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Chemoenzymatic synthesis of sialooligosaccharides on arrays for studies of cell surface adhesion†‡

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Sialooligosaccharides were generated by direct enzymatic glycosylation on arrays and the resulting surfaces were suitable for the study of carbohydrate-specific cell adhesion.

Sialic acid conjugates play a dominant role in many cellular recognition processes in health and disease. They are frequently exploited by viruses and bacteria as sites of attachment to cells and extracellular proteins and they can function as recognition determinants for host cell lectins, including the selectin and siglec families involved in inflammatory and immune responses.¹ Pioneering work over the last ten years has established glycan arrays as central tools to understanding many diverse carbohydrate–protein interactions in cells and organisms.^{2,3} However, both the complex preparation of oligosaccharide arrays and the lengthy biochemical generation of the labelled proteins have precluded more genome wide investigations of sialic acid binding proteins beyond first examples. Here we present some dramatic shortcuts to the present methods of analysis with three key components to our overall strategy: Firstly, the complex sialooligosaccharides are generated directly on the array using chemoenzymatic synthesis. Secondly, the efficiency of synthesis is monitored *in situ* using MALDI-ToF mass spectrometry. Thirdly, the sialooligosaccharides are directly interrogated by recombinant mammalian cells expressing siglecs on their surfaces.

The overall strategy is shown in Fig. 1. Self-assembled monolayers (SAMs) of alkanethiols on gold surfaces provide well established platforms for the study of enzymatic reactions and protein–ligand and cell–ligand interactions^{4–7} and are ideally suited for our studies.⁸ We have previously shown how glycan arrays on SAM-gold can be used to study the specificity of glycosyltransferases by using MALDI-ToF MS⁹ as a label free read-out method. Importantly, we have been able to show that the reactions can be quantitative, thus

allowing for the direct enzymatic synthesis and analysis of saccharides on array surfaces, without the need for multistep solution phase synthesis and subsequent linkage to array.^{6,7}

A concern for the synthesis of sialooligosaccharides was the use of harsh MALDI techniques employed in the analysis of enzymatic reactions on the arrays, given the known sensitivity of sialosides to this technique.¹⁰ Indeed, when we first attempted the enzymatic transfer of sialic acid from fetuin onto immobilised lactose **4**¹¹ (Fig. 2) using *trans*-sialidase from *Trypanosoma cruzi* (TcTS)¹² analysis by previously published^{6,7} MALDI-ToF MS analysis failed. Only peaks corresponding to the starting material **4** were seen in the MS spectra which could have been due to lack of enzymatic activity or cleavage of labile sialic acid groups.^{10,13}

The degradation of sialosides during mass spectrometric analysis was further investigated with immobilised 2-aminoethyl and 3-aminopropyl glycoside of Neu5Ac (**1** and **2** respectively) together with methyl ester **3**.¹¹

We first focussed on optimising the MALDI matrix and found 2,4,6-trihydroxyacetophenone (THAP) to be best in our hands, although significant degradation was still observed for **1** (See supplementary information, Fig. S1). Interestingly, with the propyl compound **2** only a very minor and with the ester **3** no fragmentation was observed during the MS experiments (Fig. S2, S3 in supporting information), indicating that the linker plays a significant role in stability and that methylation increases stability of the sialoside.



Fig. 1 Overall strategy for the chemoenzymatic synthesis and cell-based detection of sialooligosaccharides.

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Fig. 4 Binding of CFSE-labelled CHO cells expressing sialoadhesin to arrays displaying 3'-sialyllactose (top panel) and lactose (bottom panel) on gold at three surface dilutions.

by mass spectrometry provides an additional and essential step of quality control to confirm composition of the array surface. We have also shown that the SAM-gold platform is an ideal surface for studying cellular adhesion through protein-carbohydrate interactions with low background observed and should provide a useful tool for the study of recombinant cell-surface receptors of glycans and other ligands.

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