



Addendum

The Complete Genome Sequence of Cucumopine-Type *Agrobacterium rhizogenes* Strain K599 (NCPB2659), A Nature's Genetic Engineer Inducing Hairy Roots

Xiaxiu Tong¹, Yafei Li¹, Taihe Xiang^{1,2*}, Zhehao Chen¹, Lilin Wang¹, Chengchao Zhang¹, Mi Feng¹ and Laiyang Wu¹

¹College of Life and Environment Sciences, Hangzhou Normal University, Hangzhou 310036, China

²Zhejiang Provincial Key Laboratory for Genetic Improvement and Quality Control of Medicinal Plants, Hangzhou 310036, China

*For correspondence: xthen@163.com; xthen@hznu.edu.cn

Addendum in the Acknowledgement Section of Int. J. Agric. Biol., 20: 1167–1174

The K599 genome was sequenced and assembled once again by us. The complete genome sequence of K599 was corrected and updated in GenBank (accession numbers versions: CP019701.2, CP019702.2 and CP019703.3). The K599 genome has 5,482,241 bp and consists of one circular chromosome of 3,003,350 bp, one linear chromosome of 2,276,589 bp, and one Ri plasmid (pRi2659) of 202,302 bp.

Especially, it should be noted that the sequence size of chromosome 1 of our report (GenBank accession number version: CP019701.2) is consistent with that of the draft genome sequence reported by Valder Franco *et al.* (2016). We would like to thank Dr. Roger Thilmony of USDA-Agriculture Research Service, Western Regional Research Center, Crop Improvement and Genetics Research Unit, Albany, California, USA, for discussion on the K599 genome.

Reference

Valdes Franco, J.A., R. Collier, Y. Wang, N. Huo, Y. Gu, R. Thilmony and J.G. Thomson, 2016. Draft genome sequence of *Agrobacterium rhizogenes* strain NCPB2659. *Genome Announc.*, 4: e00746-16