



Correction

Correction: Jacobson et al. Innovative Methodology for Antimicrobial Susceptibility Determination in *Mycoplasma* Biofilms. *Microorganisms* 2024, 12, 2650

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In the original publication [1], there was an error in the notation for the model on the bottom of page 8 (Section 2.15: Statistical Analysis), where it reads

$$\log(\text{FSC}_i) \sim \beta_1 X_{\text{Treatment}_i} + \epsilon_i \\ \epsilon_i \sim \text{IG}(\mu, \lambda)$$

where i represents the observation level.

The correct notation for the model fit by the statistical package we used is as follows:

$$\log(\mu_i) = \beta_0 + \beta_1 X_{\text{Treatment}_i} \\ \text{FSC}_i \sim \text{IG}(\mu_i, \lambda)$$

where i represents the individual observation, and IG refers to the inverse Gaussian distribution.

As this is an issue with the notation of a model in the methods section, it does not affect the results of the study. It does, however, affect the reproducibility of our methods, as the current notation is not equivalent to what was fitted into R.

This correction was approved by the Academic Editor. The original publication has also been updated.



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Reference

1. Jacobson, B.T.; DeWit-Dibbert, J.; Selong, E.T.; Quirk, M.; Throolin, M.; Corona, C.; Sonar, S.; Zanca, L.; Schwarz, E.R.; Bimczok, D. Innovative Methodology for Antimicrobial Susceptibility Determination in Mycoplasma Biofilms. *Microorganisms* **2024**, *12*, 2650. [\[CrossRef\]](#) [\[PubMed\]](#)

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