

Structural and functional insights into an archaeal L-asparaginase obtained through the linker-less assembly of constituent domains. Corrigendum

Rachana Tomar,^a Pankaj Sharma,^{b†} Ankit Srivastava,^a Saurabh Bansal,^a Ashish^b and Bishwajit Kundu^{a*}

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^aKusuma School of Biological Sciences, Indian Institute of Technology Delhi, New Delhi, India, and ^bCSIR – Institute of Microbial Technology, Chandigarh, India. *Correspondence e-mail: bkundu@bioschool.iitd.ac.in

A correction is made to Fig. 7 in the article by Tomar *et al.* [(2014). *Acta Cryst. D70*, 3187–3197].

Keywords: L-asparaginase; *Pyrococcus furiosus*; linker-less assembly; corrigendum

In the article by Tomar *et al.* (2014), the product formula in Fig. 7 was incorrect. A new version of the figure with the correct formula is given here.

References

Tomar, R., Sharma, P., Srivastava, A., Bansal, S., Ashish, & Kundu, B. (2014). *Acta Cryst. D70*, 3187–3197.

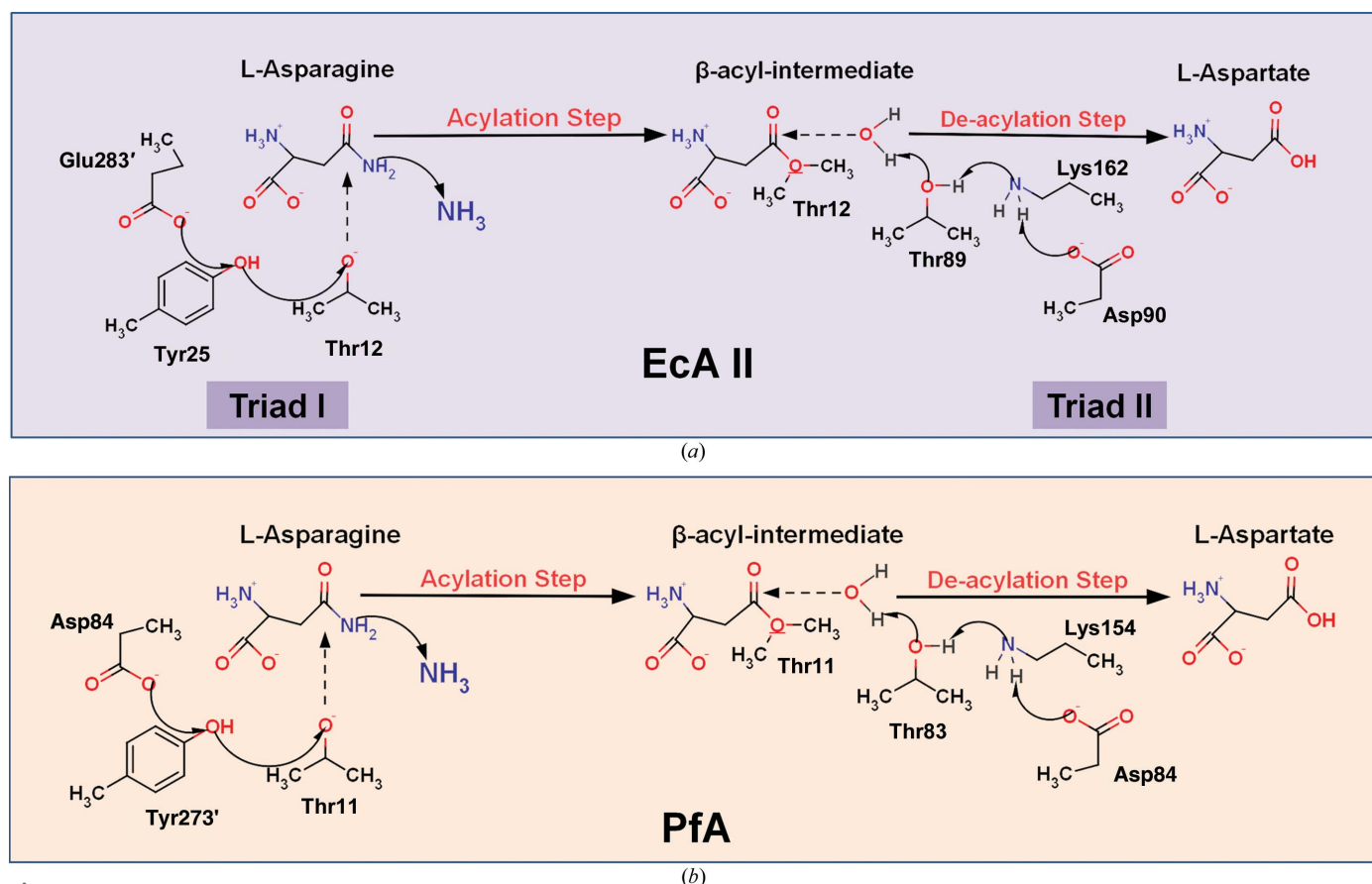


Figure 7

Reaction mechanism at the EcAII (a) and PfA (b) catalytic triads. A common acidic residue Asp84 is present in both triads I and II of PfA, with Tyr273' in triad I as the basic residue.