

OPEN ACCESS

APPROVED BY
Frontiers Editorial Office,
Frontiers Media SA, Switzerland

*CORRESPONDENCE
Jianchao Wei

☑ jianchaowei@shvri.ac.cn
Zhiyong Ma
☑ zhiyongma@shvri.ac.cn

[†]These authors have contributed equally to this work

RECEIVED 29 February 2024 ACCEPTED 01 March 2024 PUBLISHED 19 March 2024

CITATION

Hameed M, Wahaab A, Shan T, Wang X, Khan S, Di D, Xiqian L, Zhang J-J, Anwar MN, Nawaz M, Li B, Liu K, Shao D, Qiu Y, Wei J and Ma Z (2024) Corrigendum: A metagenomic analysis of mosquito virome collected from different animal farms at Yunnan–Myanmar border of China. *Front. Microbiol.* 15:1393495. doi: 10.3389/fmicb.2024.1393495

COPYRIGHT

© 2024 Hameed, Wahaab, Shan, Wang, Khan, Di, Xiqian, Zhang, Anwar, Nawaz, Li, Liu, Shao, Qiu, Wei and Ma. This is an open-access article distributed under the terms of the Creative Commons Attribution License (CC BY). The use, distribution or reproduction in other forums is permitted, provided the original author(s) and the copyright owner(s) are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.

Corrigendum: A metagenomic analysis of mosquito virome collected from different animal farms at Yunnan—Myanmar border of China

Muddassar Hameed[†], Abdul Wahaab[†], Tongling Shan[†], Xin Wang, Sawar Khan, Di Di, Liu Xiqian, Jun-Jie Zhang, Muhammad Naveed Anwar, Mohsin Nawaz, Beibei Li, Ke Liu, Donghua Shao, Yafeng Qiu, Jianchao Wei* and Zhiyong Ma*

Shanghai Veterinary Research Institute, Chinese Academy of Agricultural Sciences, Shanghai, China

KEYWORDS

mosquito, mosquito virome, metagenomics, viral community, animal farm

A corrigendum on

A metagenomic analysis of mosquito virome collected from different animal farms at Yunnan-Myanmar border of China

by Hameed, M., Wahaab, A., Shan, T., Wang, X., Khan, S., Di, D., Xiqian, L., Zhang, J.-J., Anwar, M. N., Nawaz, M., Li, B., Liu, K., Shao, D., Qiu, Y., Wei, J., and Ma, Z. (2021). *Front. Microbiol.* 11:591478. doi: 10.3389/fmicb.2020.591478

In the published article, there was an error in Figure 5 as published. When preparing Figure 5A, since JEV-C1 and JEV-G1 are on two PPT pages, when copying JEV-G1, in order to keep the zoom ratio of the two pages, JEV-C1 is directly copied to the second page as a contrast, so there is confusion when inserting the JEV-G1 DAPI picture. The corrected Figure 5 and its caption appear below.

The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

Publisher's note

All claims expressed in this article are solely those of the authors and do not necessarily represent those of their affiliated organizations, or those of the publisher, the editors and the reviewers. Any product that may be evaluated in this article, or claim that may be made by its manufacturer, is not guaranteed or endorsed by the publisher.

Hameed et al. 10.3389/fmicb.2024.1393495

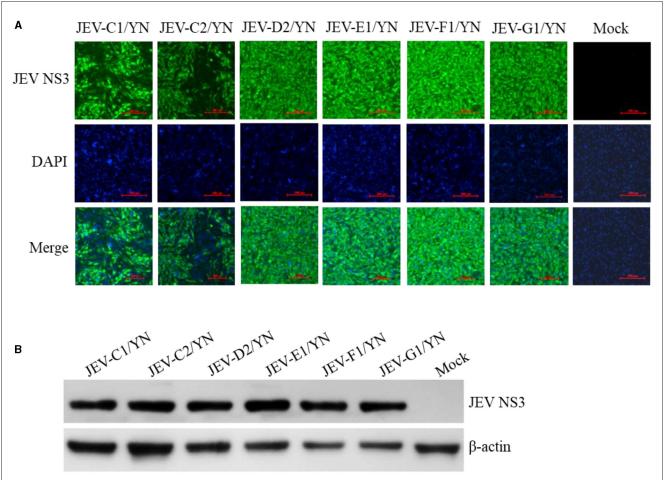


FIGURE 5

Detection of JEV in BHK-21 cells. BHK-21 cells were inoculated with the supernatants harvested from the CPE-positive BHK-21 cells inoculated with mosquito samples and incubated for 36 h. The presence of JEV was detected by IFA (A) and Western blot (B) with antibodies specific to JEV NS3 protein. Nuclei were stained with DAPI.