

A Multifunctional Fluorescence Probe for the Detection of Cations in Aqueous Solution: the Versatility of Probes Based on Peptides

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Abstract We synthesized a tetra-functional fluorescence probe based on dansyl and peptide motif, dansyl-Gly-Trp (DGT, **1**), that efficiently bound several metal ions and showed distinguishing optical properties. The probe **1** could respond to Hg^{2+} with enhanced and blue-shifted fluorescence emission but to Cu^{2+} with obvious fluorescence quenching. In addition, **1** was sensitive to pH ranging from 2.0 to 5.0 and precipitated in the presence of Pb^{2+} at neutral conditions. The combination of these intrinsic properties with the selective responses to different chemical inputs allows this system to be implemented as an ionic switch. Furthermore, **1** could penetrate the cell membrane and accumulated well in intracellular region. The underlying mechanisms of the probe to different kind of metal ion were explored successfully by using either ^1H NMR, NOESY, electron paramagnetic resonance (EPR) or FT-IR spectra. In addition, to investigate the binding model of 1/Hg^{2+} and 1/Cu^{2+} , simulations were also performed by using density functional theory (DFT) and reasonable binding configurations were achieved for these two complexes.

Keywords Peptide · Fluorescence probe · Multifunction · Mercury · Copper

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Introduction

The design and synthesis of optical probes with high selectivity and sensitivity to heavy transition-metal ions (HTMs), particularly mercury, lead, and copper ions, have attracted special attention due to the toxicity of these ions, both in physiological systems and in the environment. In numerous successful applications, the design of probes, has been essentially based on structure-activity relationship [1–3]. However, most probe designs were based on the assumption that one probe would bind to one target. Thus, the majority of probes have been restricted to measuring the concentration of one specific target ion. However, quite often, the contamination of physiological and/or environmental conditions results from the synergistic contribution of two or more infectants. Therefore, it would be beneficial to develop probes with multifunctional capabilities that could identify two or more infectants in a system. Moreover, it would be very useful, both in chemistry and material sciences, to develop fluorescence probes for metal ions that could be integrated into molecular level devices, like logic gates and/or ionic switches [4–7].

In the previous studies, the concept that introducing amino acids as binding site of probe have been proved successful [8–10]. Highly sensitive fluorescence probes derived from pyrene butyric acid for Pb^{2+} [8] and dansyl moiety for Hg^{2+} [9, 10] have been achieved, which displayed rapid and specific optical responses to target metal ion in buffered aqueous solution. In the present study, instead of using single amino acid, the motif of peptide will be introduced to design fluorescence probe for metal ions. By tuning the number and sequence of participant amino acids several attractive features of probe will be achieved as: (i) felicitous hydrophilicity and hydrophobicity, in keeping the balance between water-solubility and cell

permeability; (ii) easily optimized high selectivity and sensitivity to different metal ions; and essentially (iii) possessing more binding site in comparison to a single amino acid, which supply a possibility to be used as a logic gate or molecular switch to different kinds of metal ions. Although some probes based on peptide motif had been developed for metal ions [11–14], they could not response to different metal ions simultaneously. In the present study, two dansyl labeled peptides, dansyl-Gly-Trp (DGT, **1**) and dansyl-Gly-Gly-Trp (DGGT, **2**) (Fig. 1) were synthesized, in possessing of multiple binding sites that would confer the potential to be used as a logic gate or molecular switch for different kinds of metal ions.

Experimental

Materials and Apparatus

All chemicals were obtained from commercial suppliers and used without further purification. Mass spectra were measured on a QTRAP mass spectrometer (Applied Biosystems Sciex Foster city, USA). Infrared (IR) spectra were recorded on a Bruker Vertex 80 V FTIR spectrophotometer.

General Method

Fluorescence emission spectra were recorded on a Shimadzu (Japan) RF-5301PC fluorescence spectrophotometer. To reduce the fluctuation in the excitation intensity during measurement, the lamp was kept on for 1 h and the samples were stock-still for 3 min prior to the experiment. Samples for emission measurement were placed in a 1 cm × 1 cm quartz cuvettes (4 mL volume). All spectroscopic measurements of **1** and **2** were performed in 10.0 mM HEPES buffer solution (pH 7.4), and the concentrations of **1** and **2** in all the fluorescent experiments are 20.0 μM. A

fixed excitation wavelength at 330 nm was used. Fluorescence titration was performed in buffer solution using respective chloride salt of metal ions except for AgNO₃.

UV–vis absorption spectra were recorded on a Shimadzu (Japan) UV-3100 spectrophotometer. To reduce the fluctuation of baseline in absorbance spectra, the lamp was kept on for 0.5 h before the measurement. Sample for measurement was placed in a 1 cm × 1 cm quartz cuvette (4 mL volume). The absorption spectra of **1** (20.0 μM) were measured in 10.0 mM HEPES buffer solution (pH 7.4).

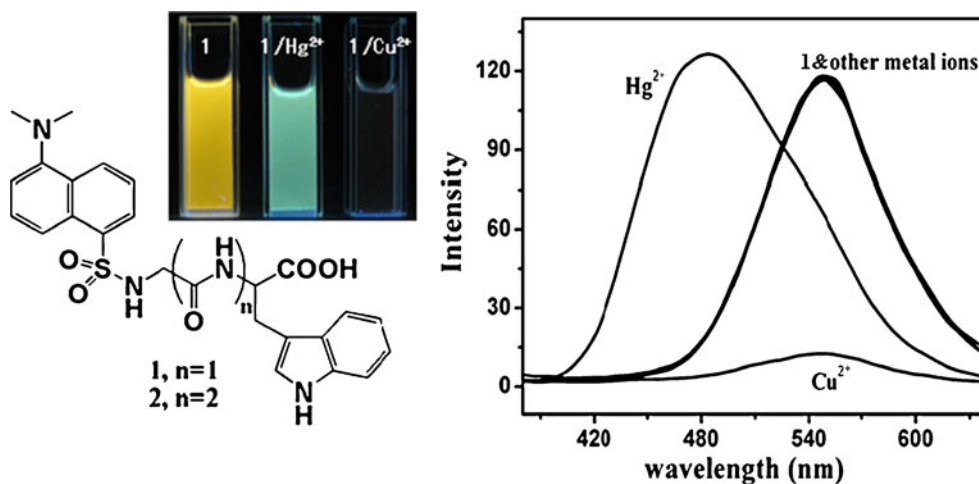
Turbidity measurements of **1** were recorded on an Ocean Optic usb-4000 spectrophotometer upon addition of Pb²⁺. The lamp was kept on for 0.5 h before measurements. Sample for test was placed in a 1 cm × 1 cm cuvette (4 mL volume). The absorption spectra of **1** (20 μM) with addition of Pb²⁺ were recorded in 10.0 mM HEPES buffer solution (pH 7.4) with stirring.

¹H NMR titration experiments were conducted on a Bruker Varian Unity-500 MHz spectrometer. We carried out ¹H NMR titration experiments on a 5 × 10⁻⁴ M solution of **1** in D₂O containing 0.3% (v/v) d₆-DMSO. The spectra were recorded in the presence of 0.5 and 1 equiv. Hg²⁺, respectively at room temperature. The NOESY experiment was performed at 25 °C in the presence of 1 equiv. Hg²⁺.

The EPR spectra were recorded on a JES-FA 200 EPR spectrometer. The instrumental parameters were as follows: scanning frequency, 9.45 GHz; central field, 3301.85 G; scanning width, 8,000 G; scanning power, 0.998 mW; scanning temperature, 25 °C. 2,2-Diphenyl-1-picrylhydrazyl (DPPH) was used as a reference for the calculation of spin concentration. The precipitate for EPR test was obtained from the solution of **1** and equivalent CuCl₂.

Density functional theory simulations were performed with the B3LYP functional and the mixed “Double-ζ” quality basis sets 6–31 G(d) for C, H, O, N, S and

Fig. 1 Tetra-functional probes (**1** and **2**) based on a dansyl moiety and a peptide motif. (Left) Chemical structure of **1** and **2**; (middle) fluorescence images of **1**, **1**/Hg²⁺, and **1**/Cu²⁺; (right) fluorescence spectra of 20 μM **1** in 10 mM HEPES solution (pH 7.4) in the presence of 40 μM Hg²⁺, 80 μM Cu²⁺, and 100 μM of other metal ions (K⁺, Na⁺, Mg²⁺, Al³⁺, Zn²⁺, Fe³⁺, Mn²⁺, Co²⁺, Cd²⁺, Ca²⁺, Ag²⁺, Ni²⁺) (λ_{ex} = 330 nm)



LANL2DZ for Hg and Cu. Optimization of the complexes was performed without any constraints for all the complexes. All calculations were carried out using Gaussian 09 [15].

Cell Permeability

Cells Culture

HEK-293 cells were cultured according to the reported protocol [16]. Briefly, cells were cultured at 37 °C in Dulbecco's modified Eagle's medium (DMEM, Sigma) containing 10% fetal bovine serum (FBS, Gibco) in a 5% CO₂/95% air incubator. Cells were plated 24 h before study into 6 well plates and at the time of study, the medium was changed with a fresh 2% serum one.

Fluorescence Imaging Measurements of Probe in Cells

Imaging experiments were performed with a FV1000 confocal laser-scanning fluorescent microscope (Olympus, Japan) with a 20× objective lens. Excitation of **1** and **2**-loaded cells at 405 nm were carried out with an argon-ion laser, and emission was collected in a window from 480 to 580 nm. **1** or **2** (from a 1.0 mM stock solution in DMSO) was added to each well except the control of cultured cells and a 20 μM of **1** or **2** was obtained in final. The cells were incubated with **1** or **2** for 1 h at 37 °C, and then the medium was removed and the cells were washed thoroughly twice by PBS (pH 7.0). After that, the fluorescence microscopic image was recorded for each case.

General Procedure for the Synthesis of **1** and **2**

Phthalimidoglycine (**4**)

Firstly, the mixture of phthalic anhydride (1.63 g, 11 mM) and glycine (0.75 g, 10 mM) was fused at 140 °C for 0.5 h. And then it was cooled to room temperature and the resulted solid was recrystallized from ethanol/water solution. After filtrating, drying in vacuum, we obtained compound **4** (1.95 g, 95%) as a light yellow solid. ¹H NMR (500 MHz, d₆-DMSO): δ=4.32 (s, 2H, CH₂), δ=7.89–7.96 (m, 4H, Pht-H).

Methyl 2-(2-(1,3-dioxisoindolin-2-yl)acetamido)-3-(1H-indol-3-yl) propanoate (**6**)

Thionyl chloride (1.5 equiv.) was added dropwise to the solid state of **4** (1 g, 4.87 mM), and the mixture was heated under reflux for 6 h. The excess of thionyl chloride was

evaporated in vacuum, and the residue was recrystallized from n-hexane three times. The produced white solid was dissolved in CH₂Cl₂ (20 mL), and then was added dropwise to an ice-cooled mixture of L-tryptophan methyl ester (1.1 equiv.) and triethylamine (1.2 equiv.) in CH₂Cl₂ (20 mL). The mixture was stirred overnight at room temperature. After dilution with CH₂Cl₂, the mixture was washed with saturated NaHCO₃ aqueous solution, diluted HCl, and brine step by step. The organic phase was dried over anhydrous MgSO₄, filtered and evaporated under reduced pressure. The resulted residue was purified by column chromatography to produce a yellow solid **6** (1.68 g, 85%). ¹H NMR (500 MHz, d₆-DMSO): δ=3.05–3.19 (m, 2H, CHCH₂), δ=3.58 (s, 3H, COOCH₃), δ=4.24 (s, 2H, CH₂CONH), δ=4.50–4.55 (m, 1H, CHCOOCH₃), δ=6.98–7.49 (m, 5H, indole-H), δ=7.87–7.93 (m, 4H, Pht-H), δ=8.76–8.77 (d, 1H, CONH), δ=10.93 (s, 1H, indole-NH).

Methyl 2-(2-aminoacetamido)-3-(1H-indol-3-yl)propanoate (**7**)

To a solution of pseudodipeptide **6** (1 g, 2.5 mM) in EtOH, hydrazine monohydrate (200 μL) was added. The mixture was stirred at room temperature for 72 h, and then it was cooled to 0 °C and phthalhydrazide was removed by filtering. The filtrate was concentrated and the resulted oil was redissolved in a minimum amount of EtOH. The solution was cooled to 0 °C, filtered and evaporated under reduced pressure. The resulted pale yellow oil **7** was immediately used for the next reaction, without further purification.

Methyl 2-(2-(5-(N,N-dimethylamino)naphthalene-1-sulfonamido)acetamido)-3-(1H-indol-3-yl)propanoate (**8**)

In an ice bath, a solution of dansyl chloride (0.62 g, 2.08 mM) in CH₂Cl₂ (20 ml) was added slowly to a mixture of **7** and triethylamine in CH₂Cl₂ (30 ml) and then was stirred for 0.5 h. The reaction was kept stirring for 12 h at room temperature, and the progress of the reaction was monitored by thin layer chromatography (TLC). After dilution with CH₂Cl₂, the mixture was washed with saturated NaHCO₃ aqueous solution, diluted HCl, and brine one by one. The organic phase was dried over anhydrous MgSO₄, filtered and evaporated under reduced pressure. The resulted residue was purified by column chromatography to produce a yellow solid **8** (0.9 g, 86%). ¹H NMR (500 MHz, d₆-DMSO): δ=2.81 (s, 6H, CH₃NCH₃), δ=2.90–3.05 (m, 2H, CHCH₂), δ=3.51–3.53 (m, 5H, CH₂CONH + COOCH₃), δ=4.39–4.43 (m, 1H, CHCOOCH₃), δ=6.94–8.27 (m, 11H, dansyl-H + indole-H), δ=8.27–8.29 (d, 1H, CONH), δ=8.43–8.44 (d, 1H, SO₂NH), δ=10.87 (s, 1H, indole-NH).

2-(2-(5-(N,N-dimethylamino)naphthalene-1-sulfonamido)acetamido)-3-(1H-indol-3-yl)propanoic acid (**1**)

To a stirred solution of **8** (0.5 g) in methanol, 1 M NaOH (50 mL) was added, and the mixture was stirred overnight at room temperature. The aqueous phase was acidified to pH 5–6 by using an aqueous solution of HCl, and then extracted with EtOAc three times. The combined organic phase were washed with brine, dried over Na₂SO₄, filtered and evaporated under reduced pressure step by step. Purification of **1** was performed by using column chromatography on silica gel. Target molecule **1** was obtained as a yellow solid (0.42 g, 85%). ¹H NMR (500 MHz, d₆-DMSO): δ=2.81 (s, 6H, CH₃NCH₃), δ=2.90–3.10 (m, 2H, CHCH₂), δ=3.43–3.48 (m, 2H, CH₂CONH), δ=4.33–4.37 (m, 1H, CHCOOH), δ=6.91–8.9 (m, 11H, dansyl-H + indole-H), δ=8.27–8.29 (d, 1H, CONH), δ=8.42–8.44 (d, 1H SO₂NH), δ=10.82 (s, 1H, indole-NH). ¹³C NMR (500 MHz, d₆-DMSO): δ=28.14 (CHCH₂), δ=42.68 (NCH₃), δ=45.93 (CH₂CONH), δ=55.0 (CHCOOH), δ=110.97–152.2 (dansyl-C + indole-C), δ=167.74 (CH₂CONH), δ=174.6 (CHCOOH). IR (KBr) 3,394, 2,932, 2,863, 2,834, 2,785, 1,720, 1,663, 1,615, 1,587, 1,573, 1,458, 1,410, 1,326, 1,160, 1,144, 1,098 cm⁻¹. Mass spectral data (ESI-MS): for C₂₅H₂₆N₄SO₅, calcd. 494.16; found 495.1.

Methyl 2-(2-(2-(1,3-dioxisoindolin-2-yl)acetamido)acetamido)-3-(1H-indol-3-yl)propanoate (**9**)

The obtained pale yellow oil **7** was dissolved in a mixture of 1:20 (v/v) DMF and CH₂Cl₂. To the solution, 1.5 equiv. Et₃N and 1.1 equiv. **4** were added with stirring. When all the reagents dissolved, 1.5 equiv. 1-Ethyl-3-(3-dimethylaminopropyl)carbodiimide (EDC) was added, and the mixture was stirred for 5–6 h at room temperature and the progress of the reaction was monitored by TLC. After dilution with CH₂Cl₂ the mixture was washed with saturated NaHCO₃ aqueous solution, diluted HCl, and brine. The organic phase was dried over anhydrous MgSO₄, filtered and evaporated under reduced pressure. The resulted residue was recrystallized from ethanol to give a pale yellow solid **9**. ¹H NMR (500 MHz, d₆-DMSO): δ=3.03–3.18 (m, 2H, CHCH₂), δ=3.57 (s, 3H, COOCH₃), δ=3.70–3.81 (m, 2H, CONHCH₂CONH), δ=4.26 (s, 2H, CH₂CONH), δ=4.51–4.55 (m, 1H, CHCOOCH₃), δ=6.97–7.45 (m, 5H, indole-H), δ=7.86–7.92 (m, 4H, Pht-H), δ=8.33–8.35 (d 1H, CONHCH), δ=8.48–8.50 (t, 1H, CONHCH₂), δ=10.86 (s, 1H, indole-NH).

The target molecule **2** and the intermediates in the synthesis route were obtained under the identical conditions as mentioned, and the corresponding assayed results by ¹H NMR and Mass Spectrum are described as follows:

Methyl 2-(2-(2-(5-(N,N-dimethylamino)naphthalene-1-sulfonamido)acetamido)acetamido)-3-(1H-indol-3-yl)propanoate (**11**)

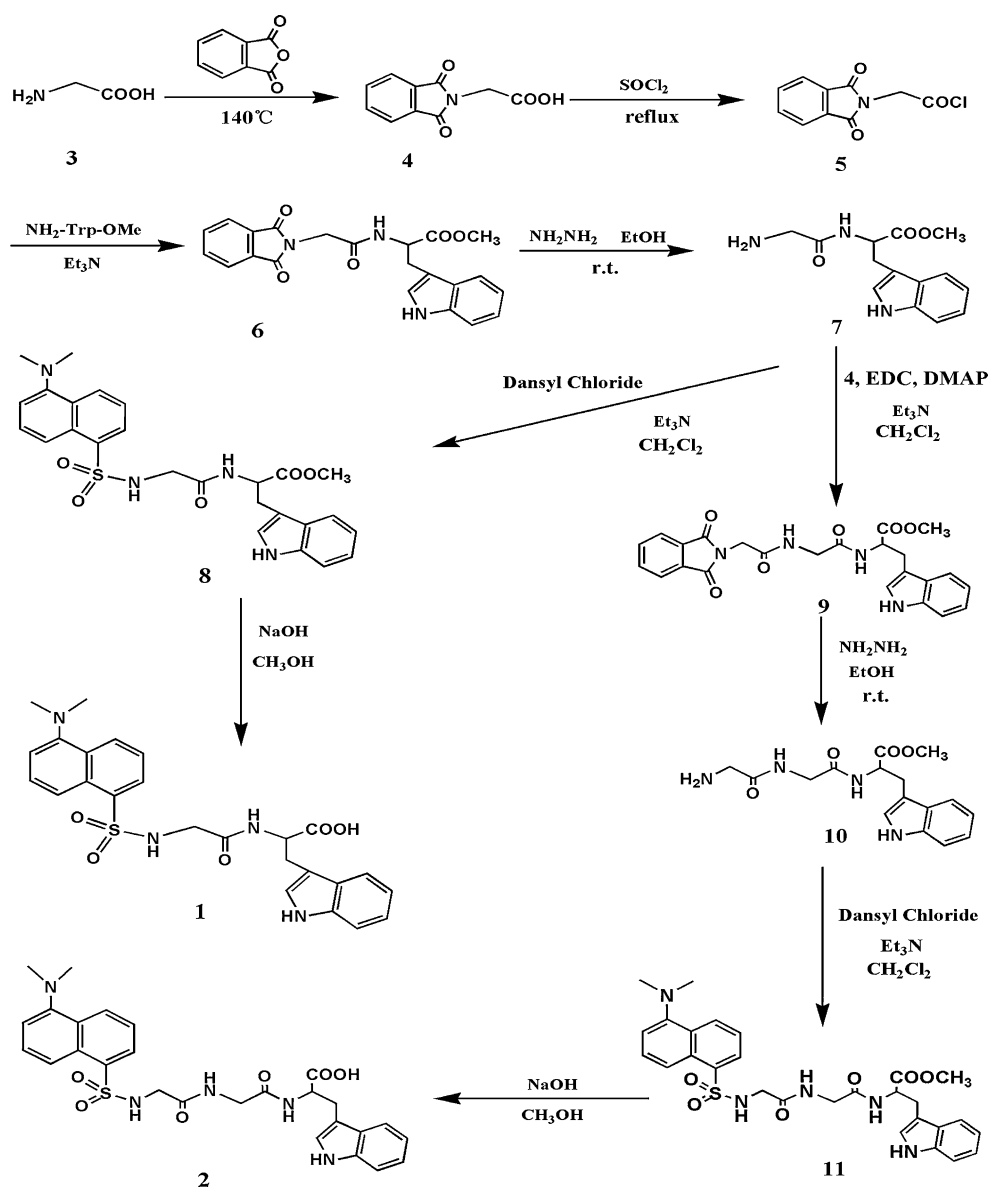
¹H NMR (500 MHz, d₆-DMSO): δ=2.82 (s, 6H, CH₃NCH₃), δ=3.01–3.17 (m, 2H, CHCH₂), δ=3.48 (s, 2H, SO₂NHCH₂CONH), δ=3.60–3.71 (m, 2H, CONHCH₂CONH), δ=4.49–4.53 (m, 1H, CHCOOCH₃), δ=6.91–8.28 (m, 11H, dansyl-H + indole-H), δ=8.29–8.31 (m, 2H, CONHCH₂CONH), δ=8.44–8.46 (d, 1H SO₂NH), δ=10.86 (s, 1H, indole-NH).

2-(2-(2-(5-(N,N-dimethylamino)naphthalene-1-sulfonamido)acetamido)acetamido)-3-(1H-indol-3-yl)propanoic acid (**2**)

¹H NMR (500 MHz, d₆-DMSO): δ=2.82 (s, 6H, CH₃NCH₃), δ=2.97–3.19 (m, 2H, CHCH₂), δ=3.47 (s, 2H, SO₂NHCH₂CONH), δ=3.56–3.69 (m, 2H, CONHCH₂CONH), δ=4.33–4.41 (m, 1H, CHCOOH), δ=6.94–8.11 (m, 11H, dansyl-H + indole-H), δ=8.27–8.31 (m, 2H, CONHCH₂CONH), δ=8.44–8.46 (d, 1H SO₂NH), δ=10.78 (s, 1H, indole-NH). ¹³C NMR (500 MHz, d₆-DMSO): δ=28.14 (CHCH₂), δ=42.68 (NCH₃), δ=45.93 (CONHCH₂CONH), δ=46.11 (SO₂NHCH₂CONH) δ=54.40 (CHCOOH), δ=110.97–152.2 (dansyl-C + indole-C), δ=168.77 (CONHCH₂CONH), δ=168.90 (CONHCH₂CONH), δ=174.6 (CHCOOH). IR (KBr) 3,361, 2,974, 2,865, 2,789, 1,720, 1,660, 1,533, 1,455, 1,407, 1,348, 1,330, 1,230, 1,143 cm⁻¹. Mass spectral data (ESI-MS): for C₂₇H₂₉N₅SO₆, calcd. 551.18; found 552.1.

Results and Discussion

We started the synthesis with free glycine (**3**), which was firstly phthaloylated with phthalic anhydride to give N-phthalimido-protected amino acid **4** in a high yield. The resulted pure **4** was reflux in excess of thionyl chloride to give carboxyl chloride **5** which was coupled with C-protected L-tryptophan to give methyl esters **6** directly without further purification. Hydrazinolysis of the phthalimido protected **6** produced the crucial amine intermediate **7**, which was unstable to heat and prolonged storage at room temperature. Free amine **7** was immediately mixed with dansyl chloride in CH₂Cl₂ solution. The resulted compound **8** was converted by alkaline hydrolysis to the first target product, dansyl-Gly-Trp (**1**). The dansyl labeled tripeptide probe, dansyl-Gly-Gly-Trp (**2**), was obtained by using the similar method (Scheme 1). After synthesis, probe **1** and **2** were assayed with Mass, Infrared, ¹H NMR and ¹³C NMR spectroscopy.

Scheme 1 Synthesis route of probes **1** and **2**

pH Effect of DGT

In solutions with pH values that ranged from 7.4 to 5.0, probe **1** did not exhibit any detectable changes in either the UV-visible absorption or the fluorescence emission spectrum. However, when the pH was lower than 5.0, with pH decrease, the intensity of a absorption band at 284 nm increased with a simultaneous decreases of the intensity of the absorption bands at 250 and 330 nm (left in Fig. 2). In addition, the corresponding fluorescence emission spectra showed unilateral quenching of a band at 550 nm (right in Fig. 2). Both results were ascribed to the protonation of the dimethylamino group in **1**, because the absorption band at 284 nm was characteristic of the protonated form of the dansyl moiety [6, 17, 11].

Absorption and Fluorescence Titration with Hg^{2+}

When Hg^{2+} ($1\sim 80\ \mu\text{M}$) was added to **1**, we observed a significant decrease in the emission band at 550 nm and a simultaneous increase in a new band at 485 nm (left in Fig. 3). The plot of intensity ratios (I_{485}/I_{550}) to Hg^{2+} concentration (right in Fig. 3) indicates a detection limit around $5\ \mu\text{M}$ of **1** to Hg^{2+} . In addition, the UV-vis absorption spectra displayed a decrease of the intensity of absorption band at 330 nm and an increase at lower wavelengths (Fig. S1†). Both changes were attributed to the formation of a **1**/ Hg^{2+} complex, which reduced the ability of the sulfonamide group to donate electrons to the dansyl moiety. This inhibited the internal charge transfer between the amide group and the dansyl moiety, and

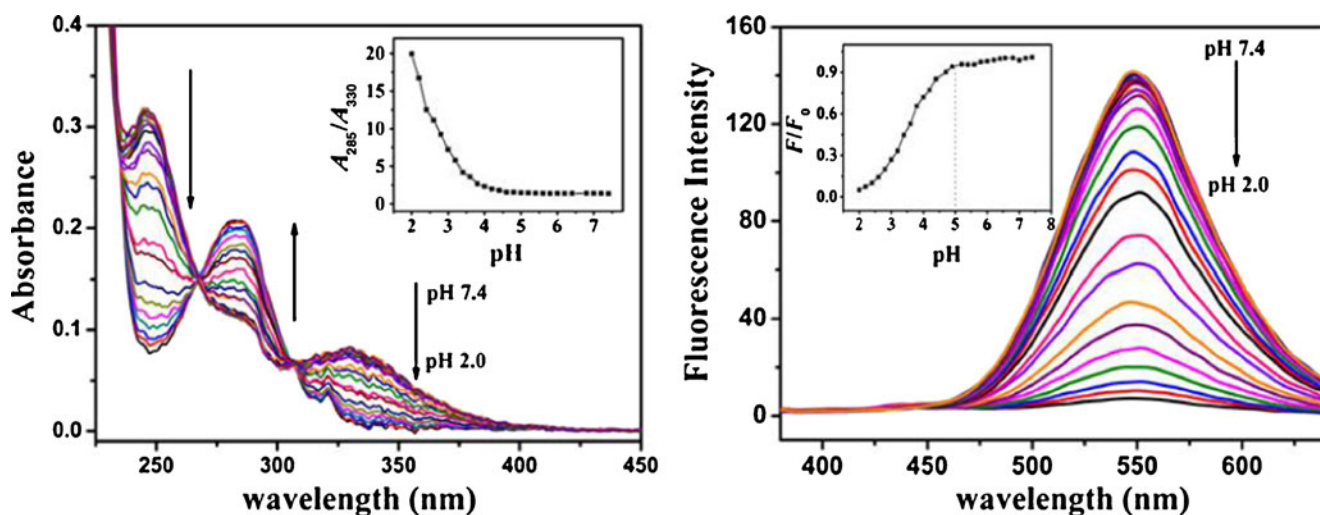


Fig. 2 (left): Variations of absorbance spectra of **1** (20 μM) at different pH values (2.0–7.4); *Inset*: the plot of absorbance band ratio at 285 to 330 nm (A_{285}/A_{330}) as a function of pH. (right): Fluorescence

emission spectra of **1** (20 μM) at different pH values (2.0–7.4); The emission intensity at 550 nm quenched with pH decrease when pH is lower than 5.0 ($\lambda_{\text{ex}}=330$ nm)

consequently, both the absorption and emission bands shifted toward the blue region of the spectra. The changes in the fluorescence emission ratio of **1** with Hg^{2+} concentration were fit with the B-H plot [18]. The resulting good linear fit demonstrated that the $1/\text{Hg}^{2+}$ complex formed at a 1:1 stoichiometry with a $1.18 \times 10^3 \text{ M}^{-1}$ binding constant (Fig. S2 \dagger). Moreover, **1** exhibited a 0.5 M fraction maximum on the Job's plot, which further confirmed a 1:1 stoichiometry for the $1/\text{Hg}^{2+}$ complex (Fig. S3 \dagger).

Next, we investigated the fluorescence of **1** to Hg^{2+} in different pH conditions. Probe **1** exhibited adequate sensitivity for Hg^{2+} at pH between 6.0 and 8.0, with a maximum I_{485}/I_{550} value at pH=7.4 (Fig. S4 \dagger). This

indicated that **1** was most sensitive to Hg^{2+} at physiological pH.

Binding Model of DGT/ Hg^{2+}

We explored the coordination model between **1** and Hg^{2+} with both ^1H NMR titration (Fig. 4 and Table S1 \dagger) and 2D Nuclear Overhauser Effect Spectroscopy (NOESY) (Fig. 5). The addition of 1 equiv. of Hg^{2+} caused large chemical shifts in the H(8) ($\Delta\delta=0.08$ ppm), H(5) ($\Delta\delta=0.04$ ppm), H(3) ($\Delta\delta=0.1$ ppm), and H(9) ($\Delta\delta=0.03$ ppm) of **1**, indicating that Hg^{2+} interacted with **1** via the sulfonamide moiety. In addition, the chemical shifts of H

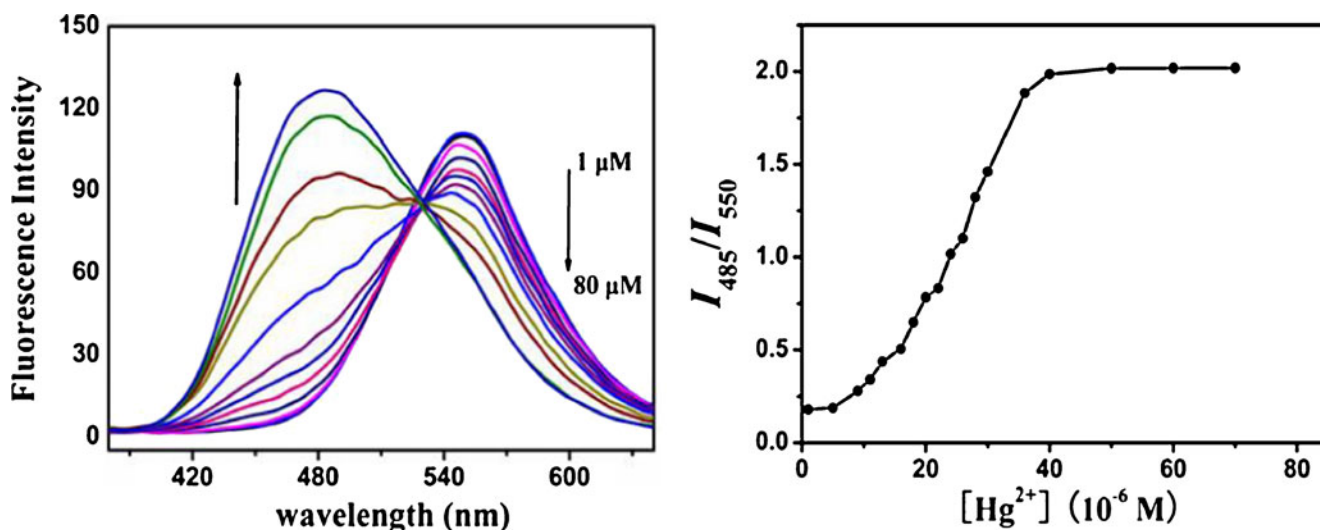
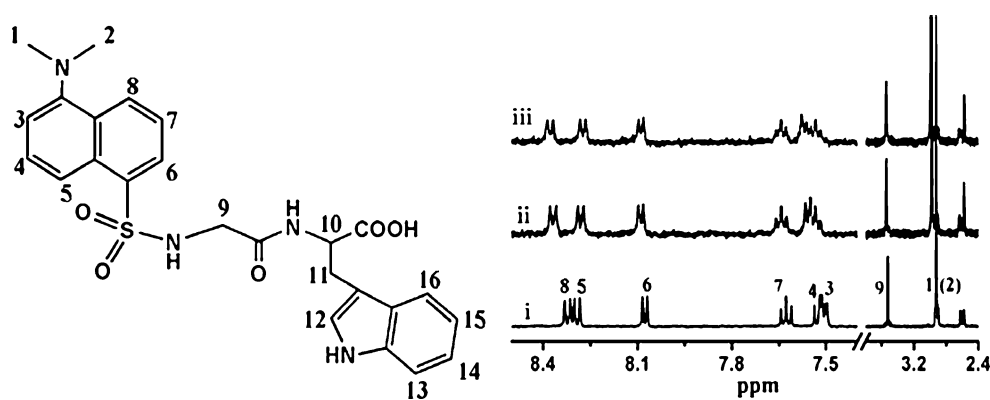


Fig. 3 Formation of a complex between probe **1** and Hg^{2+} . (left) Fluorescence titration spectra of **1** (20 μM) in buffer with different concentrations of Hg^{2+} ; (right) the corresponding plot of intensity ratios (I_{485}/I_{550}) vs. Hg^{2+} concentration ($\lambda_{\text{ex}}=330$ nm)

Fig. 4 Typical ^1H NMR spectrum of **1** (i) alone and in the presence of (ii) 0.5 and (iii) 1 equiv. of Hg^{2+} in D_2O solutions containing 0.3% DMSO-d_6



(12) ($\Delta\delta=0.027$ ppm) and H(14) ($\Delta\delta=0.01$ ppm) demonstrated that the indole ring may have a cooperative effect on Hg^{2+} complexation. In the NOESY spectrum, the strong cross peaks between $\text{H}_5\text{--H}_9$, $\text{H}_6\text{--H}_9$, $\text{H}_1\text{--H}_{12}$, $\text{H}_1\text{--H}_{15}$, and $\text{H}_1\text{--H}_{16}$ (Fig. 5) showed that the indole ring and methylene were spatially close to naphthalene. Due to the flexibility of peptides, it is reasonable that Hg^{2+} interact with the multisites of **1** to form a stable complex. A density functional theory (DFT) simulations of possible structures provide a verification of our assumption and interpret ^1H correlation data well. A simulated structure with the binding configuration revealed that the sulfonamide in **1** is the primary binding sites for Hg^{2+} (left in Fig. 6).

Absorption, Fluorescence Spectrum and Binding Model of DGT/ Cu^{2+}

In contrast to the changes observed with Hg^{2+} , the addition of Cu^{2+} to **1** induced a gradual fluorescence quenching. This was mainly attributed to the paramagnetic nature of Cu^{2+} and factors induced either by the formation of a $1/\text{Cu}^{2+}$ complex or by the close proximity of free Cu^{2+} to the dansyl moiety [19–21]. The latter possibility of close proximity was

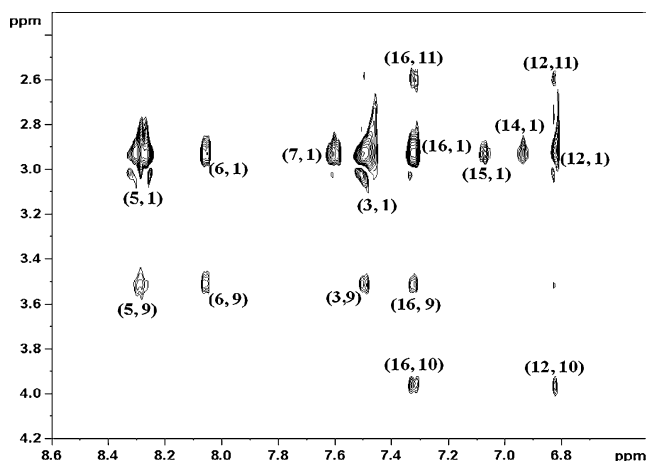


Fig. 5 The coordination between probe **1** and Hg^{2+} . Partial 500 MHz $^1\text{H}\text{--}^1\text{H}$ Nuclear Overhauser Effect Spectroscopy (NOESY) spectra of **1** in the presence of 1 equiv. of Hg^{2+} in D_2O solution with 0.3% DMSO-d_6

ruled out by the changes observed in the absorption spectra of **1** after the addition of Cu^{2+} (left in Fig. 7). In particular, the large differences in the X-band electron paramagnetic resonance (EPR) spectra between CuCl_2 and $1/\text{Cu}^{2+}$ (Fig. 8) indicated that there was a strong interaction between **1** and Cu^{2+} . The large contribution of the carboxyl in **1** to Cu^{2+} binding was confirmed by using a control compound, **8** (Fig. S5 \dagger), which had the same structure as **1** except the carboxyl was esterified with a methyl group. Upon the addition of Cu^{2+} , **8** exhibited only very slight fluorescence quenching. Moreover, because Cu^{2+} could deprotonate the amide nitrogen when the $\text{pH}>4$ [19], we proposed that **1** became a loop structure after Cu^{2+} formed a complex with the sulfonamide moiety, the amide group, and the carboxylate group. DFT simulation was also conducted to optimize the configuration of $1/\text{Cu}^{2+}$ (right in Fig. 6), which confirmed the proposed structure for $1/\text{Cu}^{2+}$. The fluorescence intensity ratio, F/F_0 , in decreasing with Cu^{2+} concentration (inset in Fig. 7), revealed a 1 μM detection limit of **1** to Cu^{2+} in aqueous solution. This detection limit is to be satisfactory to the maximum concentration of Cu^{2+} allowed in drinking water by the U. S. EPA (~ 20 μM) [22]. When the S-V equation [23] was fit to the fluorescence titration data of Cu^{2+} to **1**, we found that the complex was formed at a 1:1 stoichiometry with a binding constant of $1.58 \times 10^5 \text{ M}^{-1}$ (Fig. S6 \dagger). The stoichiometry was further supported by the Job's plot (Fig. S7 \dagger). Of not is that the further addition of EDTA or histidine could fully snatch **1** from the $1/\text{Cu}^{2+}$ complex (Fig. S8 \dagger), which

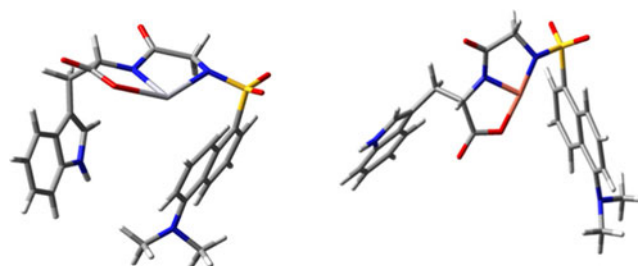


Fig. 6 The simulated structural configuration of $1/\text{Hg}^{2+}$ (left) and $1/\text{Cu}^{2+}$ (right) complexes

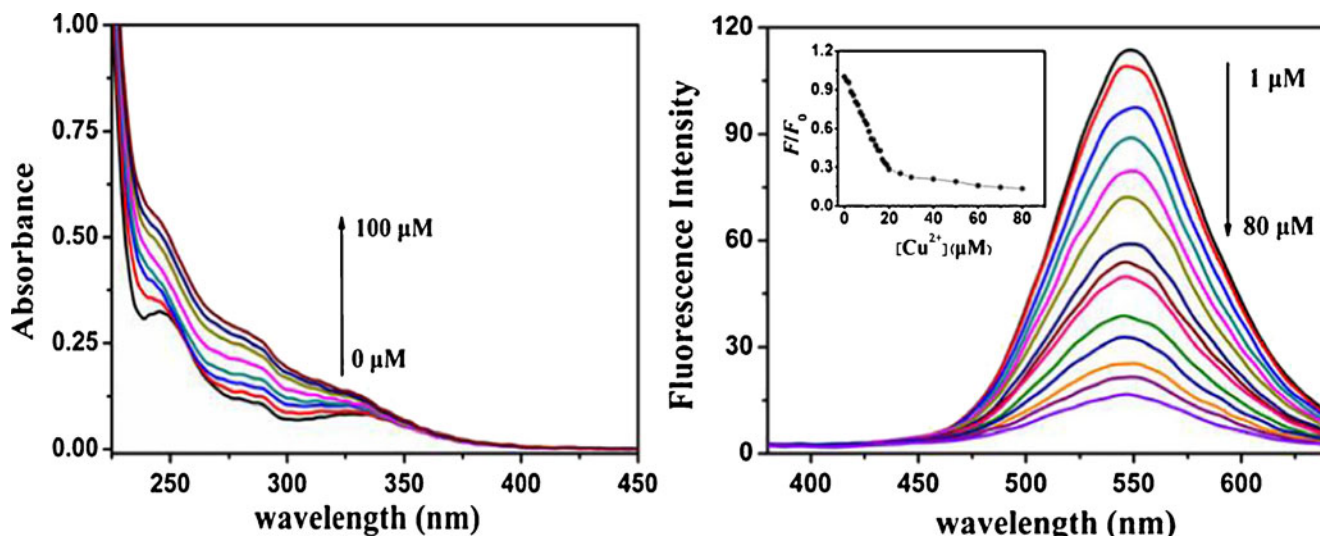


Fig. 7 (left) Variations of absorbance spectra of **1** (20 μM) in buffer solution upon addition of Cu²⁺ from 2.0 to 100 μM. (right) Fluorescence titration spectra of **1** (20 μM) in buffer solution with

Cu²⁺. The fluorescence emission intensity at 550 nm decreased upon gradually addition of Cu²⁺ from 1.0 to 80 μM (λ_{ex}=330 nm)

supply a potential application of **1**/Cu²⁺ complex in protein detection.

Design of Logic Gate

The larger binding constant of **1**/Cu²⁺ (1.58 × 10⁵ M⁻¹) vs. that of **1**/Hg²⁺ (1.18 × 10³ M⁻¹) made it possible to replace

Hg²⁺ in the **1**/Hg²⁺ complex with Cu²⁺, but not the reverse. That is, when Cu²⁺ was added to a solution that contained **1** and Hg²⁺, we observed obvious fluorescence emission quenching (Fig. S9†). This indicated that **1** could be acted as a potential logic gate, or molecular switch, in the detection of metal ions. We utilized our system as a NOR logic gate with a YES logic function (Fig. 9). The truth table values for output 1 formed a NOR logic gate, and the truth table values for output 2 formed a YES logic gate. The combination of these intrinsic properties with the selective actions of different chemical inputs allows this system to be implemented as a complex ionic switch.

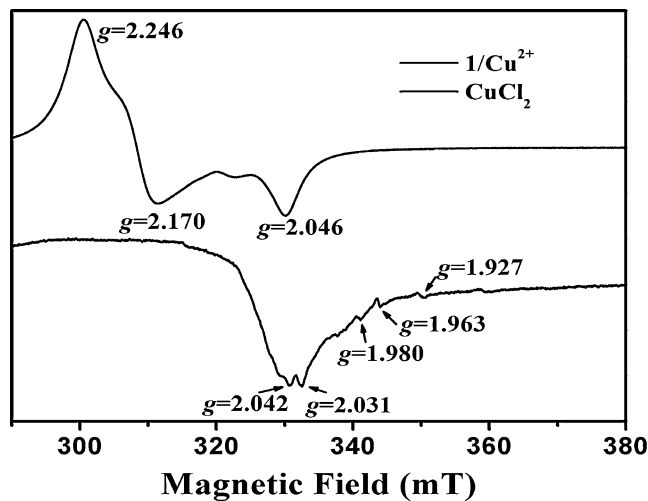


Fig. 8 Electron paramagnetic resonance (EPR) spectra indicated a strong interaction between CuCl₂ and probe **1**. A representative region of the X-band EPR spectra of CuCl₂ and its complex with **1** equiv. of **1** in the solid state. In the presence of **1**, the peaks at 300 and 325 mT disappeared, and a multitude of obvious spectral features (hyperfine splittings of the ¹⁴N nuclear spins of the peptide with the unpaired electron spin of Cu²⁺) appeared at 325–350 mT, proving that a complex formed between Cu²⁺ and **1**

Absorption Spectrum and Binding Model of DGT/Pb²⁺

The addition of Pb²⁺ to a solution that contained **1** produced a precipitate; this was further confirmed by measuring the solution turbidity while stirring (A₅₀₀, Fig. S10†). The solution turbidity increased gradually with the addition of small amounts of Pb²⁺ between 3.0 and 90 μM, which indicated that an insoluble substance was generated

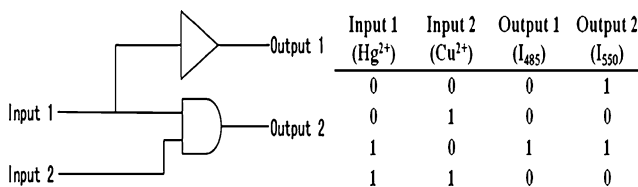


Fig. 9 (left) Combinatorial logic circuit of a dual output molecular switch; Truth table for logic gate; (right) Hg²⁺ and Cu²⁺ are inputs to the system; I₄₈₅ and I₅₅₀ are the output signals

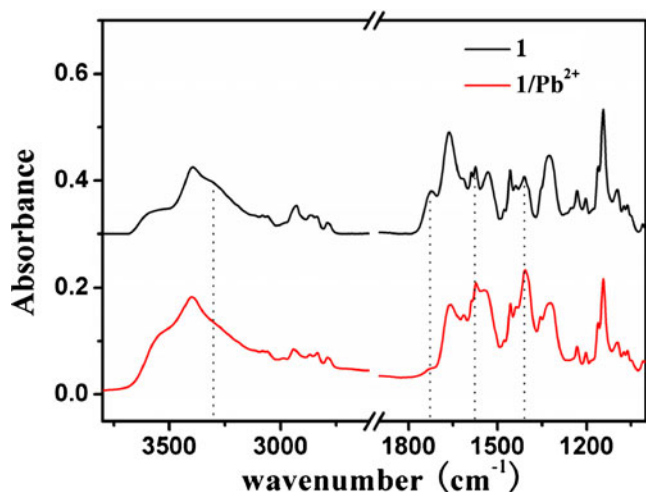


Fig. 10 Typical FT-IR spectra region of **1** in solid and that of the precipitate of $1/\text{Pb}^{2+}$

gradually. When the concentration of Pb^{2+} exceeded $120\ \mu\text{M}$, the turbidity increased dramatically to the extent that the precipitate could be observed with the naked eye. To explore the structural origin of the precipitate, it was filtrated and analyzed by using FT-IR spectroscopy and the spectra were recorded (Fig. 10). The strong peak at $1,724\ \text{cm}^{-1}$ was assigned to a carboxyl ($\text{C}=\text{O}$) stretching model, which disappeared with the formation of a $1/\text{Pb}^{2+}$ complex. Two strong peaks at $1,573$ and $1,409\ \text{cm}^{-1}$ were assigned to antisymmetric and symmetric $\text{C}=\text{O}$ stretching models of the carboxylate. It is reasonable to propose that the carboxyl of **1** bound directly to Pb^{2+} , and no other functional group interacted with Pb^{2+} . Furthermore, based on previous reports [24–29], the large difference between $\nu_{\text{COO}^{\text{as-}}}$ and $\nu_{\text{COO}^{\text{s-}}}$ ($\Delta\nu=164\ \text{cm}^{-1}$) was ascribed to a bidentate chelation model between the carboxylate and Pb^{2+} . In addition, the response of the control compound

8 was not similar to that of **1** to Pb^{2+} , which confirmed the direct binding between Pb^{2+} and the carboxyl in **1**.

Comparison DGT with DGGT

We also investigated the tri-peptide probe, **2**, under the same conditions that we used for **1** to explore whether including another Gly in the probe would affect the recognition of metal ions. Probe **2** showed similar, but slightly different behaviors from **1** toward metal ions (Fig. S11–13†). The direct connection between the dansyl moiety and the tryptophan produced a probe, dansyl-L-Trp, which showed a modest blue-shift and quenching with the addition of Hg^{2+} , but no response with other metal ions [10]. Thus, the incorporation of either one or two Gly residues between the dansyl moiety and the Trp resulted in two tetra-functional dipeptide probes, **1** and **2**, which displayed distinct optical responses to Hg^{2+} , Cu^{2+} , Pb^{2+} , and H^+ in aqueous solution. The different responses of **1**, **2**, and dansyl-L-Trp to Hg^{2+} reflected the different binding modes between them; thus, a peptide can provide more diversity in the binding site than a single amino acid. Despite the slight differences in fluorescence responses, both **1** and **2** interacted with Hg^{2+} , which suggested that a sulfonamide was necessary for the recognition of Hg^{2+} . This conclusion was further supported by the control molecules, **8** and **11**, which both had esterified moieties and also showed blue-shift or fluorescence enhancement in response to Hg^{2+} (Fig. S5†).

Cell Permeability of DGT and DGGT

Finally, we investigated the cell-permeability of **1** and **2** (Fig. 11). Cultured HEK-293 cells were incubated with $10\ \mu\text{M}$ of **1** or **2** for 1 h at room temperature. The overlay of bright-field transmission and fluorescence images

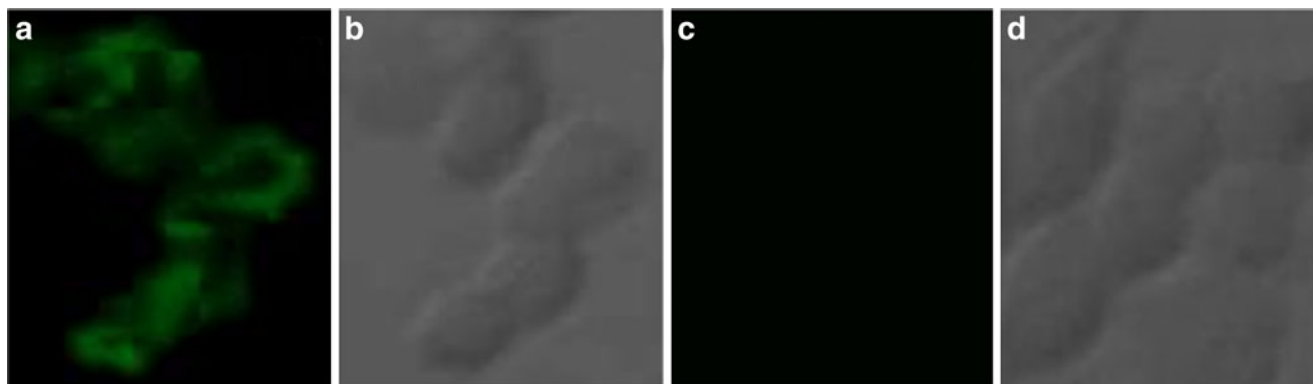


Fig. 11 Cell permeability of probes **1** and **2**. Confocal laser microscope images (**a**, **c**) and bright field images (**b**, **d**) of HEK-293 cells after incubating in the presence of **1** (**a**, **b**) and **2** (**c**, **d**). The

difference in fluorescence brightness between **1** and **2** demonstrated that **1** showed more cell-permeability than **2**

revealed that **1** was localized in intracellular region, which demonstrated that **1** penetrated the cell membrane and accumulated inside the cell. Then we tried the fluorescence response of **1** to Hg^{2+} inside cell. However, as the inclusion of dansyl moiety of **1** in the hydrophobic cavity of BSA existed both in the cell and the culture medium induced the emission of **1** blue-shifted to the wavelength exactly as same as that of 1/Hg^{2+} , the detection of Hg^{2+} in cell was interfered strongly [9]. In addition, the toxicity of Hg^{2+} resulted in the cell readily to die for the long time incubation with Hg^{2+} . Therefore, we did not succeed. In contrast to **1**, **2** could not be detected in the intracellular region under the same conditions, which suggested that **2** was less cell permeable than **1**. This may due to the fact that **2** harbored an extra hydrophilic amino acid, glycine.

Conclusion

In summary, we synthesized a tetra-functional probe based on a dipeptide motif and a dansyl moiety, which responded to Hg^{2+} and Cu^{2+} in distinct modes and distinguishing optical properties. In addition, **1** was sensitive to pH ranging from 2.0 to 5.0 and precipitated in the presence of Pb^{2+} in neutral conditions. The concepts presented here will contribute to the future development of miniaturized molecular probes with multi-functions.

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