



OPEN ACCESS

Edited by:

Kwangcheol Casey Jeong,
University of Florida, United States

Reviewed by:

Si Hong Park,
Oregon State University, United States

***Correspondence:**

Jiehong Fang
figo0726@163.com
Cheng Zhu
pzhch@cjlu.edu.cn

Specialty section:

This article was submitted to
Food Microbiology,
a section of the journal
Frontiers in Microbiology

Received: 30 June 2020

Accepted: 09 July 2020

Published: 17 August 2020

Citation:

Jiang H, Cheng H, Liang Y, Yu S, Yu T,
Fang J and Zhu C (2020)
Corrigendum: Diverse Mobile Genetic
Elements and Conjugal Transferability
of Sulfonamide Resistance Genes
(*sul1*, *sul2*, and *sul3*) in *Escherichia*
coli Isolates From *Penaeus vannamei*
and Pork From Large Markets in
Zhejiang, China.
Front. Microbiol. 11:1793.
doi: 10.3389/fmicb.2020.01793

Corrigendum: Diverse Mobile Genetic Elements and Conjugal Transferability of Sulfonamide Resistance Genes (*sul1*, *sul2*, and *sul3*) in *Escherichia coli* Isolates From *Penaeus vannamei* and Pork From Large Markets in Zhejiang, China

Han Jiang, Hui Cheng, Yi Liang, Shengtao Yu, Ting Yu, Jiehong Fang* and Cheng Zhu*

Key Laboratory of Marine Food Quality and Hazard Controlling Technology of Zhejiang Province, College of Life Sciences, China Jiliang University, Hangzhou, China

Keywords: sulfonamide resistance genes, *Escherichia coli*, mobile genetic elements, insertion sequences, conjugation

A Corrigendum on

Diverse Mobile Genetic Elements and Conjugal Transferability of Sulfonamide Resistance Genes (*sul1*, *sul2*, and *sul3*) in *Escherichia coli* Isolates From *Penaeus vannamei* and Pork From Large Markets in Zhejiang, China

by Jiang, H., Cheng, H., Liang, Y., Yu, S., Yu, T., Fang, J., et al. (2019). *Front. Microbiol.* 10:1787. doi: 10.3389/fmicb.2019.01787

In the original article, there was a mistake in **Figure 2C** as published. **Figure 2C** and **Figure 2B** were repeated. The corrected **Figure 2** appears 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.

Copyright © 2020 Jiang, Cheng, Liang, Yu, Yu, Fang and Zhu. 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.

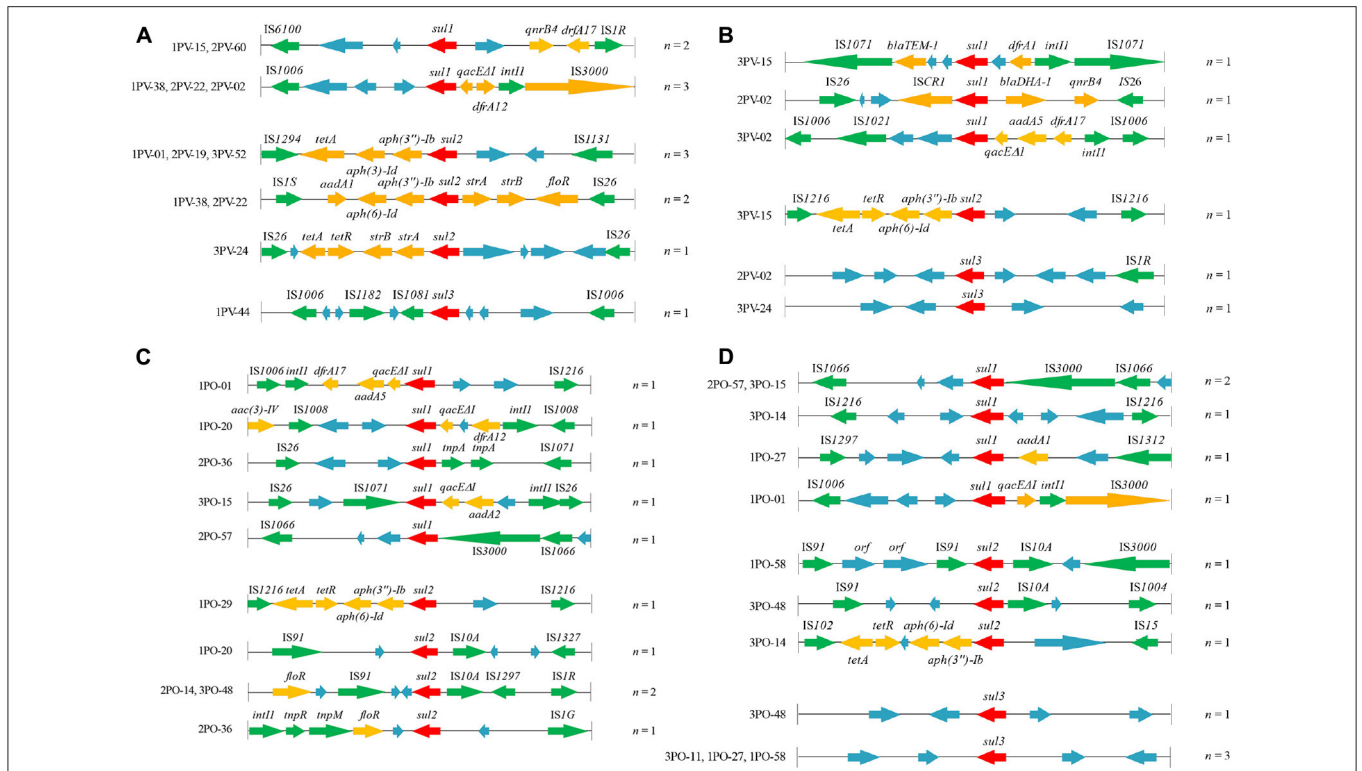


FIGURE 2 | Genetic organization of *sul* gene-associated regions in **(A)** plasmids of 12 *sul*-positive *Escherichia coli* isolates from *Penaeus vannamei*; **(B)** chromosomes of 12 *sul*-positive *E. coli* isolates from *P. vannamei*; **(C)** plasmids of 12 *sul*-positive *E. coli* isolates from pork products; and **(D)** chromosomes of 12 *sul*-positive *E. coli* isolates from pork products presented with their isolate numbers. The orientation of each gene and insertion element is indicated by arrows. The same units are shown in the same color. The same functional units or unknown functional units are shown in the same color (red, *sul* genes; yellow, antibiotic resistance genes other than *sul* genes; green, mobile genetic elements; blue, unknown functional unit). Names of sequence units are indicated above or below the arrows, and sequence units with unknown functions have been left blank.