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**ERRATUM**

**Genome-Based Cluster Deletion Reveals an Endocrocin Biosynthetic Pathway in *Aspergillus fumigatus***

Fang Yun Lim, Yanpeng Hou, Yiming Chen, Jee-Hwan Oh, Inhyung Lee, Tim S. Bugni, and Nancy P. Keller 6395

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Cover photograph (Copyright © 2012, American Society for Microbiology. All Rights Reserved.): Proline release by high-salinity stressed *Bacillus subtilis* cells. Osmotically stressed *B. subtilis* cells synthesize large amounts of the compatible solute proline (Pro) as a defense against the detrimental effects of high osmolarity on cellular water content and physiology. Cells of *B. subtilis* were spotted onto a lawn of *lac*<sup>+</sup> proline-auxotrophic *Escherichia coli* cells seeded on high-salinity minimal agar plates in the presence of IPTG (isopropyl-β-D-thiogalactopyranoside) and the LacZ indicator dye X-Gal (5-bromo-4-chloro-3-indolyl-β-D-galactopyranoside). *B. subtilis* cells carrying a defect in the osmotically inducible proline uptake system OpuE release proline, and this is visualized by a blue halo resulting from the growth of the cross-feed Lac<sup>+</sup>, proline-negative (Pro<sup>-</sup>) *E. coli* cells surrounding the *B. subtilis opuE* mutant. In contrast, the *B. subtilis opuE*<sup>+</sup> parent strain does not cross-feed the Pro<sup>-</sup> *E. coli* indicator strain, and this manifested by the absence of the blue halo. (See related article in August 2012, vol. 78, no. 16, p. 5753.)