Errata and Corrigenda

FEBS 16806

Corrigendum to: Compositional compartmentalization of the nuclear genomes of *Trypanosoma brucei* and *Trypanosoma equiperdum* (FEBS 13319)

[FEBS Letters 335 (1993) 181–183]

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In the Discussion of this paper it was mentioned that the present results corrected "an old report based on preparative CsCl banding [18] according to which the main band of nuclear DNA of *T. brucei* is centered at 1.703 g/cm³, with a shoulder at 1.702 g/cm³." The reference quoted, Borst, P., Fase-Fowler, F. Frash, A.C.C., Hoeijmakers, J.H.J. and Weijers, P.J. (1980) Mol. Biochem. Parasitol. 1, 221–246, was, however, wrong.

The authors apologize to P. Borst et al. for misquoting them. So far, the authors have been unable to retrieve the right reference.

FEBS 16807

Corrigendum to: Two distinct inwardly rectifying conductances are expressed in long term dibutyryl-cyclic-AMP treated rat cultured cortical astrocytes (FEBS 15660)

[FEBS Letters 367 (1995) 319-325]*

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The following corrections should have been made to the manuscript prior to publication.

Section 2.3. Electrophysiological recordings

The standard external solution of the KCl content is lacking at line 9. The composition of the used solution is (mM): 140 NaCl, 4 KCl, 2 CaCl₂, 2 MgCl₂, 5 TES, 5 EGTA, 5 glucose, buffered with NaOH at pH 7.3.

Section 3. Results

On page 323, lines 6-7 (right column) the used sodiumbutyrate concentration is 250 μ M.

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^{*}SSDI 0014-5793(95)00588-9

Erratum to: Lipopolysaccharide treatment in vivo induces tissue expression of GTP cyclohydrolase mRNA (FEBS 15761)

[FEBS Letters 368 (1995) 336-338]*

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As the result of a clerical error Fig. 2 was published wrongly.

Please see below for the correct Fig. 2 plus legend.

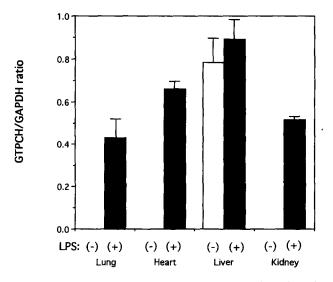


Fig. 2. Effects of LPS on GTPCH mRNA expression in various tissues. The intensity ratio of GTPCH to GAPDH was shown. Data are mean \pm S.E.M. of 3 animals (control: open columns, LPS-treated: closed columns). GTPCH/GAPDH in control lung, heart and kidney was not shown as the GTPCH signal was too low to be quantified.

^{*}SSDI 0014-5793(95)00689-3

Corrigendum to: Effect of polyethylene glycol on the activity, intrinsic fluorescence, and oligomeric structure of castor seed cytosolic-1,6-bisphophatase (FEBS 15816)

[FEBS Letters 368 (1995) 559-562]*

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On page 360, section 3.1, the last sentence of the first paragraph should have read: The I_{50} for F-2,6-P₂ at 50 and 5 μ M F-1,6-P₂ was 500 and 60 nM, respectively, in the presence of 0%, 10%, or 20% (w/v) PEG. (500 instead of 50 nM)

*SSDI 0014-5793(95)00744-X

FEBS 16810

Erratum to: A mutation in the 5' untranslated region of the human α -galactosidase A gene in high-activity variants inhibits specific protein binding (FEBS 15963)

[FEBS Letters 371 (1995) 181-184]*

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Affiliation a was assigned to the wrong Medical School and should have read:

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^{*}SSDI 0014-5793(95)00891-8

Corrigendum to: Classification of multi-helical DNA-binding domains and application to predict the DBD structures of σ factor, LysR, OmpR/PhoB, CENP-B, Rap1, and XylS/Ada/AraC (FEBS 16060)

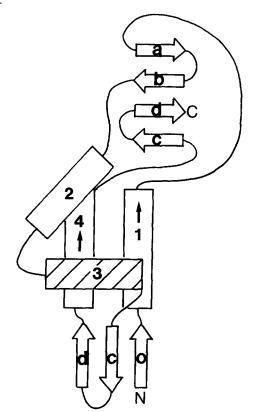
[FEBS Letters 372 (1995) 215-221]*

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Fig. 2j of this paper was wrongly drawn. One of the important points of this figure was to show that helices 1 and 4 are nearly parallel and can substitute for each other. The published figure satisfies this point but the two helices are placed the wrong way around. Helix 1 should be on the right and helix 4 should be on the left.

Please see below for the correct Fig. 2j.



^{*}SSDI 0014-5793(95)00988-4

Corrigendum to: Molecular cloning, functional expression, and signal transduction of the GIP-receptor cloned from a human insulinoma (FEBS 16078)

[FEBS Letters 373 (1995) 23-29]*

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A low-quality computer-generated Fig. 1 was published in the above paper.

Please see below for a better quality reproduction of the Northern blot to clearly demonstrate that the authors were able to determine the transcript size of 5.5 kb from total RNA of the insulinoma based upon a clear cut result in a Northern Blot rather than by simple guessing.

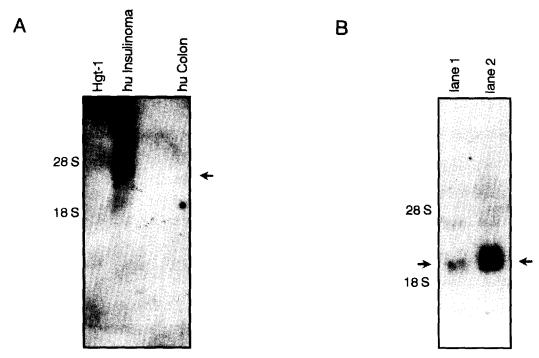


Fig. 1. Northern blot analysis of GIP receptor expression in (A) tissue of a human insulinoma, colon, stomach cancer HGT 1 cells, and (B) transfected CHL cells. A 1.0 kb fragment of the insulinoma-derived GIP receptor cDNA was utilized as probe. In (A) only total RNA from the insulinoma showed a transcript of 5.5 kb. In transfected CHL cells stably expressing the GIP receptor (B) the expected signal at 2.6 kb was detected (arrows) for both the intact and the cDNA containing the alternatively spliced II \(\textit{II} \) 62 fragment.

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^{*}SSDI 0014-5793(95)01006-8

Erratum to: Stimulation of cloned human glucagon-like peptide 1 receptor expressed in HEK 293 cells induces cAMP-dependent activation of calcium-induced calcium release (FEBS 16142)

[FEBS Letters 373 (1995)182-186]*

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As the result of a handling error during the final stages of printing Figs. 2 and 4 were interchanged.

Figures 2 and 4, therefore, should be interchanged without interchanging the legends.

*SSDI 0014-5793(95)01070-X

FEBS 16814

Corrigendum to: Characterisation of a synergohymenotropic toxin produced by Staphylococcus intermedius (FEBS 16331)

[FEBS Letters 376 (1995) 135-140]*

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In the original manuscript the legends to figures 1, 2 and 3 were interchanged.

The legends of Figures 1, 2 and 3 should be read as Figures 3, 1 and 2, respectively.

^{*}SSDI 0014-5793(95)01260-5