REGULAR ARTICLE

Bis(pyrrole-benzimidazole) conjugates as novel colorimetric sensor for anions

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Abstract. Novel bis(pyrrole-benzimidazole) (PYBI) conjugates were investigated as colorimetric anion recognition motif by employing multiple donor sites. In this regard, PYBI derivatives where pyrrole unit is connected to the C2 position of benzimidazole *via* its α -position were synthesized and their anion recognition behavior were evaluated by UV-Vis, fluorescence and ^1H NMR spectroscopy. PYBI selectively interacts with fluoride ion, whereas introduction of nitro group on the benzimidazole moiety enhances the binding affinity by at least one order, albeit at the expense of the selectivity. Bridging of two PYBIs led to interesting cooperative effect between the two subunits, which gets enhanced upon changing the spacer between the two from non-conjugating type ($sp^3\text{-C}$) to conjugating one (quinoxaline). This also affects the way they interact with anions, with the latter moiety displaying very interesting stepwise double deprotonation of benzimidazole protons upon addition of fluoride ions with strong colorimetric as well as fluorometric response. Further, acidity of the H-bond donor sites was found to play pivotal role in the anion complexation mechanism and selectivity.

Keywords. Pyrrole; benzimidazole; anion recognition; cooperative; proton migration.

1. Introduction

Considering the important roles of anions in biological, environmental and industrial processes, anion recognition chemistry has developed into an independent research area focused on the structure and binding modes of anions.¹ The goals of these studies have been to understand specific binding behaviour, to derive structure-function relationships, to search for new anion binding motifs and, last but not the least, to find useful application as sensors.² In the last three decades, several neutral receptors containing polarizable H-bond donor units such as pyrrole, imidazole, phenol, amide, urea, *etc.*, and their conjugates such as amidopyrroles, amidoureas, pyrrole-ureas were reported.^{2,3} Anionic guest interacts with these receptors either *via* H-bond formation or *via* deprotonation mechanism.¹ In case of H-bonded system, the binding constant or the formation constant can be directly correlated to the stability of the complex between the receptor and anion. However, deprotonation is a Brønsted acid-base reaction and the equilibrium constant depends on the stability of the corresponding conjugate base of the receptor. With highly basic anions such as fluoride and acetate, receptors with moderately acidic H-bond donor unit initially

forms H-bonded complex, but in presence of excess anion, deprotonation is favoured owing to the formation of stable H-bonded anion dimers.³ The mode of interaction, *i.e.*, H-bonding or deprotonation, generally depends on the acidity of the H-bond donor site of the receptor and basicity of the anion. In principle, increasing the acidity of the H-bond donor moieties increases both H-bond donor affinity as well as the deprotonation tendency. Moreover, receptors where the negative charge delocalization can be extended to the entire molecular framework favours deprotonation.⁴ Deprotonation increases electron density at the receptor sites and eventually decreases the H-bonding tendency of the remaining H-bond donor sites of the receptor.

Pyrrole and benzimidazole are widely used as H-bond donor components in the design of artificial receptors.^{5,6} While pyrrole acts as an H-bond donor, benzimidazole endowed with both N-H and imine N functionalities can respectively act as both H-bond donor and acceptor. The acidic hydrogen of benzimidazole is involved in tautomerism and thereby shuttles between the two N atoms. Therefore, benzimidazole binding sites could be easily oriented on the demand of the complementary binding sites of the guest. Bordwell reported pK_a of benzimidazole as 16.4 in DMSO, which is lower than that of pyrrole ($pK_a = 23.0$), suggesting a rel-

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Figure 1. Structure of the synthesized pyrrole-benzimidazole (PYBI) derivatives.

atively stronger H-bonding as well as deprotonation tendency of the former.⁷ Literature reports suggest that pyrrolic receptors interact with anions preferably *via* H-bonding with some exceptions.⁵ Gale and co-workers reported that diamidopyrrole derivatives with electron withdrawing groups interact with fluoride *via* deprotonation mechanism.⁸ In 1999, Sessler *et al.*,^{9b} reported that 2,3-dipyrrol-2-ylquinoxaline receptor interacts with fluoride *via* H-bonding (N-H...F⁻) with subsequent deprotonation, while other anions interact predominantly *via* H-bonding.⁹ However, systems where benzimidazole units are employed as H-bond donors are comparatively less common and mostly interact with anions *via* the deprotonation mechanism.⁶ Caussey and Allen reported that benzimidazole diamides interact with anions through H-bonding, and benzimidazole tautomerism affects the composition of the H-bonding array provided to the anion.¹⁰ Furthermore, Gale *et al.*, reported that benzimidazole appended ureas bind anions through H-bonding, and the tautomeric H-bonding structures of the anion complexes depend on the H-bonding domain of the anions.¹¹ These observations clearly reflect that the mode of interaction of pyrrole and benzimidazole units with anions basically depends on the basicity of the anion and the nature of the organic scaffold where it is incorporated.

Herein, we report new pyrrole-benzimidazole (PYBI) conjugates where the pyrrole unit is directly linked to the C2 position of the benzimidazole ring *via* its α carbon, along with their detailed anion binding behaviour (Figure 1). These new motifs have two preorganized heterocyclic N-H donor sites and represent larger, more extended and flexible binding domain for anions (owing to proton transfer between benzimidazole Nitrogens). Further, we wish to explore the relative binding affinity of the two N-H donor units, along with the role of their acidity in anion selectivity. To increase the acidity difference between the two NH units and to obtain bet-

ter colorimetric response for the recognition event, nitro group is incorporated at the benzimidazole periphery. It is assumed that insertion of nitro group will lead to facile anion induced charge transfer from the binding domain to the nitro group and yield a sharp color change.¹² In addition to the basic conjugates, we have proposed two different sets of dimers (bis-PYBI) containing two PYBI units, where both the units are bridged *via* two different spacer motifs - first, an electronically non-coupled sp^3 -carbon (analogous to dipyrromethanes) and second, they are coupled by a fluorescent quinoxaline moiety (analogous to quinoxaline bipyrrrole). These dimeric motifs are expected to display cooperative binding effect towards anionic guests owing to varied interaction between the two PYBI units. This can be envisaged in view of their difference in electronic communication and spatial disposition, which may be further aided by proton shuttling ability of the two constituent benzimidazoles.

2. Experimental

2.1 Materials and Methods

4-nitro-*o*-phenylenediamine and tetrabutylammonium salts of anions employed for the study were purchased from Sigma Aldrich and were used as received. Pyrrole, *o*-phenylenediamine, phosphorous oxychloride, oxalyl chloride and nitrobenzene were purchased from Sisco Research Laboratories, India. Pyrrole, phosphorous oxychloride and solvents used in the synthesis were dried and distilled prior to use. Spectroscopic grade DMSO (Merck, India) was used for recording all spectrophotometric data. Melting points were determined with MR-Vis⁺ visual melting point range apparatus from LABINDIA Instruments Pvt. Ltd. IR spectra were recorded on a FT-IR spectrometer (JASCO FTIR model 5300 or NICOLET 5700). NMR spectra were obtained on Bruker 400 MHz FT-NMR spectrometer operating at ambient temperature. TMS was used as internal standard for ¹H NMR spectra. LCMS spectra were recorded using Shimadzu-

LCMS-2010 mass spectrometer. Elemental analyses were obtained using Thermo Finnigan Flash EA 1112 analyzer. UV-Vis spectra were recorded on a Perkin Elmer Lambda 35 UV-Vis spectrophotometer. Fluorescence spectra were recorded using Horiba Jobin Yvon Fluoromax-4 instrument.

3. Results and Discussion

3.1 Synthesis and characterization

Pyrrole-benzimidazole (PYBI) conjugates were conveniently synthesized by condensation of 1,2-phenylenediamine or *m*-nitro-1,2-phenylenediamine with the corresponding pyrrole carboxaldehyde derivative in nitrobenzene solvent at 120°C (see Supplementary Information).¹³ All compounds were characterized by standard spectroscopic techniques including ¹H, ¹³C NMR spectroscopy, mass and elemental analysis (in SI). ¹H NMR spectra recorded in DMSO-*d*₆ at 298 K show two NH signals above 12 ppm while pyrrole α- and β-Hs resonate around 6 and 7 ppm, respectively, for all the compounds (see Supplementary Information: Figures S1, S3, S5, S7 and S9). In the case of **2b**, benzimidazole NH signal splits into a doublet, which may be attributed

to the NH tautomerism of benzimidazole moiety that becomes slower in presence of the nitro group. On the other hand, proton shuttling between the two benzimidazole nitrogens is probably too fast in the case of **1a**, **b** and **2a** to be detected in the NMR time scale; therefore, it appears as a broad signal (see SI, Figures S1, S3 and S5). The absorption and emission spectra of compounds are shown in Figure 2 (Table 1). UV-Vis spectrum of compound **1a** in DMSO revealed two absorption maxima at 312 and 327 nm (not much resolved). In addition, it shows fluorescence with emission maxima at 336 and 349 nm. On the other hand, insertion of nitro derivative (*i.e.*, **1b**) results in well-resolved absorption bands with maxima at 299 and 387 nm indicating charge transfer nature of the transition. However, this molecule did not display any detectable fluorescence owing to the intramolecular photoinduced electron transfer (PET) from the excited fluorophore to the electron withdrawing nitro group (donor-excited PET; d-PET).¹⁴ Absorption spectrum of quinoxaline-bridged dimeric conjugate **3**, in DMSO, exhibits two absorption bands with maxima at 318 and 404 nm and it shows a relatively large red shifted emission at 560 nm. This indicates facile conjugation between the two PYBI conjugates *via* quinoxaline moiety both in the ground and the excited state.

Figure 2. Normalized absorption (solid curves) and emission (dashed curves) spectra of the receptors in DMSO.

Table 1. Absorption and emission maxima along with log ($\epsilon/M^{-1}cm^{-1}$) and Stoke's shift values for PYBI conjugates **1–3** in DMSO at 25°C.

Receptors	Absorption in DMSO λ_{max} (nm)	log ($\epsilon/M^{-1}cm^{-1}$)	Emission in DMSO λ_{max} (nm)	Stoke's Shift (nm)
1a	312, 327	4.56, 4.49	337, 349	10
1b	299, 387	4.35, 4.13	–	–
2a	313, 338	4.68, 4.58	361 (broad)	23
2b	303, 398	4.55, 4.36	–	–
3	318, 404	4.49, 4.27	563	159

3.2 Anion binding study

3.2a Absorption and Emission Spectroscopy: Initial anion binding studies of the receptors were carried out at 298 K, in DMSO, using UV-Vis spectroscopic titration method. The studies were done with 20 μM solution of the receptors. Titrations were performed by adding known quantities of concentrated solution of the anions to the receptor solution. The anion solutions used in the titrations contained the receptor at the same concentration as the receptor solutions so as to nullify the dilution effect. The absorption data were fitted in the Benesi-Hildebrand plot of Connor equation (eq. 1) and from the ratio of the slope and intercept, binding constant K_a was evaluated.¹⁵

$$\frac{1}{[A - A_0]} = \frac{1}{S_t K_a b \Delta \epsilon [L]} + \frac{1}{S_t b \Delta \epsilon} \quad (1)$$

Here, A° , A : Measured absorbance of the substrate in absence and in presence of anion respectively; S_t : Total receptor concentration (M); $[L]$: Anion concentration (M); K_a : Binding constant; $\Delta \epsilon$: Difference in molar absorptivity of the free receptor solution and the receptor-anion solution, and b : Path length.

Anion screening experiment of **1a** showed an appreciable change in the absorption spectrum for fluoride ion only, among the tested anions. Addition of TBAF resulted in ~ 9 nm red shift of the two absorption bands (Figure 3). In order to evaluate the affinity of fluoride ion towards **1a**, systematic spectrophotometric titration was carried out in DMSO with varying concentration of fluoride ion. During initial part of the titration (0–10 mole equiv.) a decrease in intensity of the band at 327 nm was observed. However, on further increase in F^- concentration (>10 mole equiv.), a new band appeared at 336 nm with concomitant decrease in the

absorbance of the peak at 327 nm. These observations suggested the possible involvement of two equilibrium process during the titration with varying concentration of fluoride. It is envisaged that initially for lower effective $[\text{F}^-]$, H-bonding interaction ($\text{N-H} \cdots \text{F}^-$) could be attributed for the initial set of spectral changes. Further increase in $[\text{F}^-]$ might trigger the deprotonation of benzimidazole moiety. Comparatively higher acidity of the benzimidazole N-H and the high thermodynamic stability of the HF_2^- ion have favoured the deprotonation process. Deprotonation increases the electron density in **1a** and consequently reduces the strength of the H-bonding interaction of the pyrrolic N-H with the fluoride ion.¹⁶

Anion screening experiment of receptor **1b** displayed significant changes in UV-Vis spectrum, upon addition of F^- , CH_3COO^- and H_2PO_4^- ions, and consequently colour of the solution changed from very light yellow to dark yellow with varying intensities (Figure 4). UV-Vis spectral titration revealed that with gradual addition of fluoride and acetate ions, the intensities of absorption bands at 299 and 387 nm decreased, while concomitantly, two new peaks appeared at 305 and 439 nm (Figure 5). The titrations revealed two clear isosbestic points at 324 and 404 nm. Whereas, dihydrogenphosphate ion displayed relatively moderate response with slight change in the peak maxima (at 304 and 431 nm). The interaction of **1b** with anions resulted in increase in the electron density at the PYBI moiety and, as a consequence, facile charge transfer towards the acceptor $-\text{NO}_2$ moiety stabilizes the excited state and hence causes a bathochromic shift in the absorption maxima, resulting in the change in color.¹² This clearly reflects that nitro substitution at the benzimidazole periphery led to the colorimetric change upon addition of anions, however,

Figure 3. Left: UV-Vis spectra of **1a** (20 μM) in DMSO in the absence and presence of different anions; right: UV-Vis spectra of **1a** (20 μM) upon gradual addition of TBAF (0 to 15 mM).

Figure 4. Left: UV-Vis spectra of **1b** (20 μ M) in DMSO in absence and presence of different anions; right: UV-Vis spectra of **1b** (20 μ M) in DMSO upon gradual addition of TBAF (0 to 0.35 mM).

Figure 5. Left: Evolution of UV-Vis spectra of **1b** (20 μ M) in DMSO upon gradual addition of TBA (CH_3COO) (4.3×10^{-4} M); right: UV-Vis spectra of **1b** (20 μ M) in DMSO upon gradual addition of TBAH_2PO_4 (0 to 1.3 mM).

at the expense of reduced selectivity towards anions. Job's plot analysis indicated the formation of 1:1 stoichiometric interaction, in the case of all the three anions.

Anion binding study of the bis-PYBI receptors shows very interesting host-guest chemistry. For example, depending upon the basicity and coordination ability of the anions, either positive or negative cooperative effect between the two constituent PYBIs could be clearly noticed. Calculated equilibrium constants (K_a) value from the UV-Vis data are given in Table 2. For calculation of equilibrium constant, the reaction between **1a**, X^- and X^- (where X^- is anion) is considered. Although the values do not show any generalized trend w.r.t. basicity of the anions and the number of H-bond donor sites, a closer inspection reflects a few notable trends. First, the presence of two PIBYs, while it marginally enhances the binding affinity when bridged through insulating

sp^3 carbon, it is enhanced by more than one order when bridged through conjugating quinoxaline moiety, upon complexation with fluoride ion. For example, compared to **1a**, K_a of fluoride complexation is increased by more than two fold in the case of **2a**; on the other hand, in the case of **3**, K_a is increased by more than twelve fold. Similar trend was also noticed between **1b** and **2b** (*i.e.*, K_a increased by three fold). This can be attributed to lack of extended conjugation in the case of **2a** and **2b**, which could not stabilize the deprotonated hosts much. On the other hand, the positive cooperative effect between the two PYBIs during fluoride complexation by **3** could be ascribed to its extended conjugation that effectively stabilizes the negative charge upon decomplexation. Second, incorporation of electron withdrawing nitro group in the benzimidazole periphery (**2b**) and increased conjugation between the two PYBIs (**3**) led to reduced anion selectivity with additional guest

Table 2. Acid-base equilibrium constant, K_a (M^{-1}) of the receptors towards the interacting anions as determined by UV-Vis spectroscopy in DMSO at 25°C.

Host	F^-	CH_3COO^-	$H_2PO_4^-$
1a	8.45×10^2	–	–
1b	1.20×10^4	6.03×10^4	6.30×10^3
2a	1.90×10^3	–	–
2b	3.62×10^4	1.24×10^4	7.84×10^3
3	2.01×10^4	8.72×10^4	9.4×10^3

binding towards acetate and dihydrogen phosphate ions by these hosts (Figures S37 and S49 in Supplementary Information). Surprisingly, complexation of acetate ion with **2b** resulted in reduced affinity compared to that of **1b** by about five times, whereas dihydrogen phosphate ion complexation led to marginal increase in K_a (ratio of **2b/1b** $\sim$ 1.25). This type of negative cooperative effect between the two PYBIs could be explained on the basis of competition between them to bind the bidentate acetate ion. Whereas, the relatively larger size and presence of additional binding sites in the case of $H_2PO_4^-$ ion possibly reduced the negative cooperative effect.

Fluorescence spectroscopic titration studies were also performed in order to check the ability of these receptors to function as fluorometric anion sensors. Among the receptors synthesized, nitro analogues are non-fluorescent (fluorescence is below detection limit). Emission studies were carried out with 1 μ M solution of the remaining three receptors; *i.e.*, **1a**, **2a** and **3**. Like the UV-Vis spectroscopy observation, remarkable quench-

ing of fluorescence of receptors **1a** and **2a** was observed upon addition of F^- ion, whereas no significant change in the fluorescence intensity could be noticed in the case of other anions (see Supplementary Information). This observation further suggests interaction of receptors **1a** and **2a** with fluoride ion only. In the case of receptor **3**, addition of F^- , CH_3COO^- and $H_2PO_4^-$ ion resulted in almost complete quenching of its fluorescence but with different affinity. This may be ascribed to intramolecular photoinduced electron transfer (PET) and anion complexation triggered the electron transfer from the benzimidazole moiety to the ground state of the quinoxaline moiety.

3.2b 1H NMR study: To obtain further insight regarding the nature of interaction involved in the complexation process, 1H NMR titrations were carried out in DMSO- d_6 , with fluoride and acetate ions. In this experiment, the change in the 1H NMR signals upon gradual addition of the anionic analyte was monitored.

Addition of 0.17 equiv. of TBAF to **1a** in DMSO revealed disappearance of both the NH signals and instead a new broad peak arises at $\sim$ 12.1 ppm, which undergoes further broadening leading to complete disappearance upon subsequent additions of TBAF (Figure 6). After addition of $\sim$ 1.2 equiv. of fluoride ion, a new broad triplet starts appearing at 16 ppm, indicating the formation of HF_2^- ion, which indicates deprotonation of benzimidazole NH at higher concentration of F^- . Aromatic proton signals displayed upfield shift upon addition of fluoride ion, indicating an increase in the electron density on both the pyrrole and benzimidazole rings due to deprotonation. The strong electrostatic repulsion from **1b** $^-$ may be preventing the F^- ion to

Figure 6. Changes in the 1H NMR spectra of **1a** (5 mM) in DMSO- d_6 upon gradual addition of TBAF (0 to 12 mM).

Figure 7. Changes in the ^1H NMR spectra of **1b** (5 mM) in $\text{DMSO}-d_6$ upon gradual addition of TBAF (0 to 30 mM).

Figure 8. Changes in the ^1H NMR spectra of **1b** (5 mM) in $\text{DMSO}-d_6$ upon gradual addition of TBAAc (0 to 12 mM).

come closer to form H-bonding. The little downfield shift of the pyrrole NH proton indicates the involvement of pyrrole NH in weak H-bonding with the anion. However, broadening and subsequent disappearance of the signal upon increasing fluoride ion concentration might indicate the fast shuttling of the NH proton between the pyrrole and benzimidazole nitrogens, which is probably too fast to be detected in the NMR time scale (SI). On the other hand, titration of **1b** with fluoride ion showed deprotonation of the benzimidazole NH, while pyrrolic NH gradually broadens with slight upfield shift (Figure 7). The upfield shift of the pyrrolic NH is attributed to the increase in the electron density in the PYBI moiety, owing to the deprotonation of the benzimidazole NH. This probably reduces the hydrogen bonding interaction between the pyrrole NH and fluoride ion. This observation confirmed that the selection of the NH unit for interaction by anion is driven by the acidity of the NH. Titration of **1b** with acetate ion revealed disappear-

ance of the benzimidazole NH and downfield shift of the pyrrole NH with concomitant upfield shift of the aromatic protons (Figure 8). This observation indicates that more acidic benzimidazole N-H proton gets deprotonated and pyrrole N-H is involved in a relatively stronger H-bonding (than with fluoride ion) with the protonated guest, possibly owing to its complementary bidentate coordination mode (Figure S29, in SI). Furthermore, titration of **2b** with fluoride and acetate ions displayed similar binding patterns, as observed in the case of **1b** (SI). ^1H NMR titration of receptor **3** with fluoride ion showed two new peaks at 20.5 and 16 ppm, along with a very broad peak between 12 and 16 ppm indicating the very slow kinetics of the binding and deprotonation equilibrium (Figure 9). Interestingly, gradual addition of TBAF up to ~ 2 equiv. resulted in sharpening of the peaks at 20.5 and 16 ppm with concomitant loss of the broad peak between 12 to 16 ppm. Further addition of TBAF resulted in upfield shift of the peak at 20.5 ppm

Figure 9. Changes in the ^1H NMR spectra of **3** (5 mM) in $\text{DMSO-}d_6$ upon gradual addition of TBAF (0 to 25mM).

Figure 10. Probable H-bonding structures of mono- and di-deprotonated forms of **3**.

to 19 ppm, while the broad peak at 16 ppm resolved into a triplet without undergoing shift. The peak at 20.5 ppm emerging after addition of 1 equiv. of fluoride ion (with slight upfield shift of both uncomplexed benzimidazole and pyrrole signals), may possibly be attributed to deprotonation of one of the benzimidazole moiety, followed by strong complexation of one of the pyrrolic NH with fluoride ion and benzimidazole Nitrogen (Figure 10, Complex A). On the other hand, evolution of the signal corresponding to HF_2^- ion indicates the deprotonation of the benzimidazole protons, and the appearance of the broad peaks in between 12 and 16 ppm signifies the intramolecular proton migration in the dianionic $\mathbf{3}^{2-}$ species. Possibly, the pyrrolic NHs migrated towards the benzimidazole nitrogens and trapped inside the dianionic binding domain, by forming strong $\text{N-H}\cdots\text{NH}$ -bonding within the cavity resulting in the peak at 19 ppm (Figure 10, Complex B).¹⁷ Further, titration of **3** with acetate ion showed similar type of

changes in the β -CH pyrrolic region while no new signal was observed in the downfield region, as seen in the case of fluoride ion binding (see SI). Upon increasing the amount of acetate ion, pyrrole NH peaks get broadened without any noticeable shift. The results of ^1H NMR titration experiment corroborated the observations of the UV-Vis study, which suggested the presence of two equilibrium processes in the case of **1a** and **2a**, namely, the H-bonded adduct formation for relatively lower F^- concentration, whereas at higher concentration, Bronsted acid-base equilibrium prevails. However, in the case of nitro derivatives (**1b** and **2b**), addition of fluoride ion leads to immediate deprotonation of the benzimidazole NH and subsequent charge transfer to the nitro group leading to color change. Incorporation of nitro group in the benzimidazole periphery drastically enhances the acidity of the benzimidazole NH compared to the pyrrole NH, thereby increasing its deprotonation tendency. Receptor **3** also follows the deprotonation mechanism.

4. Conclusions

A family of colorimetric chemosensors based on pyrrole-benzimidazole conjugates (PYBI) were synthesized and their anion recognition behaviour was investigated in detail. Receptor **1a** and **2a** act as colorimetric sensors for fluoride ion only. The ^1H NMR titration experiment revealed that the binding of the fluoride ion is through N-H...F⁻ interaction. However, addition of more than one equivalent of anion eventually led to the deprotonation. Further, we have observed that incorporation of nitro group upon benzimidazole periphery induces the colorimetric response (in case of **1b** and **2b**) but at the expense of selectivity by exhibiting tendency towards binding with acetate and dihydrogenphosphate ions. This observation reveals that increasing the acidity of the H-bonding proton without affecting the receptor geometry sometimes leads to a drastic change in anion selectivity. Addition of fluoride ion to **1b** and **2b** deprotonates benzimidazole NH, while reducing H-bonding interaction with the pyrrolic NH. Acetate ion deprotonates benzimidazole NH and the corresponding conjugate acid; *i.e.*, acetic acid forms strong H-bonding with the receptor *via* complementary bidentate coordination mode. Like **1b** and **2b**, receptor **3** interacts with fluoride and acetate ions, and relatively weakly with dihydrogenphosphate ion. Interestingly, fluoride ions abstract, stepwise, both the benzimidazole NHs of receptor **3** in a unique way and as a consequence, possibly forcing delocalization of pyrrolic NHs in the binding domain, as indicated by the highly deshielded proton signal ($\sim 19\text{ppm}$), as well as a stronger colorimetric and fluorometric response.

Supplementary Information (SI)

Synthetic procedures, ^1H and ^{13}C NMR spectra, absorption and emission spectra, anion binding data and computational data are available in Supplementary Information at www.ias.ac.in/chemsci.

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