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A H-bond strategy to develop acid-resistant photoswitchable rhodamine spirolactams for super-resolution single-molecule localization microscopy†

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Rhodamine spirolactam based photoswitches have been extensively applied in super-resolution single-molecule localization microscopy (SMLM). However, the ring-opening reactions of spirolactams are cross-sensitive to acid, limiting their photoswitch use to neutral pH conditions. In addition, the ring-closing reactions of spirolactams are environment-sensitive and slow (up to hours), virtually making rhodamine spirolactams caged fluorescent dyes instead of reversible photoswitches in SMLM. Herein, by introducing hydrogen bonds to stabilize spirolactams, we report a series of acid-resistant rhodamine spirolactams with accelerated ring-closing reactions from fluorescent xanthyliums to non-fluorescent spirolactams, endowing them with good photoswitchable properties even in acidic environments. By further substitution of 6-phenylethynyl naphthalimide on the spirolactam, we shifted the photoactivation wavelength into the visible region (>400 nm). Subsequently, we have successfully applied these dyes in labeling and imaging the cell surface of *Bacillus subtilis* at pH 4.5 using SMLM.

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Introduction

Super-resolution fluorescence microscopy breaks the diffraction limit and achieves nanometre-scale resolution.¹ The fundamental principle underlying this technique relies on the “dark” and “fluorescent” pair states of fluorescent markers, enabling temporal separation of adjacent molecules in a sub-diffraction-sized region. The requirements for both dark/emitting state-switching and outstanding emission properties (including high brightness and photostability), particularly for super-resolution single-molecule localization microscopy (SMLM),² have driven the creation of improved photoswitchable

fluorescent probes *via* two means: combining a photochromic moiety with excellent fluorophores³ and incorporating fluorescent fragments in photochromic compounds.⁴

Among various photoswitchable fluorophores, rhodamine spirolactams have attracted considerable research interest, owing to their excellent photophysical properties and biocompatibility. These dyes were widely applied in constructing various chemosensors for pH, metal ions and bioactive molecules, based on ring-opening reactions from colorless and non-fluorescent spirolactams to colored and strongly fluorescent xanthyliums.⁵ Since their photoswitching properties were first reported in 1977, rhodamine spirolactams have been used as photochromic materials.⁶ Recently, owing to dark/emitting state switching, rhodamine spirolactams have also been successfully deployed in SMLM.⁷ While many exciting super-resolution imaging applications have been demonstrated,⁸ two significant challenges limit the application of rhodamine spirolactams. First, rhodamine spirolactams are strictly limited to neutral pH environments, due to their considerable acid-activated switching to fluorescent xanthyliums (the open form) across a wide acidic pH range (usually 2.0 < pH < 6.5) (Fig. 1a).⁵ As these dyes enter an acidic environment (such as in lysosomes (pH 4.5–6.0), endosomes (pH 5.0–6.5), or locations next to proton-donating groups in proteins), acid-activation becomes substantial, leading to the disablement of photo-activation. Yet, low pH or acidic environments are ubiquitous in

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biological systems. Intracellular acidification is associated with many critical diseases (such as cancer).⁹ Second, environmental changes greatly alter the lifetime of the open form, which ranges from milliseconds to hours.^{6,7a,c} With a long “open form” lifetime in aqueous solutions, photobleaching reactions could considerably compete with thermal fading ring-closing reactions, compromising the photostability of rhodamine spirolactams.^{6a,10} Moreover, on the timescale of seconds or longer, the photoswitching reversibility of rhodamine spirolactams is virtually negligible for image acquisition purposes; these dyes effectively behave as typical caged fluorescent compounds: they are activated, imaged, and then bleached (Fig. 1a).^{7a} By introducing a lag-time in between frames to allow relaxation, these dyes could be used as localized markers several times. In a landmark paper, a spontaneously blinking fluorophore based on an intramolecular spirocyclization reaction was reported to increase reversible photoswitching cycles and allow long-term dynamic studies in live cells.¹⁰ In a recent study, the photo-switch of rhodamine spirolactam was combined with fluxionality to enable long time-lapse SMLM imaging in live cells.⁷¹ However, these fluorophores were still acid sensitive or pH-dependent, limiting their application to neutral pH environments. Consequently, dedicated photoswitchable rhodamine spirolactams with acid immunity and short “open form” lifetime are required for SMLM.

Herein, we report a series of acid-resistant rhodamine spirolactams with a dedicated photoswitchable property and short “open form” lifetime. These characteristics are attributed to the

substitution of 3-amino groups in the carboxyphenyl ring and associated intramolecular hydrogen bonding in spirolactams (Fig. 1b). Inspired by Moerner's work,^{7c} we further extended the photoactivation wavelength of our dyes into the visible region (>400 nm), *via* conjugation with 6-phenylethynyl naphthalimide. The excellent acid-resistance and visible-light photoswitching make our rhodamine spirolactams suitable for SMLM imaging, and more importantly, generate reliable fluorescence signals even in acidic environments.

Results and discussion

Molecular design strategy

Our strategy of introducing intramolecular hydrogen bonding to increase the acid-resistance of spirolactams and shorten the “open form” lifetime originated from three facts: (1) the acid-activated ring-opening reaction is initiated with the protonation of the carbonyl oxygen (Fig. 1a);¹ (2) the resulting ring opening reaction is associated with a rotation of the amide group (–CONR) with respect to the phenyl ring (Fig. 1a, S1; Tables S1 and S2†); (3) introducing an electron-donating group into spirolactams reduces the driving force of ring-opening reactions.¹¹ We anticipated that the intramolecular hydrogen bonding would reduce the reactivity of carbonyl oxygen with exterior acid and lock the spirolactam group. In addition, both the suppressed rotation of the amide group and the incorporation of an electron-donating group contributed to the stabilization of the closed form. Accordingly, we synthesized



Fig. 1 (a) The switch of rhodamine spirolactams from the non-fluorescent closed form to fluorescent open form activated by light or acid. (b) The intramolecular hydrogen bonding enables acid-resistant photoswitchable rhodamine spirolactams. (c) The designed rhodamine spirolactams without and with intramolecular hydrogen bonding.



compounds **P1–P6** to verify our hypothesis (Fig. 1c and Schemes S1–S3†). Compounds **P4** and **P5** possessed 3-substituted NHR groups which were expected to form intramolecular hydrogen bonds with the lactam moiety, while the rest served as control compounds without intramolecular hydrogen bonds.

Characterization of the acid-resistance of rhodamine spirolactams

We first examined the acid resistance of **P1–P6**. With the addition of CF_3COOH , a new emission band centered at 560 nm in the absorption spectra of **P1–P3** and **P6** was observed (Fig. 2a, b, S2 and S3†). Accordingly, the solutions of **P1–P3** and **P6** displayed noticeable color changes from colorless to pink, indicating the acid-controlled xanthylium formation. In stark contrast, the UV-vis absorption spectra of **P4** and **P5** did not show obvious changes in the visible region, and the solutions almost remained colorless (Fig. 2a, b and S2†). These results indicated that the fluorescence switching of **P4** and **P5** was far less acid-sensitive, while that of **P1–P3** and **P6** was acid-sensitive.

Verification of the H-bond mechanism

We next conducted NMR and crystallographic experiments to examine the existence of intramolecular hydrogen bonding (Fig. 2c, d and Table S3†). ^1H NMR spectra show large downfield shifts of the resonance of NH protons in **P4** (5.30) and **P5** (6.75), compared with those in **P2** (3.83) and **P3** (3.83) (Fig. 2c). The crystal structure of **P4** directly revealed strong intramolecular hydrogen bonding between NH and carbonyl oxygen (distance: 2.347 Å; angle: 128.61°) (Fig. 2d), which explained the downfield shift of NH in **P4**. Though we failed to obtain the crystal structure of **P5**, we believed that the much larger downfield shift of NH in **P5** indicated even stronger intramolecular hydrogen bonding in **P5**. According to the absorption and emission spectral evolution of **P4** and **P5** (after the addition of CF_3COOH), **P5** was more acid-resistant than **P4**, which should be ascribed to the stronger intramolecular hydrogen bonding (Fig. S2–S4†). In addition, we also noticed weak intramolecular $\text{CH}\cdots\text{O}$ interactions in **P6** (distance: 2.249 Å; angle: 130.80°). The above results clearly demonstrate that intramolecular hydrogen bonding plays a key role in enhancing the acid-resistance of rhodamine spirolactams.

We also performed density functional theory (DFT) calculations on **P1–P6** to rationalize the hydrogen-bond induced acid-resistance. As expected, optimized molecular geometries show that intramolecular hydrogen bonds are present only in **P4** and **P5** (Fig. S5†). Moreover, by modelling eight representative protonation configurations for each compound to fully simulate their responses in an acidic aqueous environment (Fig. S6†), our results show that among the control compounds, fluorescent open-ring structures possess significantly lower Gibbs free energy than non-emissive closed-ring structures by up to 0.19 eV (Table S4 and Fig. S7–S12†). Therefore, DFT calculations suggest that the emissions of these control compounds could easily turn on in an acidic environment. In contrast, the closed-ring structures of **P4** and **P5** are relatively more stable than

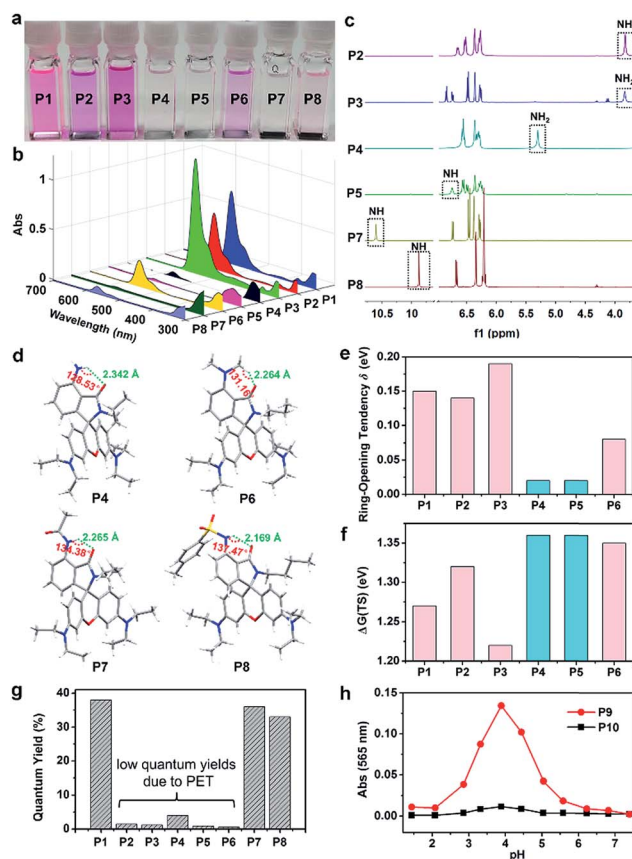


Fig. 2 (a) Photograph and (b) UV-vis absorption spectra of **P1–P8** in $\text{CH}_2\text{Cl}_2/\text{CH}_3\text{OH}$ (9/1, v/v), two hours after the addition of 2.3 μL of CF_3COOH (1000 equiv.). (c) Local ^1H NMR spectra of **P2–P5** and **P7–P8**. (d) Single-crystal structures and intramolecular hydrogen bonds of **P4**, **P7** and **P8** and weak $\text{CH}\cdots\text{O}$ interactions in **P6**. (e) Calculated ring-opening tendencies of **P1–P6** in acidic solution. (f) Calculated energy barriers during the ring-opening process of neutral **P1–P6**. (g) Fluorescence quantum yields of **P1–P8** in $\text{CH}_2\text{Cl}_2/\text{CH}_3\text{OH}$ (9/1, v/v) solution. (h) The absorption intensities at 565 nm of **P9** and **P10** as a function of pH in water.

those of **P1–P3** and **P6** by ~ 0.12 eV, suggesting that these structures are resistant to acid-activation. We further calculated the difference in Gibbs free energy between the most stable closed-ring and open-ring protonated structures (δ) to quantify the tendency of ring-opening in **P1–P6**. Herein, a large δ corresponds to a strong thermodynamic driving force for ring-opening reactions to proceed. Indeed, our results showed that **P4** and **P5** exhibited a significantly lower ring-opening tendency than **P1–P3** and **P6** did (Fig. 2e). We also computed the energy barriers during the ring-opening reactions of **P1–P6** (Table S5 and Fig. S13–S19†). Our results show that the barriers in **P4** and **P5** are higher than those of the control compounds, due to the presence of intramolecular hydrogen bonds (Fig. 2f). All these computational results rationalize that **P4** and **P5** possess significantly enhanced acid-resistance towards fluorescence activation. At the same time, we expected that **P4** and **P5** remained photoswitchable, given the high energy of photo-activation light.



Improving quantum yields and water solubility of rhodamine spirolactams

Unfortunately, the quantum yields of **P2–P6** are low (Fig. 2g), possibly quenched by photo-induced electron transfer (PET) from the 3-aminophenyl moiety to the xanthene scaffold. To enhance brightness and preserve acid-resistance, we synthesized **P7** and **P8** (Scheme S4†) with 3-acetamido and 3-toluenesulfonamide substituents to inhibit PET, respectively. As a result, the quantum yields of **P7** and **P8** were significantly improved (Fig. 2g). At the same time, **P7** and **P8** remained acid-resistant (Fig. 2a and b). ¹H NMR analysis and the single crystal structures of **P7** and **P8** clearly showed the existence of intramolecular hydrogen bonds (Fig. 2c and d). Notably, the stronger intramolecular hydrogen bonding in **P7** than that in **P8** resulted in stronger acid-resistance of **P7** than that of **P8** (Fig. S2–S4†). Owing to its high brightness and excellent acid-resistance, we chose **P7** for further derivations and applications.

Subsequently, we synthesized water-soluble compounds **P9** and **P10** with polyethylene glycol (PEG) groups to verify the applicability of our strategy in water (Scheme S5†). As shown in Fig. 2h and S20,† the UV-vis absorption and emission intensities of the open form of **P9** substantially increased when the pH was decreased from 7.40 to 3.89, and then decreased when the pH was decreased from 3.89 to 1.44. In contrast, **P10** did not show any noticeable absorption and fluorescence bands in the visible region in the pH range from 1.44 to 7.40 (Fig. 2h and S20†). These results indicated that our strategy to increase the acid-resistance of rhodamine spirolactams worked well in aqueous solutions and was suitable for biological systems.

Evaluation of open form lifetime

We next evaluated the lifetime of the open forms *via* laser flash photolysis (Table S6 and Fig. S21†). It's reported that the lifetime of the open form was greatly affected both by the environment and molecular structures.^{6,7a,c} The open form has a lifetime of a few milliseconds in polar solvents.⁶ In polyvinylalcohol, the lifetime of the open form was reported to lengthen to hours.^{7a} By introducing aromatic substituents on the lactam nitrogen, the open isomer with the charge at the amide nitrogen atom can be stabilized and then it has a longer lifetime.^{6b,7a,c} In our experiments, the lifetime of **P1** *vs.* **P7** in CH₂Cl₂ was too short to be examined (Table S6†). But in dioxane, the lifetime was detected to be in the range of 60 ns. Importantly, we found that the intramolecular hydrogen bonding could shorten the lifetime of the open forms (62.5 ns for **P1** and 59.7 ns for **P7**). The effect of intramolecular hydrogen bonding on lifetime becomes more significant when an aromatic substituent is introduced on the lactam nitrogen. Compared with the 200 ms lifetime of *N*-phenyl-rhodaminelactame in dioxane,^{6a,b} the lifetime of the reference compound **P11** was much shorter, measured to be ~31.5 ns in the same solvent. These results indicated that the thermal fading processes in our dyes may compete with photobleaching.

In vitro bioimaging experiments

With this progress, we continued to synthesize **P12** and **P13** to validate our design strategy with *in vitro* bioimaging experiments (Fig. 3 and Scheme S7†). The “control” probe **P12** and the “experimental” probe **P13** are coupled with a morpholine moiety, which enables these probes to become protonated and accumulated in lysosomes. The pH value of lysosomes is in the range of 4.0–6.0, which provides a good platform to test the acid-resistance of rhodamine spirolactams. After incubation for 30 min, **P12** emitted strong fluorescence in MCF-7 cells (Fig. S22†). Subsequently, we performed co-localization experiments of **P12** (the red channel) with LysoTracker Green DND-26 (LTG; the green channel), a commercially available lysosome probe (Fig. 3a). The good overlap of fluorescence signals in the red and green channels demonstrates that **P12** was distributed in the lysosomes of MCF-7 cells. Overall, these results show that the acidic environment in lysosomes has activated intensive fluorescence in the control probe **P12**.

In contrast, after incubating MCF-7 cells with **P13** for 2 hours, we did not observe any fluorescence signals (Fig. S22†). To ensure the presence of **P13** in lysosomes, continuous UV light of 375 nm (~1 mW cm^{−2}) was applied to activate the fluorescence of **P13**. By increasing the illumination time, we noticed a gradual increase of fluorescence signals in the red channel (Fig. 3b). These signals overlapped with the fluorescence of the lysosome probe LTG in the green channel. It is thus

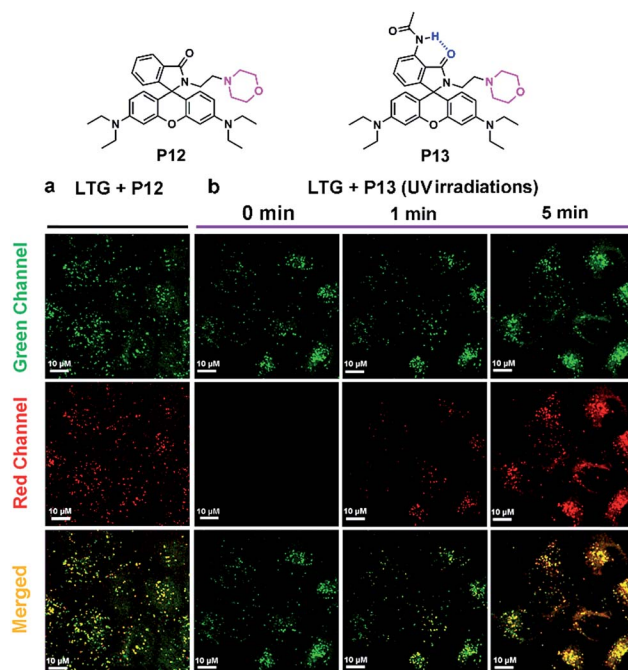


Fig. 3 Confocal images of MCF-7 cells stained with LTG (0.1 μM) and (a) **P12** (10 μM) or (b) **P13** (10 μM); green channel: excitation at 488 nm, emission collected from 500 to 550 nm; red channel: excitation at 561 nm, emission collected from 580 to 653 nm. Insets: chemical structures of **P12** and **P13**. The Pearson coefficients are high, 0.905 (LTG with **P12**) and 0.925 (LTG with **P13** activated by UV light for 5 min), respectively.

Photoactivation of rhodamine spirolactams using visible light

Encouraged by the success of **P13** in bioimaging applications, we moved on to extend the photoactivation wavelength of rhodamine spirolactams into the visible region for cell imaging. We thus synthesized **P14–P17** by conjugating a 6-phenylethynyl naphthalimide moiety to the spirolactam (Fig. 4a and Scheme S8[†]). The enlarged π -conjugation extends the UV-vis absorption spectra of **P14–P17** into the visible region, with a peak absorption wavelength at ~ 400 nm (Fig. 4b). It is thus possible to photoactivate these dyes with a 405 nm laser and read out the

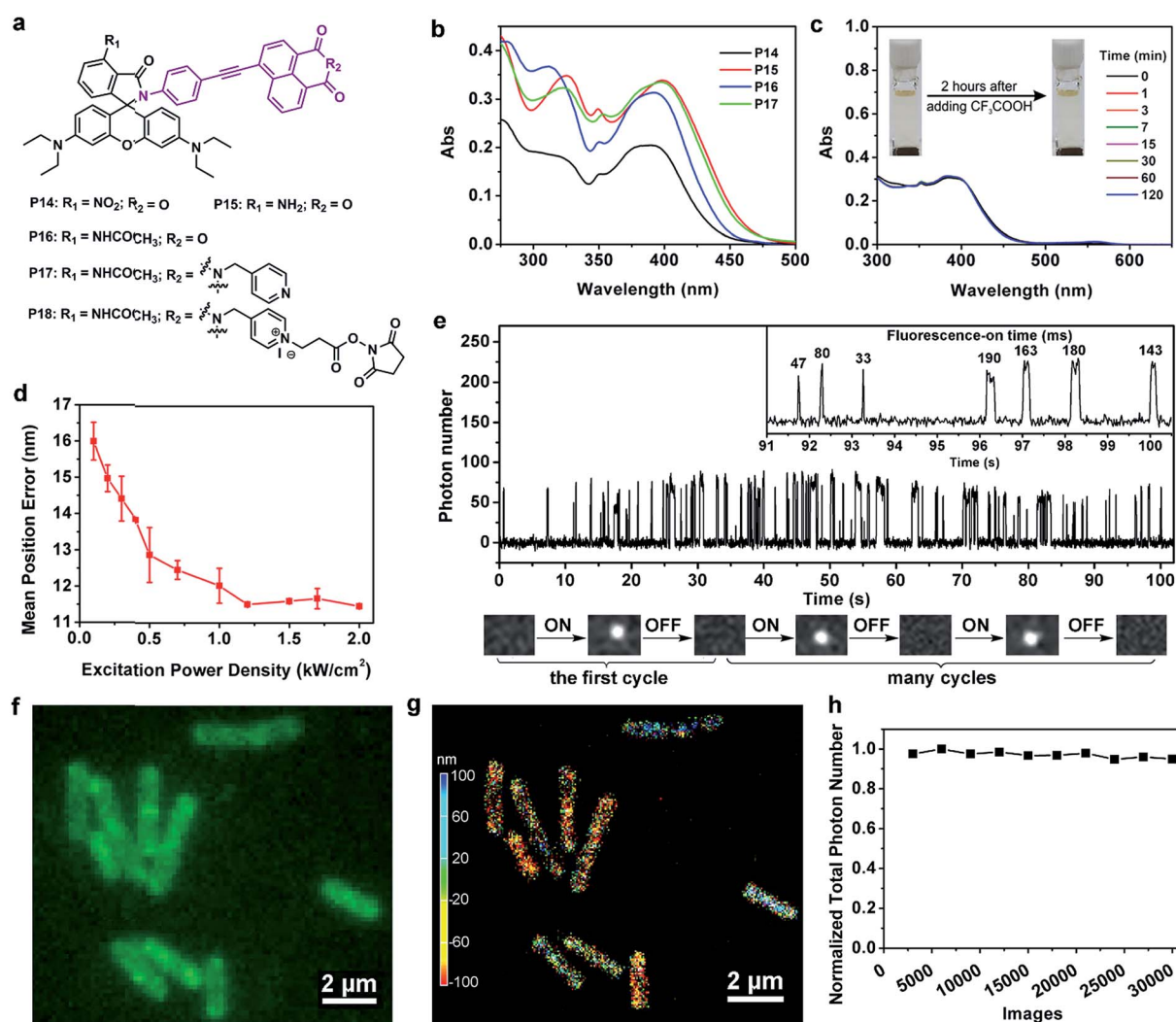


Fig. 4 (a) Chemical structures of **P14–P18**. (b) UV-vis absorption spectra of **P14–P17** (10^{-5} M) in DMSO. (c) Time-dependent UV-vis absorption spectra of **P17** (10^{-5} M) in $\text{CH}_2\text{Cl}_2/\text{CH}_3\text{OH}$ (9/1, v/v) before and after the addition of CF_3COOH (2.3 μL , 1000 equiv.); the insets show the photographs of **P17** before and two hours after adding CF_3COOH . (d) Single molecule localization precision of **P17** at various excitation laser power densities. (e) Single-molecule fluorescence time traces of **P18**. (f) Conventional diffraction-limited fluorescence image of live *Bacillus subtilis* cells stained with **P18** in pH 4.5 buffer solution showing average fluorescence over 30 000 frames (20 ms integration time per frame). (g) 3D-SMLM image of live *Bacillus subtilis* cells stained with **P18** in pH 4.5 buffer solution; the SMLM image contains ca. 5.7×10^5 localized positions, each plotted as a Gaussian with the full width at half-maximum (FWHM) in the X and Y direction. (h) Normalized total photon output in a set of 30 000-frame images.

fluorescence signals with an excitation laser of 561 nm. The visible photoactivation wavelength is critical for minimizing phototoxicity in live-cell imaging.

Among **P14**–**P17**, we selected **P17** for further experiments, foreseeing its excellent acid-resistance and good quantum yield. Indeed, **P17** remained colorless in acidic medium (Fig. 4c). We then doped **P17** into a PMMA film at a low concentration to characterize its single-molecule fluorescence properties. We applied a continuous 405 nm laser light (60 W cm^{-2}) to photoactivate **P17** (Fig. S24†), and a 561 nm laser light to generate fluorescence from activated **P17**. Our experiments showed that a laser excitation power of $\sim 1.2 \text{ kW cm}^{-2}$ at 561 nm led to a localization precision of the single molecule of $\sim 12 \text{ nm}$ in PMMA (Fig. 4d and S25†). Subsequent experiments showed that good imaging quality could be achieved by reducing the photoactivation light intensity at 405 nm down to 1.8 W cm^{-2} in a cellular environment.

Super-resolution imaging of *Bacillus subtilis* in acidic environments

To fully demonstrate the imaging applications of our photo-switchable dyes, we linked *N*-hydroxysuccinimide ester to **P17**, resulting in the formation of **P18** (Fig. 4a and Scheme S9†). Laser flash photolysis showed that the open form lifetime of **P18** was 105 ns in aqueous solutions with 5% DMSO (Table S6 and

Fig. S21†). In a single molecule fluorescence imaging experiment with a 561 nm laser (0.36 kW cm^{-2}) as an excitation source, **P18** labeled on a glass coverslip showed reversible fluorescence blinking ~ 100 times (Fig. 4e). Moreover, in this experiment the fluorescence-on time of **P18** increased to ~ 100 ms-scale, probably owing to the environmental sensitivity of the open form lifetime. For example, this lifetime of a rhodamine spirolactam increased from milliseconds in polar solvents to hours in polyvinyl alcohol.^{7a} The reversible blinking and good fluorescence signals inspired us to label **P18** on the membrane surface of Gram-positive *Bacillus subtilis* cells in PBS solution (pH 4.5), since these bacteria can maintain satisfactory activity in an acidic pH environment. We expected that **P18** could specifically label bacterial membranes for the following reasons: (1) the positively charged **P18** could bind with the anionic bacterial membranes through electrostatic interactions, and these interactions could further inhibit the diffusion of **P18** through cell membranes; (2) the NHS ester of **P18** could rapidly label the amino groups in the membrane proteins; and (3) the hydrophobic 1,8-naphthalimide group may intercalate into the membrane bilayer. We then imaged acid-resistant *Bacillus subtilis* cells, using a continuous 405 nm laser (1.8 W cm^{-2}) as photoactivation light and a 561 nm laser (1.2 kW cm^{-2}) as an excitation source. The fluorescence of **P18** exhibited an excellent photo-blinking property, allowing the reconstruction of the super-resolution image *via* single-molecule localization.

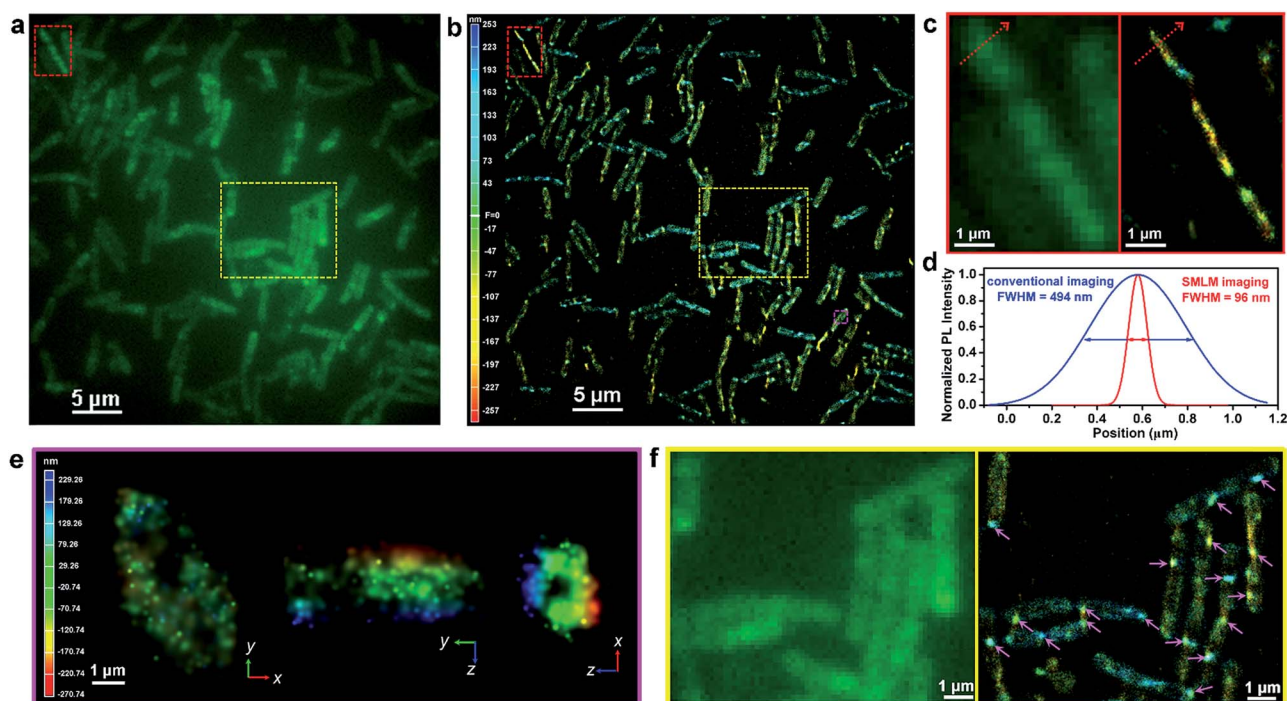


Fig. 5 (a) Conventional diffraction-limited fluorescence image and (b) 3D-SMLM image of live *Bacillus subtilis* cells stained with **P18** in pH 4.5 buffer solution; the SMLM image contains ca. 1.24×10^6 localized positions, each plotted as a Gaussian with the full width at half-maximum (FWHM) in the X and Y direction. (c) Comparison of the imaging resolution of the three adjacent *Bacillus subtilis* cells in the enlarged region, as highlighted by red rectangles. (d) Gaussian fittings to the normalized PL intensities across a randomly selected bacterium (cross-sectional analysis along the red arrow). (e) xy, yz, and xz views of the bacterial cell membranes shown in the pink square in (b), revealing the hollow cylindrical structure of the bacterial cell membrane. (f) Comparison of the imaging resolution in the enlarged region, as highlighted in yellow rectangles; the right SMLM image shows many fluid microdomains, as indicated by the pink arrows.



In comparison to a conventional diffraction-limited fluorescence image (Fig. 4f), the 3D-SMLM image of *Bacillus subtilis* displayed a greatly improved spatial resolution (Fig. 4g). Besides, owing to the low intensity of the 405 nm excitation laser, and slow photobleaching rates of the 561 nm laser (as a result of fast ring-close reactions and a short open-form lifetime), **P18** exhibited excellent photostability. The number of collected photons per 3000 frames (20 ms per frame) remained almost constant throughout our experiments of >10 min (Fig. 4h), demonstrating the potential of **P18** to perform long time-lapse nanoscopy imaging in future studies.

We continued to investigate the staining properties and labeling specificity of **P18**. As shown in Fig. 5a, b and Movie 1,[†] **P18** stained bacteria evenly with high density, and exhibited excellent photoblinking properties. Interestingly, the divisions of three *Bacillus subtilis* cells were clearly visible in a 3D-SMLM image (Fig. 5b and c), while they were barely resolved in a diffraction-limited fluorescence image (Fig. 5a and c). The full-width at half-maximum (FWHM) of a single *Bacillus subtilis* is ~96 nm in the super-resolution images, cf. ~494 nm in the conventional fluorescence images (Fig. 5d). Cross-sections of a single *Bacillus subtilis* highlighted in the pink box show a hollow structure from the *xz* view, indicating that our probes localized on the cell membranes of bacteria (Fig. 5e). The flipped 3D-SMLM image of bacterial cell membranes is further shown in Movie 2.[†] We thus demonstrated that **P18** enabled SMLM studies of the membranes of bacteria cells at nanometric resolution (without using additional targeting strategies). Moreover, we found that the membranes of most *Bacillus subtilis* bacteria did not show uniform fluorescence intensities in the 3D-SMLM image shown in Fig. 5b. The slight irregularity of the membrane staining likely indicates the presence of many fluid microdomains (as highlighted by pink arrows in the right image of Fig. 5f). These fluid microdomains had been well studied by several research groups.¹² However, these fluid microdomains were hardly resolved in conventional diffraction-limited fluorescence images (in the left image of Fig. 5f). Besides, we also discovered that the majority of these fluid microdomains appeared in certain areas, especially around cell poles.

Conclusions

In summary, this study presents an effective molecular engineering approach for developing acid-resistant rhodamine spirolactams with a dedicated photoswitchable property and short open-form lifetime. These unique characteristics resulted from intramolecular hydrogen bonding between the 3-amino substituents in the carboxyphenyl ring and the lactam moiety. Moreover, we have extended the photoactivation wavelengths of rhodamine spirolactams into the visible region (>400 nm), by conjugating a 6-phenylethynyl naphthalimide moiety to the spirolactams. The visible photoactivation and dedicated photoswitchable properties of our dyes make them highly reliable fluorescent probes for SMLM, as demonstrated by the successful labeling and imaging of *Bacillus subtilis* at pH 4.5 using 3D-SMLM. We believe that our work sheds light on rational

engineering of fluorescent properties of rhodamine dyes to facilitate the advancement of super-resolution imaging techniques.

Experimental section

Single-crystal structures

Single crystals of **P4**, **P6**, **P7** and **P8** that were suitable for X-ray structural analysis were obtained by slow evaporation from a chloroform/*n*-hexane solution at room temperature. Single crystal X-ray diffraction data were collected on an Xcalibur, Atlas, Gemini ultra-diffractometer using the ω -scan mode with a graphite-monochromator and Mo K α radiation. The crystal structures were solved with the Superflip¹³ structure solution program and refined with the SHELXL¹⁴ refinement package, as implemented in OLEX2.¹⁵ All H atoms were placed geometrically and treated with riding constraints, and displacement parameters were derived from the C atoms to which they were attached. All CH₂ groups had $U_{\text{iso}}(\text{H})$ values fixed at 1.2 times the U_{eq} value of the attached C atom. CH₃ groups were idealized as freely rotating groups, with $U_{\text{iso}}(\text{H})$ values fixed at 1.5 times the U_{eq} value of the attached C atom. H atoms on N atoms were located *via* Fourier difference map inspection and positionally refined isotropically with thermal parameters based upon the N atom to which they are bonded ($U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{N})$). These data can be obtained free of charge from the Cambridge Crystallographic Data Centre (CCDC reference numbers: 1582847, 1901370–1901372[†]).

Measurement of single molecule optical properties

P17 was immobilized in polymer films to investigate the effects of the excitation laser power density on (1) the photon output from dyes, (2) background noise, and (3) localization precision. Polymer films were prepared using 1% (by mass) solution of polymethyl methacrylate (PMMA) in toluene. The polymer films were doped with **P17** (10 nM) and then spin-cast onto argon-plasma-etched coverslips. After that, the coverslips were placed in the sample holder of an SMLM instrument for imaging. The intensity of the 405 nm photoactivation laser was adjusted to activate only a few molecules in each frame. The fluorescence images of the polymer film were acquired using an electron-multiplying charge-coupled device (EMCCD) camera at room temperature.

Sample preparation and data analysis for single-molecule fluorescence imaging experiment

Coverslips were cleaned by sonication for 10 min in each of the following: Milli-Q water, absolute ethyl alcohol, and acetone. The above coverslips were further boiled in a solution mixture of concentrated sulfuric acid and hydrogen peroxide (v/v, 7/3) for 1 h and then washed with Milli-Q water and dried using nitrogen. 20 μL of very dilute **P18** (10 nM) in DMSO solution was dripped on the dry coverslips. Then, **P18** with the *N*-hydroxysuccinimide ester adsorbed and reacted with the hydroxyl groups on the surface of the cleaned coverslip at a low density such that individual dye molecules could be clearly resolved.



Then the surface of the cleaned coverslip was rinsed with PBS (pH 7.4) to remove unbound **P18** before imaging. Finally, the coverslip was placed on the sample holder of the SMLM instrument and illuminated with 561 nm laser light (0.36 kW cm^{-2}). 5000–10 000 frames were recorded at a frame rate of 20 ms per frame. Single-molecule fluorescence time trace was calculated for a single molecule by analyzing the total integrated signal in a 2×2 pixel region centered on the single molecule during each frame of the movie. The number of photons was calculated as the total intensity count minus the background intensity count, which was converted to photon number using the calibrated parameter for electron multiplication and analog-to-digital conversion gain settings used during acquisition.

Cell culture and CLSM imaging

MCF-7 cells were grown on DMEM (Gibco) supplemented with 10% FBS (Gibco). In CLSM imaging, the MCF-7 cells were incubated with 1 mL of DMEM containing $10 \mu\text{M}$ **P11** (experimental group) or **P12** (control group) and the images were collected at 0, 30, 60 and 120 min by using an ANDORTM live cell imaging system. Then, the adherent cells were further incubated with 1 mL of fresh DMEM containing LysoTracker Green DND-26 (LTG) ($0.1 \mu\text{M}$) for another 30 min. The adherent cells were washed 3 times with PBS (pH 7.4) and utilized for CLSM imaging. In the experimental group, the CLSM images of MCF-7 cells were taken under 0–5 min continuous UV (375 nm) irradiation at 37°C .

Live bacteria sample preparation and labeling

A single colony of *Bacillus subtilis* was inoculated in 5 mL of Luria–Bertani (LB) medium with shaking at 37°C . Subsequently, 1 mL of the *Bacillus subtilis* suspension was inoculated in 10 mL of fresh LB media, which was incubated for 5 h with shaking (250 rpm) at 37°C to achieve mid-log phase growth. After the *Bacillus subtilis* had grown to the mid-log phase, 1 mL of the suspension was washed at least 3 times by centrifuging for 3 minutes at 10 000 RPM and resuspending the pellet in 1 mL of clean PBS (pH 4.5). To label the surface of *Bacillus subtilis*, 50 μL of **P17** in DMSO solution ($\sim 10 \text{ nM}$) was added slowly to the suspension of the *Bacillus subtilis* cells and left to incubate for 30 minutes. The unreacted **P17** was rinsed 6 times with PBS (pH 4.5) before imaging. Then, the cleaned *Bacillus subtilis* cells were resuspended in a small amount of PBS (pH 4.5) to produce a concentrated cell suspension. Finally, 2 μL of this cell suspension was deposited onto an agarose pad (1.5% (by mass) low melting point agarose in PBS) and mounted onto an Ar plasma etched glass slide (size: $24 \times 50 \text{ mm}$, thickness: 0.13–0.16 mm) and immediately used for the SMLM imaging.

3D-SMLM imaging and data processing

The super-resolution imaging was carried out using a Nikon N-STORM 5.0 super-resolution microscope system with a motorized inverted microscope ECLIPSE Ti2-E, a $100\times/\text{NA } 1.49$ oil immersion TIRF objective lens (CFI HP) and an ORCA-Flash 4.0 sCMOS camera (Hamamatsu Photonics K.K.). During super-resolution imaging, a 405 nm laser (1.8 W cm^{-2}) was

employed to photoactivate rhodamine spirolactams from ring-closed isomers to ring-opened isomers. At the same time, a 561 nm laser (1.2 kW cm^{-2}) was used as the excitation source to produce sparse emissions from ring-opened rhodamines. Repeating this cycle many times (*i.e.*, to collect $\sim 30\,000$ images with the sCMOS camera with an integration time of 20 ms per frame) allowed us to reconstruct a super-resolution image. Astigmatism was used in the 3D-imaging experiments. The software NIS-Elements Ar and N-STORM Analysis were used to analyze the collected images and computationally reconstruct the 3D-SMLM image.

Conflicts of interest

The authors declare no conflict of interest.

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Notes and references

- (a) S. W. Hell, *Science*, 2007, **316**, 1153; (b) A. von Diezmann, Y. Shechtman and W. E. Moerner, *Chem. Rev.*, 2017, **117**, 7244; (c) E. Betzig, *Angew. Chem., Int. Ed.*, 2015, **54**, 8034.
- G. Patterson, M. Davidson, S. Manley and J. Lippincott-Schwartz, *Annu. Rev. Phys. Chem.*, 2010, **61**, 345.
- (a) M. Heilemann, S. van de Linde, M. Schüttelpeiz, R. Kasper, B. Seefeldt, A. Mukherjee, P. Tinnefeld and M. Sauer, *Angew. Chem., Int. Ed.*, 2008, **47**, 6172; (b) J. B. Grimm, T. Klein, B. G. Kopeck, G. Shtengel, H. F. Hess, M. Sauer and L. D. Lavis, *Angew. Chem., Int. Ed.*, 2016, **55**, 1723.
- (a) O. Nevskiy, D. Sysoiev, A. Oppermann, T. Huhn and D. Wöll, *Angew. Chem., Int. Ed.*, 2016, **55**, 12698; (b) B. Roubinet, M. L. Bossi, P. Alt, M. Leutenegger, H. Shojaei, S. Schnorrenberg, S. Nizamov, M. Irie, V. N. Belov and S. W. Hell, *Angew. Chem., Int. Ed.*, 2016, **55**, 15429; (c) B. Roubinet, M. Weber, H. Shojaei, M. Bates, M. L. Bossi, V. N. Belov, M. Irie and S. W. Hell, *J. Am. Chem. Soc.*, 2017, **139**, 6611.
- (a) H. N. Kim, M. H. Lee, H. J. Kim, J. S. Kim and J. Yoon, *Chem. Soc. Rev.*, 2008, **37**, 1465; (b) X. Chen, T. Pradhan, F. Wang, J. S. Kim and J. Yoon, *Chem. Rev.*, 2012, **112**, 1910.
- (a) K. H. Knauer and R. Gleiter, *Angew. Chem., Int. Ed.*, 1977, **16**, 113; (b) H. Willwohl, J. Wolfrum and R. Gleiter, *Laser Chem.*, 1989, **10**, 63; (c) H. Montenegro, M. D. Paolo, D. Capdevila, P. F. Aramendia and M. L. Bossi, *Photochem. Photobiol. Sci.*, 2012, **11**, 1081; (d) K. Li, Y. Xiang, X. Wang, J. Li, R. Hu, A. Tong and B. Z. Tang, *J. Am. Chem. Soc.*, 2014, **136**, 1643.



- 7 (a) J. Fölling, V. Belov, R. Kunetsky, R. Medda, A. Schonle, A. Egner, C. Eggeling, M. Bossi and S. W. Hell, *Angew. Chem., Int. Ed.*, 2007, **46**, 6266; (b) V. N. Belov, M. L. Bossi, J. Fölling, V. P. Boyarskiy and S. W. Hell, *Chem.-Eur. J.*, 2009, **15**, 10762; (c) M. K. Lee, P. Rai, J. Williams, R. J. Twieg and W. E. Moerner, *J. Am. Chem. Soc.*, 2014, **136**, 14003; (d) B. Roubinet, M. Bischoff, S. Nizamov, S. Yan, C. Geisler, S. Stoldt, G. Y. Mitronova, V. N. Belov, M. L. Bossi and S. W. Hell, *J. Org. Chem.*, 2018, **83**, 6466; (e) M. Bossi, J. Fölling, V. N. Belov, V. P. Boyarskiy, R. Medda, A. Egner, C. Eggeling, A. Schonle and S. W. Hell, *Nano Lett.*, 2008, **8**, 2463; (f) V. N. Belov, C. A. Wurm, V. P. Boyarskiy, S. Jakobs and S. W. Hell, *Angew. Chem., Int. Ed.*, 2010, **49**, 3520; (g) J. Fölling, V. Belov, D. Riedel, A. Schonle, A. Egner, C. Eggeling, M. Bossi and S. W. Hell, *ChemPhysChem*, 2008, **9**, 321; (h) H. Aoki, K. Mori and S. Ito, *Soft Matter*, 2012, **8**, 4390; (i) E. A. Halabi, D. Pinotsi and P. Rivera-Fuentes, *Nat. Commun.*, 2019, **10**, 1232; (j) E. A. Halabi, Z. Thiel, N. Trapp, D. Pinotsi and P. Rivera-Fuentes, *J. Am. Chem. Soc.*, 2017, **139**, 13200; (k) E. A. Halabi, S. Püntener and P. Rivera-Fuentes, *Helv. Chim. Acta*, 2018, **101**, e1800165.
- 8 V. N. Belov and M. L. Bossi, *Isr. J. Chem.*, 2013, **53**, 1.
- 9 (a) R. A. Gottlieb, H. A. Giesing, J. Y. Zhu, R. L. Engler and B. M. Babior, *Proc. Natl. Acad. Sci. U. S. A.*, 1995, **92**, 5965; (b) R. A. Gottlieb, J. Nordberg, E. Skowronski and B. M. Babior, *Proc. Natl. Acad. Sci. U. S. A.*, 1996, **93**, 654; (c) M. A. Barry, J. E. Reynolds and A. Eastman, *Cancer Res.*, 1993, **53**, 2349.
- 10 (a) S. N. Uno, M. Kamiya, T. Yoshihara, K. Sugawara, K. Okabe, M. C. Tarhan, H. Fujita, T. Funatsu, Y. Okada, S. Tobita and Y. Urano, *Nat. Chem.*, 2014, **6**, 681; (b) H. Takakura, Y. Zhang, R. S. Erdmann, A. D. Thompson, Y. Lin, B. McNellis, F. Rivera-Molina, S. N. Uno, M. Kamiya, Y. Urano, J. E. Rothman, J. Bewersdorf, A. Schepartz and D. Toomre, *Nat. Biotechnol.*, 2017, **35**, 773.
- 11 G. Xi, L. Sheng, I. Zhang, J. Du, T. Zhang, Q. Chen, G. Li, Y. Zhang, Y. Song, J. Li, Y. Zhang and S. X. Zhang, *ACS Appl. Mater. Interfaces*, 2017, **9**, 38032.
- 12 (a) K. Matsumoto, J. Kusaka, A. Nishibori and H. Hara, *Mol. Microbiol.*, 2006, **61**, 1110; (b) A. S. Johnson, S. van Horeck and P. J. Lewis, *Microbiology*, 2004, **150**, 2815; (c) F. Kawai, M. Shoda, R. Harashima, Y. Sadaie, H. Hara and K. Matsumoto, *J. Bacteriol.*, 2004, **186**, 1475; (d) I. Fishov and C. L. Woldringh, *Mol. Microbiol.*, 1999, **32**, 1166; (e) A. Nishibori, J. Kusaka, H. Hara, M. Umeda and K. Matsumoto, *J. Bacteriol.*, 2005, **187**, 2163.
- 13 O. V. Dolomanov, L. J. Bourhis, R. J. Gildea, J. A. K. Howard and H. Puschmann, *J. Appl. Crystallogr.*, 2009, **42**, 339.
- 14 L. Palatinus and G. Chapuis, *J. Appl. Crystallogr.*, 2007, **40**, 786.
- 15 G. M. Sheldrick, *Acta Crystallogr., Sect. A: Found. Crystallogr.*, 2008, **64**, 112.

