

Acyclic indole and carbazole-based sulfate receptors

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The anion complexation properties of a series of acyclic receptors consisting of diindolylurea groups appended with amide, amidoindole or amidocarbazole groups have been studied. The receptors selectively bind and encapsulate sulfate *via* either six or eight hydrogen bonds. Receptors containing eight hydrogen bond donors perturb the pK_a of bound dihydrogen phosphate and bicarbonate to the extent that they are deprotonated by free anion in solution.

Introduction

The development of sulfate selective anion receptors is currently an area of intense interest due the important roles this anion plays in biological systems and disease,¹ in hydrometallurgy² and as a pollutant.³ Tripodal tris-urea based systems have been developed by Custelcean and co-workers as selective receptors for SO₄²⁻ with the goal of precipitating the anion from solution.⁴ These species form 2:1 receptor:sulfate complexes binding the anion *via* twelve NH...O hydrogen bonding interactions. Additionally Custelcean, Hay and co-workers have developed self-assembling urea containing cage systems to arrange six urea groups around sulfate.⁵ Sulfate has also been employed by Beer and co-workers in to assemble a variety of interlocked structures.⁶ Macrocyclic sulfate receptors include Sessler's cyclo[8]pyrrole⁷, Bowman-James' cyclic tetraamide/amine based receptor⁸ that forms a sandwich complex with SO₄²⁻ and Kubik's cyclic peptide based molecular oysters.⁹ Despite the anion complexation properties of indole being recognised in biological systems,¹⁰ it was not until 2004¹¹ and 2005¹² that the first reports of the use of indole in synthetic anion receptors appeared. Since then indole has been employed in a number of receptor systems¹³ including a variety of foldamers reported by Jeong and co-workers that exhibit conformational changes in the presence of particular anions.¹⁴ Taking inspiration from this latter work, we decided to synthesise linear receptors that could wrap around tetrahedral oxo-anions such as sulfate but that would possess lower affinities for anions possessing other geometries.

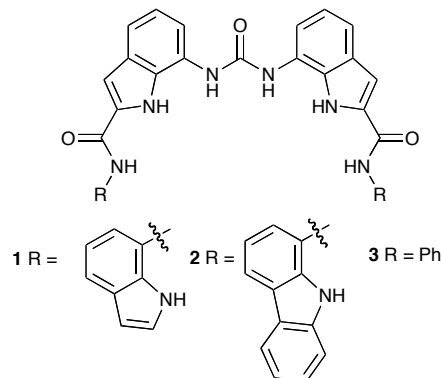
We have previously shown that diindolylureas form 3:1 complexes with PO₄³⁻ (binding the anion *via* twelve hydrogen bonds)¹⁵ in the solid state and in the process drive the deprotonation of the anion.¹⁶ In this paper we report the synthesis of receptors **1** and **2** that contain pendant amidoindole and amido-carbazole groups and their interaction with a range of anionic guests. These compounds are designed to selectively bind sulfate over chloride (due to the large size of the binding site) or carboxylates (due the the carboxylate anion being saturated by four hydrogen bonds from the diindolylurea skeleton¹⁵). We compared the anion complexation properties of these species with compound **3**¹⁶

which contains pendant phenyl amides and a total of six hydrogen bond donor groups.

Results and Discussion

Synthesis

7-Nitroindole-2-carboxylic acid was coupled to either 7-aminoindole or 1-aminocarbazole using pyBOP as an amide coupling reagent in DMF. The resulting amides were reduced from nitro- to amine derivatives using 10%Pd/C under an hydrogen atmosphere in DMF and the resulting amines coupled using triphosgene in a mixture of chloroform, DMF and saturated sodium bicarbonate solution to afford compounds **1** and **2** in 29% and 55% respective overall yields. Compound **3** was synthesised using literature methods.¹⁶



Crystallography

Initial anion complexation studies were conducted using single crystal X-ray diffraction. Crystals of the tetrabutylammonium benzoate complex of receptor **1** were obtained by slow evaporation from a DMSO solution containing excess tetrabutylammonium benzoate. The structure (Figure 1) reveals that receptor **1** binds three equivalents of benzoate in the solid state. The central diindolylurea group binds one benzoate anion *via* four hydrogen bonds N3...O6 2.719(4)Å; N4...O6 2.717(4)Å; N5...O7 2.820(4)Å; N6...O7 2.770(4)Å. The pendant amidoindole groups are oriented out of the cavity with each binding a single equivalent of benzoate *via* two or three hydrogen bonds N1...O4 2.870(4)Å; N1...O5 3.144(4)Å;

N2...O5 2.811(4)Å and N7...O8 2.841(4); N8...O9 2.766(5)Å.

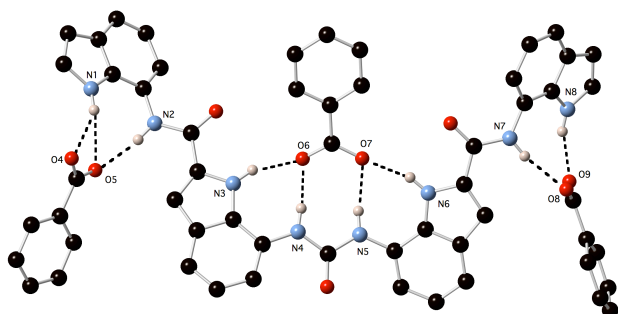


Figure 1 The X-ray crystal structure of **1**.(TBA)₃(C₆H₅CO₂).H₂O. Tetrabutylammonium counter cations and water omitted for clarity.

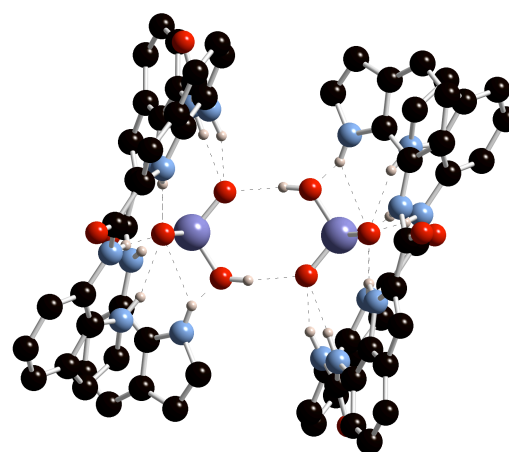
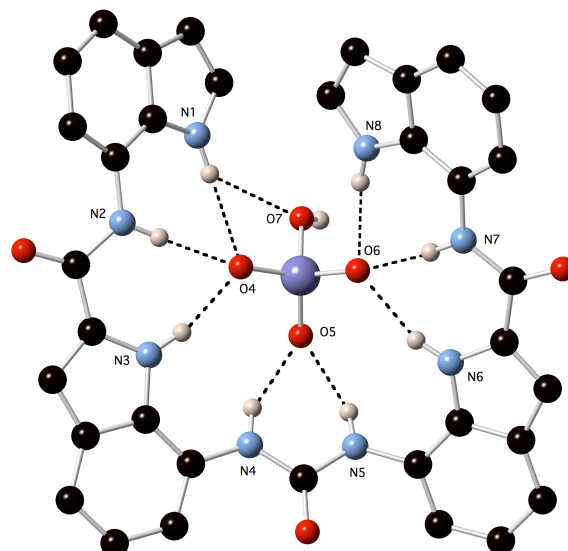


Figure 2 Two views of the X-ray crystal structure of **1**.(TBA)₂(HPO₄). Tetrabutylammonium counter cations have been omitted for clarity.

Crystals of the monohydrogen phosphate complex of receptor **1** were obtained by slow evaporation of a DMSO solution of the receptor in the presence of excess tetrabutylammonium dihydrogen phosphate. The structure (Figure 2) shows the monohydrogen phosphate anion bound by nine hydrogen bonds from the receptor: N1...O7 2.970(8)Å; N1...O4 3.109(7)Å; N2...O4 2.790(8)Å; N3...O4 2.736(8)Å; N4...O5 2.818(8)Å; N5...O5 2.764(7)Å; N6...O6 2.815(7)Å; N7...O6 2.838(8)Å; N8...O6 2.782(7)Å. Additionally the monohydrogen phosphate dimerise in the solid state via two hydrogen bonds O7...O5ⁱ and O7ⁱ...O5 2.645(7)Å the symmetry transformation -x+1, -y+1, -z generates atoms labelled (i). We have previously observed deprotonation of dihydrogenphosphate bound to neutral diindolylurea based receptors due to the multiple hydrogen bonding interactions lowering the pK_a of the bound guest to the extent that it is deprotonated by free dihydrogen phosphate in solution and we suggest that the same process occurs here.¹⁶

Crystals of the tetrabutylammonium sulfate complex of receptor **1** were obtained by slow evaporation of a DMSO solution of the receptor in the presence of excess tetrabutylammonium sulfate.¹⁷ The structure (Figure 3) reveals that the anion is bound by eight hydrogen bonds from the receptor: N1...O6 2.927(4)Å; N2...O4 2.953(3)Å; N3...O4 2.798(3)Å; N4...O5 2.788(3)Å; N5...O5 2.896(3)Å; N6...O7 2.863(3)Å; N7...O7 2.896(3)Å; N8...O6 2.856(3)Å.

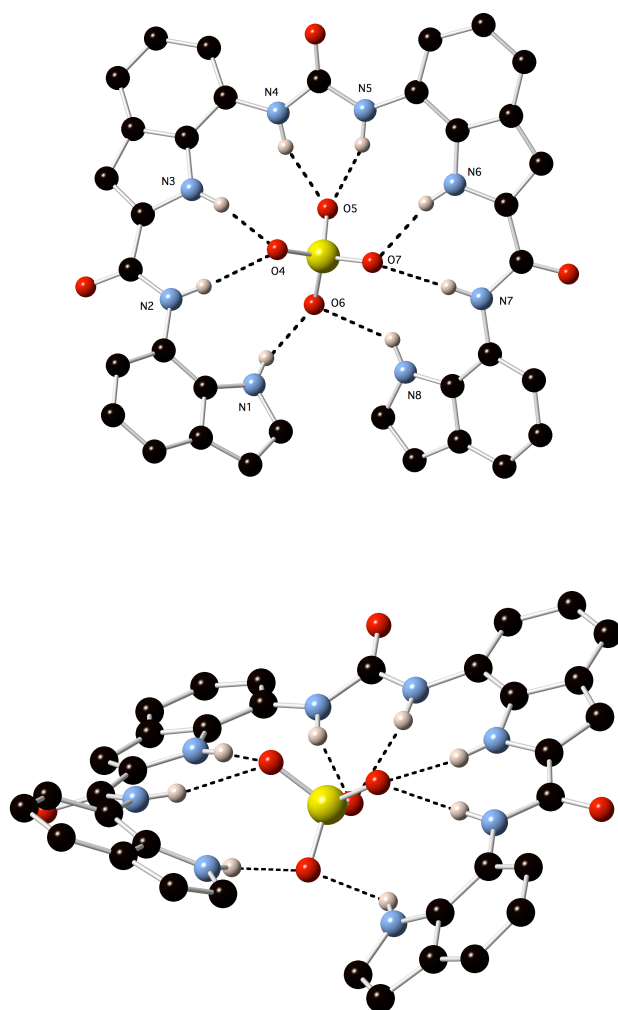


Figure 3 Two views of the X-ray crystal structure of **1**(TBA)₂SO₄. Tetrabutylammonium counter cations have been omitted for clarity.

Solution studies

The solution anion binding properties of compounds **1** and **2** were studied by ¹H NMR titration techniques in DMSO-*d*₆/water mixtures and compared to those of compound **3**. Initial ¹H NMR titration studies were conducted in DMSO-*d*₆/0.5% water. The NMR titrations with compounds **1** – **3** with acetate are shown in Figure 4. The results show that the NH groups in the central diindolylurea unit in these three compounds bind the first equivalent of acetate added to solution. Further aliquots of acetate bind to the pendant amide or pendant amide and/or indole or carbazole groups (as shown in Scheme 1 for compound **1**). This can be seen clearly in the NMR titration curves for compounds **1** and **2** wherein the pendant NH groups begin to shift downfield after the addition of 1.0 equivalents of acetate. In the case of compound **2**, it appears that more complex equilibria are present as after the addition of three equivalents of acetate, the central indole groups begin to shift downfield again after having previously reached a plateau. This process does not occur in DMSO-

*d*₆/10% water (Figure 5). Similar sequential binding of anions to simple urea-amide receptors containing multiple hydrogen bond donor groups has been observed previously by Gunnlaugsson and co-workers.¹⁸ Analogous titration experiments with compounds **1** and **2** in DMSO-*d*₆/0.5% water resulted in precipitation.

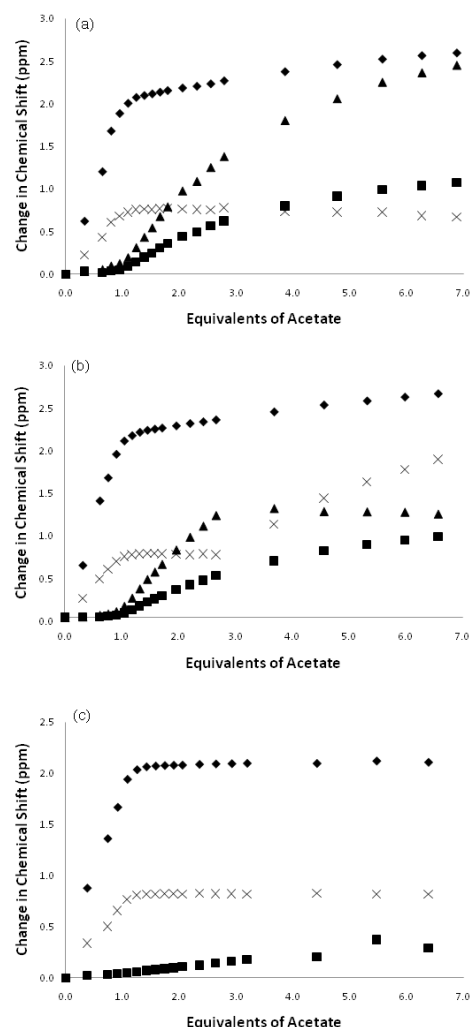
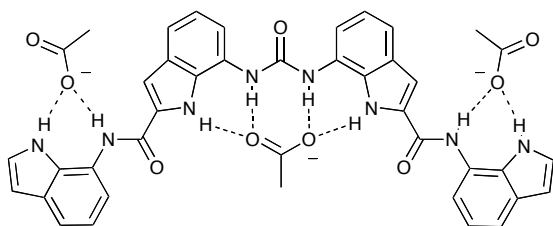
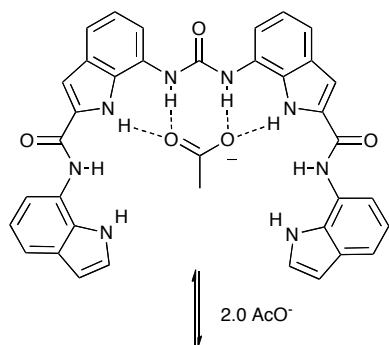
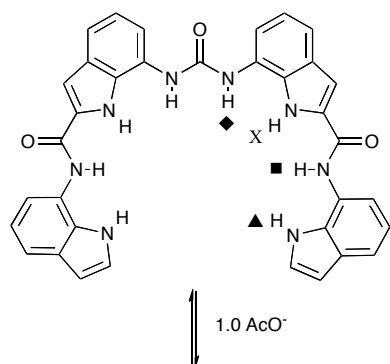


Figure 4 Proton NMR titration data for compounds **1** (a), **2** (b) and **3** (c) upon addition of tetrabutylammonium acetate in DMSO-*d*₆/0.5% water solution. (Key: ◆ = urea NH; ■ = amide NH; × = indole NH (adjacent to urea); ▲ = pendant indole or carbazole NH).



Scheme 1 Proposed solution equilibria of compound **1** with acetate in DMSO- d_6 /water solution.

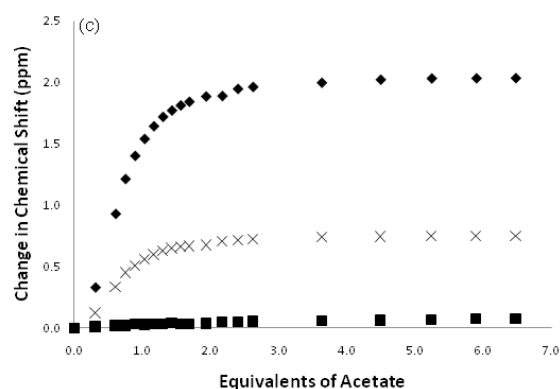
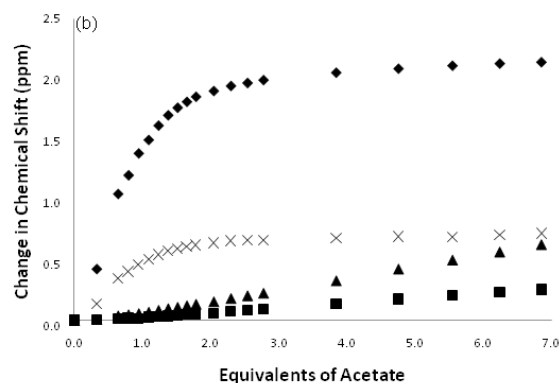
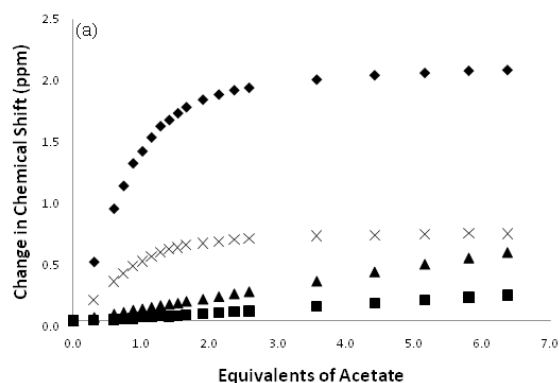


Figure 5 Proton NMR titration data for compounds **1** (a), **2** (b) and **3** (c) upon addition of tetrabutylammonium acetate in DMSO- d_6 /10% water solution. (Key: \blacklozenge = urea NH; \blacksquare = amide NH; \times = indole NH (adjacent to urea); \blacktriangle = pendant indole or carbazole NH).

- In contrast to the results found with acetate, addition of tetrabutylammonium sulfate causes all the NH groups in compounds **1**, **2** and **3** to shift downfield upon addition substoichiometric quantities of the anion. The NMR titration data conducted in DMSO- d_6 /0.5% water is shown in Figure 6.
- Compound **3** binds sulfate strongly in 1:1 stoichiometry with a stability constant $> 10^4 \text{ M}^{-1}$ under these conditions. Upon addition of sulfate to compounds **1** and **2**, more complex equilibria are observed. The NMR titration curves show a discontinuity at one equivalent of sulfate which may be indicative of initial formation of a 1:1 complex followed by a conformational rearrangement and formation of higher order complexes. However, just as was observed with acetate,

moving to DMSO-*d*₆/10% water reduces the complexity of the equilibria in solution with predominant 1:1 complex formation and an apparent stability constant $> 10^4 \text{ M}^{-1}$ for sulfate with compound **2** under these competitive conditions (Figure 7).

Interestingly both compounds **1** and **3** crystallise over the course of about 20 minutes upon addition of tetrabutylammonium sulfate to DMSO-*d*₆/10% water solutions of the receptors. This unfortunately precluded conducting NMR titrations under these conditions. The same process occurs more slowly with compound **2**. The DMSO-*d*₆/10% water solution of compound **1** after addition of 15 equivalents of tetrabutylammonium sulfate is shown in Figure 8.

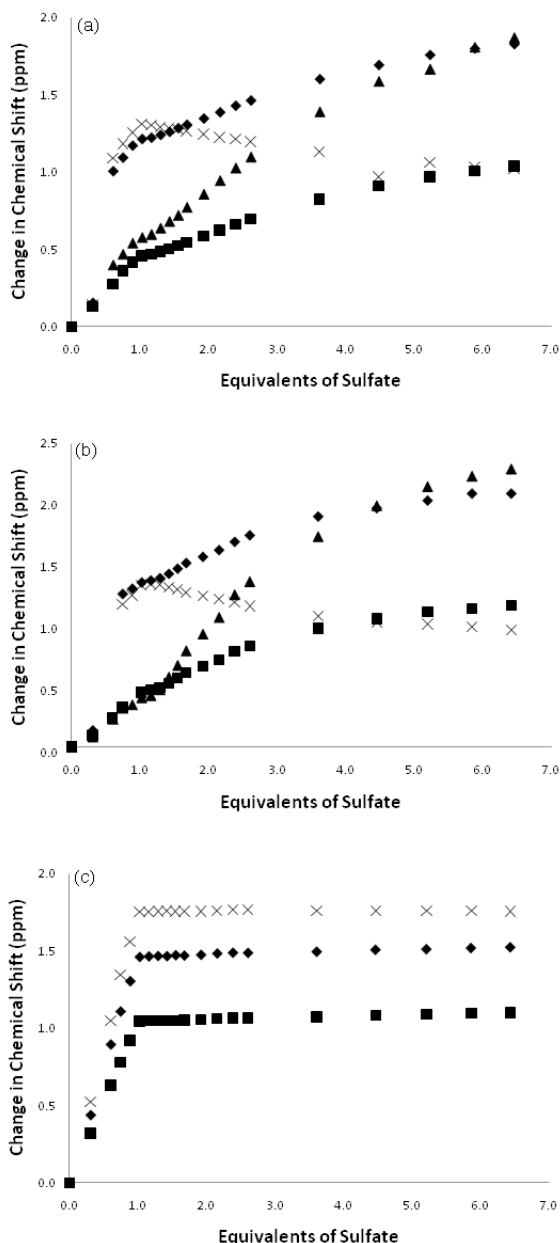


Figure 6 Proton NMR titration data for compounds **1** (a), **2** (b) and **3** (c) upon addition of tetrabutylammonium sulfate in DMSO-*d*₆/0.5% water solution. (Key: ◆ = urea NH; ■ = amide NH; × = indole NH (adjacent to urea); ▲ = pendant indole or carbazole NH).

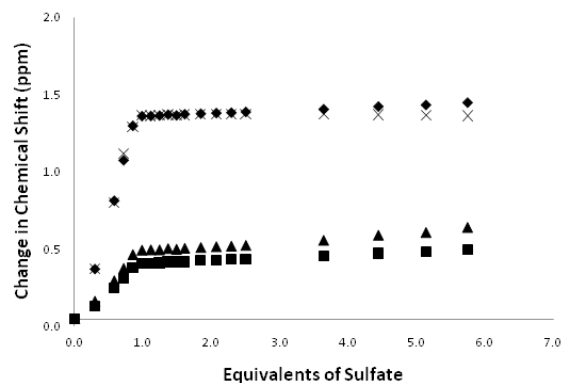


Figure 7 Proton NMR titration data for compound **2** upon addition of 20 tetrabutylammonium sulfate in DMSO-*d*₆/10% water solution. (Key: ◆ = urea NH; ■ = amide NH; × = indole NH; ▲ = pendant carbazole NH).

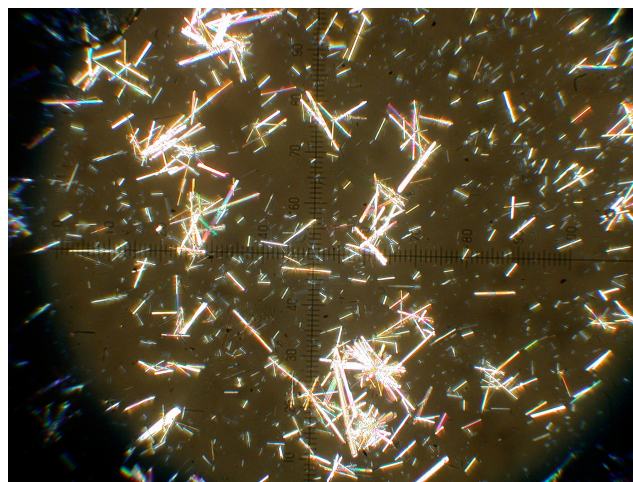


Figure 8 A view of the DMSO-*d*₆/10% water solution of compound **1** in the presence of 15 equivalents of tetrabutylammonium sulfate. Crystals form approximately 20 minutes after the addition of sulfate.

By following the change in chemical shift of the urea NH groups in DMSO-*d*₆/10% it is possible to calculate apparent stability constants 600 M^{-1} and 690 M^{-1} for the complexation of the first equivalent of acetate by compounds **1** and **2**. Stability constants of 120 M^{-1} and 315 M^{-1} for compounds **1** and **2** with tetrabutylammonium dihydrogen phosphate in DMSO-*d*₆/10% water were also measured. Addition of tetrabutylammonium chloride or hydrogen sulfate to both compounds in DMSO-*d*₆/0.5% water resulted in only very small perturbations of the NMR spectra. Compound **3** binds dihydrogen phosphate with a stability constant of 107 M^{-1} in 0.5% water and acetate with a stability constant of 8460 M^{-1} .¹⁶ In all cases these stability constants were calculated following the shift of the urea NH groups using the EQNMR computer program.¹⁹

In DMSO-*d*₆/0.5% water, behaviour indicative of deprotonation of the bound anionic guest was observed with compounds **1** and **2** upon addition of dihydrogen phosphate and bicarbonate (with the latter anion added as the tetraethylammonium salt), i.e. new peaks appear in the NMR spectrum shifted significantly downfield from the free ligand due to formation of monohydrogen phosphate or carbonate

complexes. This results from multiple hydrogen bonding interactions to the bound guest from the receptor reducing the pKa of the anion and subsequent deprotonation by free anion in solution.¹³ Deprotonation was confirmed by addition of one equivalent of the anionic guest followed by hydroxide which results in the appearance of the same new NMR resonances. This is shown in Figure 9 for compound **1** and tetrabutylammonium dihydrogen phosphate. In 10% water this was observed only upon addition of HCO₃⁻ but not H₂PO₄⁻ (see ESI for more details).

Table 1 Stability constants (K_a/M^{-1}) of compounds **1**, **2** and **3** with a variety of putative anionic guests^a determined by ¹H NMR titration techniques in DMSO-*d*₆/10% water mixtures at 298K following the urea NH group resonance. Errors < 15%.

Anion	Compounds 1	2	3 ^g
Cl ⁻	<i>b</i>	<i>b</i>	n.d.
HSO ₄ ⁻	<i>b</i>	<i>b</i>	n.d.
AcO ⁻	600 ^c	690 ^c	1422
BzO ⁻	n.d.	n.d.	481
H ₂ PO ₄ ⁻	120	315	<i>d</i>
HCO ₃ ⁻	<i>e</i>	<i>e</i>	<i>f</i>

^a Anions added as tetrabutylammonium salts except bicarbonate which was added as the tetraethylammonium salt. ^b Only small perturbations were observed in the ¹H NMR spectrum upon addition of these anions. ^c K_I values reported only. ^d Isotherm could not be fitted to a 1:1 or 1:2 binding model. ^e Deprotonation of bound guest was observed. ^f Peak broadening prevented a stability constant from being determined in this case (possibly due to deprotonation of the bound guest). ^g Data from reference 16.

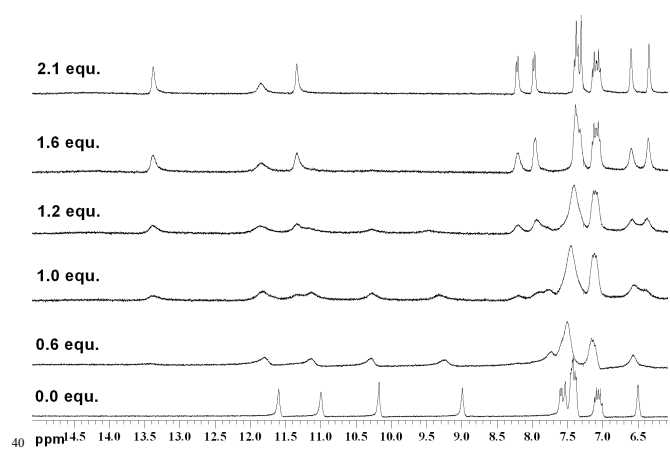


Figure 9. Proton NMR titration for compound **1** and tetrabutylammonium dihydrogen phosphate in DMSO-*d*₆/0.5% water.

Conclusions

Compounds **1** – **3** all bind sulfate strongly in DMSO-*d*₆/0.5% water. Under these conditions compound **3** forms a 1:1 complex whilst the NMR titration evidence leads us to suggest

that compounds **1** and **2** also form 1:1 complexes but at higher sulfate concentrations there are multiple equilibria present. These additional processes do not occur in DMSO-*d*₆/10% water to the same extent with predominant 1:1 complex formation. Hence by changing the solvent conditions from 0.5% water to 10% water we have reduced the complexity of the equilibria present. Compound **2** binds sulfate with a stability constant > 10⁴ M⁻¹ under these conditions whilst compounds **1** and **3** crystallise upon addition of sulfate. It's interesting to note that compound **3** binds sulfate strongly despite having two fewer hydrogen bond donors than compounds **1** and **2**. The receptors are also capable of perturbing the pKa of bound dihydrogen phosphate or bicarbonate to the extent that they are deprotonated by free anions in solution. The crystal structures show that compound **1** is capable of encapsulating tetrahedral oxo-anions *via* eight hydrogen bonds. We are currently exploring the extraction properties of this new family of sulfate selective anionophores. The results of these studies will be reported in due course.

Acknowledgements

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Notes and references

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- † Electronic Supplementary Information (ESI) available: [experimental details, characterisation data for compounds **1** and **2**, NMR and titration data]. See DOI: 10.1039/b000000x/
- ‡ Crystal data for compound (1.(TBA)₂(HPO₄)) C₆₇H₉₉N₁₀O₇P, Mr = 1187.53, T = 120(2) K, Monoclinic space group P2₁/n, *a* = 14.7860(6), *b* = 17.7836(6), *c* = 25.6491(10) Å, β = 103.512(2)° *V* = 6557.7(4) Å³, ρ_{calc} = 1.203 Mg / m³, μ = 0.102 mm⁻¹, Z = 4, reflections collected: 63501, independent reflections: 11548 (*R*_{int} = 0.1856), final *R* indices [*I* > 2σ(*I*): *R*1 = 0.1595, *wR*2 = 0.2338, *R* indices (all data): *R*1 = 0.2672, *wR*2 = 0.2775.
- § Crystal data for compound (1.(TBA)₃(C₆H₅CO₂)).H₂O) C₁₀₄H₁₅₁N₁₁O₁₀, Mr = 1715.36, T = 120(2) K, Triclinic space group P1, *a* = 8.9320(1), *b* = 12.4500(2), *c* = 22.5830(4) Å, α = 93.6790(9)°, β = 100.5490(12)°, γ = 98.6400(11)°, *V* = 2429.96(6) Å³, ρ_{calc} = 1.172 Mg / m³, μ = 0.075mm⁻¹, Z = 1, reflections collected: 38462, independent reflections: 8540 (*R*_{int} = 0.0637), final *R* indices [*I* > 2σ(*I*): *R*1 = 0.0613, *wR*2 = 0.1183, *R* indices (all data): *R*1 = 0.0811, *wR*2 = 0.1288.
- ¶ Crystal data for compound (1.(TBA)₂(SO₄)) C₆₇H₉₈N₁₀O₇S, Mr = 1187.61, T = 120(2) K, Triclinic space group P-1, *a* = 8.78020(10), *b* = 16.9156(4) *c* = 22.1638(5) Å, α = 88.1703(11)°, β = 85.7351(12)°, γ = 82.7653(13)°, *V* = 3255.72(11) Å³, ρ_{calc} = 1.211 Mg / m³, μ = 0.110 mm⁻¹, Z = 2, reflections collected: 61520, independent reflections: 11472 (*R*_{int} = 0.0907), final *R* indices [*I* > 2σ(*I*): *R*1 = 0.0734, *wR*2 = 0.1616, *R* indices (all data): *R*1 = 0.1081, *wR*2 = 0.1822.
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