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| Table S1. Genome Assembly |
| Genome Assembly size (Mbp} | 30.14 |
| Sequencing read coverage depth | 144.9 |
| # of contigs | 1195 |
| # of scaffolds | 573 |
| # of scaffolds >= 2Kbp | 506 |
| Scaffold N50 | 72 |
| Scaffold L50 (Mbp) | 0.12 |
| # of gaps | 622 |
| % of scaffold length in gaps | 1.5% |
| Three largest Scaffolds (Mbp) | 0.61, 0.54, 0.46 |