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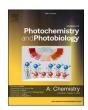
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A copper complex receptor for nanomolar sulfide sensing and applications in DNA/BSA binding

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ABSTRACT

Traditionally toxic sulfide species (S^{2-} , HS^- , and H_2S) play indispensable roles in various biological, environmental, and industrial processes. However, excessive exposure to these species can result in numerous chronic diseases. Considering the importance of sulfide recognition, we have synthesized and characterized a simple 1:1 Cu(II) complex of **L** (CuCl₂**L** receptor) based on a substituted 2,6-dipyrazinylpyridine (dppy) moiety (L = 4-(2,6-di(pyrazin-2-yl)pyridin-4-yl)-N,N-dimethylaniline). The complex was analyzed both experimentally and computationally using DFT and TD-DFT calculations. The*in-situ*prepared CuCl₂L receptor is employed to selectively recognize sulfide species (detection limit: 82.50 nM) in the presence of other anions in higher concentration (ten equivalents of F⁻, Cl⁻, Br⁻, Γ, CH₃CO₂, HCO₃, H₂PO₄, NO₃, SO₃², SO₄², SO₃², CO₃², and HPO₄² (10 mM)) in aqueous acetonitrile (2:1 v/v, 10 mM HEPES buffer, pH 7.4) through absorption spectral studies. Furthermore, the receptor is successfully utilized for various real-time detection applications and demonstrated significant intercalative binding interaction with Calf–Thymus DNA (CT-DNA) and good binding interactions with bovine serum albumin (BSA), supported by viscometry and molecular docking studies.

1. Introduction

Adequate, effective receptors with optical responses have gathered immense attention for instant anion sensing, as these species play indispensable roles in various environmental, biological, and chemical processes [1–5]. Among them, sulfide species have drawn particular concern due to their widespread dispersion in the environment as a traditional toxic and hazardous pollutant. Sulfide accumulates in water, including domestic wastewater [6,7], from various sources such as crude petroleum, volcanic eruptions, and the reduction of sulfate by anaerobic bacteria [8,9]. It also frequently emerges as a by-product of many industrial processes [10], while offering diverse applications across sectors [11] such as cosmetics [12], dyes [13], sulfuric acid, and wood pulp manusfacturing [14].

Once hydrolyzed in water, sulfide is primarily converted into hydrogen sulfide (H_2S) or hydrosulfide (HS^-) [6,7], as aqueous S^2 – does not exist under typical conditions. This has been confirmed by recent experiments involving Na_2S dissolved in hyper-concentrated NaOH(aq)

and CsOH(aq) [15], and further supported by molecular dynamics simulations [16]. The implications of this hydrolysis are profound, as H_2S plays crucial physiological roles [17], including modulation of blood pressure [18], brain neuromodulation [19], regulation of inflammation [20], and suppression of oxidative stress [21]. Furthermore, H_2S has been shown to protect against ischemia/reperfusion injury [22,23], to act as an antioxidant [24], and to scavenge reactive oxygen species [25]. Alterations in cellular H_2S levels have been linked to various diseases, such as Down's syndrome [26], Alzheimer's disease [27], diabetes [22], and liver cirrhosis [28].

Exposure to excessive concentration of sulfide can cause several metabolic and physiological issues, ranging from irritation in mucous membranes [29] and unconsciousness [30] to respiratory paralysis [31]. While $\rm H_2S$ is essential in low concentrations, its toxicity increases at higher levels, leading to permanent brain damage [32–34] and even death through asphyxiation [35]. Given its dual role as both a necessary biological molecule and a toxic hazard, the detection of sulfide species is of critical importance across biological, environmental, and industrial

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fields.

In recent years, the metal displacement approach utilizing copper receptors for sulfide detection has attracted extraordinary attention [36]. This method is based on the principle that sulfide can displace Cu (II) ions from a copper receptor. This displacement results in the formation of a highly stable CuS compound, characterized by its remarkably low solubility product constant $(K_{sp} = 6.3 \times 10^{-36})$ [37]. The formation of this stable moiety is a key indicator in the successful detection of sulfide [8,12,38]. Furthermore, copper complexes possess less toxic side effects than those based on exogenous metals [39]. In addition to their role in sulfide detection, they also exhibit potential antioxidant, antibacterial, and antitumor/anti-cancer activities [40-42]. Among the various receptor systems currently available for sulfide recognition, several encounter specific limitations, such as multistep synthetic procedure requiring stringent synthetic conditions [43–46], notably low solubility in aqueous media [12,47,48], potential interference from other S-containing moieties [49], among others. These challenges underscore the need for more streamlined and selective approaches in the development of sulfide detection methodologies.

Nitrogen-based heterocyclic ligands such as 2,2':6',2''-terpyridine, 2,6-dipyrazinylpyridine (dppy), and their transition metal complexes have received considerable interest for their integration into a diverse array of applications. These range from halogen bonding [50,51] and protein interaction to drug-displacement studies [52], hydrogelation [53,54], self-assembly [55], anions sensing [56,57], and amino acids sensing [58], to name a few. The growing popularity of these compounds can be attributed to their straightforward synthesis, high stability, and their exceptional electrochemical and photophysical properties [59-61]. Considering practicality and convenience, various detection techniques have been utilized to recognize sulfide species [33,62-65]. Among these, colorimetric and spectroscopic methods, particularly UV-Visible (UV-Vis) spectroscopy, have attracted the most attention. Their popularity stems from several key advantages: high accuracy, cost-effectiveness, rapid response, swift processing, operational simplicity, and the ability for on-site detection [66-68].

Herein, we have prepared a simple Cu-dppy receptor (dppy: 2,6dipyrazinylpyridine), specifically a Cu(II) complex (CuCl₂L) of the 4-(2,6-di(pyrazin-2-yl)pyridin-4-yl)-N,N-dimethylaniline ligand Similar to other dppy-based ligands, L demonstrates strong binding interactions with various 3d transition metal ions, including Cu(II) and Zn (II). We selected CuCl₂L for sulfide ion detection due to the pronounced interactions between Cu(II) and sulfide ions (vide supra). The CuCl₂L receptor was characterized using UV-Vis, Fourier-transform infrared (FT-IR), and electron spin resonance (ESR) spectroscopies, along with high-resolution mass spectrometry (HRMS). The in-situ synthesized CuCl₂L receptor demonstrates exceptional selectivity in detecting sulfide species at nanomolar concentrations, with a detection limit of 82.50 nM. This was achieved even in the presence of various other anions at higher concentrations (ten equivalents) such as F⁻, Cl⁻, Br⁻, Γ, CH₃CO₂, HCO_3^- , $H_2PO_4^-$, NO_3^- , SO_3^{2-} , SO_4^{2-} , $S_2O_3^{2-}$, CO_3^{2-} , and HPO_4^{2-} , in an aqueous acetonitrile environment (2:1 v/v, 10 mM HEPES) maintained at a physiological pH of 7.4. Additionally, this in-situ prepared CuCl₂L receptor is effectively utilized in various real-time sulfide detection applications. Particularly, it is employed to study Calf-Thymus DNA (CT-DNA) and bovine serum albumin (BSA) interactions with the help of absorption, emission, and molecular docking study.

2. Experimental section

2.1. Materials and methods

Spectroscopic and analytical grade chemicals used in the synthesis and spectral analyses were obtained commercially. ¹H NMR, ESI-HRMS, ESR, and FT-IR spectra were acquired using the JEOL ECZ500R/S1 NMR spectrometer, the Impact HD UHR-TOF mass spectrometer, the JES-FA200 (JEOL) ESR spectrometer, and the 3000 Hyperion FT-IR

spectrometer (Bruker, Germany), respectively. Absorption and emission spectra were recorded using the Mortas Scientific UV plus MSGUI3.1.0 absorption spectrophotometer and the HITACHI-F4600 Fluorescence spectrophotometer, respectively.

2.2. Synthesis of the Cu(II) complex of L (CuCl₂L)

The 4-(2,6-di(pyrazin-2-yl)pyridin-4-yl)-N,N-dimethylaniline ligand L was synthesized following an available synthetic procedure [69] (see the Supplementary Information, SI, for the synthetic procedure and synthetic scheme of L). An ethanolic solution of $CuCl_2$ (26.8 g, 0.2 mmol, 10 mL) was added to a dichloromethane solution of the synthesized L (0.148 g, 0.4 mmol, 10 mL), and stirring the reaction mixture at room temperature for 2 h yielded a dark red-colored precipitate. The precipitate was collected and washed with ethanol (30 mL) several times, resulting in 1:1 Cu(II) complex of L ($CuCl_2L$). Yield: 85%. ESI-MS: $[CuClL]^+$ m/z 452.0571 (found); 452.0577 (calculated). Elemental analysis ($C_{21}H_{18}Cl_2CuN_6$ g mol^{-1}): C 51.60; H 3.71; N 17.19 (calculated); C 51.41; H 3.55; N 17.02 (found). FT-IR (v_{max}/cm^{-1}): 3047, 3002, 2820, 1580, 1462, 1399, 1357, 1232, 1170, 1075, 834 (see Figs. S1–S6 in the SI for 1H NMR, HRMS, FT-IR, and EPR spectral plots of the ligand L and $CuCl_2L$).

2.3. Preparation of experimental solutions

2.3.1. Uv-vis absorption spectral studies

An acetonitrile solution of L (115 μ M, 25 mL) was prepared, along with aqueous solutions of various anions. These included sodium salts of S²⁻ (1 mM Na₂S, 50 mL), F⁻, Cl⁻, Br⁻, I⁻, CH₃CO₂, HCO₃, H₂PO₄, NO₃, SO₃²⁻, SO₄²⁻, S₂O₃³⁻, CO₃²⁻, and HPO₄²⁻ (10 mM, 50 mL). The **CuCl₂L** receptor (38 μ M, 10 mL) solution was prepared *in-situ* by mixing an aqueous CuCl₂ solution (1 mM, 1.15 mL) and an acetonitrile solution of L (115 μ M, 10 mL), followed by dilution with an aqueous HEPES buffer (10 mM, 20 mL, pH 7.4).

2.3.2. DNA binding studies

A stock solution of CT-DNA was prepared by dissolving it in aqueous HEPES buffer (10 mM, pH 7.4) and allowing it to stand overnight at 4 °C. This stock solution exhibited an absorption band intensity ratio of 1.90:1 (for absorption bands 260 nm and 280 nm), indicating it was sufficiently free from protein. The concentration of CT-DNA was determined from the intensity of the 260 nm absorbance band and extinction coefficient $\varepsilon_{260} = 6600~\text{M}^{-1}~\text{cm}^{-1}~\text{[70]}$. Absorption spectral titration was carried out by gradually adding a CT-DNA (0–260 μ M) solution to an *in-situ* prepared CuCl₂L receptor (38 μ M, 1.5 mL) in aqueous HEPES buffer (10 mM, pH 7.4). The CuCl₂L (38 μ M, 1.5 mL) receptor was prepared as in section 2.3.1. The binding constant (K_b) [71] was determined from the absorption spectral titration using the Wolfe-Shimmer Equation (1) [72]:

$$[\text{CT-DNA}]/(\varepsilon_a - \varepsilon_f) = [\text{CT-DNA}]/(\varepsilon_b - \varepsilon_f) + 1/K_b(\varepsilon_b - \varepsilon_f)$$
(1)

where [CT-DNA] is the CT-DNA concentration, ε_a is the absorption intensity/[CuCl₂L] (with [CuCl₂L] being the CuCl₂L receptor concentration), ε_f is the extinction coefficient of free CuCl₂L, and ε_b is the extinction coefficient of the fully bound CuCl₂L to DNA.

Ethidium bromide (EB), a standard DNA intercalating reagent, was used for the CT-DNA emission spectral titration study. In this experiment, an *in-situ* prepared CuCl_2L (100 μM , 50 μL) receptor was gradually added to an equimolar mixture of CT-DNA and EB (45 μM , 2 mL). The fluorescence quenching constant (K_{SV}) [73] and the binding constant (K_{app}) [74] were determined from the emission spectral titration using the Stern–Volmer equation (2) and equation (3), respectively:

$$\frac{F_0}{F} = 1 + K_{SV}[\mathbf{CuCl_2L}] \tag{2}$$

$$K_{app} \times [\mathbf{CuCl_2L}]_{1/2} = K_{EB} \times [\mathbf{EB}]$$
 (3)

where F_0 is the fluorescence intensity in the absence of $\mathbf{CuCl_2L}$ receptor, F is the fluorescence intensity in the presence of $\mathbf{CuCl_2L}$ receptor, K_{SV} is the linear Stern–Volmer constant, $[\mathbf{CuCl_2L}]_{1/2}$ is the $\mathbf{CuCl_2L}$ receptor concentration at 50 % reduction of fluorescence intensity, K_{EB} is 10 \times 10⁶ M⁻¹, and [EB] is 40 μ M.

2.3.3. BSA binding study

A BSA stock solution was prepared for a binding interaction study with the $CuCl_2L$ receptor. For absorption and emission spectral studies, the $CuCl_2L$ receptor (90 μ M, 20 μ L) was gradually added to BSA (2.5 μ M, 2 mL) in phosphate buffer (100 mM, pH 7.4). The quenching constant (K_{SV}) [75] and affinity constant (K) [76] were calculated by using equation (2) and the Scatchard equation (4), respectively:

$$\log(F_0 - F)/F = \log K + n\log[\mathbf{CuCl_2L}] \tag{4}$$

where K is the affinity constant of the $CuCl_2L$ receptor with BSA, and n is the number of binding sites per albumin or binding stoichiometry.

2.3.4. Viscometry titration study

A stock solution of CuCl_2L receptor (100 μM , 0.7 mL) was gradually added to a CT-DNA solution (25 μM , 28 mL), and a viscometry titration study was carried out using an Ostwald viscometer. The average flow time of the solutions was measured using a digital stopwatch. The relative viscosity values were determined using the expression $\eta = (t-t_0)/t_0$, where t and t_0 represent the observed flow time of the CT-DNA and buffer solution, respectively. These values were presented as $(\eta/\eta_0)^{1/3}$ vs. [CuCl₂L]/[CT-DNA] binding ratio, where η denotes the viscosity of CT-DNA in the presence of CuCl₂L, and η_0 represents the viscosity of CT-DNA alone [71,77].

2.4. Molecular docking study

The Autodock 4.2 software [78] was employed for docking studies, wherein the ground state energy-optimized structure of the $CuCl_2L$ receptor was docked with the crystal structures of CT-DNA (PDB ID: 1BNA) and BSA (PDB ID: 4F5S). The following line was added to the parameters files to include the copper atom parameters in AutoDock 4.2: "atom_par Cu 3.50 0.005 12.000—0.00110 0.0 0.0 0—1 -1 4 # Non H-bonding". The docking results were then visualized and analyzed using the Discovery Studio 3.5 software [79].

2.5. DFT calculations

Quantum chemical analyses for our study were performed using the Gaussian 16 software, Revision C.01 [80]. Building on insights from our previous work [81] with another dppy-based Cu(II) complex, we optimized geometries using the PBEO [82,83] functional, integrating the D3 (BJ) [84,85] approach for dispersion corrections. The def2-TZVP basis set was employed on the Cu atom, while the def2-SVP basis set was applied to all the others [86]. This selection is labelled as bs1, making the theory level PBEO-D3(BJ)/bs1. The unrestricted Kohn-Sham method was used for open-shell systems. Gibbs free energy corrections, obtained in calculations at the PBE0-D3(BJ)/bs1 level, were applied to singlepoint energy computations using the PBE0-D3(BJ) functional but with an expanded basis set. Here, def2-TZVP was used for all elements except Cu, which was treated using the def2-QZVP basis set, a combination referred to as bs2. Solvation effects were incorporated via the solvent model based on density (SMD) [87], considering a 2:1 water/acetonitrile solvent mixture. Consequently, free energy calculations were conducted at the SMD/PBE0-D3(BJ)/bs2 level of theory. To ensure accuracy in associative and dissociative steps, a concentration correction of $\Delta G^{0\to *} = 1.89$ kcal/mol was applied to the free energy values of all computed species, compensating for standard state variations from gas phase (1 atm) to condensed phase (1 M) [88–90]. All optimized geometries were verified as minima on their respective potential energy surfaces, confirmed by vibrational frequency calculations that showed only positive eigenvalues in the Hessian matrices. Finally, time-dependent DFT (TD-DFT) calculations were performed using the SMD/PBE0-D3(BJ)/bs1 level of theory. A total of 50 excited electronic states were considered in the calculations. The nature of key electronic transitions was examined through the charge density difference (CDD) analysis and the interfragment charge transfer (IFCT) method, as implemented in Multiwfn 3.8 [91]. Further insights into the excited state properties were obtained using the TheoDORE program [92]. A Gaussian envelope was used to shape the vertical excitations in all TD-DFT spectra, as described by the following equation:

$$\epsilon(\lambda) = \sum_{i} \epsilon_{i}^{max} \exp\left[\left(\frac{1/\lambda - 1/\lambda_{i}}{1/\sigma}\right)^{2}\right]$$
(5)

where

$$\epsilon_i^{max} = \frac{\sqrt{\pi} \times e \times N}{1000 \times \ln 10 \times c \times m_e} \times \frac{f_i}{10^7/\sigma} \tag{6}$$

In Equation (5), $\epsilon(\lambda)$ is the molar absorptivity, which depends on the wavelength, and is expressed in units of L mol $^{-1}$ cm $^{-1}$. The variables λ and λ_i refer to the wavelength (independent variable) and wavelength of the i-th transition (the center of the Gaussian), respectively, with both measured in nm. The Gaussian standard deviation, σ , is also expressed in nm, ensuring that the exponent remains dimensionless. Equation (6) rescales the oscillator strength f_i (dimensionless) for each i-th electronic transition into molar absorptivity. The factor of 10^7 is included to convert $1/\sigma$ to cm $^{-1}$, while the factor of 1000 converts cm 3 to L. In this formula, e represents the elementary charge (in esu), N is Avogadro's constant (in mol $^{-1}$), e is the speed of light (in cm s $^{-1}$), and m_e is the electron mass (in g). The value of $\sigma=6200$ nm, which corresponds to 0.2 eV, was used in the calculations.

3. Results and discussion

3.1. Synthesis and sulfide sensing analysis of CuCl₂L

The ligand L, upon reaction with CuCl2 in a dichloromethane-ethanol solvent mixture, successfully yielded the CuCl2L complex at room temperature, as outlined in Scheme 1. The complex was characterized using a range of techniques including ¹H NMR, HRMS, elemental analysis, ESR, and FT-IR (see Figs. S1-S7 in the SI). This CuCl₂L complex is expected to have a distorted square-pyramidal geometry. In this structure, the Cu(II) ion is penta-coordinated, involving three nitrogen atoms from the substituted dppy moiety and two chloride anions. Our DFT calculations confirm this geometric configuration (see Scheme 1), revealing that the optimized structure of CuCl2L exhibits a structural parameter τ value [93] of 0.098. A τ value of 0 indicates a perfect square-pyramidal geometry, while a value of 1 corresponds to a perfect trigonal bipyramid. Additionally, the ESR spectrum of the complex, displaying a $g|| > g_{\perp} > 2.002$ value [34], further corroborates this geometry (see Fig. S7 in the SI). Similar distorted square-pyramidal geometrical arrangements have been observed in other Cu(II)-dppy complexes [94-96].

The UV–Vis spectra of L (38 μ M, 1.5 mL) in aqueous acetonitrile (2:1 v/v, 10 mM HEPES buffer, pH 7.4) before and after the addition of CuCl₂ is depicted in Fig. 1a. L exhibits two prominent absorption bands at 233 nm and 295 nm, accompanied by a shoulder at 364 nm. The computed UV–Vis spectrum of L obtained from TD-DFT calculations (see Fig. 1b and additional Figs. S9–S10 and Tables S1 and S2 in the SI) shows excellent agreement with the experimental spectrum, with the three main electronic excitations occurring at 235 nm, 282 nm, and 378 nm. The first excitation is attributed to a mixed $\pi \to \pi^*$ and $n \to \pi^*$ transition

Scheme 1. Preparation of CuCl₂L, the Cu(II) complex of L, and its optimized structure.

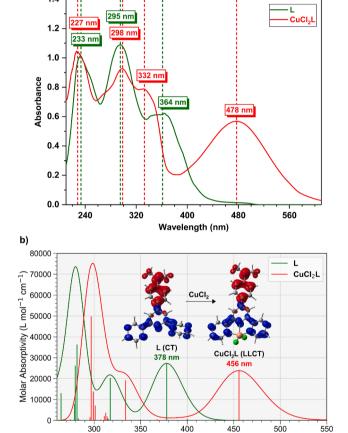


Fig. 1. (a) Absorption spectra of L (38 μ M, 1.5 mL) and the *in-situ* prepared CuCl₂L receptor (38 μ M, 1.5 mL) in aqueous acetonitrile (2:1 v/v, 10 mM HEPES, pH 7.4). (b) Computed TD-DFT absorption spectra of L and CuCl₂L at the SMD/PBE0-D3(BJ)/bs1 level. The insets show the charge density difference (CDD) plots of the charge transfer (CT) excitation of L at 378 nm and the corresponding ligand-to-ligand charge transfer (LLCT) excitation of CuCl₂L at 456 nm. In the CDD plots, charge flows from red to blue. Isovalues: 0.002 a.u.

Wavelength (nm)

with local excitation (LE) character, primarily involving the pyrazole and pyridine rings. The latter two excitations are associated with charge transfer (CT) transitions from the aniline ring to the nitrogen-bearing rings.

Upon the addition of aqueous $CuCl_2$ to L, the *in-situ* prepared $CuCl_2L$ complex exhibited two prominent absorption bands at 227 nm and 298 nm, with a shoulder at 332 nm and a new absorption band emerging at 478 nm. TD-DFT calculations reveal that the band at 478 nm (computed

at 456 nm) shares the same characteristics as the 364 nm band of the free ligand **L** but is redshifted due to metal coordination. The IFCT analysis indicates that this transition possesses an 86% CT character, with 89% of the hole density localized on the aniline group and 10% on the pyrazole/pyridine rings. The electron density is predominantly localized on the nitrogen-bearing rings (91%), with only 3.5% at the CuCl $_2$ center. A similar distribution is observed in the electron-hole correlation plots (see Figs. S9–S10 in the SI). Thus, the transition observed at 478 nm is identified as a ligand-to-ligand charge transfer (LLCT), rather than metal-to-ligand charge transfer (MLCT). The presence of this transition in the experimental spectrum confirms the successful formation of the *in-situ* CuCl $_2$ L complex in aqueous acetonitrile. Moreover, DFT calculations further corroborate the favorable formation of the CuCl $_2$ L complex from L and CuCl $_2$ in aqueous acetonitrile, with a computed free energy change (ΔG) of -32.5 kcal mol $^{-1}$.

To evaluate the selectivity of the CuCl_2L receptor for anion sensing, separate aqueous solutions of various anions were tested. These included $S^{2^-}(\text{Na}_2\text{S}\ 1\ \text{mM},\ 40\ \mu\text{L})$ and a range of other anions such as F^- , Cl $^-$, Br $^-$, Γ , CH $_3\text{CO}_2^-$, HCO $_3^-$, H2PO $_4^-$, NO $_3^-$, SO $_3^{2^-}$, SO $_4^{2^-}$, S2O $_3^{2^-}$, and HPO $_4^{2^-}$ (each at 10 mM, 57 μL concentration). Each anion solution was individually added to a solution of the CuCl_2L receptor (38 $\mu\text{M},\ 1.5\ \text{mL})$ prepared in aqueous acetonitrile (2:1 v/v, 10 mM HEPES, pH 7.4). Following the addition of these anions, the absorption spectra were recorded to assess the receptor's response and its ability to selectively detect each anion under these conditions.

Upon adding the anions (ten equivalents) specified earlier (excluding Na_2S), there was no significant alteration observed in the absorption spectrum of the $CuCl_2L$ receptor, as depicted in Fig. 2. Contrarily, the introduction of just one equivalent of Na_2S to the *in-situ* prepared $CuCl_2L$ receptor led to a notable change: the near disappearance of the LMCT absorption band at 478 nm. Subsequently, the absorption spectrum of the $CuCl_2L$ receptor shifted to closely resemble that of the free ligand L, also shown in Fig. 2.

The observed absorption spectral behavior indicates that the addition of Na₂S effectively facilitates the decomplexation of the Cu(II) moiety from the CuCl₂L receptor. This decomplexation occurs due to the strong binding interaction between the sulfide ions and the Cu(II) ion, leading to the displacement of Cu(II) from the complex. Consequently, L is released in the solution, while the sulfide precipitates as CuS given its very low solubility product constant. This mechanism highlights the selective sensing of sulfide by the *in-situ* prepared CuCl₂L receptor in an aqueous acetonitrile medium, demonstrating its potential as a practical sensor for this specific species.

In order to investigate the $CuCl_2L$ receptor's interaction dynamics and sensitivity to sulfide species, an absorption spectral titration study was performed. This involved the incremental addition of S^{2-} (Na₂S 0.1 mM, 4 μ L) to the *in-situ* prepared $CuCl_2L$ receptor solution (38 μ M, 1.7 mL) in aqueous acetonitrile (2:1 v/v, 10 mM HEPES, pH 7.4), as illustrated in Fig. 3. Each addition of the anion induced noticeable hypochromic shifts in the 298 nm absorption band, including its shoulder at

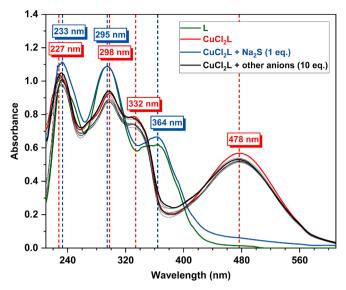


Fig. 2. Absorption spectra of the *in-situ* prepared **CuCl₂L** receptor (38 μM, 1.5 mL) upon addition of one equivalent S^2 (Na₂S 1 mM, 57 μL) and ten equivalent different anions (F⁻, Cl⁻, Br⁻, Γ⁻, CH₃CO₂⁻, HCO₃⁻, H₂PO₄⁻, NO₃⁻, SO₃⁻⁻, SO₄⁻, S₂O₃⁻, CO₃⁻, and HPO₄⁻) (10 mM, 57 μL) in aqueous acetonitrile (2:1 v/v, 10 mM HEPES, pH 7.4).

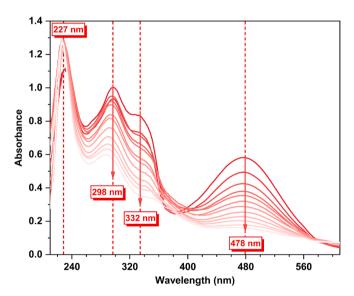


Fig. 3. Absorption spectral titration of *in-situ* prepared **CuCl₂L** receptor (38 μ M, 1.7 mL) upon gradual addition of S²⁻ (Na₂S 0.1 mM, 4 μ L) in aqueous acetonitrile (2:1 v/v, 10 mM HEPES, pH 7.4). Arrows indicate the effect of the gradual addition of sulfide on the absorption spectrum.

332 nm, and notably in the 478 nm band, which almost vanished upon reaching saturation with the added aqueous sulfide species (totaling up to 40 μ L, 0.1 mM).

This experiment revealed that the addition of nearly 0.1 equivalent of Na_2S was sufficient to displace the Cu(II) ion from the $CuCl_2L$ receptor. The receptor's absorption spectrum showed significant sensitivity to even minor variations in sulfide concentration, indicating its high efficacy in detecting the presence of sulfide species. Through this analysis, the detection limit [97,98] of the $CuCl_2L$ receptor for sulfide was determined as 82.50 nM, underscoring its potential as a sensitive tool for sulfide detection.

The selective sensing capability of the $CuCl_2L$ receptor for sulfide species was further validated through a competition experiment. In this experiment, aqueous solutions of various anions, each at a higher

concentration (ten equivalents) including F⁻, Cl⁻, Br⁻, I⁻, CH₃CO₂, HCO₃, H₂PO₄, NO₃, SO₃²-, SO₄²-, S₂O₃²-, CO₃²-, and HPO₄²- (10 mM, 57 μ L), were individually added to the CuCl_2L receptor solution (38 μ M, 1.5 mL). Subsequently, one equivalent of S²⁻ (Na₂S 1 mM, 57 μ L) was introduced into each mixture.

Initially, the presence of ten equivalents of other anions (excluding Na_2S) did not significantly impact the absorption spectrum of the $CuCl_2L$ receptor. However, upon the addition of one equivalent of Na_2S , there was a notable alteration in the receptor's absorption spectrum. This was marked by the near disappearance of the 478 nm absorption band, leading the spectrum to resemble that of the free ligand L, as shown in Fig. 4.

Furthermore, when a mixture containing ten equivalents of all the aforementioned anions (except Na₂S) was added to the CuCl₂L receptor, followed by the introduction of one equivalent of the sulfide anion, similar results were observed. This outcome from the competition experiments decisively confirms the receptor's ability to selectively recognize sulfide, even in the presence of a mixture of other anions at concentrations ten times higher.

The conditions for sulfide detection established in our study not only align well with those reported in existing literature but also demonstrate a notably improved detection limit and response time compared to the majority of previously documented receptor systems. These enhancements are clearly evidenced in our comprehensive comparison, as summarized in Table 1, emphasizing the advanced efficiency and sensitivity of our receptor system in sulfide sensing applications.

3.2. Real-time applications of sulfide detection

3.2.1. Effect of pH variation and water tolerance on the in-situ prepared $CuCl_2L$ receptor for sulfide detection

We conducted a thorough analysis of the in-situ prepared CuCl_2L receptor's response to pH variations in relation to sulfide detection. Utilizing UV–Vis absorption spectroscopy, the receptor's behavior was monitored across a pH range of 1 to 12 in an aqueous acetonitrile solution (2:1 v/v, 10 mM HEPES). The findings reveal that the CuCl_2L receptor maintains stability within the pH range of 4 to 9 and is, therefore, most effective for sulfide detection under these conditions (see Fig. 5). Outside this range, the receptor undergoes hydrolysis, regenerating the free ligand L, which compromises its functionality.

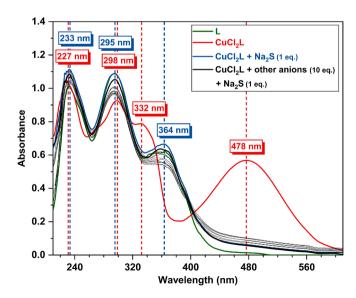


Fig. 4. Absorption spectra of *in-situ* prepared CuCl₂L receptor (38 μM, 1.5 mL) upon addition of different anions (F^- , CI^- , Br^- , I^- , $CH_3CO_2^-$, HCO_3^- , $H_2PO_4^-$, NO_3^- , SO_3^{2-} , SO_3^{2-} , SO_3^{2-} , CO_3^{2-} , and HPO_4^{2-}) (10 mM, 57 μL) followed by S^{2-} (Na₂S 1 mM, 57 μL) addition in aqueous acetonitrile (2:1 v/v, 10 mM HEPES, pH 7.4).

Table 1Important analytical parameters of a few reported receptors for sulfide sensing.

Probe	Receptor Probe	Detection Conditions/ Medium	Detection Limit (approximate)	Response Time (approximate)	Ref.
01 02	Benzimidazole-based benzoate Cu(II) receptor 2-methylquinoline-based benzohydrazide Cu(II) receptor	HEPES buffer (20 mM, pH 7.4) DMSO/HEPES (9:1 v/v, 20 mM, pH 7.22)	2.51 μM 0.949 μM	_	[9] [12]
03	Azo-dye-based bis-Schiff based receptor	DMF/ H ₂ O (1:1 v/v, 10 mM, pH 7.4)	1.7 μM 0.335 μM	<10 s	[14]
04	Imidazole-based Cu(II) receptor	Isopropanol/H ₂ O (1:1 v/v, <i>Bis-Tris</i> buffer, pH 7)	1.2 μΜ	_	[28]
05	Benzimidazole-based Cu(II) receptor	HEPES buffer (20 mM, pH 6.0)	1.35 μΜ	30 s	[29]
06	Pyridoxal Schiff base-based Cu(II) receptors	MeOH/H ₂ O	3.4 μM	_	[34]
			3.2 μΜ		
07	Quinoline-pyrazoline based Cu(II) receptor	DMSO/PBS buffer (1:9 v/v, pH 7.4)	0.2 μΜ	_	[36]
08	Coumarin-dipicolylamine-based Cu(II) receptor	CH ₃ CN/HEPES buffer (1:19 v/v, 10 mM, pH 7.4)	14 nM	Few seconds	[43]
09	Schiff-base-based receptor	DMSO/bis-Tris buffer (9:1 v/v)	4.0 μM	Few seconds	[48]
10	Terephthalic acid-coordinated Cu(II) MOF	H_2O	10.0 μΜ	20 min	[66]
11	Silver nanoparticles capped with carbon dots (AgNPs-CDs)	H_2O	0.01 μΜ	_	[67]
12	Peptide-based Hg(II) receptor	HEPES buffer (10 mM, pH 7.4)	60.9 nM	40 s	[99]
13	Silver/nitrogen-doped carbon nanoparticles (Ag/ NCNPs)	H_2O	0.18 μΜ	1 min	[100]
14	Cyclam-functionalized carbon dot-based Cu(II) receptor	HEPES buffer (pH 7.4)	0.13 μΜ	_	[101]
15	Quinolimide-based Cu(II) receptor	CH ₃ CN/H ₂ O (1:1 v/v, pH 7.2)	0.178 μΜ	0.5 min	[102]
16	Peptide-based Cu(II) and Hg(II) receptor	HEPES buffer (10 mM, pH 7.4)	0.27 μM and 0.12 μM	50 s	[103]
17	dppy-based Cu(II) receptor (CuCl ₂ L)	CH ₃ CN/HEPES buffer (1:2 v/v, 10 mM, pH 7.4)	82.50 nM	Immediate	Present Work

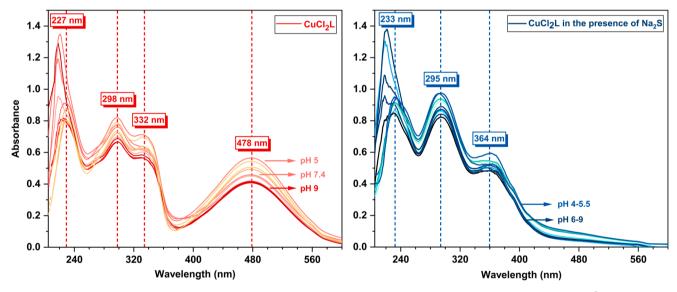


Fig. 5. UV–Vis absorption spectra of (left) in-situ prepared CuCl₂L (38 μ M, 1.5 mL) and (right) in-situ prepared CuCl₂L (38 μ M, 1.5 mL) with S²⁻ (Na₂S, 1 eq., 1 mM, 57 μ L) in aqueous acetonitrile (2:1 v/v, 10 mM HEPES, 4 \leq pH \leq 9).

The ligand ${\bf L}$ is highly soluble in acetonitrile but insoluble in water, leading to precipitation as the water content in the acetonitrile solution increases. However, in aqueous acetonitrile (2:1 v/v) within the optimal pH range of 4 to 9, the *in-situ* prepared ${\bf CuCl_2L}$ receptor exhibited good water tolerance and retained its effectiveness for sulfide detection. Upon the introduction of sulfide ions in solutions with higher water content, ${\bf Cu(II)}$ was displaced from the ${\bf CuCl_2L}$ complex, leading to the formation of ${\bf CuS}$, but this also caused the precipitation of ${\bf L}$ due to its insolubility. This precipitation interfered with the absorption spectra. To minimize this interference and maintain clear spectral readings, we recommend using a 2:1 (v/v) water-to-acetonitrile ratio.

3.2.2. Naked eye colorimetric test using glass vials

For a naked-eye colorimetric test, we added one equivalent of S2-

(Na₂S 1 mM, 57 μL) and ten equivalents of other anions (10 mM, 57 μL) to separate glass vials containing the *in-situ* prepared $CuCl_2L$ receptor (38 μM, 1.5 mL) in aqueous acetonitrile (2:1 v/v, 10 mM HEPES, pH 7.4), as depicted in Fig. 6. Notably, the addition of just one equivalent of Na₂S to the light brownish-red colored $CuCl_2L$ receptor solution resulted in an immediate color change to almost colorless, mirroring the colorless nature of **L** in acetonitrile. Conversely, the addition of other anions (ten equivalents) did not induce any color change in the $CuCl_2L$ receptor solution. This stark contrast in color response distinctly highlights the $CuCl_2L$ receptor's exceptional capability for naked-eye colorimetric detection of sulfide in an aqueous acetonitrile medium. Furthermore, for colorimetric detection, a higher water content than the recommended 2:1 (v/v) water-to-acetonitrile ratio can be used without compromising the detection capability. In this case, the $CuCl_2L$ receptor remains



Fig. 6. Colorimetric picture of the *in-situ* prepared $CuCl_2L$ (38 μ M, 1.5 mL) upon addition of S^2 –(Na₂S 1 mM, 57 μ L) and 10 equivalents of different anions (F¯, Cl¯, Br¯, l¯, CH₃CO $_2$, HCO $_3$, H₂PO $_4$, NO $_3$, SO $_3$, SO $_3$, SO $_3$, SO $_3$, and HPO $_4$; 10 mM, 57 μ L) in aqueous acetonitrile (2:1 v/v, 10 mM HEPES, pH 7.4).

effective for sulfide detection, even with the additional precipitation of \mathbf{I}

3.2.3. Spray test on tissue papers

A spray test on tissue papers was conducted to demonstrate the practical application of the $CuCl_2L$ receptor in sulfide detection. In this experiment, one equivalent of Na₂S (1 mM), ten equivalents of various other anions (10 mM), and the *in-situ* prepared $CuCl_2L$ receptor (38 μ M) were used in an aqueous acetonitrile solution (2:1 v/v, 10 mM HEPES, pH 7.4).

Initially, tissue papers were immersed in different analyte solutions and dried. These papers were then sprayed with the CuCl₂L receptor solution (Fig. 7a). In a complementary approach, other tissue papers were first soaked with the in-situ prepared CuCl₂L receptor, dried, and then sprayed with solutions of different analytes (Fig. 7b).

In both experimental setups, the results were consistent with those obtained from the naked-eye colorimetric tests in glass vials. The application of the CuCl_2L receptor, whether sprayed onto the analyte-soaked papers or used as a spray on receptor-soaked papers, led to an immediate and discernible color change. This change was a clear indicator of the presence of sulfide, demonstrating the receptor's capability for effective and low-cost nanomolar-level sulfide detection. The simplicity and rapid response of this method hold great promise for practical applications in sulfide monitoring.

3.2.4. Sulfide detection in various water samples

Next, we conducted sulfide estimation experiments on a diverse range of water samples, including river, pond, tap, and supply water, using our <code>in-situ</code> prepared <code>CuCl_2L</code> receptor (38 μM). The findings, detailed in <code>Table 2</code>, demonstrate that the sulfide detection efficacy of the <code>CuCl_2L</code> receptor is on par with previously reported receptors [100]. This suggests its potential for broad application in analyzing various types of water samples.

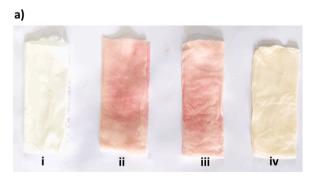
3.2.5. DNA binding study

Anti-cancer drugs function by binding to DNA, altering the DNA replication process, thereby inhibiting the growth of cancer cells and inducing apoptosis [41,104]. In this context, copper complexes have garnered significant interest due to their unique non-covalent DNA binding interactions and mechanisms of action, which differ markedly from those of platinum complexes [39,105]. To evaluate the DNA-binding capability of our *in-situ* synthesized CuCl₂L receptor, we conducted absorption and emission spectral titrations. These experiments were performed in an aqueous acetonitrile HEPES buffer at a pH of 7.4.

In our UV–Vis absorption spectral studies, (Fig 8a) the *in-situ* prepared $CuCl_2L$ receptor exhibited notable changes upon the gradual addition of DNA. Specifically, the characteristic LMCT absorption band at 478 nm displayed hypochromic shifts and a significant bathochromic shift of approximately 33 nm. This observation suggests a strong noncovalent intercalative binding, characterized by the stacking interaction between the planar aromatic chromophore of the $CuCl_2L$ receptor and DNA base pairs [73,106]. This binding mechanism is further substantiated by a high intrinsic binding constant, $K_b = 7.74 \times 10^5 \, \mathrm{M}^{-1}$, calculated through application of equation (1). A plot of $[CT-DNA]/(\varepsilon_a - \varepsilon_f)$ versus [CT-DNA] is shown in Fig. S11a.

Additionally, this absorption spectral study is corroborated by the EB displacement assay in our emission spectral study (Fig 8b). The gradual addition of the **CuCl₂L** receptor (90 μ M, 50 μ L, non-fluorescent) to the fluorescent EB-CT-DNA adduct (40 μ M, 1.8 mL) resulted in steady hypochromic shifts in the emission band of the adduct, indicative of intercalation by the **CuCl₂L** receptor [70,107]. The robustness of this intercalative binding interaction is further evidenced by the high values of $K_{qpp} = 2.381 \times 10^7 \, \mathrm{M}^{-1}$ and $K_{SV} = 7.065 \times 10^4 \, \mathrm{M}^{-1}$, calculated using equations (3) and (2), respectively, from the linear fit plot of F_0/F vs [**CuCl₂L**] (μ M) (see Fig. S11b).

Further elucidation of the binding interactions between the $CuCl_2L$ receptor and CT-DNA were observed through a viscometry titration experiment. This study revealed that the specific viscosity of CT-DNA (10 mM HEPES, pH 7.4) gradually increased upon gradual addition of



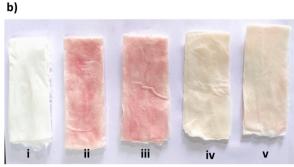


Fig. 7. a) Tissue papers initially immersed in various analyte solutions and dried, followed by treatment with the $CuCl_2L$ receptor solution by spraying. i) Fresh tissue paper; ii) no analytes; iii) mixture of other anions (except Na_2S), iv) mixture of other anions and Na_2S . b) Tissue papers pre-treated with the *in-situ* prepared $CuCl_2L$ receptor (38 μ M), followed by spraying with solutions of different analytes. i) Fresh tissue paper; ii) no analytes, iii) mixture of other anions (excluding Na_2S), iv) Na_2S , v) mixture of other anions including S^{2-} (Na_2S).

 Table 2

 Estimation of sulfide concentration in various water samples.

Water Sample	Added (µM)	Found (µM)	Recovery (%)	Water Sample	Added (μM)	Found (µM)	Recovery (%)
Supply water	2	1.88	94.00	Lab water	2	1.80	90.00
	4	3.77	94.25		4	3.61	90.25
	6	5.66	94.33		6	5.37	89.50
Pond water	2	2.17	108.50	River water	2	2.16	108.00
	4	4.11	102.75		4	4.10	102.50
	6	5.89	98.17		6	5.93	98.83

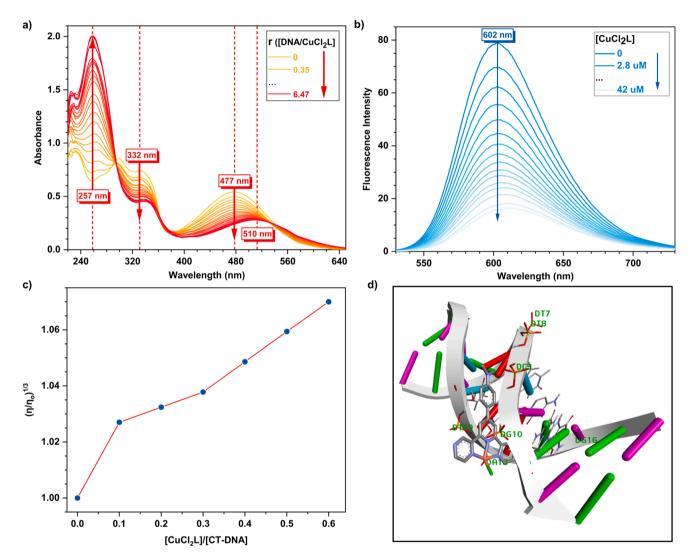


Fig. 8.A. a) Absorption spectra of the *in-situ* prepared CuCl₂L receptor (32 μM, 1.5 mL) upon gradual addition of CT-DNA (15 μL) in aqueous acetonitrile (2:1 v/v, 10 mM HEPES, pH 7.4). b) Emission spectra of EB-CT-DNA (40 μM, 1.8 mL) in aqueous HEPES buffer (10 mM, pH 7.4) upon gradual addition of *in-situ* prepared CuCl₂L receptor (90 μM, 50 μL). c) Effect of the CuCl₂L receptor addition on the relative viscosity of CT-DNA. d) Molecular docking model of the CuCl₂L receptor's interaction with CT-DNA (PDB ID: 1BNA).

CuCl₂L receptor. This increase is attributed to the intercalation of CuCl₂L receptor into the CT-DNA, leading to an elongation of its double helix structure, and, consequently, an increase in specific viscosity. This observation strongly supports the hypothesis of intercalative binding between the CuCl₂L receptor and CT-DNA. The specific viscosity values of CT-DNA were derived from the $(\eta/\eta_0)^{1/3}$ vs [CuCl₂L/CT-DNA] plot (Fig 8c) [108,109].

Finally, a molecular docking study was also carried out to investigate the mechanism of the binding interactions of the CuCl₂L receptor with CT-DNA. The lowest-energy structure obtained from this investigation (Fig 8d) revealed that the CuCl₂L receptor binds to a minor groove of

DNA. This binding is characterized by multiple interactions: strong conventional N–H bonding between a free N atom of one of the pyrazine moieties of the receptor and nucleobase DG 16 (2.40 Å); strong C–H bonding between the C atom of the amine moiety (NMe₂) with nucleobases DT 19 (2.93 Å) and DT 7 (3.08 Å); C–H bonding between the C atom of a pyrazine moiety with nucleobase DC 9 (3.28 Å); and weak π –alkyl interaction between the methyl groups of the receptor's amino moiety and the aromatic π -electron cloud of nucleobases DA 18 (5.29 Å), DT 8 (5.28 Å), and DT 19 (5.17 Å). Additionally, a strong π –anion interaction was observed between the π -electron cloud of a pyrazine moiety of the CuCl₂L and the phosphate anion of DG 10 (3.36 Å),

indicating strong intercalative binding. A 3-D graphical representation depicting the CT-DNA binding interaction of the $CuCl_2L$ receptor, illustrating both the hydrogen bonding and π -stacking interactions, is shown in Fig. S14. These molecular docking findings align well with the results obtained from our absorption, emission spectral studies, and viscometry titrations, thereby corroborating the $CuCl_2L$ receptor's intercalative binding mechanism with CT-DNA.

3.2.6. BSA binding interaction study

In the absorption spectral experiment, the gradual addition of the in- $\it situ$ prepared $\it CuCl_2L$ receptor to BSA (in a 100 mM PB buffer, 2 mL) led to the emergence of a new absorption band at 460 nm. This band displayed a consistent hyperchromic shift, indicating a binding interaction between the CuCl₂L receptor and BSA (Fig 9a). Additionally, in our emission spectral study, which focused on tryptophan quenching in BSA, a gradual decrease in the intrinsic fluorescence intensity of BSA was observed (evidenced by a hypochromic shift) with the increasing concentration of the in-situ prepared CuCl₂L receptor (Fig 9b) [110]. This quenching effect suggests a conformational change in BSA's hydrophobic cavity, particularly in the subdomain IIA containing the Trp-214 residue [111], caused by the receptor binding to BSA. This hypothesis is further supported by a comparatively high dynamic quenching constant, $K_{SV} = 1.6 \times 10^5 \,\mathrm{M}^{-1}$, calculated from the linear fit plot of F_0/F vs [CuCl₂L] (equation (2), see Fig. S12a). Moreover, the binding stoichiometry, approximately $n \sim 1$ (0.82), suggests a 1:1 binding ratio of the CuCl₂L receptor to BSA. The affinity constant, $K = 2.158 \times 10^4 \text{ M}^{-1}$,

calculated from the linear fit plot of $log(F_0 - F)/F$ vs $log[CuCl_2L]$ (equation (4), see Fig. S12b) reinforces the binding affinity of the $CuCl_2L$ receptor to BSA.

The molecular docking study also revealed that CuCl₂L receptor interacts with BSA, exhibiting a binding energy of -5.99 kcal mol⁻¹. The major interactions of the CuCl₂L receptor with the protein include a conventional hydrogen bond between the free N atom of one of its pyrazine moieties and ARG427 (2.20 Å), this pyrazine moiety also engaging in a π -alkyl interaction with ARG427 (5.41 Å). In turn, the other pyrazine moiety engages in a π -cation interaction with HIS145 (4.83 Å) and ARG458 (4.19 Å). Additionally, the pyridine moiety of the $CuCl_2L$ receptor is involved in π -anion interaction with the GLU424 (3.38 Å). The benzene ring of the receptor exhibits π -alkyl interactions with ALA193 (4.65 Å) and ILE455 (5.48 Å). Furthermore, two alkyl moieties of the receptor's amine group form π -alkyl interaction with TYR451 (4.32 Å and 4.84 Å). These diverse N-H bonding, π -cation, π -anion, and π -alkyl interactions collectively stabilize the CuCl₂L receptor within the hydrophobic cavity of BSA, A 3-D graphical representation depicting the BSA binding interaction of the CuCl₂L receptor in more detail is shown in Fig. S15. The molecular docking results corroborate the binding study findings of the CuCl₂L receptor with BSA, as observed in the UV-Vis absorption and fluorescence emission spectral studies. Details of the binding energy of various conformations of the BSA-CuCl₂L receptor and active L-amino acids residues nearby the CuCl₂L receptor are provided in Table S3.

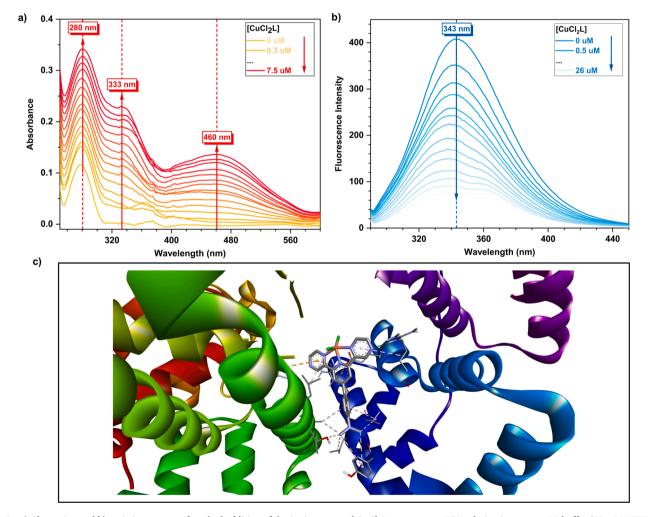


Fig. 9.A. a) Absorption and b) emission spectra of gradual addition of the *in-situ* prepared CuCl₂L receptor to a BSA solution in aqueous PB buffer (10 mM HEPES, pH 7.4). c) Molecular docking model of the CuCl₂L receptor's interaction with BSA (PDB ID: 4F5S).

4. Conclusions

In summary, recognizing the critical need for sulfide detection from biological, chemical, and environmental perspectives, we have developed and characterized a simple Cu(II)-dppy complex, CuCl2L. This complex is a 1:1 Cu(II) coordination of 4-(2,6-di(pyrazin-2-yl)pyridin-4yl)-N,N-dimethylaniline ligand L, whose properties were elucidated through a combination of spectroscopic techniques, including UV-Vis, FT-IR, HRMS, elemental analysis, and ESR, alongside quantum chemical calculations at the DFT and TD-DFT levels. UV-Vis absorption spectral studies confirmed the capability of the in-situ prepared CuCl₂L receptor to selectively detect sulfide at nanomolar levels, even in the presence of a tenfold excess of various other anions, namely F⁻, Cl⁻, Br⁻, I⁻, CH₃CO₂, HCO_3 , H_2PO_4 , NO_3 , SO_3^2 , SO_4^2 , $S_2O_3^2$, CO_3^2 , and HPO_4^2 , in aqueous acetonitrile (2:1 v/v, 10 mM HEPES, pH 7.4). The receptor exhibited an impressive detection limit of 82.50 nM. Additionally, the in-situ prepared CuCl₂L receptor presented a cost-effective approach for various real-time applications in sulfide anion detection. Further studies. including absorption, emission spectral analysis, viscometry, and molecular docking, have substantiated the CuCl₂L receptor's significant intercalative binding interaction with CT-DNA (binding constants: $K_b =$ $7.74 \times 10^5 \, \text{M}^{-1}$, $K_{app} = 2.381 \times 10^7 \, \text{M}^{-1}$ and $K_{SV} = 7.065 \times 10^4 \, \text{M}^{-1}$) and its effective binding interactions with BSA (binding constants, $K_{ab} =$ $2.158 \times 10^4 \,\mathrm{M}^{-1}$ and $K_{SV} = 1.6 \times 10^5 \,\mathrm{M}^{-1}$).

CRediT authorship contribution statement

Dipankar Das: Writing – original draft, Visualization, Methodology, Investigation, Formal analysis, Conceptualization. Aritra Roy: Software, Investigation, Formal analysis. Cauê P. Souza: Supervision, Investigation, Formal analysis. Somnath Mondal: Methodology, Formal analysis. Sourav Sutradhar: Software, Formal analysis. Parnashabari Sarkar: Software, Investigation, Formal analysis. Felipe Fantuzzi: Writing – original draft, Supervision, Software, Funding acquisition, Conceptualization. Biswa Nath Ghosh: Writing – review & editing, Visualization, Supervision, Funding acquisition, Conceptualization.

Declaration of competing interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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Appendix A. Supplementary data

Supplementary data to this article can be found online at https://doi.org/10.1016/j.jphotochem.2024.116154.

Data availability

Data will be made available on request.

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