Nucleoside Recognition by a Fluorescent Macrolactam

Hae-Jo Kim* and Jong-In Hong*,*

Department of Chemistry, College of Natural Sciences, Kyonggi University, Suwon 443-760, Korea. *E-mail: haejkim@kgu.ac.kr

*Department of Chemistry, College of Natural Sciences, Seoul National University, Seoul 151-747, Korea

*E-mail: jihong@smu.ac.kr

Received August 16, 2007

Key Words: Fluorescence, Macrolactam, Nucleoside, Recognition

Molecular recognition of nucleosides or nucleotides is attracting a great deal of interest due to their genetic functions in living organisms. Hydrophilic nature of nucleosides and nucleotides allows only a conformationally well defined receptor to form a hydrogen-bonded, electrostatic or hydrophobic complex with nucleosides or nucleotides in water. Recently, an anthracene derivative was reported to show a higher affinity toward GTP over ATP owing to cooperative interactions of hydrogen bonding and electrostatic interactions between an imidazolium moiety and a phosphate unit. A property of the state of the st

We have developed various sugar receptors with hydrogen-bonding acceptors and donors.⁴ Herein, we report a novel D_2 -symmetric fluorescent macrolactam. This host possesses not only an aromatic cavity for π - π interaction, but also hydrogen-bonding donors/acceptors in the peripheral site of the macrolactam for effective nucleoside recognition.

Macrolactam host was synthesized via the typical acid chloride coupling method⁵ in which 2,5-dimethyl-p-xylyl-diamine was treated with 2,5-dimethoxyterephthaloyl chloride in a high dilute condition⁶ to afford the desired 2:2 macrocyclization product (H). The calculated structure shows that the host has a large cavity with dimension of $10.5 \text{ Å} \times 6.9 \text{ Å}$ (Fig. 1. left). The global minimum structure clearly indicates that π - π stacking interaction exists between the dimethoxy aryl groups of H and the uracil base of uridine with aromatic-aromatic surface distances of 3.56 and 3.55 Å, and one intermolecular H-bonding interaction also exists between the carbonyl group of H and 2-OH group of uridine (Fig. 1. right). ⁷

Owing to the characteristic fluorescence property of \mathbf{H} , fluorescence titration was carried out in chloroform. Fluorescence emission intensities at $\lambda_{max} = 384$ nm were recorded after excitation at $\lambda_{ex} = 331$ nm (Fig. 2). Fluorescence intensities of the host-guest complex increase upon addition of sugars or nucleosides presumably due to the restricted rotation of \mathbf{H} . The resulting fluorescence enhancements at 384 nm are shown in the inset of Figure 3. The binding stoichiometry between \mathbf{H} and guests was also confirmed to be 1:1 by Job's plot (Fig. 4).

Curve fitting of the host signals to a 1:1 binding isotherm gives apparent dissociation constants of up to $K_d = 10^{-4}$ M, which are summarized in Table 1.

While the dissociation constants between **H** and anomers of D-glucose were found to be similar $(3.99 \times 10^{-4} \text{ M for } \beta)$

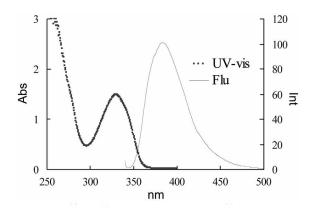


Figure 2. UV-vis and fluorescence spectra of H in CHCl₃ at 298 K.

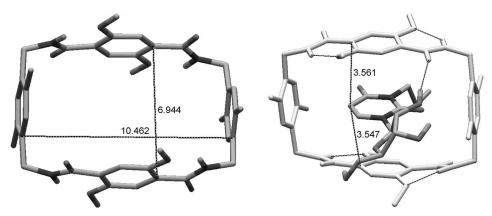


Figure 1. Global minimum structures of H (left) and its uridine complex (right).

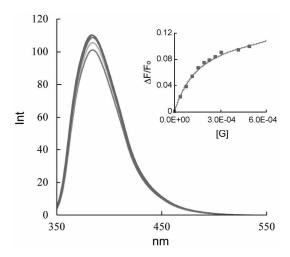


Figure 3. Fluorescence titration of **H** and uridine in CHCl₃ at 298 K. $[\mathbf{H}] = 2.0 \, \mu M$.

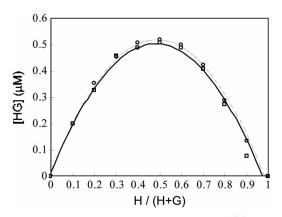


Figure 4. Job's plot between **H** and D-glucopyranosides at 298 K. [H] + [G] = 2.0 μ M, each in 2.0 mL. Rectangular and circle represent β -glucose and uridine, respectively.

and 5.38×10^{-4} M for α anomer), the binding affinity of H to β -galactose is three times lower than that of β -D-glucose (1.31 \times 10⁻³ M for β -D-galactose). This diastereoselectivity for sugars plausibly results from the slight energetic difference in the intermolecular H-bonding patterns due to the varying degree of steric interaction between sugars and H. This indicates that geometrical complementarities of H-bonding partners are crucial in hydrogen bond-based molecular recognition system.

It is noticeable that nucleosides, deoxythymidine (d-Thy) and uridine (Uri) show the comparable binding affinities although they have fewer number of hydroxyl groups compared with the pyranosides. Uridine shows much higher binding affinity (1.72 × 10⁻⁴ M) than β -D-glucose. Enhancement in the binding affinity for nucleosides probably results from the presence of π -surface and H-bonding donors and acceptors in the guests.

It is assumed that π - π stacking interaction between **H** and nucleosides plays an important role in host-guest binding. We have chosen several commercially available aromatic guests to test this assumption. While benzene is weakly bound to \mathbf{H} ($K_d = 4.05 \times 10^{-2}$ M), the binding affinity of a π

Table 1. Dissociation constants between H and guests"

entry guest structure name $K_d(M)$			
entry	guest structure	name	Kd (IVI)
1	HOH HOH OOct	β-D-Glucose	3.99(±0.70) × 10 ⁻⁴
2	HOH HOH HOOCt	α-D-Glucose	$5.38(\pm 3.22) \times 10^{-4}$
3	HOOH HOH OH OOct	β-D-Galactose	$1.31(\pm 0.58) \times 10^{-3}$
4	HO OH	Thymidine	$6.44(\pm 5.88) \times 10^{-4}$
5	HO OHOH	Uridine	$1.72(\pm 0.23) \times 10^{-4}$

°Fluorescence (itration of constant host concentration (2.0 μ M) in CHCl₃ at 298 K. Fluorescence intensity at $\lambda_{em} = 384$ nm ($\lambda_{ex} = 331$ nm) was monitored after each addition of guest,

basic guest 1,4-dimethoxybenzene was c.a. hundred times enhanced (5.26×10^{-4} M). The binding affinity of a π acidic guest dimethylterephthalate, however, was too small to determine.

In conclusion, we have developed a novel fluorescent macrolactam as an artificial receptor for nucleosides. The receptor has shown high diastereoselectivity for sugars and even higher affinities for nucleosides due to the intermolecular π - π stacking interaction as well as H-bonds between the macrolactam and sugars/nucleosides.

Experimental

Acid chloride synthesis. To a solution of 400 mg (1.77 mmol) of 2,5-dimethoxyterephthalic acid in 20 mL of dichloromethane was added cat. amount of DMF and 2.0 mL of 2 M oxalic acid chloride in dichloromethane (2 eq. ex., 4.0 mmol). Resulting white suspension was stirred at rt

Scheme 1. Synthetic scheme of macrolactam.

under nitrogen for 5 hrs to afford a yellow clear solution. All volatiles were removed under the reduced pressure, dried in vacuum.

Cyclization. To a solution of p-xylyl diamine (1 eq. 1.77 mmol) and TEA (2 eq ex.) in 500 mL of dichloromethane was dropwise added a solution of above crude 2,5-dimethoxyterephthaloyl chloride in 50 mL of dichloromethane at 0 °C under nitrogen over a period of 2 hrs. Resulting yellow solution was stirred for additional 24 hrs under nitrogen. All volatiles are removed under reduced pressure and purified by column chromatography. Column chromatography on silica gel (CH₂Cl₂:MeOH = 10:1, $R_{\rm f}$ = 0.48) gave a greenish mixture. Additional column chromatography on silica gel (EtOAc, $R_{\rm f}$ = 0.30) gave the desired product, **H** as a white solid in a 4.2% yield.

¹H-NMR (300 MHz, CDCl₃): 8.09 (t, J = 6.3 Hz, 4H of NH), 7.68 (s, 4H of ArH₄ in 2,5-dimethoxybenzene), 7.06 (s, 4H of ArH in *p*-xylylene), 4.51 (d, J = 6.3 Hz, 8H of ArCH₂N), 3.90 (s, 12H of OCH₃), 2.29 (s, 12H of ArCH₃).

UV-vis (CHCl₃): $\varepsilon_{331nm} = 3997 \text{ M}^{-1}\text{cm}^{-1}$, Fluorescence (CHCl₃): $\lambda_{em} = 384 \text{ nm}$ ($\lambda_{ex} = 331 \text{ nm}$) in 2.0 μ M Mass (FAB⁺, m-NBA): m/z 709 ([M+H], 50%)

Acknowledgement. Financial support from Korea Research Foundation (KRF-2006-312-C00592) is greatly acknowledged.

References

- (a) Yang, F.; Belitsky, J. M.; Villanueva, R. A.; Dervan, P. B.; Roth, M. J. *Biochemistry* 2003, 42, 6249.
 (b) Gearhart, M. D.; Dickinson, L.; Ehley, J.; Melander, C.; Dervan, P. B.; Wright, P. E.; Gottesfeld, J. M. *Biochemistry* 2005, 44, 4196.
- (a) Urbach, A. R.; Dervan, P. B. Proc. Nat'l Aca. Sci. 2001, 98, 4343.
 (b) Chenoweth, D. M.; Viger, A.; Dervan, P. B. J. Am. Chem. Soc. 2007, 129, 2216.
- Kwon, J. Y.; Singh, N. J.; Kim, H. N.; Kim, S. K.; Kim, K. S.; Yoon, J. J. Am. Chem. Soc. 2004, 126, 8892.
- (a) Cho, H. -K.; Kim, H. -J.; Lee, K. H.; Hong, J.-I. Bull. Korean Chem. Soc. 2004, 25, 1714. (b) Lee, D. H.; Kim, H.-J.; Hong, J.-I. Supram, Chem. 2007, 19, 251.
- Kim, H.-J.; Kim, Y.-H.; Hong, J.-I. Tetrahedron Lett. 2001, 42, 5049.
- 6. Kim, T. W.; Hong, J.-I. Bull. Korean Chem. Soc. 1995, 16, 781.
- Conformation searches were performed with the program MacroModel 7.0, Amber* force field in chloroform solvent: Mohamadi, F.; Richards, N. G. J.; Guida, W. C.; Liskamp, R.; Lipton, M.; Caufield, C.; Chang, G.; Hendrickson, T.; Still, W. C. J. Comput. Chem. 1990, 11, 440.
- (a) Jang, Y. J.; Jun, J. H.; Swamy, K. M. K.; Nakamura, K.; Koh, H. S.; Yoon, Y. J.; Yoon, J. Bull. Korean Chem. Soc. 2005, 26, 2041. (b) An, B.-K.; Kwon, S.-K.; Park, S. Y. Bull. Korean Chem. Soc. 2005, 26, 1555
- (a) Lee, D. H.; Im, J. H.; Lee, J.-H.; Hong, J.-I. Tetrahedron Lett.
 2002, 43, 9637. (b) Mcfarland, S. A.; Finney, N. S. J. Am. Chem. Soc. 2002, 124, 1178.
- 10. Job. P. Ann. Chim. 1928, 9, 113.