

## CORRECTION

[View Article Online](#)  
[View Journal](#) | [View Issue](#)Cite this: *Chem. Sci.*, 2024, 15, 1162**Correction: Synthetic ramoplanin analogues are accessible by effective incorporation of arylglycines in solid-phase peptide synthesis**Edward Marschall,<sup>abc</sup> Rachel W. Cass,<sup>abc</sup> Komal M. Prasad,<sup>abc</sup> James D. Swarbrick,<sup>d</sup> Alasdair I. McKay,<sup>e</sup> Jennifer A. E. Payne,<sup>abc</sup> Max J. Cryle<sup>\*abc</sup> and Julien Tailhades<sup>\*abc</sup>

DOI: 10.1039/d3sc90245e

[rsc.li/chemical-science](https://rsc.li/chemical-science)Correction for 'Synthetic ramoplanin analogues are accessible by effective incorporation of arylglycines in solid-phase peptide synthesis' by Edward Marschall *et al.*, *Chem. Sci.*, 2024, 15, 195–203, <https://doi.org/10.1039/D3SC01944F>.

The authors regret that the number of distance restraints used to obtain the final structure ensemble was incorrectly stated on the sixth page of the manuscript. The corrected sentence should read: "The final structure ensemble was calculated in CYANA<sup>67</sup> and is based on 330 assigned NOEs (Table S3 and Fig. S6†) which included 104 long-range NOEs and had good precision (backbone RMSD = 0.17 Å, heavy atom RMSD = 0.49 Å) and residual restraint violations."

The Royal Society of Chemistry apologises for these errors and any consequent inconvenience to authors and readers.

<sup>a</sup>Department of Biochemistry and Molecular Biology, The Monash Biomedicine Discovery Institute, Monash University, Clayton, VIC 3800, Australia. E-mail: max.cryle@monash.edu; julien.tailhades@monash.edu

<sup>b</sup>EMBL Australia, Monash University, Clayton, VIC 3800, Australia

<sup>c</sup>ARC Centre of Excellence for Innovations in Peptide and Protein Science, Clayton, VIC 3800, Australia

<sup>d</sup>Department of Microbiology, Monash University, Clayton, VIC 3800, Australia

<sup>e</sup>Department of Chemistry, Monash University, Clayton, VIC 3800, Australia

