



Synthesis of fluorescently labelled nucleoside diphosphate 6-amino-6-deoxy-β-L-galactoses

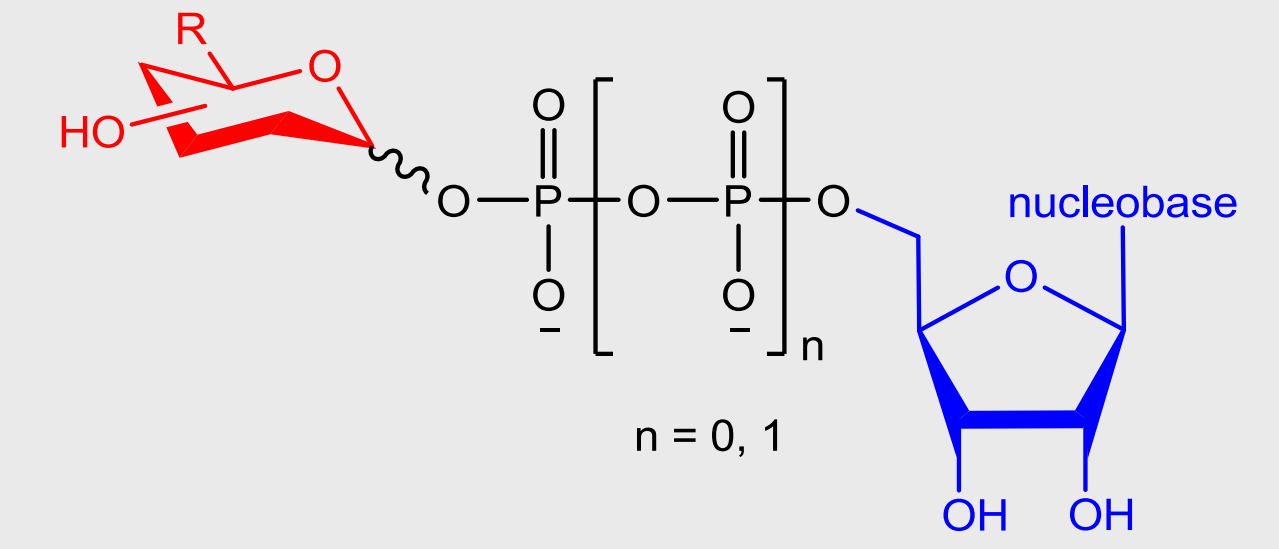
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Introduction

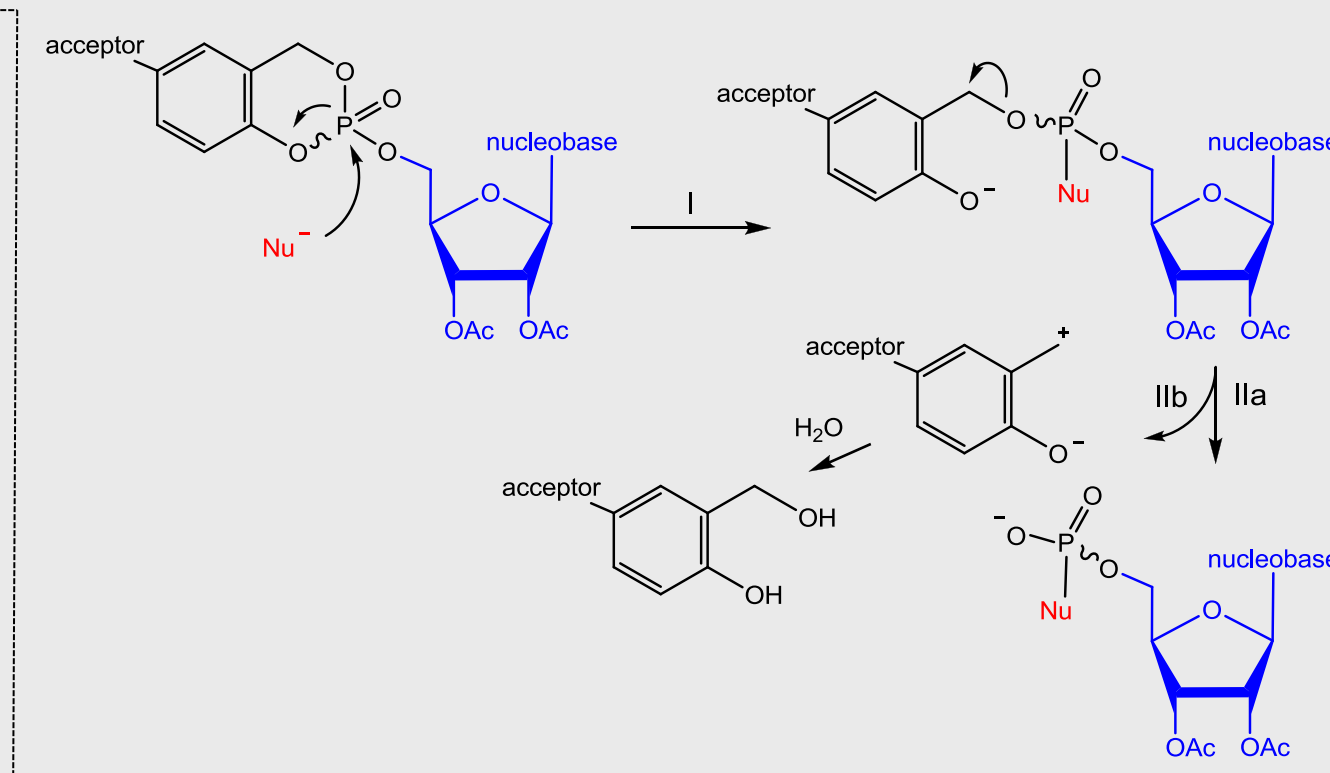
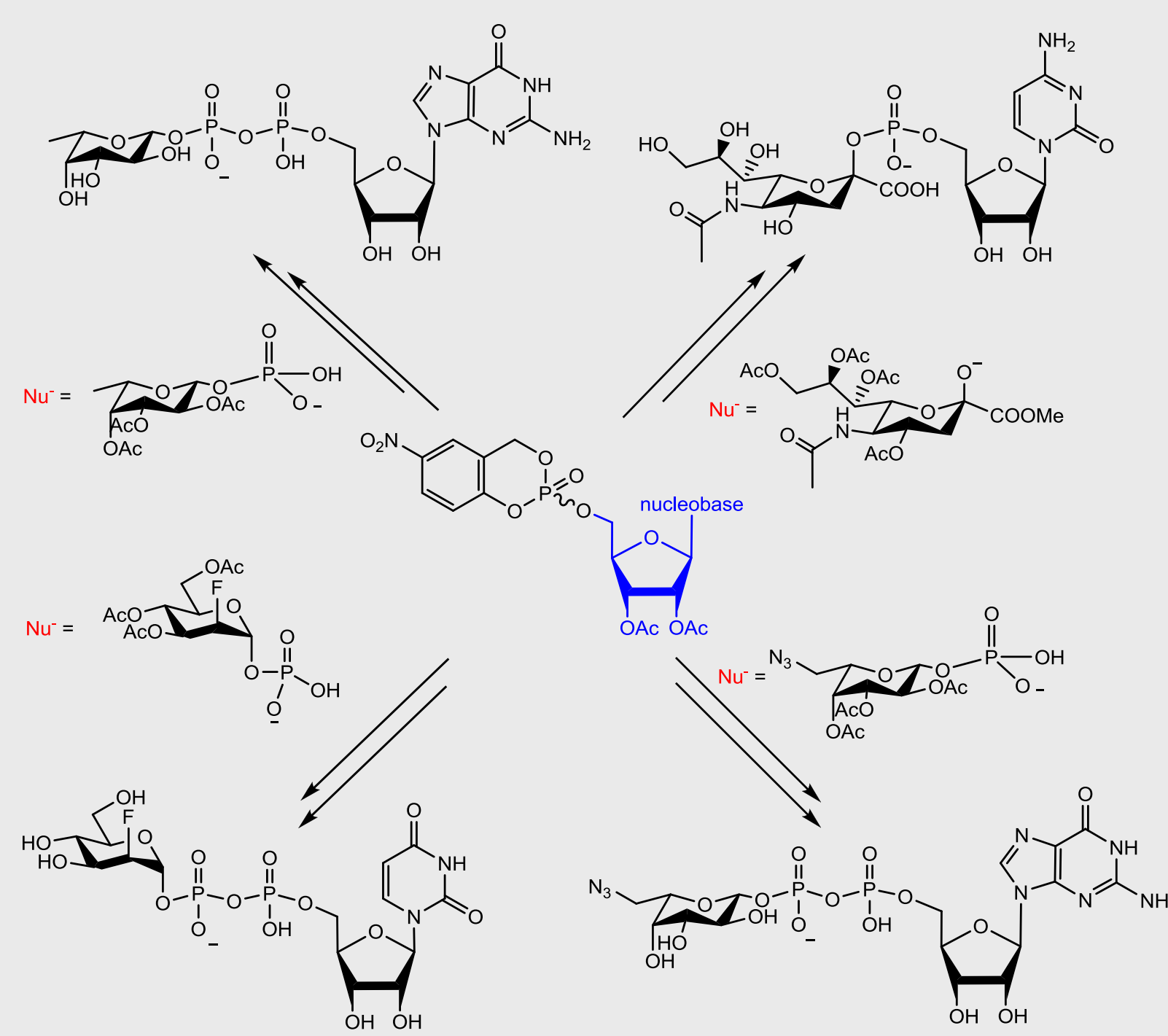
Glycosylation is one of the most complex posttranslational modifications of proteins and shows an enormous structural diversity. Furthermore, it influences several essential biological processes from storage to signaling.¹ Nucleoside monophosphate (NMP) and nucleoside diphosphate (NDP) sugars are glycosyldonors and serve therefore as substrates in various biochemical processes. The sugar transfer to specific acceptor molecules is catalysed by enzymes, generally glycosyltransferases.² Fucosylation, mediated by fucosyltransferases (FucT) from GDP-L-fucose, is often the final glycosylation step in the biosynthesis of biologically significant glycans. It is important for basic physiological and pathological procedures such as cell-cell-communication, immune response or tumor metastasis. Since fucosyltransferases show a great tolerance to modification in position C6 of the substrate³, it is possible to label a NDP fucose derivative with a dye in this position.



The cycloSal-method

The cycloSal-approach has been proven to be a suitable method for the synthesis of a great number of biological significant biomolecules, including the formation of natural and modified nucleoside diphosphate sugars.^{4,5} The cycloSal-triesters serve here as activated phosphate donors and can be attacked by a vast number of nucleophiles, for example sugar phosphates.⁵ By using 5-acceptor substituted cycloSal-nucleotides the electrophilicity of the phosphorous atom can be increased.⁶

Examples of previously prepared NM(D)P-conjugates

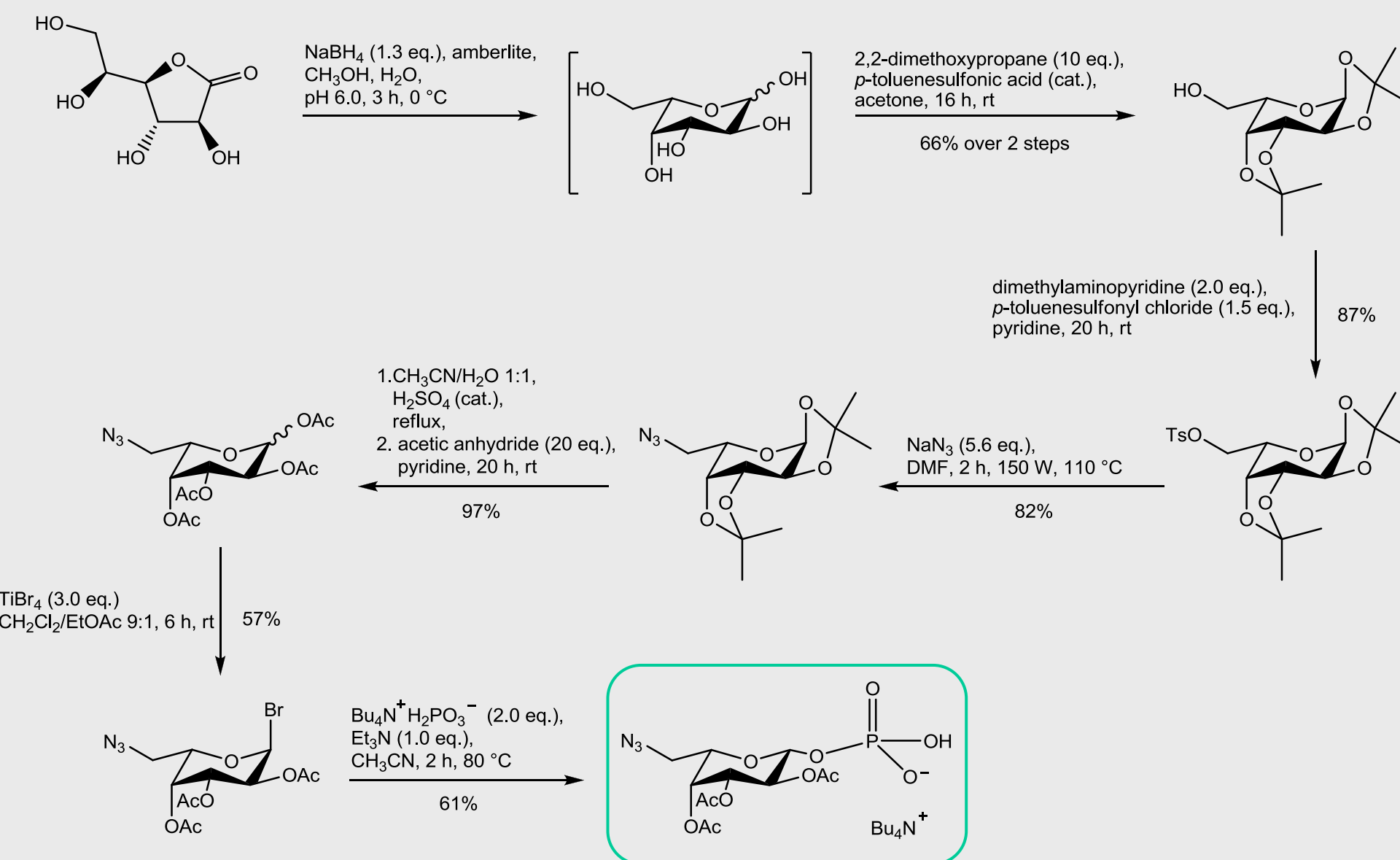


- I: The nucleophile attacks the cycloSal-triester. Since the phenolic ester bond is the most labile one, it breaks at first.
- IIa: The second step is spontaneous, since the phenolate destabilises the benzylic ester bond. The product is released.
- IIb: After hydrolysis, the released cycloSal-mask is converted to the corresponding salicyl alcohol.

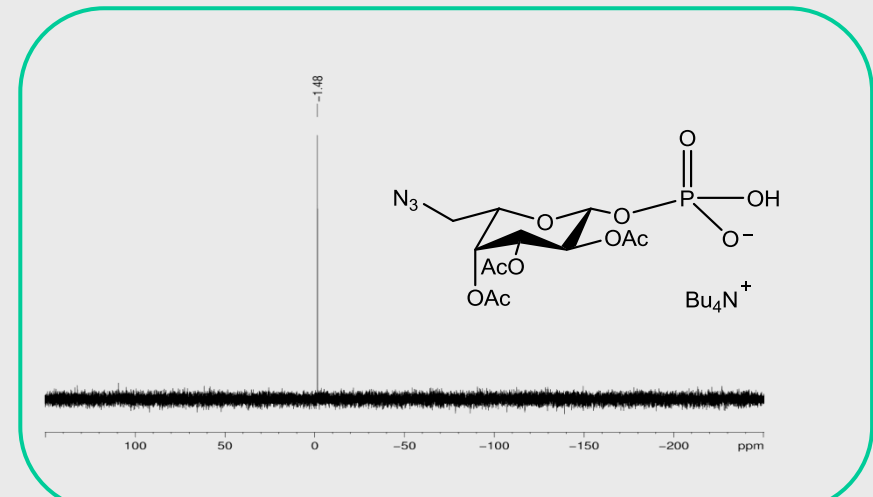
Objectives

Since detailed information about the reaction processes for most fucosyltransferases are still uncertain, the aim of this project was to synthesise fluorescently labelled NDP fucose derivatives and using them as substrates in different biochemical assays, e.g. monitoring fucosyltransferase activity. Furthermore, the protein fetuin ought to be labelled with a dye to visualise fucosylated glycans later.

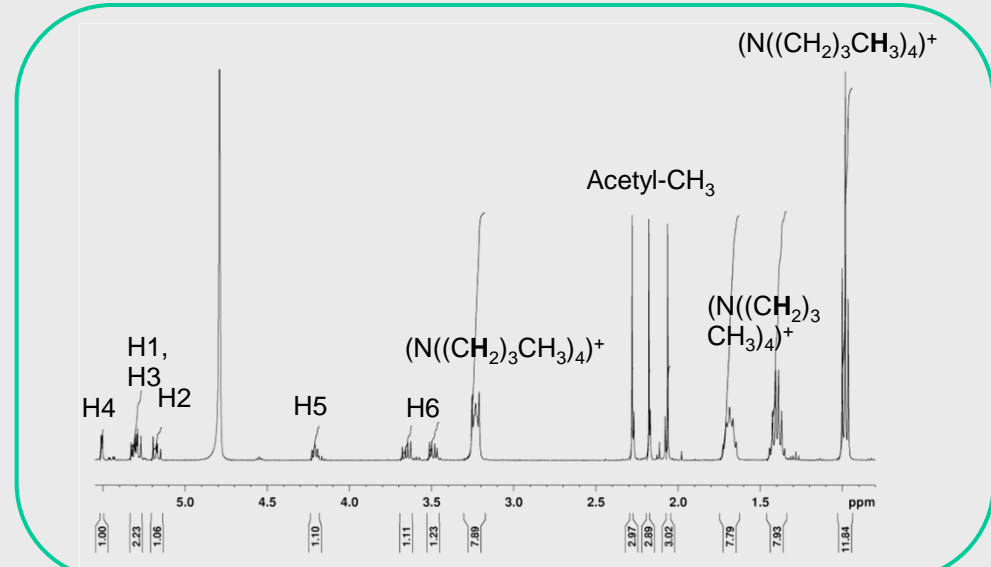
Synthesis of the sugar phosphate



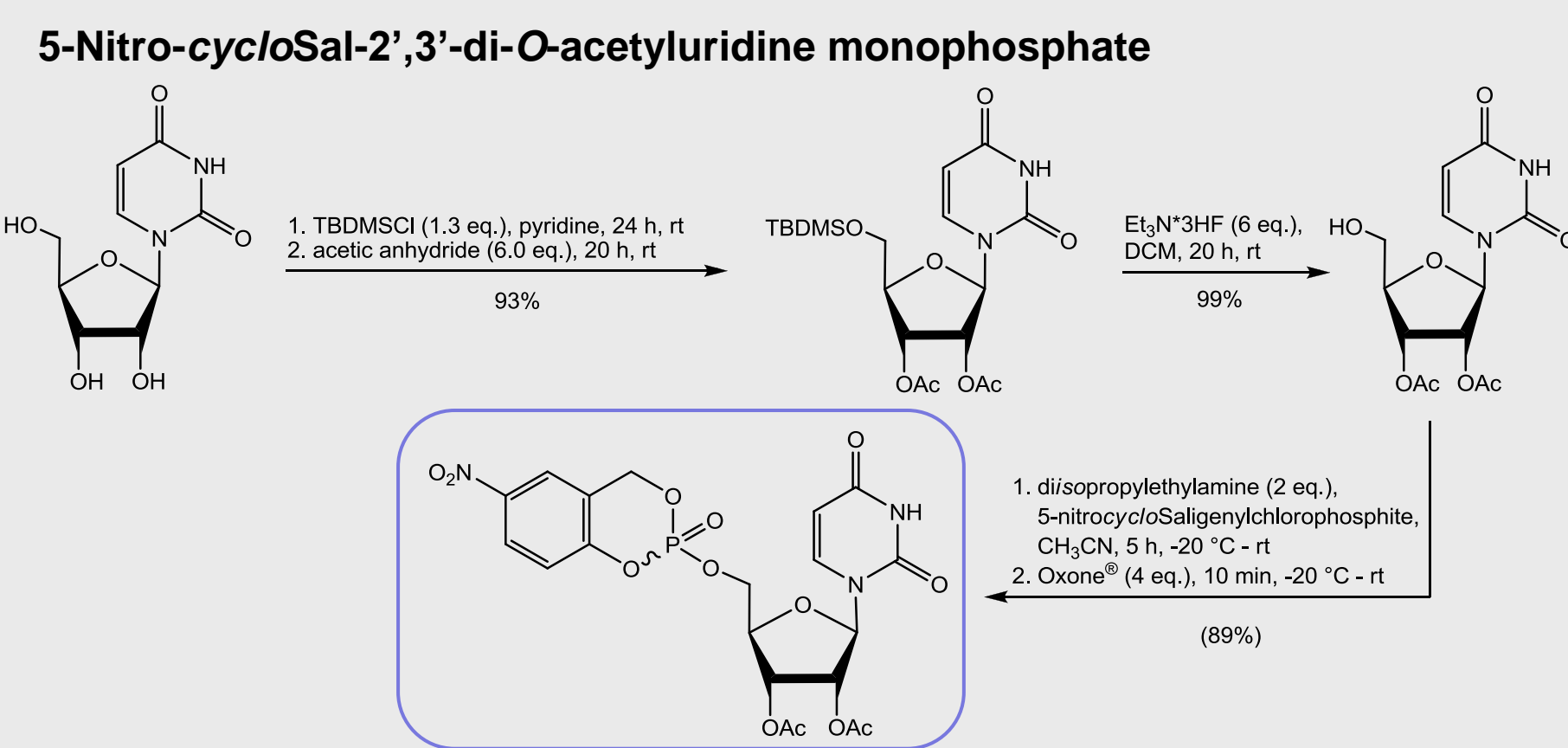
31P-NMR



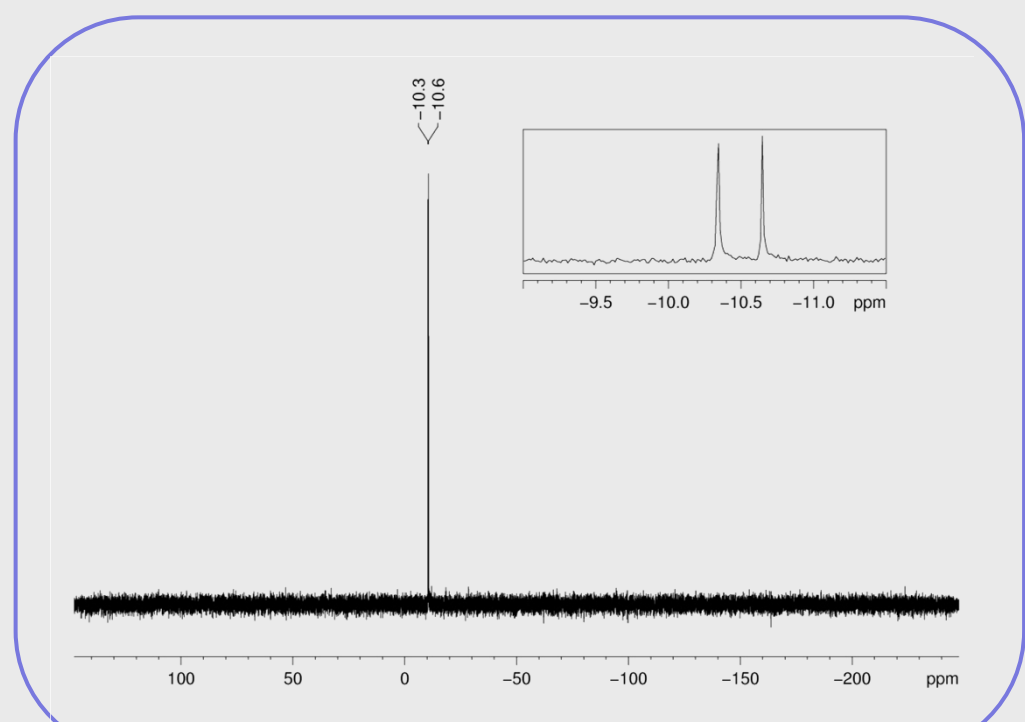
1H-NMR



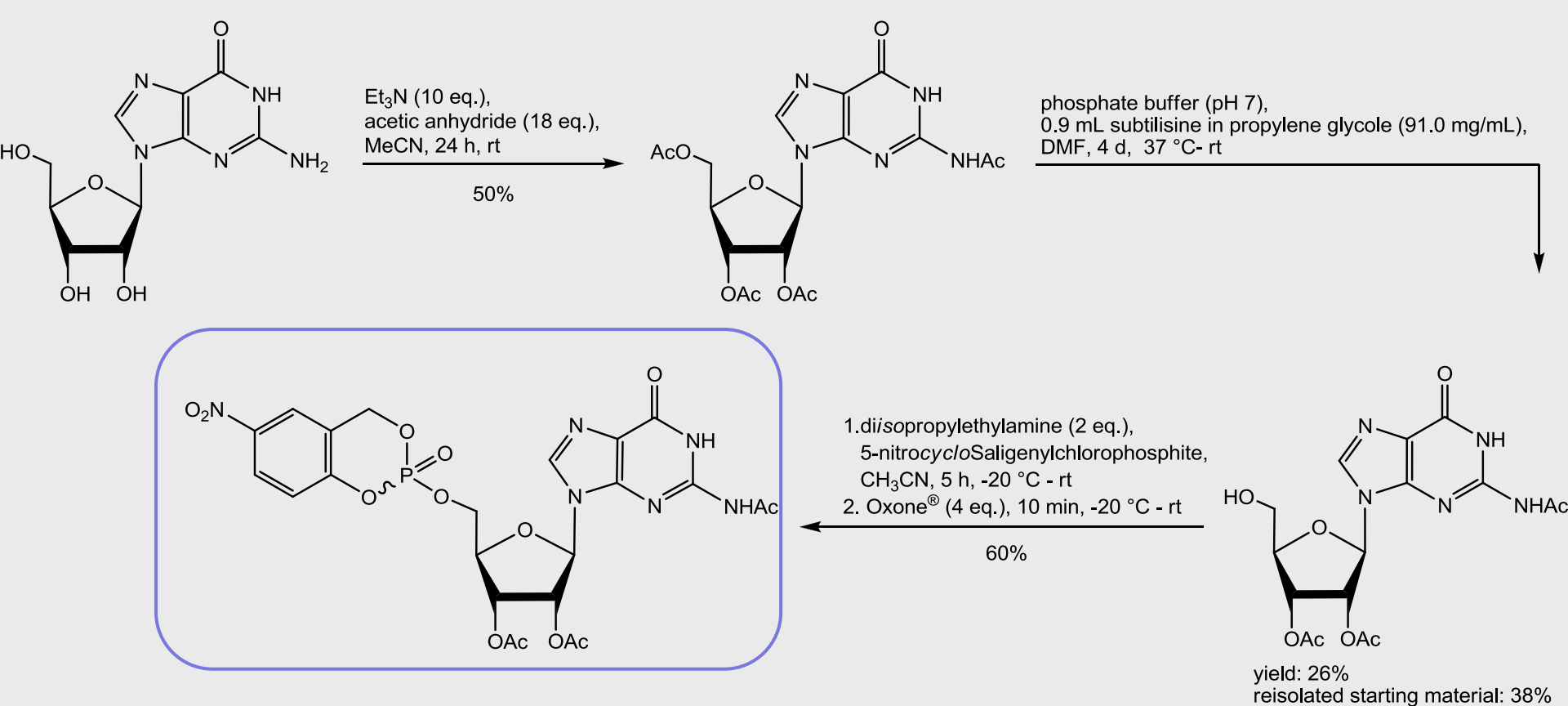
Synthesis of the cycloSal-triesters



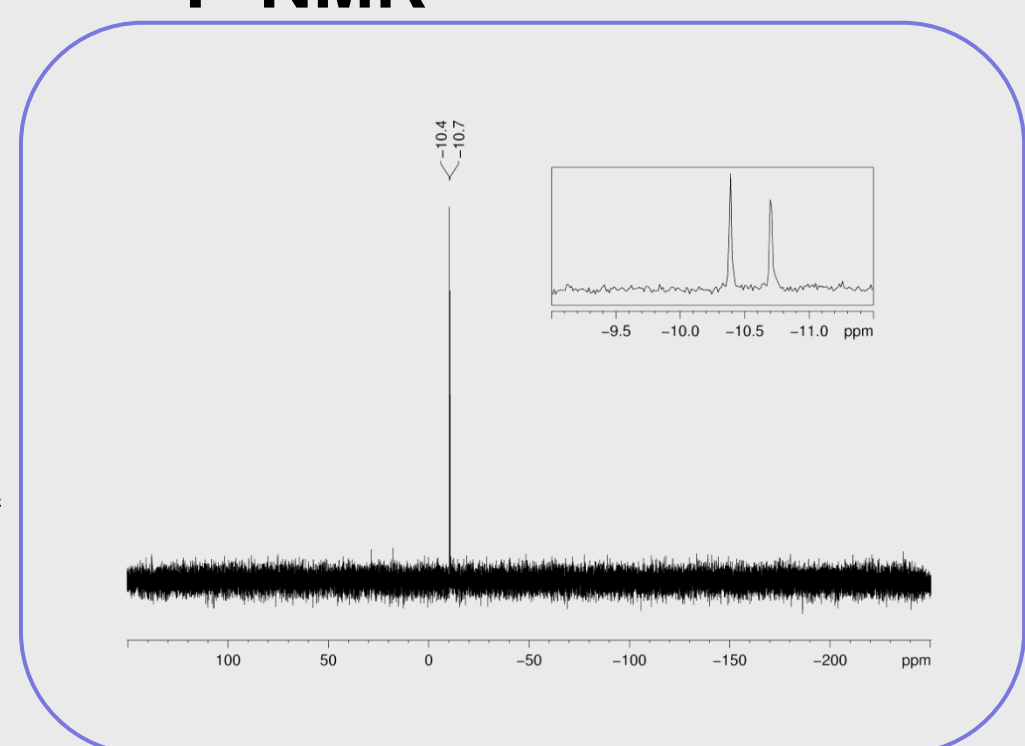
31P-NMR



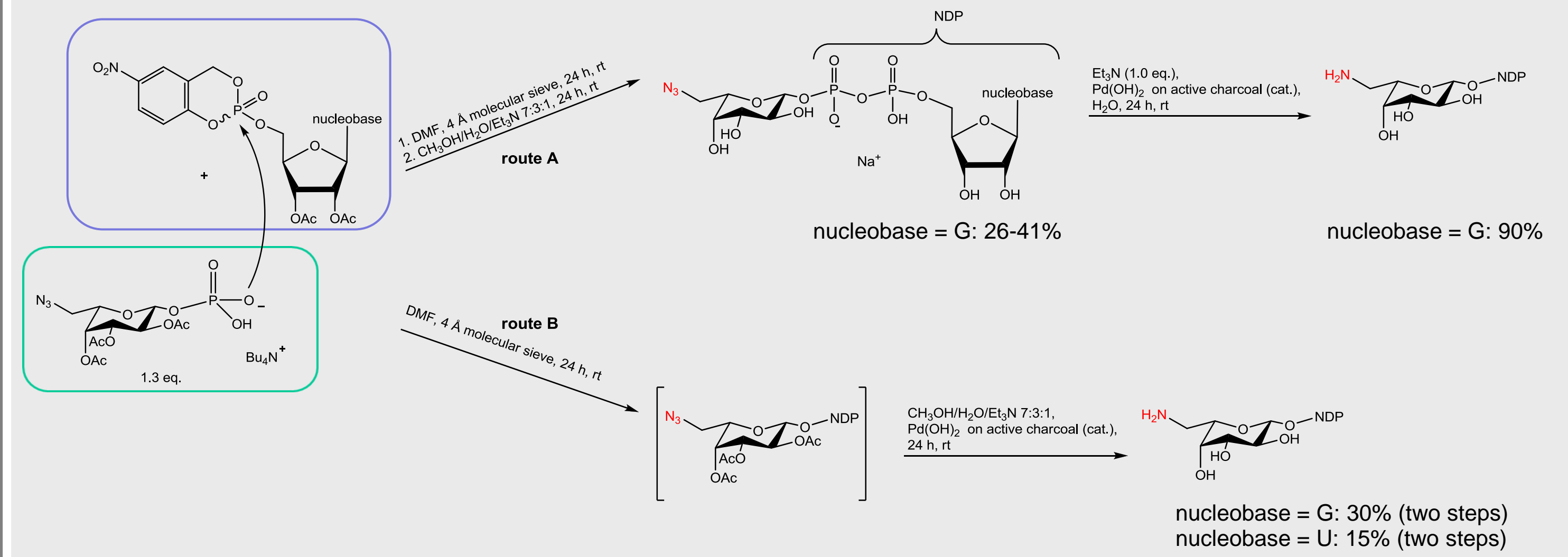
5-Nitro-cycloSal-N2-acetyl-2',3'-di-O-acetylguanosine monophosphate



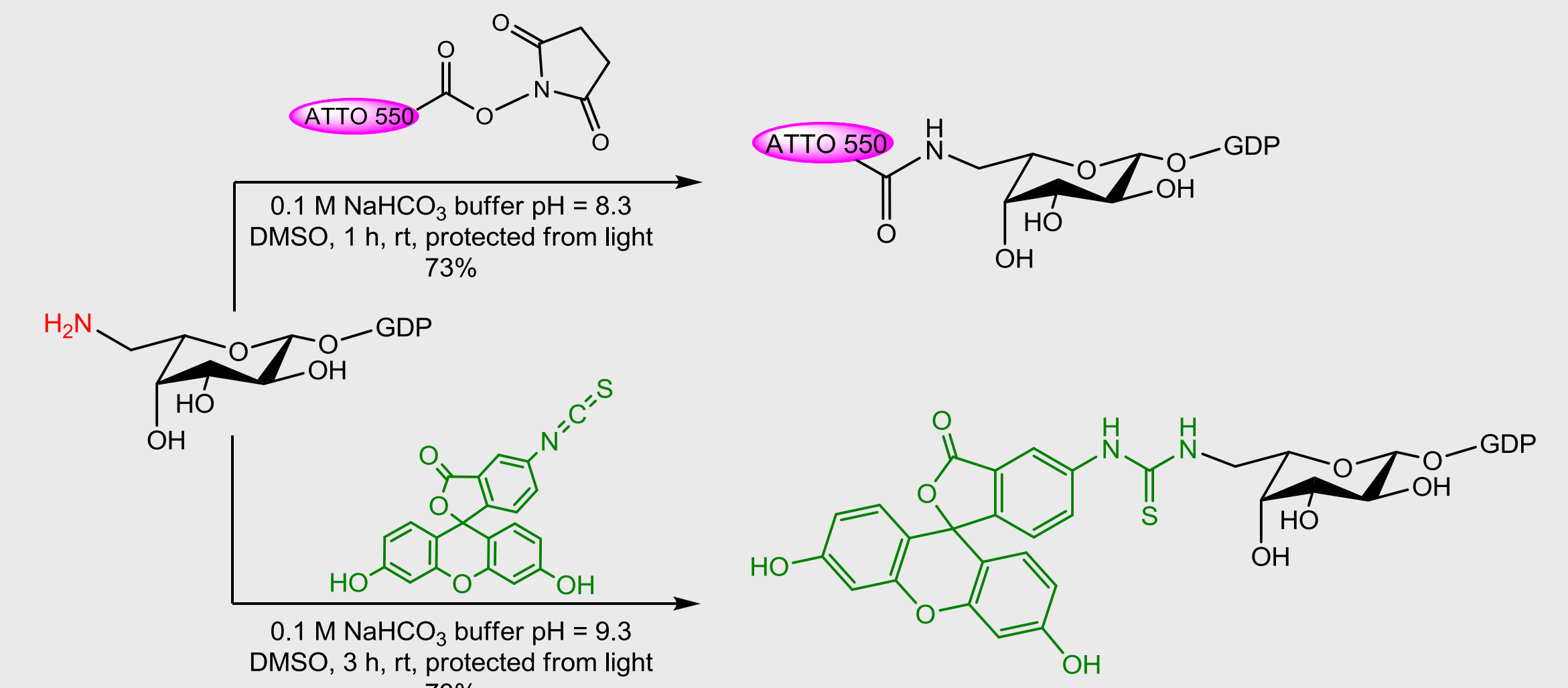
31P-NMR



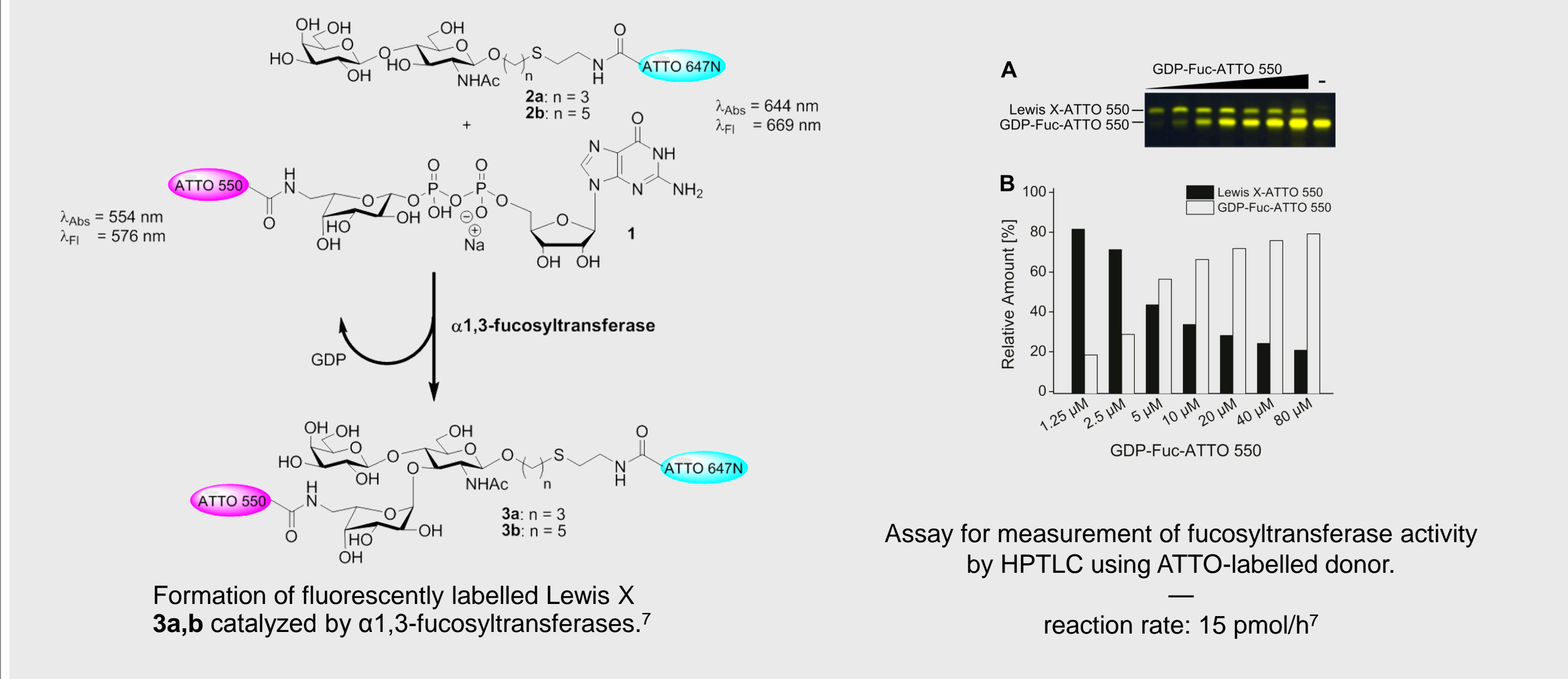
Synthesis of the modified nucleoside diphosphate sugars



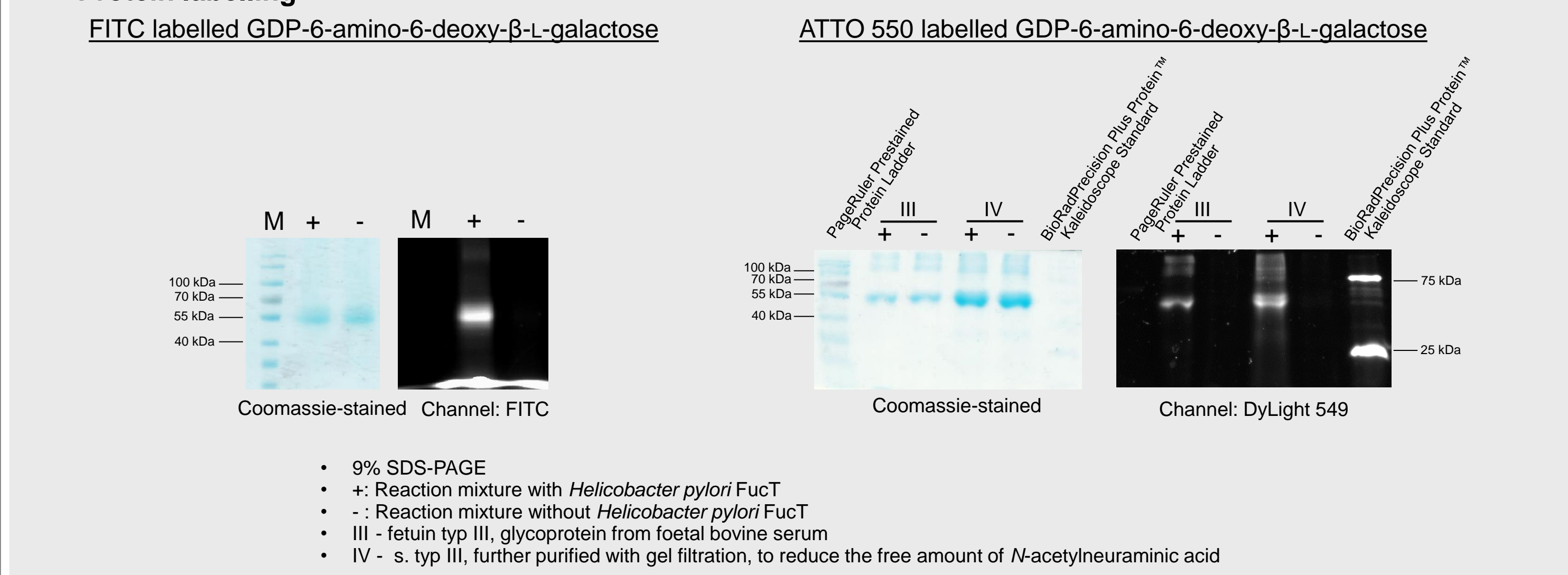
Synthesis of the fluorescently labelled nucleoside diphosphate sugars



Biochemical studies^b



Protein labelling



Conclusion

We successfully synthesised two different fluorescently labelled nucleoside diphosphate 6-amino-6-deoxy-β-L-galactoses. For the coupling to give the NDP sugars, the cycloSal-procedure was the method of choice. With the fluorescently labelled GDP sugars we were able to prove the tolerance of hFucT-IX towards substratmodifications with a dye. Moreover, the formation of fluorescently labelled Lewis X and the measurement of fucosyltransferase activity was successful.⁷ Finally, we succeeded in labelling the protein fetuin with ATTO 550 as well as with fluorescein isothiocyanate (FITC) using helicobacter pylori alpha1,3-FucT. These results open access to further biochemical assays like a high-throughput screening of fucosyltransferase IX variants using fluorescence polarisation.

References

- [1] C. Haase, O. Seitz, *Top. Curr. Chem.* **2007**, *267*, 1-36. [2] G. K. Wagner, T. Pesnot, R. A. Field, *Nat. Prod. Rep.* **2009**, *26*, 1172-1194. [3] T. Maeda, S.-I. Nishimura, *Chem. Eur. J.* **2008**, *14*, 478-487. [4] S. Wolf, R. M. Berrio, C. Meier, *Eur. J. Org. Chem.* **2011**, 6304-6313. [5] S. Wolf, T. Zismann, N. Lunau, S. Warnecke, S. Wendicke, C. Meier, *Eur. J. Cell Biol.* **2010**, *89*, 63-75; S. Wolf, T. Zismann, N. Lunau, C. Meier, *Chem. Eur. J.* **2009**, *15*, 7656-7664. [6] S. Wendicke, S. Warnecke, C. Meier, *Angew. Chem. Int. Ed.* **2008**, *47*, 1500-1502. [7] N. Lunau, K. Seelhorst, S. Kahl, K. Tschersch, C. Stacke, S. Rohn, J. Thiem, U. Hahn, C. Meier, *Chem. Eur. J.*, submitted.

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