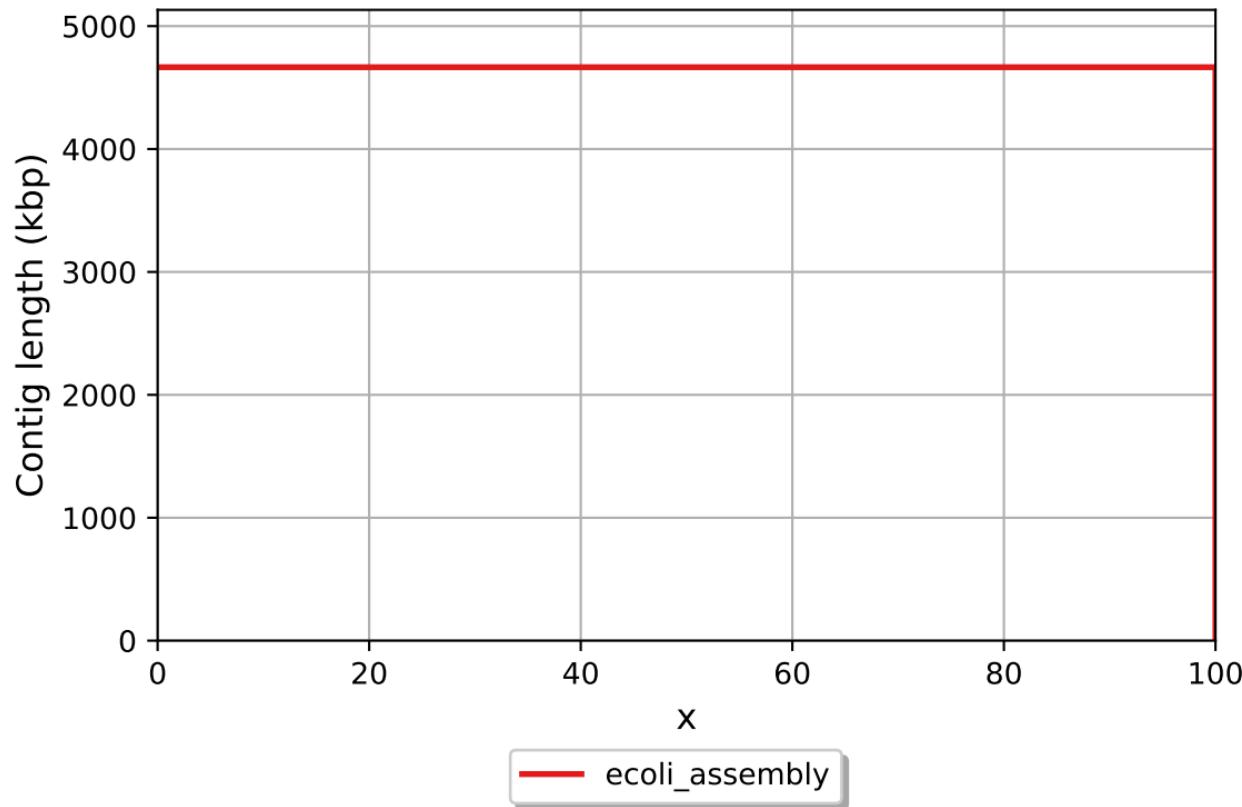


# Report

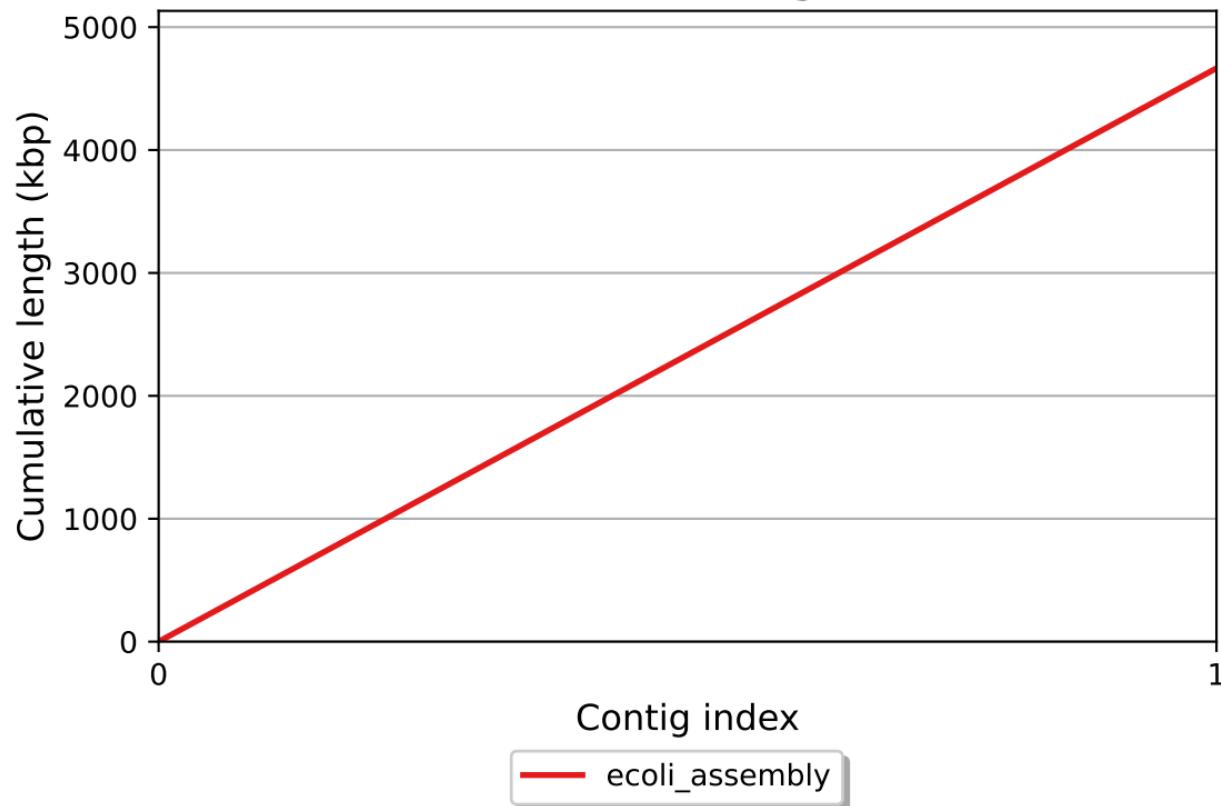
|                            | ecoli_assembly |
|----------------------------|----------------|
| # contigs (>= 0 bp)        | 1              |
| # contigs (>= 1000 bp)     | 1              |
| # contigs (>= 5000 bp)     | 1              |
| # contigs (>= 10000 bp)    | 1              |
| # contigs (>= 25000 bp)    | 1              |
| # contigs (>= 50000 bp)    | 1              |
| Total length (>= 0 bp)     | 4665559        |
| Total length (>= 1000 bp)  | 4665559        |
| Total length (>= 5000 bp)  | 4665559        |
| Total length (>= 10000 bp) | 4665559        |
| Total length (>= 25000 bp) | 4665559        |
| Total length (>= 50000 bp) | 4665559        |
| # contigs                  | 1              |
| Largest contig             | 4665559        |
| Total length               | 4665559        |
| GC (%)                     | 50.77          |
| N50                        | 4665559        |
| N90                        | 4665559        |
| auN                        | 4665559.0      |
| L50                        | 1              |
| L90                        | 1              |
| # N's per 100 kbp          | 0.00           |

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

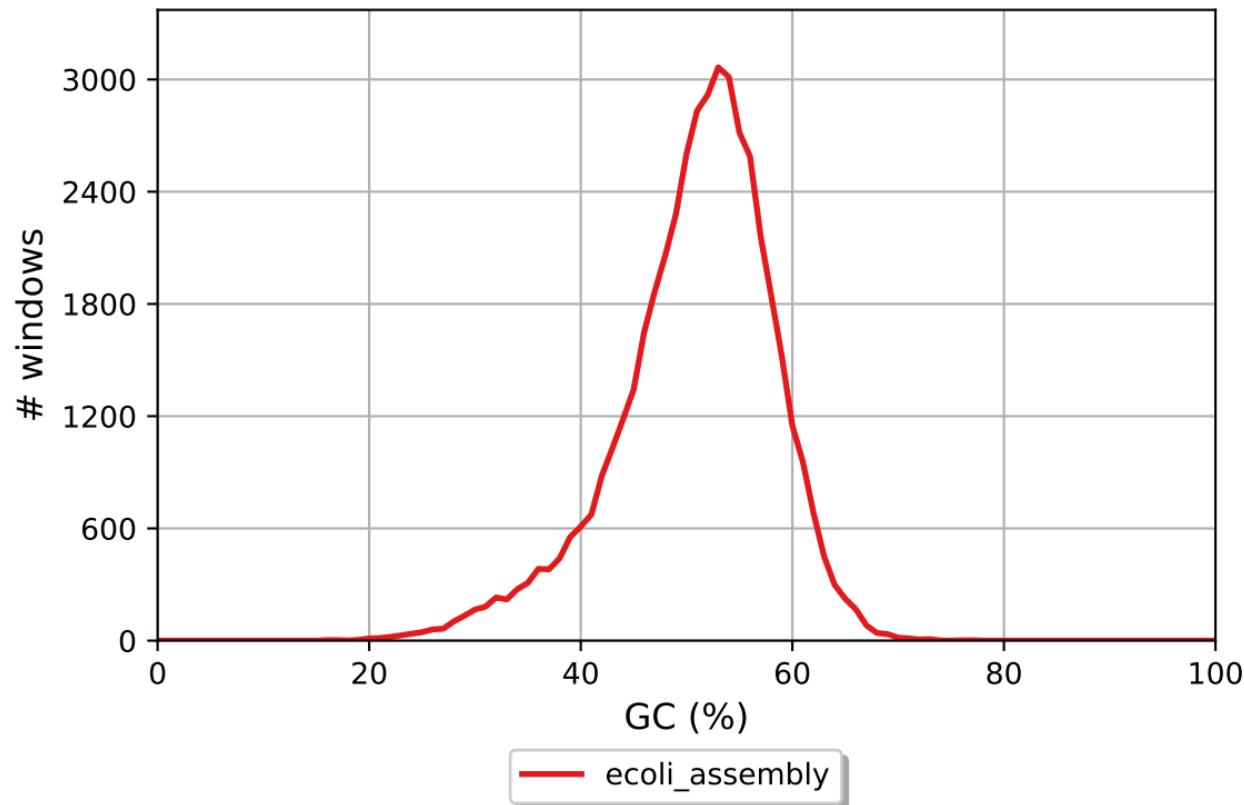
Nx



Cumulative length



## GC content



*ecoli\_assembly* GC content

