## Encapsulation of Et<sub>3</sub>N<sup>+</sup>-H···OH<sub>2</sub> in a hydrogen-bonded resorcarene capsule

## Alexander Shivanyuk,\* Kari Rissanen\* and Erkki Kolehmainen

Department of Chemistry, University of Jyväskylä, PO Box 35, Jyväskylä, FIN-40351, Finland. E-mail: shivan@jyu.fi and kari.rissanen@jyu.fi

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In the crystalline state a resorcarene dimer linked by ten hydrogen-bonding water molecules encapsulates the hydrogen-bonded complex  $Et_3N^+$ -H···OH<sub>2</sub> while bromide anions are positioned outside the cavity.

Supramolecular encapsulation is a topic of considerable current interest.<sup>1</sup> Designed self-complementary molecules form dimeric<sup>2</sup> or tetrameric<sup>3</sup> hydrogen-bonded capsules including charged and neutral guest species. Resorcarenes **1** linked by smaller hydrogen-bonding molecules assemble into dimeric<sup>4</sup> and hexameric<sup>5</sup> hollow structures in the solid state. Molecular encapsulation of dimeric hydrogen-bonded complexes has been discovered recently in apolar solutions<sup>6</sup> however much less is known about such complexation in the crystalline state.<sup>7</sup>

Herein we describe the first single crystal X-ray structure of a hydrogen-bonded molecular capsule encapsulating the hydrogen-bonded heterocomplex  $Et_3N^+$ -H···OH<sub>2</sub>.



Co-crystallization of 1a with a ten-fold excess of Et<sub>3</sub>NHBr from aqueous EtOH results in cubic crystals which, although unstable without solvent, were suitable for single crystal X-ray analysis.† The resorcarene molecules adopt a perfect cone conformation stabilized by four intramolecular hydrogen bonds between neighboring hydroxy groups. Two molecules of 1a, sitting on a two fold crystallographic rotation axis, form a dimer held together by ten hydrogen-bonded water molecules disordered over 16 sites (Fig. 1). One  $2^+$  and one hydrogen-bonded water molecule [N1A...O12a 2.735(7) Å] are encapsulated in such a dimeric cavity.<sup>8</sup> Two opposite orientations occur for the encapsulated  $Et_3N^+$ –H···OH<sub>2</sub> with occupancy factors of 0.7 and 0.3. Furthermore, in each orientation the ethyl groups are strongly disordered owing to the fact that the three-fold axis of  $2^+$  coincides with the two-fold axis of the capsule. Encapsulated  $2^+$  is shifted by 0.28 Å from the center of the cavity<sup>9</sup> towards resorcarene molecule B. Several close non-bonded contacts (3.49–3.67 Å ) were found between carbon atoms of  $\rm Et_3N^+\!\!-\!H$ and resorcinol rings of **1a** which manifest C-H $\cdots$  $\pi$  attractions. The encapsulated water molecule (O12a) is located deep in the resorcarene socket of molecule A forming four close non-bonded contacts to the aromatic C-H atoms of the resorcinol rings with O-C distances of 3.327(4) and 3.335(4) Å.<sup>10</sup> Such an arrangement allows O-H···π interactions between the encapsulated water molecule and the resorcinol rings. The hydrogen bonding in Et<sub>3</sub>N<sup>+</sup>-H···OH<sub>2</sub> should theoretically increase the positive charge on the hydrogen atoms of the water molecule and thus facilitate O–H··· $\pi$  attractions.



**Fig. 1** Single-crystal X-ray structure of  $[(1a_2\cdot 10H_2O)-@(2^{+}\cdot H_2O)]Br^{-}\cdot 4.8H_2O$ . All hydrogen atoms are omitted for clarity. Hydrogen bonds are indicated by dotted lines. The most populated orientation of the disordered cation and water molecule is indicated. The nitrogen atom of  $2^+$  and oxygen atom of encapsulated water molecule are shown with arbitrary radii. Distances between hydrogen-bonding oxygen atoms are between 2.606(3) and 2.791(5) Å. Distances between the hydrogen-bonded Br anion and water molecules are between 3.323(6) and 3.348(5) Å.

The bromide anion is positioned outside the cavity and lies within the shallow pocket formed by four pendant  $CH_2$  groups of molecule  $A^{11}$  and is solvated by four disordered hydrogenbonded water molecules (Fig. 1). The shortest N···Br distance [8.212(6) Å] is *ca.* 2.5 times longer than in  $2^+$  Br<sup>-</sup>.<sup>12</sup> The distance of the bromide anion to the carbon atom of the  $CH_2$  group is 4.004(4) Å reflecting weak C–H···Br<sup>-</sup> interactions. The hydrogen-bonded array of the resorcarene dimer is not saturated and therefore four symmetry related bromide anions are hydrogen bonded to the bridging water molecules (Fig. 1).

<sup>1</sup>H NMR spectroscopy shows that no complex is formed between 1 and  $2^+$  in polar solvents such as methanol-d<sub>4</sub> or DMSO-d<sub>6</sub>. However, in CDCl<sub>3</sub> saturated with H<sub>2</sub>O, lipophilic resorcarenes 1b,c strongly interact with  $2^+ X^- (X^- = Cl^-)$ , Br<sup>-</sup>, ClO<sub>4</sub> <sup>-</sup>). At  $[2^+ X^-]/[1b,c] \ge 0.5$ , <sup>1</sup>H NMR spectra show sharp signals of methyl and methylene protons of the complexed cation which are both shifted upfield by ca. 1.3 ppm (Fig. 2). The exchange of complexed and free  $2^+$  is slow on the NMR timescale at 303 K (500 MHz) and the integration shows that two resorcarene molecules complex one cation  $2^+$ . The signal of the water protons (5-6 H per 1c) is shifted downfield  $(\Delta \delta = 1.9 \text{ ppm})$  and the multiplet of resorcinol OH groups is centered at  $\delta$  9.5 owing to strong hydrogen-bonding. These results suggest that hydrogen-bonded water molecules participate in the formation of the 2:1 capsule also in  $\text{CDCl}_3$ .<sup>13</sup> These results are also in agreement with the fact that in dry CDCl<sub>3</sub> the systems  $1b,c + n 2^+X^-$  exhibit broad <sup>1</sup>H NMR spectra, most probably owing to the formation of ill defined hydrogen-bonded aggregates. Detailed NMR investigations of these complex systems in apolar solvents are currently in progress and will be reported in due course.

In conclusion, the solid-state encapsulation of ammonium cations by a water linked dimer of **1a** is strongly dependent on



**Fig. 2** <sup>1</sup>H NMR spectrum of 2  $\mathbf{1b} + \mathbf{2}^+Cl^- \{[\mathbf{1b}] = 3 \times 10^{-2} \text{ M}, 500 \text{ MHz}, 303 \text{ K}, CDCl_3\}$ . The signals of the complexed cation are: a, NH; b, CH<sub>2</sub>; c, CH<sub>3</sub>.

the structure of the guest. For the previously studied complex with  $Et_4N^{+14}$  no anion is found while the negative charge was distributed over all oxygen atoms of the capsule. The smaller volume of  $Et_3NH^+$  and the presence of the hydrogen bonding N–H group seem to be crucial for the encapsulation of  $Et_3N^+$ –  $H\cdots OH_2$  in the cavity of the resorcarene dimer. The incomplete saturation of the hydrogen-bonded array within the hydrogen bonded shell determines the strong hydrogen-bonding interaction of the capsule with external bromide anions. This feature is being currently explored for the rational design of externally tuned supramolecular containers.

## Notes and references

† *X-Ray crystal structure analysis*: Data were recorded with a Kappa CCD difractometer using graphite monochromatised Mo-Kα radiation [ $\lambda$ (Mo-Kα) = 0.71073 Å]. The data was processed with Denzo-SMN v0.93.0 [Z. Otwinowski and W. Minor, *Processing of X-ray Diffraction Data Collected in Oscillation Mode, Methods in Enzymology*, vol. 276: *Macromolecular Crystallography*, Part A, ed. C. W. Carter, Jr. and R. M. Sweet, Academic Press, New York, 1997, pp. 307–326]. The structures were solved by direct methods (SHELXS-97: G. M. Sheldrick, *Acta Crystallogr., Sect. A*, 1990, **46**, 467) and refinements, based on *F*<sup>2</sup>, were made by full-matrix least-squares techniques (SHELXL-97, G. M. Sheldrick, *A program for crystal structure refinement*, 1997, University of Göttingen, Germany). The hydrogen atoms were placed in calculated idealized positions with isotropic temperature factors (1.2 times the C temperature factor) and refined as riding atoms. The hydrogen atoms of water molecules and disordered cation

were not located. The atoms of the cation in the less populated orientation as well as two strongly disordered water molecules were treated isotropically.

*Crystal data*: [(**1a**<sub>2</sub>·10H<sub>2</sub>O)@(**2**<sup>+</sup>·H<sub>2</sub>O)]Br<sup>-</sup>·4.8H<sub>2</sub>O: crystal size 0.4 × 0.3 × 0.25 mm, tetragonal, space group  $I4_1/a$ , a = 20.2620(5), c = 43.1040(7), V = 17696.3(7) Å<sup>3</sup>, Z = 8,  $D_c = 1.214$  g cm<sup>-3</sup>,  $\mu = 0.542$  mm<sup>-1</sup>,  $2\theta_{\text{max}} = 49.44^{\circ}$ , F(000) = 6824, R1 = 0.0539, wR2 = 0.15875 [for 5042 reflections with  $I > 2\sigma(I)$ ],  $R_{\text{int}} = 0.0525$ , R1 = 0.0792, wR2 = 0.1800 (for all 7537 reflections), 590 parameters, S = 1.106.

CCDC 182/1639. See http://www.rsc.org/suppdata/cc/b0/b002144j/ for crystallographic files in .cif format.

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