Development of Dipicolylamine-Modified Cyclodextrins for the Design of Selective Guest-Responsive Receptors for ATP

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Academic Editor: Bernard Martel
Received: 27 February 2018; Accepted: 11 March 2018; Published: 12 March 2018

Abstract: The construction of supramolecular recognition systems based on specific host–guest interactions has been studied in order to design selective chemical sensors. In this study, guest-responsive receptors for ATP have been designed with cyclodextrins (CyDs) as a basic prototype of the turn-on type fluorescent indicator. We synthesized dipicolylamine (DPA)-modified CyD–Cu$^{2+}$ complexes ($\text{Cu·}1\alpha$, $\text{Cu·}1\beta$, and $\text{Cu·}1\gamma$), and evaluated their recognition capabilities toward phosphoric acid derivatives in water. The UV-Vis absorption and fluorescence spectra revealed that $\text{Cu·}1\beta$ selectively recognized ATP over other organic and inorganic phosphates, and that $\beta$-CyD had the most suitable cavity size for complexation with ATP. The 1D and 2D NMR analyses suggested that the ATP recognition was based on the host–guest interaction between the adenine moiety of ATP and the CyD cavity, as well as the recognition of phosphoric moieties by the Cu$^{2+}$–DPA complex site. The specific interactions between the CyD cavity and the nucleobases enabled us to distinguish ATP from other nucleoside triphosphates, such as guanosine triphosphate (GTP), uridine triphosphate (UTP), and cytidine triphosphate (CTP). This study clarified the basic mechanisms of molecular recognition by modified CyDs, and suggested the potential for further application of CyDs in the design of highly selective supramolecular recognition systems for certain molecular targets in water.

Keywords: supramolecular chemistry; cyclodextrin; ATP recognition; host–guest interaction; NMR analyses

1. Introduction

In the field of supramolecular chemistry, many kinds of molecular recognition systems have been developed that function by the specific interactions between host and guest molecules [1–4]. Cyclodextrins (CyDs), which are macrocyclic oligosaccharides composed of D-glucopyranose units, are well-known host molecules [5]. As CyDs include organic molecules in their hydrophobic cavities and increase the solubility and stability of the guest molecules in water, these molecules are utilized in various chemical fields, including analytical chemistry, organic chemistry, and biochemistry [6–8]. In addition, the application of specific host–guest interactions of chemically modified CyDs has resulted in such innovations as sensors, catalysts, functional materials, and carriers for drug delivery.
systems [9–13]. Therefore, it is very important to develop a design strategy to obtain selective CyD-based inclusion complexes. Recent reports of supramolecular recognition systems based on modified CyDs have revealed that CyDs possess the potential for use as highly selective receptors for various kinds of organic molecules [14–16]. For example, positively charged CyD bearing aminomethyl groups bound to adenosine monophosphate (AMP), adenosine diphosphate (ADP), or adenosine triphosphate (ATP) [17,18]. The affinity of these CyD hosts for nucleotides or nucleosides has been well investigated, although there are still few applications of CyDs to chemical sensors [19–24]. On the other hand, ATP recognition has been studied because of its significant roles in biological systems, and thus a number of ATP-targeting fluorescent probes have been reported over the last few decades [25–27]. However, there have been few examples of turn-on type chemical indicators that can distinguish nucleoside triphosphate from similar derivatives. Previously, we reported that Cu\(^{2+}\) complexes of dipicolylamine (DPA)-modified \(\gamma\)-CyD that possess an azobenzene unit selectively recognize ATP over other phosphoric acid derivatives [28]. The study demonstrated that multipoint recognition systems enable us to design selective receptors for certain organic molecules such as ATP. Here, we report a rational design strategy to obtain selective supramolecular recognition systems for ATP in water, where we used the affinity of chemically modified CyDs for nucleobases as a basic prototype of the turn-on type fluorescent indicators.

Our receptors (Figure 1) have two recognition sites: the DPA unit tethering with coumarin fluorophore (guest-targeting probe) and the CyD cavity. The DPA unit binds with heavy metal ions to form chelate complexes, and these metal complexes are known to recognize phosphoric acid derivatives [29]. The distance between the above two recognition sites was designed to enable selective host–guest interactions between the adenine moiety of ATP and the CyD cavity in addition to the recognition of phosphoric moieties by the Cu\(^{2+}\)–DPA complex site. Although many studies use the Zn\(^{2+}\)–DPA complex as the recognition site of phosphoric acid derivatives because of its high binding affinity, we dared to choose Cu\(^{2+}\) as the coordination center to design turn-on type fluorogenic receptors by the photo-induced electron transfer (PET) mechanism [30]. The Cu\(^{2+}\) complex of the guest-targeting probe is almost quenched but is activated by the coordination of the phosphoric moiety of ATP to the Cu\(^{2+}\)–DPA complex site. We evaluated the recognition capabilities of the metal complexes of 1α, 1β, and 1γ (Scheme 1) toward phosphoric acid derivatives. Three CyDs of different sizes were used to confirm the importance of the CyD cavity in this supramolecular recognition system.

![Figure 1. Structure of dipicolylamine-modified cyclodextrins (CyDs).](image-url)

![Scheme 1. Synthesis of 1α, 1β, and 1γ (n: the number of glucose units).](image-url)
2. Results and Discussion

2.1. Synthesis of 1α, 1β, and 1γ

Fluorescent probe 1, which possesses coumarin as the fluorophore and DPA as the recognition site, was synthesized by the Mannich reaction [30]. Probe 1 and 3-NH2-(α, β, or γ)-CyD were coupled by a condensation reaction [31]. The crude product was purified by reverse-phase column chromatography to obtain 1α, 1β, and 1γ. All the samples were identified by 1H NMR measurement and elemental analysis.

2.2. Metal Ion Recognition by 1α, 1β, and 1γ

First, to evaluate the response function of modified CyDs toward metal ions, the fluorescence spectra of 1α, 1β, and 1γ were measured in the presence of Mg2+, Fe3+, Co2+, Ni2+, Cu2+, Zn2+, Cd2+, Pt2+, and Pb2+. The addition of Zn2+ and Cd2+ possessing full d-orbitals enhanced the fluorescence intensity on the basis of the PET mechanism (Figure 2) [32]. On the other hand, the addition of Co2+, Ni2+, and Cu2+ possessing free d-orbitals quenched the fluorescence intensity due to the mechanisms of energy transfer (ET) or ligand to metal charge transfer (LMCT) [33,34]. Probes 1α, 1β, and 1γ exhibited the same fluorescence behavior toward the metal ions, indicating that the CyD cavity has little influence on metal ion recognition. The complexation of the DPA unit to the metal ion was confirmed by metal ion titration experiments.

![Fluorescence spectra of 1β in the presence of metal ions in water, pH 6.4 adjusted by piperazine-1,4-bis(2-ethanesulfonic acid) (PIPES)/NaOH buffer, at 25 °C (λex = 369 nm).](image)

2.3. Phosphoric Acid Derivatives Recognition by 1α, 1β, and 1γ Metal Ion Complexes

In order to examine the recognition functions for phosphoric acid derivatives, the UV-Vis absorption spectra of Cu·1α, Cu·1β, and Cu·1γ were measured in the presence of monophosphate (Pi), pyrophosphate (PPI), triphosphate (Tri), adenosine monophosphate (AMP), adenosine diphosphate (ADP), and ATP. The addition of ATP markedly decreased the absorbance at 404 nm derived from the coumarin moiety (Figure 3). The coordination of phosphate anion to Cu2+ ion weakens the bond between the phenol oxygen atom and Cu2+ ion [35]. In addition, it is known that adenine interacts with aromatic compounds, including coumarin derivatives [36,37]. Both coordination and interaction are considered to change the characteristics of the electronic state on the coumarin moiety, which results in a specific absorbance change. The fluorescence spectra of Cu·1α, Cu·1β, and Cu·1γ were also examined to investigate the potential use of these molecules as fluorescent sensors. The fluorescence intensity at 450 nm was increased upon the addition of ATP (Figure 4). At first, the Cu2+ ion induced
fluorescence quenching via ET or LMCT to the Cu\(^{2+}\) ion. However, the coordination of the phosphate anion to the Cu\(^{2+}\) ion weakened the bond between the DPA unit and the Cu\(^{2+}\) ion, which prevented the probe from inducing fluorescence quenching [38]. Therefore, the fluorescence intensity was recovered. In the presence of ATP, Cu·\(1\beta\) exhibited the most significant absorbance decrease and fluorescence enhancement. This selectivity for ATP suggested two mechanisms: the host–guest interaction between the adenine moiety of ATP and the CyD cavity, and the recognition of the phosphoric moiety of ATP by the Cu\(^{2+}\)–DPA complex site. The host–guest interaction stabilized supramolecular complex ATP/\(\text{Cu·1}_\beta\), which led to the specific response. These results indicated that the CyD cavity played an important role in the recognition system.

Figure 3. UV-Vis responses of (a) Cu·\(1\alpha\), (b) Cu·\(1\beta\), and (c) Cu·\(1\gamma\) in the presence of phosphoric acid derivatives in water, pH 6.4 adjusted by PIPES/NaOH buffer, at 25 °C. [\(1\alpha, 1\beta, \text{or } 1\gamma\)] = 0.010 mM, [Cu(NO\(_3\))\(_2\)] = 0.010 mM, [phosphoric acid derivative] = 1.0 mM.

Figure 4. Fluorescence responses of (a) Cu·\(1\alpha\), (b) Cu·\(1\beta\), and (c) Cu·\(1\gamma\) in the presence of phosphoric acid derivatives in water, pH 6.4 adjusted by PIPES/NaOH buffer, at 25 °C (\(\lambda_{\text{ex}}\) = 404, 408, 408 nm). [\(1\alpha, 1\beta, \text{or } 1\gamma\)] = 0.010 mM, [Cu(NO\(_3\))\(_2\)] = 0.010 mM, [phosphoric acid derivative] = 1.0 mM.

We also evaluated the UV-Vis absorption spectra and the fluorescence spectra of Zn\(^{2+}\) complexes Zn·\(1\alpha\), Zn·\(1\beta\), and Zn·\(1\gamma\). The absorbance of Zn·\(1\alpha\), Zn·\(1\beta\), and Zn·\(1\gamma\) was decreased upon the addition of ATP but these responses were weaker than those of the Cu\(^{2+}\) complexes. The addition of ATP decreased the fluorescence intensity of Zn·\(1\alpha\), Zn·\(1\beta\), and Zn·\(1\gamma\). The Zn\(^{2+}\) complexes exhibit fluorescence emission by suppressing the intramolecular PET mechanism. The coordination of the phosphate anion to the Zn\(^{2+}\) ion weakens the bond between the DPA unit and the Zn\(^{2+}\) ion, thereby inducing fluorescence quenching. However, the quenching was not suitable for our strategy because we wanted to focus on the application to turn-on type indicators.
To compare the complex formation constants to ATP, the UV-Vis absorption spectra of Cu·1α, Cu·1β, and Cu·1γ in the presence of various concentrations of ATP were measured. The complex formation constants of Cu·1α, Cu·1β, and Cu·1γ to ATP were determined to be (1.7 ± 0.06) × 10^3 M\(^{-1}\), (3.3 ± 0.15) × 10^3 M\(^{-1}\), and (2.3 ± 0.12) × 10^3 M\(^{-1}\), respectively, by the Benesi–Hildebrand method [39]. These complex formation constants confirmed that Cu·1β formed the most stable supramolecular complex with ATP in water.

The UV-Vis and fluorescence responses revealed that Cu·1β selectively recognized ATP over other phosphoric acid derivatives. In order to evaluate the recognition selectivity of Cu·1β for nucleoside triphosphates, UV-Vis absorption spectra and fluorescence spectra were measured in the presence of each nucleoside triphosphate: ATP, guanosine triphosphate (GTP), uridine triphosphate (UTP), and cytidine triphosphate (CTP). The addition of each nucleoside triphosphate decreased the absorbance and enhanced the fluorescence intensity of Cu·1β (Figure 5). In particular, Cu·1β selectively recognized ATP over the other nucleoside triphosphates. This selectivity is considered to be based on the host–guest interaction between the nucleobase and the CyD cavity. Adenine is known to interact well with the CyD cavity compared with the other nucleobases [17]. This result also disclosed that the CyD cavity is involved in the recognition. Moreover, Cu·1β selectively recognized a purine base over a pyrimidine base, indicating the potential for use as receptors capable of distinguishing between purine and pyrimidine type nucleoside triphosphates.

Figure 5. (a) UV-Vis and (b) Fluorescence responses of Cu·1β in the presence of nucleoside triphosphates in water, pH 6.4 adjusted by PIPES/NaOH buffer, at 25 °C (\(\lambda_{ex} = 408\) nm). [1β] = 0.010 mM, [Cu(NO\(_3\))\(_2\)] = 0.010 mM, [nucleoside triphosphate] = 1.0 mM.

2.4. 1D and 2D NMR Analyses

In order to clarify the coordination and interaction between Cu·1β and ATP, the \(^1\)H NMR spectra of complex Zn·1β were measured in the presence and absence of ATP. In general, it is difficult to evaluate the NMR spectra of the Cu\(^{2+}\) complex owing to the paramagnetism of the Cu\(^{2+}\) ion. Upon the addition of ATP, shifts of proton peaks assigned to h and i derived from the DPA site were noted (Figure 6), indicating that the Zn\(^{2+}\)-DPA complex site recognized the phosphoric moiety of ATP [40–42]. Shifts of proton peaks assigned to b and c derived from the coumarin moiety were also observed, revealing that the hydroxyl group is associated with the coordination to Zn\(^{2+}\) ion.

We also analyzed the supramolecular conformation of the ATP/Cu·1β complex by conducting nuclear Overhauser effect spectroscopy (NOESY) measurements of complex Cu·1β. In the NOESY measurements, protons in close proximity to each other give rise to correlation peaks [43,44]. Correlations between the protons derived from the adenine moiety of ATP (α and β) and the protons derived from the inside cavity of CyD (H3 and H5) were observed (Figure 7). The results suggested that the adenine moiety interacted with the inside cavity of CyD due to host–guest interactions. If the
NOESY spectra of Cu·1β could be measured, much stronger correlations would be expected because the Cu$^{2+}$ complex exhibited the higher binding ability to ATP.

Figure 6. $^1$H NMR spectra of complex ATP/Zn·1β and ATP (500 MHz, solvent: D$_2$O). (i) [Zn·1β] = 0.5 mM, (ii) [Zn·1β] = 0.5 mM, [ATP] = 0.5 mM, (iii) [Zn·1β] = 0.5 mM, [ATP] = 5.0 mM, (iv) [ATP] = 10 mM.

Figure 7. NOESY spectrum of complex ATP/Zn·1β (400 MHz, solvent: D$_2$O). [Zn·1β] = 5.0 mM, [ATP] = 25 mM.

2.5. Suggested Supramolecular Conformation

To sum up the obtained results, the supramolecular conformation shown in Figure 8 is suggested. First, the coordination of the DPA site and the hydroxyl group to the metal ion was confirmed by UV-Vis and $^1$H NMR measurements. Secondly, the coordination of the phosphoric moiety of ATP to the
metal ion–DPA complex site was revealed by $^1$H NMR experiments. Thirdly, the interaction between the adenine moiety of ATP and the coumarin fluorophore was implied by the UV-Vis responses, although this interaction was weak. Finally, the host–guest interaction between the adenine moiety and the CyD cavity was suggested by NOESY measurements. Thus, Cu·1$\beta$ was found to recognize ATP through the host–guest interaction between the adenine moiety and the CyD cavity in addition to the recognition of the phosphoric moiety by the metal ion–DPA complex site. In particular, the cavity size of CyD is a major contributor to the difference in selectivity for ATP. Therefore, the selective ATP recognition in this system is considered to markedly depend on the multipoint interaction in water.

![Figure 8. Suggested supramolecular conformation of the ATP/Cu·1$\beta$ complex.](image)

### 3. Materials and Methods

#### 3.1. Reagents

All organic solvents and reagents including 3-NH$_2$-CyDs were commercially available with guaranteed grades and used without further purification. Water was doubly distilled and deionized by a Milli-Q water system before use.

#### 3.2. Apparatus

UV-Vis absorption spectra were measured with a UV-Vis spectrophotometer equipped with a Peltier thermocontroller and a 10 mm quartz cell. Fluorescence spectra were measured with a fluorescence spectrometer with a 10 mm quartz cell.

#### 3.3. Metal Ion Recognition by 1$a$, 1$\beta$, and 1$\gamma$

To evaluate the metal ion recognition ability of 1$a$, 1$\beta$, and 1$\gamma$, fluorescence spectral measurements were performed. Solutions containing 1$a$, 1$\beta$, and 1$\gamma$ (0.010 mM), metal ion (0.010 mM), and piperazine-1,4-bis(2-ethanesulfonic acid) (PIPES)/NaOH buffer (5.0 mM, pH 6.4, adjusted using NaOH) were prepared and spectra were recorded at 25 °C.

#### 3.4. Phosphoric Acid Derivatives Recognition by 1$a$, 1$\beta$, and 1$\gamma$ Metal Ion Complexes

To evaluate the phosphoric acid recognition ability of 1$a$, 1$\beta$, and 1$\gamma$ metal ion complexes, UV-Vis absorption and fluorescence spectral measurements were performed. Solutions containing 1$a$, 1$\beta$, and 1$\gamma$ (0.010 mM), Cu(NO$_3$)$_2$ or Zn(NO$_3$)$_2$ (0.010 mM), phosphoric acid derivatives (1.0 mM, pH 7.4, adjusted using NaOH), and PIPES/NaOH buffer (10 mM, pH 6.4, adjusted using NaOH) were prepared and spectra were recorded at 25 °C.

#### 3.5. Calculation of Binding Constants of Cu·1$a$, Cu·1$\beta$, and Cu·1$\gamma$ to ATP

To calculate the binding constants of Cu·1$a$, Cu·1$\beta$, and Cu·1$\gamma$ to ATP, UV-Vis absorption spectral measurements were performed. Solutions containing 1$a$, 1$\beta$, and 1$\gamma$ (0.010 mM), Cu(NO$_3$)$_2$ (0.010 mM),
and PIPES/NaOH buffer (10 mM, pH 6.4, adjusted using NaOH) were prepared and spectra were recorded at 25 °C, while ATP concentrations varied from 0 to 4.0 mM. The binding constants were calculated by the Benesi–Hildebrand method.

3.6. NOESY Measurements of Zn·1β/ATP Complex

To evaluate the supramolecular conformation of the Cu·1β/ATP complex, NOESY spectral measurements of the Zn·1β/ATP complex were performed. Solutions containing 1β (5.0 mM), Zn(NO3)2 (5.0 mM), and ATP (25 mM, pD 7, adjusted using NaOD) were prepared and NOESY spectra were recorded.

3.7. General Procedure for the Synthesis of 1

Aqueous formaldehyde (305 mg, 3.76 mmol) was added to a solution of DPA (729 mg, 3.66 mmol) in 15 cm3 of acetonitrile. After 3 h of refluxing, 7-hydroxycoumarin-3-carboxylic acid (499 mg, 2.42 mmol) dissolved in 30 cm3 of acetonitrile was added dropwise. The entire reaction mixture was stirred at 40 °C for 21 h. The yellow precipitate was filtered and washed with acetonitrile. The yellow solid was washed with water and extracted with chloroform, and the extract was concentrated to obtain 1 as a yellow solid in 71% yield. The sample was identified by 1H NMR and FAB-MS. The characterization data are shown in the Supplementary Materials (Figures S1 and S2, and Table S1).

3.8. General Procedure for the Synthesis of 1α, 1β, and 1γ

A mixture of 1, N,N'-dicyclohexylcarbodiimide (DCC), and 1-hydroxybenzotriazole (HOBt)-H2O (Table 1) in 4.0 cm3 of super dehydrated N,N-dimethylformamide (DMF) was stirred in an ice bath for 20 min. The compound 3-NH2-(α, β, or γ)-CyD in 2.0 cm3 of super dehydrated DMF was added dropwise over a period of 5 min. The entire reaction mixture was again stirred in an ice bath for 30 min and then at room temperature for 24 h. The solution was distilled under reduced pressure until half of the DMF was removed. The sample was kept in a refrigerator for one day. The resulting white solid was removed by filtration. The filtrate was poured into 800 cm3 of acetone and stirred for 1 h to obtain a yellow precipitate. The crude product was charged on a column of CHROMATOREX ODS-DM1020T (reverse phase) and eluted with 10% acetonitrile-90% water (v/v, 1.0 dm3) and then with 20% acetonitrile-80% water (v/v, 0.5 dm3). The second spot was collected as the desired product. The fraction was lyophilized under reduced pressure to obtain a yellow solid. The compounds 1α, 1β, and 1γ were obtained in 52%, 46%, and 49% yield, respectively. All the samples were identified by 1H NMR measurement and elemental analysis. The characterization data are shown in the Supplementary Materials (Figures S3–S5 and Tables S2–S7).

| Compound | DCC  | HOBT·H2O | 3-NH2-(α, β, or γ)-CyD |
|----------|------|----------|-----------------------|
| 1α       | 24 mg, 0.11 mmol | 18 mg, 0.12 mmol | 70 mg, 0.072 mmol |
| 1β       | 28 mg, 0.14 mmol | 21 mg, 0.14 mmol | 102 mg, 0.090 mmol |
| 1γ       | 22 mg, 0.11 mmol | 16 mg, 0.10 mmol | 82 mg, 0.063 mmol |

4. Conclusions

We have developed fluorogenic DPA-modified CyDs to obtain supramolecular recognition systems for ATP as a basic prototype of the selective guest-responsive indicator. The absorption and fluorescence responses revealed that Cu·1β recognized ATP selectively over other phosphoric acid derivatives possessing similar phosphoric moieties or nucleobases. The β-CyD cavity was found to be the most acceptable for the formation of the supramolecular complex with ATP. NMR analysis suggested that the selective ATP recognition not only depended on the recognition of phosphoric moieties by the Cu2+-DPA complex site but also on the host–guest interactions between the adenine
moiety of ATP and the CyD cavity. The results demonstrated that the CyD cavity played an important role in the selectivity over other nucleoside triphosphates in the supramolecular recognition system. Therefore, this system enabled us to discriminate a purine base from a pyrimidine base as well as the lengths of the phosphoric moieties. This study clarified the basic mechanisms of modified CyDs as selective indicators, and suggested the potential for further application in the design of supramolecular recognition systems based on CyDs with high selectivity for certain molecular targets in water.

**Supplementary Materials:** Supplementary materials are available online. Compound characterization data, the pH profiles, phosphoric acid recognition by UV-Vis and fluorescence measurements of 1α, 1β, or 1γ metal ion complexes, calculation of binding constants, competitive experiments, and copies of 1H-1H COSY and NOESY spectra of Zn·1β/ATP complex.

**Acknowledgments:** This work was supported by JSPS KAKENHI Grant Numbers JP15K05548 and JP26248038.

**Author Contributions:** T. Hayashita conceived, designed, and supervised the project; T.Y. and S.F. performed the experiments; K.F., Y.T., and T. Hashimoto analyzed the data; T.Y. and K.F. wrote the manuscript.

**Conflicts of Interest:** The authors declare no conflict of interest.

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**Sample Availability:** Samples of the compounds are not available from the authors.

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