

Protonation of electroneutral *p*-*tert*-butylcalix[4]arenetetraacetic acid

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Received: 4 September 2009 / Accepted: 22 October 2009 / Published online: 20 November 2009
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Abstract Using ^1H NMR spectroscopy together with density functional theoretical calculations, it is shown that electroneutral *p*-*tert*-butylcalix[4]arenetetraacetic acid forms an equimolar complex with a proton in the form of the H_3O^+ ion in nitrobenzene- d_5 . Protons were offered by hydrogen bis(1,2-dicarbollyl)cobaltate and converted to hydroxonium ions by traces of water. In the resulting complex, the H_3O^+ cation is bound by strong hydrogen bonds to two phenoxy oxygen atoms of the parent calix[4]arene ligand and to one carbonyl oxygen of the corresponding COOH group of this ligand.

Keywords Calixarenes · Macrocycles · Protonation ·
Ab initio calculations · Complex structure

Introduction

Calix[*n*]arenes are a well-known family of macrocyclic molecules with many potential applications in various branches of chemistry. Because of their simple one-pot preparation, easy derivatization, and unique complexation abilities, calix[*n*]arenes are widely used as building blocks

for the construction of more sophisticated molecular systems. Their unique three-dimensional pre-organization make them very attractive as the receptors for the complexation of cations, anions, and even neutral molecules. Calix[*n*]arenes find applications as selective binders and carriers, as analytical sensors, as catalysts, and as model structures for biomimetic studies [1, 2]. In the field of host-guest chemistry, many studies have focussed on the binding ability of calixarene derivatives with carbonyl groups at their lower rims toward metal ions, predominantly alkali and alkaline-earth, but also transition and heavy metal cations [3–11], and even toward H_3O^+ [12–19].

Recently, deprotonation of calix[*n*]arenes in which $n = 4, 6,$ and 8 has been studied using acetonitrile as the solvent [20]. In this communication, on the basis of ^1H NMR spectroscopy and DFT quantum mechanical calculations, we suggest the most probable structure of protonated *p*-*tert*-butylcalix[4]arenetetraacetic acid.

Results and discussion

NMR measurements

Similarly to our previous studies on protonation of valinomycin [21, 22], we used hydrogen bis(1,2-dicarbollyl)cobaltate (HDCC) as a versatile proton source [23]. The protons were converted to hydroxonium ions H_3O^+ by traces of water in the system under study. In order to ensure full ionization of HDCC we had to use a solvent with a rather high dielectric constant, e.g. nitrobenzene ($\epsilon = 35.6$), acetonitrile ($\epsilon = 36.6$), or dimethylformamide ($\epsilon = 38.3$). Among these, nitrobenzene- d_5 proved to give least equivocal results in our previous works as it does not form inclusion complexes with calixarenes and does not

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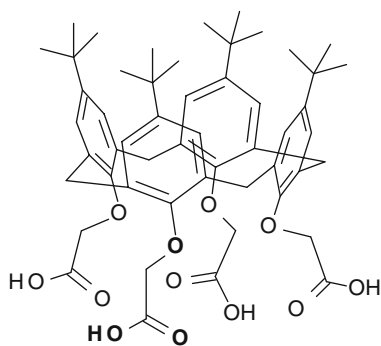
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compete perceptibly with their proton binding. We therefore chose this solvent for use in the present study.

^1H NMR spectra of free electroneutral ligand **1** (Scheme 1) and its different mixtures with HDCC in nitrobenzene- d_5 are depicted in Fig. 1. The spin-echo technique was used in order to remove obscuring broad signals of HDCC, which was present in large excess. In Fig. 1, Ar means aromatic protons, CH_a and CH_e are axial and equatorial protons of the macrocycle CH_2 groups, CH_2 is the signal of carboxyl-bearing CH_2 groups, and *t*-Bu denotes *p*-*tert*-butyl protons. The signal marked by the asterisk corresponds to the protons involved in fast proton exchange between free and bound hydrated protons, water molecules, and protons of COOH groups.

It is necessary to emphasize that the formation of a complex of **1** with the hydrated proton, H_3O^+ , leads to a relative chemical shift of virtually all the signals, indicating a slight change in the conformation of the parent calix[4]arene ligand **1**. In particular, the mutual nearing of the signals of the axial and equatorial protons in the macrocyclic CH_2 groups indicates slight opening of the calix[4]arene cup [24], probably because of the shorter



Scheme 1

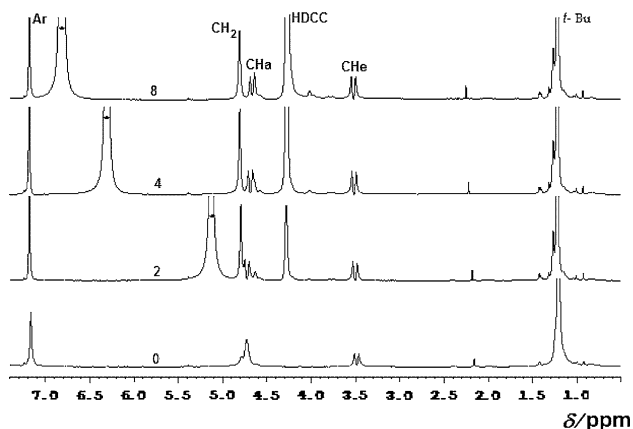


Fig. 1 300.13 MHz ^1H NMR spin-echo spectra of **1** (5×10^{-3} M) and its mixtures with the indicated number of equivalents of HDCC (nitrobenzene- d_5 , 294 K)

distance of the coordinating groups at the lower rim. The increase of the individual shifts between $\beta = 1$ and 10 ($\beta = [\text{H}_3\text{O}^+]_0/[\mathbf{1}]_0$) shows, however, that the equilibrium $\mathbf{1} + \text{H}_3\text{O}^+ \rightleftharpoons \mathbf{1} \cdot \text{H}_3\text{O}^+$ is not entirely on the side of the $\mathbf{1} \cdot \text{H}_3\text{O}^+$ complex. Assuming that the resulting complex is equimolecular ($\mathbf{1} \cdot \text{H}_3\text{O}^+$), the considered equilibrium should follow the usual relationship (Eq. 1):

$$K = \frac{[\mathbf{1} \cdot \text{H}_3\text{O}^+]}{[\mathbf{1}][\text{H}_3\text{O}^+]} \quad (1)$$

Under conditions of fast exchange between free and coordinated **1**, the actual chemical shift of any signal is a weighted average of the shifts corresponding to the exchanging sites. Thus, the actual relative shift is $\Delta\delta = \alpha\Delta\delta_{\text{max}}$, where $\alpha = [\mathbf{1} \cdot \text{H}_3\text{O}^+]/[\mathbf{1}]_0$ and $\Delta\delta_{\text{max}}$, obtained by extrapolation, is the maximum relative shift corresponding to the pure complex. Using Eq. (1), $\Delta\delta$ can be expressed by means of Eq. (2), where β was already defined above:

$$\Delta\delta = \Delta\delta_{\text{max}} \times \frac{1 + K[\mathbf{1}]_0(1 + \beta) - \sqrt{\{1 + K[\mathbf{1}]_0(1 + \beta)\}^2 - 4K^2[\mathbf{1}]_0^2\beta}}{2K[\mathbf{1}]_0} \quad (2)$$

It is evident that both K and $\Delta\delta_{\text{max}}$ can be obtained by fitting Eq. (2) to the experimental values of $\Delta\delta$. Moreover, Fig. 2 presents $\Delta\delta$ of individual signals in the proton spectra of **1**/HDCC mixtures and the fitting curves. As one can see, the fitting is quite good and offers values of K in a relatively narrow range. Thus, the average value of K is $113.6 \pm 0.8 \text{ M}^{-1}$.

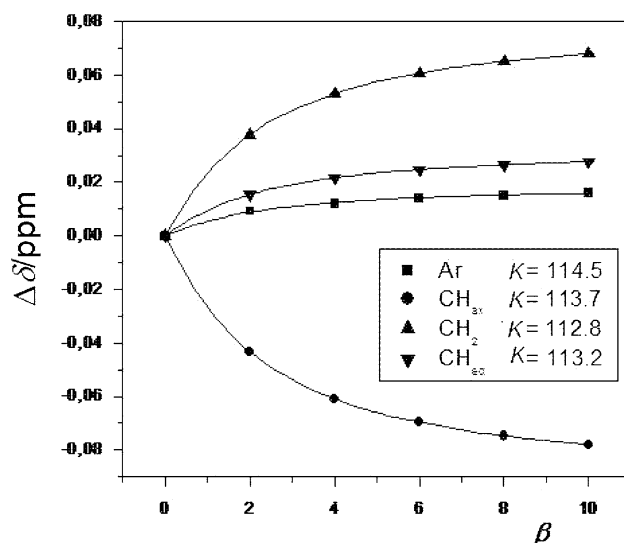


Fig. 2 Experimental relative shifts $\Delta\delta$ of individual signals of **1** in its mixtures with HDCC and the fits obtained using Eq. (2) (the fitted values of K in the graph)

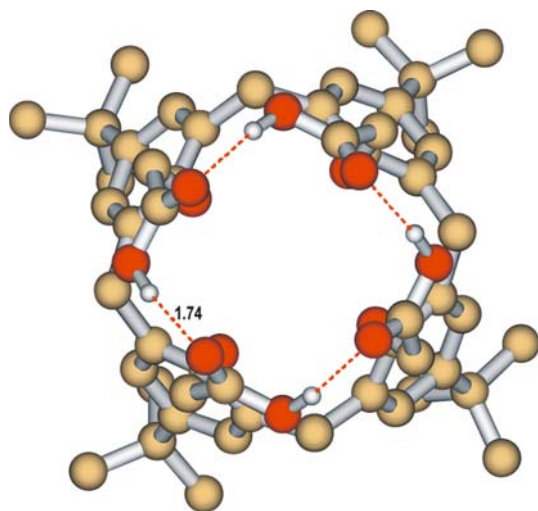


Fig. 3 DFT optimized structure of free ligand **1** (B3LYP/6-31G(d)) (hydrogen atoms omitted for clarity except the four hydrogens taking part in four internal hydrogen bonds between the nearest COOH groups)

Quantum mechanical calculations

The quantum mechanical calculations were carried out at the density functional level of theory (DFT, B3LYP functional) using the Gaussian 03 suite of programs [25]. The 6-31G(d) basis set was used and the optimizations were unconstrained. Although a possible effect of the polar solvent on the detailed structures of **1** and **1**·H₃O⁺ could be imagined, our quantum calculations in similar cases, performed analogously, showed very good agreement of experiment with theory [21, 26–30].

In the model calculations, we optimized the molecular geometry of the parent calix[4]arene ligand **1** and its complex with H₃O⁺. The optimized structure of **1** is depicted in Fig. 3. From this figure it follows that the most

stable conformation of the mentioned ligand **1** is a *cone* structure with C₄ symmetry. It is necessary to emphasize that this structure is supported by four strong internal hydrogen bonds between the nearest COOH groups of free ligand **1**.

In Fig. 4, two projections of the **1**·H₃O⁺ complex obtained by the full DFT optimization are illustrated together with the lengths of the corresponding hydrogen bonds (in Å). It is evident that the *cone* structure of the parent calix[4]arene ligand **1** in the cationic complex species **1**·H₃O⁺ is preserved; however, the four internal hydrogen bonds between the nearest COOH groups are broken in the resulting **1**·H₃O⁺ complex. In this complex, the H₃O⁺ ion is bound by strong hydrogen bonds to two phenoxy oxygen atoms of **1** (1.76 and 1.59 Å) and to one carbonyl oxygen of the respective COOH group of **1** (1.67 Å).

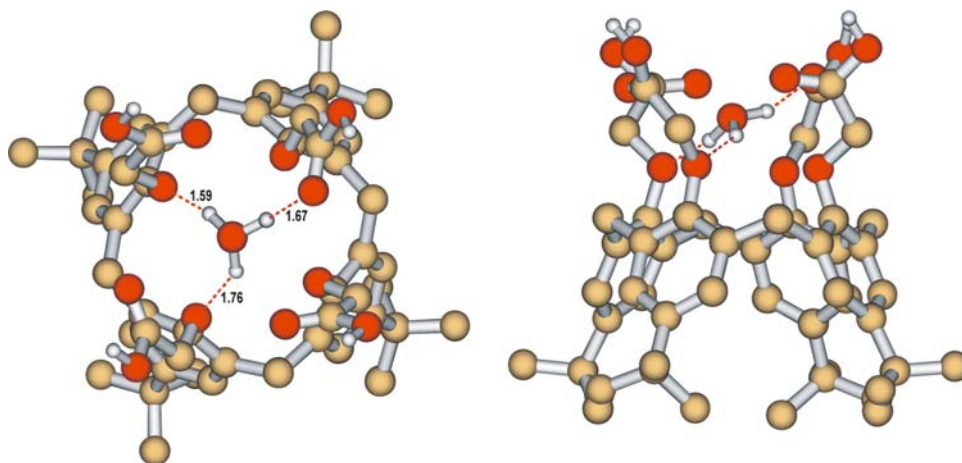
Finally, the calculated binding energy of the **1**·H₃O⁺ cationic species is −368.3 kJ mol^{−1}, which confirms formation of the complex.

Experimental

The electroneutral macrocyclic compound *p*-*tert*-butylcalix[4]arenetetraacetic acid (**1**, Scheme 1) was kindly supplied by Prof. V. I. Kalchenko, Institute of Organic Chemistry, NASU, Kiev, Ukraine. Nitrobenzene-*d*₅ was purchased from Fluka, Buchs, Switzerland. Preparation of hydrogen bis(1,2-dicarbollyl)cobaltate (HDCC) [23] was described elsewhere [21].

¹H NMR spectra in nitrobenzene-*d*₅ were measured in a quadrature detection mode at 300.13 MHz with an upgraded Bruker Avance DPX300 spectrometer collecting 32 kpoints in 128 scans. All NMR measurements were carried out at a temperature of 294 K.

Fig. 4 Two projections of DFT optimized structure of the **1**·H₃O⁺ complex (B3LYP/6-31G(d)) (hydrogen atoms omitted for clarity except the four hydrogens of the corresponding COOH groups and those of H₃O⁺). H-bond lengths of H₃O⁺ to phenoxy oxygens of **1**: 1.76 and 1.59 Å; H-bond length of H₃O⁺ to carbonyl oxygen of **1**: 1.67 Å



Acknowledgments This work was supported by the Academy of Sciences of the Czech Republic, project T 400500402, and by the Czech Ministry of Education, Youth, and Sports, projects MSM 4977751303, MSM 6046137301, and MSM 6046137307.

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