

CORRECTION

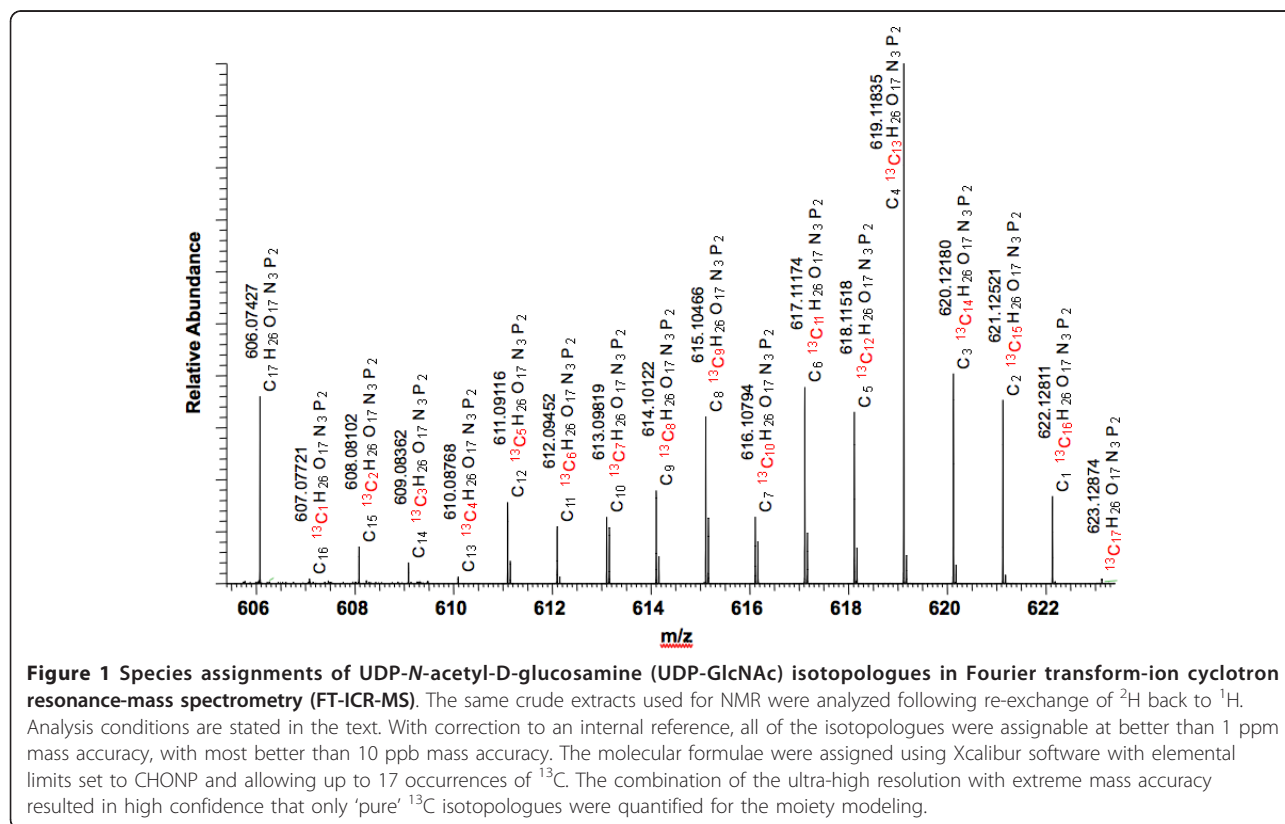
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# A novel deconvolution method for modeling UDP-*N*-acetyl-D-glucosamine biosynthetic pathways based on $^{13}\text{C}$ mass isotopologue profiles under non-steady-state conditions

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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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