# A Caged, Localizable Rhodamine Derivative for Superresolution Microscopy

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ⓢ *Supporting Information*

ABSTRACT: A caged rhodamine 110 derivative for the specific labeling of SNAP-tag fusion proteins is introduced. The caged rhodamine 110 derivative permits the labeling of cell surface proteins in living cells and of intracellular proteins in fixed cells. The probe requires only a single caging group to maintain the fluorophore in a non-fluorescent state and becomes highly fluorescent after uncaging. The high contrast ratio is confirmed both in bulk and at the single molecule level. This property, together with its high photon yield makes it an excellent dye for photoactivated localization microscopy (PALM), as we demonstrate here.
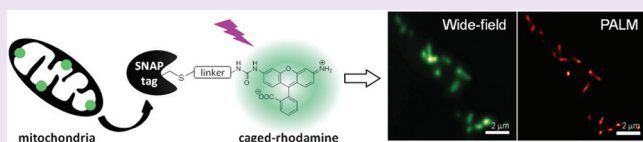
The caging of a fluorophore is defined as its derivatization with a photocleavable group that converts the fluorophore into a non-fluorescent state. As the uncaging and resulting recovery of fluorescence through a light pulse can be achieved with high spatial and temporal resolution, caged fluorophores have become important tools in studying numerous biological processes, including cell lineage and protein trafficking.[1,2] Another attractive application of caged fluorophores is in photoactivated localization microscopy (PALM), which permits imaging of cellular components with nanometer resolution.[3] In PALM, sparse subsets of dye molecules are stochastically activated or uncaged, imaged, and bleached over the course of thousands of raw images. Since the density of activated molecules depends on the intensity of the activation light, we can ensure that molecules in each image are spatially well-separated and thus can be localized by fitting to the point-spread function.[4] The certainty with which a molecule can be localized depends inversely on the square-root of the number of photons collected, so the high photon yields of synthetic fluorophores offer a significant advantage over fluorescent proteins (FPs) in this respect. Furthermore, there are few options for good photoactivatable FPs in the green part of the spectrum for single-molecule-based imaging, and those that do exist suffer from high background and low contrast ratios. Caged fluorophores that could be coupled to selected proteins in living or fixed cells therefore have the potential to become important tools for superresolution microscopy. Caged versions of several fluorophores such as fluorescein and rhodamine have been reported, but caging of these fluorophores required the attachment of two photocleavable groups.[1,2,5,6] The removal of two caging groups necessitates a longer irradiation time to recover the whole fluorescence from the illuminated sample. Moreover, attempts to selectively uncage only a subset of a population of caged fluorophores will mostly generate fluorophores where only one of the two caging groups has been removed. In the case of caged fluorescein and rhodamine

derivatives, the partially uncaged fluorophore displays only relatively weak fluorescence compared to the fully uncaged fluorophore, and therefore this approach does not exploit the full potential of synthetic fluorophores for PALM.[6] Caged versions of the fluorophore Tokyogreen with only one photocleavable group and of rhodamine with a 2-diazoketone (COCNN) caging group have been reported,[7,8] but the specific coupling of these probes to individual proteins has not been described yet. Recently, a caged version of a dicyanomethylenedihydrofuran (DCDHF) fluorophore has been described that can be selectively coupled to Halo-tag fusion proteins.[9] DCDHF was efficiently caged by replacing a crucial amino group with a photolabile azide; the probe is then uncaged by photoconversion of the azide back to the amino group. The potential of this caged DCDHF was demonstrated by performing PALM of Halo-tag fusion proteins in bacteria and mammalian cells. In summary, there is a generally acknowledged need for new localizable caged fluorophores.

Here we present a caged version of rhodamine 110 (Rh_{110}) that can be coupled to SNAP-tag fusion proteins and that is suitable for PALM (Figure 1). The large fluorescence increase upon uncaging and the brightness and stability of the SNAP-tag-bound fluorophore make it an attractive probe for superresolution imaging and the study of dynamic cellular processes.

The design of our caged rhodamine probe **1** (BG-cRhod) was based on the observation that ureated derivatives of Rh_{110} retained much of the fluorescence of the parental compound.[10] Particularly, the relative fluorescence intensity (a product of extinction coefficient and quantum yield) of the ureated derivative is 35% that of Rh_{110}. The relative ease with which
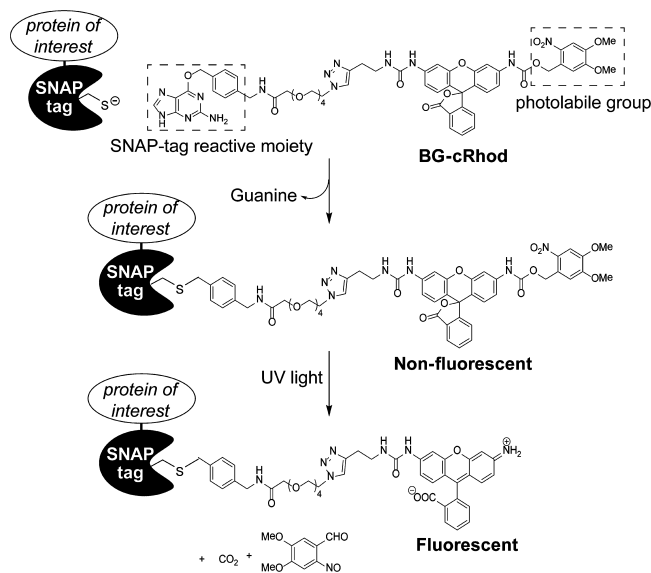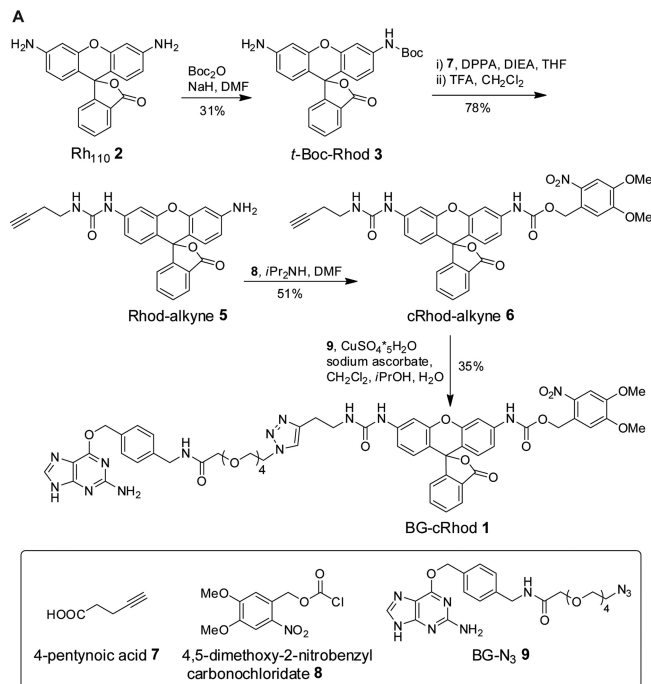
Figure 1. Design of the caged rhodamine BG-cRhod and its use for the labeling of SNAP-tag fusion proteins.

Figure 2. (A) Synthesis of BG-cRhod. (B) *In vitro* characterization of BG-cRhod. Fluorescence increase of free BG-cRhod (triangles) and its SNAP-tag conjugate (circles) before (open) and after (filled) uncaging. An enlargement of the fluorescence spectrum of caged BG-cRhod and caged SNAP-tag-bound cRhod is shown in Supporting Information. Boc$_2$O = di-*tert*-butyldicarbonate, DIEA = *N*,*N*-diisopropylethylamine, DPPA = diphenylphosphoryl azide, *i*Pr$_2$NH = diisopropylamine, TFA = trifluoroacetic acid, DMF = dimethylformamide, THF = tetrahydrofuran.

ureated Rh$_{110}$ derivatives can be synthesized and their good spectroscopic properties have triggered the generation of a number of Rh$_{110}$-based probes for various applications.[11−13] On the basis of these reports, we envisioned the synthesis of a caged Rh$_{110}$ derivate for the reaction with SNAP-tag fusion proteins. SNAP-tag is a protein tag that reacts with O^6-benzylguanine (BG) derivatives carrying different probes *in vitro* and *in vivo*.[14,15] In BG-cRhod, a BG is attached *via* a urea linkage to Rh$_{110}$, while the attachment of a photocleavable 4,5-dimethoxy-2-nitrobenzyl group (DMNB) to the other amino group of Rh$_{110}$ would force the probe in its non-fluorescent lactone configuration (Figure 1). We anticipated that SNAP-tag fusion proteins could be labeled with BG-cRhod and subsequently uncaged through irradiation with UV light.

Starting from commercially available Rh$_{110}$, BG-cRhod was synthesized in five steps in a total yield of 4.3% (Figure 2, panel A). The fluorescence properties of BG-cRhod were studied *in vitro* before and after reaction with SNAP-tag. First, we confirmed by HPLC and mass spectroscopy that the photolysis of BG-cRhod generated the corresponding BG Rh$_{110}$ derivative BG-Rhod (Supplementary Figure 1). As expected, BG-cRhod is non-fluorescent ($\Phi_{fl} < 0.001$) and became fluorescent upon irradiation with 365 nm light, although the fluorescence quantum yield of BG-cRhod was relatively low ($\Phi_{fl} = 0.074$) (Figure 2, panel B). We then demonstrated that BG-cRhod permits the labeling of SNAP-tag with cRhod (Supplementary Figures 3 and 4). As observed on BG-cRhod, SNAP-tag-bound cRhod is virtually non-fluorescent ($\Phi_{fl} < 0.002$). However, the fluorescence quantum yield of uncaged SNAP-tag-bound cRhod increased to 0.36, which is about 5 times higher than that of the uncaged free substrate (Figure 2, panel B). The observed increase of fluorescence upon uncaging of SNAP-tag-bound cRhod is at least 200-fold. We assume that the background fluorescence observed prior to uncaging is most likely due to the presence of small amounts of already uncaged BG-cRhod. The low fluorescence quantum yield of BG-cRhod is most likely due to quenching through benzylguanine.[16,17] Similar behavior has already been observed for other fluorescent BG derivatives used for SNAP-tag labeling.[18−20] The observed quenching in uncaged BG-cRhod relative to that

of SNAP-tag-bound uncaged cRhod is an advantageous feature of our probe because it reduces potential background fluorescence from unreacted, uncaged probe.

We then investigated the use of BG-cRhod for the labeling of cell surface receptors and subsequent uncaging of labeled protein. We chose the *β*-adrenergic receptor *β*2AR as a model protein as its SNAP-tag fusion has previously been shown to be functional.[21] HEK293T cells were transiently transfected with a plasmid for expression of SNAP-*β*2AR and consecutively labeled with BG-cRhod (0.3 *μ*M, 1 h) and BG-Cy5 (2 *μ*M, 10 min). Simultaneous labeling with BG-Cy5 was performed to identify transfected cells expressing SNAP-*β*2AR prior to uncaging. Using a 488 nm laser for excitation, no significant fluorescence of BG-cRhod-labeled SNAP-*β*2AR could be detected prior to uncaging (Figure 3, panel A). However, irradiation with a 405 nm laser resulted in uncaging of labeled SNAP-*β*2AR, as demonstrated by the strong fluorescence signal observed at the plasma membrane of transfected cells (Figure 3,
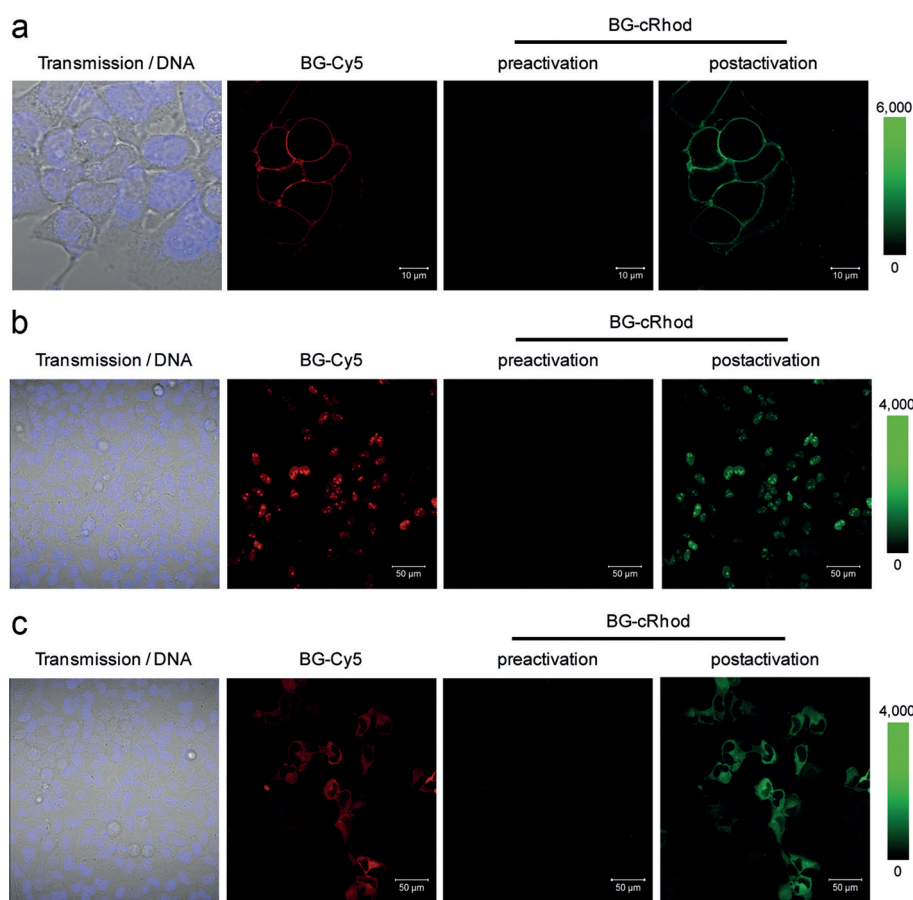
Figure 3. (A) Confocal fluorescence images of living HEK293T cells expressing SNAP-β2 adrenergic receptor. Transiently expressed SNAP-β2AR was labeled with BG-cRhod (green) and then with BG-Cy5 (red). Nuclear DNA staining with Hoechst 33342 was used as a reference (blue). Images were taken before and after UV irradiation. (B, C) Confocal fluorescence images of fixed U2OS cells expressing SNAP-NLS and SNAP-MEK1. (B) Transiently expressed SNAP-NLS (nucleus) and (C) SNAP-MEK1 (cytoplasm) were labeled with the SNAP-tag substrates BG-cRhod (green) and then with BG-Cy5 (red). Nuclear DNA staining with Hoechst 33342 was used as a reference (blue). Images were taken before and after uncaging at 405 nm. Cells were fixed with paraformaldehyde prior to staining.

panel A). The possibility to uncage BG-cRhod-labeled proteins with a 405 nm laser is important for practical applications as this laser line is found on most confocal microscopes and illumination at 405 nm is less phototoxic than illumination with UV light.

We then tested if BG-cRhod permits the labeling of intracellular SNAP-tag fusion proteins. For these experiments, a fusion protein of SNAP-tag with the kinase MEK1 (SNAP-MEK1) was chosen as an example of a cytosolic protein and transiently expressed in U2OS cells. Although labeling of SNAP-MEK1 with the cell-permeable fluorescent SNAP-tag substrate TMR-star was achieved, no significant labeling with BG-cRhod could be observed (Supplementary Figure 5). The failure to label intracellular SNAP-tag fusion proteins with BG-cRhod in living cells limits the application range of this substrate. However, the labeling of intracellular SNAP-tag fusion proteins with BG-cRhod could be achieved with invasive approaches such as bead-loading, microinjection or electroporation; all of these approaches have been previously been shown to permit the labeling of SNAP-tag fusion proteins with cell-impermeable substrates.[22−24]

Superresolution imaging using point-based localization (PALM, STORM, *etc.*) for the elucidation of biological structures is generally performed on fixed cells.[25,26] We

therefore investigated if the selective labeling of SNAP-tag fusion proteins with BG-cRhod can be achieved after fixation. To this end, we performed labeling experiments with BG-cRhod on fixed U2OS cells expressing SNAP-tag fusion proteins localized either in the nucleus (SNAP-NLS) or in the cytoplasm (SNAP-MEK1) (Figure 3, panels B and C). Cells were fixed with paraformaldehyde and consecutively labeled with BG-cRhod (0.3 μM, 1 h) and then BG-Cy5 (2 μM, 10 min) to facilitate the identification of transiently transfected cells. After labeling with BG-cRhod and uncaging with a 405 nm laser, cells showed the expected localized fluorescence signals. In contrast, no significant fluorescence was detected prior to photoactivation. These experiments demonstrate that BG-cRhod is suitable for the labeling of SNAP-tag fusion proteins in fixed cells and suggest that the probe is suitable for PALM.

For PALM imaging, we used BG-cRhod to label SNAP-tag fusion proteins localized to mitochondria (Mito-SNAP). U2OS cells expressing Mito-SNAP were fixed with paraformaldehyde and labeled with BG-cRhod (0.3 μM, 2 h). Before PALM imaging, cells were exposed to a high level of 488 nm light to photobleach any previously uncaged molecules. The sum of all raw images corresponds to a time-integrated epifluorescence image (Figure 4, panel A), which is shown for comparison with the rendered super-resolution PALM image (Figure 4, panel
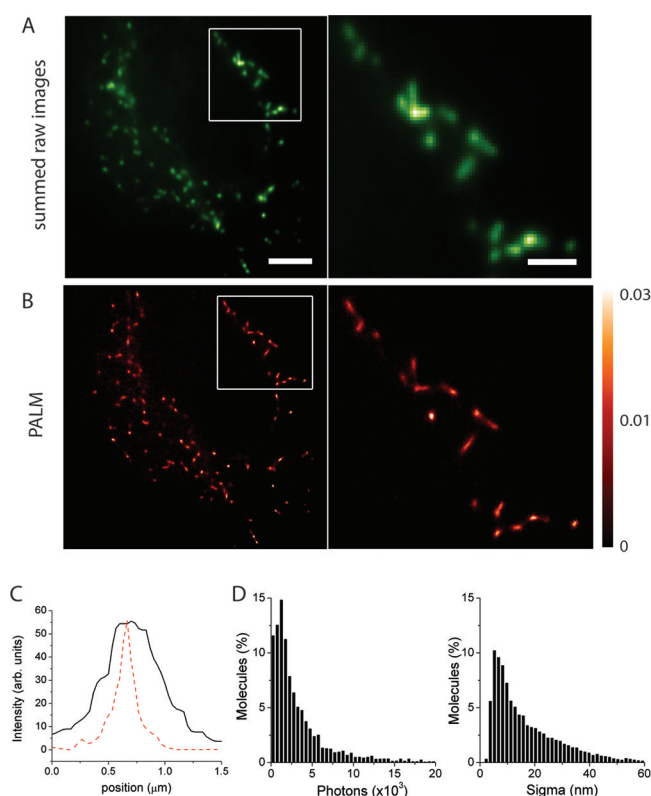
Figure 4. (A) Summed raw images of fixed U2OS cells expressing SNAP-Mito (mitochondria). Quasi-stable cells expressing a mitochondrial targeting sequence were labeled with BG-cRhod, and a stack of 5000 images was collected with sparse activation using a 405 nm laser and then summed to create the equivalent of an epi-fluorescence image. (B) PALM images. The same stack of images was processed for PALM. Color bar corresponds to the probability density of finding a molecule in units of molecules per nm^2. Number of molecules: 142,433 (left), 16,960 (right). (C) Intensity profiles across a single mitochondria were analyzed to extract the full width half max from a Gaussian fit for summed raw (solid line, fwhm = 517 nm) and PALM (dashed line, fwhm = 135 nm) images. (D) Histograms of the number of photons and molecular localization uncertainty. The molecules shown in panel B were analyzed for their photon yield (mean = 3488; median = 2027) and localization uncertainties, σ (mean = 16 nm, median = 11.7 nm). Scale bar (left) is 5 μm, scale bar (right, zoom) is 2 μm.

B). As expected, features that are obscured in wide-field imaging are sharply resolved in the PALM image. We used the information from the raw data to obtain statistics on molecular photon yields and localization precision. In comparison with the best red FPs or PA-GFP, which give on average 300−800 photons per molecule (ref 27 and Supplementary Figure S7), we found a mean value of 3488 photons/molecule, an increase in photon yields of a factor of ∼5−10. This increased molecular brightness is translated into an improvement in localization; because there is a rather long tail in the distribution this is demonstrated by the median value of uncertainty in position, which is 11.7 nm, a factor of ∼2 better than for the red FPs. The mode of the distribution is even lower, 6.1 nm, a factor of ∼3 better than for the red FPs. Thus, cRhod is an excellent label for PALM imaging in general.

Stochastic optical reconstruction microscopy (STORM), direct STORM (dSTORM), and "ground state depletion microscopy followed by individual molecule return" (GSDIM) are alternative approaches toward superresolution microscopy.[28−30] These methods rely on synthetic fluorophores subjected to reduction−oxidation or triplet-state cycling to induce blinking and therefore temporally separated signals as required by point-localization superresolution microscopy. Compared to PALM, these approaches have been able to exploit the higher photon counts available from synthetic dyes, resulting in higher molecular localization precision. However, for high densities of molecules, it can be difficult to fully initialize the system into the dark state. Furthermore, it is not straightforward to control the rate of molecules cycling, to maintain their temporal separation. Caged dyes such as cRhod provide an advantage in that their initial state is largely the caged one, with the possibility of easily photobleaching any small uncaged population. Moreover, their uncaging is well-controlled as a function of the intensity of the UV uncaging light. Another interesting aspect of cRhod for future practical applications is the possibility to use it in combination with red FPs or synthetic photoswitchable probes for multicolor imaging. In light of these general considerations and the favorable spectroscopic properties of cRhod, we expect that this dye will become a useful addition to the already available dyes for superresolution microscopy.

■ ASSOCIATED CONTENT

Ⓢ Supporting Information

Detailed experimental procedures and characterization of compounds. This material is available free of charge *via* the Internet at http://pubs.acs.org.

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Author Contributions

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■ ACKNOWLEDGMENTS

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