

Synthesis and Binding Studies of Bowl-Shaped Hosts for Quaternary Ammoniums

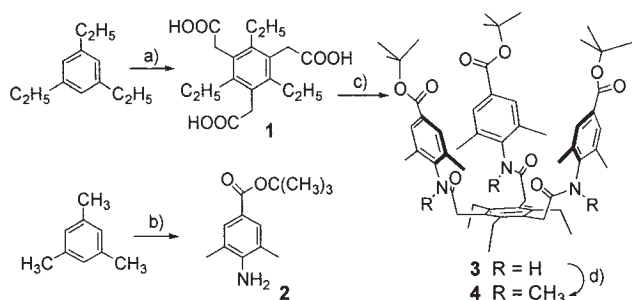
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Two bowl-shaped hosts for quaternary ammonium salts were synthesized and their binding properties, along with the anion effect, were systematically examined and compared in CDCl₃.

A large number of artificial hosts for quaternary ammoniums (QA's) have been studied for the last decade, and provided much experimental evidence for the nature of noncovalent forces, especially the cation- π interaction.^{1,2} The hosts contain a rigid cavity for the QA binding surrounded by aromatic surfaces, like in calixarenes and cyclophanes.² Because QA salts exist as ion pairs in organic solvents, the coexisting counteranions have a significant influence on the binding of QA to the host. The electrostatic interaction of the ion pair may have either a negative or a positive effect on the QA binding as recently demonstrated by several groups including ours.³ We herein prepared two new bowl-shaped hosts for QA and investigated systematically their binding properties with QA salts in CDCl₃.

The basic scaffold of the hosts is a hexasubstituted benzene that has been widely used as a molecular building block for the construction of artificial hosts.⁴ The synthesis of hosts is outlined in Scheme 1. 2,4,6-Triethyl-1,3,5-tris(acetic acid) (**1**)⁵ and *tert*-butyl 4-amino-3,5-dimethylbenzoate (**2**)⁶ were prepared from commercially available 1,3,5-triethylbenzene and 1,3,5-trimethylbenzene, respectively. After reaction with oxalyl chloride, the acid **1** was coupled with the amine **2** to give the N-H host **3**, which was in turn converted into N-Me host **4**.⁷



Scheme 1. a) i) HBr/(CH₂O)_n/ZnBr₂, 42% ii) NaCN/EtOH-H₂O, 95% iii) 50% aqueous H₂SO₄, 72%, b) i) fuming HNO₃/CH₃COOH, 97% ii) CrO₃-H₂SO₄/H₂O-CH₃COOH, 14% iii) SOCl₂, then *t*-BuOH-DMAP/CHCl₃, 83% iv) H₂, Pd-C/MeOH, 96%, c) (COCl)₂, then **2**/DMAP(cat), 45%, d) MeI-KOH/DMSO, 71%.

According to molecular modeling studies (Tripos, Sybyl 6.3), the hosts present the following features. First, three benzoate walls are all positioned to the same side as the result of alternative up and down conformation of adjacent substituents in the central benzene. Second, 3,5-dimethyl substituents of the benzoate walls enforce the aryl plane to be perpendicular to the amide plane and consequently the cavity becomes surrounded by the aryl surfaces.

Finally three amide carbonyl groups are directed to the cavity of the host in a convergent way (Figure 1).

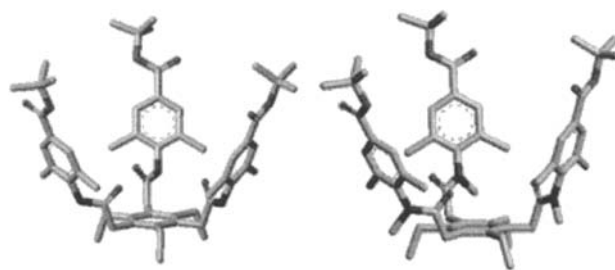


Figure 1. Energy-minimized (Tripos, Sybyl 6.3) structures of the N-H host **3** (left) and the N-Me host **4** (right).

The binding studies were first performed with N-H host **3** and benzyltrimethylammonium (BTMA⁺) halides (**5**, X⁻ = Cl⁻, Br⁻, and I⁻). Upon addition of the host **3** (~3 equiv) in CDCl₃, the signals for N⁺CH₃ and N⁺CH₂ of BTMA⁺·Cl⁻ were significantly upfield shifted ($\Delta\delta$ = 1.5 and 1.8 ppm, respectively). This is a strong evidence for the QA guest locating inside the cavity surrounded by aryl surfaces of the host. However, the magnitude of the shifts is highly anion-dependent; $\Delta\delta$ = ~1 ppm for BTMA⁺·Br⁻ and <0.3 ppm for BTMA⁺·I⁻ when equal amounts of the host **3** were added. Quantitative binding affinities were determined by nonlinear least squares fitting of ¹H NMR chemical shift changes observed by diluting a 1 : 1 mixture of host and guest in CDCl₃ at 23 ± 1 °C. The association constants (*K*_a's) were calculated to be >3 × 10⁴, 3.4 (±0.2) × 10³, and 1.7 (±0.2) × 10³ M⁻¹ for BTMA⁺·Cl⁻, BTMA⁺·Br⁻, and BTMA⁺·I⁻, respectively.

As suggested previously,³ this anion dependence may be rationalized by hydrogen bonds between the amide NH hydrogen of the host **3** and the anion X⁻ of the guest.⁸ The hydrogen-bonded anions may exert electrostatic interactions with the oppositely charged ammonium ions, especially in a less polar organic solvent, CDCl₃. Second, the hydrogen bond may increase the electron density on the amide carbonyl oxygens and consequently strengthen the cation-dipole and/or C-H...O interactions.⁹

Next, we have prepared the N-Me host **4** that cannot directly interact with anions through hydrogen bonds. As seen in Figure 1, the energy-minimized structure¹⁰ of the host **4** is very close to that of the N-H host **3**. When the host **4** was added to a CDCl₃ solution of various QA salts, the ¹H NMR signals of the N⁺CH_n hydrogen were gradually upfield shifted ($\Delta\delta$ = 0.3–0.5 ppm), but the degree of the shifts is smaller compared with the N-H host **3**. The association constants are calculated by nonlinear least-square fitting of ¹H NMR titration curves¹¹ and the results are summarized in Table 1. Job's plots also confirm a 1 : 1 complex formation between the host **4** and each of the guests **5**–**8**.¹²

Trends in the binding affinity are as follows. The magnitude

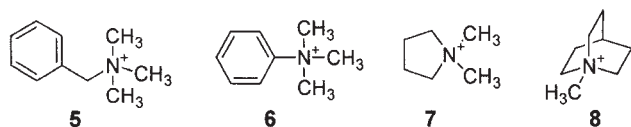


Table 1. Association constants ($K_a \pm 10\%$) between the N-Me host **4** and ammonium guests in CDCl_3 at $23 \pm 0.5^\circ\text{C}$

Guest	K_a/M^{-1}	Guest	K_a/M^{-1}
5 · Cl^-	570	8 · I^-	490
5 · Br^-	1160	$\text{Et}_4\text{N}^+\cdot\text{I}^-$	140
5 · I^-	1780	$\text{Bu}_4\text{N}^+\cdot\text{I}^-$	no binding
6 · I^-	3300	<i>t</i> -BuOH	no binding
7 · I^-	2200		

of the association constants between host **4** and $\text{BTMA}^+\cdot\text{X}^-$ is the order of $\text{Cl}^- < \text{Br}^- < \text{I}^-$, exactly opposite to that seen with the N-H host **3**. This is expected because tightening the ion pair weakens the cation binding to the cavity, like in the host **4** having no hydrogen-bonding site for anion. Second, two hosts **3** and **4** show nearly identical association constant ($\approx 1700 \text{ M}^{-1}$) toward $\text{BTMA}^+\cdot\text{I}^-$, implying that iodide ion is poor hydrogen-bonding acceptor and exerts negligible anion effect. Third, the host **4** binds strongly to the small QA guests but negligibly either to a large guest $\text{Bu}_4\text{N}^+\cdot\text{I}^-$ or to a neutral guest *tert*-butanol.

In conclusion, the coexisting counteranions have a large influence on the QA binding in organic solvents, but the magnitude depends on the nature of the host as well as the anion.

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References and Notes

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- Physical and spectroscopic properties of the N-H host **3**: mp $> 300^\circ\text{C}$; ^1H NMR (250 MHz, CDCl_3) δ 7.67 (br s, 3NH), 7.59 (s, 6H), 3.93 (s, 6H), 2.80 (q, $J = 7.3$ Hz, 6H), 2.13 (s, 18H), 1.53 (s, 27H), 1.29 (t, $J = 7.3$ Hz, 9H); ^{13}C NMR (63 MHz, CDCl_3) δ 169.2, 165.3, 143.1, 137.6, 135.1, 130.8, 130.4, 129.5, 81.2, 28.3, 18.6, 15.0; FABMS $m/z = 968.5$ (7%) for $[\text{M}\cdot\text{Na}]^+$, 944.5 (12%) for $[\text{M}\cdot\text{H}]^+$; Anal. Calcd for $\text{C}_{57}\text{H}_{75}\text{N}_3\text{O}_9$: C, 72.35; H, 7.99; N, 4.44. Found: C 72.38; H, 7.95; N, 4.45. Physical and spectroscopic properties of the N-Me host **4**: mp > 192 – 193°C ; ^1H NMR (250 MHz, CDCl_3) δ 7.79 (s, 6H), 3.17 (s, 6H), 3.14 (s, 9H), 2.36–2.29 (m, 24H), 1.61 (s, 27H), 0.92 (t, $J = 7.2$ Hz, 9H); ^{13}C NMR (63 MHz, CDCl_3) δ 171.1, 165.3, 144.9, 141.1, 136.1, 131.7, 130.4, 129.4, 81.6, 34.7, 28.3, 17.5, 14.3; FABMS $m/z = 988.7$ (100%) for $[\text{MH}]^+$, 987.5 (45%) for $[\text{M}]^+$; Anal. Calcd for $\text{C}_{60}\text{H}_{81}\text{N}_3\text{O}_9$: C, 72.92; H, 8.26; N, 4.25. Found: C 72.96; H, 8.26; N, 4.25.
- The formation of the hydrogen bond is evident in the ^1H NMR spectra. When ≈ 3 equivalents of $\text{BTMA}^+\cdot\text{X}^-$ were added to a CDCl_3 solution of the host (1 mM), large downfield shifts of the NH signal were observed, $\Delta\delta = 2.4$, 1.9, and 0.7 ppm for $\text{X} = \text{Cl}$, Br, and I, respectively. For hydrogen bonding interaction between anions and amide NH^+ 's, see: a) F. P. Schmidtchen and M. Berger, *Chem. Rev.*, **97**, 1609 (1997). b) K. Choi and A. D. Hamilton, *J. Am. Chem. Soc.*, **123**, 2456 (2001).
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- This structure is the energy-minimized one of the host **4** when complexed with tetramethylammonium guest inside the cavity. In the ^1H NMR spectroscopy, the free host **4** contains a small amount ($\approx 5\%$) of conformational isomer in CDCl_3 , but only one set of ^1H NMR signals was observed upon the complex formation. Therefore, the guest binding seems to induce the host structure in a way of maximizing intermolecular noncovalent interactions.
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