

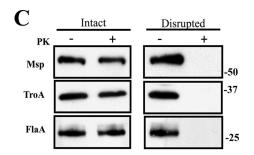
## **AUTHOR CORRECTION**

## Correction for Anand et al., The Major Outer Sheath Protein (Msp) of *Treponema denticola* Has a Bipartite Domain Architecture and Exists as Periplasmic and Outer Membrane-Spanning Conformers

Arvind Anand,<sup>a</sup> Amit Luthra,<sup>a</sup> Maxwell E. Edmond,<sup>a,b</sup> Morgan LeDoyt,<sup>a</sup> Melissa J. Caimano,<sup>a,c,f</sup> Justin D. Radolf<sup>a,c,d,e,f</sup>

Department of Medicine, Pediatrics, Genetics and Developmental Biology, Immunology, and Molecular Microbiology and Structural Biology, and the Health Careers Opportunity Program, University of Connecticut Health Center, Farmington, Connecticut, USA

Volume 195, no. 9, pages 2060 – 2071, 2013. Page 2066: Figure 4C should appear as shown below. An image containing the two rightmost lanes from Fig. 4B was inadvertently used to generate the top left panel in Fig. 4C (Intact T. denticola — /+ PK treatment immunoblotted with anti-Msp). In addition, the middle and bottom right panels (Disrupted T. denticola — /+ PK treatment immunoblotted with anti-TroA and anti-FlaA, respectively) were inadvertently switched in the composite figure. The corrected figure shows that, as stated in Results, Msp is largely inaccessible to proteinase K because it is not on the surface of the treponeme as opposed to being protease resistant.



Copyright © 2014, American Society for Microbiology. All Rights Reserved. doi:10.1128/JB.02013-14