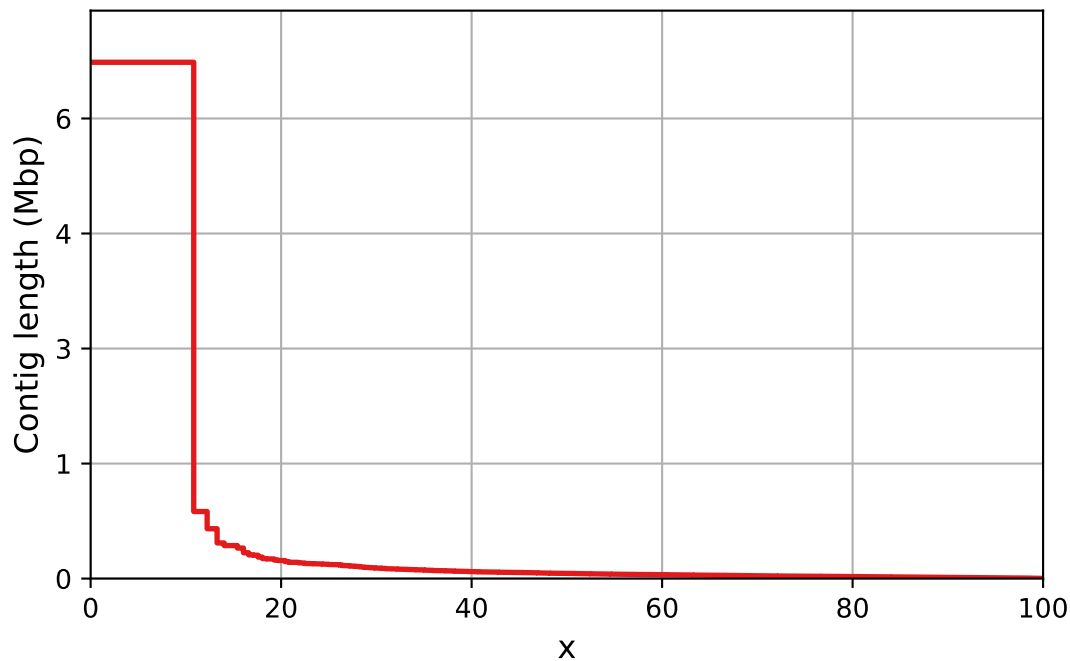


# Report

|                            | genome.nextpolish |
|----------------------------|-------------------|
| # contigs (>= 0 bp)        | 1679              |
| # contigs (>= 1000 bp)     | 1674              |
| # contigs (>= 5000 bp)     | 1530              |
| # contigs (>= 10000 bp)    | 1197              |
| # contigs (>= 25000 bp)    | 661               |
| # contigs (>= 50000 bp)    | 301               |
| Total length (>= 0 bp)     | 62186396          |
| Total length (>= 1000 bp)  | 62182713          |
| Total length (>= 5000 bp)  | 61678583          |
| Total length (>= 10000 bp) | 59161295          |
| Total length (>= 25000 bp) | 50309639          |
| Total length (>= 50000 bp) | 37689394          |
| # contigs                  | 1679              |
| Largest contig             | 6733065           |
| Total length               | 62186396          |
| GC (%)                     | 51.80             |
| N50                        | 68422             |
| N90                        | 15523             |
| auN                        | 824926.1          |
| L50                        | 187               |
| L90                        | 945               |
| # N's per 100 kbp          | 0.00              |
| Complete BUSCO (%)         | 0.00              |
| Partial BUSCO (%)          | 0.00              |
| # predicted rRNA genes     | 97 + 5 part       |

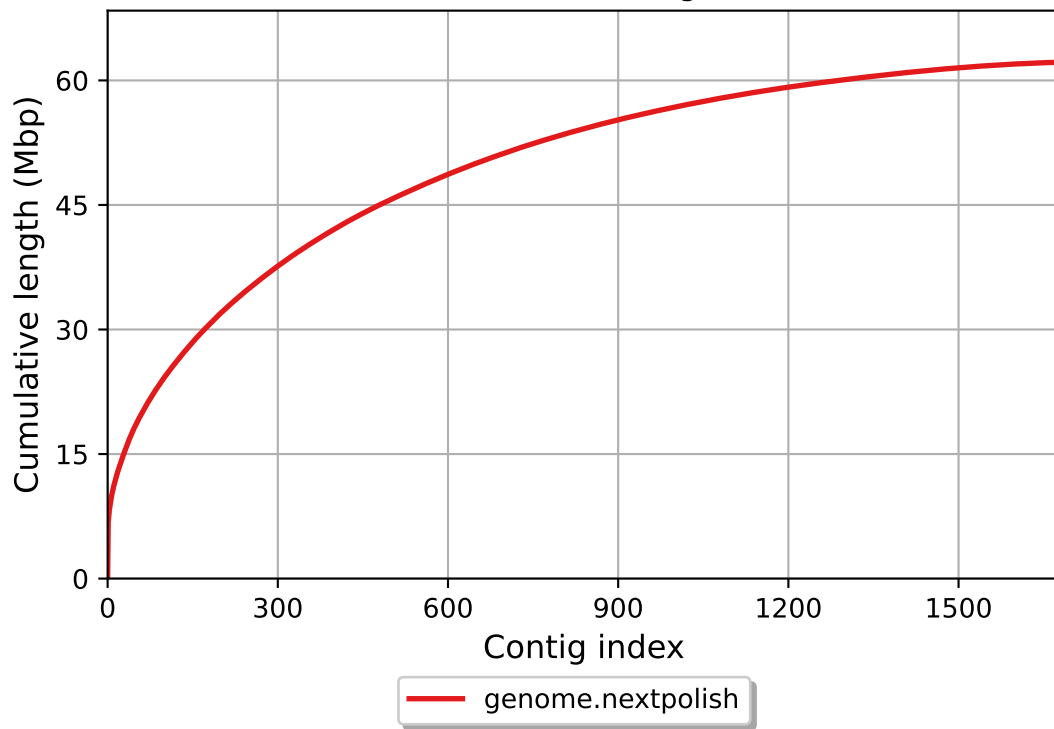
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

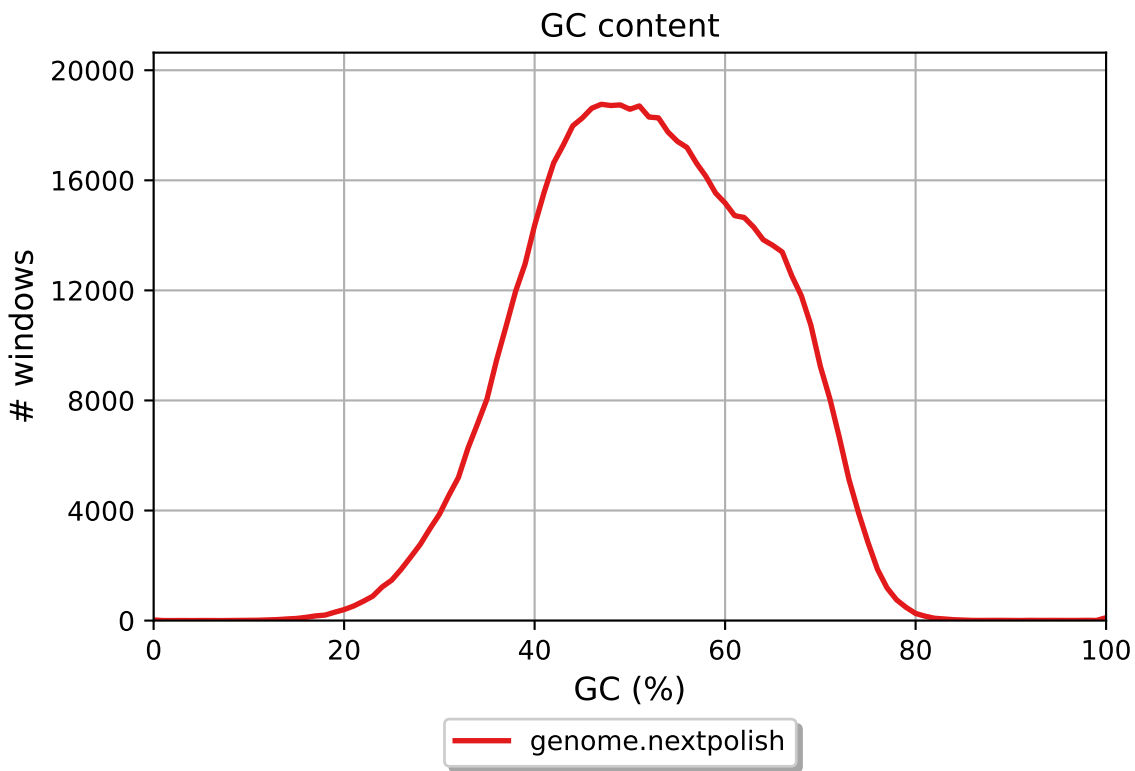
Nx



— genome.nextpolish

Cumulative length





genome.nextpolish GC content

