

## Crystallization and preliminary X-ray study of the deaminase AmnE from *Pseudomonas* sp. AP-3. Corrigendum

Dan Yu,<sup>a,b</sup> Yongji Jiang,<sup>a,b</sup> Jianfeng Hou,<sup>c</sup> Shuai Chen,<sup>b</sup>  
Guofang Zhang,<sup>a,b</sup> Xiang Liu,<sup>b</sup> Hui Dong<sup>b\*</sup> and Bo Yu<sup>c\*</sup>

<sup>a</sup>Tianjin University of Science and Technology, Tianjin 300222, People's Republic of China, <sup>b</sup>Tianjin International Joint Academy of Biotechnology and Medicine, Tianjin 300457, People's Republic of China, and <sup>c</sup>Institute of Microbiology, Chinese Academy of Sciences, Beijing 100101, People's Republic of China

Correspondence e-mail: hui.dong@htmdc.org, yub@im.ac.cn

The article by Yu *et al.* (2013, *Acta Cryst.* **F69**, 812–814) is corrected.

---

In the article by Yu *et al.* (2013) the Matthews coefficient and solvent content are incorrect. The correct Matthews coefficient should be  $2.10 \text{ \AA}^3 \text{ Da}^{-1}$  instead of  $3.15 \text{ \AA}^3 \text{ Da}^{-1}$ , corresponding to a solvent content of 41.53% instead of 61.02%. In Table 1 the resolution range should be 50.00–2.09 Å. The number of observed reflections should be 319580 (10109) instead of 22932 (752), and the number of unique reflections should be 23327 (1087) instead of 14970 (708).

### References

Yu, D., Jiang, Y., Hou, J., Chen, S., Zhang, G., Liu, X., Dong, H. & Yu, B. (2013). *Acta Cryst.* **F69**, 812–814.