

FEBS 20373

Corrigendum to: **Molecular characterisation of *meaB*, a novel gene affecting nitrogen metabolite repression in *Aspergillus nidulans*** (FEBS 17512)

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It has been brought to our attention [1] that a number of errors were included in the published sequence of *meaB*. We have re-analysed the sequence data and re-sequenced regions of ambiguity. The principal discrepancy noted at the 5' end of the gene arose due to the omission of a 54 nucleotide intron during compilation of the sequence. This intron occurs immediately downstream of the first ATG of the coding region in the published genomic sequence. There is therefore no ambiguity relating to the N-terminus of the MEAB protein. The most significant error was a 1 bp frameshift that affects the C-terminus of the putative MEAB protein, which is 37 residues shorter than previously reported. None of the errors

significantly alters the general conclusion of the paper nor does it disrupt the homology of MEAB to the Yap proteins reported by Fernandes et al. [2]. A fully revised sequence is available from the EMBL database (X98065).

References

- [1] Lamb, H.K., Dodds, A.L., Swatman, D.R., Cairns, E. and Hawkins, A.R. (1997) *J. Bacteriol.* 179, 6649–6656.
- [2] Fernandes, L., Rodrigues-Pousada, C. and Struhl, K. (1997) *Mol. Cell. Biol.* 17, 6982–6993.

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Corrigendum to: **Expression, purification and preliminary crystal analysis of the human low M_r phosphotyrosine protein phosphate isoform 1** (FEBS 20075)

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The authors wish to correct an error in Table 1. The correct table is reproduced below.

Second, on page 55, the sentence beginning on line 9, left column, should read: IF1 and IF2 specific activities, about 80 IU/mg and 120 IU/mg, respectively, were slightly higher than those reported for the same enzymes purified from natural sources.

Table 1
IF1 and IF2 purification parameters^a

Step	Volume (ml)	Total protein (mg)	Total activity (IU)	Purification (fold)	Yield (%)
Cell lysate (IF1)	150	200.0	2500	1.0	100.0
Cell lysate (IF2)	150	200.0	7000	1.0	100.0
Affinity (IF1)	20	18.7	1492	4.8	59.6
Affinity (IF2)	20	35.9	4378	3.5	62.5

^aFrom a 2-l culture.