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Near-Infrared Fluorescent Probes with Amine-Incorporated Xanthene Platforms for the Detection of Hypoxia

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Supporting Information

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NMR, mass, absorption, and fluorescence spectra; computation results; MTT assays of the probes, their fluorescence cellular imaging, and fluorescence imaging of *D. melanogaster* first-instar larvae (PDF)

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Abstract

Three fluorescent probes **A**, **B**, and **C** that function in the near-infrared wavelengths and utilize pseudo xanthene platforms with an oxygen atom at the 10-position replaced by a $[Me-N]^{2-}$ group have been created to identify hypoxia via nitroreductase determinations at the 0.04, 0.10, and 0.19 ng/mL levels. Theoretical calculations suggest that the probes are not planar due to steric interactions. Absorptions of photons result in the transition of electron density from the indoline moieties to delocalized orbitals on the anthranilic section, ending up on the nitro groups of the electron-poor (i.e., nonreduced) probes (i.e., **A**, **B**, and **C**), whereas those for the more electron-rich (i.e., reduced) probes consisted of movement from the indoline groups to the right side of the anthranilic sections, resulting in a shift in absorption.

Graphical Abstract



Keywords

fluorescent probes; nitroreductase; tunable sensitivity; hypoxia; ultrasensitivity

1. INTRODUCTION

Hypoxia, an inadequate oxygen level in tissues and cells, is often present with many illnesses. Tumors metastasize and deteriorate because vascular growth factors from tumor cells result in abnormal blood vessels under hypoxic conditions.¹⁻³ Therefore, it is important to accurately determine hypoxia with noninvasive, specific, and timely quantitative approaches.^{1,2,4} Optical fluorescence imaging fulfilled these criteria by visualizing the detailed morphology of cells and tissues with subcellular resolution and high sensitivity.¹⁻³ There have been recent reports of fluorescent probes functioning in the near-infrared region for hypoxic identification as these possess advantages of minimum interference from water Raman peaks, deep-tissue penetration, and low autofluorescence from the cells and tissues.^{1,2,4,5} As the level of hypoxia is closely associated with the concentration of reductive enzymes, such as nitroreductase, many fluorescent probes have been reported that can detect hypoxia by measuring nitroreductase. This is accomplished by introducing a nitro moiety to different fluorophores as photoinduced electron transfer acceptors or as a part of the fluorophores, allowing for the quantitative reduction of the nitro moiety and

thus the detection of nitroreductase.¹⁻³ These fluorescent probes display a low fluorescence background because of an electron-withdrawing feature of the nitro moiety.⁶ The enzyme nitroreductase transforms the nitro group into an amine and significantly turns on the probes' fluorescence.^{7,8} However, with some probes, the enzyme reaction can take a few minutes or up to an hour for the fluorescent probes to be reduced by nitroreductase.^{1,2}

Herein, three fluorescent probes **A**–**C** that are capable of signaling hypoxia in the nearinfrared region and incorporating nitro groups into new pseudo xanthene platforms, where the oxygen atoms at the 10-position are replaced by $[Me-N]^{2-}$ groups, are detailed (Scheme 1).⁹ As expected, these probes show extremely weak fluorescence since the nitro group quenches the fluorescence of the fluorophores. The effect of electron-withdrawing (i.e., Cl⁻) or electron-donating (i.e., piperidine) groups attached to the 9 position on the fluorescent probes on the response time to nitroreductase was also determined. Probe **A**, bearing the electron-with-drawing chloro group, exhibits the fastest response to nitroreductase in less than 1 min, presumably because the electron-withdrawing chloro group facilitates reduction of the nitro group by nitroreductase, while probe **C**, bearing an electron-donating piperidine moiety, has the slowest response to nitroreductase. Probe **B**, containing a neutral H atom at the 9-position, shows a slower response to nitroreductase than probe **A** but is faster than probe **C**. The probes have been used to visualize hypoxia in cells caused by treatment of different concentrations of CoCl₂ and to detect nitroreductase activity in first-instar larvae of *Drosophila melanogaster*.

2. EXPERIMENTAL SECTION

2.1. Chemicals and Reagents.

2.1.1. 9-Chloro-6-nitro-1,2,3,4-tetrahydroacridine (Compound 3).—2-Amino-4nitrobenzoic acid (364 mg, 2 mmol) and POCl₃ (6 mL) were mixed with cyclohexanone (500 mg, 2.46 mmol). The mixed solution was heated at 105 °C, stirred for 12 h under an argon atmosphere, cooled to ambient temperature, and dumped into to 200 mL of ice-cold water. The pH was adjusted to 7.0 by adding saturated NaHCO₃. Dichloromethane (2 × 100 mL) was used to extract the mixture which was further subjected to a saturated NaCl solution. Following collection and removal of the organic layer, the resulting yellow solid was chromatographed on a silica gel packed column (dimension, 0.063–0.2 mm Merck) through gradient elution with ethyl acetate and dichloromethane using ratios from 1:1 to 1:0 to yield compound **3** as a pale-yellow solid (419 mg, 80%). ¹H NMR (400 MHz, CDCl₃) δ 8.78 (d, *J* = 7.9 Hz, 1H), 8.30–8.15 (m, 2H), 3.11 (d, *J* = 5.9 Hz, 2H), 3.07–2.82 (d, *J* = 5.9 Hz, 2H), 1.92 (s, 6H). ¹³C NMR (101 MHz, CDCl₃) δ 22.57, 28.12, 34.73, 119.88, 125.09, 125.72, 128.67, 132.65, 141.42, 145.59, 148.00, 162.64.

2.1.2. Synthesis of compound 4—Methyl trifluoromethanesulfonate (3.12 g, 19 mmol) was placed into a solution of compound **3** (1 g, 3.8 mmol) dissolved into 6 mL of chloroform. This mixture was stirred at 60 °C with stirring for 10 h. Following removal of the solvent, the concentrate was washed with diethyl ether (2×50 mL) yielding compound **4** as a yellow solid.

2.1.3. Synthesis of Probe A.—Fisher's aldehyde (**5**) (0.56 mmol, 112 mg) and compound **4** (0.56 mmol, 200 mg) were placed into acetic anhydride (6 mL). This was heated and stirred at 50 °C for 2 h. The solvent was removed and the residue components were separated through chromatography on a column of nature and size similar to the one used in the procedure in section 2.1.1 under gradient elution with dichloromethane and MeOH from a 100:1 ratio to 30:1 and finally 20:1, yielding probe **A** as blue crystals (138 mg, 60%). ¹H NMR (400 MHz, CD₃OD) δ 8.69 (s, 1H), 8.35 (s, 1H), 7.75 (d, *J* = 13.9 Hz, 1H), 7.47 (d, *J* = 7.7 Hz, 1H), 7.39 (d, *J* = 7.3 Hz, 1H), 7.33 (d, *J* = 8.2 Hz, 1H), 7.29–7.23 (m, 2H), 6.20 (d, *J* = 13.9 Hz, 1H), 4.25 (s, 3H), 3.67 (s, 3H), 3.03 (t, *J* = 6.0 Hz, 2H), 2.85 (t, *J* = 6.1 Hz, 2H), 1.99 (t, *J* = 6.0 Hz, 2H), 1.65 (s, 6H). ¹³C NMR (101 MHz, CD₃OD) δ 19.87, 27.55, 27.80, 29.10, 30.75, 46.18, 49.45, 99.57, 102.29, 110.93, 113.77, 117.31, 119.74, 122.07, 125.30, 126.58, 127.14, 128.54, 137.79, 141.48, 142.87, 143.37, 148.68, 149.24, 157.71, 174.53. MS (ESI): calcd for [C₂₇H₂₇ClN₃O₂]⁺ 460.9815, found 460.1779.

2.1.4. Synthesis of Probe B.—Hydrazine 60% (43 mg, 0.86 mmol) and probe A (200 mg, 0.43 mmol) were placed into 6 mL dimethylformamide (DMF) and then stirred at 50 °C for 4 h. A separatory funnel containing 100 mL of dichloromethane was mixed with this solution, and 50 mL of saturated saline solution was used thrice to remove ionic materials. The organic layers were then obtained and dried using anhydrous Na₂SO₄. The solvents were obtained by filtration and then evaporated. The residue components were separated on an aforementioned column under gradient elution with dichloromethane and MeOH from a 70:1 ratio to 20:1, affording probe **B** as blue crystals (128 mg, 70%). ¹H NMR (400 MHz, CDCl₃) δ 8.69 (s, 1H), 8.23 (q, *J* = 9.1 Hz, 2H), 7.24 (d, *J* = 10.1 Hz, 3H), 7.04 (t, *J* = 7.4 Hz, 1H), 6.92 (d, J = 7.8 Hz, 1H), 5.67 (d, *J* = 12.9 Hz, 1H), 4.32 (s, 3H), 3.43 (s, 7H), 2.76–2.80 (m, 4H), 1.91–1.95 (m, 2h), 1.77 (s, 6H), 1.55 (s, 6H). ¹³C NMR (101 MHz, CDCl₃) δ 21.37, 24.28, 27.04, 27.98, 29.22, 29.52, 45.75, 48.12, 53.67, 109.04, 114.85, 118.88, 119.63, 122.17, 123.42, 127.84, 127.99, 128.46, 131.80, 141.64, 142.75, 148.77, 153.68, 161.69, 169.05. MS (ESI): calcd for [C₃₂H₃₇N₄O₂]⁺ 509.2911, found 509.2913.

2.1.5. Synthesis of Probe C.—Piperidine (55 mg, 0.65 mmol) and probe A (200 mg, 0.43 mmol) were placed into 6 mL of dimethylformamide (DMF) and stirred at 50 °C for 2 h and then placed into a separatory funnel with 100 mL of dichloromethane used to clean out the reaction flash. This solution was thrice washed with 50 mL of a saturated saline mixture. The relevant layers were collected and dried by anhydrous Na₂SO₄, filtered followed by evaporation to dryness. The dried solids were purified on a column (described above) under gradient elution of CH₂Cl₂ and MeOH at a 70:1 ratio to 25:1, affording probe **C** as blue crystals (175 mg, 80%). ¹H NMR (400 MHz, CD₃OD) δ 8.29 (s, 1H), 7.89 (dd, J = 8.6, 2.0 Hz, 1H), 7.59 (d, J = 8.6 Hz, 1H), 7.52 (s, 1H), 7.25–7.18 (m, 1H), 6.97 (t, J = 7.7 Hz, 2H), 6.86–6.75 (m, 2h), 5.56 (d, J = 13.6 Hz, 1H), 3.92 (s, 3H), 3.20 (s, 3H), 2.44–2.48 (m, 2H), 2.38–2.42 (m, 2H), 1.53–1.58 (m, 2H), 1.23 (s, 5H). ¹³C NMR (101 MHz, CDCl₃/DMSO = 1:1) δ 20.40, 27.82, 28.93, 29.50, 31.24, 45.93, 48.84, 97.85, 110.43, 113.82, 116.42, 120.20, 122.03, 124.92, 128.57, 129.21, 129.45, 132.89, 140.10, 140.55, 142.48, 147.38, 148.40, 157.95, 172.61. MS (ESI): calcd for [C₂₇H₂₈N₃O₂]⁺ 426.5395, found 426.2182.

2.1.6. Synthesis of Probe AH.—SnCl₂ (163 mg, 0.86 mmol) was mixed with 6 mL of ethanol containing probe **A** (200 mg, 0.43 mmol) and stirred at 80 °C for 6 h. The solvents were removed and the residue components were separated by chromatography (using the same type of column mentioned above) with solutions of dichloromethane and MeOH in ratios of 30:1 to 10:1, yielding **AH** as blue crystals (111 mg, 60%). ¹H NMR (400 MHz, CDCl₃) δ 7.87 (d, *J* = 9.1 Hz, 1H), 7.77 (s, 1H), 7.24 (d, *J* = 1.2 Hz, 1H), 7.21–7.18 (m, 1H), 7.13–7.09 (m, 1H), 7.08 (s, 1H), 7.01–6.95 (m, 1H), 6.81 (d, *J* = 7.9 Hz, 1H), 5.47 (d, *J* = 12.8 Hz, 1H), 4.24 (s, 3H), 3.30 (s, 3H), 2.85 (t, *J* = 6.5 Hz, 2H), 2.70 (t, *J* = 6.5 Hz, 2H), 1.93–1.85 (m, 2H), 1.51 (s, 6H). ¹³C NMR (101 MHz, CDCl3) δ 12.45, 21.63, 27.47, 27.95, 29.83, 30.18, 42.27, 45.45, 47.43, 53.92, 93.10, 98.66, 108.13, 116.67, 117.03, 119.62, 122.20, 122.63, 126.89, 127.26, 128.33, 139.73, 143.59, 143.94, 155.28, 156.74, 166.60. MS (ESI): calcd for [C₂₇H₂₉ClN₃]⁺ 430.9995, found 430.2043.

2.2. General Procedure to Detect Nitroreductase.

All absorption and fluorescence spectra were conducted in a mixture of 0.05 M PBS buffer (pH 7.4) containing 1.0% DMSO. Twenty microliters of a solution containing 1 mM probe **A**, **B**, or **C** was added to 1 mL of PBS buffer in separate 2 mL test tubes, and 500 μ M of NADH was prepared by adding NADH to these solutions. Nitroreductase was added to this solution followed by more PBS buffer to attain a final volume of 2 mL with nitroreductase concentrations of 0.4 ng/mL for probe **A**, 0.7 ng/mL for **B**, and 1.75 ng/mL for **C**. Once solutions were mixed completely, optical measurements were conducted by transferring the solution to a quartz cuvette while maintaining the temperature at 37 °C.

2.3. Cell Culture.

The A549 material was purchased from ATCC. A549 material were cultured in Dulbecco's modified Eagle's medium (DMEM) in the presence of 10% fetal bovine serum at 37 °C under a 5% CO₂ atmosphere for the normoxic circumstance (20% O₂), while the cells were cultivated for 5.5 h at 37 °C under a hypoxic condition with 1% O₂ and 5% CO₂ concentrations.

2.4. Confocal Fluorescence Imaging for Living Cells.

A549 cells were cultured for 12 h with initially 1×10^5 cells in each 35 mm confocal dish (MatTek). The cells were scrubbed with 0.05 M pH 7.04 PBS buffer and further subjected to 5 μ M of probe **A**, **B**, or **C** in 0.05 M pH 7.4 PBS buffer containing 1.0% DMSO for 10 min at 37 °C. For experiments (i.e., of the colocalization variety), A549 cells were subjected to 5 μ M MitoView blue and 5 μ M of compound **A**, **B**, or **C** in DMEM cell culture media in the presence of 1% DMSO for 10 min, and then scrubbed with PBS buffer thrice before fluorescence imaging using an Olympus IX 81 confocal fluorescence microscope. Fluorescence with samples containing MitoView Blue was obtained from 425 to 475 nm at 405 nm excitation, while that for the near-infrared fluorescence of the probes was obtained in the range 625 to 675 nm at 559 nm excitation. An FV10-ASW 3.1 viewer (Olympus) equipped with *Photoshop* and *ImageJ* software was employed to differentiate the pictures.

2.5. Theoretical Calculations.

Optimization and frequency calculations (no imaginary frequencies observed) were done at the APFD/6-31+G(d) level with *Gaussian 16*¹⁰ and *GausView*¹¹ following previously described procedures.¹² Excited states (ten) were computed using TD-DFT optimizations¹³ in water employing the Polarizable Continuum Model (PCM).¹⁴ Data for the probes are in Supporting Information.

3. RESULTS AND DISCUSSION

3.1. Synthetic Approach and Characterization.

Herein we detail the development of nitroreductase-based hypoxia probes as hypoxia can result in the overexpression of reductase enzymes.^{1,5} We introduced a nitro group to near-infrared fluorophores based on the pseudo xanthene platform, 10-methyl-1,2,3,4tetrahydro-10-4-acridine, at the 6-position (Scheme 1).⁹ This was accomplished by conducting a condensation reaction of 4-nitroanthranilic acid, 1, with cyclohexanone, 2, in fresh phosphorus(V) oxychloride at high heat,¹⁶⁻¹⁸ yielding 9-chloro-6-nitro-1,2,3,4tetrahydroacridine, 3, (Scheme 2). We then converted 3 into 9-chloro-10-methyl-6nitro-1,2,3,4-tetrahydroacridin-10-ium iodide, 4, by a methylation reaction on compound **3** (Scheme 2). Probe A was synthesized by reacting compound 4 with Fisher's aldehyde (5) through a condensation reaction. In order to assess the effect of different electronwithdrawing or electron-donating groups on the probe's sensitivity to nitroreductase, we replaced the chlorine atom on **A** with a hydrogen atom (i.e., **B**) and a nitrogen atom (i.e., in piperidine, C). Probe B was synthesized by reacting probe A with hydrazine in dimethylformamide, while probe \mathbf{C} was synthesized by a substitution reaction of probe \mathbf{A} with piperidine. The reduction product (probe AH) of probe A was obtained by mixing probe A with tin(II) chloride in dichloromethane (Scheme 2).

3.2. Optical Responses of the Probes to Nitroreductase.

The responses of the probes verified by spectroscopy to nitroreductase in pH 7.4 PBS buffer containing 1% DMSO solution (Figure 1) were examined. Probes A, B, and C absorbed at 633, 623, and 603 nm in the absence of nitroreductase, respectively (Figure 1). In the presence of nitroreductase (0.4 ng/mL for probe A, 0.7 ng/mL for B, and 1.75 ng/mL for C), they were blue-shifted by 30, 36, and 49 nm in their absorption spectra, and they exhibited absorptions at 603, 567, and 574 nm, respectively. The shifts in the probes' absorption to the blue spectral region arose from changes in the electron delocalization in the HOMO due to an electron-donating amine group formed through the reduction of the nitro group on the compounds by the enzyme. Solutions of the probes did not generate fluoresce due to quenching presumably from the nitro group. However, the fluorescence intensities of the probes significantly increased upon the gradual addition of nitroreductase with significantly enhanced fluorescence peaks at 666, 655, and 656 nm for probes A, B, and C, respectively. The detection limit was calculated by eq 3σ /S. All three probes show good linearity within a stated range of nitroreductase concentrations. For example, probe A (Figure S18) exhibits a good linear fluorescence relationship in the 0.04–0.2 μ g range with added nitroreductase resulting in the equation of F = 8.00 [nitroreductase] (μ g) + 0.00895 ($R^2 = 0.98363$), where

F signifies the intensity of the reaction solution. Probe **A** showed the fastest response to nitroreductase in less than one minute and has a limit of detection of 0.04 ng/mL. This is presumably because the electron-withdrawing chlorine atom at the 9-position facilitates the reduction of the nitro group by nitroreductase. Probe **C**, bearing the electron-donating piperidine moiety at the 9-position of the xanthene platform, displayed the slowest response to nitroreductase and required 10 min and can detect down to 0.19ng/mL (Figure 2). Probe **B**, bearing an H atom at the 9-position of the xanthene platform, exhibited a slower response to nitroreductase than probe **A**, but faster than probe **C** (Scheme 1 and Figure 2). Probe **B** possesses a 0.10 ng/mL limit of detection. Temperature increases from 25 to 37.5 °C also resulted in a slight fluorescence enhancement. However, a further increase in temperature to 40 °C resulted in a slight fluorescence decrease.

3.3. Theoretical Calculations of the Probes.

The geometries of the probe and their electronic transitions were estimated via theoretical calculations. Current density diagrams for the probes and their reduced forms are displayed in Figure 3. The probes are not planar and contain interplanar angles between the 1,3,3trimethyl-2-(prop-1-en-1-yl)indoline and the anthranilic moieties of 42.8°, A; 42.9°, AH; 40.5° , **B**; 41.6° , **BH**, 47.4° , **C**; and 43.3° , **CH**. This is presumably due to the reduction of steric interactions between the methyl group attached to the anthranilic N atom and the two methyl groups attached to the indoline five-membered ring. The calculated absorption values compared to those obtained experimentally are within the theorized values of 0.20-0.25 eV (i.e., A 0.04, AH 0.23, B 0.26, BH 0.22, C 0.03, CH 0.17 eV), except for probe B.¹⁵ This suggests that the level of theory at APFD/6-31+G(d) was useful in calculating the absorption spectra. The spectra for probes A, B, and C consisted of two main absorptions in the visible region arising from HOMO to LUMO and HOMO to LUMO+1, as shown in Figure 3. Both of these consisted of a movement of electron density from the indoline moieties to largely delocalized orbitals over the anthranilic section and localized onto the nitro groups. In contrast, absorptions for the reduced probes consisted of transitions from the indoline groups to the right side of the anthranilic moieties, and this is the reason for the absorption shift upon reduction of probes A-C.

3.4. Determination of the Probe Product by a Mass Spectrometer.

To confirm that probe **AH** is produced from the action of nitroreductase on probe **A**, we conducted high-resolution electrospray mass spectrometric analysis on the product. A peak at m/z of 430.20427 in the electrospray mass spectrum corresponding to probe **AH** ([M+] = 430.21) was observed (Figure S14). Furthermore, the fluorescence spectra of the enzyme-catalyzed product and that of synthesized probe **AH** exhibit similar characteristics (Figure S22A). These results confirm that nitroreductase effectively reduces probe **A** to probe **AH**.

3.5. Study Sensing Mechanism of Probe A to Nitroreductase.

We chose probe **A** to study the probe response function to nitroreductase by fluorescence spectroscopy. Probe **A** shows weak fluorescence in the presence of 500 μ M NADH, indicating that NADH is unable to reduce probe **A** (Figure S22B). Probe **A** exhibits

fluorescence increases with both 0.4 μ g/mL nitroreductase and 500 μ M NADH at 37 °C (Figure S22B). An enzyme inhibition reaction established that this increased fluorescence response resulted from the enzyme-catalyzed reduction. When probe **A** was treated with either 1 mM or 2 mM of dicoumarin (DM) followed by addition of nitroreductase, the probe fluorescence intensities reduced significantly compared to those observed without added dicoumarin (Figure S22B).⁷ This suggests that dicoumarin effectively inhibits the enzyme activities and hinders the reduction of the probe. Enzyme inhibition was also observed at 4 °C, as we observed a very weak fluorescence when probe **A** and nitroreductase were mixed at that temperature (Figure S22B).⁷ These enzyme inhibition results convincingly demonstrated that the enhanced fluorescence responses result from enzyme-facilitated reduction reactions.⁷ Measurements of solutions, using high-resolution ESI mass spectrometry, also confirmed the enzyme-reduced product, probe **AH** (Figures S14 and S15).

3.6. Probe Selectivity to Nitroreductase.

The probes are selective and reactive to nitroreductase over CoCl₂, CaCl₂, glucose, vitamin C, hydrogen peroxide, tyrosine, glycine, glutamate, BSA, and KO₂ (Figure S19). The probes showed low cytotoxicity and biocompatibility through MTT assays, and cell viability was higher than 84% with probe concentrations up to 20 μ M (Figure S23), indicating that the probes can be useful potential imaging agents for nitroreductase detection without cytotoxic effects.

3.7. Fluorescence Cellular Imaging Applications of the Probes.

The probes can selectively stain mitochondria by electrostatically interacting with the negatively charged mitochondrial membrane.¹⁶ This was confirmed in a colocalization experiment, where A549 cells were incubated with mitochondria-specific Mitoview blue and one of either probe **A**, **B**, or **C** under a 1% O₂ level for 5.5 h, respectively. A correlation coefficient of 0.948 (i.e., Pearson coefficient) for probe **A** and Mitoview demonstrated that **A** specifically targets mitochondria. Further, probes **B** and **C** also have similar specificities to mitochondria in live cells (Figures S24-S26).

To investigate the result of a hypoxic condition for enzyme capability, we visualized the spectroscopic reactions of probes **A-C** in live A549 cells under normoxia and hypoxia conditions. Hypoxia conditions can be achieved either by reducing oxygen levels to 1% or by chemical treatment with varying concentrations of Co^{2+} ions for 6 h.^{6,17} Induced hypoxia via Co^{2+} treatment is easily accomplished in the laboratory and is a well-known effective strategy in the biology realm. It is known that the Hypoxia Inducible Factor-1 (HIF-1) can be induced by adding Co^{2+} ions as this inhibits the enzyme prolyl hydrolase which regulates HIF.^{6,17} A549 cells under hypoxic conditions through cobalt chloride treatment displayed a much stronger cellular fluorescence with increased cobalt chloride concentrations from 50 to 100 μ M, while the cells under normoxia conditions without chemical treatment showed no fluorescence (Figure 4). We find that the strong fluorescence signal in A549 cells after treatment with cobalt chloride, can be reduced significantly by pretreatment of the cells with N-acetyl cysteine (NAC). This is known to reverse cell damage by consuming ROS species resulting in increases in GSH levels and protecting the H9c2 cardiomyocytes from injuries

3.8. Fluorescence Imaging of Drosophila melanogaster First-Instar Larvae.

Probes **A**–**C** can be used to detect nitroreductase in *D. melanogaster* first-instar larvae. As is evident in Figure 5 for probe **A** and Figure S29 for probes **B** and **C** (all labeled blank), larvae do not fluoresce if incubated with these probes. However, there is strong fluorescence of the larvae after they were hatched with the probes and then further treated with nitroreductase (Figure 5 and Figure S29), suggesting that the probes can be employed to image nitroreductase.

4. CONCLUSIONS

We have successfully synthesized amine-incorporated pseudo xanthene platforms that can be used to identify hypoxia in live cells. The probes have many advantages, including NIR emission, fast reaction times, and determinative fluorescence responses to nitroreductase.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Figure 1.

Absorption and fluorescence spectra for $10 \,\mu\text{M}$ probes –-**C** in pH 7.4 0.05 M tris buffers with 1.0% DMSO containing different concentrations of nitroreductase and 500 μ M NADH under 580 nm excitation and 3 min incubation time for probe **A**, 5 min for probe **B**, and 10 min for probe **C**, respectively.



Figure 2.

Time-dependent fluorescence intensities at 655 nm for 10 μ M probe **A**, at 655 nm for 10 μ M probe **B**, and at 656 nm for 10 μ M probe **C** in pH 7.4 0.05 M. The solution consisted of tris buffer and 1% DMSO with 0.2 μ g/mL nitroreductase and 500 μ M NADH under 580 nm excitation.



Figure 3.

Diagrams of the electron density variance as 3D surface representations of the probes A (top row), B (middle), and C (bottom) for two excited states (ES) and the reduced forms in the column to the right, respectively. The Supporting Information contains diagrams of the numbered MOs.



Figure 4.

Confocal fluorescence microscopy pictures of A549 cells subjected to 5 μ M probe **A** with various hypoxic conditions for 10 min. From left to right: (1) control cells without treatment of CoCl₂. (2) The cells were treated with 50 μ M CoCl₂ for 6 h, and further incubated with 5 μ M probe **A** for 10 min. (3) The cells were treated with 100 μ M CoCl₂ for 6 h, and further incubated with 5 μ M probe **A** for 10 min. (4) The cells were pretreated with N-acetyl cysteine (1.5 mM) for 1 h, further treated with 100 μ M CoCl₂ for 6 h, and finally incubated with 5 μ M probe **A** for 10 min. (5) The cells were cultured under a hypoxic condition with 1% oxygen level for 6 h, and then further incubated with 5 μ M probe **A** for 10 min. The images were obtained in the ranges detailed in the Experimental Section. Scale bar = 20 μ m.



Figure 5.

Images of *D. melanogaster* first-instar larvae subjected to incubation with 10 μ M probe **A** for 2 h, and further incubation with 10 μ g/mL NTR, and 500 μ M NADH at 37 °C for 120 min. Scale bar: 200 μ m. (650 ± 30 nm, λ_{ex} = 580 nm semiconductor laser).



Scheme 1.

Chemical and Structural Changes of the Probes in the Responses to Nitroreductase



Scheme 2. Synthetic Approach for the Probes