



Cite this: *RSC Adv.*, 2019, 9, 40263

# Monitoring glycosidase activity for clustered sugar substrates, a study on $\beta$ -glucuronidase†

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Determination of glycosidase hydrolysis kinetics for a monovalent sugar substrate is relatively straightforward and classically achieved by monitoring the fluorescence signal released from the sugar-conjugated probe after enzymatic hydrolysis. Naturally occurring sugar epitopes are, however, often clustered on biopolymers or at biological surfaces, and previous reports have shown that glycosidase hydrolytic rates can differ greatly with multivalent presentation of the sugar epitopes. New probes are needed to make it easier to interpret the importance of substrate clustering towards a specific enzyme activity. In this work, we developed multivalent glucuronide substrates attached to fluorescent amino-coumarines through self-immolative linkers to enable real time-monitoring of the hydrolysing activity of *E. coli*  $\beta$ -glucuronidases (GUS) towards clustered substrates. GUS are exoglycosidases of considerable therapeutic interest cleaving  $\beta$ -D-glucuronides and are found in the lysosomes, in the tumoral microenvironment, and are expressed by gut microbiota. GUS showed a much lower catalytic efficiency in hydrolysing clustered glucuronides due to a significantly lower enzymatic velocity and affinity for the substrates. GUS was 52-fold less efficient in hydrolysing GlcA substrates presented on an octameric silsequioxane (COSS) compared with a monovalent GlcA of similar chemical structure. Thus, kinetic and thermodynamic data of GUS hydrolysis towards multivalent glucuronides were easily obtained with these new types of enzymatically-triggered probes. More generally, adapting the substrate nature and valency of these new probes, should improve understanding of the impact of multivalency for a specific enzyme.

Received 28th October 2019  
 Accepted 18th November 2019

DOI: 10.1039/c9ra08847d

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## Introduction

Glycoclusters bearing several copies of a sugar ligand on a chemical or biological core have been extensively developed to target carbohydrate-binding-proteins (lectins).<sup>1–3</sup> In numerous reports, the tethered ligands are shown to provide synergistic affinity enhancement compared with their corresponding monovalent references. Large multivalent effects with glycoclusters have been reported on a wide range of lectins, which are often multimeric in nature or multivalently displayed at a cell surface. The enhanced selectivity and avidity is generally explained by an increased local concentration (effective

molarity) of the additional ligands close to the binding sites and in theoretical models by a statistical component (degeneracy prefactor) growing non-linearly with increased multivalency of both ligands and receptors.<sup>4</sup>

Although, mostly exemplified on carbohydrate-binding proteins, multivalency effects have also been demonstrated on carbohydrate-processing enzymes (glycosidases and glycosyltransferases). Indeed, these enzymes were previously shown to display very different hydrolytic profiles towards multimerized substrates compared with monovalent analogues. *N*-acetylglucosamine-capped glycolipids can form microdomains in phospholipid bilayers. Webb and co-workers showed an accelerated galactosylation of these clustered sugars by bovine  $\beta(1,4)$ -galactosyltransferase. This was explained by a lower apparent  $K_M$  attributed to multivalent binding.<sup>5</sup> Boons and co-workers observed that bacterial sialidases flanked by a lectin domain hydrolyse polyvalent sialosides with much higher catalytic efficiency than their monovalent counterparts.<sup>6</sup> In stark contrast, *O*-glucosides and *O*-mannosides multivalently displayed on nanodiamond particles were shown to be fully stable and even to inhibit their matching enzymes (glycosidases and mannosidases, respectively).<sup>7</sup>

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† Electronic supplementary information (ESI) available. See DOI: 10.1039/c9ra08847d





by the liver for passivation but GUS from the microbiome cleaves the sugar in the gastrointestinal tract, restoring the initial drug toxicity.<sup>28</sup>

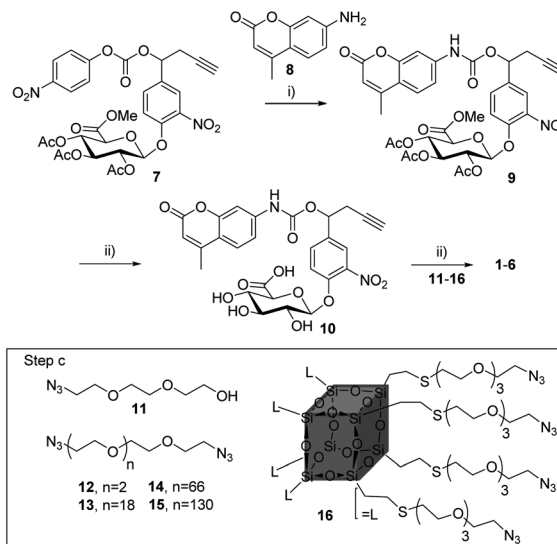
We designed mono-, di- and octavalent FluoGlcA (Fig. 2) bearing a glucuronide trigger (in red) which, upon GUS cleavage uncap a self-immolative linker (in black) releasing an amino-coumarin dye (in blue,  $\lambda_{\text{ex}} = 380 \text{ nm}$ ,  $\lambda_{\text{em}} = 450 \text{ nm}$ ). The self-immolative linker presented Fig. 2 has been selected due to its widespread use in the design of enzyme-responsive prodrugs.<sup>29–31</sup> Such aromatic linkers bearing a nitro group allow a fast liberation kinetics of the drugs.<sup>32</sup>

Amino-coumarin functionalized by a propargyl-oxo-carbonyl group was previously shown to be virtually non-fluorescent but was converted into a highly fluorescent amino-coumarin after palladium-triggered depropargylation.<sup>33</sup> Here, this responsive fluorescence is exploited for the kinetic measurement of GUS activity. Divalent p-FluoGlcA with varied PEG spacer arm length were designed to assess whether the spanning of two binding sites in GUS by the GlcA leads to a specific hydrolytic profile. Molecular dynamics experiments were performed to assess the average distance between the glucuronides ligands of compounds 2–5, and we found distances ranging from 8 to 59 Å (Chart 1). This is in accordance with previous calculations from polymer theory performed with a different set of compounds bearing PEG linkers of similar lengths.<sup>17,34</sup> Compound 5 should therefore cover the distance of around 50 Å between two adjacent GUS binding sites, as determined from the crystal structure of *E. coli* GUS bound to an urea inhibitor (PDB code = 3LPG).<sup>35</sup>

Compound 6 was designed from a cubic octameric silsesquioxane (COSS) scaffold. COSS are emerging as versatile core for the design of multivalent sugar ligands,<sup>36–38</sup> the cubic geometry allowing an homogeneous spatial distribution of the grafted ligands.

## Chemical synthesis

Compounds 1–6 (Scheme 1) were readily accessible in only three synthetic steps from the glucuronide 7<sup>29,39</sup> (Synthesis in ESI†).



Scheme 1 Synthesis of compounds 1–6. (i) 8, HOBT, DMF, (*i*-Pr)<sub>2</sub>NEt, 50 °C, 24 h, 38%; (ii) LiOH, H<sub>2</sub>O/MeOH, 0 °C, 2 h, 58%; (iii) CuSO<sub>4</sub>, NaAsc, dioxane–water, r.t., 12 h, with 11 for 1 (95%), 12 for 2 (78%), 13 for 3 (67%), 14 for 4 (61%), 15 for 5 (62%), 16 for 6 (58%).

First, the 7-amino-4-methyl-2H-chromen-2-one 8 was introduced *via* nucleophilic substitution in the presence of hydroxybenzotriazole (HOBT) and (*i*-Pr)<sub>2</sub>NEt to give the carbamate 9 (38%). Full deprotection of the glucuronide moiety was then carried out using LiOH, giving the alkyne 10 with 58% yield after purification by reverse phase column chromatography. Finally, triazoles 1–6 were obtained by coupling 10 with the corresponding azide derivatives 11, using the well-known copper(I)-catalysed azide–alkyne 1,3-cycloaddition.<sup>40–43</sup> After size-exclusion column purification, 1–6 were obtained with yields ranging from 58 to 95%.

## Enzymatic assays

Compounds 1–6 were assessed against the commercially available  $\beta$ -glucuronidase from *E. coli*. *E. coli* GUS shows a low sequence similarity with human GUS (45%) but both enzymes are tetravalent with a conserved three-dimensional fold.<sup>35</sup> The fluorescence signal was directly monitored during the reaction and converted into concentration by a calibration curve. The Michaelis constant  $K_m$  and the maximal velocity  $V_{\text{max}}$  values were determined from the Michaelis–Menten equation at a GUS concentration of 9.6 nM. The  $K_m$ ,  $V_{\text{max}}$  and catalytic efficiency ( $C_{\text{eff}} = k_{\text{cat}}/K_m$ ) are presented in Fig. 3 together with the fitted curves of GUS hydrolysis velocity as a function of the molecule concentration of 1–6 (average from quadruplicates). To improve comparison of the enzymatic efficiency per mol of GlcA ligand (instead of molecule) we calculated the  $R_{(C_{\text{eff}})}$  representing the fold-decrease of GUS catalytic efficiency per GlcA ligand of a multivalent compound X of valency  $V_X$ , compared with the GlcA ligand of monovalent reference 1.  $R_{(C_{\text{eff}})}$  was obtained from the equation  $R_{(C_{\text{eff}})} = V_X \times (k_{\text{cat}1}/K_{m1}) / (k_{\text{cat}X}/K_{mX})$ . GUS showed an apparent decreased affinity (higher  $K_m$ ) and lower hydrolysis

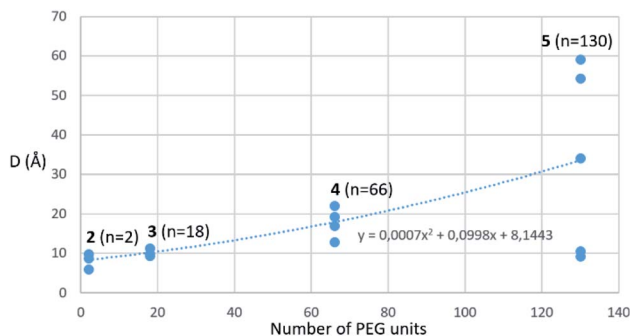


Chart 1 Average distance  $d$  between the glucuronide ligands of compounds 2–5 plotted against the PEG number of the compounds. The distances were extracted from several 20 ns long, independent molecular dynamics simulations (for more details see Sup. Mat.) A nonlinear least squares regression using a power fitting function was used to fit the data points and is shown as a dotted line in the graph.



Cpd	$K_M$ ( $\mu\text{M}$ )	$V_{\text{max}}$ ( $\mu\text{M}\cdot\text{min}^{-1}$ )	$k_{\text{cat}}/K_M$ ( $\text{s}^{-1}\cdot\mu\text{M}^{-1}$ )	$R_{(\text{C}_{\text{eff}})}$
1 <sup>l</sup>	$4.6 \pm 0.9$	$1.03 \pm 0.04$	0.39	1
2	$17.7 \pm 3.0$	$0.84 \pm 0.05$	0.08	10
3	$8.1 \pm 0.6$	$0.83 \pm 0.02$	0.18	4
4	$14.4 \pm 1.5$	$0.68 \pm 0.02$	0.08	10
5	$18.5 \pm 2.6$	$0.51 \pm 0.02$	0.05	16
6	$11.5 \pm 2.0$	$0.38 \pm 0.02$	0.06	52

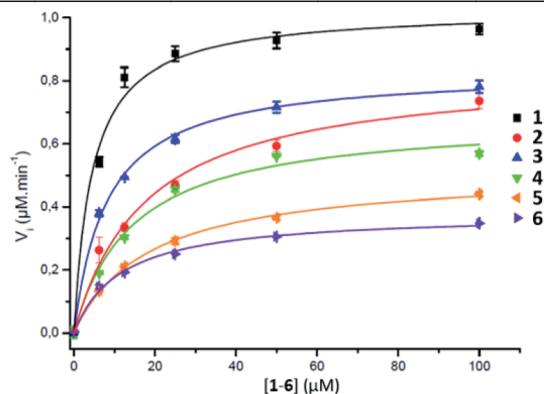


Fig. 3 Kinetic parameters obtained from the Michaelis–Menten equation on GUS activity with compounds 1–6.

rate (lower  $V_m$  or  $k_{\text{cat}}$ ) for the multivalent GlcA 2–6 than monovalent 1. It follows that the relative hydrolytic catalytic efficiency  $R_{(\text{C}_{\text{eff}})}$  for a GlcA ligand was reduced by 4 to 52-fold when multimerized on scaffolds 2–6. This significant decrease may not be explained by different interactions between 1 and 2–6 in close binding site proximity because the GlcA environment is the same independently of the substrate valency.

The PEG length elongation from  $n = 2$  to  $n = 130$  for the divalent serie 2–5 also affected  $R_{(\text{C}_{\text{eff}})}$ , suggesting that the scaffold and the GlcA–GlcA average distance play a role in the reduced  $C_{\text{eff}}$ . This scenario was enhanced with COSS 6 where the GlcA are also tethered through EG spacer arms. Despite an eight fold higher GlcA concentration due to the octavalency, the GUS  $C_{\text{eff}}$  of 6 was only  $0.06 \mu\text{mol min}^{-1}$ , each of the GlcA being hydrolysed with 52-fold lower efficiency compared with 1. Although it is difficult at present to hypothesize about a specific binding mode, the data clearly show that a multivalent display of the sugar substrates strongly affects GUS activity.

## Conclusions

In summary we developed multivalent glucuronides with self-immolative linkers releasing a fluorescent amino-coumarin after GUS activation. The multivalent probes allowed easy monitoring of the kinetic parameters for GUS enzymatic activity. GUS was shown to hydrolyze the clustered glucuronide probes with much less efficiency than the monovalent glucuronide analogues. The scope of the multivalent probes developed here could potentially be broadened to any glycosidase by adapting the sugar nature and valency. This may help to improve understanding the intriguing function of multivalency in carbohydrate processing.

## Conflicts of interest

There are no conflicts to declare.

## Acknowledgements

This work was carried out with financial support from the Conseil Régional des Pays de La Loire (MAGGIC project), the Centre National de la Recherche Scientifique (CNRS) and the French Ministère de l'Enseignement Supérieur et de la Recherche. E. M. K. is a postdoctoral researcher of the Fonds de la Recherche Scientifique de Belgique (F.R.S.-F.R.N.S.), Belgium.

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