Correction to: Identification and molecular characterization of tea-oil camellia-associated totivirus 1

Z.-Y. Zhang¹ · H. Huang¹ · X. X. Han · R. Li² · L.-P. Wu¹ · L. Wu¹

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Authors would like to correct the error in their publication.

The accession number [NC 2384770] is invalid. The correct version is [MW_025466]. The modified phylogenetic tree (Fig. 2) is also updated here.

The original article has been corrected.

The original article can be found online at <https://doi.org/10.1007/s00705-021-05058-2>.

✉ L.-P. Wu
lpwu000@ncu.edu.cn

¹ Key Laboratory of Poyang Lake Environment and Resource, School of Life Science, Ministry of Education, Nanchang University, Nanchang 330031, Jiangxi, China

² National Germplasm Resources Laboratory, USDA-ARS, Beltsville, MD 20705, USA

Fig. 2 Phylogenetic trees generated by sequences of tea-oil camellia associated totivirus 1 (TOCaTV 1) and selected members of the family *Totiviridae* by maximum likelihood method. Two representative members of the family *Chrysoviridae* were included as an outgroup. Sequences used for phylogenetic analyses were the whole genomic sequences (a), the amino acid sequences of CP (b) and RdRp (c). The solid dot indicates the virus characterized in this study. Bootstrap values from 1000 replicates are given at each node

Fig. 2 (continued)

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