



**Highly effective yet simple transmembrane anion transporters based upon ortho-phenylenediamine bis-ureas**

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## COMMUNICATION

# Highly effective yet simple transmembrane anion transporters based upon *ortho*-phenylenediamine bis-ureas

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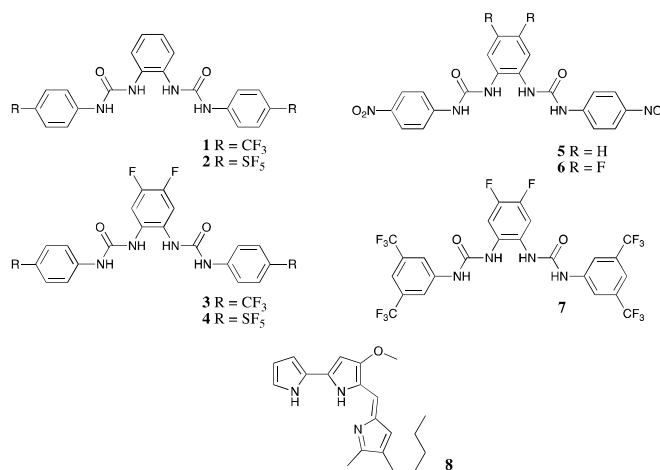
**Simple, highly fluorinated receptors are shown to function as highly effective transmembrane anion antiporters with the most active transporters rivalling the transport efficacy of natural anion transporter prodigiosin for bicarbonate.**

The transport of ions, and in particular chloride and bicarbonate anions, across cell membranes is important to numerous biological processes, and is often mediated by proteins embedded within cellular lipid bilayers.<sup>1</sup> Malfunction of these proteins, through genetic mutation or otherwise, can lead to diseases such as cystic fibrosis.<sup>1</sup> With this in mind, much focus is being placed on the development of small synthetic molecules that can mimic the role of such proteins. With the potential to bind or encapsulate ions, such synthetic “carriers” could be developed for therapeutic applications.<sup>2</sup>

The prodigiosin family of compounds are highly potent natural anionophores, produced by bacteria from the *Streptomyces* and *Serratia* genera.<sup>3</sup> Prodigiosins have been shown to symport H<sup>+</sup>/Cl<sup>-</sup>,<sup>4, 5</sup> and function as Cl<sup>-</sup>/NO<sub>3</sub><sup>-</sup><sup>6</sup> and Cl<sup>-</sup>/HCO<sub>3</sub><sup>-</sup><sup>7</sup> antiporters, demonstrating an ability to act as antibiotic and anticancer agents, as well as exhibiting immunosuppressive behaviour.<sup>8</sup> Prodigiosin itself, **8**, is amongst the most potent anion transporters known.<sup>7</sup>

Synthetic approaches to the design of anion transporters include the attachment of hydrogen bond donor groups to a lipophilic scaffold, an approach to anion transporter design pioneered by Davis and co-workers, with the development of cholic acid derived transporters and more recently systems based upon decalin and cyclohexane.<sup>9</sup> Our approach has focused on developing new hydrogen bonding arrays for transport, with systems containing thiourea, squaramide and thiosquaramide groups showing excellent anion transport properties.<sup>10</sup> Recently we began to study the anion

transport properties of bis-urea receptors based on *ortho*-phenylenediamine, a system that we developed as an anion receptor in 2005.<sup>11</sup> We found that these compounds had excellent anion transport properties but in our initial study the compounds (including compounds **1** and **5**)<sup>12</sup> had not been optimised for transport. Here we report a more complete series of *ortho*-phenylenediamine-based bis ureas functionalised with the NO<sub>2</sub>, F, CF<sub>3</sub> and -SF<sub>5</sub><sup>13</sup> group that are structurally simple, can be synthesised in one step and yet possess exceptional anion transport properties (i.e. low EC<sub>50</sub> values – see below) some of which are more effective chloride/bicarbonate antiport agents than the natural product prodigiosin that is capable of mediating anion transport at very low concentration.<sup>14</sup>

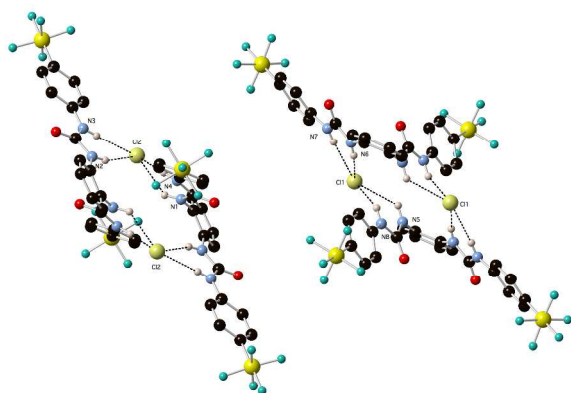


We have previously reported the synthesis and transport properties of receptors **1** and **5**<sup>11</sup>. Reaction of 1,1'-carbonyl diimidazole

(CDI) with 4-aminophenyl sulfur pentafluoride and the appropriately functionalised *ortho*-phenylenediamine in dichloromethane yielded receptors **2** (86% yield) and **4** (23% yield). Reaction of the appropriate isocyanate with the appropriately substituted phenylenediamine in a mixture of dichloromethane and pyridine gave compounds **3**, **6** and **7** in 68%, 59% and 32% yields respectively.

X-ray crystal structures of the chloride complexes of compounds **2** and **6** were elucidated using single crystal X-ray diffraction techniques. Crystals were obtained by slow evaporation from DMSO solutions (a mix with MeOH in the case of receptor **6**) in the presence of an excess quantity of tetrabutylammonium chloride or tetramethylammonium chloride.

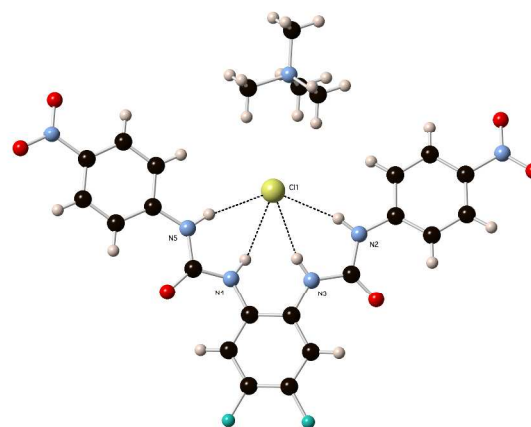
Receptor **2** was found to form a 2:2 receptor:anion complex in the solid state with tetrabutylammonium chloride (N-Cl distances 3.205(4)-3.270(4) Å, N-H...Cl angles 133.2(2)-163.9(2)°). Each chloride anion is bound by four hydrogen bonds, two from each of the participating receptors (Figure 1). Receptor **6** forms a 1:1 receptor:anion complex with chloride (N-Cl distances 3.205(2)-3.315(2) Å, N-H...Cl angles 151.5(1)-167.1(1)°). The chloride anion is bound by each of the four possible hydrogen bonds from the receptor (Figure 2).



**Figure 1:** X-ray crystal structure of **2**.TBACl showing the packing arrangement. Counter ions, DMSO solvent and non-acidic hydrogen atoms have been omitted for clarity.

Proton NMR anion-binding studies (Table 1) were conducted for receptors **1-7** with chloride, bicarbonate and nitrate anions, to determine their affinity and stoichiometry of binding. Stability constants,  $K_{11}$ , were determined by following the shift of the most deshielded N-H proton and fitting the data to a 1:1 receptor:anion binding isotherm using WinEQNMR2.<sup>15</sup>

Stability constants measured for the interaction between the receptors and chloride are quite moderate (30-100 M<sup>-1</sup>). Many of the measured stability constants between the receptors and bicarbonate were very large, particularly that of receptor **3** (>10<sup>6</sup>), and in some cases the data could not be fitted to a binding isotherm due to peak splitting and broadening within the <sup>1</sup>H NMR spectra. This behaviour can be attributed to strong binding interactions and possible secondary equilibrium processes in solution. In general, the receptors did not significantly interact with nitrate anions, and the majority of the stability constants were measured at <10 M<sup>-1</sup>.



**Figure 2:** X-ray crystal structure of **6**.TMACl.

Anion transport studies were carried out for receptors **2**, **3**, **4**, **6**, **7** and prodigiosin **8** using vesicle based methods,<sup>16</sup> using the same procedures employed to study compounds **1** and **5** previously.<sup>2</sup>

To determine chloride/nitrate exchange transport capabilities a sample of unilamellar POPC vesicles were prepared containing 489 mM NaCl buffered to pH 7.2 with 5 mM sodium phosphate salts. The vesicles were suspended at a lipid concentration of 1 mM in 489 mM NaNO<sub>3</sub> buffered to pH 7.2 with 5 mM sodium phosphate salts. The experiment was initiated by the addition of a small amount of a DMSO solution of the receptor (mol% with respect to lipid), and the resulting chloride efflux was monitored using a chloride ion selective electrode (ISE). At the end of the experiment (300 s), the vesicles were lysed by adding a detergent, and the final electrode reading was used to calibrate 100% chloride release. Figure 3 shows a comparison of the transport properties of receptors **1-8** (excluding **1** and **5**) for chloride/bicarbonate antiport at 0.02 mol% loading of transporter.

In order to determine the mechanism of chloride release by the receptors, we performed a second experiment using POPC vesicles containing 451 mM NaCl buffered to pH 7.2 with 20 mM sodium phosphate salts. The vesicles were suspended in 150 mM Na<sub>2</sub>SO<sub>4</sub> buffered to pH 7.2 with 20 mM sodium phosphate salts. On addition of a sample of receptor, there was little significant chloride release. This indicates that these receptors function by an anion exchange mechanism. In the first assay, the receptors were able to facilitate Cl<sup>-</sup>/NO<sub>3</sub><sup>-</sup> exchange, but in this second experiment their activity was diminished due to the high hydrophilicity of the SO<sub>4</sub><sup>2-</sup> anion,<sup>17</sup> which in most cases prevents its transport by synthetic transporters.<sup>18</sup> After 120 s, we added a pulse of NaHCO<sub>3</sub> solution, and found that the receptors were mediating chloride efflux, evidence in support of the hypothesis that these receptors are also able to facilitate a Cl<sup>-</sup>/HCO<sub>3</sub><sup>-</sup> antiport mechanism.

In order to quantify these anion transport processes, we performed a Hill analysis for chloride/nitrate<sup>-</sup> and chloride/bicarbonate antiport by the receptors.<sup>19</sup> These experiments allowed us to calculate the EC<sub>50</sub> for these processes, a measure of activity defined as the effective receptor concentration required to mediate 50% chloride efflux after a specified time period. As is standard in our studies, this period is 270 s for chloride/nitrate experiments and 390 s for

chloride/bicarbonate experiments (corresponding to 270 s after the  $\text{HCO}_3^-$  spike). We were also able to determine the Hill coefficient ( $n$ ), which can be used to give an indication of the number of monomers that form the active transport system.<sup>20</sup> These values, and the previously reported data are shown in Table 1.

A study of retention times of the compounds by a reversed phase HPLC<sup>21</sup> showed that there is not a simple relationship between lipophilicity and the  $\text{EC}_{50}$  values of this series of compounds.

An increase in transport activity is observed when going from bisurea receptor **1** to receptor **2**. These results may be rationalised by considering the effect of increased fluorination,<sup>10,22</sup> in that receptor **2** is expected to have higher lipophilicity than receptors **1**, and hence are expected to partition into the POPC membrane more effectively.

Receptors **3** and **4** are analogous to receptors **1** and **2** respectively, except that the central phenyl rings are fluorinated. Unlike their simpler counterparts, receptors **3** and **4** do not follow the same trend in activity. The  $\text{SF}_5$  containing compound **4** was found to be less active than the  $\text{CF}_3$  containing compound **3** for chloride/nitrate antiport. An efficacious transporter must strike a balance between aqueous solubility and lipophilicity to allow for successful delivery to the POPC vesicles and to allow partitioning into and out of the membrane to transport anions from one side of the membrane to the other. In the case of receptor **4**, one possibility is that the balance of aqueous solubility and lipophilicity has been pushed too far in favour of lipophilicity, and so the receptor resides more within the membrane, hence chloride/nitrate transport is slightly hindered when compared to that of receptor **3**. The transport efficiency of **3** and **4** are roughly equal for chloride/bicarbonate antiport, but perhaps the most surprising thing here is that this seemingly minor structural change results in receptors that can outperform natural product prodigiosin, **8**, at chloride/bicarbonate antiport (Table 1).

To explore this idea further we synthesised receptor **6**. Attaching additional electron-withdrawing fluorine substituents to the central ring boosts the transport properties of receptor **6** relative to receptor **5**. In fact receptor **6** is almost 2.5 times better at promoting chloride/nitrate and chloride/bicarbonate antiport processes than the parent compound. However, unlike with receptors **3** and **4**, receptor **6** does not outperform prodigiosin **8** at chloride/bicarbonate antiport.

The strategy of introducing higher degrees of fluorination onto a receptor scaffold to enhance its transport properties is further illustrated by compound **7** which contains a fluorinated central ring and 3,5-bis(trifluoromethyl) substituents on its terminal phenyl rings. This compound has a lower  $\text{EC}_{50}$  value for chloride/bicarbonate exchange than prodigiosin **8** (Table 1) and is thus a more effective antiporter for this anion pair than the natural product (Figure 3).

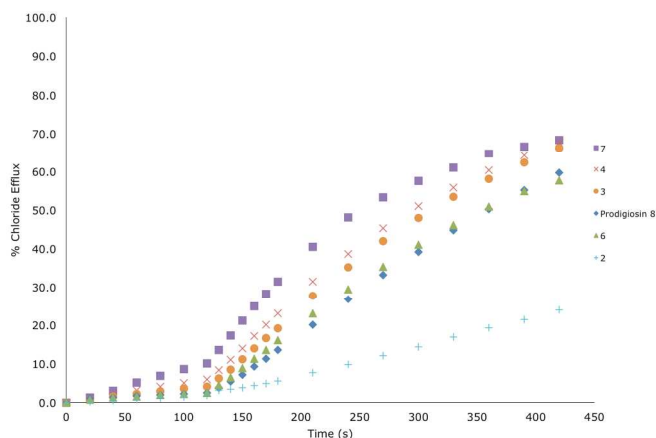


Figure 3: Chloride efflux as a function of time, promoted by the addition of 0.02 mol% (with respect to lipid) of receptors **2**, **3**, **4**, **6**, **7** and prodigiosin **8**, from unilamellar POPC vesicles containing 451 mM NaCl buffered to pH 7.2 with 20 mM sodium phosphate salts. The vesicles were dispersed in 150 mM  $\text{Na}_2\text{SO}_4$  buffered to pH 7.2 with 20 mM sodium phosphate salts. The receptor was loaded as a DMSO solution at 0 s, and a spike of  $\text{NaHCO}_3$  (33 mM) added as 120 s. At the end of the experiment the vesicles were lysed to calibrate the ISE to 100% chloride efflux.

## Conclusions

We have demonstrated that fluorinated receptors based on the bisurea motif are highly effective anion transporters and function by an antiport mechanism of anion transport. Modification of existing transporters, by adding fluorination at the central phenyl ring or by increasing fluorination at the peripheral phenyl groups, yielded highly potent anion transporters capable of surpassing the activity of the natural anion transporter prodigiosin, **8**, for chloride/bicarbonate exchange. This may be due to the affinity of the bisureas for carboxylates<sup>11</sup> including bicarbonate.

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## Notes and references

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† Electronic Supplementary Information (ESI) available: details of receptor synthesis, characterisation, anion transport and anion binding studies as well as additional X-ray diffraction information can be found in the ESI. See DOI: 10.1039/c000000x/

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**Table 1:** EC<sub>50</sub> (mol%) values **1-8** for the release of chloride from POPC vesicles in Cl<sup>-</sup>/NO<sub>3</sub><sup>-</sup> and Cl<sup>-</sup>/HCO<sub>3</sub><sup>-</sup> antiport systems at 270 s and 390 s respectively. EC<sub>50</sub> values are the average of three runs, unless otherwise stated, and Hill coefficient (n) values are also averaged. Stability constant K<sub>i</sub> (M<sup>-1</sup>) for receptors **1-7**, measured in DMSO-d<sub>6</sub>/H<sub>2</sub>O 0.5% at 298K. Guest anions Cl<sup>-</sup> and NO<sub>3</sub><sup>-</sup> were added as tetrabutylammonium salts, while HCO<sub>3</sub><sup>-</sup> was added as the tetraethylammonium salt. Stability constants were determined by <sup>1</sup>H NMR titrations following the most deshielded N-H resonance. All data sets were fitted to a 1:1 receptor:anion binding isotherm using WinEQNMR.<sup>14</sup>

	EC <sub>50</sub> <sup>a</sup> (mol%) Cl <sup>-</sup> /NO <sub>3</sub> <sup>-</sup>	n <sup>b</sup> Cl <sup>-</sup> /NO <sub>3</sub> <sup>-</sup>	EC <sub>50</sub> <sup>c</sup> (mol%) Cl <sup>-</sup> /HCO <sub>3</sub> <sup>-</sup>	n <sup>d</sup> Cl <sup>-</sup> /HCO <sub>3</sub> <sup>-</sup>	K <sub>i</sub> , Cl <sup>-</sup> (M <sup>-1</sup> ) <sup>k</sup>	K <sub>i</sub> , HCO <sub>3</sub> <sup>-</sup> (M <sup>-1</sup> ) <sup>k</sup>	K <sub>i</sub> , NO <sub>3</sub> <sup>-</sup> (M <sup>-1</sup> ) <sup>k</sup>
<b>1</b> <sup>f</sup>	0.018	1.2	0.15	0.9	78	2090	<10
<b>2</b>	0.0099	1.1	0.077	1.0	65	<sup>i</sup>	<10
<b>3</b>	0.0015 <sup>h</sup>	1.1	0.012	1.1	50	>10 <sup>d</sup>	<10
<b>4</b>	0.0030	1.1	0.012	1.0	71	<sup>i</sup>	13
<b>5</b> <sup>f</sup>	0.0048	0.9	0.038	1.1	78	7330	<10
<b>6</b>	0.0019	1.1	0.016	1.2	94	<sup>i</sup>	<10
<b>7</b>	0.0016 <sup>h</sup>	1.1	0.011	1.3	<sup>j</sup>	<sup>i</sup>	<10
<b>8</b>	0.00064	1.3	0.014	1.0	not determined	not determined	not determined

<sup>a</sup> Concentration of receptor (mol% receptor to lipid) required to achieve 50% chloride efflux at 270 s during the Cl<sup>-</sup>/NO<sub>3</sub><sup>-</sup> exchange experiment. <sup>b</sup> Hill coefficient for Cl<sup>-</sup>/NO<sub>3</sub><sup>-</sup> exchange experiment. Standard deviations are shown in brackets where available. <sup>c</sup> Concentration of receptor (mol% receptor to lipid) required to achieve 50% chloride efflux at 390 s during the Cl<sup>-</sup>/HCO<sub>3</sub><sup>-</sup> exchange experiment. <sup>d</sup> Hill coefficient for Cl<sup>-</sup>/HCO<sub>3</sub><sup>-</sup> exchange experiment. Standard deviations are shown in brackets where available. <sup>e</sup> From published results.<sup>10</sup> <sup>f</sup> From previously published results.<sup>11</sup> <sup>g</sup> Accurate Hill analysis was not possible due to low receptor activity. <sup>h</sup> Average of four runs. <sup>i</sup> Data could not be fitted due to resonance broadening upon addition of anions. <sup>j</sup> Data could not be adequately fitted to a binding model due to multiple equilibria in solution (see ESI). <sup>k</sup> See ESI for stability constant errors

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