

# 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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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,<sup>1</sup> and their interactions with the positively charged amino acids, lysine and arginine, has been investigated.<sup>2</sup> In the case of *p*-sulfonatocalix[4]arene, 1:1 complexes with lysine and arginine are observed, with binding constants derived from <sup>1</sup>H 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.<sup>3</sup> 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.<sup>4</sup> 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 *p*-sulfonatocalix[4]arene with transition metal compounds<sup>5</sup> 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,<sup>‡</sup> and consists of two [calix[4]arenesulfonate]<sup>4-</sup> 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.<sup>4,5</sup> 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 sulfonate 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 [ $\chi_1 = -68(1)^\circ$ ,  $\chi_2, \chi_3, \chi_4$  values close to  $180^\circ$ ], 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 L-lysine 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 [ $\chi_1 = -66(1)^\circ$ ,  $\chi_2, \chi_3, \chi_4$  values close to  $180^\circ$ ], which is that most usually observed.<sup>6</sup> 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  $-63.5(7)^\circ$ ,  $\chi_2, \chi_3$  values close to  $180^\circ$ ]. This unusual conformation<sup>6</sup> 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

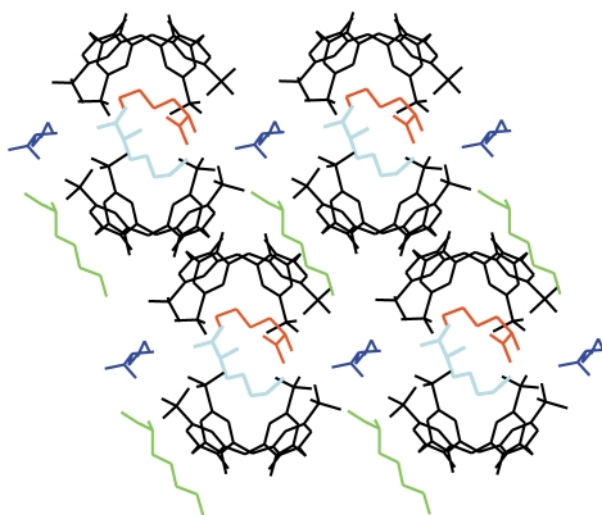


Fig. 1 Packing view along the *a*-axis of the structure demonstrating the spanning of one L-lysine molecule (**L1**, green) within the bilayer formed with the **A** (up) and **B** (down) *p*-sulfonatocalix[4]arenes. The other three L-lysine molecules are shown within the chiral layer which separates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

