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Conformational selectivity and high-affinity binding in the complexation of N-phenyl amides in water by a phenyl extended calix[4]pyrrole

We report the synthesis of a tetrapyridinium phenyl extended calix[4]pyrrole showing high selectivity and binding affinity for the cis-conformers of N-phenyl-amides in water. The conformational selectivity is remarkable owing to the energetic preference of the amides to adopt the trans-conformation. In turn, the high binding affinities derived from the combination of hydrogen bonding interactions and the hydrophobic effect. We used two binding models for the mathematical analyses of the titration data and rationalized the different magnitudes calculated for the corresponding binding constants (apparent and intrinsic).
Conformational selectivity and high-affinity binding in the complexation of \( N \)-phenyl amides in water by a phenyl extended calix[4]pyrrole†

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We describe the synthesis of a tetrapyridinium phenyl extended calix[4]pyrrole that is soluble in neutral water solution at mM concentrations. We show that, in pure water, the synthesized calix[4]pyrrole receptor selectively binds the cis-(E) conformers of secondary \( N \)-phenyl-amides and tertiary \( N \)-methyl-\( N \)-phenyl-formamide with binding affinities larger than \( 10^{3} \) M\(^{-1}\). The conformational selectivity is remarkable owing to the energetic preference of amides to adopt the trans-(Z) conformation in solution. In this respect, we used two binding models for the mathematical analyses of the titration data and calculated apparent and intrinsic binding constants. The combined action of hydrogen bonding and the hydrophobic effect that operates in the binding of the amides in water is responsible for the large affinities displayed by the receptor.

**Introduction**

Carboxamides (a.k.a organic amides) are important functional groups in both natural (e.g. peptides) and synthetic (e.g. nylons) polymers. The existence of many methods for their synthesis and their high thermodynamic stability under a wide range of chemical conditions render amides very useful covalent linkers in many chemical constructs. The pseudo-double-bond character of the carbon(C=O)–nitrogen \( \sigma \)-bond restricts its rotation, and therefore, secondary and tertiary amides exist as two isomeric cis–trans rotamers (Fig. 1). In solution, secondary amides show a marked preference for the trans-rotamer (Z-isomer). On the one hand, the energetic difference between the two rotamers lies in the range of 0.5 to 2.5 kcal mol\(^{-1}\) for both secondary and tertiary amides. On the other hand, because the free energy barrier for the cis–trans interconversion is moderate (\( \Delta G = 16–22 \) kcal mol\(^{-1}\)), the process is typically slow on the NMR time-scale but fast on the human time-scale. Steric contributions on either side of the amide group seem to be the most significant factors for easing the barrier crossing, as well as controlling the relative population of cis–trans isomers.

The cis–trans isomerism of the amide bond plays an important role in determining folding, functional and ligand interactions in proteins and peptides. Likewise, the photophysical behaviour of simple phenanthrene carboxamides was shown to be dependent on the amide conformation. The equilibrium between secondary amide rotamers is known to be solvent dependent. In addition, cis-amide rotamers have been stabilized by intramolecular CH–π interactions, lone pair–π interactions, and the hydrophobic effect. In organic solvents, the formation of host-guest hydrogen bonded complexes has also been used to stabilize cis-amides. In water, the combination of hydrogen bonding and the hydrophobic effect has provided thermodynamically stable complexes of synthetic receptors with small linear and cyclic peptides. However, to the best of our knowledge, examples of synthetic receptors displaying selective binding of the cis-rotamers of mono-amides in water and producing thermodynamically highly stable complexes are not known.

Sessler et al. described the binding of small tertiary amides (i.e. DMF and DEF) using octa-methyl calix[4]pyrrole in benzene solution. The resulting complexes displayed binding affinities of the order of \( 10^{3} \) M\(^{-1}\) and were mainly stabilized by the establishment of hydrogen-bonding interactions between the

![Fig. 1 Equilibrium between the cis-(E) and trans-(Z) conformers of a secondary amide. The structural analogy of the cis-(E) isomer with the lactam tautomeric form of creatinine is highlighted.](Image)

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pyrrole NHs and the oxygen atom of the amide’s carbonyl group. More recently, we reported the use of phosphonate calix[4]pyrrole cavities, featuring a polar aromatic cavity, for the efficient binding of creatinine in organic solvents. In solution and in the solid-state, the inclusion of creatinine in the cavity of the receptor was mainly driven by the establishment of multiple hydrogen bonds. Nevertheless, additional intermolecular CH–π interactions were also present in the above creatinine complex. To us, the lactam tautomer of the bound creatinine resembled a creatinine complex. To us, the lactam tautomer of the bound creatinine complex. To us, the lactam tautomer of the bound creatinine complex.

Herein, we report the synthesis of a water soluble tetra-arylextended calix[4]pyrrole receptor 1. We also describe the results of the binding studies of receptor 1 with a series of acyclic mono-amides in water. The conformational selectivity displayed by receptor 1 in the exclusive binding of the cis-rotamers of N-phenyl-amides with affinity constants larger than 10^3 M^-1 is highlighted.

Results and discussion

Synthesis of tetra-pyridinium calix[4]pyrrole receptor 1

Water soluble aryl-extended calix[4]pyrroles bearing ionizable carboxylic acid functions have been previously reported in the literature. These receptors required basic media for their water solubilisation. In order to avoid these conditions, not compatible with biological standards, we designed α,α,α,α-tetra-phenyl calix[4]pyrrole 1 (Scheme 1a) featuring four meso-(3-pyridinium-propyl) substituents at its lower rim. The water solubilization of calix[4]pyrrole 1 was inspired by previous studies on water soluble deep cavitands bearing benzimidazolones at their upper rims. In these examples, the deep cavitands were rendered water soluble by converting the lower-rim tetra-chloride derivatives into quaternary ammonium salts by treatment with an excess of pyridine. Receptor 1 was synthesized in two steps from commercially available starting materials (Scheme 1a). To start with, the α,α,α,α-tetra-chloro calix[4]pyrrole derivative 2 was prepared using a synthetic methodology recently developed by us. Treatment of tetra-chloride 2 at 110 °C in an excess of pyridine provided the tetra-pyridinium calix[4]pyrrole salt 1 as a yellow precipitate in excellent yield and purity. Calix[4]pyrrole 1 was soluble in neutral water at concentrations up to 15 mM. A dilution experiment (1 mM to 0.4 mM) showed negligible changes in the proton signals of the acquired 1H NMR spectra. At diluted concentrations, the proton signals of 1 are sharp and well-defined and their number is in agreement with C2v symmetry (ESS^2). Most likely, both alternate and cone conformations of 1 are present in solution. The two conformers are in equilibrium and display a fast chemical exchange on the chemical shift time-scale.

Binding studies of 1 with formamides in water

Molecular modelling. Earlier studies with aryl-extended calix[4]pyrroles related to 1 and their cavitand derivatives demonstrated that, in the solid-state and organic media, N,N-dimethylformamide and creatinine were suitable guests for inclusion in their polar aromatic cavities defined by the four meso-phenyl substituents. Based on these precedents, we envisaged that α,α,α,α-aryl-extended calix[4]pyrrole 1 might show conformational selectivity in the recognition of size and shape complementary secondary and tertiary small amides in water. Simple molecular modelling studies (Fig. 2) revealed that the inclusion of N-phenyl-formamide 3 in the aromatic cavity of the cone conformation of 1 was energetically more favorable (ΔE = 7.3 kcal mol^-1) for the cis-rotamer (Fig. 2a).

The cis-3·1 complex is stabilized by four hydrogen-bonding interactions established between the amide carbonyl oxygen

![Diagram](image-url)

**Scheme 1** (a) Synthesis of tetra-phenyl tetra-pyridinium calix[4]pyrrole receptor 1 and (b) line-drawing structures of the formamides (3–7) and acetamides (8–14) discussed in this work. When applicable only the cis-rotamer is depicted. See Fig. 1 for the cis/trans equilibrium.

![Diagram](image-url)

**Fig. 2** Energy minimized structures (PM3, water COSMO model as implemented in SCIGRESS Version FJ 2.6 (EU 3.1.9)) of the inclusion complexes: (a) cis-3·1 and (b) trans-3·1. A difference of energy of 7.3 kcal mol^-1 was calculated in favor of the cis-3·1 complex. Calix[4]pyrrole 1 is shown in stick representation and amides (cis-3 and trans-3) are shown as CPK models. Meso-(3-pyridinium-propyl) groups were pruned to methyl groups to ease the calculations.
atom and the NHs of the four pyrrole units. Additional CH–π and NH–π interactions are also established between the included cis-3 amide and the meso-phenyl walls of 1. On the other hand, the inclusion of trans-3 in the cone conformation of 1, trans-3⊂1 complex, evidenced significant steric clashes between the N-phenyl substituent and two of the four meso-phenyl groups of the receptor (Fig. 2b).

\[ 1^H \text{NMR spectroscopy titrations.} \] We decided to experimentally probe the interaction of 1 with N-phenyl-formamide 3 in water using \( ^1 \)H NMR titration experiments. We mentioned above that the steric factors are usually the most important contributors to the relative populations of the cis/trans rotamers in small secondary and tertiary amides, and that the isomer-ratio is also affected by the solvent.\[8\] In this respect, the \( ^1 \)H NMR spectrum of a 10 mM solution of N-phenyl-formamide 3 in water showed a 32 : 68 cis/trans ratio, based on the integral values of the two singlets resonating at \( \delta = 8.65 \) and 8.28 ppm, respectively, which are assigned to the hydrogen atoms of the formyl group in each rotamer (ESI†).

The incremental addition of formamide 3 to a 1 mM water solution of the tetra-pyridinium receptor 1 induced significant chemical changes to several proton signals of the receptor. In particular, the triplet of the para-aromatic proton, \( H_c \), of the meso-phenyl groups of 1 moved upfield (\( \Delta \delta_{\text{max}} = -0.21 \) ppm). In contrast, the \( \beta \)-pyrrole protons (\( H_b \)) moved downfield (\( \Delta \delta_{\text{max}} = +0.09 \) ppm).\[9\] These observations indicated the existence of a fast chemical exchange on the \( ^1 \)H NMR time-scale between free and bound receptor 1. Remarkably, after the addition of 0.6 equiv. of 3 only the sharp singlet corresponding to the hydrogen atom of the formyl group for the free trans-isomer became visible. At this point, we also observed a complex signal of aromatic protons resonating at \( \delta = 7.5 \) ppm that was assigned to the phenyl group of the formamide. The signals assigned to free trans-3 grew in intensity as the concentration of 3 was increased (Fig. 3). In contrast, the signal of the formyl proton in the free cis-rotamer was not observed when 2 equiv. of 3 were added.

The titration data were mathematically analyzed using the HypNMR 2008 software Version 4.0.66,67 The fit of the chemical shift changes experienced by the selected proton signals of receptor 1 (\( H_n, H_c, H_d, \) and \( H_e \), see Scheme 1 for proton assignment) to a binding isotherm of a theoretical 1 : 1 binding model was good. We estimated an apparent binding constant value \( K_a > 10^4 \text{M}^{-1} \) for the formed complex and determined the chemical shift values of the protons of the receptor in the complex (ESI†). These latter values allowed the calculation of the corresponding complexation induced shifts (CIS).

At first sight, the determined magnitude of the apparent binding constant might be considered as the weighted-average of two putative 1 : 1 inclusion complexes: cis-3⊂1 and trans-3⊂1. In order to assess the conformational selectivity exhibited by receptor 1 in the binding of 3, we decided to titrate it against tertiary \( N,N,N \)-diphenyl-formamide 4. The hydrogen-bonding inclusion of formamide 4 in the cavity of 1 should produce the 4⊂1 complex exhibiting a binding geometry closely resembling that of the trans-3⊂1 complex (Fig. 2b). Our idea was to use the binding constant value of 4⊂1 as the reference for the trans-3⊂1 counterpart.

The extensive sonication of a suspension of \( N,N,N \)-diphenyl-formamide 4 with a 1.7 mM water solution of 1 led to the dissolution of 2 equiv. of 4 but did not induce appreciable changes in the chemical shift values of the receptor’s protons. Thus, we estimated a binding constant value \( K_a(4⊂1) < 10^{-1} \text{M}^{-1} \).

On the one hand, this result strongly supports a conformational selectivity towards the cis-rotamer in the binding of \( N \)-phenyl-formamide 3 with receptor 1. On the other hand, it also requested a change in the mathematical analysis of the titration data in order to assess a more accurate value of the binding constant for the cis-3⊂1 complex.

Therefore, we reanalyzed the titration data of 1 with 3 using a theoretical binding model that considers the existence of the equilibrium between the two rotamers of 3 and the exclusive formation of a 1 : 1 complex with the cis-rotamer and receptor 1.

In doing so a theoretical binding isotherm was calculated from the simulated speciation profile produced with the Specfit Software Version 3.0.40 (differential kinetics module). We considered the equilibrium constant between the two rotamers of 3 and the determined chemical shift values for the protons in the free and bound receptor 1 as fixed values. The value of \( K(cis-3\text{}/trans-3) = 32/68 = 0.47 \) was determined from the \( ^1 \)H NMR spectrum of 3. The complexation induced shift values (CISs) for the protons of 1 were those calculated from the previous fit (vide supra and ESI† for details). The fit of the experimental titration data to the more elaborate theoretical binding model only served to estimate that the binding constant value \( K_a(cis-3⊂1) \) was larger than \( 10^4 \text{M}^{-1} \). The magnitude of this estimate represents a remarkable binding affinity for the complexation of a small polar molecule in water using a synthetic receptor.\[38,39\]

The fact that the cis-rotamer is present in solution to a significant extent (32%) is the main reason for the observed coincidence between the binding constant values estimated using the two bindings models, that is \( K_a(cis-3⊂1) \sim K_{\text{app}} \). We will show below that when the cis-rotamer is present in solution at low levels, the mathematical analyses of the titration data using the two binding models will produce very different values for \( K_a(cis-3⊂1) \) and \( K_{\text{app}} \).

Next, we performed a reverse titration of \( N \)-phenyl-formamide 3 with receptor 1 (Fig. 4). The \( ^1 \)H NMR spectrum
of a 1.2 mM water solution of 3 containing just 0.05 equiv. of 1 (Fig. 4b) showed the formyl proton of the trans-3 rotamer as a sharp singlet resonating at the same chemical shift as in the absence of 1. On the contrary, the formyl singlet assigned to the cis-3 rotamer was not detectable at this stage of the titration. This observation indicated that while the equilibrium involving the two free rotamers of 3 displayed slow chemical exchange on the chemical shift time-scale, the cis-3 rotamer was also involved in a binding equilibrium, probably with bound cis-3, featuring intermediate chemical exchange on the chemical shift time-scale. The intermediate kinetics of the latter chemical exchange processes produced broadening beyond detection for the formyl proton in the cis-3 rotamer. This result also provides irrefutable support to the selective binding of the cis-3 rotamer by receptor 1.

In complete agreement with the previous statement, an increase in the concentration of 1 did not affect the appearance and chemical shift value of the formyl proton signal for the trans-3 rotamer, however, its intensity decreased. We also observed an increase in the intensity of three aromatic signals, two triplets moving downfield and one doublet that shifted upfield. We assigned these three signals to the protons of the N-phenyl group of the cis-3 rotamer, which are involved in a fast chemical exchange on the chemical shift time-scale between its free and bound states. The different kinetics of the chemical exchanges experienced by the protons of the cis-3 rotamer, that is, the formyl proton is involved in an intermediate exchange and the aromatic protons are involved in a fast exchange, result from notably different CISs. The formyl proton of bound cis-3 is included in the aromatic cavity of 1 experiencing the strong shielding exerted by the meso-phenyl substituents.[1]

On the other hand, the N-phenyl unit of bound cis-3 resides almost completely outside of the aromatic cavity of 1, with its ortho-protons being the only ones affected by the shielding effect (Fig. 2a and 4).

In order to evaluate the scope of the conformational selectivity featured by tetra-phenyl calix[4]pyrrole receptor 1 in the binding of other formamides, we investigated the complexation of 1 with N-methyl-N-phenyl-formamide 5. Based on the integral values of the two separate formyl protons, formamide 5 is present in water solution as an 82 : 18 cis/trans mixture of rotamers (Kcis-5/trans-5 = 4.56).

The direct $^1$H NMR titration of 1 with incremental amounts of 5 showed an analogous behavior with respect to the chemical shift changes of the proton signals of receptor 1 to the one previously described for 3. The mathematical analysis of the titration data using the binding model that considers the equilibrium between rotamers and the exclusive formation of a 1 : 1 complex with the cis-5 isomer allowed us to estimate that the affinity constant value $K_a(cis-5=C1)$ must be > $10^4$ M$^{-1}$. We also performed an ITC titration experiment for formamide 5 and calix[4]pyrrole 1 (ESI†). The calculated apparent binding constant was 8.6 ± 0.1 × 10$^3$ M$^{-1}$, a value that is in line with the estimated one from the $^1$H NMR titration analysis. The ITC results showed that the binding process was enthalpically driven (−7.1 kcal mol$^{-1}$) and entropically opposed (1.75 kcal mol$^{-1}$). Clearly, this thermodynamic signature is not the expected one for purely hydrophobic binding.

We also performed a reverse titration by adding increasing amounts of 1 to a 2.9 mM water solution of 5. The proton signals assigned to the trans-5 rotamer did not experience noticeable chemical shift changes. In contrast, those of the cis-5 counterpart, especially the formyl proton and the methyl protons, moved significantly upfield. From these titration data, we could estimate the CISs experienced by these protons because, although their signals broadened beyond detection in the initial and middle phases of the titration, they became observable again in the presence of 2 equiv. of 1. In the cis-5$\subset$1 complex, the formyl group resonates at $\delta = 5.01$ ppm (Δ$\delta = -3.39$ ppm) and the methyl resonates at $\delta = 1.25$ ppm (Δ$\delta = -2.07$ ppm). The large calculated CISs confirm the deep inclusion of cis-5 in the aromatic cavity of 1.

A 2D NOESY experiment showed intermolecular cross-peaks between the singlet of the methyl protons of bound cis-5 and the aromatic protons (H$_c$ and H$_d$) of the meso-phenyl substituents of the calix[4]pyrrole 1.**

We also assessed the affinity constant of calix[4]pyrrole 1 for N-methyl-formamide 6. Formamide 6 is present in water as an 8 : 92 mixture of cis/trans rotamers. Nevertheless, molecular modelling studies and the previous results obtained in the binding of the secondary and tertiary formamides 3 and 5 strongly supported the inclusion of both isomers of 6 in receptor 1. The energy minimized 1 : 1 inclusion complexes, cis-6$\subset$1 and trans-6$\subset$1, featured close to isoenergetic values (ESI†).

Therefore, the $^1$H NMR titration data for the interaction of 1 with 6 were analyzed using a simple 1 : 1 binding model obtaining a good fit and returning an affinity constant value of $K_a(6$/$C1$) = 4.4 × 10$^3$ M$^{-1}$. The removal of the phenyl group in 6 compared to 5 slightly diminished the binding affinity for receptor 1 (Table 1). Probably, this difference is caused by a combination of reduction in hydrophobicity and the hydrogen-bonding accepting character of the oxygen atom in formamide 6 (secondary vs. tertiary).

In agreement with this hypothesis, the addition of an extra methyl group into $N,N$-dimethyl-formamide 7 increased its...
binding affinity for 1, compared to 6, placing it at the same level observed for the analogous but more lipophilic N-phenyl derivatives 3 and 5.

It is worth noting that in the reverse $^1$H NMR titration of N-methyl-formamide 6 with receptor 1, after the first addition of the receptor, the separate methyl proton signals of the two rotamers of 6 coalesce into a broad singlet that shifted upward upon increasing the concentration of 1 (ESI†). This result clearly supports the binding of two rotamers of 6, which are involved in a chemical exchange equilibrium with the free counterparts featuring fast/intermediate dynamics on the chemical shift time-scale.

Also note that, in the examples of N-phenyl-formamides 3 and 5, only the proton signals assigned to the cis-rotamer experienced broadening and chemical shift changes. We used this observation as evidence of the conformational selectivity in the binding process. The results described above indicate that receptor 1 might be considered as a minimal synthetic chaperone selecting the cis-conformation of the bound N-phenyl-formamides and increasing their relative concentration in solution in the bound form.

**Binding studies of 1 with acetamides in water**

**Molecular modelling.** Molecular modelling studies suggested that acetamide 8 and the cis-rotamer of N-methyl-acetamide 9, as well as those of other secondary N-aryl amide derivatives like 10–13, were suitable candidates for inclusion in the aromatic cavity of 1 (Fig. 5a). The modelled acetamides closely resembled the cis-rotamer of N-methyl-formamides (ESI†) in terms of size complementarity with the dimensions of the aromatic cavity of 1 in the cone conformation.

In contrast, tertiary N,N-dimethyl acetamide 14 presents an additional methyl group to be included in the cavity of 1. The

![Energy minimized structure of the 14⊂1 complex displayed a severe distortion of the cone conformation of the receptor owing to steric clashes between the meso-phenyl substituents and the two included methyl groups of the amide (Fig. 5b).]

**$^1$H NMR spectroscopy titrations.** The titration of receptor 1 with increasing amounts of acetamide 8 afforded exclusively the 8⊂1 complex and, accordingly, it was analyzed using a simple 1 : 1 binding model returning $K_a(8⊂1) = 6.8 \times 10^4 \text{ M}^{-1}$. The magnitude of the binding constant is in agreement with the value determined for the isosteric N-methyl-formamide, $K_a(6⊂1) = 4.4 \times 10^3 \text{ M}^{-1}$.

In water, N-methyl-acetamide 9 and N-phenyl-acetamide 10 displayed cis/trans isomeric ratios of 2 : 98 and 1 : 99, respectively, based on the integral values of their methyl acetamide proton signals. In particular, the assignment of the methyl proton signal for the cis-10 rotamer to a small singlet resonating at $\delta = 1.90$ ppm was confirmed by magnetization transfer experiments (ESI†). The titration data of 1 with both acetamides, 9 and 10, were analyzed in an analogous manner. For the sake of brevity, we will only describe in detail the titration experiments performed with 10 and the corresponding data analyses.

The incremental addition of 10 (up to 15 equiv.) to a 1.2 mM solution of 1 in water produced very small changes in the chemical shift values of the diagnostic protons of the receptor (H$_b$, H$_c$, H$_d$ and H$_e$) used for signaling of complex formation in the formamide series.

We also performed a reverse $^1$H NMR titration experiment adding incremental amounts of calix[4]pyrrole 1 to a 1.5 mM solution of 10 (ESI†). As expected the proton signals of trans-10 did not experience chemical shift changes and only a small reduction of their intensities was observed. Unfortunately, it was not possible to accurately quantify the amount of trans-10 rotamer that has been isomerized into the cis-10 rotamer using the integral values of selected signals in the acquired $^1$H NMR spectra due to its selective complication.

We rationalized these results assuming that, as expected, receptor 1 does not bind the trans-10 rotamer and that, owing to

![Energy minimized structure of the 14⊂1 complex displayed a severe distortion of the cone conformation of the receptor owing to steric clashes between the meso-phenyl substituents and the two included methyl groups of the amide (Fig. 5b).]
the low concentration of the cis-10 rotamer in solution, the formation of the cis-10-C1 complex (in the direct and reverse titrations) takes place to a reduced extent. To verify our hypothesis, we simulated the speciation profile of a direct titration considering $K_{cis-10/10} = 0.01$ and $K_{cis-10/C1} = 1 \times 10^4$ M$^{-1}$. Surprisingly to us, the simulated profile indicated a significant formation of the cis-10$\subset$1 complex in the concentration range used for the experimental titration (ESI†). Owing to the low saturation levels of complex formation attained during the direct and reverse titration of 1 with 10 we could not determine an accurate binding constant value for the cis-10$\subset$1 complex. Nevertheless, we obtained a reasonable fit of the experimental data using the theoretical binding model that considers the equilibrium between rotamers and the exclusive formation of a 1:1 complex with the cis-isomer. The fit was performed by fixing the values of $K_{cis-10/10}$, $\delta_{free}$ and $\delta_{bound}$ (0.01, 5.97 ppm, 6.08 ppm). The only variable to refine was $K_{cis/C1}$. The best fit was obtained assuming $K_{cis-10/10} = 5.0 \times 10^5$ M$^{-1}$. This result suggests a reduction in the binding constant of the inclusion complex of the cis-rotamer of acetamide 10 compared to the isosteric N-methyl-N-phenylformamide 5. Remarkably, the mathematical analysis of the same titration data using the simple 1:1 binding model provided an apparent binding constant of 51 M$^{-1}$.

This example showcases the significant difference in the calculated values of $K_{cis/C1}$ and $K_{app}$ when the cis-rotamer is present in solution at very low concentration. It also serves to justify the use of the elaborate binding model in the estimation or accurate determination of $K_{cis/C1}$ values.

We also determined the cis/trans ratios for the N-aryl acetamide derivatives 11-13 in water. We expected to observe an increase in the amount of the cis-rotamer compared to 10 as was previously described in organic solvents.18 Unfortunately, in water, the percentage of the cis-isomers for the series of acetamides 11-13 did not increase over 3%. This limitation precluded the undertaking of the experimental quantification of the corresponding $K_{cis/C1}$ values for these N-aryl acetamides.

Finally, the affinity constant value of receptor 1 for N,N-dimethyl-acetamide 14 was determined to be $K_{app}(14/C1) = 27$ M$^{-1}$ using a simple 1 : 1 binding model. This value supports the conformational selectivity of the receptor in the binding of cis-9.

It also serves to quantify in three orders of magnitude the binding selectivity expressed by receptor 1 in the binding of N,N-dimethyl-formamide 7 compared to the homologated and more hydrophobic N,N-dimethyl-acetamide 14.

**Conclusions**

In summary, we report the synthesis of a tetra-$\alpha$ aryl-extended calix[4]pyrrole receptor 1 bearing four meso-(3-pyridinium-propyl) groups providing water solubility. We studied the binding of a series of primary, secondary and tertiary formamides and acetamides with receptor 1 in water. The mathematical analyses of the $^1$H NMR titrations of N-phenyl-formamides 3 and 5 using a simple 1 : 1 binding model, $K_{app}$ or a more elaborate theoretical model including the cis/trans isomerization and the exclusive formation of the cis$\subset$1 complex, $K_{cis/C1}$, returned similar values, which were typically larger than 10$^3$ M$^{-1}$. We demonstrated that calix[4]pyrrole 1 selectively binds the cis-rotamers of these two formamides. In contrast, both rotamers of formamides 6 and 7, the N-methyl analogues of 3 and 5, are bound by receptor 1 without substantial changes in affinity values.

**Conflicts of interest**

There are no conflicts to declare.

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**Notes and references**

† The results of the binding studies of related water soluble aryl-extended calix[4] pyroles with creatinine and a homologous series of lactams will be reported elsewhere.

§ Some foam appeared in the solutions after vigorous shaking at high concentrations indicating the amphiphilic nature of tetra-pyridinium calix[4]pyrrole 1. In order to avoid misleading results, we performed our experiments at concentrations below 5 mM.
The pyridinium proton signals also displayed chemical shift changes upon addition of incremental amounts of the guest. However, we decided not to use these nuclei to determine binding constants from the $^1$H NMR titration data because they are not close to the receptor’s binding site.

A broad signal resonating at $\delta = 5.32$ ppm was assigned to the formyl proton of bound cis-3. The computationally determined chemical shift value (DFT) of this proton in the inclusion complex cis-3 $\subset$ 1 is in complete agreement with the assignment.

** Using DFT calculations we computed the chemical shift values of the protons for bound cis-5. We were gratefully surprised to find a nice agreement with the experimental ones (ESI).

†† The chemical shift value of the beta-pyrrole protons in the cis-5 $\subset$ 1 complex was considered to be a good estimate for the $\delta_{\text{bound}}$ value in cis-10 $\subset$ 1.

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