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LETTER

Selective recognition of biogenic amine hydrochlorides by heteroditopic dihomooxacalix[4]arenes†

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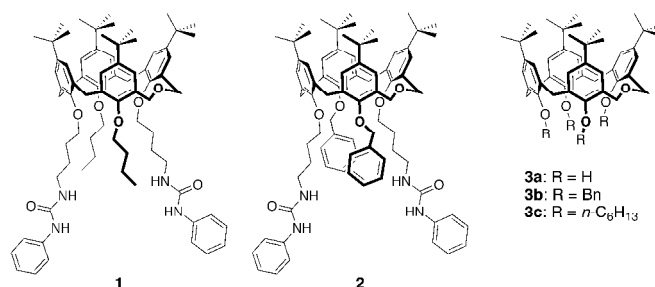
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Ion-pair recognition/complexation of monoamine neurotransmitter and trace amine hydrochlorides takes place –with outstanding efficiency and selectivity– within the π -rich cavity and the pocket formed at the lower rim by the two ureido moieties of tailor-made dihomooxacalixarene receptors **1 and **2**. The solid-state structure of **1** is also reported.**

The so-called ‘monoamine neurotransmitters’ (e.g. dopamine, histamine, norepinephrine, and serotonin) and ‘trace amines’ (e.g. 2-phenylethylamine and tyramine) are structurally related species that share an arylethylamine moiety as a common structural feature. The ability to selectively detect such target molecules is of enormous significance in medical and clinical research/analysis as these molecule play a key role in the physiological nervous system’s signaling processes, as well as in several psychiatric disorders and neurodegenerative pathologies (e.g., schizophrenia and Parkinson’s disease).¹

In recent years, a number of different artificial receptors, based on calixarenes,² homooxacalix[3]arenes,³ hemicyptophanes⁴ and other molecular frameworks⁵ have been prepared, but selective recognition nevertheless remains an open challenge. We have a long-standing interest in the fine-tuning of macrocyclic compounds of different type/size as artificial receptors⁶ and now, urged by the very recent findings that 18-membered *p-tert*-butyldihomooxacalix[4]arenes **3b,c** can efficiently bind linear alkylammonium substrates, providing a ‘superweak’ counterion is used,⁷ we wish to report our latest results on the ion-binding properties and selectivities of new heteroditopic dihomooxacalix[4]arenes **1** and **2** towards *n*-butylammonium halides and a number of biogenic amine hydrochlorides (see Fig. 3). Based on our earlier findings⁷ on **3b,c** –where the mandatory presence of a very loose anion (such as tetrakis[3,5-bis(trifluoromethyl)phenyl]-borate⁸) was required for alkylammonium cation *endo*-cavity complexation to occur– two of the lower rim alkyl/benzyl moieties of these receptors were replaced⁹ with ureido pendant groups with the specific aim of obtaining ditopic host molecules.¹⁰

Derivatives **1** and **2**,⁹ owing to their additional ureido anionic recognition sites,¹¹ were then expected to simultaneously bind the two ionic counterparts of alkylammonium salts and, as a result, minimize the detrimental ion-pairing effect¹² typically encountered in the recognition/binding process of a charged guest by a neutral host.



To verify this hypothesis, ¹H NMR titration experiments were carried out by adding increasing amounts of *n*-butylammonium chloride, bromide and iodide (0.5, 1.0 and 2.0 equiv.) to a 1.0 mM solution of receptors **1** and **2** in CDCl₃ to preliminary assess slow vs fast mode of complexation on the NMR timescale. Doubling of all the receptor resonances together with a new set of signals compatible with the presence of a guest molecule bound to the host were observed right from the addition of the first salt aliquot. *n*-Butylammonium cation inclusion, inside the homooxacalixarene cavity, is unequivocally substantiated by the appearance of the alkyl-chain resonances in the –1.4 to 0.0 ppm region of the ¹H NMR spectrum.^{3a,7,13} Owing to the chiral environment –arising from the asymmetric substitution pattern present at the lower rim of the hosts **1** and **2**– the pairs of enantiotopic hydrogen atoms belonging to the α - and β -CH₂ groups of the included guest display chemically non-equivalent signals once the alkylammonium ion is firmly anchored within the dihomooxacalixarene cavity.‡ On the other hand, simultaneous halide binding to the ureido moieties is underlined by the downfield shift observed for all the resonances assigned to the NHs (Fig. 1 and Fig. S4; ESI†). Collectively, these observations are

fully consistent with a slow complexation mode on the NMR timescale. Further addition of salt aliquots caused the disappearance of the free hosts' resonances.

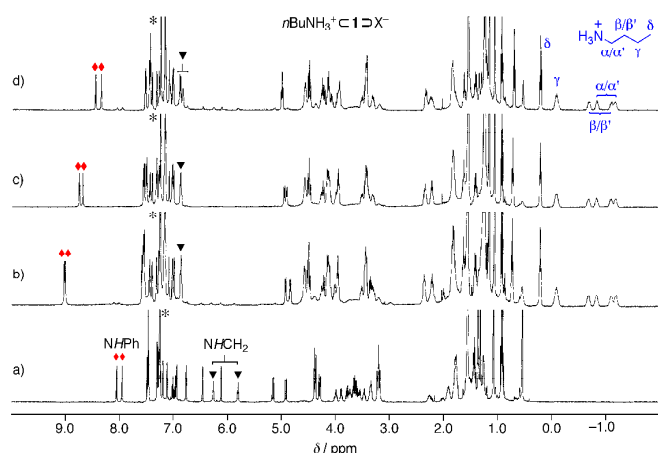


Fig. 1 ^1H NMR spectra (500 MHz, CDCl_3 , 298 K) of: a) $[\mathbf{1}] = 1.0$ mM, b) $[\mathbf{1}] = [n\text{-BuNH}_2 \cdot \text{HCl}] = 1.0$ mM; c) $[\mathbf{1}] = [n\text{-BuNH}_2 \cdot \text{HBr}] = 1.0$ mM and d) $[\mathbf{1}] = [n\text{-BuNH}_2 \cdot \text{HI}] = 1.0$ mM. The asterisk indicates the resonance of the residual solvent signals.

All host-guest ternary systems under investigation (1.0 mM, 1:1 host/guest salt solutions in CDCl_3) displayed very high percentages of complexation ($\geq 95\%$, corresponding to $K_{\text{ass}} > 10^9 \text{ M}^{-2}$), preventing us from reliably calculating the pertinent association constants by ^1H NMR analysis. The complexation induced shifts (CISs) of the receptor NHPh and the guest α - to γ - CH_2 resonances (acting as anion and cation binding probes, respectively), observed upon dihomooxalixarene/ n - $\text{BuNH}_2 \cdot \text{HX}$ binding, indicate that halide ions interact with the ureido binding pocket of $\mathbf{1}$ and $\mathbf{2}$ in a similar fashion, while the alkylammonium chain, irrespective of its counterion, is accommodated at the same depth inside the macrocycle cavity. Accordingly, the CIS values of the latter probe-resonances do not vary in the presence of different anions, while those of the NHPh s decrease with an anion dependent pattern ($\text{I}^- < \text{Br}^- < \text{Cl}^-$) which may suggest a correlation with the halogen electronegativity scale (Table S4; ESI †).¹⁴ The solid-state structure \ddot{s} of receptor $\mathbf{1}$ (Figs 2, S1, S2 and Tables S1–S3; ESI †) provides the means to explain the marked efficiency in the recognition of n -butylammonium halides. Similarly to what has been observed with calix[5]arenes,¹⁵ dihomooxalix[4]arene $\mathbf{1}$ possesses a highly preorganized aromatic cavity, ready to accommodate the incoming butylammonium cation. In addition, the proclivity of the ureido moieties to act as anion-binding sites is highlighted by the fact that, even in the absence of guest molecules, they interact with each other, self-assembling into an infinite one-dimensional H-bonded array.

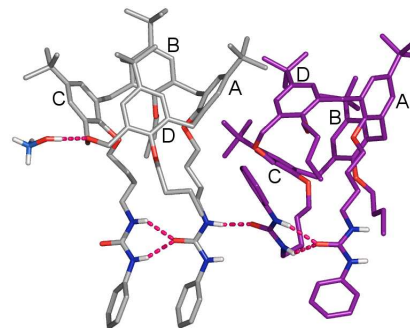


Fig. 2 The solid state structure of dihomooxalix[4]arene $\mathbf{1}$, showing the presence of molecules \mathbf{I} and \mathbf{II} (depicted in the figure in grey and violet, respectively with the pertinent ring labeling) in the asymmetric unit. Molecules \mathbf{I} and \mathbf{II} are seen to adopt similar *cone-in* conformations with their phenylureido moieties being involved in an infinite array of intra- and intermolecular $\text{N-H}\cdots\text{O}$ hydrogen bonds.

Having assessed the heteroditopic nature of hosts $\mathbf{1}$ and $\mathbf{2}$, we then turned our attention to monoamine neurotransmitters and trace amines (as the corresponding hydrochlorides) $\mathbf{4}$ – $\mathbf{10}$ (Fig. 3).

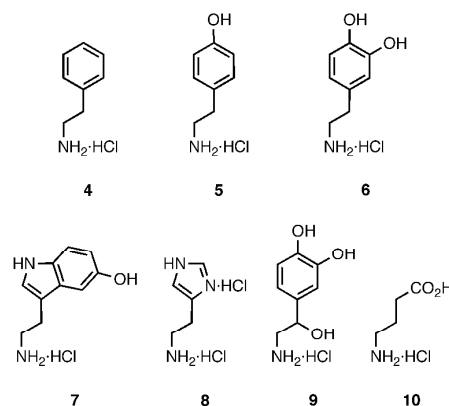


Fig. 3 Structures of the monoamine neurotransmitters and trace monoamines (tested as the corresponding hydrochlorides): 2-phenylethylamine (Pea-HCl, $\mathbf{4}$), tyramine (Tyrm-HCl, $\mathbf{5}$), dopamine (Dopa-HCl, $\mathbf{6}$), serotonin (Sert-HCl, $\mathbf{7}$), histamine (Hist-2HCl, $\mathbf{8}$), norepinephrine (Nore-HCl, $\mathbf{9}$) and \square -aminobutyric acid (Gaba-HCl, $\mathbf{10}$).

^1H NMR selectivity screening experiments were carried out in a $\text{CDCl}_3/\text{CD}_3\text{OD}$ solvent mixture (10:1, v/v) to ensure adequate solubility of both host and guest. Preliminary tests, carried out by adding 1 equiv. of ammonium guests $\mathbf{4}$ – $\mathbf{7}$ and $\mathbf{10}$ to a 1.0 mM solution of the host at 298 K caused a severe broadening of all ^1H NMR signals, indicating a strong host-guest interaction, but gave no clues as to the mode and/or the stoichiometry, nor to the extent of binding taking place in solution. For a slow –on the NMR timescale– host-guest complexation/decomplexation regime to be fully established, lowering of the temperature to 233 K was necessary and, as a result, analysis of the complexation process became possible (see Figs 4, S5 and S6 in the case of $\mathbf{1}$ and Tyrm-HCl; ESI †).

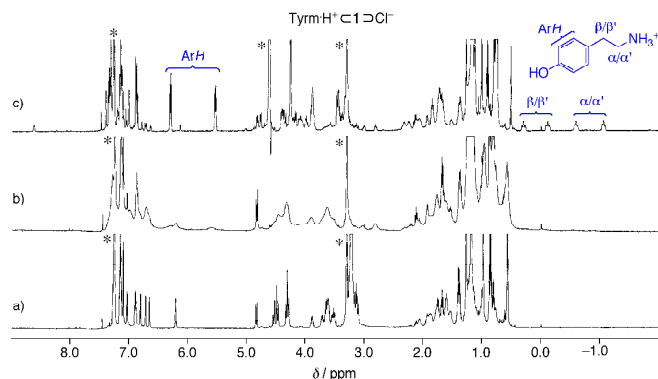


Fig. 4 ^1H NMR spectra (500 MHz, $\text{CDCl}_3/\text{CD}_3\text{OD}$ 10:1 v/v) of: a) **1** = 1.0 mM at 298 K, b) **1** = $[\text{Tyrm}\cdot\text{HCl}] = 1.0$ mM at 298 K and c) **1** = $[\text{Tyrm}\cdot\text{HCl}] = 1.0$ mM at 233 K.

In analogy with the butylammonium halide cases, the α - and β - CH_2S of the guests are recognized and included inside the asymmetric cavity of **1** and **2** and as a result display four distinct resonances in the high field region of their ^1H NMR spectrum.¹⁶ The percentages of complex formation and the association constants (K_{ass}), determined by ^1H NMR, are summarized in Table 1.

Table 1 Percentages of complex formation and corresponding association constants, K_{ass} ($\times 10^3 \text{ M}^{-2}$), determined^a by ^1H NMR (500 MHz, in $\text{CDCl}_3/\text{CD}_3\text{OD}$ (10:1, v/v), at 233 K) between dihomooxalix[4]arenes **1** and **2** and amine hydrochlorides **4–10**.

	Pea·H Cl (4)	Tyrm·H Cl (5)	Dopa·H Cl (6)	Sert·H Cl (7)	Hist·2H Cl (8)	Nore·H Cl (9)	Gaba·H Cl (10)
1	86% 41±4.5	85% 36±4.3	67% 6.3±0.9	61% 4.1±0.6	n.o. ^b	n.o. ^b	63% 4.5±0.2
2	83% 30±4.4	81% 23±2.9	63% 4.5±0.6	60% 3.7±0.5	n.o. ^b	n.o. ^b	60% 3.7±0.3

^aPercentage values derive from an average of three independent measurements. ^bN.o. stands for no complexation observed.

A closer inspection of these data shows that host **1** is marginally more efficient than **2**. Moreover, K_{ass} values indicate a comparable affinity trend between arylalkyl amines and both hosts (Pea·HCl > Tyrm·HCl > Dopa·HCl > Sert·HCl). It is reasonable to speculate that guests **4** and **5** best fit inside the calixarene cavities as a result of their less bulky aryl moiety. According to our data, the lower association constants observed in the case of the aliphatic neurotransmitter Gaba suggest that π - π interactions, very often operating in the binding pockets of biological receptors,¹⁷ play a key role in the recognition/binding process between *p*-*tert*-butyldihomooxalix[4]arenes **1** and **2** and phenylethylamine or tyramine. Interestingly, no interaction at all was detected with the β -substituted norepinephrine·HCl or the dicationic histamine·2HCl, either at 298 or at 233 K (see Fig. S7; ESI[†]). Despite the structural similarity of the neurotransmitters tested (**4–9** contain an arylethylammonium moiety), selectivity data –calculated as the ratio of two given K_{ass} values, Table 2– indicate that receptors **1** and **2** show a higher affinity (in the 5–10-fold range) for Pea·HCl and Tyrm·HCl over the other amine hydrochlorides.

Table 2 Selectivities of dihomooxalix[4]arenes **1** and **2** for monoamine hydrochlorides **4–7** and **10**.

	$S_{\text{Pea/Dopa}}$	$S_{\text{Pea/Sert}}$	$S_{\text{Pea/Gaba}}$	$S_{\text{Tyrm/Dopa}}$	$S_{\text{Tyrm/Sert}}$	$S_{\text{Tyrm/Gaba}}$
1	6.6	10.1	9.2	5.8	8.9	8.1
2	6.6	8.1	8.0	5.0	6.1	6.0

In conclusion, our studies demonstrate that dihomooxalix[4]arenes **1** and **2** behave as heteroditopic receptors efficiently binding organic ammonium halides in organic media. Their preorganized π -rich cavities mimic the hydrophobic binding pockets of the G protein-coupled receptors, displaying high association constants and a good degree of selectivity towards 2-phenylethylamine, tyramine, dopamine and serotonin neurotransmitters. These features make receptor **1** and **2** promising candidates for the development of sensing devices capable of monitoring the level of biogenic amines in biological fluids.

Experimental

General

Dihomooxalix[4]arenes **1** and **2** were available from previous studies.⁹ Amine hydrochlorides [**4–10**] were purchased from Sigma-Aldrich and were used as received.

^1H NMR Complexation Experiments

All spectra were recorded at 500 MHz at either 298 or 233 K. Percentages of the host-guest ternary complex formation (required for the calculation of the corresponding association constants, K_{ass}) were determined by direct ^1H NMR integration analysis of the ‘free’ and ‘complexed’ resonances of the guest and/or the host, present at equilibrium, in a equimolar host-guest solution. A set of three different equimolar sample solutions was examined for each K_{ass} determination. In the case of dihomooxalix[4]arenes **1** and **2** with *n*-BuNH₃⁺X[−] (X[−] = Cl[−], Br[−] and I[−]), where percentages of complexation were found to be $\geq 95\%$, K_{ass} values were estimated to be $>10^9 \text{ M}^{-2}$.

All samples were prepared by mixing together aliquots of stock solutions of host (600 mL) and guest (60 mL) to obtain a final equimolar host-guest solution (1.0 mM). The following stock solutions were used: [**1–2**] = 1.1 mM in CDCl_3 or $\text{CDCl}_3/\text{CD}_3\text{OD}$ (10:1, v/v), [*n*-BuNH₃⁺X[−]] = 11.0 mM in CDCl_3 and [**4–10**] = 11.0 mM in $\text{CDCl}_3/\text{CD}_3\text{OD}$ (10:1, v/v). For the assessment of the NH resonance complexation induced shifts a $\text{CDCl}_3/\text{CH}_3\text{OH}$ (10:1, v/v) solution of [**1**] = $[\text{Tyrm}\cdot\text{HCl}] = 1.0$ mM was used.

X-ray crystallography

Colourless very small single crystals (typical size $0.05 \times 0.08 \times 0.1$ mm) suitable for X-ray analysis by synchrotron radiation were obtained from slow diffusion of CH_3OH into a CHCl_3 solution of dihomooxalix[4]arene **1** at room temperature. Data collection was carried out at the X-ray diffraction beamline of the Elettra Synchrotron, Trieste, Italy, employing the rotating-crystal method with the cryo-cooling technique. The crystal dipped in Paratone, as cryo-protectant, was mounted in a loop and flash frozen to 100 K under a stream of nitrogen. Diffraction data were indexed and integrated using MOSFLM¹⁸ and scaled with SCALA.^{19,20}

The structure was solved by direct methods using SIR2011.²¹ Non-hydrogen atoms were refined by full-matrix least-square methods on F^2 using SHELXL-13²² with the contribution of H atoms placed at the geometrically calculated positions and refined using the riding model. The crystallographic model is affected by disorder and a detailed description is provided in

the ESI†. Crystallographic data and refinement details are reported in Table S1, see the ESI†.

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Notes and references

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† Electronic supplementary information (ESI) available: additional ¹H NMR spectra and crystal data for **1**. CCDC reference number 1009330. See DOI: 10.1039/x0xx00000x

‡ Signal assignment follows from COSY experiments (see Fig. S3).

§ The asymmetric unit of the triclinic cell contains two crystallographically independent receptor molecules (**I** and **II**, as seen in Fig. 2) and a methanol solvent molecule. See the ESI† for a full description.

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