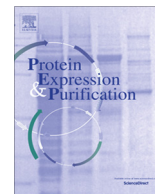




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Corrigendum

Corrigendum to “High-level over-expression, purification, and crystallization of a novel phospholipase C/sphingomyelinase from *Pseudomonas aeruginosa*” [Protein Expr. Purif. 90 (2013) 40–46]Daphné Truan ^{a,1}, Adriana Vasil ^b, Martin Stonehouse ^b, Michael L. Vasil ^{b,*}, Ehmke Pohl ^{c,*}^a Swiss Light Source, Paul Scherrer Institute, 5232 Villigen, Switzerland^b Department of Microbiology, University of Colorado, School of Medicine, Anschutz Medical Center, Aurora, CO 80045, USA^c Department of Chemistry & School of Biological and Biomedical Sciences, Durham University, Durham DH1 3LE, UK

The authors regret that the reported microcrystals of L-selenomethionine PlCHR2 (space group C2 with $a = 157.9 \text{ \AA}$, $b = 75.4 \text{ \AA}$, $c = 141.0 \text{ \AA}$, $\beta = 93.2$) presented in Table 1 on page 44 have been solved and identified as the putative cysteine hydrolase YcaC from *Pseudomonas aeruginosa*. YcaC has presumably been co-purified as minor contaminant and crystallized from the L-selenomethionine protein preparations. Details of the structure determination and analysis of YcaC will be described elsewhere (Groftehauge, Truan, Vasil, Denny, Vasil & Pohl, 2014). The crystal structure analysis of native PlCHR2 (space group C222₁, $a = 175.5 \text{ \AA}$, $b = 196.4 \text{ \AA}$, $c = 325.3 \text{ \AA}$) also reported in Table 1 is currently underway. The authors would like to apologize for any inconvenience caused.

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