PLOS ONE

CORRECTION

Correction: Expanding our Understanding of Sequence-Function Relationships of Type II Polyketide Biosynthetic Gene Clusters: Bioinformatics-Guided Identification of Frankiamicin A from *Frankia* sp. EAN1pec

Yasushi Ogasawara, Benjamin J. Yackley, Jacob A. Greenberg, Snezna Rogelj, Charles E. Melançon, III

The following information is missing from the Funding section: This work was supported by NIH NM-INBRE grant P20 GM103451 (CEM) and by start-up funds from University of New Mexico (CEM).

There are errors in the legend for Fig 1, "Structures of prototypical type II polyketides". Please view Fig 1 and its complete, correct legend here.

Citation: Ogasawara Y, Yackley BJ, Greenberg JA, Rogelj S, Melançon CE, III (2015) Correction: Expanding our Understanding of Sequence-Function Relationships of Type II Polyketide Biosynthetic Gene Clusters: Bioinformatics-Guided Identification of Frankiamicin A from *Frankia* sp. EAN1pec. PLoS ONE 10(6): e0129408. doi:10.1371/journal.pone.0129408

Published: June 1, 2015

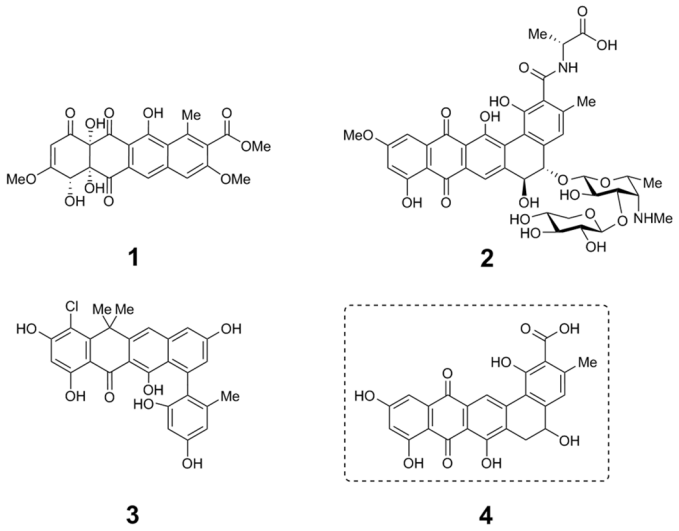

Fig 1. Structures of prototypical type II polyketides. Structures of tetracenomycin C (**1**), pradimicin A (**2**), fasamycin A (**3**), and the pentangular polyketide frankiamicin A (**4**) identified in this study.

doi:10.1371/journal.pone.0129408.g001

Reference

1. Ogasawara Y, Yackley BJ, Greenberg JA, Rogelj S, Melançon CE III (2015) Expanding our Understanding of Sequence-Function Relationships of Type II Polyketide Biosynthetic Gene Clusters: Bioinformatics-Guided Identification of Frankiamicin A from *Frankia* sp. EAN1pec. PLoS ONE 10(4): e0121505. doi: 10.1371/journal.pone.0121505 PMID: 25837682