



Corrigendum: Functional Repertoire of Antibiotic Resistance Genes in Antibiotic Manufacturing Effluents and Receiving Freshwater Sediments

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A Corrigendum on

Functional Repertoire of Antibiotic Resistance Genes in Antibiotic Manufacturing Effluents and Receiving Freshwater Sediments

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In the original article, the reference for Sugimoto et al. (2017) was incorrectly written as Sugimoto, A., Maeda, A., Itto, K., and Arimoto, H. (2017). Deciphering the mode of action of cell wall-inhibiting antibiotics using metabolic labeling of growing peptidoglycan in *Streptococcus pyogenes*. *Sci. Rep.* 7:1129. doi: 10.1038/s41598-017-01267-5. The correct reference appears in the Reference List 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.

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Sugimoto, Y., Suzuki, S., Nonaka, L., Boonla, C., Sukpanyatham, N., Chou, H.-Y., and Wu, J.-H. (2017). The novel *mef* (C)-*mph*(G) macrolide resistance genes are conveyed in the environment on various vectors. *J. Glob. Antimicrob. Resist.* 10, 47–53. doi: 10.1016/j.jgar.2017.03.015

Conflict of Interest Statement: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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