The first example of a substrate spanning the calix[4]arene bilayer: the solid state complex of *p*-sulfonatocalix[4]arene with L-lysine[†]

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Received (in Columbia, MO, USA) 6th August 1999, Accepted 23rd November 1999

The complex of *p*-sulfonatocalix[4]arene with L-lysine shows a new type of intercalation behaviour with regard to the achiral hydrophobic bilayer assembly of the calixarenes, and represents the first structural example of a cationic organic substrate spanning such a bilayer.

It has recently been shown that the *p*-sulfonatocalix[*n*]arenes can act as heparin mimics,¹ and their interactions with the positively charged amino acids, lysine and arginine, has been investigated.² In the case of *p*-sulfonatocalix[4]arene, 1:1 complexes with lysine and arginine are observed, with binding constants derived from ¹H NMR spectroscopy at pH 5 of 600 and 1700. Undoubtedly such electrostatic interactions are also important in the activity of the sulfonatocalixarenes as chloride ion blockers.³ In recent years, *p*-sulfonatocalix[4]arene salts have been extensively studied; these complexes crystallize with a large variety of cations in bilayer-type structures.⁴ The basic molecular motif is achiral and formed by two calixarenes related by crystallographic symmetry. They are arranged in the up-down fashion with the sulfonate groups covering the surfaces of the bilayers which are separated by a hydrophilic layer containing the majority of the water molecules and cationic counterions. The structures of complexes of psulfonatocalix[4]arene with transition metal compounds⁵ have demonstrated that an aromatic organic substrate can be intercalated into the bilayer. However, no example of substrates traversing the bilayer have been observed. We report here on a new type of intercalation within the calixarene bilayer system showing that a chiral cationic organic molecule possessing a flexible aliphatic side chain, in this case L-lysine, can span the bilayer. The crystalline title compound also exhibits a chiral hydrophilic layer, containing three other L-lysine molecules, which separates the bilayers.

The crystal structure belongs to the triclinic *P*1 space group,‡ and consists of two [calix[4]arenesulfonate]^{4–} anions (**A** and **B**), four L-lysine counterions and 17.5 water molecules distributed over 20 sites. The calix[4]arene pattern is strongly pseudo-centrosymmetrical and similar to those already described in other complexes.^{4,5} The four L-lysine molecules and the sulfonate groups interact largely with water molecules. No water sites have been detected within the calixarene cavities.

For all the L-lysine molecules, the α - and ε -amino groups show contacts indicative of N–H···O hydrogen bonds with oxygen atoms on the sulphonate groups of the calixarenes, but they display different types of interactions. One L-lysine molecule (L1) is seen to traverse the hydrophobic bilayer with the main chain directed towards the sulfonate groups of the 'up', (A), calixarenes. As its side chain is in the fully extended conformation [$\chi_1 = -68(1)^\circ$, χ_2 , χ_3 , χ_4 values close to 180°], the ε -amino group points towards the opposite edge of the bilayer, nearly at the level of the S atoms of the 'down' (**B**), calixarenes (Fig. 1). While the α -amino group is in short contact with one sulfonate group of an **A** calixarene [N···O separation of 2.75(1) Å], the ε -amino group is connected to three **B** calixarenes, related by crystallographic translations along the *a*-and *b*-axes [N···O separations ranging from 2.853(6) to 3.013(6) Å]. This L-lysine molecule is thus the first example of a ligand which truly spans the hydrophobic calixarene bilayer.

Of further interest in the structure is the chiral hydrophilic layer separating the bilayers which contains the other three Llysine molecules. One L-lysine molecule (L2) is placed in the core of this layer§ and aligned parallel to the a-axis. As with L1, the side chain adopts the fully extended conformation [χ_1 = $-66(1)^{\circ}$, χ_2 , χ_3 , χ_4 values close to 180°], which is that most usually observed.⁶ The other two L-lysine molecules (**L3** and L4) lie just above the calixarene macrocycles (A and B for L3 and L4, respectively). Their arrangement is illustrated in Fig. 2. For both, the α -amino and α -carboxylate groups are directed into the chiral layer and the side chain is nearly at the level of the S atoms of their respective calixarene, with the ε -amino group pointed towards the exterior of the cavity. A common structural characteristic of L3 and L4 is that their side chains exhibit a folded conformation ($\chi_1 = 65.7(6)$ and $54.2(7)^\circ$, $\chi_4 = -46.2(7)$ and $-63.5(7)^\circ$, χ_2 , χ_3 values close to 180°]. This unusual conformation⁶ allows N–H…O contacts between one sulfonate group of the parent calixarene and the two amino groups of the same L-lysine molecule (Table 1). This conformation has been previously observed in the structure of the L-lysine



Fig. 1 Packing view along the *a*-axis of the structure demonstrating the spanning of one L-lysine molecule (L1, green) within the bilayer formed with the A (up) and B (down) *p*-sulfonatocalix[4]arenes. The other three L-lysine molecules are shown within the chiral layer which separates the bilayers (L2, dark blue; L3, light blue; L4, red).

[†] A figure showing the arrangement of the three L-lysine molecules within the chiral layer is available from the RSC web site, see http://www.rsc.org/ suppdata/cc/a9/a906546f/



Fig. 2 A view of a section of the structure illustrating the arrangement of L3 and L4 within the chiral layer and representing the N(amino)…O(sulfonate) contacts (dashed lines) referred to in the text. The position of the ε -amino group of L1 (NZ1) is also shown, the contacts between NZ1 and the sulfonate groups of two B calixarenes are indicated in dashed lines.

Table 1 Selected contacts (Å) between sulphonate oxygen atoms and the L-lysine molecules L3 and L4

L3		L4	L4	
N3…O41A	2.988(8)	N4…O43B	2.939(6)	
NZ3…021A	2.864(8)	NZ4···O33B	3.071(8)	
OB3····O42Ba	2.672(6)	NZ4····O42B	2.899(8)	
N3…O31B	2.890(7)	OB4····O41A	2.635(7)	
NZ3····O33B ^a	2.961(6)	NZ4····O23A ^b	2.806(10)	
Symmetry code ε-amino groups	s: $a x + 1$, y, z. $b x - are not considered.$	- 1, y, z. The minor comp	oonents of the	

sulfate,⁷ with similar specific interactions between the L-lysine molecule and the sulfate anion. L3 and L4 are connected by an N–H···O contact between the α -amino group of L3 and the carbonyl oxygen of L4 [N···O separation 2.928(7) Å]. These dimers are only interconnected indirectly through one sulfonate oxygen atom (O42B) of the B calixarene, which is simultaneously in contact with the ε -amino group of L4 and the hydroxy group of L3. The dimers are stacked along the *a*-axis. Thus, the L-lysine network within the chiral layer is composed of alternating rows of dimers and monomers. The bilayers are cross-linked through the chiral layer as L3 and L4 are also in contact with sulfonate groups of two calixarenes (B and A) lying in an adjacent bilayer (Table 1).

Thus, in contrast to the situation in solution, where a 1:1 complexation is observed, in the solid state a lysine–calixarene complex of 2:1 stoichiometry may be obtained. Actually, the system may be considered as two similar 1:1 complexes, each involving a folded L-lysine molecule (L3 and L4), which would correspond to the situation observed in solution.² The other two L-lysine molecules, both with the side chain in the fully extended conformation, would be implicated in the solid state structure. Of the four independent L-lysine molecules, three are found within the hydrophobic layer separating the typical sulfonatocalix[4]arene bilayer, while the remaining molecules (L1) spans this bilayer in a manner resembling biomolecules

traversing a lipid bilayer.¶ Further studies are underway to investigate the structural behaviour of the sulfonatocalix[n]arenes with peptides containing lysine and arginine.

Notes and references

‡ *Crystal data* for 2(C₂₈H₂₀O₁₆S₄)·4(C₆H₁₆O₂N₂)·17.5H₂O: M_r = 2389.5; suitable crystals for X-ray diffraction were grown in sealed tubes from two to one lysine–calix[4]arene aqueous solution, at room temperature during six months; triclinic *P*1, *a* = 13.599(4), *b* = 14.369(2), *c* = 15.169(3) Å, α = 111.02(1), β = 100.02(2), γ = 99.02(2)°, V = 2646(1) Å³, *Z* = 1, ρ_{calc} = 1.50 g cm⁻³, $2\theta_{max}$ = 46°, μ (Mo-K α) = 0.28 mm⁻¹, *T* = 293 K; 7673 independent reflections, 6530 with *I* > 2 σ (I); *R*₁ = 0.0863, *wR*₂ = 0.2553. The structure was solved by direct methods (SHELXS-86) and refined using the program SHELX-97. 17.5 water molecules were located, distributed over 20 sites. For three sulfonate groups, two O atoms are positionally disordered (two partially occupied sites for each). The ε-amino groups of **L3** and **L4** are disordered over two sites (0.75 and 0.25 occupancy factors). The parameters of the **A** and **B** calixarenes were refined in separate blocks. CCDC 182/1505. See http://www.rsc.org/suppdata/cc/a9/a906546f/ for crystallographic data in .cif format.

§ The L2 molecule is situated in a large intermolecular space and all the atoms show high isotropic displacement parameters. Nevertherless, the amino groups of L2 show contacts having relatively large separation distances with sulfonate oxygen atoms of two B calixarenes related by the *a*-translation [N···O separations of 3.02(1) and 3.07(1) Å]. A contact with a somewhat long N···O separation distance [3.09(1) Å] is also seen between the ε -amino group of L3 and the carbonyl oxygen of L2.

¶ It is of interest to note that there was debate concerning whether the original sulfonatocalix[4]arene structures resembled those of bilipid membranes or clay minerals.

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Communication a906546f