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## COMMUNICATION

## Cooperativity of H-bonding and anion $-\pi$ interaction in the binding of anions with neutral $\pi$ -acceptors<sup>†</sup>

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A rare anion– $\pi$  complex between bromide and a neutral receptor is reported and related receptor systems are studied with a series of anions. The interaction is observed in the solid state and in solution, and further evidence for it is obtained by a computational study.

Due to their crucial role in biological and chemical processes anions have become the focus of intense recognition and sensing studies.<sup>1</sup> Common anion receptors are based on electrostatic attraction, hydrogen-bridges, hydrophobic effects or ion pair recognition and are more or less specific.<sup>2</sup> In recent years anion– $\pi$ interaction was identified as an interesting new binding motif.<sup>3</sup> Many theoretical<sup>4</sup> and structural studies<sup>3</sup> have proven the existence of this weak interaction. However, the relevance and strength of anion– $\pi$  interactions in solution still remain elusive.<sup>5</sup>

In 2008 we started systematic studies on anion– $\pi$  interactions in pentafluorophenylammonium and -phosphonium salts.<sup>6</sup> Initial results revealed a certain flexibility in the position of the anion above the electron-deficient arene, which can be controlled by using directing substituents.<sup>7</sup> The spatial structure of the anion seems not to have an effect on the anion– $\pi$  interaction.<sup>8</sup> However, anions which are of low stability or even unstable in the gas phase like the tetraiodide dianion can be stabilized in the crystal lattice by the support of electron-deficient arenes.<sup>9</sup> Just recently our results showed for the first time that the anion position depends on the fluorination degree of the arene in the solid state.<sup>8</sup>

In the present work anion– $\pi$  interactions are studied by co-crystallizing a series of pentafluorophenyl derivatives with tetrabutylammonium halides (TBAX, X = Cl<sup>-</sup>, Br<sup>-</sup>, I<sup>-</sup>, PF<sub>6</sub><sup>-</sup>, BF<sub>4</sub><sup>-</sup>). Pentafluorobenzamide with tetraethylammonium bromide (1) crystallizes in the monoclinic space group  $P2_1/n$  (Fig. 1). In the solid state the bromide is fixed by N–HBr<sup>-</sup> hydrogen bonds [N···Br<sup>-</sup> = 3.431(3) Å] close to the  $\pi$ -system of the uncharged receptor. On the other hand, this directing



**Fig. 1** Part of crystal structure of **1** showing the pentafluorobenzamide and the bromide (the TBA cation was removed for clarity).

hydrogen bond withdraws the Br<sup>-</sup> from the center of the electron-deficient arene (Fig. 1). Furthermore, there is an additional intermolecular N–H···Br<sup>-</sup> hydrogen bond from the adjacent molecule [N···Br<sup>-</sup> = 3.392(3) Å]. The contacts between the carbon atoms of the pentafluorophenyl group and the bromide are in between 3.650(3) and 4.164(3) Å and the distance to the centroid of the C<sub>6</sub>F<sub>5</sub> substituent is 3.67 Å.

Due to the low solubility of pentafluorobenzamide in chloroform the *N*-phenylbenzamides  $2\mathbf{a}-\mathbf{c}$  were synthesized (see ESI<sup>†</sup>) by reacting amine  $3\mathbf{a}$  or  $3\mathbf{b}$  with pentafluorobenzoyl chloride ( $4\mathbf{a}$ ) or 3,5-dichlorobenzoyl chloride ( $4\mathbf{b}$ ) in the presence of pyridine.

In order to determine the binding constants for **2a** or **2b** in chloroform the stoichiometry was checked using the Job plot (Fig. 2a).<sup>10</sup> By following the amide proton signal in the <sup>1</sup>H NMR spectra a 1 : 1 complex for **2a** as well as for **2b** was observed for various anions (Cl<sup>-</sup>, Br<sup>-</sup>, I<sup>-</sup>, BF<sub>4</sub><sup>-</sup>). Corresponding results can be obtained by following other signals in the <sup>1</sup>H or <sup>19</sup>F NMR spectra. The titration with TBABF<sub>4</sub> shows a different (2 : 1) ratio of receptor to anion (Fig. 2b and c).

The NMR titrations were performed in  $CDCl_3$  using 0.01 M solutions of the receptors **2a** and **2b**. The TBA salts were added as 0.04 M solutions in chloroform. The binding constants were calculated by analyzing the titration curves *via* non-linear regression and are summarized in Table 1 (for full details see ESI†).<sup>10</sup>

A diagram for **2a** showing the binding constants for various anions reveals a decreasing binding constant in the order  $Cl^- > Br^- > I^-$  (Fig. 3a). The binding constants for  $BF_4^-$ (135–153 M<sup>-1</sup>) are somewhat higher than for iodide while  $PF_6^-$  has very similar values. It should be noticed that the complex stoichiometry of the  $PF_6^-$  complex is not clear and that this value is estimated for a 1 : 1 complex. A deviation from the general trend is observed for the binding constants calculated by following the *ortho*-fluorine signals. This might

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**Fig. 2** (a) Representative Job plots for **2a** with various anions (Cl<sup>-</sup>, Br<sup>-</sup>, I<sup>-</sup>, PF<sub>6</sub><sup>-</sup>) added as *n*-Bu<sub>4</sub>N salts in CDCl<sub>3</sub>. (b) Selected <sup>1</sup>H-NMR spectra for the addition of *n*-Bu<sub>4</sub>NCl to a solution of **2a** in CDCl<sub>3</sub>. (c) Resulting titration curves obtained by following the shifting of the NH-signal for **2a** during the addition of TBA salts.

**Table 1** Binding constants  $K_a$  [M<sup>-1</sup>] for the 1 : 1 complexes of **2a** and **2b** with various anions TBAX,  $X = Cl^-$ ,  $Br^-$ ,  $I^-$ ,  $PF_6^-$ ). The binding constants were determined by following the amide proton in the <sup>1</sup>H NMR in CDCl<sub>3</sub> (errors are estimated to be lower than 20%)

	Cl <sup>-</sup>	$\mathrm{Br}^{-}$	$I^-$	$\mathrm{BF_4}^{-a}$	$PF_{6}^{-}$
2a	237	173	99	135	89
2b	163	114	82	102	74
<sup>a</sup> Estim	ated for a 1	: 1 complex	stoichiomet	ry.	

be due to intramolecular interactions of the *ortho*-fluorine atoms with the amide protons. For **2b** an analogous trend was observed. The binding constants are slightly higher for receptor **2a** (Table 1, Fig. 3b). There are two possible explanations for this behaviour: the cooperativity of NH–anion and anion– $\pi$  interaction or the acidification of the NH-group by the electron-withdrawing C<sub>6</sub>F<sub>5</sub>—compared to the C<sub>6</sub>H<sub>5</sub> unit. Proton NMR is able to confirm the H-bonding, but we cannot definitely confirm anion– $\pi$  interactions in solution.

Attempts to perform similar studies in solution for 2c failed due to the very low solubility of 2c in chloroform. Interestingly, 2c becomes soluble in chloroform by adding 1.0 eq. of TBAC1. Solid state structure of 2c revealed the stacking of the molecules by N-H···O=C hydrogen bonds of the amide groups (see ESI†). The strong double N-H···O=C H-bonds form robust columns resulting in the low solubility of 2c. By adding the tetrabutylammonium salt to the solution the



**Fig. 3** (a) Binding constants for **2a** obtained by following the signal in the NMR for various anions (Cl<sup>-</sup>, Br<sup>-</sup>, I<sup>-</sup>, BF<sub>4</sub><sup>-</sup>, PF<sub>6</sub><sup>-</sup>). (b) Comparison of the binding constants of **2a** and **2b** for various anions (Cl<sup>-</sup>, Br<sup>-</sup>, I<sup>-</sup>, BF<sub>4</sub><sup>-</sup>, PF<sub>6</sub><sup>-</sup>).

hydrogen bonds are broken and the phenylbis(pentafluorobenzamide) complex becomes soluble.

Job plots confirm a 1 : 1 ratio for 2c with chloride, bromide and iodide in deuterated acetonitrile. For hexafluorophosphate no significant shifting of the signals could be observed in the NMR spectra. The binding constants obtained for 2c are summarized in Table 2.

The binding constants for 2c and various anions confirm the expected binding order of  $Cl^- > Br^- > I^-$  (Table 2). In this case the ortho-fluorine signals lead to slightly higher values. The binding constants for 2c in CD<sub>3</sub>CN are significantly higher than for 2a in CDCl<sub>3</sub>. This mainly arises from the strong anion binding capability of two amide hydrogens, which is further enhanced by the electronwithdrawing effect of the pentafluorophenyl groups. In addition, cation solvation is stronger in CD<sub>3</sub>CN leaving a "free" anion (and not an ion pair) for binding to the receptor. Moreover, binding of the anions is further strengthened by direct anion $-\pi$  interactions. It should be noticed that due to the different solubility of 2a and 2b the binding constants were determined in different solvents. Nevertheless, the higher binding constant for 2c in CD<sub>3</sub>CN compared to the binding constants of 2a and 2b determined in CDCl<sub>3</sub> is even more surprising due to the stronger solvation of the anion in polar acetonitrile. However, there are two different opportunities for 2c to interact with an anion (see Scheme 1, A and B). In A the anion is bound to a cleft by hydrogen bonds via the two amidic protons and anion- $\pi$  interactions to the two C<sub>6</sub>F<sub>5</sub> moieties, while **B** allows the

**Table 2** Binding constants  $K_a$  [M<sup>-1</sup>] for the 1 : 1 complexes of 2c with various anions (Cl<sup>-</sup>, Br<sup>-</sup>, I<sup>-</sup>). The binding constants were determined by following the proton signals in the <sup>1</sup>H NMR in CD<sub>3</sub>CN (errors are estimated to be lower than 20%)

	$Cl^{-}$	$\mathrm{Br}^-$	$I^-$
NH	3345	637	182
Hortho	3198	645	203
H <sub>meta</sub>	3060	606	148
Fortho	4683	800	134
F <sub>meta</sub>	3401	602	156
F <sub>para</sub>	3660	625	186



Scheme 1 Binding motifs for the interaction of 2c with an anion.



Fig. 4 Conformers A01 ( $C_1$ ) and A02 ( $C_{2v}$ ) of 2c with bromide optimized at the HF/6-311++G\*\* level of theory.

anion to interact only with one H-bond and one electron-deficient unit. In order to determine whether A or B is the preferred structural motif for the anion binding, low-temperature NMR studies of 2c Br as well as computational studies were performed. The low-temperature NMR measurements show highly symmetric NMR spectra and in addition with the computational results A seems to be the favoured binding mode for the anion (see ESI<sup>†</sup>). The shielding of the anion in A also could be the reason for enhanced solubility of 2c by addition of TBACl in chloroform solution. However, it should be mentioned that the DFT studies are performed in the gas phase and no solvent effects are considered. Based on the observed binding constants and the low-temperature measurements, the interaction between the anion and the pentafluorophenyl groups can be considered as an attractive force. To further support this assumption, both conformers A and B (Scheme 1) were geometrically optimized with Gaussian  $09^{11}$  at the MP2/6-31G\* level of theory.

Conformer A (A02) adopting  $C_{2v}$  symmetry is 5.90 kcal mol<sup>-1</sup> lower in energy than B with  $C_1$  symmetry. At the HF/6-311+++G\*\* level an even more stable conformer (A01) could be found (Fig. 4). The energy difference between conformer A01 and A02 is 3.85 kcal mol<sup>-1</sup>. Normal mode analyses resulted in two imaginary frequencies for conformer A02 and none for conformer A01. Therefore, 2c would most likely adopt the structure of conformer A01.

In conclusion we were able to show that pentafluorobenzamides are appropriate systems for studying anion– $\pi$  interactions in the solid state as well as in solution. To the best of our knowledge the crystal structure of  $1 \cdot Br^-$  is the first example of anion– $\pi$ interactions between an uncharged pentafluorophenyl derivative and an anion. Moreover, our investigations in solution show a slight difference between an electron-rich and -poor system, which can be explained by a cooperative effect of N–H···anion and anion– $\pi$  interaction and the enhanced acidity of the amide proton by the electron-withdrawing  $C_6F_5$ -unit. So far we were not able to clarify this point sufficiently. However, the differences in the binding constants between **2a** and **2b** are quite small and might be insignificant. Nevertheless, for receptor **2c** with various anions, an attractive interaction between the anion and the pentafluorophenyl moieties can be expected in solution.

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