

## Multivalent Carbohydrate Ligands Assembled on a Metal Template

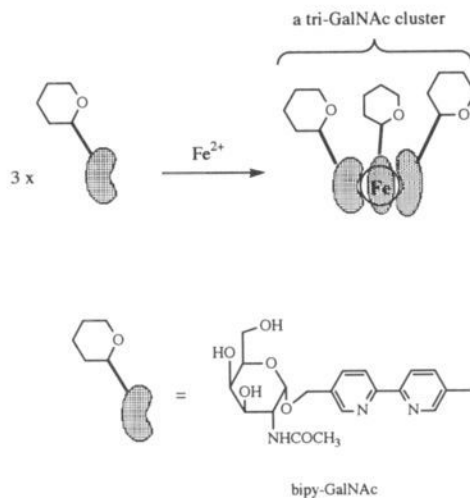
Shin Sakai† and Tomikazu Sasaki\*

Department of Chemistry, BG-10  
University of Washington  
Seattle, Washington 98195

Received November 8, 1993

Multiple saccharide units are often involved in molecular recognition of biologically active carbohydrates. A simple mucin-type carbohydrate antigen, Tn, contains 2-acetamido-2-deoxy- $\alpha$ -D-galactopyranosyl serine or threonine residues and is reportedly a good marker of malignant transformation in several epithelial tissues.<sup>1,2</sup> Recent binding studies<sup>3,4</sup> of GalNAc-modified peptides with Tn-specific antibodies showed that repeating units of at least two GalNAc-Ser or -Thr residues were necessary for the specific antigen-antibody interactions. Tn-specific lectins, *Vicia villosa* B4 lectin<sup>5,6</sup> and *Salvia sclarea* lectin,<sup>7</sup> also appear to recognize multiple GalNAc residues in their binding to the antigen. Simple synthetic carbohydrates that mimic native structures of the cancer-specific antigens would be extremely useful in diagnostic and therapeutic applications.<sup>8</sup> The synthesis of native carbohydrate clusters, however, usually requires a tedious multistep synthetic strategy.<sup>9–13</sup> We have for the past several years been interested in self-assembly of biomolecules on a metal template.<sup>14–16</sup> Carbohydrate clusters could be assembled by the metal-assisted association of carbohydrate components modified with a metal-binding ligand. We report here the synthesis of bipyridine-modified GalNAc and its metal-assisted association to form tridentate GalNAc clusters (Figure 1).

Bipy-GalNAc<sup>17,18,19</sup> trimerizes to give a tridentate GalNAc cluster in the presence of Fe(II). Four isosbestic points (227, 250, 268, and 296 nm) were observed during the titration, consistent with the large third binding constant of Fe(bipy)<sub>3</sub> complexes relative to the first and second ones.<sup>16</sup> The stoichiometry of the complex formation was determined to be 3.2:1 for



**Figure 1.** Schematic representation of the assembly of a tri-GalNAc cluster by metal-assisted self-association of bipy-GalNAc.

bipy-GalNAc:Fe(II). The absorption spectrum of the red iron complex (Fe<sup>II</sup>(bipy-GalNAc)<sub>3</sub>,  $\lambda_{\max}$  = 303, 351, 481 (sh), and 516 nm) is comparable to that of the iron(II) tris(5,5'-dimethylbipyridine) complex [Fe(Me<sub>2</sub>bipy)<sub>3</sub>]<sup>2+</sup> ( $\lambda_{\max}$  = 296, 348, 480 (sh), and 519 nm). Since the bipyridine moiety of bipy-GalNAc is unsymmetrical, four stereoisomers, *fac*- $\Delta$ , *fac*- $\Delta$ , *mer*- $\Delta$ , and *mer*- $\Delta$ , are possible at the Fe(II) center.<sup>16</sup> The structural differences between the *fac* and *mer* templates are significant; for example, in the *fac* isomers, the pyridyl carbons that are attached to the glycosidic oxygen of the GalNAc unit are positioned 7 Å apart, while in the *mer* isomers, two GalNAcs are 11 Å distant from one another and each is 7 Å away from the third GalNAc. Each GalNAc cluster would, therefore, exhibit considerably different affinity to a lectin or antibody that recognizes multiple GalNAc residues. All four isomers were separated by reverse-phase HPLC, using a linear gradient of acetonitrile and water containing 0.1% TFA on a Vydac C4 analytical column (Figure 2). The isomer ratio was 26:42:12:20, which deviates slightly from the expected statistical ratio, 37.5:37.5:12.5:12.5, for the two *mer* isomers and two *fac* isomers. The CD spectrum of Fe<sup>II</sup>(bipy-GalNAc)<sub>3</sub> showed only weak peaks at 296 nm ( $[\theta]$  =  $7.5 \times 10^4$  deg cm<sup>2</sup>/dmol) and 311 nm ( $[\theta]$  =  $-1.5 \times 10^5$  deg cm<sup>2</sup>/dmol), in accord with the formation of four diastereomers in almost equal amounts.<sup>20</sup>

We studied the binding of bipy-GalNAc, Fe<sup>II</sup>(bipy-GalNAc)<sub>3</sub> and asialoglycophorin A (MN) to *Vicia villosa* B<sub>4</sub> lectin. All synthetic carbohydrates inhibited the binding of peroxide-modified

\* To whom correspondence should be addressed.

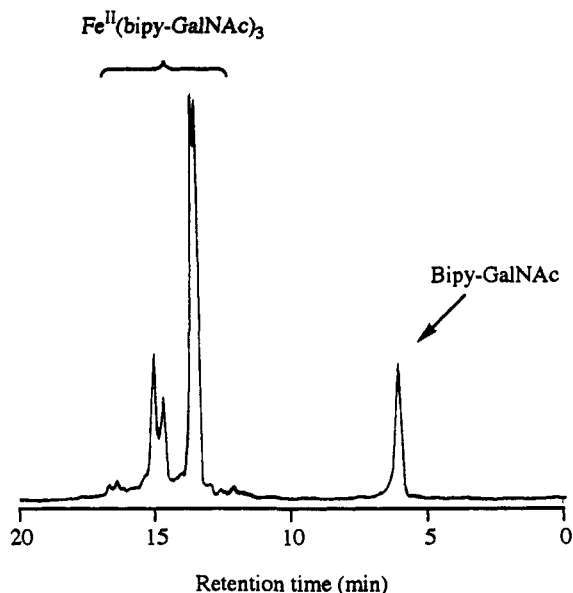
† Present address: Research & Development Department, Tomez Technology Corporation, 2-11-33, Noritakeshimachi, Nishi-ku, Nagoya, 451 Japan.

- (1) Springer, G. F. *Science* **1984**, *224*, 1198–1206.
- (2) Springer, G. F.; Tegtmeyer, H.; Taylor, C. R. *Proc. Am. Assoc. Cancer Res.* **1986**, *27*, 5–65.
- (3) Nakada, H.; Numata, Y.; Inoue, M.; Tanaka, N.; Kitagawa, H.; Funakoshi, I.; Fukui, S.; Yamashina, I. *J. Biol. Chem.* **1991**, *266*, 12402–12405.
- (4) Nakada, H.; Inoue, M.; Numata, Y.; Tanaka, N.; Funakoshi, I.; Fukui, S.; Mellors, A.; Yamashina, I. *Proc. Natl. Acad. Sci. U.S.A.* **1993**, *90*, 2495–2499.
- (5) Tollefsen, S. E.; Kornfeld, R. *J. Biol. Chem.* **1983**, *258*, 5166–5171.
- (6) Tollefsen, S. E.; Kornfeld, R. *J. Biol. Chem.* **1983**, *258*, 5172–5176.
- (7) Piller, V.; Piller, F.; Cartron, J.-P. *Eur. J. Biochem.* **1990**, *191*, 461–466.
- (8) Hakomori, S. *Curr. Opin. Immunol.* **1991**, *3*, 646–653.
- (9) Ferrari, B.; Pavia, A. A. *Int. J. Pept. Protein Res.* **1983**, *22*, 549–559.
- (10) Kunz, H.; Birnbach, S. *Angew. Chem., Int. Ed. Engl.* **1986**, *25*, 360–361.
- (11) Kunz, H.; Birnbach, S.; Wernig, P. *Carbohydr. Res.* **1990**, *202*, 207–223.
- (12) Kunz, H.; Wernig, P.; Schilling, M.; Marz, J.; Unverzagt, C.; Birnbach, S.; Lang, U.; Waldmann, H. *Environ. Health Perspect.* **1990**, *88*, 247–249.
- (13) Toyokuni, T.; Dean, B.; Hakomori, S. *Tetrahedron Lett.* **1990**, *31*, 2673–2676.
- (14) Lieberman, M.; Sasaki, T. *J. Am. Chem. Soc.* **1991**, *113*, 1470–1471.
- (15) Lieberman, M.; Tabet, M.; Tahmassebi, D.; Zhang, J.; Sasaki, T. *Nanotechnology* **1991**, *2*, 203–205.
- (16) Sasaki, T.; Lieberman, M. *Tetrahedron* **1993**, *49*, 3677–3689.
- (17) Fiandor, J.; Garcia-Lopez, M. T.; de las Heras, F. G.; Mendez-Castrillon, P. P. *Synthesis* **1985**, 1121–1123.

(18) The fully acetylated GalNAc was treated with ammonia in acetonitrile to selectively remove the acetyl group at the C-1 hydroxyl group.<sup>17</sup> The resulting tri-*O*-acetyl-*N*-acetylgalactosamine (Ac<sub>3</sub>-GalNAc) was reacted with 5-(bromomethyl)-5'-methylbipyridine in dry DMF in the presence of Ag<sub>2</sub>O. After deacetylation with NaOCH<sub>3</sub> in methanol, bipyridine-modified *N*-acetylgalactosamine (bipy-GalNAc) was purified with C4 reverse-phase HPLC. The small coupling constant of the C-1 proton (3.81 Hz) in <sup>1</sup>H NMR is consistent with the  $\alpha$ -configuration<sup>19</sup> of bipy-GalNAc. Bipy-GalNAc: <sup>1</sup>H NMR (300 MHz, D<sub>2</sub>O)  $\delta$  8.84 (br s, 1H), 8.70 (br s, 1H), 8.45 (br s, 2H), 8.31 (d, 1H,  $J$  = 8.4 Hz), 8.22 (d, 1H,  $J$  = 8.4 Hz), 5.04 (d, 1H,  $J$  = 3.8 Hz, H-1), 4.94 (d, 1H,  $J$  = 13.2 Hz, benzyl), 4.79 (d, 1H,  $J$  = 13.2 Hz, benzyl), 4.20 (dd, 1H,  $J$  = 10.5, 3.7 Hz), 4.05–3.96 (m, 3H), 3.76 (d, 2H,  $J$  = 6.0 Hz, H-6,6'), 2.60 (s, 3H, CH<sub>3</sub>), 2.02 (s, 3H, COCH<sub>3</sub>); MS(FAB)  $m/e$  = 404 (M + H)<sup>+</sup>; UV (20 mM Tris-HCl, 150 mM NaCl, pH 7.2)  $\lambda_{\max}$  240 ( $\epsilon$  = 15 000), 286 nm (19 500).

(19) Pavia, A. A.; Ferrari, B. *Int. J. Pept. Protein Res.* **1983**, *22*, 539–548.

(20)  $\Delta$ -[Fe(bipy)<sub>3</sub>]<sup>2+</sup> complex exhibits peaks at 285 nm ( $[\theta] \geq 4.95 \times 10^5$  deg cm<sup>2</sup>/dmol) and 300 nm ( $[\theta] \leq -1.34 \times 10^6$  deg cm<sup>2</sup>/dmol) in its CD spectrum, see: Milder, S. J.; Gold, J. S.; Kliger, D. S. *J. Am. Chem. Soc.* **1986**, *108*, 8295–8296.

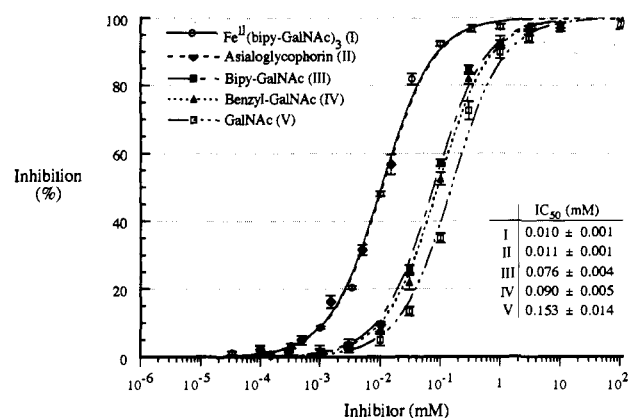


**Figure 2.** HPLC separation of four diastereomers of the  $\text{Fe}^{\text{II}}(\text{bipy-GalNAc})_3$  complex. The complex was analyzed with C4 reverse-phase HPLC, using a solvent system of water- $\text{CH}_3\text{CN}$  containing 0.1% TFA. A linear gradient of 10%–30%  $\text{CH}_3\text{CN}$  over 15 min, with a flow rate of 1.0 mL/min, was employed to separate the four isomers. The chromatography was monitored at 296 nm, which is an isosbestic point for the formation of the  $\text{Fe}^{\text{II}}(\text{bipy-GalNAc})_3$  complex.

$\text{B}_4$  lectin to GalNAc-agarose by competition. Figure 3 shows the binding curves of the synthetic GalNAc derivatives to  $\text{B}_4$  lectin. The calculated relative binding constants<sup>21</sup> for GalNAc, benzyl- $\alpha$ -GalNAc, bipy-GalNAc, and  $\text{Fe}^{\text{II}}(\text{bipy-GalNAc})_3$  are 1.0,  $1.7 \pm 0.1$ ,  $2.0 \pm 0.1$  and  $15.3 \pm 1.0$ , respectively.  $\text{Fe}^{\text{II}}(\text{Me}_2\text{bipy})_3$  neither binds to  $\text{B}_4$  lectin nor enhances the binding of benzyl- $\alpha$ -GalNAc. The increased binding affinity of  $\text{Fe}^{\text{II}}(\text{bipy-GalNAc})_3$  over bipy-GalNAc toward  $\text{B}_4$  lectin must, therefore, be due to the cluster formation of GalNAc residues on the metal template. Asialoglycophorin A (MN) showed a similar binding constant to  $\text{Fe}^{\text{II}}(\text{bipy-GalNAc})_3$ . Asialoglycophorin A (MN) contains several repeating units of GalNAc-modified Ser or Thr residues. Recently, Pepe et al. reported<sup>22</sup> the conformational analysis of a pentapeptide taken from the sequence of asialoglycophorin. The energy-minimized structure of the pentapeptide, Ser-Ser\*-Thr\*-Thr\*-Gly, where \* indicates *O*-linked  $\alpha$ -D-GalNAc residue, shows a cyclic conformation in which the distances between  $\text{C}\alpha$  carbons in this model Tn antigen are 5.3 and 6.7 Å, comparable to inter-GalNAc distances of  $\text{Fe}^{\text{II}}(\text{bipy-GalNAc})_3$  complexes.<sup>23</sup>

(21) Data presented in Figure 3 were fitted to the standard binding isotherm, % inhibition =  $100 \times [\text{inhibitor}] / ([\text{inhibitor}] + \text{IC}_{50})$ . Dividing the  $\text{IC}_{50}$  value for GalNAc by  $\text{IC}_{50}$  values for other inhibitors gave the relative binding constants.

(22) Pepe, G.; Siri, D.; Odon, Y.; Pavia, A. A.; Reboul, J.-P. *Carbohydr. Res.* **1991**, *209*, 67–81.



**Figure 3.** Binding of peroxidase-labeled  $\text{B}_4$  lectin to GalNAc-modified agarose measured in the presence of synthetic GalNAc ligands and asialoglycophorin A as an inhibitor. All inhibition experiments were carried out at 22 °C in an aqueous buffer (20 mM Tris-HCl, 150 mM NaCl, 0.25 or 0.5% bovine serum albumin, pH 7.2). A mixture of lectin (5  $\mu\text{M}$ ), agarose-bound GalNAc (30 mg of packed gel), and an inhibitor of varying concentrations was gently shaken for 2 h to establish the binding equilibrium. The concentration of unbound lectin was determined by a colorimetric assay, using *o*-phenylenediamine and hydrogen peroxide as substrates for peroxidase. Each inhibition experiment was repeated more than three times to ensure reproducibility.

In conclusion, we have demonstrated a novel metal-assisted assembly process of a tridentate GalNAc cluster that is recognized by a Tn-specific lectin. Multivalent carbohydrate-protein interactions have been shown<sup>24–27</sup> in binding of several other lectins to their carbohydrate ligands. A similar metal-assisted assembly process could be used to synthesize analogues of these multivalent carbohydrate ligands. The approach could also be extended to construct heteromeric carbohydrate clusters, using a combination of various bipyridine-modified carbohydrates. A series of such compounds would be useful in generating a library of synthetic carbohydrate ligands for screening of specific inhibitors for various biologically important lectins and receptors.

**Acknowledgment.** This work was supported in part by grants from the American Cancer Society and Tomey Technology Corporation. We thank Dr. Gerald Pepe for providing us with atomic coordinates of the model for carbohydrate-modified pentapeptides including Tn antigen. We also thank Prof. Sen-itiroh Hakomori, Dr. Kohei Yokoyama, and Ms. Marya Lieberman for valuable discussions.

(23) Binding selectivity of four diastereomers of  $\text{Fe}^{\text{II}}(\text{bipy-GalNAc})_3$  was tested using agarose-bound  $\text{B}_4$  lectin. Preliminary experiments showed that isomers nos. 1 and 4 bind to the agarose-bound  $\text{B}_4$  lectin with much higher affinities than to other isomers. Structural characterizations of these isomers are currently underway.

(24) Glick, G. D.; Knowles, J. R. *J. Am. Chem. Soc.* **1991**, *113*, 4701–4703.

(25) Lee, R. T.; Lin, P.; Lee, Y. C. *Biochemistry* **1984**, *23*, 4255–4261.

(26) Lee, R. T.; Lee, Y. C. *Biochem. Biophys. Res. Commun.* **1988**, *155*, 1444–1451.

(27) Wu, A. M.; Sugii, S. *Adv. Exp. Med. Biol.* **1988**, *228*, 205–263.