



Corrigendum: Genetic and Phenotypic Characterization of the Etiological Agent of Canine Orchiepididymitis Smooth *Brucella* sp. BCCN84.3

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A Corrigendum on

Genetic and Phenotypic Characterization of the Etiological Agent of Canine Orchiepididymitis Smooth *Brucella* sp. BCCN84.3

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In the original article, there was an error. In the Funding statement is written that MZ was granted with a fellowship from SEP, Universidad de Costa Rica. The correct Initials are MS-E.

A correction has been made to the **Funding** statement:

This work was supported by FEES-CONARE, Costa Rica; Fondo Institucional de Desarrollo Académico (FIDA), Universidad Nacional; Wellcome Trust; CITA-INIA, Spain (project Bru-Epidia 291815-FP7/ERANET/ANIHWA); MINECO (AGL2014-58795-CA), and Aragon Government (Consolidated Group A14). Authors from the Sanger Institute were supported by Wellcome Trust (098051). NR-V was partially sponsored by a scholarship from the University of Costa Rica. KB was founded by a Wellcome Trust Postdoctoral Training Fellowship for Clinicians (106690/Z/14/Z). MS-E was granted with a fellowship from SEP, Universidad de Costa Rica.

Additionally, an inaccurate sentence was included in Author Contributions stating that MZ kept the bacterium in ISP, INRA collection.

A correction has been made to **Author Contributions** statement:

EM, CG-V, and AC conceived the study. EM, CG-V, IM, NT, and JB obtained funding. EC and EM performed the isolation of the bacterium. CC-D performed fatty acid analysis. CG-V, MS-E, KB, AC, NR-V, MZ, EV-R, and MG performed genomics analyses. RC-Á, BA-A, and IM performed the LPS and lipid characterization. JB, EC, PM, and IM performed the bacteriological analysis. EM, CG-V, MS-E, AC, NR-V, NT, MG, and CC-D performed data interpretation. EM and CG-V wrote the paper. All authors read and approved the manuscript content.

The authors apologize for these errors and state that they do not change the scientific conclusions of the article in any way. The original article has been updated.

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