



A new water-soluble cavitand with deeper guest binding properties

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| Complete List of Authors: | Rahman, Faiz-Ur; Shanghai University Feng, Hai-Na; Shanghai University Yu, Yang; Shanghai University, |
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imidazole at 90 °C and the water-soluble cavitant **1** was isolated in quantitative yield. Cavitant **1** was highly soluble (>25 mM) in water and showed a well-resolved ¹H NMR spectrum at 25 mM in D₂O. The cavitants **2** and **1** were obtained in pure forms from the reaction mixture without

chromatography. All precursors and the water-soluble cavitant were characterized by ¹H, ¹³C NMR spectroscopy and high-resolution electrospray mass (HR ESI-MS) spectrometry (ESI Fig. S1-8).

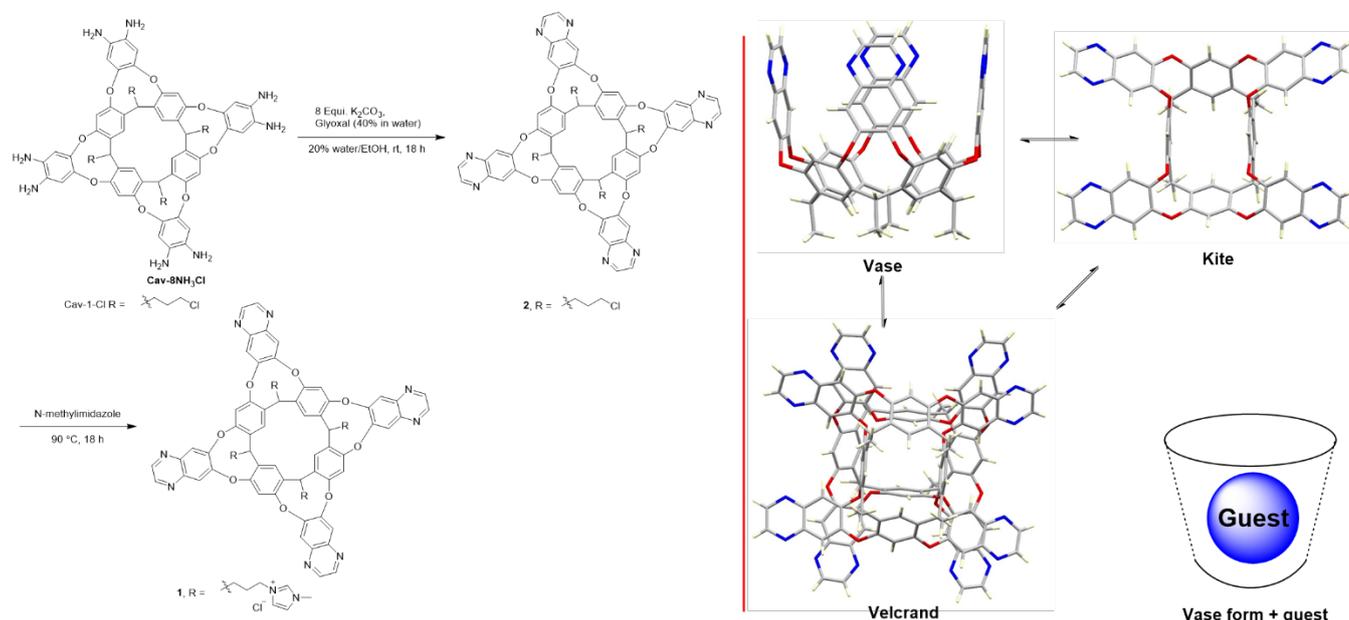


Fig. 2 Route for the synthesis (left) and structural conformations (right) of **1**, and a cartoon showing cavitant/guest complex, velcrand is formed by dimerization of the two molecules in kite form

The vase/kite/velcrand conformations of cavitants can be determined by ¹H NMR spectroscopy. The vase form has its characteristic methine protons chemical shift observed around 5.5 ppm, while the kite form the methine is reported to be below 4 ppm^{20, 29, 30}. The new water-soluble cavitant **1** showed characteristic vase and kite conformation in different solvents as revealed by the ¹H NMR spectra in various solvents such as DMSO, methanol and water (Fig. 3 and ESI Fig. S9-12). In DMSO the characteristic methine proton chemical shift was observed at 5.51 ppm showing only the vase conformation while the other protons showed single sets of chemical shifts. In D₂O the characteristic chemical shift was observed at 3.91 ppm showing exclusively the kite conformation of this cavitant. The other protons showed multiple sets of chemical shifts showing the presence of this conformation in water. In deuterated methanol **1** showed a complex mixture of different conformations where the methine protons chemical shifts could be observed around 5.5, 5.2, 4.4, 4.2 and 3.9 ppm. This showed the existence of multiple conformations (vase, vase intermediate, velcrand and kite) in methanol.

Cavitant **1** was found exclusively in the kite or dimeric kite conformation (velcrand) in water. The most fascinating property of this cavitant was its complexation with a number of various cyclic guests. Upon binding with the guest, the kite form of this cavitant is converted to vase form while forming a cavity in which the hydrophobic guest (cyclic alkane guests) or hydrophobic part of the guest (cyclic acid guests) is captured. As mentioned earlier, upon binding with the guest the characteristic vase conformation methine protons chemical shift of **1** appeared downfield around 5.5 ppm and that is observed at 3.91 ppm for kite conformation. We found that cyclic alkanes could bind to **1** resulting in the vase form of the cavitant that

was studied by ¹H NMR spectroscopy. The binding of the guest is achieved by folding and unfolding of the walls of the cavitant. This uptake and release of the guest is influenced by several factors that are well documented in previous studies with other related cavitants (e.g. **3**) including

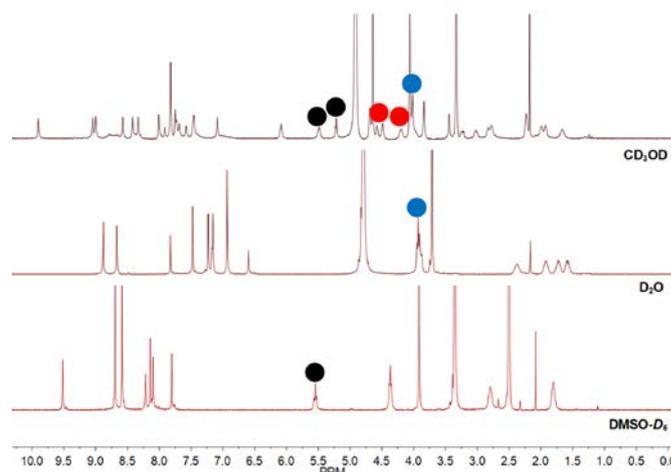


Fig. 3 Effect of solvents on the vase kite form of **1**; 1 mg of **1** was dissolved in 0.5 mL of particular deuterated solvent (from bottom to top, DMSO-*d*₆, D₂O, CD₃OD) and analysed by ¹H NMR spectroscopy. Black dots represent methine protons of vase form or vase intermediate form, red dots represent methine protons of velcrand form and blue dots represent methine protons of kite form.

the hydrophobicity of the guest, solvent size and polarity. The sparingly or non-soluble cyclic alkane guests showed good occupation of the small cavity formed by **1**. The protons

chemical shifts for all these bound guests were observed upfield in the negative region of the spectrum caused by the shielding of the aromatic panels of the walls of **1**. Each guest acquired a specific position in the cavity depending on its size. A single chemical shift was observed for all protons of these guests indicating fast rotation of each cyclic compound in the cavity. This also showed that the guest is not fixed in the cavity but remains in constant motion. Bound cyclohexane signals were observed at - 3.34 ppm. Free cyclohexane's chemical shift in the absence of cavitand is assigned at 1.40 ppm while upon binding in **1** it is shifted up field $\Delta\delta = - 4.74$ ppm. This is a larger value as compared to binding with **3**³¹ (i.e. $\Delta\delta = - 3.58$ ppm) (ESI Fig. S21-22). This larger upfield shift in **1** is possibly due to the deeper binding of the guest or the higher aromatic walls of the host. With an increase in the size of the cycloalkane, the chemical shifts were shifted upfield. In the case of cyclodecane, it was observed at - 2.24 ppm, showing a higher average position of the guest methylenes in **1** as compared to cyclohexane. The integration of the host methine protons and the guest methylene protons showed a 1:1 complex stoichiometry (Fig. 4 and ESI Fig. S11-14).

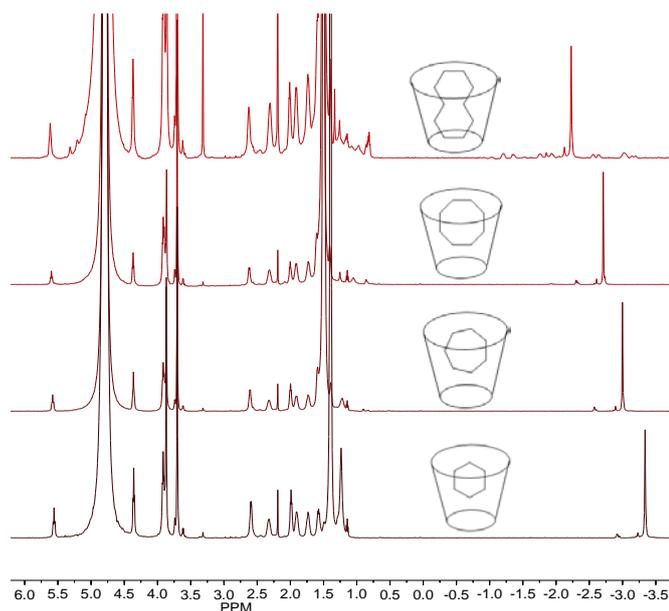


Fig. 4 Spectra of cavitand **1** in D₂O with bound guests. From bottom to top: **1** + cyclohexane, **1** + cycloheptane, **1** + cyclooctane, **1** + cyclodecane, excess of pure guest was added to 0.5 mL, 1 mM solution of **1** in D₂O, sonicated for 1 h and analyzed by ¹H NMR spectroscopy at rt.

We also tested the binding of various carboxylic acid guests and found that even water-soluble acids would be taken up by **1**. The proton signals for these bound guests were observed in the upfield region due to the anisotropic shielding provided by the aromatic walls. Each guest bound in a similar fashion with alkyl part inside the cavity at a depth corresponding to its size, while keeping the most hydrophilic function (COOH) exposed to the water solvent. This was also observed previously with the cavitand **3**³¹. The terminal methyl groups of 3-methylbutanoic acid showed a chemical shift at - 4.12 ppm showing their averaged position deep in the cavity, while for other acid guests the deepest methylene chemical shift was found above - 4.0 ppm. Accordingly, the methylenes are unable to reach as deep as the methyls of 3-methylbutanoic acid. The integration of the host methine proton and the guest protons showed a strict 1:1 complex formation (Fig. 5 and ESI Fig. S15-20). Cavitand **3** was extensively studied in capturing different small organic molecule guests in D₂O and formed 1:1 host guest

complexes. The size and hydrophobicity of the guest determined the depth of binding of the guest in the cavity of **3** (ESI Fig. S21-31). Interestingly, we found the binding of these organic small molecule guests showed further upfield shifts in the cavity of **1** when compared to those observed for **3**. In addition, water-soluble cyclopentyl carboxylic acid bound to **1** but not to **3** (ESI Fig. S26), indicating a somewhat stronger affinity of **1**. This property of **1** will be explored for other related guests and may bring applications in selective recognition of such small molecules.

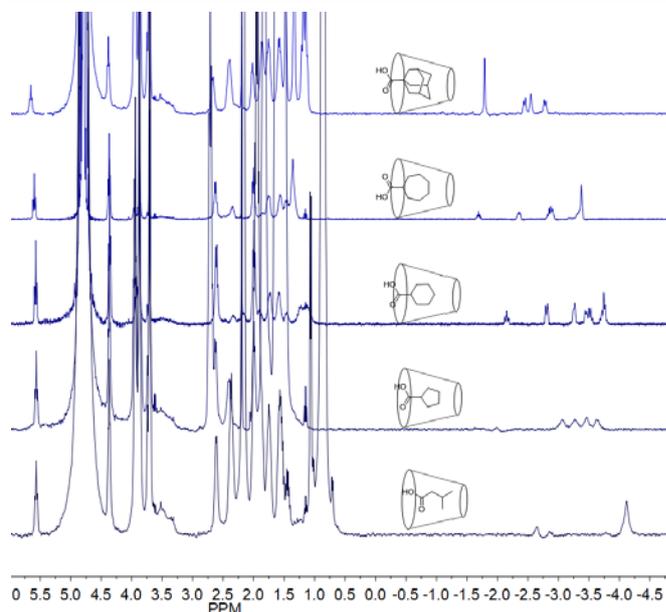


Fig. 5 Spectra of complexes of **1** in D₂O and guests from bottom to top: **1** + 3-methylbutanoic acid, **1** + cyclopentylcarboxylic acid, **1** + cyclohexylcarboxylic acid, **1** + cycloheptylcarboxylic acid, **1** + adamantancarboxylic acid, excess of pure guest was added to 1 mM solution of **1** in D₂O, sonicated for 1 h and analyzed by ¹H NMR spectroscopy at rt.

In summary, a new and deeper cavitand that showed good solubility in water was prepared and characterized. Depending on the solvent, this cavitand existed in vase, kite or multiple forms. This water-soluble cavitand showed binding properties for small organic molecule guests such as cyclic alkanes and carboxylic acids. The easy synthesis, purification and good water solubility make it an improved platform for catalysis, sensing and molecular recognition in aqueous environments. The presence of N-donor atoms near the top of the quinoxaline rim may further be used for the coordination of metals for catalysis reactions of bound guest.

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Conflicts of interest

The authors declare no conflict of interest.

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