

## A Strategy for the Selective Imaging of Glycans Using Caged Metabolic Precursors

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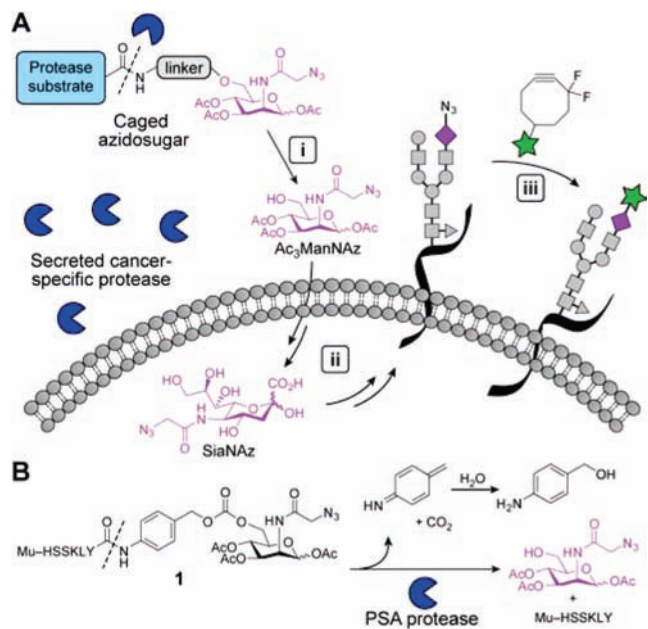
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**Abstract:** Glycans can be imaged by metabolic labeling with azidosugars followed by chemical reaction with imaging probes; however, tissue-specific labeling is difficult to achieve. Here we describe a strategy for the use of a caged metabolic precursor that is activated for cellular metabolism by enzymatic cleavage. An *N*-azidoacetylmannosamine derivative caged with a peptide substrate for the prostate-specific antigen (PSA) protease was converted to cell-surface azido sialic acids in a PSA-dependent manner. The approach has applications in tissue-selective imaging of glycans for clinical and basic research purposes.

A fundamental challenge in molecular imaging is the design of reagents that report on cell- or tissue-specific molecular processes. A popular approach to achieving such selectivity is the design of caged probes that are activated by enzymes produced by the target cells, a prototypical example being cancer-associated proteases.<sup>1</sup> Alternatively, probes can be conjugated to targeting elements that bind, noncovalently or covalently, to cell-surface markers.<sup>2</sup> We have been exploring this latter approach in the context of imaging cell-surface glycans. These biopolymers, collectively termed the glycome, change in structure and expression during development and cancer progression as well as many other physiological processes.<sup>3</sup> We previously reported that broad sectors of the glycome can be imaged in model organisms using a two-step procedure: (1) metabolic labeling of the glycans with azidosugar precursors and (2) chemical reaction of azide-labeled cell-surface glycans *in vivo* via Cu-free click chemistry with difluorinated cyclooctyne (DIFO) probes.<sup>4</sup>

The tissue selectivity of this approach is limited by the breadth of glycans targeted by a single metabolic precursor. For example, treatment of mice with peracetylated *N*-azidoacetylmannosamine (Ac<sub>4</sub>ManNAz) leads to widespread labeling of glycans with *N*-azidoacetyl sialic acid (SiaNAz) in numerous tissues (e.g., liver, heart, kidney, and intestines) as well as serum glycoproteins.<sup>5</sup> The broad distribution of the azidosugar can undermine experiments that seek to probe glycomic changes in a single organ as a function of time or disease. One means to achieve tissue selectivity in glycan imaging is to restrict azidosugar metabolism to the cells of interest. Tissue-specific enzyme activities such as those mentioned above might be exploited to achieve this goal. Here, we describe a strategy for cell-selective glycan imaging using a caged metabolic precursor that is activated by a protease (Figure 1A).

We synthesized a variant of Ac<sub>4</sub>ManNAz in which the 6-hydroxy group was conjugated through a self-immolating linker to a peptide substrate for the prostate-specific antigen (PSA) protease (**1**, Figure 1B). PSA is a serine protease that is secreted at low levels by normal prostatic glandular cells but is highly upregulated by prostate cancer

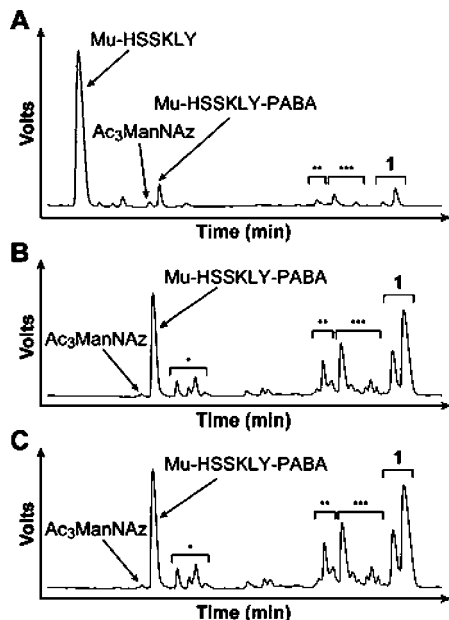


**Figure 1.** Strategy for tissue-specific release of Ac<sub>3</sub>ManNAz via enzymatic activation. (A) (i) A nonmetabolizable caged azidosugar serves as a substrate for a secreted, cancer-specific protease, releasing Ac<sub>3</sub>ManNAz. (ii) This azidosugar is then metabolized by the cell and incorporated into cell-surface glycans. (iii) The azide-labeled glycans are detected via Cu-free click chemistry using DIFO reagents. (B) Caged azidosugar used in this study (**1**). Cleavage of the indicated bond (dashed line) by the prostate-specific antigen (PSA) protease results in the release of a linker, the peptide, and Ac<sub>3</sub>ManNAz.

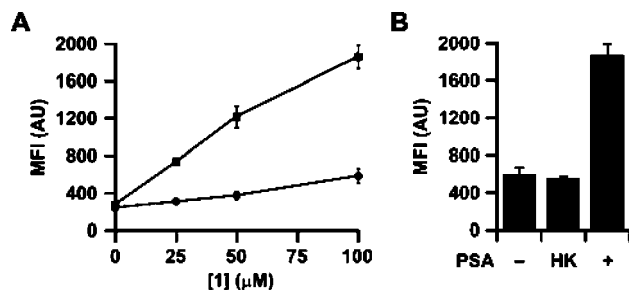
cells.<sup>6</sup> The enzyme has been widely targeted with caged drugs and imaging reagents in cell culture and in animal models.<sup>6</sup> The hexapeptide Mu-HSSKLY (Mu = morpholino ureidyl) was chosen as a PSA substrate based on reports of its high selectivity for this enzyme over other ubiquitous serine proteases.<sup>6c,7</sup> Upon cleavage of the peptide, the *p*-aminobenzyl alcohol (PABA) linker in **1** spontaneously fragments to release 1,3,4-tri-*O*-acetyl-*N*-azidoacetylmannosamine (Ac<sub>3</sub>ManNAz), carbon dioxide, and an iminoquinone methide intermediate that is subsequently quenched by water (Figure 1B).<sup>8</sup> This process is known to occur rapidly and should therefore limit the diffusion of the released metabolic substrate from its target cell (for discussion see Figure S1 legend). Compound **1** was synthesized as a mixture of sugar anomers analogously to the route developed by Jones et al. (Scheme S1).<sup>6c</sup> Given the importance of hydroxy group acylation for the cellular uptake of Ac<sub>4</sub>ManNAz,<sup>9</sup> we verified that Chinese hamster ovary (CHO) and prostate cancer (PC-3) cells incubated with Ac<sub>3</sub>ManNAz could produce SiaNAz residues in their cell-surface glycans (Figure S1).

To confirm that **1** can serve as a substrate for PSA, the compound was incubated *in vitro* with active enzyme or, as negative controls, buffer only or heat-killed (HK) PSA. These enzymatic reactions were analyzed by reversed-phase HPLC and mass spectrometry. Incubation

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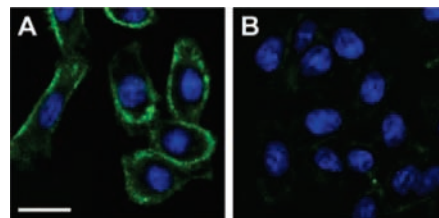
**Figure 2.** Compound **1** is a substrate for PSA *in vitro*. Shown are HPLC traces of *in vitro* 6 h enzymatic reactions of **1** (500  $\mu\text{M}$ ) in 50 mM Tris, 0.1 M NaCl, pH 7.8, with (A) active PSA (50  $\mu\text{g}/\text{mL}$ ), (B) buffer only, or (C) HK PSA (50  $\mu\text{g}/\text{mL}$ ). The identities of the various species based on mass spectrometry are indicated on the traces. \*Monoacetylated Mu-HSSKLY-PABA, \*\*Monodeacetylated **1**, and \*\*\*Isomers of **1**.



**Figure 3.** Cell-selective metabolic labeling of glycans using **1** and PSA. Flow cytometry analysis of CHO cells treated with (A) various concentrations of **1** (0–100  $\mu\text{M}$ ) and PSA (50  $\mu\text{g}/\text{mL}$ , squares) or buffer only (circles) or (B) **1** (100  $\mu\text{M}$ ) and either buffer only (–), HK PSA (50  $\mu\text{g}/\text{mL}$ , HK), or PSA (50  $\mu\text{g}/\text{mL}$ , +). Cells were then labeled with DIFO–biotin (100  $\mu\text{M}$ ) and FITC–avidin. Error bars represent the standard deviation from the mean of three replicate samples. MFI = mean fluorescence intensity in arbitrary units (AU).

of **1** with PSA resulted in the release of Mu-HSSKLY as the major peptide product, along with Ac<sub>3</sub>ManNAz (Figure 2A). Minor products were also observed. These included Mu-HSSKLY-PABA, presumably produced by nonenzymatic carbonate hydrolysis, as well as monoacetylated **1** and products formed by migration of acetyl groups within **1**. Control reactions lacking PSA (Figure 2B) or with HK PSA (Figure 2C) produced only nonenzymatic hydrolysis and acetyl migration products.

We next tested the performance of **1** as a caged substrate for metabolic glycan labeling. CHO cells were incubated with **1** at various concentrations (0–100  $\mu\text{M}$ ) in the presence of PSA, no enzyme, or HK PSA for 12 h at 37 °C. The cells were then washed and labeled with a DIFO–biotin conjugate,<sup>4b</sup> incubated with fluorescein isothiocyanate-labeled avidin (FITC–avidin), and analyzed by flow cytometry. We observed labeling that was both PSA- and substrate concentration-dependent, suggesting that the signal is due to enzymatic activation of **1** (Figure 3A). In a separate experiment, we demonstrated that the labeling intensity correlates with PSA concentration (Figure S2). Additionally, we verified that treatment of PC-3 cells with **1** resulted



**Figure 4.** Selective imaging of cells using **1** in the presence of PSA. Fluorescence microscopy analysis of CHO cells treated with **1** (100  $\mu\text{M}$ ) and (A) PSA (50  $\mu\text{g}/\text{mL}$ ) or (B) HK PSA (50  $\mu\text{g}/\text{mL}$ ), followed by DIFO–biotin (100  $\mu\text{M}$ ) and a quantum dot 605–streptavidin conjugate. Green = Texas Red channel; Blue = DAPI channel. Scale bar = 20  $\mu\text{m}$ .

in PSA-dependent metabolic labeling (Figure S3). In the absence of PSA or with HK PSA, both CHO and PC-3 cells exhibited modest background labeling that likely reflects low levels of Ac<sub>3</sub>ManNAz produced by nonenzymatic carbonate hydrolysis (Figures 3B and S3). Importantly, we verified that **1** did not cause any cytotoxicity by incubating CHO cells labeled as above with phycoerythrin-conjugated annexin V, a marker of apoptosis (Figure S4).

Finally, we tested **1** as an enzyme-activatable metabolic substrate for glycan imaging. CHO cells were incubated with **1** in the presence of PSA or HK PSA for 12 h at 37 °C. The cells were then washed and labeled with DIFO–biotin, followed by quantum-dot-conjugated streptavidin. We observed substantial cell-surface labeling of cells treated with **1** and PSA (Figure 4A) and minimal fluorescence on cells treated with **1** and HK PSA (Figure 4B).

In conclusion, we have developed a strategy for targeted metabolism of azidosugars using an enzymatically activated substrate. While we chose PSA to demonstrate proof-of-concept, it should be noted that the concentrations of PSA employed in our studies are physiologically relevant; i.e., they are similar to the levels of PSA secreted by both prostate cancer xenografts in mice and prostate tumor tissue obtained from human patients.<sup>10</sup> In addition, many cancers, including prostate cancer, are known to express elevated levels of sialic acid compared to surrounding tissue.<sup>11</sup> Thus, clinical imaging applications may be worth pursuing. More generally, however, the approach has promise for use in tissue-specific glycan imaging, a major future direction.

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**Supporting Information Available:** Synthetic procedures and additional data. This material is available free of charge via the Internet at <http://pubs.acs.org>.

## References

- (1) (a) Blum, G.; von Degenfeld, G.; Merchant, M. J.; Blau, H. M.; Bogoy, M. *Nat. Chem. Biol.* **2007**, *3*, 668–677. (b) Jiang, T.; Olson, E. S.; Nguyen, Q. T.; Roy, M.; Jennings, P. A.; Tsien, R. Y. *Proc. Natl. Acad. Sci. U.S.A.* **2004**, *101*, 17867–17872. (c) Weissleder, R.; Tung, C. H.; Mahmood, U.; Bogdanov, A., Jr. *Nat. Biotechnol.* **1999**, *17*, 375–378.
- (2) Massoud, T. F.; Gambhir, S. S. *Genes Dev.* **2003**, *17*, 545–580.
- (3) Varki, A.; Cummings, R. D.; Esko, J. D.; Freeze, H. H.; Stanley, P.; Bertozzi, C. R.; Hart, G. W.; Etzler, M. E. *Essentials of Glycobiology*, 2nd ed.; Cold Spring Harbor Laboratory Press: New York, 2008.
- (4) (a) Sletten, E. M.; Bertozzi, C. R. *Angew. Chem., Int. Ed.* **2009**, *48*, 6974–6998. (b) Baskin, J. M.; Prescher, J. A.; Laughlin, S. T.; Agard, N. J.; Chang, P. V.; Miller, I. A.; Lo, A.; Codelli, J. A.; Bertozzi, C. R. *Proc. Natl. Acad. Sci. U.S.A.* **2007**, *104*, 16793–16797.
- (5) (a) Chang, P. V.; Prescher, J. A.; Sletten, E. M.; Baskin, J. M.; Miller, I. A.; Agard, N. J.; Lo, A.; Bertozzi, C. R. *Proc. Natl. Acad. Sci. U.S.A.* **2010**, *107*, 1821–1826. (b) Prescher, J. A.; Dube, D. H.; Bertozzi, C. R. *Nature* **2004**, *430*, 873–877.
- (6) (a) Denmeade, S. R.; Nagy, A.; Gao, J.; Lilja, H.; Schally, A. V.; Isaacs, J. T. *Cancer Res.* **1998**, *58*, 2537–2540. (b) Khan, S. R.; Denmeade, S. R. *Prostate* **2000**, *45*, 80–83. (c) Jones, G. B.; Crasto, C. F.; Mathews, J. E.;

- Xie, L.; Mitchell, M. O.; El-Shafey, A.; D'Amico, A. V.; Bubley, G. J. *Bioorg. Med. Chem.* **2006**, *14*, 418–425.
- (7) Denmeade, S. R.; Lou, W.; Lovgren, J.; Malm, J.; Lilja, H.; Isaacs, J. T. *Cancer Res.* **1997**, *57*, 4924–4930.
- (8) de Groot, F. M.; Damen, E. W.; Scheeren, H. W. *Curr. Med. Chem.* **2001**, *8*, 1093–1122.
- (9) Jacobs, C. L.; Yarema, K. J.; Mahal, L. K.; Nauman, D. A.; Charters, N. W.; Bertozzi, C. R. *Methods Enzymol.* **2000**, *327*, 260–275.
- (10) Denmeade, S. R.; Sokoll, L. J.; Chan, D. W.; Khan, S. R.; Isaacs, J. T. *Prostate* **2001**, *48*, 1–6.
- (11) (a) Zhang, S.; Cordon-Cardo, C.; Zhang, H. S.; Reuter, V. E.; Adluri, S.; Hamilton, W. B.; Lloyd, K. O.; Livingston, P. O. *Int. J. Cancer* **1997**, *73*, 42–49. (b) Zhang, S.; Zhang, H. S.; Cordon-Cardo, C.; Reuter, V. E.; Singhal, A. K.; Lloyd, K. O.; Livingston, P. O. *Int. J. Cancer* **1997**, *73*, 50–56.

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