

CORRECTION

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Correction: Antimicrobial resistance and molecular typing of *Salmonella* in the chicken production chain in Hubei Province, China

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In the original publication of this article (Wu et al., 2022), the authors identified errors in Figs. 1 and 3. The correct figures are given below.

In addition, "*tetA* (40.91%, 9/22)" in the ninth line on the left column of page 7 needs to be changed into "*tetA* (45.45%, 10/22)".

The original article (Wu et al., 2022) was updated.

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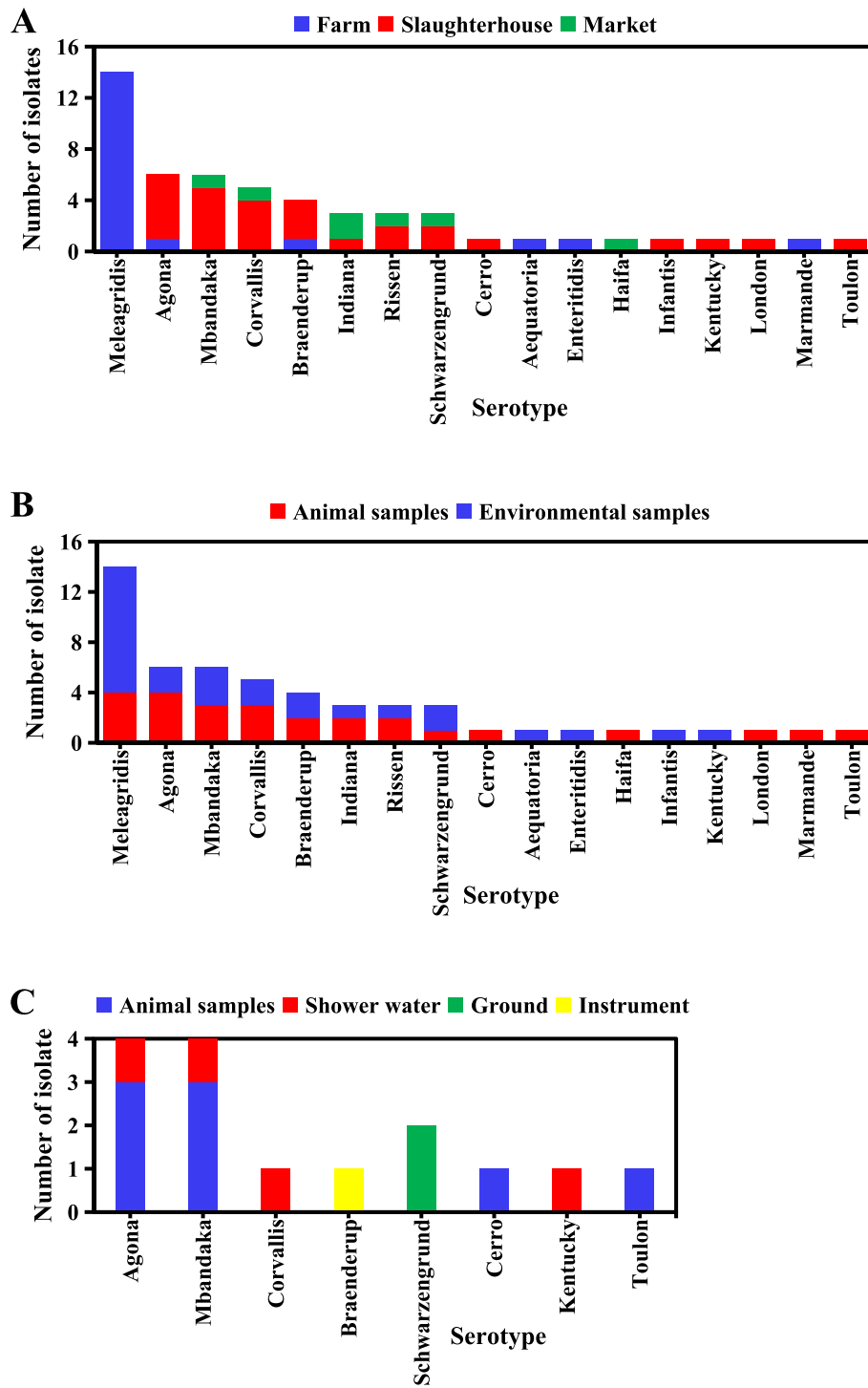


Fig. 1 Serotypes of 53 *Salmonella* isolated in the chicken production chain in Hubei Province, China. **A** Serotype distribution in different links. **B** Serotype distribution according to animal and environmental samples. **C** Serotype distribution in WYS slaughterhouse

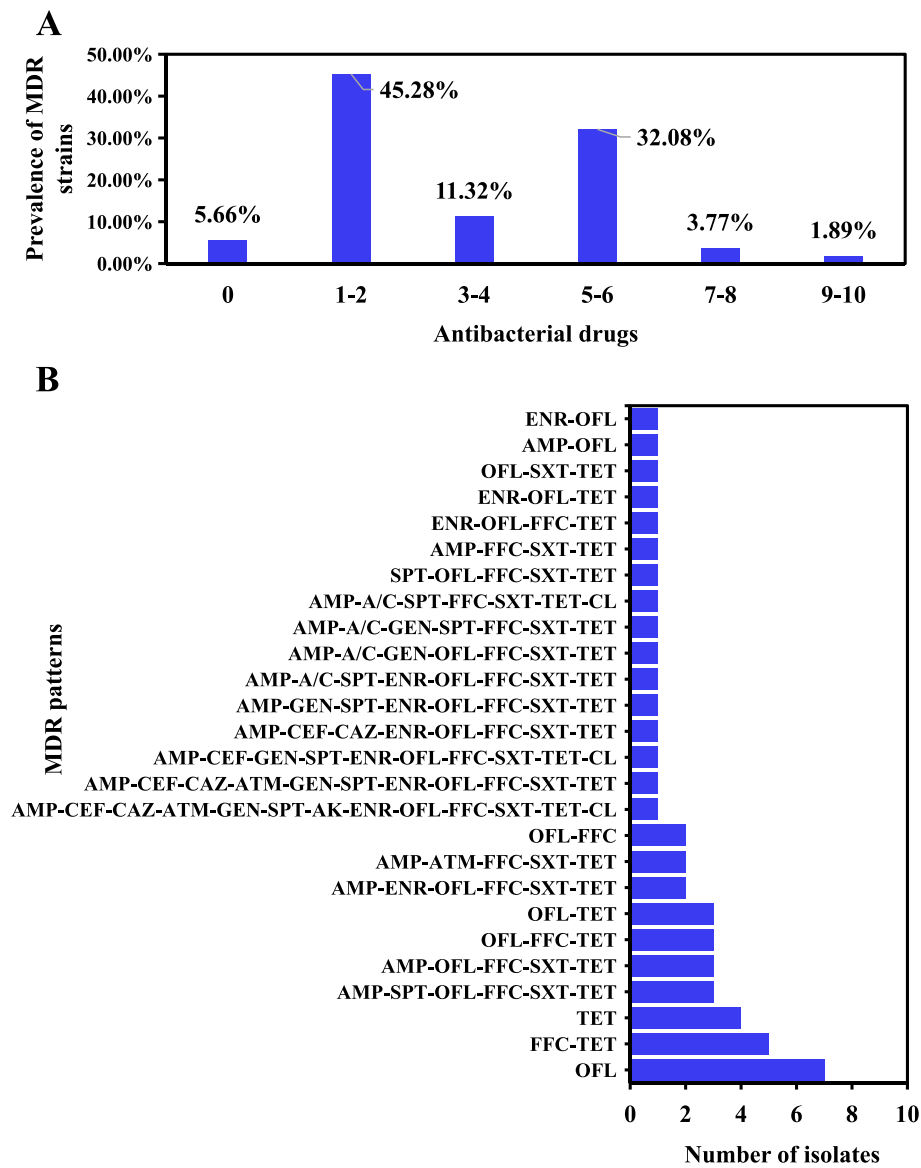


Fig. 3 Prevalence (A) and patterns (B) of multi-drug resistance of 53 *Salmonella* strains isolated from the chicken production chain in Hubei Province, China

Reference

Wu, Cuirong, et al. 2022. Antimicrobial resistance and molecular typing of *Salmonella* in the chicken production chain in Hubei Province, China. *Animal Diseases* 2: 31. <https://doi.org/10.1186/s44149-022-00063-7>.