

Minerva Access is the Institutional Repository of The University of Melbourne

Author/s:

Kordafshari, S;Shil, P;Marenda, MS;Olaogun, OM;Konsak-Ilievski, B;Disint, J;Noormohammadi, AH

Title:

Correction to: Preliminary comparative analysis of the genomes of selected field reisolates of the *Mycoplasma synoviae* vaccine strain MS-H reveals both stable and unstable mutations after passage in vivo (BMC Genomics, (2020), 21, 1, (598), 10.1186/s12864-020-06995-z)

Date:

2020-12-01

Citation:

Kordafshari, S., Shil, P., Marenda, M. S., Olaogun, O. M., Konsak-Ilievski, B., Disint, J. & Noormohammadi, A. H. (2020). Correction to: Preliminary comparative analysis of the genomes of selected field reisolates of the *Mycoplasma synoviae* vaccine strain MS-H reveals both stable and unstable mutations after passage in vivo (BMC Genomics, (2020), 21, 1, (598), 10.1186/s12864-020-06995-z). BMC Genomics, 21 (1), <https://doi.org/10.1186/s12864-020-07067-y>.

Persistent Link:

<https://hdl.handle.net/11343/251706>

License:

CC BY

CORRECTION

Open Access



# Correction to: Preliminary comparative analysis of the genomes of selected field reisolates of the *Mycoplasma synoviae* vaccine strain MS-H reveals both stable and unstable mutations after passage in vivo

Somayeh Kordafshari\*, Pollob Shil, Marc S. Marenda, Olusola M. Olaogun, Barbara Konsak-Ilievski, Jillian Disint and Amir H. Noormohammadi

**Correction to: BMC Genomics (2020) 21:598**  
<https://doi.org/10.1186/s12864-020-06995-z>

Following the publication of the original article [1], it was noted that due to a typesetting error the tables were ordered incorrectly. Tables 1, 2, and 3 were given as Table 2, 3 and 1 respectively.

The original article has been corrected.

Published online: 16 October 2020

## Reference

1. Kordafshari S, Shil P, Marenda MS, et al. Preliminary comparative analysis of the genomes of selected field reisolates of the *Mycoplasma synoviae* vaccine strain MS-H reveals both stable and unstable mutations after passage in vivo. *BMC Genomics*. 2020;21:598 <https://doi.org/10.1186/s12864-020-06995-z>.

The original article can be found online at <https://doi.org/10.1186/s12864-020-06995-z>.

\* Correspondence: [somayeh.kordafshari@unimelb.edu.au](mailto:somayeh.kordafshari@unimelb.edu.au)  
Asia Pacific Centre for Animal Health, Faculty of Veterinary & Agricultural Sciences, The University of Melbourne, Werribee, Victoria 3030, Australia



© The Author(s). 2020 **Open Access** This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit <http://creativecommons.org/licenses/by/4.0/>. The Creative Commons Public Domain Dedication waiver (<http://creativecommons.org/publicdomain/zero/1.0/>) applies to the data made available in this article, unless otherwise stated in a credit line to the data.