Erratum to: Enzymatic properties of the glycine D-alanine aminopeptidase of *Aspergillus oryzae* and its activity profiles in liquid-cultured mycelia and solid-state rice culture (rice koji)

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The original version of this article unfortunately contained spelling errors. The term “d-alanine” in the title and text should be written as “D-alanine”. The correct title is given above.

The online version of the original article can be found at <http://dx.doi.org/10.1007/s00253-011-3610-y>.

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In addition, the Fig. 1 is not presented properly. For the reader's convenience, the corrected figure and its legend are presented here:

Fig. 1 Schematic representations of the *gdaA* gene and its encoded protein. **a** Structure of the *gdaA* gene. The exon and intron of the *gdaA* gene are represented as *filled* and *open boxes*, respectively. The transcription and translation start sites deduced in the present study are indicated by -81 and +1, respectively. The in silico-predicted translation start site in *A. oryzae* genome database is indicated at position +182 by an *arrowhead*. **b** Deduced functional domains of GdaA. The serine β -lactamase fold and the D-aminopeptidase domains B and C predicted by a sequence search of the Pfam database are indicated by *dotted*, *hatched*, and *crosshatched boxes*, respectively, referring to the N-terminal Met as position 1. The Ser-Xaa-Lys motif at positions 58 to 61 in the serine β -lactamase fold is represented