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HIGHLIGHTS

Synthesis of urea-based tripodal ligands, with propeller-like appearance, is described

These structures form hydrogen bonding networks with SO_4^{2-}

These self-assemblies give rise to highly ordered layered materials

The formation of hierarchical structures, including nano-sized particles, is observed

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Sulfate-Templated 2D Anion-Layered Supramolecular Self-Assemblies

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SUMMARY

Using solution and solid-state analyses, we demonstrate that the tripodal *N*-methylated(1,3,5-benzene-tricarboxamide)-tris(phenylurea) BTA ligands, possessing urea functionalities in the *meta* position, are able to form extended self-assembly 2D networks via hydrogen bonding templated by sulfate (SO_4^{2-}). The divergence of the urea binding sites confers a propeller-like conformation to the ligands and is key to formation of the self-assemblies. Studies in solution and in the solid state as well as scanning electron microscopy (SEM) on the self-assembly properties of the ligands showed that the convergence also leads to the formation of hierarchical structures, including porous films and spherical nano-sized particles; the morphological outcome is highly solvent dependent.

INTRODUCTION

Supramolecular anion chemistry is a topical area of research with a large number of organic and coordination structures developed for the recognition, sensing, and transport of anions.^{1–3} Anions play a critical role as counter ions in the formation of coordination networks and MOFs⁴ and in directing self-assembly processes, such as in the formation of clusters,⁵ metallo-macrocyles and cages,⁶ helicates,⁷ knots,⁸ and mechanically interlocked structures.⁹ Their application in supramolecular gel formation is also well studied.^{10,11} In contrast, examples of anions directing the formation of higher order organized self-assemblies and functional hierarchical materials from charge neutral organic building blocks is limited. Noteworthy examples include that of Basarić et al.,¹² who used phosphate to mediate the formation of self-assembly structures from adamantane bisurea derivatives and that of Das et al.¹³ who used carbonate to induce the formation of dimeric capsular hexagonal assemblies. Recently, Bielawski et al.¹⁴ used bis-anions in the formation of anion cross-linked soft materials, while Granja et al.¹⁵ formed higher order organized self-assemblies of spherical aggregates from cyclic peptides. In the last decade, several examples of anion coordination polymer (ACP) have also been developed which normally give 1D structures^{16,17} and other anion based polymeric systems.^{18,19} In fact, anion supramolecular self-assembly chemistry has become an import and fast growing area of research.^{20–22} The BTA (1,3,5-benzenetricarboxamide) moiety is a tripodal building block that has been extensively used in the formation of a range of supramolecular structures and materials.²³ Recently, Stefankiewicz et al. used BTA ligands in the formation of a large enantiopure nanocapsule as a host for C_{60} .²⁴ We have recently shown that the BTA scaffold can form anion-templated self-assembly cages using SO_4^{2-} in 2:1 stoichiometry and self-sorting self-assembled clusters with 4:4 stoichiometry.²⁵ Here, we demonstrate that the high coordination requirement of SO_4^{2-} and its templating ability can be capitalized upon in the formation of a supramolecular layered polymeric 2D network, and that

The Bigger Picture

Anions are widespread in our world, and many anions carry out specific roles in nature, for both living and inanimate matter. Because these processes are often accompanied by specificity, when changes occur in these interactions, the results can be of harm to life and the environment. For this reason, the study of the noncovalent bonds of anions has been of interest to scientists for many years, with a focus on hydrogen bonding, imitating the way proteins and enzymes can interact with anions in nature.

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A deep understanding of these interactions is needed to comprehend how their activity can be actively used toward the formation of hierarchical assemblies to control their functions and activities in their environment.

The development of aniontemplated self-assemblies is of interest for their potential in the removal of hazardous and polluting anions from the environment using new improved methodologies.

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Figure 1. Structure of Tripodal Ligands 1–3

(A) Structures of 1-3.

(B) The X-ray crystal structure of **2**, viewed from the **BTA** core demonstrating the open propeller conformation with divergent urea binding sites. Hydrogen atoms and solvent molecules omitted for clarity.

these self-assemblies form nano-sized hierarchical aggregates in solution. We demonstrate by using SEM imaging that the nature of the solvent mixture dictates the way that these 2D networks aggregate to form higher-ordered hierarchical materials, the morphology of which are highly solvent dependent.

RESULTS AND DISCUSSION

Synthesis and Characterization

Inspired by the ability of the SO_4^{2-} anion to form anion-templated capsules, the N-methylated BTA structures 1-3, Figure 1A, possessing three inner-core phenyl urea groups substituted in the meta position were synthesized in three steps. Starting from 1,3,5-benzenetricarbonyltrichloride and N-methyl-3-nitroaniline, which were coupled to give a tripodal nitro precursor, which, once catalytically reduced to the corresponding amine, was reacted with the relevant phenyl isocyanate to give the desired ligands (see Scheme S1 and Supplemental Information for full details). Our objective was to investigate whether these ligands could be used to direct self-assembly formation away from the "capsule" structure¹⁸ toward the formation of higher-ordered hierarchical structures. Slow evaporation of DMSO and CH₃CN solutions of 2 and 3, respectively, resulted in the formation of diffraction-quality single crystals. The crystal structures are shown in Figures 2A-2D. Unfortunately, attempts to grow suitable crystals of 1 failed. Ligand 2 crystallized in the triclinic P-1 space group and showed the three N-methylamido groups of the BTA unit to be directed away from the plane of the central benzene core (Figures 1B, 2A, and 2B). The three amide oxygen atoms bend away from the plane defined by the central benzene ring with torsion angles of ca. 43° for two arms and 62° for the third. The coplanar conformation of two of the arms is further stabilized by low angle, ca. 120°, hydrogen bonds between aromatic C-H donors, and the acceptor oxygen atom of the ureas (Table S7). These interactions as well as $\pi - \pi$ interactions stabilize the conformation of 2,4-difluorophenyl moiety which, in principle, can rotate freely. These two ureas are also engaged in hydrogen bonding with two DMSO molecules, reflecting solvation of the potential binding site of the receptor. The third arm exhibits a wider angle because of bifurcated intermolecular hydrogen bonding of the urea with another arm of a neighboring molecule of 2. The crystals of 3 were also formed in the triclinic P-1 space group, with the asymmetric unit containing one molecule of ligand and

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Figure 2. Crystal Structures of 2 and 3

Structure of 2. Shown in (A) is a capped stick representation demonstrating the open propeller conformation, with divergent urea binding sites, that are involved in H-bond with 2 solvent molecules (DMSO), and in (B) is the packing showing hydrogen bonding from one of the urea groups NH to the BTA C=O on a neighboring molecule. Shown is the structure of 3: in (C) is a capped stick representation showing the formation of internal hydrogen bond from one urea NH and the Ortho NO₂ group and in (D) is packing showing hydrogen bonding from two of the urea groups NH (the one not involved in the intramolecular H-bond) to the BTA C=O on a neighboring molecule. Hydrogen atoms omitted for clarity.

some solvent molecules. As was seen for 2, the central BTA moiety adopts a conformation where all three amide oxygen atoms locate on the same side of the central ring, and a range of different supramolecular interactions (intermolecular hydrogen bonds, π – π stacking, and CH– π bonds) are evident in the crystal packing. Furthermore, an intramolecular hydrogen bond between the nitro-substituted distal phenyl ring and urea NHs exists for all three arms (Figure 1C). We²⁶ and others²⁷ have demonstrated such interactions in structurally similar urea-based receptors. As was observed for 2, intermolecular hydrogen bonds between the arms of neighboring molecules also exist in 3 and give rise to a 1D network in the crystallography a axis (Figure 2D).

Anion Binding Studies

Anion Binding Studies in Solution

The solid-state analysis above demonstrated the presence of the anticipated open "propeller-like" conformation for both 2 and 3 and suggests that this preorganization may be appropriate to form anion-templated networks. With this in mind, we investigated the affinity of 1–3 for different anions (tetrabutyl-ammonium salts of AcO^- , SO_4^{2-} , $H_2PO_4^-$, and CI^{-})²⁶ in CH₃CN solution using UV-vis absorption spectroscopy. The absorption spectra consisted of a broad band at high energy for the BTA core, and a shoulder for 1 and 2, or a band for 3, that was significantly red shifted, because of internal charge transfer. All the anion titrations clearly demonstrated that 1–3 participated in anion binding, with concomitant formation of

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different isosbestic points (Figures S58–S69). However, the overall changes were in some cases relatively small, and the titration data could not be fit easily to give reliable anion binding constants (see Table S1 for fitting). Therefore, titrations were carried out in DMSO- d_6 solutions and followed using ¹H NMR spectroscopy.

Full NMR characterization of 1–3 was completed using 2D NMR experiments (i.e., HMBC, C-H, and N-H HSQC) combined with selective ¹H ROESY, which allowed us to identify, and confidently monitor, the shift of each proton during the binding events. Titration of these ligands with the aforementioned anions caused de-shielding to occur in the urea proton resonances, indicating hydrogen bonding interaction with the anions. From these changes, the anion binding stoichiometry and constants could be determined using the non-linear regression analysis program HypNMR.^{28,29} The analysis showed the formation of 1:1 complexes (ligand 1: $K_{1:1} = 32,200$ and $8 M^{-1}$ for Cl⁻, $H_2PO_4^{-}$, and SO_4^{2-} , respectively; for other ligands see Table S2), while with AcO- ligands 2 and 3 formed the 1:1, and ligand 1 showed the formation of two species, 1:1 and 1:3. This behavior is in line with the Hofmeister series and the solvation of the anions. The formation of 1:1 complexes for Cl⁻, SO_4^{2-} , and $H_2PO_4^{-}$ was also confirmed by MALDI-TOF analysis (Figures S37-S41), with the isotopic distribution patterns matching those of the calculated species. Unfortunately, ligand 1 was the only host that displayed sufficient solubility in CD₃CN for ¹H NMR titrations to be carried out. In this media, all anions induced a greater downfield shift in the urea proton resonances than seen in DMSO- d_6 and less selective binding was observed (SO $_4^{2-}$ and Cl⁻ showed shifts that were 2 and 4 times those seen in DMSO- d_6) (Figure 3). This reflects their stronger interactions in less competitive media (Table S3). These titrations were fit to the same binding models as employed above, showing both 1:1 and 1:3 (L:A) stoichiometry for Cl⁻, SO_4^{2-} and $H_2PO_4^{-}$ (Table S3). Having determined the binding affinity of 1–3 and possible species distributions, we next set out to investigate if these ligands, in the presence of high-coordinating oxyanions, could be used to form extended anion-templated self-assemblies as outlined above (in either solution or in the solid state).

Solid-State Studies: Formation of Anion-Templated 2D Materials

The crystallization of 1-3 in the presence tetrabutyl-ammonium salts of SO_4^{2-} , AcO⁻, H₂PO₄⁻, and Cl⁻ was carried out in both DMSO or CH₃CN solutions, using various methodologies. However, of these, only 2 in the presence of SO_4^{2-} yielded crystals. The slow evaporation of DMSO or CH_3CN solutions of 2 (in excess SO_4^{2-}) resulted, on both occasions, in the formation of crystals that were suitable for single crystal X-ray diffraction analysis (Figure 2) and the structural details showed them to be isostructural (Table S4). The complex of 2 with SO_4^{2-} (2 · SO_4^{2-}) crystallized in the trigonal R-3 space group, with each of the SO_4^{2-} anions positioned on a hexagonal axis, within a tight hydrogen binding pocket. The asymmetric unit contains 1/3 of a molecule of 2, 1/6 of a SO_4^{2-} anion. Solvent molecules and the counter cations were present, however, these were disordered and not interacting with their host receptor or guest anion. Each SO_4^{2-} ion is surrounded by six urea groups, from six different molecules of 2 (Figures 4A-4D) with a total of 12 hydrogen bonds (Figure 4D) with N-H...O(S) bond lengths ranging from 1.923-2.240 Å, with an average N-H...O bond length of 2.22 Å (Table S10). Within the $2 \cdot SO_4^{2-}$ assembly, each ligand binds to three different SO_4^{2-} ions (Figures 4B and 4C). This demonstrates that the high coordination preference of the anion is fulfilled within this binding arrangement.³⁰ In contrast to the structure of 2 (cf. Figure 2A), in $2 \cdot SO_4^{2-}$, all three 2,4-difluorophenyl moieties are equally rotated at 77° with respect to the urea groups, the amide carbonyls and the N-methyl groups pointing away from the central C₃ axis, and it is

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Figure 3. ¹H NMR Binding Studies of Ligand 1

(A–D) Binding isotherms of urea protons comparing results between CD₃CN (blue) and DMSO-d₆ (green) titration of 1 with (A) Cl⁻, (B) CH₃COO⁻, (C) SO_4^{-2-} , (D) H₂PO₄⁻.

clear that the preorganization observed in 2 is to greater extent maintained in $2 \cdot SO_4^{2-}$, where 2 maintains its "propeller-like" conformation. The anion templation results in an overall C_3 symmetry, which tessellates the ligands around the SO_4^{2-} anion, resulting in the formation of an extended 2D layer along the ab plane. These planes are stacked along the c axis, each one rotated +/- 120° with respect to surrounding planes. This network is further stabilized by inter-ligand π - π stacking and additional hydrogen bonding interactions. The π -stacking interactions differ from the ligand crystal where face-to-face interactions were seen between distal phenyl rings from the neighboring molecules. This places the ortho-fluorine atoms in one of two sites within the structure, with a similar occupancy distribution to the ligand crystal, where 32% has a potential aromatic CH...O(amide) hydrogen bond in place and 68% in the alternate conformation. The SO_4^{2-} anion can be found in two different orientations along the c axis, up or down, producing two crystallographically different oxygen atoms in axial or equatorial orientations. One of the orthofluorine atoms is situated very close to this axis, so it has two symmetry generated F at 2.54(3) Å, suggesting the possibility of a F-F interaction (Figure 4E), as we³¹ have observed in other self-assembly structures.³² The above results demonstrate that our design strategy has been successful, and that the "propeller-like"

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Figure 4. Crystal Structure of 2 SO₄²⁻

(A) Packing details viewed down the c axis (only one layer shown) and schematic representation of the 2D network formed by the SO_4^{2-} -templated self-assembly of the propeller-like molecule.

(B) Detail of a cluster formed by 6 molecules of 2 around 1 molecule of SO_4^{2-} .

- (C) Details of the interaction of 1 molecule of 2 interacting with three SO_4^{2-} molecules.
- (D) Detail of 1 molecule of SO_4^{2-} interacting with 6 different molecules of 2.

(E) Details of the F–F interactions between three different molecules.

preorganization of 2 results in the formation of an extended 2D network upon binding to SO_4^{2-} (Figure 4A). This is, to the best of our knowledge, the first example of the use of SO_4^{2-} anion to generate such a higher order 2D self-assembly in preference to the more commonly seen capsule formation.^{24,25,33}

Self-Aggregation Studies

Aggregates and hierarchical materials such as extended 2D and 3D networks are formed because of inter- and intra- molecular supramolecular interactions and solvent effects.²³ Given that the $2 \cdot SO_4^{2-}$ assembly gives rise to crystalline 2D networks,

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and that 1-3 and $1-3 \cdot SO_4^{2-}$, all have the ability to partake in such extended supramolecular network formation, we investigated the hierarchical formation for 1-3 in solution, both in the absence and presence of anions. ¹H NMR dilution studies (5 x $10^{-6} \rightarrow 1 \times 10^{-2}$ M) were first undertaken and these showed that in DMSO- d_6 only minor changes were observed for the free ligands. However, such dilution studies in the presence of SO_4^{2-} clearly showed aggregation effects occurring for $1 \cdot SO_4^{2-}$ and $2 \cdot SO_4^{2-}$ (Figures S81 and S85). In the case of 1, aggregation effects were seen in both CD₃CN and CD₃OD, with sharpening and shifts in key resonances observed at higher concentrations (Figures S82-S83). Next, solutions of 1 $(1 \text{ mM in DMSO}, CH_3CN, and CH_3OH), 2$, and 3 $(\text{in DMSO})^{34}$ were deposited on silica plates; these were dried in vacuum and imaged using scanning electron microscopy (SEM). The formation of thin films was observed in the SEM for 1-3 in DMSO (Figure S87) and for 1 CH₃CN, Figure 5A. However, for ligand 1 in CH₃OH, the formation of micro-spherical aggregates of ca. 1–2 μm size was observed, Figure 5B. 35 It is clear from the SEM imaging that the solvent has a major effect on the supramolecular interactions of these 2D structures, which results in the formation of the hierarchical structures. While it is difficult to elucidate what self-assembly pathways these 2D structures take to form these higher order structures, it is clear that the polarity, and hydrogen bonding ability, of the solvent is an important factor. As 1 is hydrophobic in nature, we investigated the effect of H_2O on the assembly process of 1 in CH_3OH by adding various aliquots of H_2O to the 1mM solution of 1 in CH_3OH above. This instantly generated off-white precipitates that would remain dispersed in solution for days. Samples of these suspensions were analyzed by SEM and all showed the formation of spheres, Figure 5C. However, unlike that seen from pure CH₃OH solution, these were more uniform and nano-sized (ca. 150-300 nm), demonstrating the effect of water on the aggregation pathway, and that the ligand alone can form hierarchical material. The formation of the spherical structures of 1 was also observed at even lower concentrations. This prompted us to monitor their formation as a function of both concentration of 1 (between $10^{-6} \rightarrow 10^{-5}$ M) and in different H₂O:CH₃OH compositions, using UV-vis absorption spectroscopy. The results showed that at 8×10^{-6} M concentration of 1, the aggregate formation instantaneously occurred upon addition of H₂O and the absorption of 1 at 263 nm, being initially slightly blue shifted (0% \rightarrow 50% H₂O), was followed by a red shift (265 nm) with the formation a broad shoulder (~300 nm) (Figure 5E). DLS studies of these samples confirmed their monodispersed nature and size distribution between 150-350 nm (Figure 5F).

Next, we investigated what effect SO_4^{2-} (1 equiv) would have on such hierarchical formations. The addition of SO_4^{2-} to DMSO solutions of 1 and 3 resulted in the formation of aggregates from deposited solutions, these clearly being formed as discrete microspheres for 3 (Figure S88), while 1, consisted of "film" covered particles (Figure 6A). In contrast, 2 showed the formation of microcrystals within a film structure (Figure 6C). Addition of H₂O (50%) to these solutions showed the formation of smooth spherical particles for 1 (Figure 6B) and 3 (Figure S88), while 2 showed a spherical aggregate with a microcrystalline texture (Figure 6D). We also added SO_4^{2-} to preformed spheres of 1 in 1:1 CH₃OH:H₂O solutions causing the precipitation of the microparticles. SEM imaging of these showed that further aggregation of the spheres occurred (Figure 5D), giving rise to the formation of "coral"-like aggregates. This clearly demonstrates that the 2D self-assembly material forms alternative hierarchical structures in highly competitive hydrogen bonding media, which could possibly be due to the combination of hydrophobic effects between the layered material and the solvent, as well as a degree of hydrogen boning interactions with the urea oxygen atoms. Although we have not been able to prove this experimentally, it is clear that the polarity of the urea arms (e.g., due to the nature

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Figure 5. Self-Assembly Studies of 1 in CH₃CN and CH₃OH

(A–D) SEM images of self-assembled nanospheres of 1 in: (A) CH₃CN (bar; 1 μ m), (B) CH₃OH (bar; 2 μ m), (C) 1:1 CH₃OH-H₂O mixture (1 wt %) (bar; 200 nm), (D) 1:1 CH₃OH -H₂O mixture (1 wt %) and 3 equiv of SO₄²⁻ (bar; 200 nm).

(E) UV-vis plot; (inset) plot of absorbance at 262 nm against H_2O content in solution. Ligand concentration 8 × 10^{-6} M.

(F) Average particle size distribution determined using DLS technique from suspensions of ligand 1, prepared in CH₃OH:H₂O in different ratios (c = 8 × 10^{-6} M, at 25° C).

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Figure 6. Self-Assembly Studies for 1 and 2 in DMSO

(A and B) SEM images of deposited solutions of **1**. SO_4^{2-} in: (A) DMSO (scale bar, 20 μ m), (B) 1:1 DMSO-H₂O (scale bar, 10 μ m, inset = 1 μ m).

(C and D) SEM images of deposited solutions of **2.** SO_4^{2-} in: (C) DMSO (scale bar, 2 µm), (D) 1:1 DMSO-H₂O (scale bar, 2 µm).

of the substituents; e.g., CF_3 versus F) allows us to gain some degree of tuning of the morphological outcome.

Using Cl⁻, AcO⁻, and H₂PO₄⁻ caused similar effect (Figure S89). In contrast, SEM images of samples where $SO_4^{2^-}$ was first added to a solution of 1 in CH₃OH solution, followed by the addition of H₂O, gave no evidence of sphere formation, and only the formation of films, similar to that seen above, was observed (Figure S90). These studies showed that the self-assembly properties of the ligand would be highly affected by the solvent, which can influence both the self-assembly of the free ligand and of its $SO_4^{2^-}$ complexes.^{36,37}

Conclusions

In summary, we have demonstrated that supramolecular self-assembly networks and aggregates, including uniform nano- and microspheres, can be formed from *N*-methylated BTA structures possessing *meta* phenyl urea groups, in both the presence and absence of SO_4^{2-} . Using X-ray crystallography, we showed that $2 \cdot SO_4^{2-}$ formed a 2D hydrogen bonded network, where each anion is bound in a symmetrical way to six 2 "hosts". To the best of our knowledge, these are the first examples of such SO_4^{2-} based or connected supramolecular self-assembled structures hierarchical materials using BTA based urea ligands, demonstrating the powerful role anions can have in generating novel functional material towing to its high coordination requirement.

EXPERIMENTAL PROCEDURES

All solvents and chemicals were purchased from commercial sources and used without further purification. Deuterated solvents used for NMR analysis (CD₃CN, DMSO- d_6) were purchased and used as received. Silica chromatography was carried out on a Teledyne Isco CombiFlash automated machine using pre-packed RediSep

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cartridges. Infrared spectra were recorded on a Perkin Elmer Spectrum One FT-IR spectrometer fitted with a universal ATR sampling accessory. Melting points were determined using an Electrothermal IA9100 digital melting point apparatus. Where applicable, elemental analysis (CHN) was carried out by the School of Chemistry, University College Dublin.

Mass Spectroscopy

Mass spectrometry was completed in the departmental mass spectrometry service of the School of Chemistry, Trinity College Dublin. Electrospray mass spectra were measured on a Micromass LCT spectrometer calibrated against a leucine enkephalin standard. MALDI Q-ToF mass spectra were recorded on a MALDI Q-TOF Premier (Waters Corporation, Micromass MS Technologies, Manchester, UK) and high-resolution mass spectrometery was performed using Glu-Fib as an internal reference (m/z = 1,570.677).

Dynamic Light Scattering

DLS measurements were made from back scatter in using Malvern Instrument Zetasizer Nano Series at 25°C in quartz cuvette with a path length of 10 mm. Statistical distribution by intensity were determined using the instrument software with standard deviations determined from triple replicate measurements. Samples were prepared using HPLC-grade CH₃OH and H₂O filtered six times through a 0.45 μ m syringe filters (Acrodisk, PTFE).

UV-Vis Absorption Studies

UV-visible absorption spectra were recorded using a Varian Cary 50 spectrophotometer, a spectroscopic window of 450–200 nm was used for all spectra with applied baseline correction from blank solvent. All UV-vis measurements were performed at 298 K in CH₃CN or DMSO:CH₃CN (spectroscopic grade) solutions. UV-vis absorption spectra were measured in 1 cm quartz cuvettes, except for the aggregation measurements for which 0.2 cm quartz cuvette was used. Baseline correction was applied for all spectra. Spectroscopic solutions were prepared from stock solutions using Pipetman Classic micropipettes (Gilson, Inc.).

Preparation of Solutions for UV-Vis Titrations with Anions

The TBA salts of the anions used in the titrations were of spectroscopic grade and were purchased from Sigma Aldrich. All TBA salts were dried over P_2O_5 at room temperature under vacuum. Solutions of these salts were prepared at varying concentrations of ca. 0.1 M, 5 × 10⁻² M, 1 × 10⁻² M, and 1 × 10⁻³ M. Host solutions of ca. 1 × 10⁻³ M were prepared and then diluted before titration (ca. 1 × 10⁻⁵ M).

¹H NMR Studies

The ¹H NMR spectra were recorded at either 400 or 600 MHz on Bruker instruments (AV3-400 or AV2-600). ¹³C NMR spectra were recorded, on the same instruments, at either 150.9 or 100.6 MHz, while ¹⁹F NMR spectra were recorded at 376.5 MHz. All ¹³C NMR spectra were decoupled from ¹H, couplings to other nuclei are specified where appropriate. All spectra were recorded in commercially sourced per-deuterated solvents and referenced to residual proton signals of those solvents. Recorded free-induction decay signals were Fourier-transformed and processed using MestreNova v.6 without apodization and chemical shifts expressed in parts per million (ppm/ δ) and coupling constants (*J*) in Hz. Residual solvent resonances were used as the internal reference, while 2D spectra were graphically referenced. All NMR spectra were carried out at 298 K.

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Preparation of Solutions for ¹H NMR Titrations with Anions

Spectroscopic solutions were prepared from stock solutions using Pipetman Classic micropipettes (Gilson, Inc.). The concentrations of these solutions were prepared in deuterated solvent such that 2 μ L would give ca. 0.1 molar equivalents of the anion. Host solutions of 7 × 10⁻³ M were prepared by dissolving a known amount of host in 0.7 mL of deuterated solvent. All measurements were performed at 25°C in CD₃CN. Receptors were titrated with a 100-fold more concentrated solution of the anion, as TBA⁺ salt. After each addition of the titrant solution, the ¹H-NMR spectrum was recorded. TBA salts were dried in vacuum over P₂O₅. The ¹H NMR spectra were recorded at 400 MHz on Bruker instruments AV3-400.

SEM

Microscopy analysis of gel-aggregate samples by SEM was carried out using the facilities of the Advanced Microscopy Laboratory (AML) in Trinity College Dublin. Samples were prepared by a drop-casting methodology onto clean silicon wafers (cleaned by sonication in HPLC-grade acetone followed by HPLC-grade propan-2-ol) from glass Pasteur pipettes.²⁴ The manually drop cast samples were dried overnight in ambient conditions (a week in the case of DMSO solutions) and under high vacuum for 2 h immediately prior to their imaging. Samples were coated with a conductive Au layer, by sputtering, in order to improve contrast where static charging interfered with the imaging. Low kV SEM was carried out using the Zeiss ULTRA Plus using an SE2 detector.

X-Ray Crystallography

X-ray data were collected on a Rigaku Saturn 724 CCD Diffractometer using graphitemonochromated Mo-K α radiation (λ = 0.71073 Å). The data were collected using Crystalclear-SM 1.4.0 software. Data integration, reduction, and correction for absorption and polarization effects were all performed using Crystalclear-SM 1.4.0 software. Space group determination was obtained using Crystal structure version 3.8. Data for structure ($2 \cdot SO_4^{2-}$).22 was collected on an APEX Duo diffractometer using Cu-K α radiation (λ = 1.54056 Å). The data collection, integration, and reduction were made with Bruker Apex II software. The structures were solved by direct methods (SHELXS-97) and refined against all F2 data (SHELXL97). All H-atoms, except for N-H protons, were positioned geometrically and refined using a riding model with d(CHaro) = 0.95 Å, Uiso = 1.2Ueq (C) for aromatic and 0.98 Å, Uiso = 1.2Ueq (C) for CH3. N–H protons were found from the difference map and fixed to the attached atoms with UH = 1.2UN. In some cases, disordered solvent molecules were removed using SQUEEZE program from the PLATON crystallographic software.³⁷

DATA AND CODE AVAILABILITY

X-ray crystallographic data can be obtained free of charge from the Cambridge Crystallographic Data Center (www.ccdc.cam.ac.uk/data_request/cif) under access numbers CCDC: Structure 2: 1874341; structure 2.SO₄⁻: 1874342, and structure 3: 1874344.

SUPPLEMENTAL INFORMATION

Supplemental Information can be found online at https://doi.org/10.1016/j.chempr. 2019.06.023.

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AUTHOR CONTRIBUTIONS

A.B.A. carried out the synthesis, anion binding, and aggregation studies. Crystal structures were solved by S.B. Samples for SEM were prepared by A.B.A. and S.A.J., and images were collected by S.A.J. A.B.A. and T.G. co-wrote the manuscript with support from all authors. All research was performed under the guidance and supervision of T.G. and P.E.K.

DECLARATION OF INTERESTS

The authors declare no competing interests.

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