

CORRECTION

Open Access



# Correction to: Complete genome analysis of a virulent *Vibrio scophthalmi* strain VSc190401 isolated from diseased marine fish half-smooth tongue sole, *Cynoglossus semilaevis*

Zheng Zhang<sup>1,2\*</sup>, Yong-xiang Yu<sup>1</sup>, Yin-geng Wang<sup>1,2\*</sup>, Xiao Liu<sup>1</sup>, Li-fang Wang<sup>1</sup>, Hao Zhang<sup>1</sup>, Mei-jie Liao<sup>1,2</sup> and Bin Li<sup>1</sup>

**Correction to: BMC Microbiol. 20, 341 (2020)**  
<https://doi.org/10.1186/s12866-020-02028-7>

Following publication of the original article [1], we were notified that Prof. Yin-geng Wang needs to be marked as the second corresponding author.

The original article has been corrected.

#### Author details

<sup>1</sup>Key Laboratory of Maricultural Organism Disease Control, Ministry of Agriculture and Rural Affairs, Yellow Sea Fisheries Research Institute, Chinese Academy of Fishery Sciences, Qingdao, Shandong 266071, PR China.

<sup>2</sup>Laboratory for Marine Fisheries Science and Food Production Processes, Pilot National Laboratory for Marine Science and Technology (Qingdao), Qingdao, Shandong 266237, PR China.

Published online: 14 December 2020

#### Reference

1. Zhang Z, et al. Complete genome analysis of a virulent *Vibrio scophthalmi* strain VSc190401 isolated from diseased marine fish half-smooth tongue sole, *Cynoglossus semilaevis*. *BMC Microbiol.* 2020;20:341. <https://doi.org/10.1186/s12866-020-02028-7>.

The original article can be found online at <https://doi.org/10.1186/s12866-020-02028-7>.

\* Correspondence: [zhangzheng@ysfri.ac.cn](mailto:zhangzheng@ysfri.ac.cn); [wangyg@ysfri.ac.cn](mailto:wangyg@ysfri.ac.cn)

<sup>1</sup>Key Laboratory of Maricultural Organism Disease Control, Ministry of Agriculture and Rural Affairs, Yellow Sea Fisheries Research Institute, Chinese Academy of Fishery Sciences, Qingdao, Shandong 266071, PR China

Full list of author information is available at the end of the article



© The Author(s). 2020 **Open Access** This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit <http://creativecommons.org/licenses/by/4.0/>. The Creative Commons Public Domain Dedication waiver (<http://creativecommons.org/publicdomain/zero/1.0/>) applies to the data made available in this article, unless otherwise stated in a credit line to the data.