

## Metallohelical Triangles for Selective Detection of Adenosine Triphosphate in Aqueous Media

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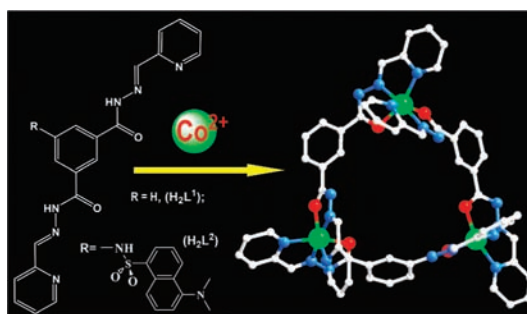
Metallohelical triangles consisting of chromophore units and hydrogen-bonding trigger sites were prepared by modulating two tridentate  $N_2O$  units containing amide groups within a central benzene ring at the meta sites, for the selective detection of adenosine triphosphate in aqueous media over other ribonucleotides.

Among biologically important anions, a great deal of interest has been focused on the molecular recognition of the nucleotide polyphosphates; in particular, adenosine triphosphate is one of the basic components in bioenergetic processes of living organisms, with its polyphosphate chain being the center for chemical energy storage and transfer.<sup>1</sup> Although several intriguing strategies have been developed to detect adenosine triphosphate (ATP), such as synthetic host–guest receptors,<sup>2,3</sup> peptides,<sup>4</sup> and RNA aptamers,<sup>5,6</sup> the complexation selectivity, particularly the selectivity to adenine over all other nucleobases, is still insufficient and remains as a challenging problem.<sup>7</sup>

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**Scheme 1.** Molecular Structures of Ligands  $H_2L^1$  and  $H_2L^2$  as Well as the Cationic Helical Triangle  $[Co_3(HL^1)_3]^{3+\alpha}$



<sup>a</sup> The metal, oxygen, nitrogen, and carbon atoms are drawn as green, red, blue, and white, respectively.

On the other hand, self-assembled, metal–ligand coordination molecular hollows are appealing as synthetic hosts for application as a fascinating class of size- or shape-selective dynamic molecular chemosensors with various responses and metal-triggered functions.<sup>8,9</sup> By using carefully functionalized amide groups as the trigger sites for efficient host–guest interactions, these Werner-type capsules and some of the organic capsules that have the potential to recognize the important biological molecules have been realized.<sup>10,11</sup> Herein, we develop a new strategy for preparing helical triangles through modulation of the tridentate  $N_2O$  units containing amide groups within a central benzene ring at the meta sites (Scheme 1). We reasoned that these amide groups in the positively charged cages matched the static,

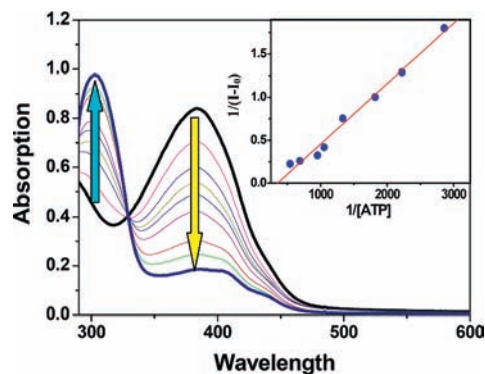
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geometric, coordinative, and functional requirements of the bowl-like capsules for recognition of nucleotide anions.<sup>12</sup> Also, the formation of hydrogen bonds of the amide groups is likely to perturb the electronic distribution of the conjugated backbone of the ligand, leading to significant changes in optical properties.<sup>13</sup>

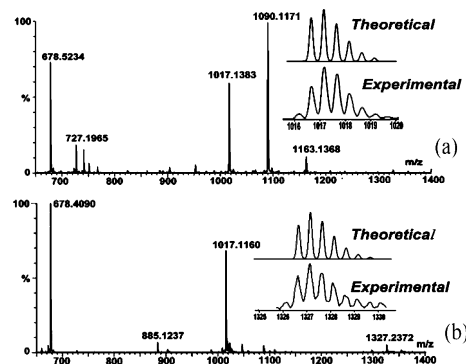
The reaction of 1,3-dicarbohydrazidobenzene with 2-pyridinecarboxaldehyde gave a ligand,  $H_2L^1$ . Adding  $NH_4PF_6$  into the methanol solution of ligand  $H_2L^1$  and  $Co(NO_3)_2 \cdot 6H_2O$  led to the formation of the cobalt(II)-based helical triangle **MC-1**.<sup>14</sup> The triangular cation  $[Co_3(HL_1)_3]^{3+}$  has an ideal  $C_3$  symmetry with three identical cobalt(II) centers exhibiting the same chiral coordination geometries. The three benzene rings of the ligands lie on the same side of the averaged plane (Scheme 1). The presence of three  $PF_6^-$  anions corresponding to each cationic cage suggests that three of the six amide groups lose their protons during coordination to metal ions.

Ligand  $H_2L^1$  exhibits an absorption band at 300 nm in  $N,N$ -dimethylformamide (DMF)/water (8:2) aqueous media, while the Co-based compound **MC-1** shows a band at about 380 nm. The absorbance of **MC-1** was a slight change over a wide concentration range of acid/base (from  $\log [HCl] = -4.2$  to  $\log [NaOH] = -2.5$ ; see Figure S9 in the Supporting Information). The addition of ATP to the solution of **MC-1** causes a significant absorbance increase at 300 nm and an obvious absorbance decrease at 380 nm. The presence of a sharp isosbestic point at 330 nm indicates that only two species coexist in the equilibrium. The individual profile of the ratio of the two absorbencies  $A_{300}/A_{380}$  demonstrates a 1:1 stoichiometric complexation, with the association constant ( $\log K_{ass}$ ) being calculated as  $2.55 \pm 0.03$ .

Upon the addition of adenosine 5'-monophosphate (AMP) and adenosine diphosphate (ADP) instead of ATP, no obvious spectral changes were found under the same experimental conditions, indicating that the affinity of **MC-1** for ATP was stronger than that for the other adenosine-based nucleotides. In the meantime, the addition of other ribonucleotides (XNP, where X = U (uridine), C (cytidine), and G (guanosine), N = mono-, di-, and tri-, and P = phosphate) could not cause any significant spectral changes of the UV-vis spectra under the same experimental conditions. Compared to the selectivity of ATP with respect to the less charged ADP, AMP, or inorganic phosphate anions, the selective recognition and sensing of ATP over the other triphosphate nucleotides is a much more difficult goal. The excellent selectivity of **MC-1** for ATP over the other ribonucleotides thus not only suggests that the receptor molecule **MC-1** exhibits size-selective recognition for nucle-



**Figure 1.** Family of UV-vis spectra of **MC-1** ( $10 \mu M$ ) in 8:2 DMF/ $H_2O$  aqueous solutions upon the addition of ATP. The inset shows the Benesi-Hildebrand fitting of the titration curve, where  $I$  and  $I_0$  are the ratio of the absorbance at 300 and 380 nm ( $A_{300}/A_{380}$ ) in the presence and absence of ATP, respectively.



**Figure 2.** ESI-MS of **MC-2** in a DMF solution (a) and upon the addition of 1 equiv of ATP (b). The insets exhibit the measured and simulated isotopic patterns at  $m/z = 1017.12$  and  $1327.24$  in parts a and b, respectively.

otides via electrostatic interactions between the cationic binding sites of the receptor and the negatively charged polyphosphate groups but also indicates that the amide groups in **MC-1** situate in the correct place and are able to conceivably interact with adenosine, from which the chromomolecular response is achieved.<sup>11c</sup>

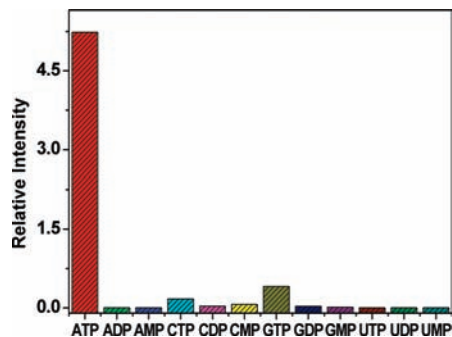
To further investigate the potential interaction sites, a fluorescent dansyl unit was introduced into another meta position, giving a new ligand,  $H_2L^2$ , which has two kinds of optical responding groups and two kinds of guest binding amide groups.<sup>15</sup> The cobalt(II) compound **MC-2** was isolated by a similar synthetic procedure using  $H_2L^2$  instead of  $H_2L^1$ . The electrospray ionization mass spectrometry (ESI-MS) spectrum of **MC-2** in a DMF solution exhibited three intense peaks at  $m/z = 678.40$ ,  $1017.12$ , and  $1090.11$ , respectively. These signals were assignable to  $[Co_3(H_3L^2_3)]^{3+}$ ,  $[Co_3(H_2L^2_3)]^{2+}$ , and  $[Co_3(H_3L^1_3)(PF_6)]^{2+}$ , respectively, reflecting the formation of a [3 + 3] triangle species and the substantial stability of **MC-2** in solution. In the presence of 1 mol equiv of ATP (Figure 2), the peak at  $m/z = 1090.11$  disappeared, indicating that the  $PF_6^-$  anion was replaced. In the meantime, two new peaks at  $m/z = 885.12$  and  $1327.24$  appeared. They were assigned to  $[Na[Co_3(H_4L^2_3) \supset ATP](DMF)(H_2O)]^{3+}$  and  $[Na[Co_3(H_3L^2_3) \supset ATP](DMF)(H_2O)]^{2+}$ ,

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(14) Crystal data of **MC-1**:  $C_{60}H_{60}Co_3F_{18}N_{18}O_{12}P_3$ ,  $M = 1834.27$ , hexagonal, space group  $R\bar{3}c$ , red block,  $a = 18.28(1) \text{ \AA}$ ,  $c = 90.71(5) \text{ \AA}$ ,  $V = 26250(16) \text{ \AA}^3$ ,  $Z = 3$ ,  $D_c = 1.394 \text{ g cm}^{-3}$ ,  $\mu(Mo K\alpha) = 0.716 \text{ mm}^{-1}$ ,  $T = 293(2) \text{ K}$ , 5049 unique reflections [ $R_{int} = 0.1177$ ]. Final  $R1$  [with  $I > 2\sigma(I)$ ] = 0.0814,  $wR2$  (all data) = 0.2092. CCDC number 678275.

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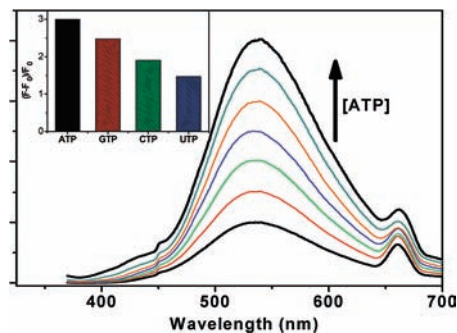


**Figure 3.** UV-vis responses of **MC-2** ( $10 \mu\text{M}$  in a 8:2 DMF/ $\text{H}_2\text{O}$  solution) upon the addition of ribonucleotide polyphosphates ( $0.4 \text{ mM}$ ). The bars represent the value of  $[(I - I_0)/I_0]$ , where  $I$  and  $I_0$  are the ratio of the absorbances at 300 and 380 nm ( $A_{300}/A_{380}$ ) in the presence and absence of nucleotide polyphosphates, respectively.

respectively, based on the exact comparison of these peaks with simulations on the basis of natural isotopic abundances, demonstrating the recognition of **MC-2** for ATP and providing evidence for a 1:1 stoichiometric host-guest complexation.

The UV-vis spectral properties of ligand  $\text{H}_2\text{L}^2$  and **MC-2** were similar to those of  $\text{H}_2\text{L}^1$  and **MC-1**, respectively. Also, **MC-2** exhibited a similar UV-vis spectral change in a DMF/water (8:2) aqueous solution upon the addition of ATP with a sharp isosbestic point at 330 nm, indicating the reversibility of the sensing of **MC-2** for ATP. The individual profile of the ratio of the absorbance  $A_{300}/A_{380}$  also demonstrated the 1:1 stoichiometric host-guest complexation behavior, with an association constant ( $\log K_{\text{ass}}$ ) being calculated as  $3.86 \pm 0.03$ . From the structure of **MC-1**, we reasoned that the three dansyl groups in **MC-2** should position in the same direction showing a calix[3]arene fashion, leading to a 10-fold enhancement of the affinity for ATP. Interestingly, the addition of other ribonucleotides (XNP, where X = A, U, C, and G, N = mono-, di-, and tri-, and P = phosphate) could not cause any significant spectral changes of the UV-vis spectra under the same experimental conditions (Figure 3). Because **MC-1** and **MC-2** exhibited almost the same UV-vis spectral changes upon the addition of ATP, it was reasonable to assign the spectral response to the formation of hydrogen bonds between the amide groups and the ATP guests. The response of **MC-1** and **MC-2** toward ATP varied slightly over a wide acid/base concentration range (Figures S10 and S12 in the Supporting Information). Also, the addition of sodium trisphosphate to the two hosts did not cause obvious UV-vis spectral changes (Figure S13 in the Supporting Information).

Furthermore, the presence of dansylsulfonamide groups allows the host-guest property of **MC-2** to be evaluated also by luminescent analysis. Compound **MC-2** exhibits a typical emission of the dansylsulfonamide group at about 530 nm in the DMF/ $\text{H}_2\text{O}$  aqueous solution ( $50 \mu\text{M}$ , excitation at 330 nm).<sup>16</sup> Upon the addition of 5 equiv of an ATP aqueous



**Figure 4.** Family of luminescence spectra of **MC-2** ( $50 \mu\text{M}$  in a 8:2 DMF/water solution) upon the addition of a standard aqueous solution of ATP. The inset shows the luminescence responses of **MC-2** ( $50 \mu\text{M}$  in a 8:2 DMF/water solution) upon the addition of ribonucleotide triphosphates ( $0.25 \text{ mM}$ ). Data are recorded at 530 nm; excitation is at 330 nm.

solution, the luminescence intensity exhibits a 4-fold enhancement, with the emission wavelength not being shifted (Figure 4). The 1:1 stoichiometric host-guest complexation was also confirmed by the individual titration profile, with an association constant ( $\log K_{\text{ass}}$ ) being calculated as  $4.08 \pm 0.03$ , which is comparable to that calculated by the UV-vis titration data. Clearly, the recognition of **MC-2** to ATP should include the contribution of the hydrogen-bonding interaction about the sulfonamide groups, which block the photoinduced electron transfer process from the amide groups to the dansyl groups, resulting in luminescence enhancement.

Interestingly, the presence of the other ribonucleotide triphosphates, CTP, GTP, and UTP, could cause similar luminescence enhancement of the dansylsulfonamide groups, but the addition of ribonucleotide monophosphate did not lead to any luminescence enhancements under the same experimental conditions. Thus, the fluorescence response is size-selective for nucleotides via hydrogen bonds between the sulfonamide groups and the negatively charged polyphosphate groups. From a mechanism viewpoint, the amide groups that are fixed at the inflexible backbone of the Werner-type macrocyclic receptors can respond to ATP through hydrogen-bonding patterns between the amide group and the nucleoside. However, the size-selective interactions between the negatively charged polyphosphate groups and the cationic triangle receptor provided the opportunity to distinguish the triphosphates from other polyphosphates but did not prompt any chromogenical responses. Thus, the UV-vis responses of both **MC-1** and **MC-2** for ATP exhibited high selectivity over all other ribonucleotides. Whereas these sulfonamide groups interacted with the polyphosphate groups, the luminescence enhancements were quite similar for all four ribonucleoside triphosphates.

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**Supporting Information Available:** CIF file of crystal data of **MC1**, experimental details, and additional spectroscopic data. This material is available free of charge via the Internet at <http://pubs.acs.org>.

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