CORRECTION

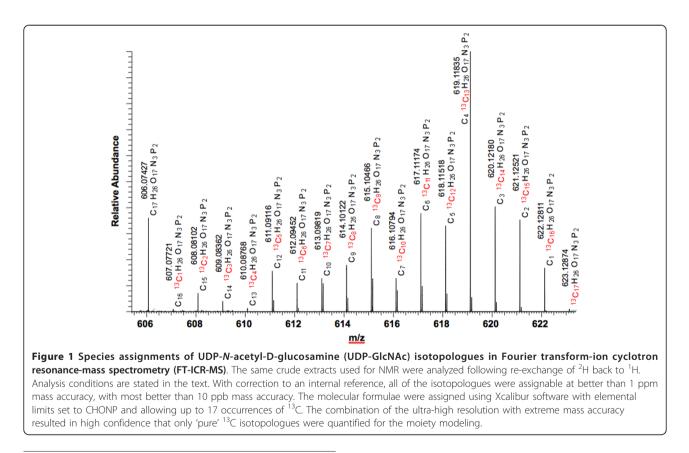


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A novel deconvolution method for modeling UDP-*N*-acetyl-D-glucosamine biosynthetic pathways based on ¹³C mass isotopologue profiles under non-steady-state conditions

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The figure published as Figure 2 in the original published version of the manuscript is in fact a duplicate of Figure 5. The correct Figure 2 is shown here (Figure 1 in this correction). Note that the legend for Figure 2 and references to it in the main text apply to the correct Figure 2. The authors and publisher regret the error.



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Received: 17 August 2012 Accepted: 17 August 2012 Published: 17 August 2012

Reference

 Moseley HNB, Lane AN, Belshoff AC, Higashi RM, Fan TWM: A novel deconvolution method for modeling UDP-N-acetyl-D-glucosamine biosynthetic pathways based on ¹³C mass isotopologue profiles under non-steady-state conditions. *BMC Biology* 2011, 9:37.

doi:10.1186/1741-7007-10-74

Cite this article as: Moseley *et al.*: A novel deconvolution method for modeling UDP-*N*-acetyl-D-glucosamine biosynthetic pathways based on ¹³C mass isotopologue profiles under non-steady-state conditions. *BMC Biology* 2012 **10**:74.

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