

Highly Selective and Sensitive Detection of Biogenic Defense Phytohormone Salicylic Acid in Living Cells and Plants Using a Novel and Viable Rhodamine-Functionalized Fluorescent Probe

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ABSTRACT: Detecting plant-derived signal molecules using fluorescent probes is a key topic and a huge challenge for scientists. Salicylic acid (SA), a vital plant-derived defense hormone, can activate global transcriptional reprogramming to systemically express a network of prominent pathogenesis-related proteins against invasive microorganisms. This strategy is called systemic acquired resistance (SAR). Therefore, monitoring the dynamic fluctuations of SA in subcellular microenvironments can advance our understanding of different physiological and pathological functions during the SA-induced SAR mechanism, thus benefiting the discovery and development of novel immune activators that contribute to crop protection. Here, detection of signaling molecule SA in plant callus tissues was first reported and conducted by a simple non-fluorescent rhodamine-tagged architecture bearing a flexible 2-amino-*N,N*-dimethylacetamide pattern. This study can markedly advance and promote the usage of fluorescent SA probes for distinguishing SA in the plant kingdom.

KEYWORDS: salicylic acid, systemic acquired resistance, rhodamine probe, plant callus tissues

INTRODUCTION

Biogenic signal molecules are attractive and important physiological activators that can conduct and determine diverse physiological processes, such as cellular homeostasis, metabolism, energy conversion, signal conduction, and immunity, throughout the cell cycle.^{1–6} The abnormal expression level of these bioactive substances can result in a series of imbalance and reverse effects and is constantly associated with various major diseases. Thus, detecting these biogenic signal molecules in subcellular microenvironments is crucial for bioanalysis, biomedicine, immunology, and related drug discovery. In the past few decades, numerous animal-derived bioactive ingredients, including reactive oxygen species (H_2O_2 , $^1\text{O}_2$, and HOCl),^{7–10} reactive sulfur species (H_2S , SO_2 , GSH, and Cys),^{11–15} reactive nitrogen species (HNO , NO , and NO_2),^{7,16–18} acetylcholine (ACh),^{19–21} adenosine triphosphate (ATP),^{22–24} dopamine,^{19,25–27} and histamine,^{19,28–30} have been elaborately monitored by exploiting fluorescent imaging techniques. This process has achieved an improved understanding of various pathological and physiological processes that potentially benefit diagnosis and treatment of animal diseases. However, detecting plant-derived signal molecules using fluorescent probes is also a key topic and a huge challenge for scientists. Salicylic acid (SA), a vital plant-derived defensive signal molecule, can activate global transcriptional reprogramming to express a network of prominent pathogenesis-related proteins in uninoculated tissues against secondary infections from invasive microorganisms systemically.^{31,32} This resistance can spontaneously spread throughout tissues of the whole plant; this mechanism is called systemic

acquired resistance (SAR).^{33–35} Through this process, the plant can obtain long-term protection in response to pathogen challenges controlled by the innate immune system of the plant. This process is considered the most economical and ideal strategy for crop protection. Many proofs show that the expression level of endogenous SA is enhanced in local and systemic tissues after the initial invasion of offensive pathogens, thus leading to the launch of SA-triggered signaling pathways.^{32,36–38} Therefore, monitoring the dynamic fluctuations of SA in subcellular microenvironments can advance our understanding of different physiological and pathological functions during the SA-induced SAR mechanism and benefit the discovery and development of novel immune activators that contribute to crop protection. This outcome will not only secure agricultural production and food supply but also provide an ideal approach to reducing the usage of traditionally harmful pesticides, thereby eliciting the concern of people. However, the accurate, dynamic, and real-time detection of SA has achieved considerable challenges for technological restraints and significant interferences from SA analogues. These distractors are methyl salicylate, 4-hydroxybenzoic acid, 2-methylbenzoic acid, 3-hydroxybenzoic acid, benzoic acid, and catechol; thus, few chemosensors can monitor SA in

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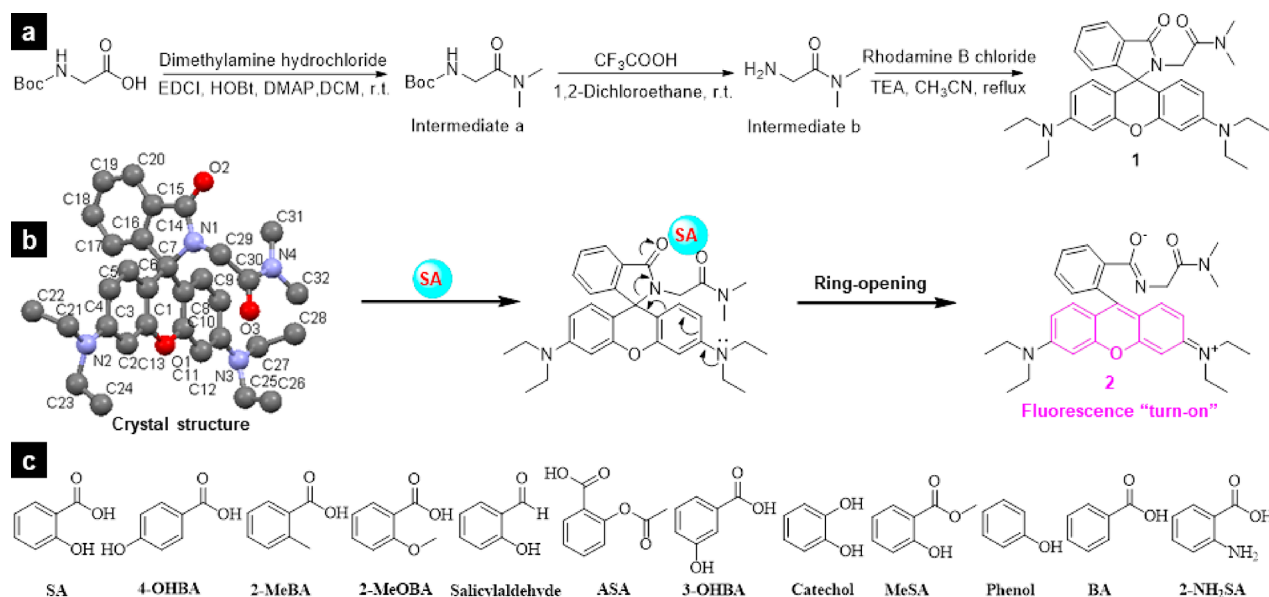


Figure 1. (a) Synthetic route for probe 1, (b) proposed mechanism for sensing SA using probe 1, and (c) chemical structures of SA analogues.

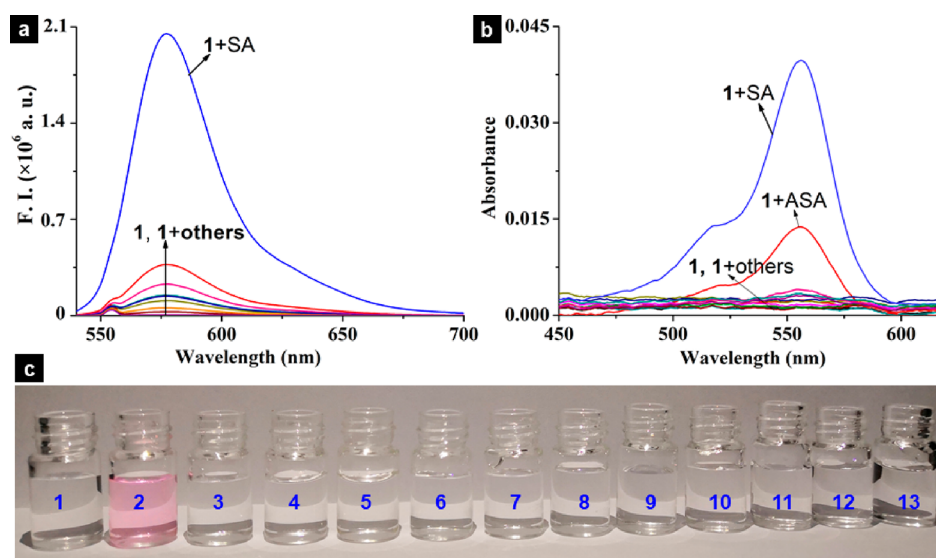


Figure 2. (a) Fluorescence and (b) absorption spectra of probe 1 (10 μM) after adding 100 μM SA and its analogues, with $\lambda_{\text{ex}} = 554$ nm, slits of 2/2 nm, and MeOH-H₂O (9:1, v/v), and (c) photographs of probe 1 (10 μM) after adding 20 equiv of SA or its analogues: (1) 1, (2) 1 + SA, (3) 1 + 4-OHBA, (4) 1 + 2-MeBA, (5) 1 + 2-MeOBA, (6) 1 + salicylaldehyde, (7) 1 + ASA, (8) 1 + 3-OHBA, (9) 1 + catechol, (10) 1 + MeSA, (11) 1 + phenol, (12) 1 + BA, and (13) 1 + 2-NH₂BA in MeOH-H₂O (9:1, v/v).

vitro^{39–43} and *in vivo*.⁴⁴ Thus, a campaign to discover and develop fluorescent probes is urgently required to monitor biological signaling molecule SA *in situ* and *in vivo*.

Rhodamine-based chemosensors, which exhibit versatile and broad detection windows, have been extensively studied and exploited for biological imaging in subcellular microenvironments.^{7,18,45–47} Numerous research results have revealed that fluorescent probes containing rhodamine skeletons display exhilarating privileges, such as excellent sensitivity and selectivity,^{48–51} real-time detection feature,^{52–54} and low detection limits,^{55,56} toward detected objects. In our previous work, a rhodamine-based sensor owning a *N,N*-dimethylhydrazinecarboxamide moiety could monitor SA in living animal cells.⁴⁴ To pursue a high-performing probe and further explore the substantial application in detecting SA on plants, herein, a simple and non-fluorescent rhodamine derivative 1 bearing a

flexible 2-amino-*N,N*-dimethylacetamide pattern was rationally fabricated to monitor SA *in vitro*, in living cells and plant callus tissues (Figure 1a). Within this framework, a free-rotated methylene bridging linker was introduced to facilitate the formation of a 1-SA binary complex in a relatively relaxed pocket promoted by hydrogen bonding. The newly formed combinations led to rearranging the electronic property of the whole molecule and subsequently caused the ring opening of the spirolactam structure accompanied by producing fluorescence for distinguishing SA (Figure 1b). An achromatous solution containing probe 1 immediately changed into pink after adding SA, thus revealing that a colorimetric probe for sensing SA with a naked-eye detection characteristic should be developed.

MATERIALS AND METHODS

Instruments and Chemicals. All chemical reagents and analytical reagent (AR) solvents were acquired from commercial suppliers and used without further purification. Distilled water was used throughout. Nuclear magnetic resonance (NMR) spectra were obtained using a JEOL-ECX 500NMR spectrometer. High-resolution mass spectroscopy (HRMS) spectra were performed on a Thermo Scientific Q Exactive mass spectrometer. Fluorescence spectra were performed by a Fluoromax-4 spectrofluorometer. Ultraviolet–visible (UV–vis) spectra were recorded by a TU-1900 spectrophotometer (Beijing Purkinje General Instrument Co., China). Fluorescence imaging was performed using a Nikon ECLIPSE Ti-S fluorescence microscope.

The stock solution of probe **1** (1.0×10^{-3} M) was prepared in methanol. Analyte stock solutions (1.0×10^{-2} M) of SA and its distractors were prepared in methanol, respectively. The preparation of samples for UV–vis and fluorescence detections follows the operations: 0.1 mL of stock solution of probe **1** was added to a 10 mL volumetric flask, and then certain amounts of detection objects were added; the mixture was filled up to 10 mL with the associated solvents; and finally, the related spectra were obtained after incubating the mixture for 10 min.

RESULTS AND DISCUSSION

Rhodamine-based chemosensor **1** was prepared, as illustrated in Figure 1a. The starting material Boc-glycine was reacted with dimethylamine through typical condensation to provide an intermediate a, which was then treated with trifluoroacetic acid to remove the protective group (–Boc) to realize an intermediate b. Finally, the target probe **1** was fabricated through a simple cyclization reaction between the intermediate b and rhodamine B chloride. The final molecular framework was confirmed using ^1H and ^{13}C NMR, HRMS, and its related crystal structures (Figures S1–S6 and Table S1 of the Supporting Information). Fluorescence and UV–vis spectra were performed to evaluate the detection competence of probe **1** toward SA and its distractors (Figure 1c and panels a and b of Figure 2). A methanol solvent and a methanol–water solution (9:1, v/v) were rationally selected for monitoring SA based on the fluorescence enhancement folds at 578 nm upon exposing the **1**–SA complex in various test fluids (Figures S7 and S8 of the Supporting Information). Probe **1** displayed superior selectivity and sensitivity to those of other SA analogues in detecting SA from the fluorescence spectrum (Figure 2a). These SA analogues included 4-hydroxybenzoic acid, *o*-methylbenzoic acid, *o*-methoxybenzoic acid, salicylaldehyde, acetylsalicylic acid (ASA), 3-hydroxybenzoic acid, catechol, methyl salicylate, benzoic acid, and anthranilic acid. The significantly produced fluorescent intensity triggered by adding SA revealed that SA can ring open the spirolactam pattern probably promoted by hydrogen-bonding interactions. This result was consistent with the newly generated UV absorption peak depicted in Figure 2b, thereby confirming that probe **1** can distinctively monitor SA. A pink color was only observed after adding SA (Figure 2c), thus suggesting that a colorimetric sensor was unexpectedly acquired for distinguishing SA with a naked-eye detection feature. Definitely, probe **1** showed improved detection behavior compared to our previously reported probe (Figure S9 of the Supporting Information).

To validate the selectivity and anti-interference functions further, competition assays should be conducted by subsequently adding SA to the premixed solution containing probe **1** with diverse interfering ingredients. Fluorescent intensity at

578 nm was substantially elevated only by the subsequent stimulation from SA, thus manifesting that the designed probe **1** possesses a superior anti-interference characteristic for recognizing SA (Figure 3). Moreover, a 1:1 binding

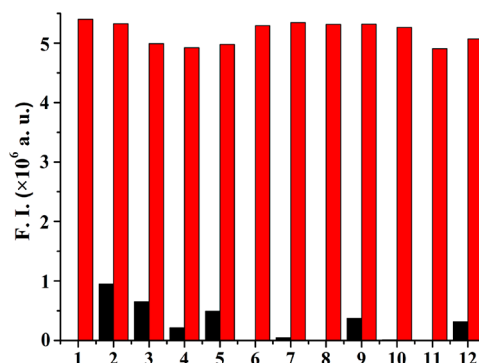


Figure 3. Competition experiments for adding 600 μM SA into the premixed solution consisting of probe **1** (10 μM) and various SA analogues (600 μM): (1) blank, (2) ASA, (3) 2-MeOBA, (4) 4-OHBA, (5) 2-MeBA, (6) catechol, (7) 2-NH₂BA, (8) MeSA, (9) BA, (10) phenol, (11) salicylaldehyde, and (12) 3-OHBA. Black bars, fluorescence intensity for probe **1** with SA analogues at 578 nm; red bars, after adding SA into the premixed solution containing probe **1** with SA analogues. Experimental conditions: λ_{ex} 554 nm; slits, 2/2 nm; and MeOH–H₂O, 9:1 (v/v).

stoichiometry between probe **1** and SA was revealed by Job's plot experiment (Figure S10 of the Supporting Information), and the binding constant was calculated as $4.31 \times 10^3 \mu\text{M}^{-1}$ (Figure S11 of the Supporting Information). The concentration-dependent titration assay demonstrated that the fluorescent intensity at 578 nm increases gradually after adding different dosages of SA. Consequently, a favorable linear detection within 10–100 μM was detected for a potential quantitative study (Figures S12 and S13 of the Supporting Information). Furthermore, the related detection limit was obtained as 1.0 nM by testing the signal-to-noise ratio. Given the above-mentioned results, a prospective probe that can selectively monitor SA *in vitro* in a naked-eye detection manner was discovered.

^1H NMR spectra were performed to explain the possible mechanism for monitoring SA in a concentration-dependent manner (Figure 4). The protons (1', 2', 3', and 4') that belong to the benzene ring of SA exhibited large chemical shifts to high fields because the molar ratio was 1:1 (Tables S2 and S3 of the Supporting Information). In contrast, further increasing the amount of SA (2 or 3 equiv) resulted in a reduced variation on the chemical shift, as illustrated by the corresponding changes in protons 1', 2', 3', and 4': –0.03, –0.08, –0.05, and –0.06 ppm for 1 equiv of SA and –0.02, –0.06, –0.04, and –0.05 ppm for 3 equiv of SA. This phenomenon might contribute to insufficient binding sites offered by probe **1** with excess SA molecules directed by hydrogen bonding. In contrast, the protons in probe **1** provided gradually increased chemical shifts by elevating the abundance of SA. The protons (11) of the methylene group showed significant low-field shifts with the changed values of 0.08 ppm for 1 equiv of SA, 0.11 for 2 equiv of SA, and 0.13 ppm for 3 equiv of SA. This finding might be due to the increased shielding effect from excess SA molecules promoted by hydrogen bonding. This action simultaneously led to rearranging the global electrons of probe **1**. The protons at the 5–10 positions presented

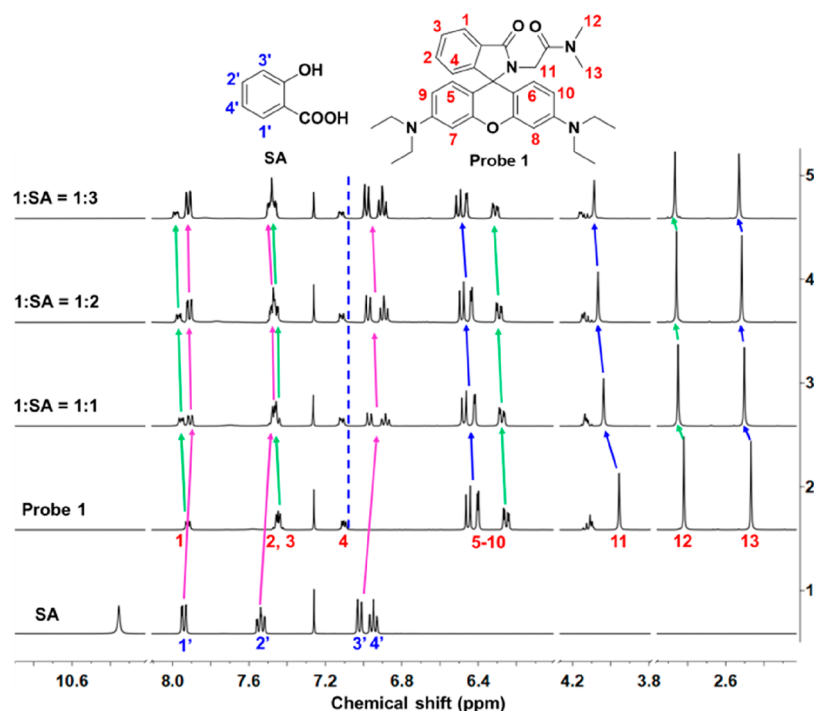


Figure 4. Partial ^1H NMR spectra for different molar ratios of probe 1 with SA (500 MHz, CDCl_3).

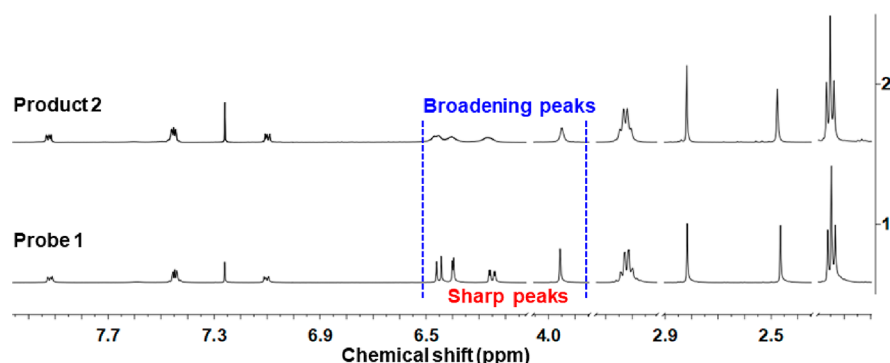


Figure 5. Comparison of partial ^1H NMR spectra of the new product 2 and probe 1 (500 MHz, CDCl_3).

considerable low-field shifts, which might be attributed to the launch of ring opening the spirolactam structure. We successfully obtained the newly fluorescent product 2 by the common separation on a silica gel column, and its molecular structure was confirmed by ^1H NMR and HRMS (Figure 5 and Figures S14 and S15 of the Supporting Information), in which broadening peaks were observed at 5–10 and 11 positions in comparison to those of probe 1 with sharp peaks. This finding further verified that SA detection was based on a fluorescence “turn-on” mode, thereby verifying the proposed mechanism (Figure 1b).

To explore the *in vivo* detection capability of probe 1 toward SA, A549 cell lines and plant callus tissues of *Malus asiatica* Nakai were randomly selected for SA imaging. For animal cell imaging, $10\ \mu\text{M}$ probe 1 was added to the prepared A549 cells and incubated for 0.5 h at $37\ ^\circ\text{C}$. These cells were washed by a phosphate-buffered saline (PBS) buffer (0.01 M, pH 7.4) thrice and subsequently treated with $100\ \mu\text{M}$ SA for 1 h at $37\ ^\circ\text{C}$. Finally, the cells were washed with PBS buffer thrice before imaging. Red fluorescent cells (Figure 6) were only observed by incubating the cells with probe 1 and SA, thereby indicating

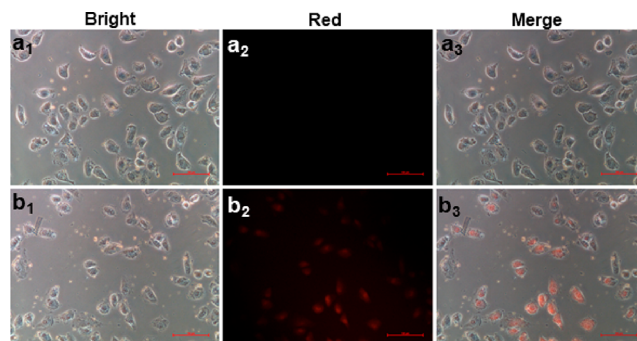


Figure 6. Fluorescence imaging of probe 1 relative to SA in living A549 cells: (a₁) bright field and (a₂) fluorescence images of A549 cells incubated using $10\ \mu\text{M}$ probe 1 for 30 min at $37\ ^\circ\text{C}$, (a₃) merged image of a₁ and a₂, (b₁) bright field and (b₂) fluorescence image of A549 cells incubated with $10\ \mu\text{M}$ probe 1 and $100\ \mu\text{M}$ SA for 1 h at $37\ ^\circ\text{C}$, and (b₃) merged image of b₁ and b₂. Red channel, 510–560 nm.

that probe 1 could penetrate the cell membrane and detect SA in animal cells. Moreover, we found that probe 1 showed low

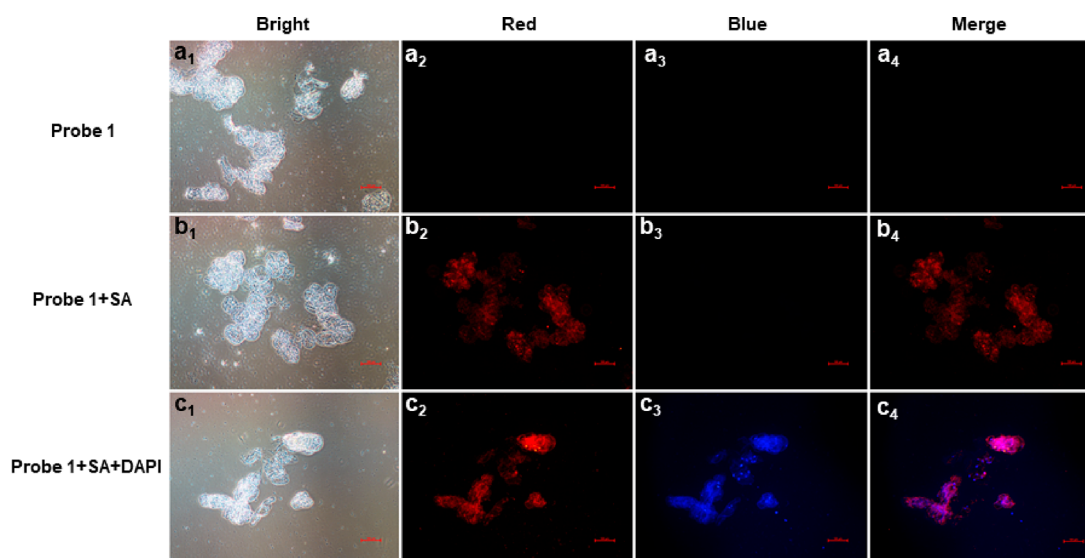


Figure 7. Fluorescence imaging of probe 1 relative to SA in living *M. asiatica* Nakai callus: (a₁) bright field and (a₂ and a₃) fluorescence images of callus tissues incubated using 10 μ M probe 1 for 25 days at 25 $^{\circ}$ C, (a₄) merged image of a₂ and a₃, (b₁) bright field and (b₂ and b₃) fluorescence images of callus tissues incubated using 10 μ M probe 1 and 100 μ M SA for 30 days at 25 $^{\circ}$ C, (b₄) merged image of b₂ and b₃, (c₁) bright field and (c₂ and c₃) fluorescence images of callus tissues incubated with 10 μ M probe 1 and 100 μ M SA for 30 days at 25 $^{\circ}$ C, and (c₄) merged image of c₂ and c₃. Red channel, 510–560 nm; blue channel, 330–380 nm.

toxicity toward A549 cells monitored by 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) assays (Figure S16 of the Supporting Information). For plant callus tissue imaging, the corresponding incubation time for probe 1 (10 μ M) and SA (100 μ M) in solid agar medium was 25 and 30 days before imaging. A strongly emerging fluorescence located at callus tissues (Figure 7) was only presented in the samples containing probe 1 and SA, thereby further confirming that probe 1 can also monitor SA in plant callus tissues (Figure 7b). Moreover, the localization of coloring was mainly in the cytoplasm (Figure 7c). Thus, probe 1, which possesses promising applications in SA signaling *in vitro* and *in vivo*, should be further developed.

In summary, a novel non-fluorescent rhodamine derivative 1 possessing a flexible 2-amino-*N,N*-dimethylacetamide pattern was designed and prepared to monitor SA. During this process, a probable mechanism for detecting SA was recommended. A binary complex 1–SA could form directed by hydrogen-bonding interaction and subsequently rearranging the electronic property of the whole molecule. This occurrence would cause the ring opening of the spirolactam structure along with producing fluorescence for distinguishing SA. This outcome was further confirmed by fluorescence, UV–vis, and 1 H NMR spectra. The results showed that probe 1 can selectively distinguish SA *in vitro* with the detection limit of 1 nM. A favorable linear detection ranging from 10 to 100 μ M was obtained for the potential quantitative study of SA. Probe 1 could be exploited to detect SA in living cells and plant callus tissues sensitively. Given these promising performances and applications, we anticipate that this probe should be further studied in monitoring SA in the plant kingdom and can launch the discovery of fresh immune activators for crop protection.

■ ASSOCIATED CONTENT

Supporting Information

The Supporting Information is available free of charge at <https://pubs.acs.org/doi/10.1021/acs.jafc.9b06771>.

Preparation of the intermediates and probe 1, NMR and HRMS spectra, Tables S1–S3, and Figures S1–S16 (PDF)

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Notes

The authors declare no competing financial interest.

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