



Cite this: *Org. Biomol. Chem.*, 2019, **17**, 215

Received 31st October 2018,
Accepted 3rd December 2018

DOI: 10.1039/c8ob02711k

rscl.li/obc

Fluorogenic probes for super-resolution microscopy

Eszter Kozma* and Péter Kele  *

Fluorogenic probes efficiently reduce non-specific background signals, which often results in highly improved signal-to-noise ratios. Although this implies improved resolution, fluorogenic probes in the context of super-resolution microscopy are somewhat overlooked. Several excellent reviews summarize recent developments in SRM techniques, labeling techniques or different aspects of small synthetic fluorophores, however there is no comprehensive review on fluorogenic probes suitable for super-resolution microscopy. Herein we wish to fill this gap by providing the readers with an up-to-date summary of fluorogenic probes applied to super-resolution imaging of cellular structures.

Introduction

Recent developments in super-resolution microscopy (SRM) techniques have transformed life sciences in the past decade (for detailed explanation of the respective super-resolution techniques the readers are directed to review papers compiled in ref. 1). Breaking Abbé's law enabled non-invasive visualization of cells at unprecedented resolutions giving us new insights into subcellular structures and the dynamics of biomolecular events. Due to the recent hardware developments, it is the unavailability of fluorescent markers suitable for site-specific intracellular labeling schemes that can currently be considered as the biggest limitation in SRM techniques.^{1b} During the past few decades, several labeling techniques have emerged.^{2,3} Fusion of engineered fluorescent proteins (FP), for example, is routinely accomplished, however their relatively large size (2–5 nm) often impairs precise localization of the protein of interest (POI) in SRM, not to mention their possible interference with the function of the POI.^{1b,3a} Synthetically tailored organic fluorophores have a much smaller size than FPs, allow greater flexibility in terms of labelling sites and often provide much better photophysical properties such as a wider spectral range, greater photostability, and, in many cases, higher brightness. Not to mention that some FPs have the tendency to polymerize.^{1b,2c,4} Furthermore, the use of FPs and other fusion tags is limited to the (mostly terminal) labeling of proteins only.^{3a} Synthetic fluorophores in contrast, can be applied to virtually any biomolecular structures (*e.g.* by means of bioorthogonal chemistry).^{3b,c,4a} Techniques using small

organic fluorophores covering the whole range of the visible spectrum in terms of excitation and emission colors represent the state-of-the-art methods for fluorescence labeling in SRM.^{4a} Besides suitable brightness and photostability, synthetic probes should also address problems such as auto- and background fluorescence in order to achieve further improvements in resolution.^{4b,c} With careful selection of fluorescent cores the first three criteria are easily met. Suppression of the background fluorescence of non-specifically adsorbed dyes, however, often requires extensive washing cycles. An alternative approach to overcome this problem uses so-called fluorogenic (or sometimes also referred to as smart) probes.^{5a,b} Fluorescent cores can be made fluorogenic by several means but they share one thing in common: they exist in a quenched, non- or weakly emissive form, which is transformed into a fluorescent species at a given wavelength solely upon specific conjugation/binding to target structures. The extent of fluorogenicity, *i.e.* the ratio of the fluorescence of the bound and unbound forms, can reach even four orders of magnitudes.^{5c} Although the larger the increase the higher the contrast and the less it is required to wash off unreacted probes, experience suggests that even one order of magnitude difference is enough to tell specifically reacted and freely diffusing or non-specifically bound probes apart.^{5d,e} Irreversibly switched-on fluorogenic probes offer low-background, thus high-contrast images. Reversibly switchable fluorogenic probes can be used to induce random fluorescent events (better known as stochastic blinking), which can be used in single-molecule-localization microscopy (SMLM) based super-resolution technologies, such as stochastic optical reconstruction microscopy (STORM) (for detailed discussion of the technique see ref. 2c and 5f). Yet, relatively few examples are found in the literature that demonstrate the robustness of fluorogenic probes in the context of super-resolution imaging.

Chemical Biology Research Group, Institute of Organic Chemistry, Research Centre for Natural Sciences, Hungarian Academy of Sciences, Magyar tudósok krt. 2, 1117 Budapest, Hungary. E-mail: kele.peter@ttk.mta.hu, kozma.eszter.erika@ttk.mta.hu



Although the set of small-molecule fluorogenic probes is much larger, herein we restrict our discussion to fluorogenic systems that found applications in super-resolution imaging of biological structures. The arbitrary classification that we follow distinguishes structural change induced, (photo)chemically modulated systems and energy transfer based fluorogenicity.

Structural fluorogenicity

This category covers the examples of fluorophores that undergo certain structural changes upon specific conjugation/binding events resulting in an increase of fluorescence intensity. Herein we list instances for probes with fluorogenic behavior due to tautomeric changes, reduction of rotational freedom or planarization of out-of-plane twisted frames, in the context of SRM imaging of cellular structures.

2'-Carboxy-rhodamines and congeners

Xanthene dyes possess several favorable (photo)physical characteristics with respect to bioimaging applications. These include high water solubility, fluorescence quantum yields and molar extinction coefficients. Consequently, xanthene probes are widely utilized as fluorescent markers in protein labeling, ion sensing, *etc.*⁶ The emission maxima of rhodamines can be chemically tailored on the one hand by varying the substituents around the framework. On the other hand, the excitation and emission maxima are also efficiently modulated, *i.e.* shifted towards the red end of the spectrum by incorporating group-14 ($X = \text{C}, \text{Si}, \text{Ge}, \text{Sn}$) elements within the rhodamine frame.⁷ The reason is the low LUMO energy of these fluorophores which is explained by conjugation between the σ^* orbitals of exocyclic X-R bonds and the π^* -system of the X-containing tricyclic system. The extent of the red shift is in the order of $\text{Si} > \text{Ge} > \text{Sn} > \text{C}$ (although Sn-rhodamines are unstable), due to the decreasing efficiency of the $\sigma^*-\pi^*$ overlap with the increasing atomic radius of X and C-X bond length. Regardless of their elemental composition, rhodamines carrying a carboxylic function at their 2' position have the unique feature of existing in two forms due to valence tautomerism, depending on the polarity of the microenvironment (Fig. 1).⁸ In less polar media they are mostly present in a non-fluorescent, spirolactone form, while in polar environments a fluorescent, zwitterionic form is the major contributor. The polarity or pH dependency of the equilibria between the two forms is readily

fine-tuned by electron withdrawing/donating substituents. In general, group-14 element modified carboxyl-rhodamine derivatives are more prone to spirocyclization, which makes them excellent polarity-sensitive fluorogenic probes. Such polarity-driven fluorogenicity of rhodamines can be harvested in fluorescence imaging as non-specific adsorption of the probes onto apolar surfaces forces the molecules to remain dark.⁹

The first report on membrane-permeable siliconrhodamine (carboxy-SiR, **1**) derivatives was published by Johnsson *et al.* (Scheme 1).¹⁰ The dye has excitation and emission maxima in the far red region ($\lambda_{\text{abs max}} = 645 \text{ nm}$; $\lambda_{\text{em max}} = 661 \text{ nm}$) with a spirocyclization $D_{0.5}$ value of 65. The $D_{0.5}$ value allows the estimation of the sensitivity of the dye to polarity and is equal to the dielectric constant, where the dye has half of the maximum absorbance. In physiologically relevant environments low $D_{0.5}$ values suggest non-fluorogenic dyes, while high values of $D_{0.5}$ (*e.g.* 60–70) indicate fluorogenic behavior. They discovered superior performance of SiR-SNAP (**2**), SiR-CLIP (**3**) and SiR-Halo (**4**) derivatives in different types of super-resolution microscopy and performed live cell direct stochastic optical reconstruction microscopy, (dSTORM) with a 640 nm laser power of $\sim 1 \text{ kW cm}^{-2}$. Labeling of H2B histone proteins with SiR-SNAP (**2**) allowed the localization of $\sim 140\,000$ molecules in 10 000 frames, at a time rate of 50 Hz, yielding a super-resolved image in just over three minutes (for an overview of dSTORM technology and key parameters of the acquisition, see ref. 2d). SiR-SNAP performed well in live cell STED imaging of centrosomal protein Cep41 as well. STED microscopy revealed that SNAP-Cep41 forms a ring-like structure along the axis of the centriole, a structural feature previously hidden in confocal microscopy (Fig. 2a). Further derivatives of SiR-carboxyl were made for visualization of microtubules (SiR-docetaxel derivative-SiR-tubulin, **5**) and actin (SiR-desbromo-desmethyl-jasplakinolide derivative-SiR-actin, **6**) in human fibroblasts and also in primary rat hippocampal neurons using live-cell structured illumination microscopy (SIM) and stimulated emission depletion (STED) super-resolution techniques.¹¹ Importantly, the authors proved that a higher resolution ($39 \pm 10 \text{ nm}$) can be achieved with SiR-tubulin (**5**) compared to SiR-SNAP (**2**) ($77 \pm 3 \text{ nm}$) in microtubule imaging, which underlines the importance of using small molecule probes that directly target the structure of interest since it enhances the resolution substantially. Mishin *et al.* used SiR-actin (**6**) to visualize endogenous actin within tissues with dSTORM and revealed that the normal tissue actin pattern differs substantially in comparison to cancer cells.¹² SiR-Hoechst (**7**) was applied to nucleus staining enabling non-toxic, SRM-compatible visualization of nuclei in various cell types. This derivative showed a 50-fold increase in fluorescence upon DNA binding.¹³ A bioorthogonally applicable SiR-tetrazine (**8**) was also developed and applied in STORM imaging of cytoskeletal proteins (actin, vimentin) previously modified genetically with a bicyclononyne-derivatized non-canonical amino acid allowing 92 nm lateral resolution.¹⁴ Just very recently, SiR-Halo (**4**) was used in studying the distribution of abundant scaffolding protein, PSD95, at the postsynaptic

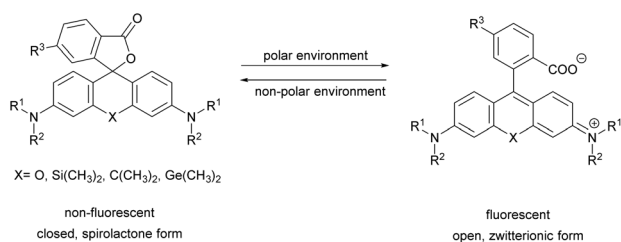


Fig. 1 General structure of 2'-carboxy-rhodamines and their spirolactonization based on the environment polarity.



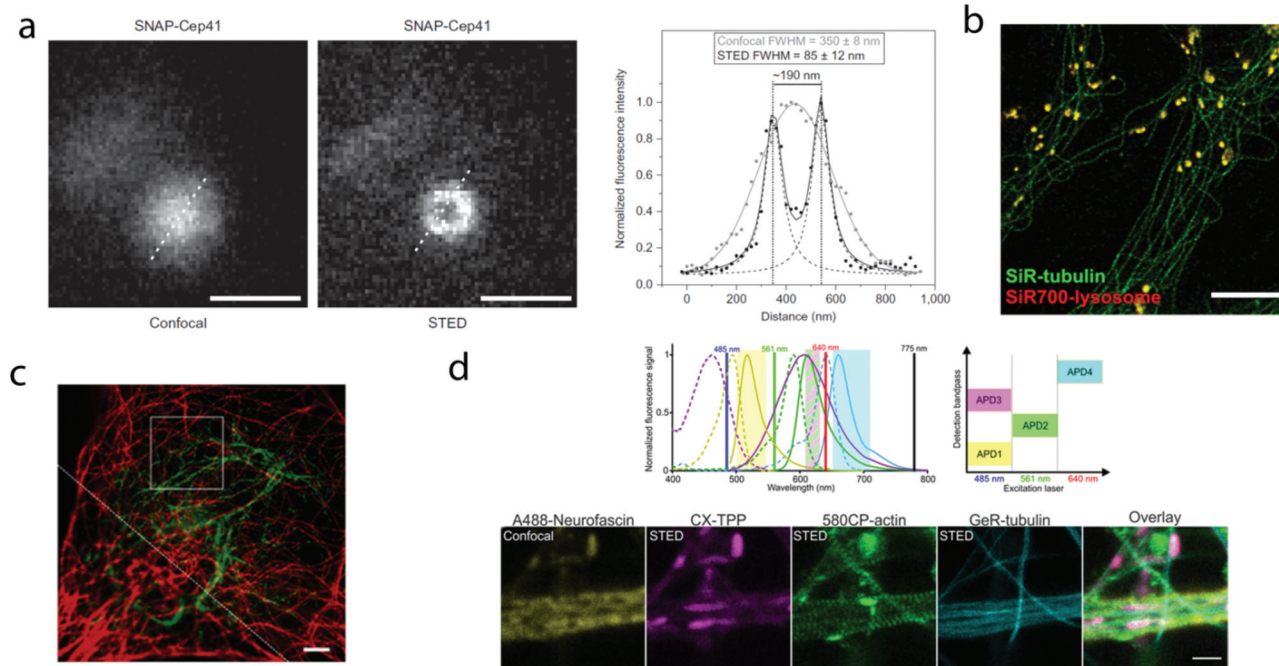


Fig. 2 Superresolution microscopy images with SiR and other carboxy-rhodamine derivatives. (a) Comparison of confocal and STED images of SiR-SNAP-Cep41 localized at the centrosome with an intensity line profile (right) along the white dotted line showing two separated Lorentz-distribution on the STED image (reprinted partly from ref. 10 with permission. Copyright 2013 Springer Nature). Copyright 2016 American Chemical Society). (b) Two-color STED nanoscopy of human primary fibroblasts stained with SiR(1)-tubulin (green) and SiR-700(9)-lysosome (red). Scale bar: 5 μm (adapted from ref. 16 with permission. Copyright 2016 American Chemical Society). (c) Two-color STED image of vimentin (green) and tubulin (red) in a living HeLa cell. Vimentin was labelled with rhodamine 580R (13) applied as a Halo-tag probe, tubulin was directly labelled with SiR(1)-tubulin. Scale bar: 2 μm . Source: ref. 20. (d) Four-colour microscopy with CX-TPP (22). Upper panel: normalized absorption and emission of Alexa Fluor 488 (yellow), CX (22) (magenta), 580CP (15) (green) and GeR (16) (cyan), related laser lines and detection windows. Lower panel: living rat hippocampal neurons stained with 580CP-actin, GeR-tubulin, CX-TPP (targeting mitochondria) and anti-neurofascin and anti-mouse AlexaFluor 488. Scale bar: 1 μm . (Adapted from ref. 29 with permission. Copyright 2017 American Chemical Society.)

gates of 510R (14) and SiR (1) were used to stain microtubule filaments of living human fibroblasts. As a result of high-contrast images obtained with the fluorogenic probes, an apparent diameter of 35 ± 13 nm was obtained corresponding to 29 ± 11 nm optical resolution after taking into account the real microtubule diameter of ~ 20 nm in STED microscopy.

The Hell group has extended two-channel imaging to an 810 nm STED setup. In their account a carbopyronine derivative (610CP, 17) and a siliconrhodamine (680SiR, 18) ensured minimal crosstalk and clear color separation. At the same time both probes were found to be efficiently depleted at 810 nm. The two dyes were used for near-IR live cell imaging of vimentin and tubulin filaments labeled with 610 CP and 680SiR, respectively using two distinct excitation lasers (*i.e.* 600/20 nm for 610 CP and 676/10 nm for 680 SiR) and one 810 nm de-excitation (STED) laser enabling apparent optical resolution values *ca.* 100 and 70 nm, respectively.²²

When a biologically applicable fluorescent or fluorogenic probe comes to intracellular live-cell applications, membrane permeability is always an issue. Although, as just discussed above, sometimes aggregation may facilitate cross-membrane trafficking, it may also impair the efficiency of labeling. Rhodamines conjugated to covalent (*e.g.* SNAP-, Halo-Tag) or non-covalent recognition units (*e.g.* taxanes) efficiently stain

intact cell structures, however their affinity and specificity largely depend on the recognition unit, the probe or the linker applied for conjugation. To address some of these limitations, Butkevich and Hell introduced hydroxylated rhodamines, together with group-14 derivatives (*e.g.* 530RH, 19). While hydroxylation of the fluorescent cores made these probes hydrophilic without compromising their cell permeability the photophysical properties remained mostly unaffected. These probes were indeed cell- and nucleus-permeant that enabled one and two-color STED imaging in live cells with high contrast. However, the authors found that all probes possessed lower Halo-tag binding affinity, and thus required increased dye loadings. Consequently, the study suggests that moderate or high values of $D_{0.5}$ are required but not sufficient to obtain desirable fluorogenic probes.²³

Single molecule localization microscopy, SMLM, generally requires intense laser irradiation (~ 1 kW cm^{-2}) and additives such as thiols (10–100 mM) in the form of so-called blinking buffers to induce on-off switching of the fluorophore, limiting live-cell applications of these techniques. Spontaneously blinking fluorophores that are able to switch on/off stochastically without high power laser irradiation or any additives allow super-resolved localization microscopy of living cells.^{24,25} Urano and co-workers developed siliconrhodamines with an



appending internal nucleophile at the 2'-position, *e.g.* HMSiR (20). Similar to 2'-carboxyrhodamines, such structures tend to form a non-fluorescent spirocyclic form, which is in thermal equilibrium with its fluorescent open form in the ground state (Fig. 3).²⁶ For SMLM applications it is required that at physiological pH only a small subset of probes exist in the fluorescent form and that this state should last for several camera frames in order to enable the detection of sufficiently enough photons for accurate localization. These two requirements are dependent on the intramolecular spirocyclization equilibrium constant (pK_{cycl}) and the lifetime of the open form (τ). They found that pK_{cycl} is strongly dependent on the electrophilicity of the fluorophore core and the nucleophilicity of the pendant group. Following the evaluation of these two properties for several fluorophore-pendant nucleophile combinations, a SiR-derivative, HMSiR (20), with an appending 2'-hydroxymethyl group ($-\text{CH}_2\text{OH}$), was found to be ideal with a pK_{cycl} of 5.8 and a τ of 0.245 s. Halo and SNAP substrate derivatives of HMSiR (20) showed spontaneous blinking and enabled live-cell time-lapse SMLM of microtubules in Vero-2 cells in sodium phosphate buffer (pH 7.4) with laser irradiation at 647 nm with an

intensity of 40 W cm^{-2} . In multichannel SMLM imaging, using conventional probes with redox-active blinking buffers, the composition of the imaging buffer has to be carefully optimized to each fluorophore. Thus it is particularly difficult to find optimal conditions that induce blinking of multiple fluorophores to visualize two different targets at the same time. Spontaneously blinking fluorophores require only regular buffers without any additives therefore suitable candidates for multi-color SMLM imaging. Urano *et al.* developed further probes based on the rhodamine scaffold. A screening process involving differently substituted rhodamines in combination with 2'-hydroxyalkyl chains of varying length resulted in the identification of a green-light emitting probe, HETetFER (21). HMSiR (20) in combination with HETetFER (21) allowed dual-color SMLM of mitochondria and the microtubular network in PBS solution without the use of any redox-buffer or photomodulation.²⁷

In a comprehensive review paper Hell *et al.* highlighted the importance of large Stokes-shift dyes in multi-color STED microscopy.²⁸ With the use of such large Stokes-shift dyes multichannel imaging becomes possible without the need for

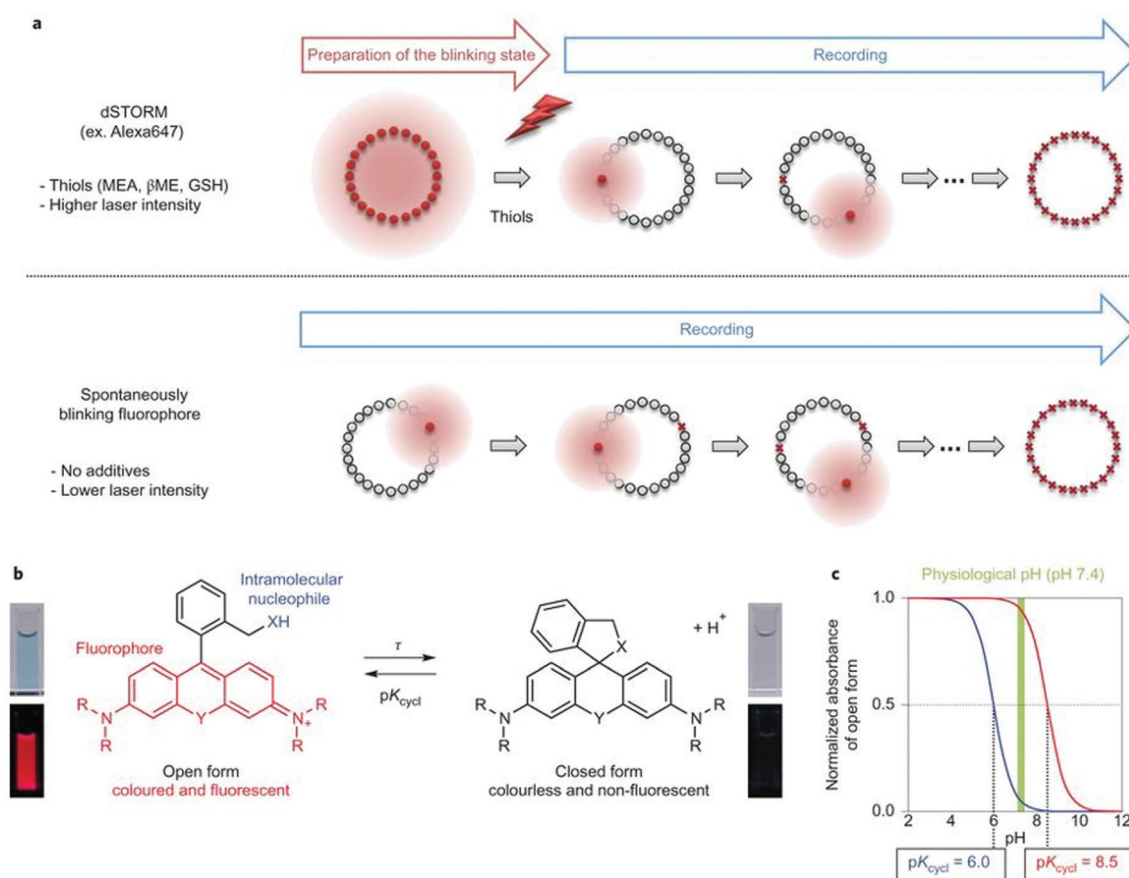


Fig. 3 (a) Single molecule localization sequence using a conventional fluorophore and the spontaneously blinking fluorophore. (b) Thermal equilibrium of intramolecular spirocyclization between the fluorescent open form and the non-fluorescent closed form. pK_{cycl} is the equilibrium constant for intramolecular spirocyclization and τ is the lifetime of the open form (the duration until the open form reverts to the closed form). (c) Schematic pH titration curves of rhodamine derivatives with a pK_{cycl} of 6.0 (blue) and 8.5 (red). For SMLM, most of the individual fluorophores should be in the closed form at the physiological pH of 7.4 to avoid overlapping signals. (Reprinted from ref. 26 with permission. Copyright 2014 Springer Nature.)



sophisticated techniques such as hyperspectral detection or fluorescence lifetime recording. In a separate account, the same group realized such multi-channel imaging using 9-iminoanthrone dyes possessing Stokes shifts between 140 and 165 nm. These probes were not fluorescent in apolar media but showed protonation-based fluorogenic behavior in polar-protic environments (CX-TPP, **22**) (Fig. 4).²⁹

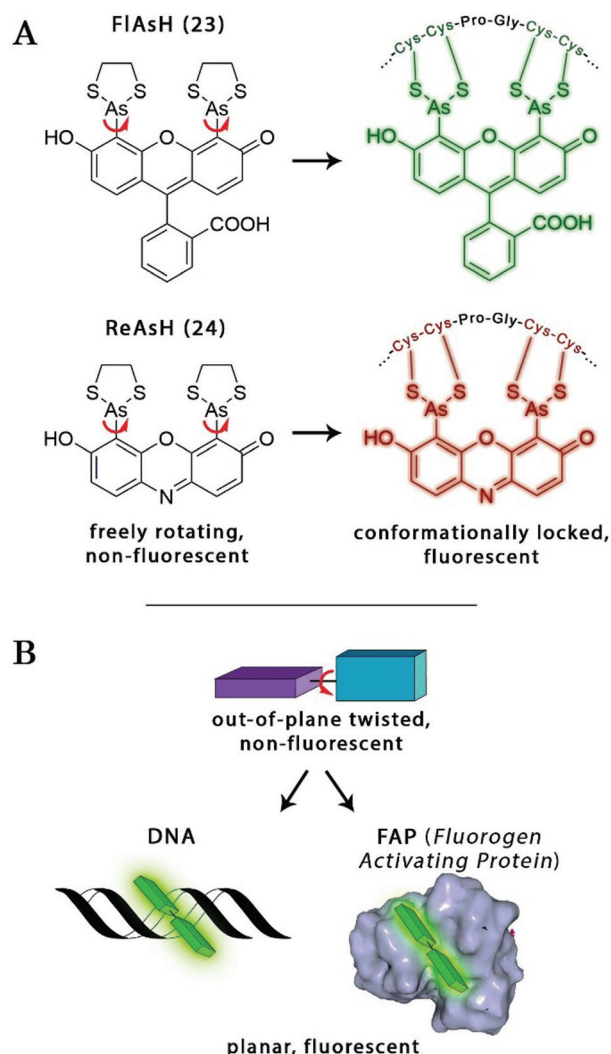
Such a set of cell-permeant dyes enabled two-, three- and even four-color imaging (3× STED, 1× confocal) of cytoskeletal frameworks (GeR-tubulin – **16**, 580CP-actin – **15**) and mitochondria (CX-TPP, **22**, TPP: triphenylphosphonium) in human fibroblasts or living rat hippocampal neurons with a single 775 nm STED laser (Fig. 2d).

Conformationally lockable fluorogenic probes

The probe–target interaction induced conformational confinement of freely rotating or out-of-plane twisted structures often results in a dramatic increase of fluorescence due to forced planarization or rigidification of the probe. Formation of conformationally locked, fluorescent forms can be the result of covalent interactions or non-covalent binding events. Such conformational dynamics related fluorescent changes are realized *e.g.* in covalent binding of biarsenical probes to tetracysteine tag fused proteins or upon intercalation of freely rotating dyes between DNA-bases or binding to DNA grooves (Scheme 2). The same phenomenon can be tracked down during non-covalent binding of quenched systems to engineered fluorogen activating proteins.

Biarsenical helix binders for protein imaging

Amongst protein fusion tags, tetracysteine (Cys₄) motifs represent the smallest ones (4–15 amino acids) that enable selective labeling of the protein of interest (POI). The non-natural arrangement of these natural functions allows highly specific interactions between target proteins and biarsenical probes.³⁰ The bis-1,2-ethanedithiol (EDT) derivative of the fluorescein arsenical helix binder (Scheme 2, FIAsh-EDT, **23**) is weakly fluorescent due to free rotation around the C–As bond (Scheme 2). However, when EDT is replaced by the Cys₄ tag, the cyclic product becomes intensely fluorescent.^{31,32} Since fluorescein can undergo photoswitching, it was a promising candidate to be applied to fluorogenic SRM techniques.³³ To confirm this, Zimmer *et al.* demonstrated FIAsh-based photoactivated localization microscopy (dubbed as FIAsh-PALM) with a spatial resolution ~30 nm.^{34,35} They studied Human Immunodeficiency Virus, HIV, by inserting the tetracysteine



Scheme 2 Schematic examples of conformational change driven fluorogenicity.

tag into integrase (IN), an enzyme that mediates the integration of viral DNA into the host genome.

FIAsh-PALM recovered subdiffraction virion morphology, imaging HIV-1 as free virions and as intracellular complexes in fixed and living cells (Fig. 5). A substantial change in virus size was detected between cytoplasmic and nuclear complexes. Their data highlighted that FIAsh-PALM preserves the biological function of a protein that otherwise would have been disrupted by larger fusion tags (*e.g.* GFP, SNAP, *etc.*) or antibody binding, and furthermore revealed many unexplored steps in the replication cycle of HIV *in vivo*. A red-emitting variant of FIAsh called ReAsH (**24**) has also been developed with an emission maximum at around 608 nm (Scheme 3).³⁰ Selvin *et al.* attached single ReAsH-EDT (Scheme 2, **24**) to Cys₄-fused calmodulin immobilized on a glass surface at low density.³⁶ They determined the center of the distribution of photons, using TIRF microscopy from the image of a single molecule in order to determine the position of the dye. They also found

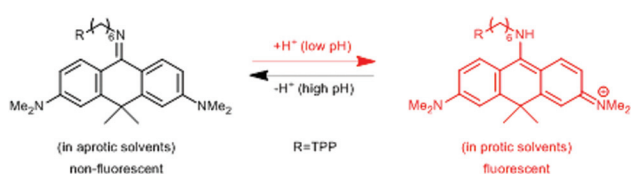


Fig. 4 Protonation–deprotonation mechanism responsible for the environment sensitivity of CX-TPP (**22**).



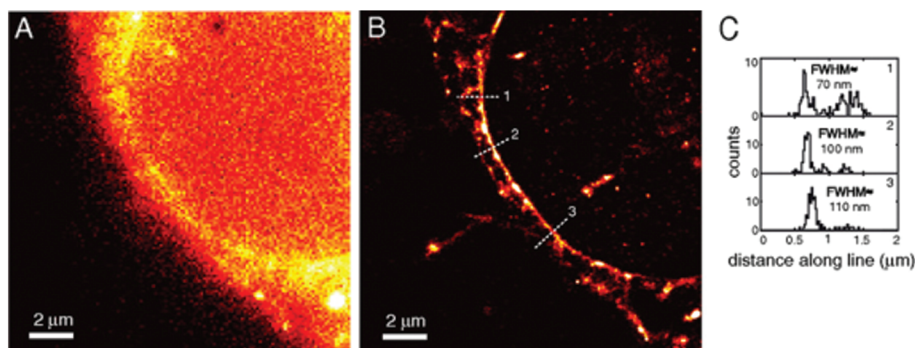
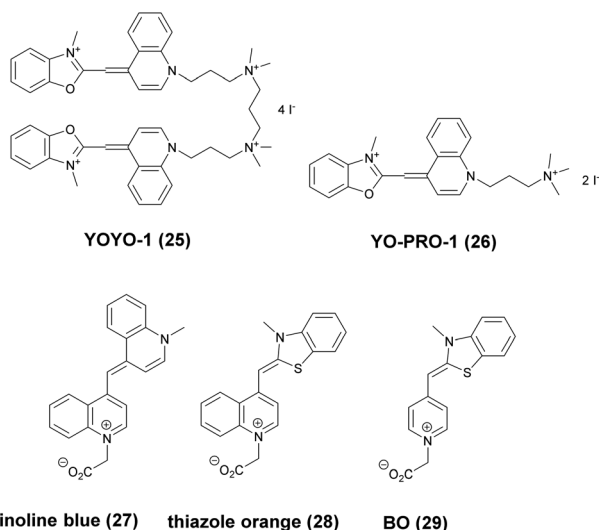


Fig. 5 FISH-PALM imaging of HIV-1 integrase (IN) enzyme in fixed cells. (a) Widefield image, (b) super-resolution reconstruction of the same area, (c) histogram of positions projected on (b). (Reprinted from ref. 34 with permission.)



Scheme 3 Structure of nucleic acid binding probes.

that by increasing photostability, 2-mercaptoethanesulphonic acid increased the number of collected photons from ReAsH molecules by a factor of two. To the best of our knowledge, live cell applications of ReAsH have not yet been presented.

Non-covalent binder probes for DNA imaging

Minor groove binder or intercalating DNA stains have the key advantage that their fluorescence emission intensity increases typically 100–1000 times upon binding to the DNA.³⁷ Minor groove binding molecules tend to be small and flexible in order to bend and fit into the minor groove, whereas intercalator dyes usually contain out-of-plane twisted heterocyclic systems, which interact with DNA bases. Some cyanines, such as TOTO or YOYO, bind to DNA in a ‘bis-intercalating mode’ meaning that the aromatic part of the cyanine intercalates, whilst the amine ‘arms’ lie in the minor groove.³⁸ DNA binder cyanines adopt an out-of-plane twisted conformation when unbound. The twist makes these dyes non-fluorescent. However, intercalation into DNA induces the confinement of

these dyes into a fluorescent planar form.³⁹ YOYO-1 is a DNA-binding cyanine derivative and is the tetracationic homodimer of oxazole yellow. In aqueous solution it is virtually non-fluorescent, however, its emission intensity greatly increases upon bis-intercalation into DNA (3200×).⁴⁰ In 2009, Flors and co-workers applied YOYO-1 (25), to image λ -DNA topology in switching buffers allowing 38 nm resolution of stained structures using SMLM (Scheme 3).⁴¹ The authors suggested that the blinking occurs through an electron transfer reaction of YOYO-1 (25) with the reducing mercaptoethylamine (MEA). YOYO-1 (25) was further applied to single DNA imaging using binding-activated localization microscopy (BALM) in various dye-base pair ratios to analyse the distance between base pairs.^{42,43} A drawback of YOYO-1 (25) and other dimeric intercalating dyes is that their binding constant to DNA is rather high ($K = 10^8$ – 10^9 M⁻¹), which influences the DNA topology and its ability to interact with DNA-binding proteins.⁴⁴

Monomeric intercalating dyes with lower affinity towards DNA can address this problem, though their brightness is about half of their dimeric congeners, and their affinity to DNA is about two magnitude lower. In a separate account Flors *et al.* tested monomeric YO-PRO-1 (26) in DNA staining and achieved 43 nm resolution using SMLM, however, the much lower intensity and staining ability of YO-PRO-1 required higher dye loadings (Scheme 3).⁴⁵ Given the photophysical properties of YOYO dyes, the deterministic nature of STED microscopy may have an inherent advantage over the stochastic methods in DNA imaging.⁴⁶ Hell *et al.* demonstrated that STED can be readily used to identify subtle structures (bends and kinks) along the DNA strand with an average FWHM = 42 nm using YOYO-1 (25) with a 568 nm depletion laser (Fig. 6).⁴⁶ Although YO-PRO-1 (26) is able to pass the cell membrane under very specific circumstances, bis-intercalating cationic dyes are considered to be impermeant to cell membranes.⁴⁷ Thus, it is necessary to use membrane permeable alternatives for live cell DNA imaging *e.g.* SYTO dyes. Like YOYO and its derivatives, SYTO dyes belong to the cyanine family, with the substantial difference of being minor groove binders. Compared to *e.g.* YOYO they produce dimmer SRM images, therefore in general, they are less efficiently localized



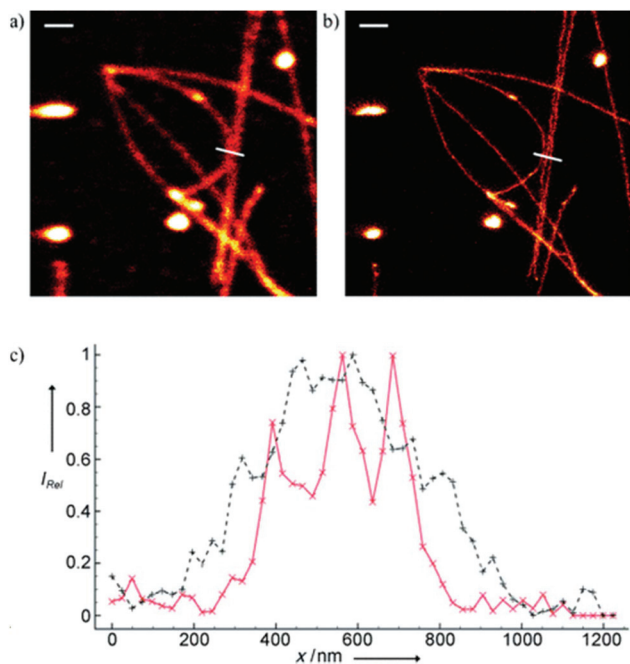


Fig. 6 Confocal image of YOYO stained λ -DNA (a) and the corresponding STED image (b). Scale bars: 1 μm . (c) Average of three line profiles from the STED (solid red line) and confocal (dotted black line) images. Line profiles extracted along the white lines in (a) and (b). The three distinct peaks belonging to different DNA molecules are only resolved by STED. (Reprinted partly from ref. 46 with permission. Copyright 2011 John Wiley and Sons.)

in stochastic methods.⁴⁸ Moreover, SYTO dyes are not DNA-specific and they also bind to RNA, however the staining pattern is dye-dependent.⁴⁹ Preliminary data suggest that SYTO-13 might be useful for imaging chromatin of fixed mitotic chicken DT40 cells in blinking buffer.⁵⁰ SYTOX Orange, whose chemical structure is proprietary, is reported to be monomeric and intercalating.⁵¹ It has just recently been found to be promising in live cell labeling applications.⁵² Moerner *et al.* studied single molecule rotational dynamics using SYTOX Orange and detected the nanoscale deformations of individual λ -DNA strands *in vitro*.^{53a} By combining image data with thousands of dye molecule orientation measurements, they developed a technique to probe the structure of individual DNA strands. In spite of the promising results, live cell DNA imaging using fluorogenic DNA-binders is still a field full of challenges.

Simultaneous imaging of multiple intracellular RNA molecules is also challenging and requires careful optimization of label and probe design. Seitz and co-workers described a set of quencher-free forced intercalation (FIT) probes, in which a single non-symmetric thiazole orange (TO) derived cyanine dye serves as an environment sensitive nucleobase surrogate.^{53b} Hybridization with a complementary target forces the dye to intercalate between predetermined base pairs, resulting in restricted torsions around the methine bridge leading to a prolonged lifetime of the excited state. The authors developed the

first red-emitting FIT probe, quinoline blue (27), showing up to 152-fold enhancement in fluorescence upon hybridization with RNA. The combined use of quinoline blue with other FIT probes (*e.g.* thiazole orange, TO, 28 and benzothiazole orange BO, 29) allowed localization of three different mRNA molecules in developing oocytes of *Drosophila melanogaster* by means of wash-free *in situ* hybridization and STED microscopy (Scheme 3).

Fluorogen activating proteins for super-resolution microscopy

A similar approach, involving conformationally lockable, non-covalent binder probes is also applied for super-resolution imaging of proteins in combination with genetically encodable fusion protein tags. Such a group of proteins termed fluorogen activating proteins (FAP) are generated *e.g.* by screening of yeast surface-displayed libraries for specific binding and efficient activation of a variety of fluorogenic probes.^{54,55} Binding of the probes results in significant enhancements in the extinction coefficient and fluorescence quantum yields, sometimes in combination with significant absorption band shifting as a result of structural confinements (Fig. 7a).

These FAPs have been fused to a variety of extra- and intracellular proteins and recently, have been applied in super-resolution microscopy. Bruchez *et al.* genetically expressed FAP-fused proteins (*e.g.* yeast cell surface and actin in HeLa cells) that selectively bind a far-red fluorogenic malachite green (30) probe (Scheme 4).⁵⁶ Free malachite green is essentially non-fluorescent due to rotational relaxation, however an enhanced fluorescence is observed in the presence of cognate FAPs (up to 18000-fold).⁵⁷ Live-cell STED microscopy revealed a sufficient signal-to-noise ratio resulting in a three-fold resolution enhancement compared to confocal microscopy. More recently, it was reported that with careful control of malachite green (30) concentration a sparse labeling distribution on

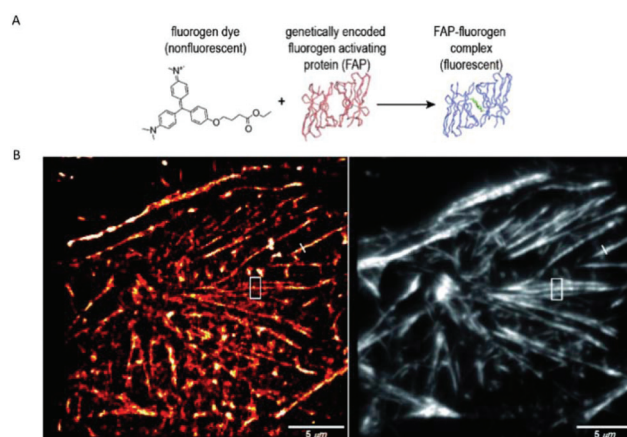
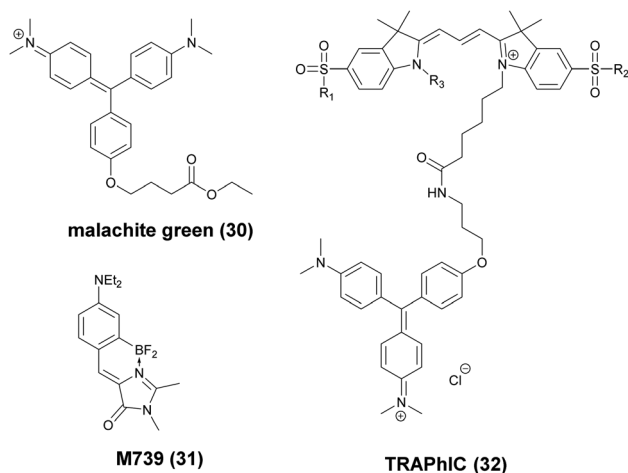


Fig. 7 (a) The fluorogen activating protein complex consists of a nonfluorescent dye (fluorogen) and a genetically encoded protein which forms a fluorescent complex upon interaction. (b) Live cell F-BALM imaging of actin in HeLa cells. Left: reconstruction of SMLM of actin, right: wide-field image of phalloidin-Alexa488. (Reprinted partly from ref. 58 with permission. Copyright 2013 John Wiley and Sons.)





Scheme 4 Structure of fluorogen activating protein (FAP)-complex ligands used in super-resolution microscopy.

densely tagged genetically encoded proteins can be achieved.⁵⁸ Stochastic SRM was enabled with equilibrium binding using concentrations below the dissociation constant (K_d) of the fluorogenic substrate. Spontaneous blinking was produced without the use of any additives or switching buffers. The developed method, dubbed as F-BALM (binding-activated localization microscopy), permitted live cell single molecule imaging of actin in HeLa cells (Fig. 7b).

Mishin *et al.* performed *in silico* mutagenesis of amino acids within the ligand binding pocket of Blc protein, belonging to the lipocalin family, in order to identify potential protein-chromophore pairs for SMLM.⁵⁹ Protein complexes (named as Dye in Blc, DIB) capable of efficiently activating M739 (31) were fused to different protein structures (histone H2B, cytokeratin, actin, vimentin) in mammalian cells (Scheme 4). As compared to F-BALM, this complex (in a technique named protein-PAINT) showed increased photostability resulting in enhanced temporal resolution at the single-molecule level. A particular FAP, dubbed as dL5, has been optimized for high binding affinity towards a membrane permeant malachite green (30) derivative with increased photostability and better spectral properties. Moerner and co-workers used this construct in the labeling scheme of fusion proteins in live bacteria *C. crescentus* and demonstrated long time-scale protein tracking and STED microscopy with a four-fold resolution enhancement compared to diffraction-limited imaging. Furthermore, the authors provided evidence that in comparison with YFP, dL5 fusion induced less perturbation in the structure and function of the intermediate filament protein CreS.^{60,61} Further attempts to perform SMLM failed since MG-ester activated upon binding to the bacterial cell surface.

Bruchez and co-workers reported on an excitation ratio-metric, activatable pH responsive tandem system based on a bichromophoric probe acquired by linking the pH sensitive Cy3 donor to a conformationally activatable malachite green acceptor (TRAPhIC) (32) (Scheme 4).⁶² The complex is acti-

vated by its cognate FAP dL5**. In the absence of FAP binding, the malachite green rather acts as a non-fluorescent quencher of the Cy3 fluorescence, however, in the presence of the FAP (fused to target receptor protein) dual emission is generated due to the efficient FRET between the chromophores. Fine changes in the pH due to receptor internalization during endocytosis are sensitively followed by the ratio of the signals. TRAPhIC labeling was found to be compatible with ratiometric STED imaging, enabling tracking of β_2 adrenergic G-protein coupled receptors (GPCR) with dual excitation (560 nm and 640 nm) and single depletion at 775 nm.

(Photo)chemically modulated systems

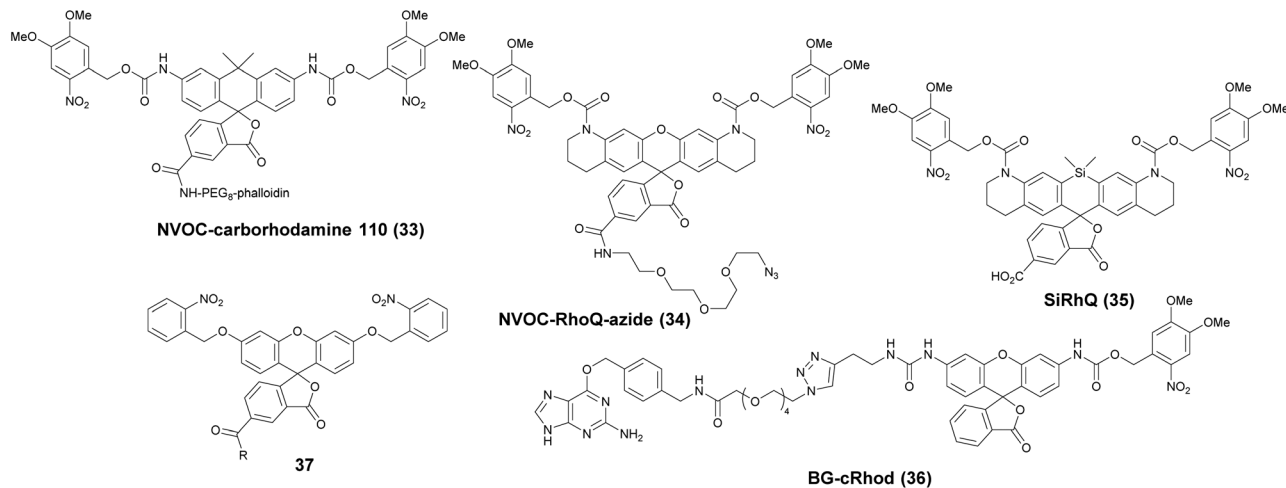
This category consists of fluorogenic systems where the fluorescence of light emitting frames is quenched either by a photochemically cleavable quencher unit or a chemically active function that efficiently modulates the fluorescent signal until it is reacted with a specific target. Furthermore, photochromic systems also belong to this category. Here the photoirradiation does not induce cleavage of quenchers but rather affects the equilibrium between dark and emissive structural isomers. Photoirradiative removal of caging units, chemical transformation of the quencher moiety or photo-switching of photochromic dyes leads to restoration or appearance of the fluorescent signal. Photocaged and photochromic probes are quite popular in super-resolution applications due to their spatiotemporally controllable switch-on characteristics.

Photoactivatable and photochromic systems

Photoactivatable fluorophores are non- or weakly fluorescent molecules that can be photochemically converted to intensively emitting species. For PALM imaging, photoactivatable fluorophores bearing photolabile protecting (caging) groups have been developed. Common caging groups are the 2-nitrobenzyl moiety and its derivatives such as 4,5-dimethoxy-2-nitrobenzyl- (DMNB) or *ortho*-nitroveratryloxycarbonyl-groups (NVOC) that are efficiently removed by UV light irradiation.^{63,64} Caged synthetic fluorophores have particular advantages in localization microscopy. They exhibit a higher photon count than fluorescent proteins resulting in better localization precision. Additionally, caged compounds are activated by a photochemical reaction, allowing a sparse, controlled turn-on of individual molecules. This eliminates the need of using harsh imaging conditions (such as buffers in dSTORM), however uses a UV-laser, which can potentially be phototoxic.

The open-closed equilibrium of 2'-carboxy-xanthene derivatives can also be controlled by installing caging groups onto the oxygens of fluoresceins or the aniline nitrogen of rhodamines, which shifts the equilibrium significantly towards the dark spirolactone form. Removal of the caging groups restores the equilibrium shifted towards the fluorescent zwitterionic form yielding a large increase in fluorescence. Lavis and co-workers used a photocaged carborhodamine NVOC-carborhodamine 110 (33, Scheme 5) derivatized phalloidin for the





Scheme 5 Photoactivatable fluorophores with photolabile groups mentioned in the text.

imaging of actin filaments by means of PALM without any redox buffers.⁶⁵ The shifted open-close equilibrium observed with carborhodamine derivatives ensures that PALM imaging can be performed even on densely labeled structures. The same authors reported a general synthetic route to access photoactivatable xanthene fluorophores.⁶⁶ In this scheme, they used reduced “leuco” form as a synthetic intermediate allowing installation of the caging groups using mild conditions. NVOC-caged azide-derivatized Q-rhodamine (34), rhodamine 110 and Oregon Green derivatives were applied for the first time in SRM imaging of DNA in the cellular context (Scheme 5). Using the NVOC-caging strategy, Grimm *et al.* reported a Si-containing analogue of Q-rhodamine (SiRhQ, 35), which exhibits higher photon counts compared to established localization microscopy dyes, such as Alexa Fluor 647, especially after the addition of oxygen-depriving Trolox.⁶⁷ The achieved high precision allowed high resolution imaging of actin-rich structures, such as filopodia (FWHM = 45 nm). SiRhQ (35) performed well in multicolor localization microscopy as well, in combination with the fluorescent protein mEos2. SiRhO, 35 and mEos2 have sufficiently different activation cross-sections to allow successive imaging of the two colors (Fig. 8). Johnsson *et al.* presented a caged rhodamine 110 derivative that can be conjugated to SNAP-tag fusion proteins (BG-cRhod, 36).⁶⁸ In BG-cRhod (36), the SNAP-tag reactive O⁶-benzylguanidine (BG) is linked *via* a urea motif to rhodamine 110. The photolabile DMNB group attached to the other amino group of rhodamine 110 forces the probe to its non-fluorescent lactone form until it is irradiated with 365 nm UV light. PALM imaging of mitochondria of U2OS cells resulted in *circa* twice as better localization as with red fluorescent proteins. Wombacher *et al.* described new fluorophores for dual color photoactivated SRM in live mammalian cells.⁶⁹ They synthesized trimethoprim and Halo-tag conjugates of NVOC-Q-Rhodamine and *ortho*-nitrobenzyl-caged fluorescein (37) and demonstrated dual-color activation following exposure to UV-light using epifluorescent microscopy and iFRAP

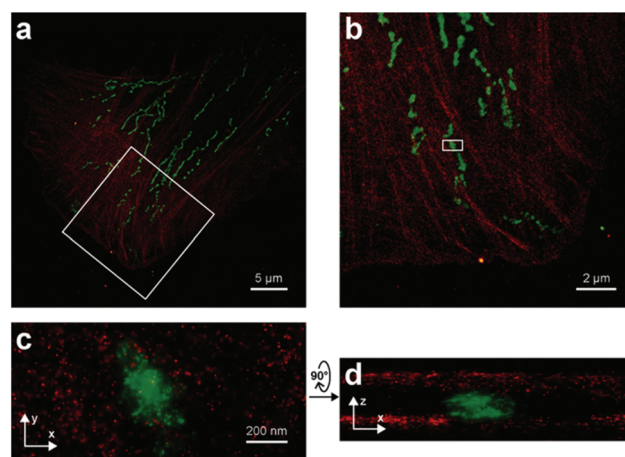


Fig. 8 Two-color iPALM images of fixed U2OS cells labelled with SiRhQ(35)-phalloidin conjugate (actin, red) and mEos2 (mitochondria, green). (b) Enlarged image of the boxed area is in (a). (c) is x, y projection of the boxed area in (b). (d) x, z projection of the boxed area in (b). (Reprinted from ref. 67 with permission. Copyright 2015 John Wiley and Sons.)

(inverse fluorescence recovery after photobleaching) as well as PALM imaging with caged RhoQ. It should be noted that the 2-nitrobenzyl group and its surrogates are bulky and generate toxic, colored and reactive 2-nitrosobenzaldehyde or 2-nitrosobenzophenone derivatives upon photolysis.⁷⁰

Dicyanomethylenedihydrofuran (DCDHF) type push-pull fluorophores contain an electron donor amine moiety connected to the DCDHF acceptor *via* a conjugated π -system. Moerner *et al.* reported azide caged fluorogen 38 that remains dark until photoactivation. A short burst of 407 nm UV light converts the azide to an amine shifting the absorption to a longer wavelength (*i.e.* 570 nm). The created photostable compound 39 could be imaged at the single molecule level (Fig. 9a).⁷¹ Later, they applied a derivative of this probe in



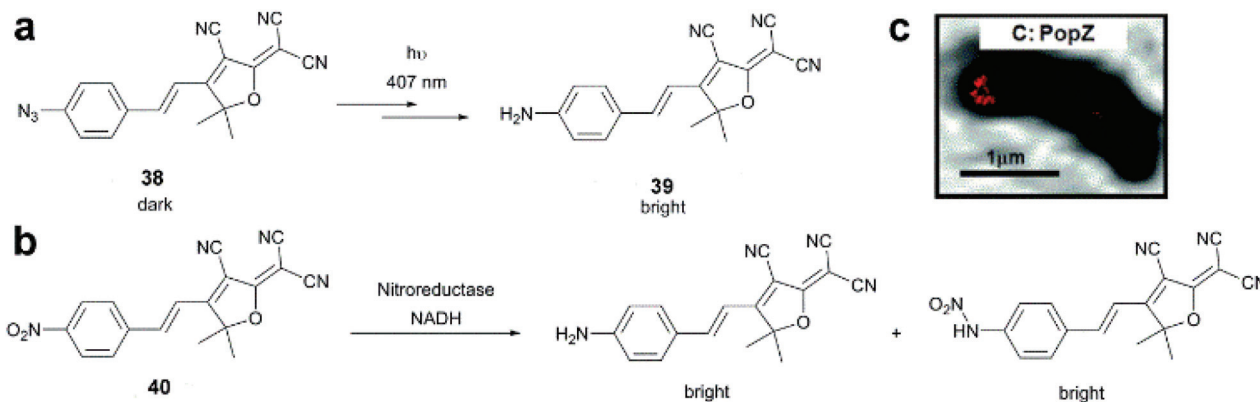


Fig. 9 (a–b) Structure of DCDHF class fluorogens and their fluorescent products. (c) SRM imaging of PopZ protein fusions inside live *Caulobacter crescentus* cells using **38** in a Halo-tag form. (Adapted from ref. 72 with permission. Copyright 2010 American Chemical Society.)

Halo-tag mediated super-resolution imaging of microtubules in fixed mammalian BS-C-1 cells with 85 ± 15 nm resolution.⁷² The cell-permeable nature of this probe allowed live *Caulobacter crescentus* cell labeling and revealed protein localization patterns below the diffraction limit (Fig. 9c). In 2013, they replaced the azide with a nitro group, (nitro-DCDHF, **40**), and subjected it to treatment with nitroreductase and electron donors (NADH) (Fig. 9b) resulting in the formation of fluorescent products without phototoxic UV irradiation.⁷³ The method, named enzymatic turnover activated localization microscopy (ETALM), enabled live cell SRM imaging of bacterial cells.

A somewhat distinct group of photoactivatable fluorophores suitable for SRM is represented by so-called photochromic probes. These fluorophores have the characteristics that they can be reversibly interconverted between a dark non-fluorescent and an emissive form. The reversible switching involves either photoirradiation in both directions or a photoirradiation-thermal back isomerization process. Either way, photoswitching does not require the use of blinking buffers. Rhodamine spirosamides exist in a non-fluorescent closed form, which becomes fluorescent upon irradiation with UV light (single photon) or two-photon laser irradiation (Fig. 10a). The open isomer has a lifetime of a few milliseconds in polar solvents before being converted back thermally to the dark form. Such behavior is ideal for SMLM applications.⁷⁴ Hell *et al.* developed SRA577 (**41**), a rhodamine B derivative with an appending 4-aminophthalimide substituent, with a thermal relaxation time of 20–100 ms in polar solvents (Fig. 10a).⁷⁵ In contrast to caged probes, which are activated and then photo-bleached, photochromes can be used to localize markers several times, thereby allowing multiple measurements on the same sample. This characteristic is quite beneficial in live-cell imaging applications. In their report, the Hell group tested SRA577 (**41**) using PALMIRA (PALM with independently running acquisition) on silica nanobeads offering non-invasive optical sectioning. Separate detection of individual molecules enabled assignment of every single molecular event to the

corresponding dye class, thus allowing multicolor localization imaging. The same authors later developed four more rhodamine spirosamide derivatives and demonstrated multicolor separation on silica nanobeads with improved resolution (15 nm).⁷⁶ Two-color labeling of microtubules and the keratin network in Ptk2 cells with SRA552 (**42**) and SRA577 (**41**) using secondary antibodies was also presented at the nanoscale. Further rhodamine spirosamides were reported spanning the whole visible spectrum for multicolor SMLM by Belov and Hell.⁷⁷ Photoswitching of photochromic probes to emitting forms is usually achieved upon UV light irradiation, however these wavelengths may damage cells. To address this, Moerner *et al.* shifted the photoswitching wavelengths of rhodamine spirolactams towards visible wavelengths >400 nm.⁷⁸ Additionally, they varied the chromophores on the lactam nitrogen, which facilitated the switching at lower photoactivation energy without substantially altering other photophysical features. An optimized derivative (**43**) was used to label surface amines of live *Caulobacter crescentus* bacterial cells for 3D super-resolution imaging (Fig. 11). Low laser intensity purple beam (405 nm, 18 W cm^{-2}) PALM imaging revealed a localization precision of 14–17 nm with enhanced specificity.

In conventional multicolor spectroscopy, the different fluorophores are usually separately excited to induce the corresponding emission wavelength. Such multiple excitations can, however, produce chromatic artefacts and overlaps. Belov, Hell *et al.* elegantly addressed this and used dyes with similar spectral characteristics but with different switching-on properties.^{70,79} The authors combined non-caged TMR, diazoketone-caged compound **45** and spirosamide-caged SRA552 (**42**). Actin, microtubule cytoskeleton, and peroxisomes were labeled using different labeling strategies (phalloidin, antibodies) with the above-mentioned derivatives. First, TMR was imaged and bleached with 546 nm light, and then TMR-NN was uncaged (420/30 nm) and imaged, followed by uncaging of SRA552 (**42**) (spirosamide of rhodamine S – RhS on the figure) (360/40 nm) (Fig. 12).

Rivera-Fuentes and co-workers developed a dual-activatable diazoketone-based probe to sense esterase activity.⁸⁰ Without



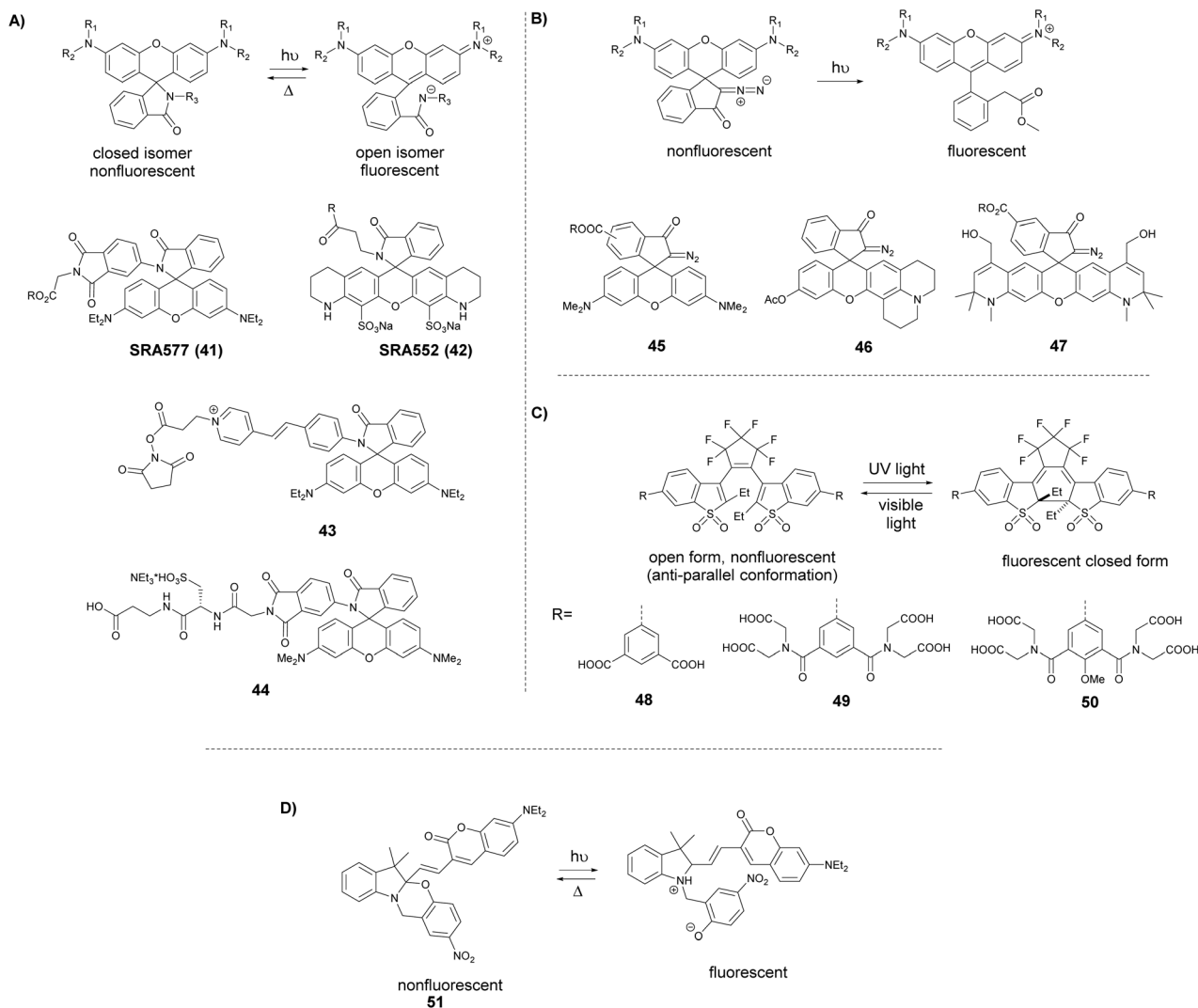


Fig. 10 Switching process and structure of photochromic labels discussed in the text: (a) spiroamides, (b) diazoketones, (c) diarylethenes, and (d) oxazine photochromes.

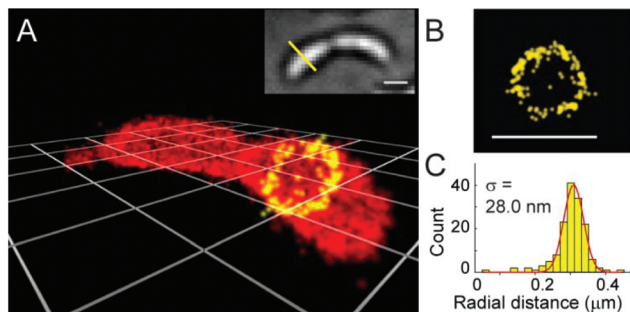


Fig. 11 3D super-resolution reconstruction of surface localization in live *Caulobacter crescentus* bacterial cells using compound 43. (a) A 75 nm slice is highlighted in yellow, and viewed in the panel of the cell axis (b). (c) Histogram of the radial distances of each of the localizations to the center of the circle fitted to the points in panel B. (Adapted from ref. 78 with permission. Copyright 2014 American Chemical Society.)

enzymatic preactivation, the diazoketone probe (46) has an electron-poor core, which, upon irradiation, undergoes Wolff rearrangement to yield a non-fluorescent ring-expanded xanthene. However, when carboxylesterases activate the fluorogen, the tricyclic core becomes electron-rich, and the photo-induced Wolff rearrangement produces a fluorescent rhodol. This probe was utilized in SMLM to sense esterase activity in live HeLa cells.

As the abovementioned examples show, photoactivatable spiroamides and diazoketones proved to be useful markers recently in SMLM, however their applicability is often limited by aggregation, non-specific adhesion and low reactivity caused by low water-solubility. Hell *et al.* applied a universal hydrophilizer (a 3-sulfo-L-alanyl – beta-alanine dipeptide) and allylic hydroxyl groups as synthetic modifications to increase the polarity (e.g. compounds 44 and 47).⁸¹

The fluorescence of 1,2-bis(2-alkyl-6-phenyl-1-benzothio-phen-1,1-dioxide-3-yl)perfluorocyclopentenes (diarylethenes)



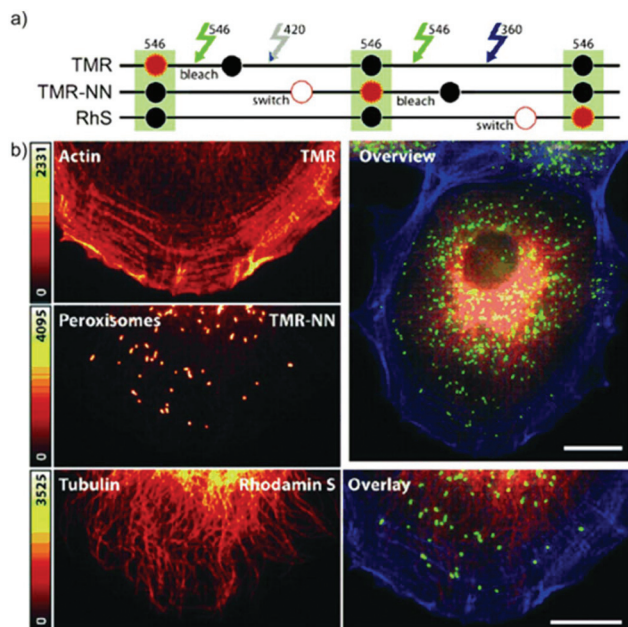


Fig. 12 Monochromatic multilabel imaging using photochromic probes TMR-NN (45) and RhS (SRA552, compound 42 in the text) in combination with conventional TMR. See details of imaging scheme (a) in the text. (b) Application on Ptk2 cells. Scale bar: 10 μm . (Reprinted from ref. 70 with permission. Copyright 2010 John Wiley and Sons.)

can be modulated by light in both directions. While UV light facilitates the formation of the fluorescent “closed” form visible light promotes isomerization to the dark “open” form (Fig. 10c). These photoswitches emit bright, green fluorescence with high photostability, however they are generally hydrophobic and non-soluble in aqueous solutions.⁸² Hell *et al.* decorated the diarylethene core with four or eight carboxylic groups to access 48 and 49, respectively, to increase the water solubility and prevent the aggregation of the free dyes.⁸³ Derivatives 48 and 49 were tested in immunostaining of tubulin filaments and nucleopore complexes in Vero cells, and robust photoswitching was observed upon irradiation with 355 nm light for 50 μs . Following 200 μs delay irradiation with 488 nm light for 1.2 ms in a doughnut shape resulted in the formation of the dark “open” form at the respective area. By means of Reversible Saturable Optical Fluorescence Transitions (RESOLFT) microscopy the authors achieved an optical resolution of 74 nm (FWHM). Attempts, however, to apply these derivatives in coordinate-stochastic approaches failed since switching properties in aqueous media were inadequate for efficient detection of single molecules. To overcome this limitation, the diarylethene structure was modified with electron donor (*p*-methoxyphenyl) and electron acceptor substituents in order to increase the push-pull effect to improve the on/off ratio.⁸⁴ These modifications resulted in octa-acid 50, which showed high aqueous solubility, high emission efficiency and low cycloconversion quantum yield and proved to be suitable for STORM imaging by applying single 488 nm light without additional photoactivation with

UV light or further additives. Raymo *et al.* studied oxazine photochromes in fluorophore-photochrome dyads.⁸⁵ They designed a set of hemicyanine-based fluorophores involving a photoswitchable oxazine unit. The 2*H*,4*H*-[1,3]-oxazine ring of these compounds opens on a subnanosecond time scale to generate a zwitterionic isomer (Fig. 10d). The photogenerated fluorescent forms revert spontaneously to the original, non-emissive species within submicroseconds. They applied derivative 51 for stochastic SRM on amphiphilic copolymer nanoparticles, demonstrating the excellent blinking effect.

Chemically activatable fluorogenic probes

The most important representatives of chemically activatable fluorogenic probes suitable for super-resolution applications possess azide or tetrazine moieties as a reactive quencher unit. Both the azide and tetrazine can participate in highly selective, biocompatible reactions, *i.e.* bioorthogonal reactions. Thus these functions are two-in-one combinations that make fluorescent cores bioorthogonally applicable and fluorogenic at the same time.^{86–88} When the azide/tetrazine moiety is transformed in a specific reaction with the target biomolecule into a triazole or dihydropyridazine, respectively, the fluorescence is restored. To the best of our knowledge, up to now, only tetrazine quenched systems were applied to SRM imaging of biological samples. Although the modular design of bioorthogonally active tetrazine quenched fluorogenic probes allows the use of various fluorophores, their potential for site-specific intracellular fluorogenic labeling was just lately exploited in super-resolution imaging techniques.

There are three ways tetrazines are able to modulate the fluorescence of aromatic cores: (i) *via* Förster-type of resonance energy transfer (FRET)^{88a–c} (ii) *via* a through-bond energy transfer (TBET) process^{5d,88d,e} and (iii) as just recently reported, in directly conjugated, monochromophoric systems.⁸⁹ In FRET systems the fluorescent core is linked to the tetrazine through a flexible linker. Besides other criteria (*e.g.* distance, alignment of transition moments *etc.*), a considerable overlap between the emission band of the fluorophore and the absorption band of the tetrazine (typically falls between 520–540 nm) is necessary. This criterion limits the applicable fluorescent cores to green emitting frames.^{88a–c} Much more efficient energy transfer occurs *via* TBET fluorogenic probes, where the fluorophore and the tetrazine are attached together *via* a rigid, conjugated, otherwise electronically decoupled (twisted) linker.^{5d,88d,e} It is important to note that energy transfer *via* TBET does not require spectral match between the fluorophore and the quencher, which allows the combination of virtually any kind of tetrazines and fluorescent cores with various emission wavelengths. In monochromophoric design where the tetrazine is directly linked to the fluorescent core the presence of an optically inactive $n-\pi^*$ transition is believed to be the reason for quenched fluorescence.⁸⁹ Despite the many advantages, there are only a few examples of super-resolution applications of tetrazine quenched systems.

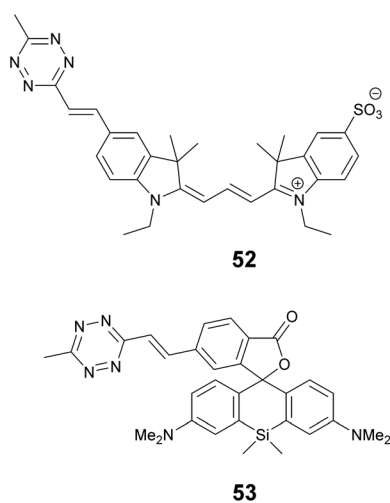
We recently applied the TBET strategy to cyanine (Cy3 and Cy5) scaffolds, where the two units were linked either *via* a



phenylene or a vinylene unit.⁹⁰ All four probes showed fluorogenic behavior, however only one of them, Cy3-derivative **52**, produced a fluorescence enhancement over one order of magnitude upon reaction (Scheme 6). The 14-fold intensity increase, however, enabled sufficient distinction between unreacted, non-specific and specifically reacted, turned-on signals even under no-wash conditions. Although indocyanines are not preferred in STED-based super-resolution techniques, we have found compound **52** to be suitable for STED imaging of actin bundles with 168 nm resolution using a 660 nm continuous wave laser for depletion (Fig. 13). It should also be mentioned that no loss of signal-to-noise ratio was observed for three cycles of STED imaging with >90% STED laser intensity.

One of the reasons tetrazine quenched fluorogenic probes are not widespread in super-resolved imaging techniques is the fact that the quenching power of the bioorthogonal units

(tetrazine and azide) is becoming less and less efficient as the excitation and emission wavelengths approach the biologically preferred red, NIR range. We believe that this is due to the more prominent role of other quenching processes that are present in more conjugated systems and the modulation power exerted by a single unit is less profound. We attempted to address this problem by introducing more than one quencher to cyanine cores and synthesized bisazides and bistetrazines.^{91–93} It was expected that multiple quenching *i.e.* more quenchers together with the formation of a constrained cyclic product with biscyclooctynylated targets enhances fluorogenicity. Indeed, some of these probes were quite promising, however, the complementing technique to produce biscyclooctynylated proteins required double-amber suppressed fusion tags, which made the whole process complicated. Thus, in our other attempt to create more efficient fluorogenic probes that are excitable in the red range we rather combined quenching mechanisms and created polarity sensitive 2'-carboxy silicon-rhodamines linked to tetrazines *via* phenylene and vinylene linkers.^{5e,94} Such polarity and reactivity based fluorogenic scaffolds were found to be quite efficient and resulted in an overall 30–40-fold increase upon conjugation, whereas separate quenching mechanisms are reported to achieve 6–7-fold enhancements.^{5d,10} The best-performing double-fluorogenic probe, SiR-derivative, **53** was applied in the labeling of intracellular vimentin genetically modified with a cyclooctynylated non-canonical amino acid. GSDIM (ground state depletion microscopy followed by individual molecule return) imaging resulted in an improved, 28 nm lateral resolution.^{5e,94}



Scheme 6 Structure of chemically activatable fluorogenic probes.

FRET-based fluorogenicity

Although Förster resonance energy transfer (FRET) systems are not considered as classical fluorogenic systems, their ability to convert one emission wavelength into a different one justifies

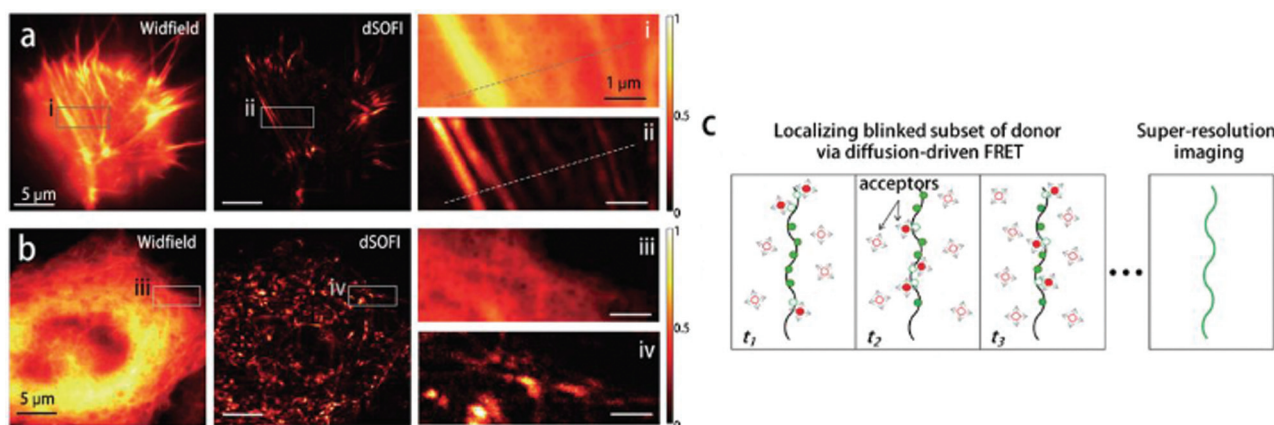


Fig. 13 Confocal (a, e) and STED images (b, f) of actin filaments tagged with phalloidin-BCN and labeled with compound **52** (3 μ M) in mammalian COS7 cells. Images were taken after (a, b) or without washing out (e, f) the dye after 30 min incubation. Enlargement of the boxed areas are shown at (c, g) (confocal, corresponding to a, e) and (d, h) panels (STED, corresponding to (b, f)). $\lambda_{\text{exc}} = 552$ nm with a 660 nm depletion laser for STED images. Color encodes the intensity values of each pixel. (Reprinted from ref. 90 with permission. Copyright 2018 American Chemical Society.)



their discussion in this context. Indeed, there are few examples where a super-resolved image was acquired through FRET signal. FRET-based SRM was first applied by Park and co-workers based on diffusion-assisted FRET.⁹⁵ In their setup, fluorescent donor molecules that label target structures could be stochastically quenched by diffusing acceptor molecules, separating temporally otherwise overlapped fluorescence signals, thus allowing SRM. The approach, named dSOFI (direct super-resolution optical fluctuation imaging) is based on SOFI reconstruction and uses general fluorophores in a conventional epifluorescent microscope setup with light-emitting device (LED) illumination.⁹⁶ Its general utility has been demonstrated in dynamic live-cell imaging of filamentous actin and α -tubulin in NIH 3T3 cells using cerulean and Citrine fluorescent proteins as FRET pairs (Fig. 14).⁹⁵

Cui *et al.* developed a FRET-based dual emission nanoprobe (FREDEN) with improved blinking behaviour for PALM/STORM imaging.⁹⁷ In their setup, Alexa Fluor 647 was attached to CdSSe/ZnS quantum dots (QD) *via* glutathione and used as

an acceptor in FRET to modulate the 610 nm emission of QDs. The FREDEN conjugate shows two distinct emission bands at 610 nm and 670 nm upon excitation of the QDs at 405 nm due to efficient FRET. The two-channel blinking of this conjugate is explained by the excited state reaction of Alexa Fluor 647 that leads to the formation of a non-fluorescent species. The reaction is reversible and is – coincidentally – induced by irradiation with 405 nm light. Thus, the 405 nm excitation wavelength excites the QDs and reforms the fluorescent form of the Alexa dye. The excited state reaction of the Alexa dye affects the FRET process as well, and thus fluctuations at both channels appear. This led to improved blinking and localization precision, compared to AF647 and QD alone, without the need for blinking buffers. Live cell STORM imaging of FREDEN conjugates inside MRC-5 cells was demonstrated.

DNA point accumulation for imaging in nanoscale topography (DNA-PAINT) was a major breakthrough in recent years in SMLM techniques.⁹⁸

DNA-PAINT achieves single-molecule switching and localization by applying freely diffusing dye-labeled single-stranded (ss) DNA (imager strand) that transiently binds to a complementary, unlabeled target-bound ss-DNA strand (docking strand) (Fig. 15a). DNA-PAINT offers several advantages over traditional SRM techniques (no bleaching, wide choice of fluorophores, multicolor-imaging up to 9 color, *etc.*), yet it still has limitations. Since the imager strand is non-fluorogenic, the technique has to rely on optical sectioning methods such as total internal reflection microscopy (TIRF) and limited image acquisition speed in order to eliminate the background fluorescence of freely diffusing imager strands. To address this, Hohng *et al.* introduced the concept of FRET-PAINT.⁹⁹ In their work they designed an unlabeled docking strand that could accommodate two labeled imager strands at a time. The imager strands were designed such that they formed a FRET pair either by using Cy3/Cy5 or AlexaFluor 488/Cy5 labeled strands. Furthermore, the imager strand carrying the acceptor fluorophore had higher binding affinity than the donor labeled strand. Transient binding of the imager strands to the docking strand resulted in stochastic blinking of the acceptor probe when exciting the donor, due to FRET. Since there was no need to eliminate the imager strands' background fluorescence, a higher loading of the freely diffusing imager strands was allowed, enabling faster image acquisition. In a coincident paper, Jungmann and co-workers applied the very same concept, however, in their setup they used an Atto674N labeled docking strand and an Atto488 conjugated imager strand.¹⁰⁰ The high efficiency of the energy transfer was proven along with the highly increased acquisition time as low as 28 s. However, the advantages came at the cost of photobleaching, originally one of the main advantages of DNA-PAINT. When the image acquisition exceeded 400 s, the Atto674N got bleached. Thus they also used a non-labeled docking strand with two recognition sites. Transient, simultaneous binding of donor and acceptor labeled imager strands leads to the same results, however, at the price of reduction in the achievable imaging speed (Fig. 15b). They demonstrated the potential of

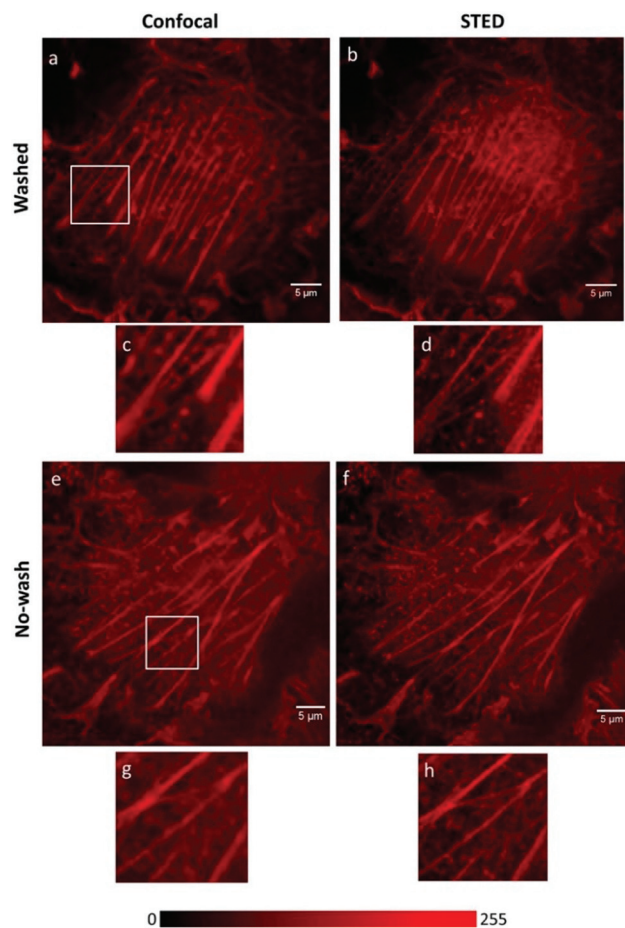


Fig. 14 Live-cell dynamic imaging using dSOFI. (a) Actin-cerulean expressed in NIH 3T3 cells. Cytoplasmic citrine acts as a FRET acceptor. (i, ii) Magnification of boxed areas. (b) Tubulin-cerulean expressed in NIH 3T3 cells. (iii, iv) Magnification of boxed areas. (c) Principle of dSOFI. (Reprinted from ref. 95. with permission. Copyright 2013 Springer Nature.)



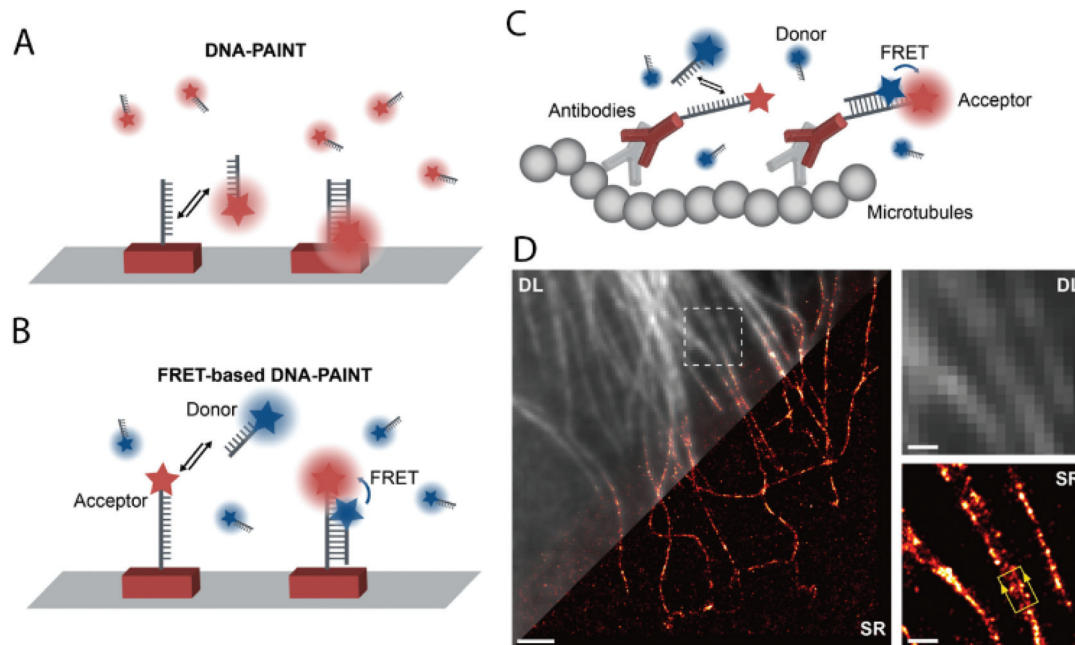


Fig. 15 High speed super-resolution imaging with FRET-PAINT. Principle of DNA-PAINT (a) and FRET-PAINT (b). (c) Labeling scheme using fixed acceptor binding strands. (d) Corresponding diffraction limited and SMLM imaging of the microtubule network in HeLa cells. Scale bar: 2 μm . (Adapted from ref. 100 with permission. Copyright 2017 American Chemical Society.)

this setup in FRET-PAINT imaging of DNA origami structures. In their work, they also utilized their original setup consisting of labeled docking and imager strands in rapid image acquisition of cellular structures. Microtubules of HeLa cells were resolved with a localization precision of 46 nm (Fig. 15c and d).

Heileman and co-workers calculated FRET efficiencies from super-resolved FRET-PAINT images and could assign FRET populations to specific target strands.¹⁰¹ Multiplexed imaging using Atto 674N and Cy3B was also demonstrated on DNA origami with high specificity. FRET-PAINT also allows the measurement of distances in the range of 1–10 nm, extending the spatial resolution of DNA-PAINT near the molecular level.

Conclusions and outlook

Recent developments in super-resolution imaging of biomolecules revealed fine details of cellular structures at sub-diffraction resolutions by means of non-invasive optical methods. The emerged SRM techniques are powerful, yet far from their limits. The biggest constraint that does not allow further improvements is the limited number of suitable probes. High contrast fluorescence imaging of (site)specifically labeled biomolecules by fluorogenic probes is a promising way to address these challenges. The additional level of complexity rationalized in fluorogenic probes that allow site-specific tagging with minimal background and autofluorescence of intracellular targets led to the development of robust labeling schemes with improved resolution. However, the increasing

need for multicolor imaging of multiple targets requires the development of further probes with orthogonal fluorogenic and/or distinct spectral properties. Combination of desirable characteristics such as membrane permeability, selective targeting by means of biologically tolerable chemistries and good spectral features make the design and development of such probes quite challenging as implementation of one such feature often comes at the cost of others. Herein we attempted to highlight the few developments that applied small molecular fluorogenic labels in SRM imaging of biologically relevant structures and listed some design principles suitable for making fluorescent frames fluorogenic. In the future it is expected that existing design principles will result in further fluorogenic probes suitable for SRM imaging and substantial developments are expected in the field of DNA-PAINT, with improved acquisition times. Although not listed herein, as no SRM application is reported so far, design principles, which rely on the *in situ* assembly/generation of fluorescent cores during the conjugation reaction, hold great promise. Such probes possess virtually infinite fluorogenicity and thus provide the users with a superb contrast.

Future developments in the field of labeling techniques in combination with new fluorogenic probes should provide us with robust techniques with unprecedented resolutions both spatially and temporally.

Conflicts of interest

There are no conflicts to declare.



Acknowledgements

The present work was supported by the Hungarian Academy of Sciences within the frames of the “Lendület” Program (LP2013-55/2013). The authors thank Ms Bianka Söveges for her help with the graphical content and Dr Attila Kormos for carefully reading the manuscript.

References

- (a) B. Huang, H. Babcock and X. Zhuang, *Cell*, 2010, **143**, 1047; (b) S. J. Sahl, S. W. Hell and S. Jakobs, *Nat. Rev. Mol. Cell Biol.*, 2017, **11**, 685; (c) A. M. Sydor, K. J. Czymmek, E. M. Puchner and V. Mennella, *Trends Cell Biol.*, 2015, **25**, 730; (d) J. Vangindertael, R. Camacho, W. Sempels, H. Mizuno, P. Dedeker and K.P. F. Janssen, *Methods Appl. Fluoresc.*, 2018, **6**, 022003.
- (a) M. Fernandez-Suarez and A. Y. Ting, *Nat. Rev. Mol. Cell Biol.*, 2008, **9**, 929; (b) Z. Yang, A. Sharma, J. Qi, X. Peng, D. Y. Lee, R. Hu, D. Lin, J. Qu and J. S. Kim, *Chem. Soc. Rev.*, 2016, **45**, 4651; (c) S. van de Linde, S. Aufmkolk, C. Franke, T. Holm, T. Klein, A. Löschberger, S. Proppert, S. Wolter and M. Sauer, *Chem. Biol.*, 2013, **20**, 8; (d) M. Heilemann, S. van de Linde, M. Schüttel, R. Kasper, B. Seefeldt, A. Mukherjee, P. Tinnefeld and M. Sauer, *Angew. Chem., Int. Ed.*, 2008, **47**, 6172.
- (a) D. Jung, K. Min, J. Jung, W. Jang and Y. Kwon, *Mol. Biosyst.*, 2013, **9**, 862; (b) K. Lang and J. W. Chin, *Chem. Rev.*, 2014, **114**, 4764; (c) X. Chen and Y. Wu, *Org. Biomol. Chem.*, 2016, **14**, 5417.
- (a) L. D. Lavis, *Biochemistry*, 2017, **56**, 5165; (b) L. D. Lavis and R. T. Raines, *ACS Chem. Biol.*, 2014, **9**, 855; (c) G. B. Cserép, A. Herner and P. Kele, *Methods Appl. Fluoresc.*, 2015, **3**, 042001; (d) C. P. Toseland, *J. Chem. Biol.*, 2013, **6**, 85.
- (a) Y. Hori and K. Kikuchi, *Curr. Opin. Chem. Biol.*, 2013, **17**, 644; (b) C. Li, A. G. Tebo and A. Gautier, *Int. J. Mol. Sci.*, 2017, **18**, 1473; (c) L. G. Meimetis, J. C. T. Carlson, R. J. Giedt, R. H. Kohler and R. Weissleder, *Angew. Chem., Int. Ed.*, 2014, **53**, 7531; (d) A. Wiczorek, P. Werther, J. Euchner and R. Wombacher, *J. Chem. Biol.*, 2017, **8**, 1506; (e) E. Kozma, G. Estrada Girona, G. Paci, E. A. Lemke and P. Kele, *Chem. Commun.*, 2017, **53**, 6696; (f) J. Vogelsang, C. Steinhauer, C. Forthmann, I. H. Stein, B. Person-Skegro, T. Cordes and P. Tinnefeld, *ChemPhysChem*, 2010, **11**, 2475.
- M. Beija, C. A. M. Afonso and J. M. G. Martinho, *Chem. Soc. Rev.*, 2009, **38**, 2410.
- Y. Koide, Y. Urano, K. Hanaoka, T. Terai and T. Nagano, *ACS Chem. Biol.*, 2011, **6**, 600.
- Y. Kushida, T. Nagano and K. Hanaoka, *Analyst*, 2015, **140**, 685.
- X. Chen, T. Pradhan, F. Wang, J. S. Kim and J. Yoon, *Chem. Rev.*, 2012, **112**, 1910.
- G. Lukinavicius, K. Umezawa, N. Olivier, A. Honigmann, G. Yang, T. Plass, V. Mueller, L. Reymond, I. R. Correa, Z. G. Luo, C. Schultz, E. A. Lemke, P. Heppenstall, C. Eggeling, S. Manley and K. Johnsson, *Nat. Chem.*, 2013, **5**, 139.
- G. Lukinavicius, L. Reymond, E. D'Este, A. Masharina, F. Göttfert, H. Ta, A. Güther, M. Fournier, S. Rizzo, H. Waldmann, C. Blaukopf, C. Sommer, D. W. Gerlich, H. D. Arndt, S. W. Hell and K. Johnsson, *Nat. Methods*, 2014, **11**, 731.
- N. V. Klementieva, L. B. Snopova, N. N. Prodanets, O. E. Furman, V. V. Dudenkova, E. Zagaynova, K. A. Lukyanov and A. S. Mishin, *Anticancer Res.*, 2016, **36**, 5287.
- G. Lukinavicius, C. Blaukopf, E. Pershagen, A. Schena, L. Reymond, E. Derivery, M. Gonzales-Gaitan, E. D'Este, S. W. Hell, D. W. Gerlich and K. Johnsson, *Nat. Commun.*, 2015, **6**, 8497.
- C. Uttamapinant, J. D. Howe, K. Lang, V. Beranek, L. Davis, M. Mahesh, N. P. Barry and J. W. Chin, *J. Am. Chem. Soc.*, 2015, **137**, 4602.
- J. Masch, H. Steffens, J. Fischer, J. Engelhardt, J. Hubrich, J. Keller-Findeisen, E. D'Este, N. T. Urban, S. G. N. Grant, S. J. Sahl, D. Kamin and S. W. Hell, *Proc. Natl. Acad. Sci. USA*, 2018, **115**, E8047.
- G. Lukinavicius, L. Reymond, K. Umezawa, O. Sallin, E. D'Este, F. Göttfert, H. Ta, S. W. Hell, Y. Urano and K. Johnsson, *J. Am. Chem. Soc.*, 2016, **138**, 9365.
- K. Kolmakov, E. Hebisch, T. Wolfram, L. A. Nordwig, C. A. Wurm, H. Ta, V. Westphal, V. N. Belov and S. W. Hell, *Chem. – Eur. J.*, 2015, **21**, 13344.
- A. N. Butkevich, G. Y. Mitronova, S. C. Sidenstein, J. L. Klocke, D. Kamin, D. N. H. Meineke, E. D'Este, P. T. Kraemer, J. G. Danzl, V. N. Belov and S. W. Hell, *Angew. Chem., Int. Ed.*, 2016, **55**, 3290.
- K. Kolmakov, V. N. Belov, C. A. Wurm, B. Harke, M. Leutenegger, C. Eggeling and S. W. Hell, *Eur. J. Org. Chem.*, 2010, **2010**, 3593.
- A. N. Butkevich, G. Y. Mitronova, S. C. Sidenstein, J. L. Klocke, D. Kamin, D. N. H. Meineke, E. D'Este, P. T. Kraemer, J. G. Danzl, V. N. Belov and S. W. Hell, *Angew. Chem., Int. Ed.*, 2016, **55**, 3290.
- G. Lukinavicius, G. Y. Mitronova, S. Schnorrenberg, A. N. Butkevich, H. Barthel, V. N. Belov and S. W. Hell, *Chem. Sci.*, 2018, **9**, 3324.
- A. N. Butkevich, H. Ta, M. Ratz, S. Stoldt, S. Jakobs, V. N. Belov and S. W. Hell, *ACS Chem. Biol.*, 2018, **13**, 475.
- A. N. Butkevich, V. N. Belov, K. Kolmakov, V. V. Sokolov, H. Shojaei, S. C. Sidenstein, D. Kamin, J. Matthias, R. Vlijm, J. Engelhardt and S. W. Hell, *Chem. – Eur. J.*, 2017, **50**, 12114.
- H. Takakura, Y. Zhang, R. S. Erdmann, A. D. Thompson, Y. Lin, B. McNellis, F. Rivera-Molina, S. Uno, M. Kamiya, Y. Urano, J. E. Rothman, J. Bewersdorf, A. Schepartz and D. Toomre, *Nat. Biotechnol.*, 2017, **35**, 773.



- 25 P. J. Macdoldnald, S. Gayda, R. A. Haack, Q. Ruan, R. J. Himmelbach and S. Y. Tetin, *Anal. Chem.*, 2018, **90**, 9165.
- 26 S. 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.
- 27 S. Uno, M. Kamiya, A. Morozumi and Y. Urano, *Chem. Sci.*, 2018, **54**, 102.
- 28 M. V. Sednev, V. N. Belov and S. W. Hell, *Methods Appl. Fluoresc.*, 2015, **3**, 042004.
- 29 A. N. Butkevich, G. Lukinavicius, E. D'Este and S. W. Hell, *J. Am. Chem. Soc.*, 2017, **139**, 12378.
- 30 S. R. Adams, R. E. Campbell, L. A. Gross, B. R. Martin, G. K. Walkup, Y. Yao, J. Llopis and R. Y. Tsien, *J. Am. Chem. Soc.*, 2002, **124**, 6063.
- 31 J. Enninga, J. Mounier, P. Sansonetti and G. T. Van Nhieu, *Nat. Methods*, 2005, **2**, 959.
- 32 B. A. Griffin, S. R. Adams and R. Y. Tsien, *Science*, 1998, **281**, 269.
- 33 J. Fölling, M. Bossi, H. Bock, R. Medda, C. A. Wurm, B. Hein, S. Jakobs, C. Eggeling and S. W. Hell, *Nat. Methods*, 2008, **5**, 943.
- 34 M. Lelek, F. Di Nunzio, R. Henriques, P. Charneau, N. Arhel and C. Zimmer, *Proc. Natl. Acad. Sci. U. S. A.*, 2012, **109**, 8564.
- 35 M. Lelek, F. Di Nunzio and C. Zimemr, *Methods Mol. Biol.*, 2014, **1174**, 183.
- 36 H. Park, G. T. Hanson, S. R. Duff and P. R. Selvin, *J. Microsc.*, 2004, **216**, 199.
- 37 C. Flors, *J. Microsc.*, 2013, **251**, 1.
- 38 A. I. Dragan, J. R. Casas-Finet, E. S. Bishop, R. J. Strouse, M. A. Schenerman and C. D. Geddes, *Biophys. J.*, 2010, **99**, 3010.
- 39 A. Larsson, C. Carlsson, M. Jonsson and B. Albinsson, *J. Am. Chem. Soc.*, 1994, **116**, 8459.
- 40 H. S. Rye, S. Yue, D. E. Wemmer and M. A. Quesada, *Nucleic Acids Res.*, 1992, **20**, 2803.
- 41 C. Flors, C. N. J. Ravarani and D. T. F. Dryden, *ChemPhysChem*, 2009, **10**, 2201.
- 42 G. Park, S. K. Chakkarapani, S. Ju, S. Ahn and S. H. Kang, *Chin. Chem. Lett.*, 2018, **29**, 505.
- 43 S. K. Chakkarapani, G. Park and S. H. Kang, *Chin. Chem. Lett.*, 2015, **26**, 1490.
- 44 X. Meng, W. W. Cai and D. C. Schwartz, *J. Biomol. Struct. Dyn.*, 1996, **13**, 945.
- 45 C. Flors, *Photochem. Photobiol. Sci.*, 2010, **9**, 643.
- 46 F. Persson, P. Bingen, T. Staudt, J. Engelhardt, J. O. Tegenfeld and S. W. Hell, *Angew. Chem., Int. Ed.*, 2011, **50**, 5581.
- 47 T. Idziorek, J. Estaquier, F. De Bels and J. C. Ameisen, *J. Immunol. Methods*, 1995, **185**, 249.
- 48 H. Miller, Z. Zhou, A. J. M. Wollman and M. C. Leake, *Methods*, 2015, **88**, 81.
- 49 D. Wlodkowic, J. Skommer and Z. Darzynkiewicz, *Cytometry, Part A*, 2008, **73**, 496.
- 50 C. Flors, *Biopolymers*, 2011, **95**, 290.
- 51 X. Yan, R. C. Habbersett, J. M. Cordek, J. P. Nolan, T. M. Yoshida, J. H. Jett and B. L. Marrone, *Anal. Biochem.*, 2000, **286**, 138.
- 52 S. Bakshi, H. Choi, N. Rangarajan, K. J. Barns, B. P. Bratton and J. C. Weisshaar, *Appl. Environ. Microbiol.*, 2014, **80**, 4977.
- 53 (a) A. S. Backer, M. Y. Lee and W. E. Moerner, *Optica*, 2016, **3**; (b) F. Hövelmann, I. Gáspár, J. Chamiolo, M. Kasper, J. Steffen, A. Ephrussi and O. Seitz, *Chem. Sci.*, 2016, **7**, 128.
- 54 L. Jullien and A. Gautier, *Methods Appl. Fluoresc.*, 2015, 042007.
- 55 M. P. Bruchez, *Curr. Opin. Chem. Biol.*, 2015, **27**, 18.
- 56 J. A. J. Fitzpatrick, Q. Yan, J. J. Sieber, M. Dyba, U. Schwarz, C. Szent-Gyorgyi, C. A. Woolford, P. B. Berget, A. S. Waggoner and M. P. Bruchez, *Bioconjugate Chem.*, 2009, **20**, 1843.
- 57 C. Szent-Gyorgyi, B. F. Schmidt, Y. Creeger, G. W. Fisher, K. L. Zakel, S. Adler, J. A. J. Fitzpatrick, C. A. Woolford, Q. Yan, K. V. Vasilev, P. B. Berget, M. P. Bruchez, J. W. Jarvik and A. Waggoner, *Nat. Biotechnol.*, 2008, **26**, 235–240.
- 58 Q. Yan, S. L. Schwartz, S. Maji, F. Huang, C. Szent-Gyorgyi, D. S. Lidke, K. A. Lidke and M. P. Bruchez, *ChemPhysChem*, 2014, **15**, 687.
- 59 N. G. Bozhanova, M. S. Baranov, N. V. Klementieva, K. S. Sarkisyan, A. S. Gavrikov, I. V. Yampolsky, E. V. Zagaynova, S. A. Lukyanov, K. A. Lukyanov and A. S. Mishin, *Chem. Sci.*, 2017, **8**, 7138.
- 60 S. Saurabh, A. M. Perez, C. J. Comerci, L. Shapiro and W. E. Moerner, *J. Am. Chem. Soc.*, 2016, **138**, 10398.
- 61 S. Saurabh, A. M. Perez, C. J. Comerci, L. Shapiro and W. E. Moerner, *Curr. Protoc. Cell Biol.*, 2017, **75**, 4.32.1.
- 62 L. A. Perkins, Q. Yan, B. F. Schmidt, D. Kolodienznyi, S. Saurabh, M. B. Larsen, S. C. Watkins, L. Kremer and M. P. Bruchez, *Biochemistry*, 2018, **57**, 861.
- 63 S. R. Adams and R. Y. Tsien, *Annu. Rev. Physiol.*, 1993, **55**, 755.
- 64 P. Klán, T. Solomek, C. G. Bochet, A. Blanc, R. Givens, M. Rubina, V. Popik, A. Kostikov and J. Wirz, *Chem. Rev.*, 2013, **113**, 119.
- 65 J. B. Grimm, A. J. Sung, W. R. Legant, P. Hulamm, S. M. Matlosz, E. Betzig and L. D. Lavis, *ACS Chem. Biol.*, 2013, **8**, 1303.
- 66 L. M. Wysocki, J. B. Grimm, A. N. Tkachuck, T. A. Brown, E. Betzig and L. D. Lavis, *Angew. Chem., Int. Ed.*, 2011, **50**, 11206.
- 67 J. B. Grimm, T. Klein, B. G. Kopek, G. Shtengel, H. F. Hess, M. Sauer and L. D. Lavis, *Angew. Chem., Int. Ed.*, 2016, **55**, 1723.
- 68 S. Banala, D. Maurel, S. Manley and K. Johnsson, *ACS Chem. Biol.*, 2012, **7**, 289.
- 69 S. Hauke, A. von Appen, T. Quidwai, J. Ries and R. Wombacher, *Chem. Sci.*, 2017, **8**, 559.
- 70 V. N. Belov, C. A. Wurm, V. P. Boyarskiy, S. Jakobs and S. W. Hell, *Angew. Chem., Int. Ed.*, 2010, **49**, 3520.



- 71 S. J. Lord, N. R. Conley, H. D. Lee, R. Samuel, N. Liu, R. J. Twieg and W. E. Moerner, *J. Am. Chem. Soc.*, 2008, **130**, 9204.
- 72 H. D. Lee, S. J. Lord, S. Iwanaga, K. Zhan, H. Xie, J. C. Williams, H. Wang, G. R. Bowman, E. D. Goley, L. Shapiro, R. J. Twieg, J. Rhao and W. E. Moerner, *J. Am. Chem. Soc.*, 2010, **132**, 15099.
- 73 M. K. Lee, J. Williams, R. J. Twieg, J. Rao and W. E. Moerner, *Chem. Sci.*, 2013, **4**, 220.
- 74 H. Willwohl, J. Wolfrum and R. Gleiter, *Laser Chem.*, 1989, **10**, 63.
- 75 J. Fölling, V. Belov, R. Kunetsky, R. Medda, A. Schönle, A. Egner, C. Eggeling, M. Bossi and S. W. Hell, *Angew. Chem., Int. Ed.*, 2007, **46**, 6266.
- 76 M. Bossi, J. Fölling, V. N. Belov, V. P. Boyarskiy, R. Medda, A. Egner, C. Eggeling, A. Schönle and S. W. Hell, *Nano Lett.*, 2008, **8**, 2463.
- 77 V. N. Belov, M. L. Bossi, J. Fölling, V. P. Boyarskiy and S. W. Hell, *Chem. – Eur. J.*, 2009, **15**, 10762.
- 78 M. K. Lee, P. Rai, J. Williams, R. J. Twieg and W. E. Moerner, *J. Am. Chem. Soc.*, 2014, **136**, 14003.
- 79 K. Kolmakov, C. Wurm, M. V. Sednev, M. L. Bossi, V. N. Belov and S. W. Hell, *Photochem. Photobiol. Sci.*, 2012, **11**, 522.
- 80 E. A. Halabi, Z. Thiel, N. Trapp, D. Pinotsi and P. Rivera-Fuentes, *J. Am. Chem. Soc.*, 2017, **139**, 13200.
- 81 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.
- 82 Y. Shoji, A. Yagi, M. Horiuchi, M. Morimoto and M. Irie, *Isr. J. Chem.*, 2013, **53**, 303.
- 83 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.
- 84 B. Roubinet, M. Weber, H. Shojaei, M. Bates, M. L. Bossi and V. N. Belov, *J. Am. Chem. Soc.*, 2017, **139**, 6611.
- 85 E. Deniz, M. Tomasulo, J. Cusido, I. Yildiz, M. Petriella, M. L. Bossi, S. Sortino and F. M. Raymo, *J. Phys. Chem. C*, 2012, **116**, 6058.
- 86 G. B. Cserép, A. Herner and P. Kele, *Methods Appl. Fluoresc.*, 2015, **3**, 042001.
- 87 (a) A. Herner, I. Nikic, M. Kállay, E. A. Lemke and P. Kele, *Org. Biomol. Chem.*, 2013, **11**, 3297; (b) P. Shieh, M. S. Siegrist, A. J. Cullen and C. R. Bertozzi, *Proc. Natl. Acad. Sci. USA*, 2014, **111**, 5456.
- 88 (a) J. Yang, J. Seckute, C. M. Cole and N. K. Devaraj, *Angew. Chem., Int. Ed.*, 2012, **51**, 7476; (b) N. K. Devaraj, S. Hildebrand, R. Upadhyay, R. Mazitschek and R. Weissleder, *Angew. Chem., Int. Ed.*, 2010, **49**, 2869; (c) D. S. Liu, A. Tangpeerachaikul, R. Selvaraj, M. T. Taylor, J. M. Fox and A. Y. Ting, *J. Am. Chem. Soc.*, 2012, **134**, 792; (d) J. C. T. Carlson, L. G. Meimetis, S. A. Hildebrand and R. Weissleder, *Angew. Chem., Int. Ed.*, 2013, **52**, 6917; (e) G. Knorr, E. Kozma, A. Herner, E. A. Lemke and P. Kele, *Chem. – Eur. J.*, 2016, **22**, 8972.
- 89 Y. Lee, W. Cho, J. Sung, E. Kim and S. B. Park, *J. Am. Chem. Soc.*, 2018, **140**, 974.
- 90 G. Knorr, E. Kozma, J. M. Schaart, K. Németh, G. Török and P. Kele, *Bioconjugate Chem.*, 2018, **29**, 1312.
- 91 O. Demeter, E. A. Fodor, M. Kállay, G. Mező, K. Németh, P. T. Szabó and P. Kele, *Chem. – Eur. J.*, 2016, **22**, 6382.
- 92 O. Demeter, A. Kormos, C. Koehler, G. Mező, K. Németh, E. Kozma, L. Takács, E. A. Lemke and P. Kele, *Bioconjugate Chem.*, 2017, **28**, 1552.
- 93 A. Kormos, C. Koehler, E. A. Fodor, Z. R. Rutkai, M. E. Martin, G. Mező, E. A. Lemke and P. Kele, *Chem. – Eur. J.*, 2018, **24**, 8841.
- 94 E. Kozma, G. Paci, G. Estrada Girona, E. A. Lemke and P. Kele, *Methods Mol. Biol.*, 2018, **1728**, 337.
- 95 S. Cho, J. Jang, C. Song, H. Lee, P. Ganesan, T. Yoon, M. W. Kim, M. C. Choi, H. Ihee, W. D. Heo and Y. Park, *Sci. Rep.*, 2013, **3**, 1208.
- 96 T. Dertinger, R. Colyer, G. Iyer, S. Weiss and J. Enderlein, *Proc. Natl. Acad. Sci. U. S. A.*, 2009, **106**, 22287.
- 97 S. Zong, X. Jiang, Z. Wang, C. Chen, J. Lu, L. Wang, D. Zhu and Y. Cui, *Nanoscale*, 2016, **8**, 19110.
- 98 (a) J. Schnitzbauer, M. T. Strauss, T. Schlichthaerle, F. Scheuder and R. Jungmann, *Nat. Protoc.*, 2017, **12**, 1198; (b) R. Jungmann, C. Steinhauer, M. Schieble, A. Kuzyk, P. Tinnefeld and F. C. Simmel, *Nano Lett.*, 2010, **10**, 4756.
- 99 J. Lee, S. Park, W. Kang and S. Hohng, *Mol. Brain*, 2017, **10**, 63.
- 100 A. Auer, M. T. Strauss, T. Schlichthaerle and R. Jungmann, *Nano Lett.*, 2017, **17**, 6428.
- 101 N. S. Deubner-Helfmann, A. Auer, M. T. Strauss, S. Malkusch, M. S. Dietz, H. D. Barth, R. Jungmann and M. Heilemann, *Nano Lett.*, 2018, **18**, 4626.

