ERRATUM Open Access



## Erratum to: An integrative Bayesian Dirichlet-multinomial regression model for the analysis of taxonomic abundances in microbiome data

W. Duncan Wadsworth<sup>1</sup>, Raffaele Argiento<sup>2</sup>, Michele Guindani<sup>3</sup>, Jessica Galloway-Pena<sup>4</sup>, Samuel A. Shelburne<sup>5</sup> and Marina Vannucci<sup>1\*</sup>

## **Erratum**

After publication of the original article [1] it was brought to our attention that author Samuel A. Shelburne was incorrectly included as Samuel A. Shelbourne. The correct spelling of the name is included in the author list of this erratum and has been updated in the original article.

## Author details

<sup>1</sup>Department of Statistics, Rice University, Houston, TX, USA. <sup>2</sup>ESOMAS Department, University of Torino and Collegio Carlo Alberto, Torino, Italy. <sup>3</sup>Department of Statistics, University of California, Irvine, CA, USA. <sup>4</sup>Department of Infectious Disease, Infection Control, and Employee Health, The University of Texas MD Anderson Cancer Center, Houston, TX 77030, USA. <sup>5</sup>Department of Genomic Medicine, The University of Texas MD Anderson Cancer Center, Houston, TX 77030, USA.

Received: 15 March 2017 Accepted: 15 March 2017 Published online: 23 March 2017

## Reference

 Wadsworth, et al. An integrative Bayesian Dirichletmultinomial regression model for the analysis of taxonomic abundances in microbiome data. BMC Bioinform. 2017;18:94. doi:10.1186/s12859-017-1516-0.

<sup>&</sup>lt;sup>1</sup>Department of Statistics, Rice University, Houston, TX, USA



<sup>\*</sup> Correspondence: marina@rice.edu