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Binding Studies on the Control of the Conformation and Self-assembly of a Calix[4]arenedicarboxylic Acid through Hydrogen Bonding Interactions

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A convenient method for measuring the dimerisation constant in solution of a calix[4]arenedicarboxylic acid is reported. This compound also co-crystallises with a bis(amidopyridyl)calix[4]arene as independent tubular calixarene stacks that are cross-linked via carboxyl– amidopyridyl hydrogen bonding and π -stacking of the amidopyridyl arms.

Keywords: Calix[4]arene; Hydrogen bonds; Carboxylic acid; Amidopyridine; Crystal engineering

INTRODUCTION

Self-organisation and self-assembly through hydrogen bonds has enabled large, highly-ordered structures to be synthesized with the minimum of synthetic effort [1]. Calix[4]arenes are attractive molecules in this respect since these bowl-shaped molecules can provide appropriate molecular recognition sites through self-organisation. For example, calix[4]resorcinarenes are known to form capsule-like dimers [2–5] and higher aggregates [6–8] through hydrogen bonds involving their OH groups. Such structures can then encapsulate neutral and/or charged guests in the cavities. Calix[4]arenes containing urea groups are also known to form dimers through the formation of intermolecular hydrogen bonds between the urea NH protons and the carbonyl group [9,10]. Recently, calix[4]arenes derivatives containing two [11–15] or four [16] carboxylic acid groups on the upper rim have been shown to form

self-assembled dimers in organic solvents. However, attempts at quantifying the strength of these assemblies have been hampered by the high stability of the dimer with respect to the monomer in nonprotic organic solvents [13]. Here we report a convenient method for measuring dimerisation constants for these assemblies in a competitive solvent mixture using the parent structure, calix[4] arenedicarboxylic acid 1 [11,12]. We also report its interaction with bis(amidopyridyl)calix[4]arene 2 [17] to yield a 1:1 co-crystal that comprises tubular calixarene stacks linked via hydrogen bonds between carboxylic acid and amidopyridine moieties and π -stacking of the amidopyridyl arms.

RESULTS AND DISCUSSION

Compound 1 was prepared as described previously [11]. Crystals of 1 suitable for X-ray diffraction were grown from methanol and reveal the formation of a symmetrical dimer $1₂$, as a result of hydrogen bonds

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FIGURE 1 Crystal structure of the calix[4]arene pinched cone dimer 12

between the carboxylic acid groups $[06 \cdots 07 =$ 2.647(4) \dot{A} , (O6) $H6 \cdot \cdot \cdot O7 = 1.83 \dot{A}$, O6–H $6 \cdot \cdot \cdot O7 =$ 162.4° ; O8 $\cdot \cdot$ O5 = 2.621(4) Å, (O8)H8 $\cdot \cdot$ O5 = 1.80 Å, O8–H8···O5 = 165.5°] (Fig. 1). These distances are very similar to those reported from a crystal structure of a related calix[4]arenedicarboxylic acid dimer containing two nitro groups [13]. The presence of these intermolecular hydrogen bonds results in the structure adopting a pinched cone conformation, where two of the phenyl rings are turned in towards one another [dihedral angle = $21.06(16)°$], whereas the unsubstituted rings point away from one another in an open conformation δ [dihedral angle = $75.51(11)°$].

The $\rm ^1H\text{-}NMR$ spectrum of 1 (2.5 mM) at 25 $\rm ^{\circ}C$ (after heating to $40-50^{\circ}$ C to effect dissolution) in CDCl₃ reveals that this dimeric structure also exists in solution, in agreement with previous studies on this and related compounds [11-15]. Specifically, the signal for the four equivalent protons (H_a) on the two substituted phenyl rings is shifted upfield $(\delta = 6.76$ ppm) relative to the expected chemical shift for these protons in a bis-substituted calixarene containing a symmetrical cone of C_{4v} symmetry (usually >7.0 ppm, e.g. 7.62 ppm for the signal for these protons in compound 2).[†] However, the addition of DMSO- d_6 resulted in a downfield shift in this signal; in CDCl₃ solutions of 1 (2.5 mM) containing 0.5%, 1% and 3% DMSO- d_6 , the H_a signal was observed at 6.78 ppm, 7.26 ppm and 7.42 ppm, respectively. These changes, along with the signals for the six protons on the other two aromatic rings of 1 undergoing upfield shifts upon addition of DMSO- d_6 (up to 3% DMSO in CDCl₃), show that DMSO competes effectively for solvation of the carboxylic acid groups. This causes the equilibrium to shift towards a monomer, resulting in the calixarene ring adopting the usual symmetrical cone conformation (Scheme 1).

In solutions containing higher amounts $($ >3%) of $DMSO-d₆$, no further changes to the NMR spectra were observed. Therefore the signal for H_a at 7.42 ppm is a limiting value and corresponds to the presence of only monomeric 1 in solution. These results gave us the opportunity to measure the dimerisation constant, K_2 , where $K_2 = [\text{dimer}]$ / [monomer]², via a NMR dilution method. An appropriate solvent ratio (1% DMSO- d_6 -CDCl₃) was chosen to enable appreciable amounts of monomer and dimer to be observed over a millimolar concentration range. The resulting data were fitted to the Spurr and Byers equation (see the Experimental section) [18], giving $K_2 = 35 \,\mathrm{M}^{-1}$ at 298 K (error $\langle 5\% \rangle$ from a linear plot (Fig. 2). In this solvent system, the H_a signal moved from 7.00 ppm at 10 mM to the limiting value of 7.42 ppm at 0.65 mM. In accordance with other studies $[13]$, in CDCl₃ alone, no downfield shift in the H_a signal was observed in this concentration range, which indicates that the pinched cone dimer 1_2 is by far the predominant species in solution in the absence of DMSO.

[†]The apparent C_{4v} symmetry of the symmetrical cone of calix[4]arenes is in fact a result of fast interconversion on the NMR timescale between two pinched cone conformers of C_{2v} symmetry [13].

SCHEME 1 Equilibrium between calix[4]arene monomer 1 and dimer 1_2 (3-0% DMSO- d_6 -CDCl₃, concentration of 1: 0.65 mM-10 mM).

Previously, we reported the bis(amidopyridyl) calix[4]arene 2 as a receptor for various dicarboxylic acids in 0.5–3% DMSO- d_6 –CDCl₃ solutions [17]. Therefore a complexation study between 2 and calix[4]arene dicarboxylic acid 1 was attempted in the same way. As expected, downfield shifts in the signal corresponding to the two NH protons of 2 $(ca. + 0.6 ppm, [2] = 0.2 mM, [1] = 1.8 mM)$ were observed in 0.5% DMSO- d_6 -CDCl₃ solution. Unfortunately, a Job plot [19] could not establish with any certainty the binding stoichiometry, due in part to

FIGURE 2 Spurr and Byers plot for 1 at 298 K in 1% DMSO- d_6 -CDCl3.

competition from the $1-1₂$ equilibrium in solution. However, the shape of the curve (Fig. 3) indicated a preference for a 2:1 (acid:amide) stoichiometry, which implies that a $[1-2-1]$ trimer may exist in the solution phase. This result is consistent with previous binding studies on 2 with aliphatic diacids $(HOOC- (CH₂)_n$ -COOH) that also show a preference for this stoichiometry for guests with larger (e.g. $n = 12$) spacer units [17]. Crystals suitable for X-ray diffraction were obtained by slow evaporation of a 0.5% DMSO- d_6 -CDCl₃ solution containing equimolar amounts of 1 and 2. The crystal structure was revealed to be a 1:1 co-crystal of the two compounds comprising parallel tube-like stacks each containing a single type of calixarene. Neighbouring tubes of 1 and 2 are linked in a 'zig-zag' arrangement via hydrogen bonds between carboxylic acids and amidopyridine units (Fig. 4). Additionally, neighbouring antiparallel tubes containing amidopyridylcalixarenes are mutually intercalated via π -stacking of the planar amidopyridyl arms (Fig. 5). Other examples of tubular arrangements involving calixarenes and resorcarenes have been observed in the solid state [20,21]. The hydrogen bond lengths in 1·2 are similar to those found in related amidopyridine–carboxylic acid complexes $[06 \cdots N2 =$ $2.684(7)$ Å, $(O6)H6 \cdot \cdot N2 = 1.86$ Å, $O6-H6 \cdot \cdot N2 =$ 167.2°; $N1 \cdots O7 = 3.046(7) \text{ Å}, \qquad (N1) H1 \cdots$ $O7 = 2.25$ Å, N1–H1··· $O7 = 149.8^{\circ}$] [22,23].

FIGURE 3 Job plot of 2 with 1 in 0.5% DMSO- d_6 -CDCl₃ solution.

These studies confirm that the calix[4]pyrrole dicarboxylic acid 1 forms a pinched cone dimer, $1₂$ both in the solid state and in solution, with the extent of the monomer(1)–dimer(1₂) equilibrium in solution dependent upon the extent of intermolecular hydrogen bonding. As well as demonstrating that this compound associates through hydrogen bonding interactions with a bis(amidopyridyl) calix[4]arene both in solution and in the solid state, we have utilised a convenient spectroscopic method for measuring the dimerisation constant of 1 in solution that may be applied to a range of related self-assembling calixarene systems.

EXPERIMENTAL

General

All reagents were used as received from chemical suppliers. $CH₂Cl₂$ was distilled from calcium hydride under inert gas. THF was distilled from sodium under inert gas. ¹H and ¹³C NMR spectra were recorded on either a Bruker AC 300 or a Bruker Avance DRX 400 spectrometer. Chemical shifts are

FIGURE 4 View of 1·2 down the tubular calixarene stacks. Molecules of dicarboxylic acid 1 are shown in blue and yellow; molecules of bis(amidopyridyl)calix[4]arene 2 are shown in green and blue. Dashed lines indicate hydrogen bonding between carboxylic acids and amidopyridyl groups that propagates a zigzag chain orthogonal to the tube axes (viz. yellow–red–yellow and blue–green–blue).

FIGURE 5 View of neighbouring antiparallel tubes comprising bis(amidopyridyl)calix[4]arene molecules 2. Mutual intercalation of amidopyridyl arms is observed. All contacts that lie within the sum of van der Waals radii are indicated with dashed lines. Note that it is not appropriate to discuss the geometry of any $C-H\cdots\pi$ contacts involving the tolyl groups since methyl hydrogen atoms were not refined without constraints.

reported in ppm (δ) . J_H values are given in Hz. 25,26,27,28-Tetrakis(1-propoxy)calix[4]arene-5,17 dicarboxylic acid 1 was prepared according to literature procedures [11].

5,17-Bis(amidopyridyl)-25,26,27,28-tetrakis (1-propoxy)calix[4]arene 2

25,26,27,28-Tetrakis(1-propoxy)calix[4]arene-5,17 dicarboxylic acid 1 (200 mg, 0.294 mmol) was dissolved in dry CH_2Cl_2 (5 ml) and $(COCl)_2$ (0.51 ml, 5.87 mmol) was added to the solution. The reaction mixture was refluxed for 3 hours. Then, the CH_2Cl_2 was distilled off and the residue was dried in vacuo for 1 hour. The obtained acyl chloride was dissolved in THF (25 ml), and the solution was added to solution of 2-amino-6-picoline (127 mg, 1.175 mmol) and Et_3N (0.245 ml, 1.175 mmol) dissolved in THF (10 ml). The reaction mixture was stirred for 1 hour. Then, the THF was distilled off, the residue was dissolved in $CHCl₃$ (30 ml) and washed with water (20 ml). The water layer was extracted with $CHCl₃$ (10 ml) and the organic layer was dried over MgSO₄. Following evaporation, the resulting yellow oil was purified by column chromatography on silica gel using hexane–ethyl acetate (3:1) as eluent. Yield 180 mg (71%), mp = 268-269°C (ethanol). ¹H NMR $(400 \text{ MHz}, \text{CDCl}_3)$: 0.94 (t, J = 7.4, 6H, OCH₂CH₂-CH₃), 1.10 (t, J = 7.5, 6H, OCH₂CH₂CH₃), 1.97 (m, 8H, OCH₂CH₂CH₃), 2.49 (s, 6H, NHpyCH₃), 3.25 (d, J = 13.6, 4H, ArCH₂Ar), 3.74 (t, J = 7.0, 4H, OCH₂CH₂-CH₃), 4.08 (t, $J = 8.0$, 4H, OCH₂CH₂CH₃), 4.49 (d, $J = 13.5$, 4H, ArCH₂Ar), 6.24 (d, $J = 7.4$, 4H, ArH), 6.32 (m, 2H, ArH), 6.91 (d, $J = 7.5$, 2H, ArpyH), 7.61 (t, $J = 7.8$, 2H, ArpyH), 7.62 (s, 4H, ArH), 8.41(s, 2H, ArCONH); ¹³C NMR (300 MHz,CDCl₃): 10.00, 10.67, 23.20, 23.46, 23.99, 31.01, 76.79, 111.00, 119.07, 122.49, 127.95, 127.99, 132.91, 136.96, 138.75, 149.48, 151.14, 155.40, 156.62, 161.35, 165.65, 221.78; MS (EI) m/z 860.45 (M⁺); Elemental analysis, found: C, 74.79; H, 7.02; N, 6.27%. Calcd. for $(C_{54}H_{60}O_6N_4)$: C, 75.32; H, 7.02; N, 6.51%.

X-ray Crystallography

The crystal of 1 was grown by slow evaporation from methanol after full dissolution by heating. Co-crystal [1·2] was grown by slow evaporation from the 0.5% $DMSO-d_6$ –CDCl₃ solution that was used in the NMR study.

Crystallographic data for 1. $C_{42}H_{48}O_8$, $M = 680.80$, $T = 120(2)$ K, monoclinic, space group $P2_1/n$, $a = 11.7592(4)$ Å, $b = 25.9714(9)$ Å, $c = 12.6411(5)$ Å, $\beta = 96.581(2)^\circ$, $V = 3835.2(2)$ \mathring{A}^3 , $Z = 4$, $\rho_{\text{calc}} =$ 1.179 Mg/m^3 . Reflections collected: 16713, independent: 6625 ($R_{\text{int}} = 0.0817$). Final R indices [$I > 2\sigma I$] : $R_1 = 0.0696$, $wR_2 = 0.1509$; R indices (all data): $R_1 = 0.1549$. $wR_2 = 0.1839$. All hydrogen atoms were located from the difference map and refined using a riding model. CCDC number: 183225.

Crystallographic data for [1-2]: $C_{96}H_{108}N_4O_{14}$, $M =$ 1541.86, $T = 120$ K, monoclinic, space group $P2/n$, $a = 19.6710(6)$ Å, $b = 9.3321(4)$ Å, $c = 27.5985(11)$ Å, $\beta = 101.504(2)^\circ$, $V = 4964.5(3) \text{ Å}^3$, $Z = 2$, $\rho_{\text{calc}} =$ 1.031 Mg/m^3 . Colourless block, crystal size $0.10 \times 0.07 \times 0.07$ mm. Reflections collected: 26307, independent: 8595 ($R_{int} = 0.142$). Final R indices $[I > 2\sigma I]$: $R_1 = 0.1141$, $wR_2 = 0.1753$; R indices (all data): $R_1 = 0.2802$. $wR_2 = 0.2185$. All hydrogen atoms were located from the difference map and refined using a riding model. CCDC number: 183226

Determination of the Dimerisation Constant of 1

The dimerisation constant (K_2) of 1 in 1% DMSO- d_6 – $CDCl₃$ was determined by ¹H NMR spectroscopy using a dilution technique. The data were analyzed by a linear graphical method for dimerisation developed by Spurr and Byers [18]. The experiment started with the most concentrated solution $(S_t = 10 \text{ mM})$ of compound 1, which was then diluted gradually until the detection limit was reached. The $1H NMR$ spectrum (δ_{obs} = the chemical shift of the H_a proton of 1) was recorded for each of these concentrations. The chemical shift of this proton in the most dilute solution was taken to be that of the monomer, $\delta_{\rm monomer}$. The dimerisation constant (K_2) of 1 was then calculated according to $\Delta = 2\Delta_2K_2S_t/(1+4K_2S_t)$ (Spurr and Byers equation), where $\Delta = \delta_{\rm obs.}$ – $\delta_{\text{monomer}}, \ \Delta_2 = \delta_{\text{dimer}} - \delta_{\text{monomer}}, \ \ S_t = \text{total}\ \text{concern}$ tration, and $1/\Delta = (1/S_t)(1/2\Delta_2K_2) + 2/\Delta_2$ (linear form), where the slope = $1/2\Delta_2K_2$, intercept = $2/\Delta_2$. Intercept/slope = $4\bar{K}_2$, $4K_2 = 1.483/0.01071$, $K_2 =$ $34.6 M^{-1}$.

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