

DBM susceptible (Fuzhou-S) strain

| Data set   |                   | Total No. | GC %   | Size   | N50 (bp) | Max (bp)  |
|------------|-------------------|-----------|--------|--------|----------|-----------|
| Version 1  | Scaffolds (>1 kp) | 1,819     | 38.34% | 394 Mb | 737,182  | 3,493,687 |
|            | Contigs (>1kbp)   | 15,632    | 38.34% | 387 Mb | 49,352   | 317,222   |
| Version 2* | Contigs (>1kbp)   | 31,979    | 38.34% | 334 Mb | 18,785   | 202,508   |

\* Version 2 of the DBM genome assembly (Fuzhou-S) was generated by masking with “n” in version 1 according to Poisson distribution-based K-mer statistics (13).