

Synthesis and Anion Binding Properties of *N,N'*-Bispyrrol-2-yl-2,5-diamidopyrrole

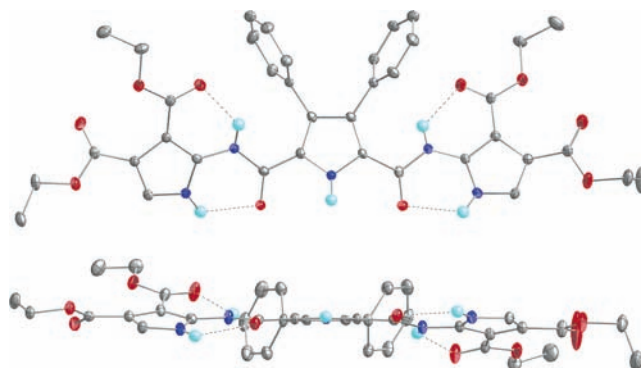
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ABSTRACT



A bispyrrol-2-yl-2,5-diamidopyrrole has been synthesized and shown to have a significantly higher affinity for oxo-anions than previous generation 2,5-diamidopyrroles.

The ubiquity of anions in nature makes the study of their interaction with natural and artificial receptors a topic of considerable current interest.^{1,2} The synthesis of artificial anion receptor systems is driven also by their potential utility in a wide range of applications as different as separations and waste remediation to biomedical analysis and therapy.³ Within the general context of supramolecular chemistry, the field of synthetic anion receptor chemistry is one of the fastest growing disciplines.⁴ To date, a wide variety of heterocyclic anion receptors have been reported in the literature.

These include systems that run the gamut from simple linear monomeric pyrroles to open chain polypyrroles, as well as cyclic pyrrolic structures.⁵ Acyclic 2,5-diamidopyrroles, such as **2**, reported by Gale and co-workers,⁶ are among the most versatile and easy to synthesize of the known pyrrole-based anion receptors. These systems have been found to interact particularly strongly with benzoate and dihydrogen phosphate anions in DMSO or DMSO/water solution.

We were thus curious to explore whether the incorporation of additional pyrrole NH hydrogen bond donor elements

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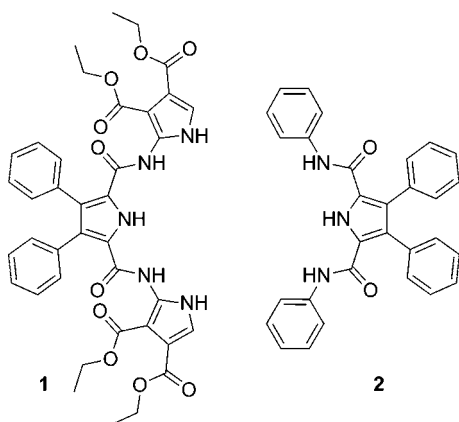


Figure 1. Receptor **1** and the “parent” compound receptor **2**.

would lead to enhancements in affinity or modifications in the inherent anion binding selectivity. Toward this end, we have prepared receptor **1**, a system whose synthesis is predicated on the availability of a functionalized 2-aminopyrrole precursor.

2-Aminopyrroles are not readily available precursors. In general, such species are hard to make and are notoriously prone to decomposition. An exception is systems bearing electron-withdrawing groups in the β -pyrrolic positions, such as diethyl 2-aminopyrrole-3,4-dicarboxylate, **3**.⁷ This latter pyrrole was prepared readily using the procedure first described by Duffy and Wibberley.⁷

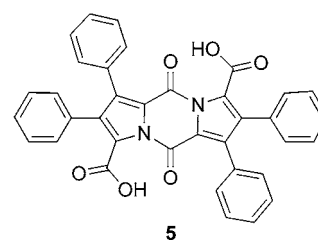
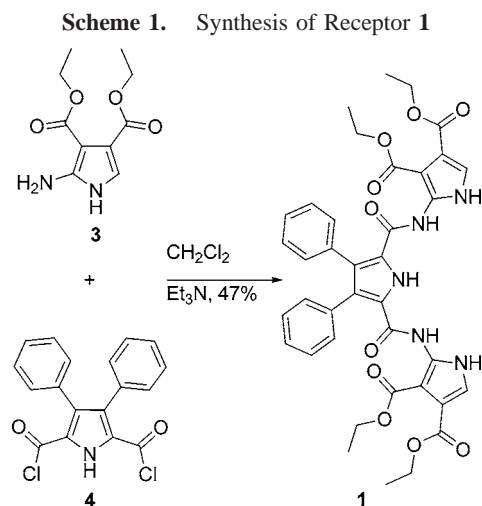
Once this key precursor was in hand, it was reacted in dry CH_2Cl_2 with 0.5 equiv of 3,4-diphenylpyrrole-2,5-dicarbonyl dichloride, **4**,⁸ in the presence of Et_3N . Following purification via column chromatography (silica gel, 3% MeOH in CH_2Cl_2 eluent), compound **1** was isolated in 47% yield (Scheme 1). The moderate yields for this reaction are ascribed to the inherent instability of the diacid chloride, **4**, and to the formation of the three-ring dimer species **5**.⁹ The formation of such dimeric species has been noted previously in the context of forming **2** and its analogues.⁶

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Structural proof for receptor **1** came from a single-crystal X-ray diffraction analysis of crystals of **1** grown by slow evaporation of a CHCl_3 solution of the receptor (see Supporting Information). Figures 2 and 3 show the top and side views of **1** and intermolecular hydrogen bonding interactions between adjacent species in the solid state, respectively.

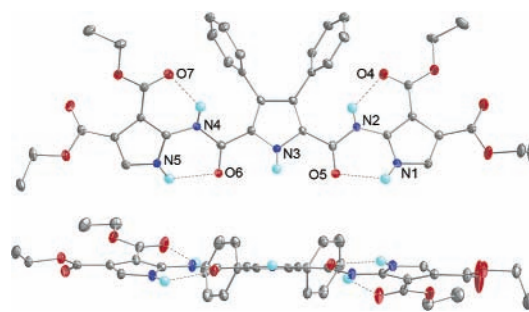


Figure 2. Top and side view of **1**. Dashed lines indicate hydrogen bonding interactions. The thermal ellipsoids are scaled to the 50% probability level. Most hydrogen atoms have been removed for clarity.

There are a number of different hydrogen bonding interactions observed in the solid-state structure of **1**. The first, depicted in Figure 2, is of an intramolecular nature and is formed between the pyrrole NH proton and the carbonyl oxygen of the amide moiety ($\text{N1H}\cdots\text{O5}$, 2.73 Å, 118°;

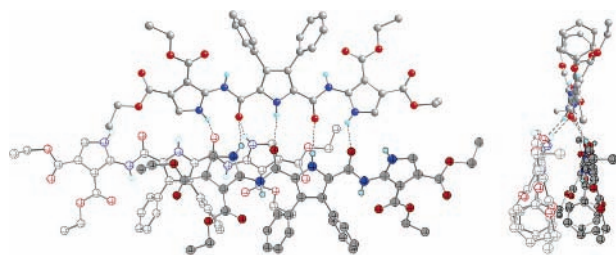


Figure 3. View of the crystal structure of **1** showing interactions between adjacent receptors in the solid state.

$N5H\cdots O6$, 2.72 Å, 116°). A second intramolecular interaction involves the amide NH proton and the carbonyl oxygen of the ester functionality on the β -position of the terminal pyrrole ($N2H\cdots O4$, 2.75 Å, 126°; $N4H\cdots O7$, 2.74 Å, 124°). In addition to these two intramolecular interactions, there are six intermolecular hydrogen bond interactions. The intermolecular interactions serve to assemble compound **1** into what can be described as an extended dual sheet structure. This unusual structure reflects the fact that half of a molecule is hydrogen bonded to one complementary molecule, while the other half is hydrogen bound to a second independent but also complementary molecule. The actual hydrogen bonding interactions involve the pyrrole NH protons and the corresponding amide carbonyl oxygen atoms from one of the molecular partners in the dual sheet. Four of the hydrogen bond interactions are considered to be fairly strong, as evidenced by the bond lengths between 2.81 and 2.92 Å, while the other two, with identical lengths of 3.22 Å, are judged to be weaker.

The fact that extensive intermolecular interactions were observed for **1** in the solid state led to concerns that aggregation might be observed in solution. Such aggregation behavior, to the extent it was observed, would complicate the anion binding behavior of this putative receptor. Accordingly, its interactions with representative anions (in the form of their tetrabutylammonium salts) were studied in DMSO- d_6 solution using 1H NMR spectroscopic techniques. This solvent, which was also the one used previously for the study of **2**, was expected to stabilize the monomeric form of **1**. Consistent with this supposition, the binding constants obtained from fits of the 1H NMR spectroscopic titration curves (cf. Figure 4) were found to be concentration independent over the concentration range of 2.2–4.9 mM (0.95–9.97 mM in the case of benzoate). Standard curve protocols, as used previously by our groups¹⁰ (and others), were used to calculate the binding constants, with the host–guest stoichiometries being determined using the Job plot method. The results of these analyses are summarized in Table 1.

Inspection of Table 1 reveals that in the series of anions studied only dihydrogenphosphate and benzoate anions are

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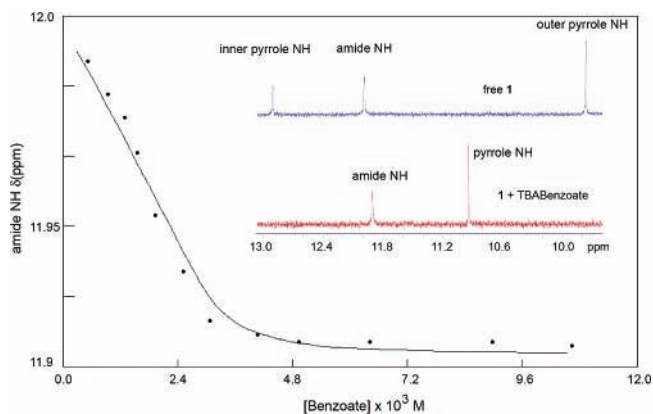


Figure 4. Curve-fitting trace of the data obtained by following the amide NH proton shift in the 1H NMR spectral titration of **1** with tetrabutylammonium benzoate. The inset shows part of the 1H NMR spectrum of **1** recorded in the absence (blue trace) and in the presence (red trace) of benzoate anion (6.14 equiv).

bound appreciably by receptor **1** in DMSO. In accord with the design expectations, receptor **1** displays a much higher affinity for these two anions than does **2**.⁶ It also appears to be even more selective for these oxyanions than these previously reported 2,5-diamidopyrrole anion receptor systems. In fact, in contrast to **2**, receptor **1** displays no detectable affinity for chloride anion and, in analogy to these analogues, likewise shows no appreciable interaction with either bromide or hydrogen sulfate anions. On the other hand, receptor **1** displays a dihydrogenphosphate/benzoate selectivity that is actually reversed compared to that seen in the case of receptor **2**.

Table 1. Affinity Constants^a for the Binding of Anions by Receptors **1** and **2** as Determined from 1H NMR Spectroscopic Titrations in DMSO- d_6 (the anions studied were in the form of their tetrabutylammonium salts)

anion	1	2 ^b
Cl ⁻	NB ^c	11
Br ⁻	NB ^c	<10
HSO ₄ ⁻	NB ^c	<i>d</i>
H ₂ PO ₄ ⁻	5500 ^e	1450
C ₆ H ₅ COO ⁻	10300 ^e	560

^a All studies were carried out at an initial receptor concentration of 3.0 mM. The tabulated values are the average of at least two independent determinations; error $\leq 15\%$. ^b From ref 6. ^c NB = no binding, as inferred from the lack of observable change in the 1H NMR spectrum upon the addition of the indicated anion. ^d Not determined. ^e 1:1 stoichiometry as confirmed by Job plots.

These results can be rationalized by the presence of the two new pyrrole rings in **1** and hence a beneficial increase in the number of hydrogen bond donor sites as compared to compound **2**. The selectivity of **1** toward oxo-anions over halide anions may be explained by the presence of intramolecular hydrogen bond interactions between the amide NH

protons and one of the β -pyrrolic ester moieties (Figure 2; N4H \cdots O7 and N2H \cdots O4). These interactions, observed in the solid state, are maintained in solution as judged by the significant difference seen in the chemical shift of the amide NH proton seen in the case of **1** (11.99 ppm) and **2** (9.37 ppm).⁶ Support for this conclusion also comes from the fact that the amide NH proton of **1** undergoes an upfield shift upon addition of an oxo-anion. Interactions with such anions serve to break up these intramolecular NH \cdots O H-bonds, replacing them with a number of NH \cdots A (A = anion) contacts. Each of these individual NH \cdots A hydrogen bonds is likely weaker than the original internal hydrogen bonds. As a result, a net upfield shift in the NH signal is observed. For the same reason, strong binding is seen only the case of substrates, such as oxo-anions (e.g., benzoate, dihydrogen phosphate), but not halides, that are of an appropriate size and shape to stabilize a large number of NH \cdots A hydrogen bonding interactions.

In summary, we have shown that the addition of supplementary NH hydrogen bond donors on to an easy-to-make pyrrole–amide anion recognition platform, such as that defined by the parent system **2**, can lead to demonstrable increases in anion affinity. The fact that in **1** these additional

NH donor moieties are constrained within a quasi-planar receptor framework via intramolecular hydrogen bond interactions in the absence of an added anion guest favors interaction with oxo-anions, such as benzoate and dihydrogen phosphate, and imparts selectivity relative to halide anions.

Current work is focused on probing further the utility of this approach, as well as studying the intermolecular interactions of systems such as **1** and **2** in less competitive solvent environments.

Acknowledgment. This work was supported in part by the NIH (Grant No. GM 58907 to J.L.S.). P.A.G. thanks the Royal Society for a University Research Fellowship and Professor Mike Hursthouse and the EPSRC for access to the crystallographic facilities at the University of Southampton.

Supporting Information Available: General methods and materials, synthetic experimental, representative Job plots, ¹H NMR titrations, corresponding binding isotherms, details of affinity constant determinations, and X-ray crystallographic information including CIF files. This material is available free of charge via the Internet at <http://pubs.acs.org>.

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