CORRIGENDUM

Corrigendum: Tripropeptins, novel antimicrobial agents produced by *Lysobacter* sp. II. structure elucidation

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Correction to: *The Journal of Antibiotics* (2004) **57**, 52–58; doi:org/10.7164/antibiotics.57.52

The authors of the above article did misjudgment the stereochemistry of tripropeptins-constituent serine residue which was determined by Marfey's method using HPLC. From our recent analysis using

advanced Marfey's method (LC/MS) revealed that the stereochemistry of serine residue was proved to be D-form as shown below. Resulting chromatograms and the conditions were shown in Supplementary Figure S1.

Adding to above, we corrected the Figures 1 and 3 as follows. Authors would like to apologize for the mistakes.

Figure S1 Extracted ion chromatograms of L-FDLA derivatives of (A) TPPC hydrolysate and (B) authentic serines. Acid hydrolysate of tripropeptin C (TPPC) and authentic serines were coupled with *N*^ε-(5-fluoro-2,4-dinitrophenyl)-L-leucinamide (FDLA). Resulting samples were subjected to LC-MS analysis. Extracted ion chromatograms, monitoring at *m/z* 400.14–400.15 (FDLA coupled serine) were shown. HPLC conditions of Supplementary Figure S1: Device: LTQ Orbitrap (Thermo Fisher Scientific, Waltham, MA). Column: Shiseido Capcell Pak UG120, 2 × 150 mm, 5 μm. Solvent system: 0–20 min, 20%A–60%A linear gradient. A: CH₃CN–0.01% HCOOH, B: 0.01% HCOOH. Flow rate: 0.3 ml min^{–1}. Temperature: 50 °C. A full color version of this figure is available at *The Journal of Antibiotics* online.

Figure 1 Structure of tripropeptins.

Figure 3 Amino acid analysis of the fragments 6 and 7.