Check for updates

OPEN ACCESS

APPROVED BY Frontiers Editorial Office, Frontiers Media SA, Switzerland

*CORRESPONDENCE Zichen Yang Zichen_yang@yahoo.com Yiming Zhang Zhangyiming@tmmu.edu.cn

[†]These authors have contributed equally to this work

RECEIVED 24 February 2024 ACCEPTED 26 February 2024 PUBLISHED 08 March 2024

CITATION

Liao D, Zhang J, Liu R, Chen K, Liu Y, Shao Y, Shi X, Zhang Y and Yang Z (2024) Corrigendum: Whole-genome sequencing, annotation, and biological characterization of a novel Siphoviridae phage against multi-drug resistant *Propionibacterium acne. Front. Microbiol.* 15:1390825. doi: 10.3389/fmicb.2024.1390825

COPYRIGHT

© 2024 Liao, Zhang, Liu, Chen, Liu, Shao, Shi, Zhang and Yang. 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: Whole-genome sequencing, annotation, and biological characterization of a novel Siphoviridae phage against multi-drug resistant *Propionibacterium acne*

Danxi Liao¹, Jian Zhang¹, Ruolan Liu¹, Kui Chen², Yuanyuan Liu³, Yuming Shao¹, Xi Shi¹, Yiming Zhang^{1*†} and Zichen Yang^{1,4*†}

¹Department of Plastic and Cosmetic Surgery, Xinqiao Hospital, The Second Affiliated Hospital, Army Medical University (The Third Military Medical University), Chongqing, China, ²Department of Clinical Laboratory, Xinqiao Hospital, The Second Affiliated Hospital, Army Medical University (The Third Military Medical University), Chongqing, China, ³Cadet Brigade 4, College of Basic Medicine, Army Medical University (The Third Military Medical University), Chongqing, China, ⁴Department of Microbiology, College of Basic Medicine, Army Medical University (The Third Military Medical University), Chongqing, China

KEYWORDS

Propionibacterium acne, bacteriophage, severe acne vulgaris, antibiotics-resistant, phage therapy

A corrigendum on

Whole-genome sequencing, annotation, and biological characterization of a novel Siphoviridae phage against multi-drug resistant Propionibacterium acne

by Liao, D., Zhang, J., Liu, R., Chen, K., Liu, Y., Shao, Y., Shi, X., Zhang, Y., and Yang, Z. (2023). *Front. Microbiol.* 13:1065386. doi: 10.3389/fmicb.2022.1065386

In the published article, there was an error in the Funding statement. The funding statement "the Foundation of Open Projects from State Key Laboratory of Trauma, Burns and Combined Injury (No. SKLKF201918), the Novel Clinical Technology Category of Army Medical University (XJSZR1206), and the Talent Training Project of Army Medical University (2019R031)" was incorrect. The correct Funding statement appears below.

Funding

This work was supported by the National Natural Science Foundation of China (grant nos. 82002051), the Natural Science Foundation of Chongqing CSTC (cstc2021jcyj-msxmX0655), and the Doctor Through Line Project of Chongqing CSTB (CSTB2022BSXM-JCX0019).

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.