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ALLOSTERIC EFFECTS IN A SYNTHETIC RECEPTOR FOR ALKALI METAL IONS AND DINITROBENZENE

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A novel host is described which contains three potential binding sites, viz. a cleft and two crown ether rings. The molecule can adopt three different conformations. One of these is able to form a complex with two alkali metal ions. The binding of potassium ions is shown to be a cooperative process. The association constant of the second ion is 128 times larger than that of the first. The compound also binds 1,3-dinitrobenzene at the third site by \( \pi-\pi \) interactions. The complex with two potassium ions binds 1,3-dinitrobenzene more strongly than does the free compound by a factor of 2–6, depending on the solvent used. This allosteric effect is caused by the conformational change which is induced by the binding of the potassium ions.

INTRODUCTION

Many biological processes and functions in nature are regulated by molecular systems which respond to changes in the environment or to outside stimulants. Very often these systems contain receptors or enzymes whose binding properties are controlled by interaction with light, small molecules or ions. Mimicking such systems is one of the most challenging goals in host–guest chemistry. Moreover, understanding the underlying principles of these interactions will be an important step towards the construction of novel molecular devices.

A number of examples of 'responsive hosts' have been described. Crown ethers that change the binding strength of ions on irradiation with light are probably the best studied examples within this class of molecules. These crown ethers contain a photoreactive functionality, e.g., a diaza group, which induces a change in conformation of the host by interaction with light and hence a change in affinity of the binding site. Beer and others have prepared redox-responsive hosts which contain an electrochemically switchable functionality, such as a quinone or a ferrocenyl group. Other examples are pH-responsive hosts and temperature-responsive hosts. Some of these molecules have been used to realize ion transport across liquid membranes, induced by, e.g., light or a pH gradient.

A very important way of regulating binding in receptors and proteins in nature is the allostERIC effect. In an allostERIC protein more than one binding site is present. Binding of a guest (substrate in biochemical terminology) at one site influences the binding strength at a remote site. The paradigmatic example of an allostERIC protein is haemoglobin, a tetrameric protein, that can bind four molecules of oxygen. Binding of the fourth molecule of oxygen is stronger than binding of the first. If the binding sites in an allostERIC protein are identical, the allostERIC effect is described as cooperative binding between homologous sites. Depending on whether the binding of successive guests is increasingly stronger or weaker, it is called positive or negative cooperativity.

Few synthetic receptors displaying allostERIC effects have been described. One of the first successful ones in this respect was reported by Rebek, et al., who synthesized bis-crown ethers capable of binding two alkali metal ions or two mercury complexes such as Hg(CN)\(_2\). For the latter guest the second binding constant was shown to be about ten times larger than the first, whereas for the alkali metal ions mutual electrostatic repulsion prevented cooperative binding of the second guest.

Only recently have organic chemists succeeded in developing synthetic receptors that show allostERIC effects or cooperative binding for organic guests. A tetracarboxylic acid has been reported by Ebmeyer and Rebek which binds two diamines in a cooperative fashion. Schneider and Ruf developed a water-soluble host which binds aromatic guests 10–100 times more strongly when copper ions are present.

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Elsewhere we have reported on the conformational properties of host compounds of type 1. These molecules are receptors for aromatic guests. In solution, compound 1 exists in three conformations, ss, sa and aa (Figure 1). The most predominant conformations, ss and sa, are non-binding. When an aromatic guest is added the equilibrium shifts to aa, and the guest is bound in a sandwich-like fashion between the naphthalene walls of the host. As an extension of this work, we describe here a bis-aza-crown ether derivative of 1 which displays allosteric binding properties (compound 2).

Space-filling models indicate that only the aa conformer of 2 is able to bind an alkali metal ion. We expected that addition of alkali metal ions would promote binding of an aromatic guest, as depicted schematically in Figure 2. We also expected that the binding of the alkali metal ions would be a cooperative process, because conversion of the receptor to the aa conformation by the uptake of one ion would facilitate the binding of the second ion. Experiments will be presented that confirm these ideas.

RESULTS AND DISCUSSION

Synthesis and characterization

The bis-aza-crown ether derivative 2 was synthesized via the route in Scheme 1. Starting compound 4 was refluxed in 1,2-dichloroethane with 2,7-dihydroxynaphthalene using SnCl₄ as a catalyst to give 1a (64% yield). Compound 1a was alkylated with the tosyl ester of chloroethoxyethanol to give 3 in 41% yield. Intermediate 3 was converted into 2 by reaction with 2·2 equiv. of benzylamine in acetonitrile in the presence of sodium iodide using sodium carbonate as a base. After column chromatography, pure 2 was isolated in 91% yield.

In CDCl₃ solution, both 2 and 3 are present as a mixture of sa, ss and aa conformers. For 3, the ratio of these conformers is 82 : 12 : 6. In the solution of 2 the ratio of sa and ss conformers is 88 : 12. The aa con-

![Figure 1. Computer-generated structures of the three conformers of 1](image)

![Figure 2. Induced binding of a guest in 2 by the addition of a metal ion](image)

Scheme 1. (a) 2,7-Dihydroxynaphthalene, SnCl₄, C₂H₅Cl, reflux; (b) TosO(CH₂)₂O(CH₂)₂Cl, DMSO, KOH, 20 °C; (c) PhCH₂NH₂, Na₂CO₃, NaI, acetonitrile, reflux
Table 1. \(^1\)H NMR signals (ppm) and coupling constants (Hz, in parentheses) in the \(^1\)H NMR spectrum of the ss, sa and aa conformers of 2 and 3 in CDCl₃

<table>
<thead>
<tr>
<th>Compound</th>
<th>NCH₂Ar syn</th>
<th>NCH₂Ar anti</th>
<th>Napht-H syn</th>
<th>Napht-H anti</th>
<th>Crown CH₂N</th>
<th>Benzyl CH₂N</th>
<th>Napht-OCH₂</th>
<th>Benzyl-H</th>
</tr>
</thead>
<tbody>
<tr>
<td>2-sa</td>
<td>4·79</td>
<td>6·12</td>
<td>7·23</td>
<td>7·75</td>
<td>3·00</td>
<td>4·00</td>
<td>4·60</td>
<td>7·43</td>
</tr>
<tr>
<td>(14·8)</td>
<td>(16·8)</td>
<td>(8·9)</td>
<td>(8·9)</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>2-ss</td>
<td>5·18</td>
<td>3·33</td>
<td>6·83</td>
<td></td>
<td></td>
<td>a</td>
<td>4·35b</td>
<td>(7·2)</td>
</tr>
<tr>
<td>(14·8)</td>
<td></td>
<td></td>
<td>(8·9)</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>3-sa</td>
<td>4·79</td>
<td>6·03</td>
<td>7·23</td>
<td>7·73</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>(14·8)</td>
<td>(16·8)</td>
<td>(8·9)</td>
<td>(8·9)</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>3-ss</td>
<td>5·13</td>
<td>6·20</td>
<td>7·25</td>
<td></td>
<td></td>
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<td></td>
</tr>
<tr>
<td>(14·8)</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>3-aa</td>
<td>3·92</td>
<td>5·78</td>
<td>7·34</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>(16·8)</td>
<td></td>
<td></td>
<td>(8·9)</td>
<td></td>
<td></td>
<td></td>
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<td></td>
</tr>
</tbody>
</table>

\(^a^\) No assignment could be made.  
\(^b^\) Tentative assignment.

The former is not detectable in the \(^1\)H NMR spectrum and hence constitutes less than 2% of the population. Because of the presence of the conformers, the spectra of 2 and 3 are complicated, but most peaks could be assigned by comparison with the spectra of 1b and with the aid of 2D NOESY and COSY spectra. Apart from cross-peaks due to the nuclear Overhauser effect, the NOESY spectra of 2 and 3 also contained cross-peaks due to conformational exchange processes between the conformers. The positions of the most important peaks are given in Table 1.

Complexation

Alkali metal ions

The effect of alkali metal ions on the conformational equilibrium of 2 was investigated by \(^1\)H NMR in CDCl₃–DMSO-d₆ (9:1,v/v) using 2 mM solutions of host and 8 mM potassium thiocyanate or sodium thiocyanate. Under these circumstances virtually all molecules of 2 were found to be in the aa conformation (Figure 3). The spectrum of the sodium thiocyanate complex is different from that of the potassium thiocyanate complex, as can be seen in Figure 3 and Table 2. In the complex with sodium ions the signals of the naphthyl groups and those of the benzyl groups have moved upfield compared with their positions in the complex with potassium. The signal of the CH₂N- protons in the crown ether moieties of 2 is a triplet in the potassium complex, but is split into two multiples in the sodium complex. These differences suggest that in the latter complex, the crown ether moieties are folded.

Figure 3. \(^1\)H NMR spectra of 2 mM solutions of 2 in CDCl₃–DMSO-d₆ (9:1, v/v) (a) in the absence of salt, (b) in the presence of 8 mM KSCN and (c) in the presence of 8 mM NaSCN.
around the sodium ions in such a way that the benzyl groups are in the shielding zone of the naphthyl groups, or perhaps are complexed in the cleft of the host.

As host 2 has two binding sites for alkali metal ions, it was expected that these ions would be bound in a $1:2$ host–guest stoichiometry. In order to ascertain this and to find out if 2 showed cooperative binding towards these ions, a $^1$H NMR titration with potassium picrate in CDCl$_3$-DMSO-$d_6$ (3:1, v/v) was performed. During this titration the fraction of molecules in the $aa$ conformation was monitored as a function of the number of equivalents of salt added (Figure 4).

The titration data were evaluated with the help of a computer program that determines the goodness of fit $^{12}$ (expressed as $\chi^2/\nu$) of a theoretical binding curve to the experimental data for any combination of $K_1$ and $K_2$. The latter constants $K_1$ and $K_2$ are defined as

$$K_1 = \frac{[2 \cdot M^+]}{[M^+] [2]} \quad K_2 = \frac{[2 \cdot 2M^+]}{[2M^+] [M^+]}$$

In order to obtain intrinsic binding constants from these thermodynamic constants, a statistical correction must be applied which takes into account the number of binding sites in the receptor. $^{13}$ For a receptor with two binding sites, $K_1$ must be divided by 2 and $K_2$ must be multiplied by 2: $K_1' = K_1/2$; $K_2' = 2K_2$.

In the calculations the assumption was made that only the $aa$ conformer of 2 binds ions. The fact that 2 is completely converted into the $aa$ conformer by addition of alkali metal salts shows that this assumption is valid.

For a number of values of $K_2'/K_1'$, the value of $K_1'$ that gives the best fit to the experimental data was calculated. The results of the calculations are shown in Figure 4. For independent binding sites the best fit curve [Figure 4(a)] has a curvature which is too strong and at higher saturation the calculated values of the $aa$ fraction are too low. A much better fit is obtained when a strongly cooperative binding process is assumed [Figure 4(b)]. The lowest value of $\chi^2/\nu$ is obtained for $K_2'/K_1' = 128$ ($K_1' = 43$; $K_2' = 5440$).

A number of models have been proposed to explain cooperative binding by allosteric interactions, one of the first being that of Monod, Wyman and Changeux (MWC). $^{14}$ In this model, the following properties are considered to be required to obtain positive cooperativity: the receptor system should contain identical binding sites at identical, symmetry-related positions; the receptor should have more than one conformation, and these conformations are in equilibrium and differ in their affinity to the substrate; and the symmetry of the receptor should be preserved on going from one conformation to another.

The present system is an excellent example of cooperative binding according to the MWC model. It displays all the properties which are required for the cooperative binding of potassium ions.

It is of interest to compare our receptor 2 with Ebmeyer and Rebek's bis-crown ether receptor $^9$ (see formula 5). In 5 cooperative binding is also the result of binding-induced conformational changes. Ebmeyer and Rebek found that the cooperativity is completely based on an entropy effect: binding of one guest in the

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### Table 2. Relevant signals (ppm) and coupling constants (Hz, in parentheses) in the $^1$H NMR spectra of the potassium and sodium thiocyanate complexes of 2 in CDCl$_3$-DMSO-$d_6$ (9:1, v/v)

<table>
<thead>
<tr>
<th>Complex</th>
<th>$NCH_2Ar$</th>
<th>Napht-$H$</th>
<th>Crown-$CH_2N$</th>
<th>Benzyl-$CH_2N$</th>
<th>Napht-$OCH_2$</th>
<th>Benzyl-$H$</th>
</tr>
</thead>
<tbody>
<tr>
<td>$2 \cdot 2K^+$</td>
<td>5.80; 4.14 (16-5)</td>
<td>7.66; 7.22 (8-9)</td>
<td>2.92 (t)</td>
<td>3.92</td>
<td>4.59</td>
<td>7.26 (br)</td>
</tr>
<tr>
<td>$2 \cdot 2Na^+$</td>
<td>5.93; 4.14 (16-8)</td>
<td>7.45; 6.97 (8-9)</td>
<td>2.93 (m)</td>
<td>4.29</td>
<td>4.68</td>
<td>7.0-7.3 (br)</td>
</tr>
</tbody>
</table>

---

Figure 4. Best fit curves to data points of titration of 2 with potassium picrate. (a) Two independent binding sites, $K_1' = K_2' = 140$; (b) strongly cooperative binding, $K_1' = 43$, $K_2'/K_1' = 128$.
first binding site freezes the conformational freedom of the second binding site, resulting in increased affinity for the second guest. A similar entropy effect may be operative in our receptor 2, but enthalpy effects probably also play an important role. The ion-binding conformer constitutes less than 2% of the total mixture of conformers in a solution of the free host 2. The free energy needed to convert 2 into the binding conformer has to be paid for completely in the first binding step. This fact alone could explain a difference in binding strength of a factor of 50 between the two binding sites. Another important difference is that in Ebmeyer and Rebek’s receptor the distance between the two binding sites is smaller than in our receptor. As a result the electrostatic repulsion between the two bound ions in 4 is much larger than in 2. Consequently, 4 does not show cooperativity in the binding of alkali metal ions, but only in the binding of neutral mercury compounds such as Hg(CN)₂.

**Ternary complexes**

The allosteric effect of potassium ions on the binding of 1,3-dinitrobenzene (DNB) in 2 was qualitatively assessed by determining the complexation-induced shifts of the protons of DNB on the addition of 2, both in the presence and absence of KPF₆. If 27·5 mM 2 was added to a 19·8 mM solution of DNB in CHCl₃-DMSO (3:1, v/v), the DNB proton signals shifted upfield by 0·03, 0·06 and 0·07 ppm for the H-2, H-4,6 and H-5 protons, respectively. In the presence of an excess of KPF₆ these induced shifts increased to 0·06, 0·14 and 0·19 ppm, respectively. In a separate experiment it was ascertained that the signals of DNB do not shift in the presence of KPF₆ alone.

For a quantitative evaluation of the allosteric effect, a series of UV titrations were performed in two solvent mixtures, viz. CHCl₃-DMSO (3:1, v/v) and CHCl₃-DMSO (9:1, v/v). In these titrations the increase in absorption of the charge-transfer band around 400 nm was monitored as a function of the concentration of DNB while the concentration of 2 was varied. Titrations were performed with both sodium and potassium thiocyanate salts. The results of the titration experiments are given in Table 3.

<table>
<thead>
<tr>
<th>Host</th>
<th>CHCl₃-DMSO (9:1, v/v)</th>
<th>CHCl₃-DMSO (3:1, v/v)</th>
</tr>
</thead>
<tbody>
<tr>
<td>2</td>
<td>1·2 ± 0·3</td>
<td>0·3 ± 0·15</td>
</tr>
<tr>
<td>2·2K⁺</td>
<td>7·2 ± 1</td>
<td>0·5 ± 0·12</td>
</tr>
<tr>
<td>2·2Na⁻</td>
<td>1·1 ± 0·3</td>
<td></td>
</tr>
</tbody>
</table>

CHCl₃-DMSO (9:1, v/v). Although sodium ions are bound by 2 [see Figure 3(c)], they are completely ineffective in increasing the binding strength of 2 towards DNB. One possible reason may be that in the sodium complex the benzyl groups are occupying the cleft of the host, thus disfavouring the complexation of DNB. Another explanation may be that in this complex the naphthyl walls are at too close a distance to allow sandwiching of a DNB molecule.

**EXPERIMENTAL**

17b,17c-Dihydro-1,6,10,15-tetraydroxy-17b,17c-diphenyl-7H,8H,9H,16H,17H,18H-7a,8a,16a,17a-tetraazapentaleno[1",6":5,6,7;3",4":5,6,7"] dicycloocta[1,2,3-de:1',2',3'-d'e'] dinaphthalene-8,17-dione (1a). A mixture of 2·48 g (5·08 mmol) of 4, 3·2 g (20 mmol) of 2,7-dihydroxynaphthalene and 5·5 ml (44 mmol) of SnCl₄ was refluxed for 30 min in 100 ml of C₂H₂Cl₂. After refluxing with dilute HCl the product was isolated from the reaction mixture by filtration and was washed with methanol. The product was purified by recrystallisation from DMSO. Yield: 2·1 g (64%) of colourless needles. FAB-MS, m/z 663 [M + H]⁺. Analysis calculated for C₉₀H₃₀Na₂O₆: 0·5H₂O, C, 71·53, H 6·465, N 8·34; found, C, 71·53, H 6·43, N 8·37%.

17b,17c-Dihydro-1,6,10,15-tetraydroxy-17b,17c-diphenyl-7H,8H,9H,16H,17H,18H-7a,8a,16a,17a-tetraazapentaleno[1",6":5,6,7;3",4":5,6,7"] dicycloocta[1,2,3-de:1',2',3'-d'e'] dinaphthalene-8,17-dione (1b). Compound 1a (1·32 g, 1·99 mmol) was dissolved in 25 ml of DMSO. The solution was degassed and placed under a nitrogen atmosphere. Dimethylsulphate (1·4 ml, 8·3 mmol) and powdered KOH (2·5 g) were added and the mixture was stirred at room temperature for 1 h. The reaction mixture was poured into 250 ml of water. The product was extracted from the aqueous suspension with CH₂Cl₂ and purified by column chromatography [chloroform-methanol (99:5:0·5, v/v)]. Yield: 1·09 g (76%) of 1b. 1H NMR (CDCl₃) (only the signals of the sa conformer are given here), δ 7·75, 7·23, 7·20 and 6·84 (4 d, 8 H, napht-H, J = 8·7 Hz), 7·05-6·30 (m, 10 H, ArH), 6·05, 5·97, 4·81 and 4·14...
A 3·71 mm stock solution of 2 in CDCl₃–DMSO-d₆ (3:1, v/v) was prepared (stock solution A). From this solution, a solution 3·71 mm in 2 and 7·74 mm in potassium picrate was prepared (stock solution B). For each data point in the titration, a precisely weighed amount of solution B was made up to 0·8 g with solution A. The last data point in the titration was obtained from solution A which had been saturated with solid potassium picrate. For the determination of the fractions of each conformer, the integral of as many non-overlapping peaks as possible was used.

Computer program for the evaluation of the association constants from the ¹H NMR titration data. A computer program was written in FORTRAN, which calculates for a range of values of K₁ and K₂/K₃, the fractions of molecules of 2 that are present as the 1:1 and 1:2 complex at each data point in the point. The program subsequently evaluates the goodness of fit (χ²/ν) for each combination of K₁ and K₂/K₃, with the help of the expression

$$\frac{\chi^2}{\nu} = \sum \frac{X_{aa\, \text{obs}} - X_{aa\, \text{calc}}}{\sigma^2(n_{exp} - 2)}$$

where Xₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐₐ¢°
REFERENCES


