



Erratum for Kastman et al., “Biotic Interactions Shape the Ecological Distributions of *Staphylococcus* Species”

Erik K. Kastman,^a Noelani Kamelamela,^a Josh W. Norville,^a Casey M. Cosetta,^a
Rachel J. Dutton,^b Benjamin E. Wolfe^a

Department of Biology, Tufts University, Medford, Massachusetts, USA^a; Division of Biological Sciences, University of California, San Diego, La Jolla, California, USA^b

Volume 7, no 5, <https://doi.org/10.1128/mBio.01157-16>, 2016. In the Results section (PDF page 7), two errors were made in Fig. 3 that might cause confusion but do not change the major conclusions of the work. In Fig. 3E, labels that were intended to indicate the base pair position of genes in the staphyloferrin B operon along a contig in the *Staphylococcus equorum* BC9 genome were floating above the figure. In Fig. 3F, the *x* axis labels “CCA + *Penicillium*” and “CCA + *Scopulariopsis*” were switched. The revised Fig. 3 contains corrected versions of those labels.

Published 21 March 2017

Citation Kastman EK, Kamelamela N, Norville JW, Cosetta CM, Dutton RJ, Wolfe BE. 2017. Erratum for Kastman et al., “Biotic interactions shape the ecological distributions of *Staphylococcus* species.” *mBio* 8:e00329-17. <https://doi.org/10.1128/mBio.00329-17>.

Copyright © 2017 Kastman et al. This is an open-access article distributed under the terms of the [Creative Commons Attribution 4.0 International license](https://creativecommons.org/licenses/by/4.0/).

