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Bioorthogonal labeling with tetrazine-dyes for super-resolution microscopy

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Genetic code expansion (GCE) technology allows the specific incorporation of functionalized noncanonical amino acids (ncAAs) into proteins. Here, we investigated the Diels-Alder reaction between trans-cyclooct-2-ene (TCO)-modified ncAAs, and 22 known and novel 1,2,4,5-tetrazine-dye conjugates spanning the entire visible wavelength range. A hallmark of this reaction is its fluorogenicity - the tetrazine moiety can elicit substantial quenching of the dye. We discovered that photoinduced electron transfer (PET) from the excited dye to tetrazine is the main quenching mechanism in red-absorbing oxazine and rhodamine derivatives. Upon reaction with dienophiles quenching interactions are reduced resulting in a considerable increase in fluorescence intensity. Efficient and specific labeling of all tetrazine-dyes investigated permits super-resolution microscopy with high signal-to-noise ratio even at the single-molecule level. The different cell permeability of tetrazine-dyes can be used advantageously for specific intra- and extracellular labeling of proteins and highly sensitive fluorescence imaging experiments in fixed and living cells.

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Single-molecule localization microscopy is a powerful method for subdiffraction-resolution fluorescence imaging of cells and tissue^{1,2}. Since the density of fluorophores controls the achievable structural resolution³, efficient and specific labeling with fluorescent probes is a decisive factor in this super-resolution microscopy technique. Despite recent progress in the development of new fluorophores with higher fluorescence quantum yield, photostability, and intrinsic photoswitching in aqueous buffer^{4–8}, specific and efficient labeling of the molecule of interest with minimal linkage error remains challenging. As the field moves toward ever higher spatial resolution using, e.g. expansion microscopy^{9,10}, the effective size of the label (fluorophore, linker and affinity reagent) will be the main limiting factor of super-resolution microscopy.

Immunolabeling with antibodies is still the method of choice for fluorescence imaging of fixed cells, but the large size of antibodies introduces a displacement of 10–15 nm of the fluorophore from the molecule of interest¹. This linkage error is smaller for fluorescent proteins and protein-based self-labeling tags but is still several nanometers¹¹. Small (1.5 × 2.5 nm) camelid antibodies (so called nanobodies) directed against green fluorescent protein (GFP) or shorter peptide epitopes have been used successfully for super-resolution microscopy¹² but the palette of targets is small¹³. Even the most optimized labeling strategies using short peptide tags labeled with bivalent nanobodies or fluorescently labeled monomeric streptavidin yield linkage errors of ~2 nm in direct stochastic optical reconstruction microscopy (dSTORM) experiments^{14–17}. This problem demands the development of efficient labeling methods with small dyes, which can be site-specifically and quantitatively attached to a protein of interest with low linkage error¹⁸.

Genetic code expansion (GCE) technology enables the introduction of noncanonical amino acids (ncAAs) with small functional groups at any position in a target protein^{19–21}. In this strategy, a native codon is replaced with a rare codon, such as the amber (TAG) stop codon, at a specific site in the gene of the protein of interest. The modified protein is then expressed in mammalian cells along with an additional tRNA–tRNA synthetase pair (tRNA-RS) that is orthogonal to the host translational machinery. The active site of the tRNA synthetase enzyme is engineered to only accept a specific ncAA, which is incorporated into a tRNA that recognizes the rare codon. The ncAA is simply added to the growth medium and thereby incorporated into the protein at a specific site^{19–21}. A particularly promising type of ncAA include strained alkenes, such as trans-cyclooct-2-ene (TCO*), that can react with a 1,2,4,5-tetrazine in an ultrafast, specific, and bioorthogonal inverse electron-demand Diels-Alder reaction. In this strategy, TCO*-modified ncAAs, such as TCO*-L-lysine (TCO*-Lys), can be efficiently and directly labeled with organic dyes with minimal linkage error^{22,23}. The high selectivity and rate of this click chemistry reaction has resulted in a large number of commercially available fluorophore-tetrazine conjugates allowing labeling of mammalian cells and whole organisms with organic dyes, even in living systems^{24–26}.

Another interesting property of this labeling strategy is the potential for fluorogenicity. It has been reported that some tetrazine-functionalized dyes (tetrazine-dyes) function as fluorogenic probes, meaning these compounds substantially increase fluorescence intensity upon reaction with the strained dienophile such as TCO^{27,28}. This makes tetrazine-dyes especially interesting for live-cell labeling and fluorescence imaging applications since the fluorogenic reaction could lower background and potentially eliminate the need for washing out excess fluorophore. Despite these apparent advantages however, the use of ncAAs and tetrazine-dyes to label proteins for super-resolution microscopy applications remain rare and unoptimized^{29–32}.

Motivated by these considerations, we studied the spectroscopic characteristics and quenching mechanism of 22 known and novel tetrazine-dyes that span the entire visible spectral range. The tetrazine-dyes included in this study are ATTO425, ATTO465, ATTO488, ATTO532, Cy3, carboxytetramethylrhodamine (TAMRA), ATTO550, ATTO565, ATTO590, ATTO594, ATTO620, Si-rhodamine (SiR)⁴, ATTO647N, Cy5, ATTO655, ATTO680, ATTO700, two large Stokes shift dyes AZ503, and AZ519, as well as the recently introduced Si-rhodamine derivative JF₆₄₆⁶, spontaneously blinking HMSiR⁵, and the bridged carbocyanine dye Cy5B⁸ (Supplementary Fig. 1).

Results

Spectroscopic characteristics of tetrazine-dyes. The absorption and emission maxima of the 22 tetrazine-dyes were measured in aqueous buffer (PBS, pH 7.4) before and after reaction with ncAA TCO*-Lys (Table 1). In TCO*-Lys isomerization to the less reactive *cis*-isomer is prevented and the compound is stable for several days under physiological conditions³³. The majority of tetrazine-dyes are commercially available as 3-methyl-6-phenyl-1,2,4,5-tetrazine (Me-Tet) derivatives and only some of them are also available as 3-phenyl-1,2,4,5-tetrazine (H-Tet) derivatives. Those dyes which are not available as tetrazine derivatives were synthesized by reacting *N*-hydroxysuccinimide derivatives of the fluorophores with 3-(*p*-benzylamino)-1,2,4,5-tetrazine (H-Tet-amine; Supplementary Fig. 1). The main difference between Me-Tet and H-Tet dye conjugates are the different reactivity and chemical stability: H-Tet exhibits an approximately 30-fold higher click reaction rate constant but a lower chemical stability³³.

Almost all tetrazine-dyes investigated exhibited a strong fluorescence increase upon reaction with TCO*-Lys (Table 1, Fig. 1). The increase in fluorescence intensity is most pronounced for the shorter wavelength absorbing dyes ATTO425, ATTO465, and ATTO488 with turn-on ratios of 15–40. Due to the broad absorption spectrum of tetrazine peaking at ~515 nm it can efficiently quench the fluorescence of dyes emitting at wavelengths ≤ 550 nm by fluorescence resonance energy transfer (FRET) (Supplementary Fig. 1)²⁷. Hence, the tetrazine chromophore can act as both quencher and bioorthogonal click-reaction group. Indeed, the turn-on ratios of the tetrazine-dyes ATTO425, ATTO465, ATTO488, and ATTO532 after coupling to TCO*-lysine scale according to the emission maxima of the dyes and their overlap with the absorption spectrum of tetrazine (Table 1, Supplementary Figs. 1 and 2).

The longer wavelength absorbing oxazine dyes ATTO655, ATTO680, and ATTO700, and the large Stoke-shift dyes AZ503 and AZ519 show turn-on ratios of 6–13 (Table 1, Fig. 1). These high turn-on ratios are not entirely reflected in the increase in fluorescence lifetime indicating that short fluorescence lifetime components are missed in our time-correlated single-photon counting (TCSPC)-experiments with a time resolution of approximately 40 ps.

Long-term spectroscopy studies revealed that some dyes such as ATTO465, ATTO488, and ATTO655 show a slight but highly reproducible decrease in fluorescence intensity after the initial strong increase upon addition of TCO*-Lys (Fig. 1), which can be attributed most likely to tautomerization of the TCO*-tetrazine conjugate resulting in the release of lysine by decarboxylative elimination³⁴. Elimination of the dye moiety from the lysine results in the formation of a new heterocyclic compound (pyridazine), which potentially quenches the fluorescence of some dyes depending on their electronic properties.

Comparative click reactions demonstrate that the click reaction of H-Tet-ATTO488 and H-Tet-Cy5 with TCO*-Lys proceeds

Table 1 Spectroscopic characteristics (absorption maxima, λ_{abs} (nm), emission maxima, λ_{em} (nm), fluorescence lifetimes, τ (ns)) of flexibly linked tetrazine-dyes after click reaction with TCO*-lysine and turn-on ratios measured in PBS, pH 7.4. Click reactions were performed by adding 25 μM TCO*-Lys to 1 μM dye solutions

Tetrazine-dye	λ_{abs} (nm)	λ_{em} (nm)	τ_{av} (ns) ^a	Turn-on ratio ^b
Me-Tet-ATTO425	441	478	1.28 (3.93)	15
Me-Tet-ATTO465	455	505	0.21 (2.82)	39
Me-Tet-ATTO488	502	522	1.39 (3.60)	25
Me-Tet-ATTO532	535	553	3.13 (3.00)	7
H-Tet-Cy3	551	564	0.37 (0.44)	2
Me-Tet-5-TAMRA	553	580	1.96 (2.38)	1
Me-Tet-AZ519	478	602	3.29 (4.08)	8
Me-Tet-AZ503	498	629	2.38 (2.79)	8
Me-Tet-ATTO550	555	573	1.95 (3.40)	4
Me-Tet-ATTO565	566	587	2.24 (3.80)	2
Me-Tet-ATTO590	598	621	1.99 (3.70)	3
Me-Tet-ATTO594	604	664	1.85 (3.74)	4
Me-Tet-ATTO620	620	637	1.47 (2.86)	3
Me-Tet-JF646	646	667	2.45 (3.01)	10
Me-Tet-ATTO647N	647	660	3.01 (3.58)	2
H-Tet-Cy5	649	664	0.76 (1.21)	3
H-Tet-SiR	649	669	2.40 (2.71)	4
H-Tet-HMSiR	654	667	2.56 (2.49)	1
Me-Tet-ATTO655	668	676	1.50 (2.12)	13
H-Tet-Cy5B	670	683	1.08 (1.91)	3
Me-Tet-ATTO680	686	694	1.38 (1.91)	6
Me-Tet-ATTO700	707	712	1.28 (1.81)	8

^aThe quality of the fit was judged by the reduced χ^2 values and the randomness of the weighted residuals. For all fits, χ^2 values between 0.900 and 1.200 were obtained. Most clicked fluorescent dyes showed nonexponential fluorescence kinetics, which could be best described using a biexponential model. Therefore, average fluorescence lifetimes $\tau_{\text{av}} = a_1\tau_1 + a_2\tau_2$ are given. The lifetime values in brackets were measured after clicking with TCO*-lysine

^bThe turn-on ratio was determined from the minimum and maximum fluorescence intensity value recorded before and after addition of 25 μM TCO*-Lys to the tetrazine-dyes (Fig. 1)

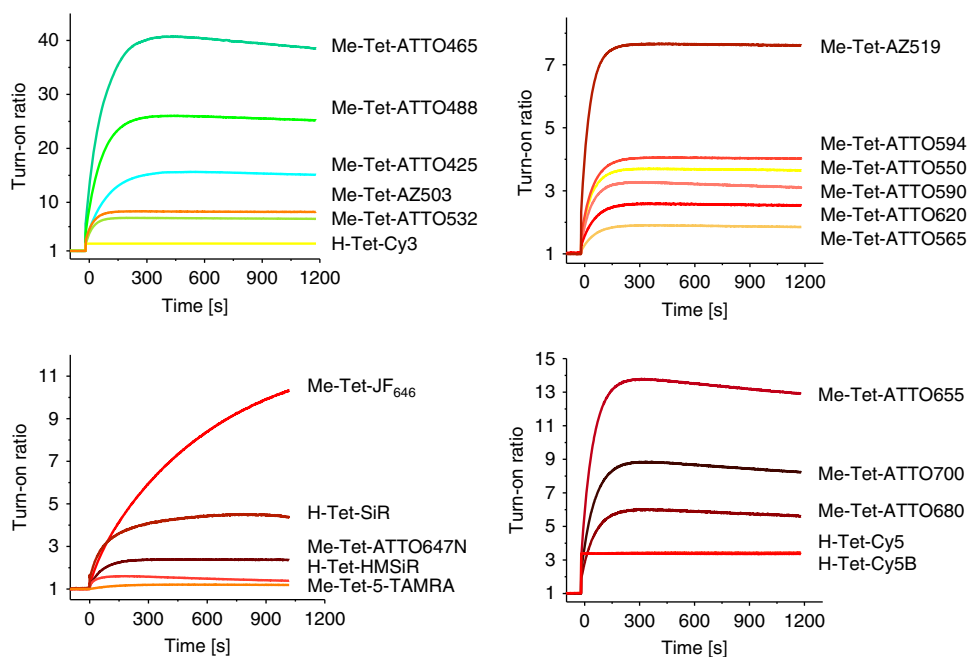


Fig. 1 Fluorescence turn-on. Fluorescence intensity increase of 1 μM solutions of different tetrazine-dyes upon addition of 25 μM TCO*-Lys. All measurements were performed in PBS, pH 7.4 at 25 $^{\circ}\text{C}$

~30-times faster than for the corresponding Me-Tet derivatives³⁵ but the turn-on ratios differ strongly especially for the two ATTO488 tetrazine derivatives (Supplementary Fig. 3). In particular, H-Tet-ATTO488 is substantially less quenched than Me-Tet-ATTO488. Accordingly, the turn-on ratio is much higher for the Me-Tet derivative. Assuming similar FRET efficiencies for

both ATTO488 conjugates the different turn-on ratios can only be explained by the reduced chemical stability of H-Tet³⁵. Obviously, a fraction of H-Tet-ATTO488 decomposes spontaneously in physiological buffer because of the higher reactivity, which results correspondingly in a lower FRET-efficiency and turn-on ratio (Supplementary Fig. 3).

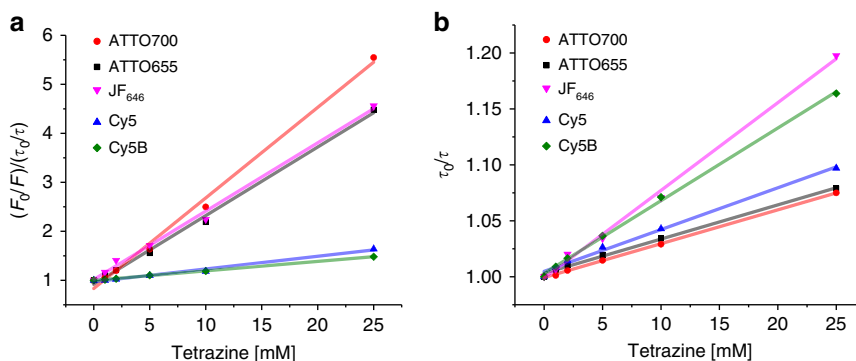


Fig. 2 Fluorescence quenching of flexibly linked tetrazine dyes. **a** Static and **b** dynamic quenching of ATTO655, ATTO700, Cy5, Cy5B, and JF₆₄₆ with 0–25 mM Me-Tet-amine. The straight lines represent a linear least squares fit to the data

Higher quenching efficiencies with turn-on rates of >1,000 have been achieved by using through-bond energy transfer (TBET) in tetrazine-BODIPY derivatives²⁸. However, the TBET quenching is decreasing with increasing absorption wavelength of the dyes³⁶. Therefore, bodipy and coumarin dyes have been used mostly in studies, which are less suited for super-resolution microscopy^{27,28,37}. Flexibly-linked tetrazine-dyes as used here are easier to access from a synthetic standpoint but exhibit lower turn-on ratios after reaction with dienophiles (Table 1)²⁹.

Fluorescence quenching in red-absorbing tetrazine-dyes. To shed light on the fluorescence quenching mechanism in the longer wavelength absorbing flexibly linked tetrazine-dyes we performed intermolecular steady-state and time-resolved fluorescence quenching experiments with free dyes and Me-Tet-amine under physiological conditions (PBS, pH 7.4) following previous studies on tryptophan quenching^{38,39} (Fig. 2, Supplementary Fig. 4). The time-resolved fluorescence decays remain almost monoexponential with a longer fluorescence lifetime, dependent on the quencher concentration, and an additional shorter decay component with lifetimes in the range of a few tens of picoseconds. For the quencher dependent longer fluorescence lifetimes, we observed linear Stern-Volmer plots for all measured dyes (ATTO655, ATTO700, JF₆₄₆, Cy5, and Cy5B) (Fig. 2) and calculated bimolecular dynamic quenching rate constants k_{dyn} of $(1.7 \pm 0.1) \times 10^9 \text{ M}^{-1} \text{ s}^{-1}$, $(1.9 \pm 0.1) \times 10^9 \text{ M}^{-1} \text{ s}^{-1}$, $(2.6 \pm 0.1) \times 10^9 \text{ M}^{-1} \text{ s}^{-1}$, $(4.1 \pm 0.2) \times 10^9 \text{ M}^{-1} \text{ s}^{-1}$, and $(3.5 \pm 0.1) \times 10^9 \text{ M}^{-1} \text{ s}^{-1}$, respectively. These values represent dynamic quenching that is approaching the diffusion limit³⁸ indicating that most collisional encounters result in fluorescence quenching.

In addition, strong static quenching was observed for the two oxazine dyes ATTO655 and ATTO700 as well as for the Si-rhodamine dye JF₆₄₆ (Fig. 2) suggesting that non- or only weakly fluorescent ground-state complexes between the longer wavelength absorbing oxazine and Si-rhodamine dyes and Me-Tet-amine form in aqueous solutions, where the lifetime of the complexes is below the time-resolution of our TCSPC setup.

From the static quenching component the association constant, K_a , for complex formation can be calculated by plotting $(F_0/F)/(\tau_0/\tau)$ versus the quencher (Me-Tet-amine) concentration (Fig. 2, Supplementary Fig. 4). We estimated K_a for complex formation between the dyes ATTO655, ATTO700, JF₆₄₆ and Me-Tet-amine to be $(135 \pm 4) \text{ M}^{-1}$, $(175 \pm 7) \text{ M}^{-1}$, and $(140 \pm 3) \text{ M}^{-1}$, respectively. Cy5 and Cy5B, on the other hand, show only weak static quenching (Supplementary Fig. 4) with K_a of $(24 \pm 1) \text{ M}^{-1}$, and $(19 \pm 1) \text{ M}^{-1}$, respectively.

Me-Tet-ATTO655, Me-Tet-ATTO680, Me-Tet-ATTO700, and Me-Tet-JF₆₄₆ show a high turn-on ratio (Table 1) and are therefore efficiently quenched by Me-Tet-amine also in

intramolecular quenching experiments. This finding indicates that the dye and the tetrazine moiety, when connected by a flexible linker, can adopt a nearly coplanar stacking conformation, which is required for efficient fluorescence quenching.

Our results indicate that the underlying quenching mechanism in longer wavelength absorbing oxazine and Si-rhodamine dyes is only efficient at very short distances, i.e. within complexes. This is also supported by the observation that the fluorescence intensity of ATTO655, ATTO680, and ATTO700 tetrazine conjugates increases strongly upon addition of denaturing agents such as guanidinium chloride while ATTO488 and Cy5 show only small effects if at all (Fig. 3). Hence, it can be concluded that hydrophobic interactions between the longer wavelength absorbing oxazine and Si-rhodamine dyes and Me-Tet with a stacked arrangement of the conjugated π -electron systems play an important role in the formation of ground-state complexes with ultrafast fluorescence quenching. Since tetrazine exhibits a very high electron affinity^{40,41}, the excited longer-wavelength absorbing dyes are most likely efficiently quenched in their stacked conformation via photoinduced electron transfer (PET). The carbocyanine dyes such as Cy5 exhibit a higher water solubility and are thus less prone to form complexes with Me-Tet. In the case of Me-Tet-ATTO488 fluorescence quenching is dominated by through-space FRET where complex formation between dye and tetrazine moiety does not play a dominant role.

Fluorescence quenching dynamics. Knowing that ATTO655 and similar dyes form nonfluorescent complexes with tetrazine in aqueous solutions driven by hydrophobic interactions, we can assume an equilibrium between two states for the dye in conformationally flexible tetrazine-dyes: an ‘open’ fluorescent state A_{open} and a ‘closed’, complexed nonfluorescent state B_{closed} . Both states are populated according to the closing and opening rate constants, k_{closing} and k_{opening} . To investigate this equilibrium in more detail, we performed fluorescence correlation spectroscopy (FCS) experiments (Fig. 4). FCS analyzes temporal fluorescence fluctuations from highly diluted samples (nanomolar concentrations), probing molecules diffusing through a confined detection volume (typically 1 femtoliter with 1–20 molecules) by Brownian motion⁴². Characteristic time scales of molecular processes that result in fluctuating fluorescence emission can be measured under thermodynamic equilibrium conditions with nanosecond time resolution^{43–45}.

Under moderate excitation conditions, no photophysical process, such as intersystem crossing, is apparent in the submillisecond time domain of FCS-curves recorded from Me-Tet-ATTO655 after reaction with TCO*-Lys (Fig. 4a). The FCS curve shows a millisecond decay corresponding to the molecules’ diffusion through the confocal excitation/detection volume.

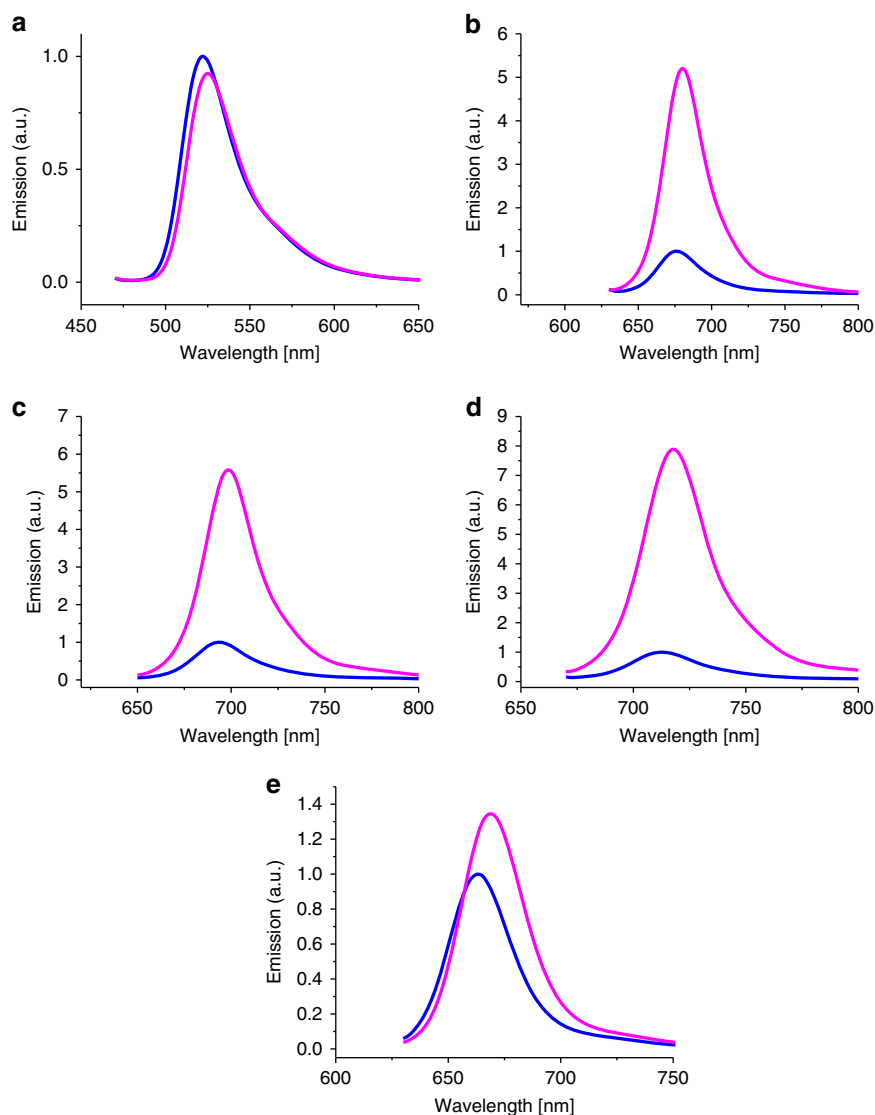


Fig. 3 Relative fluorescence emission spectra upon addition of guanidinium chloride of **a** Met-Tet-ATTO488, **b** Met-Tet-ATTO655, **c** Met-Tet-ATTO680, **d** Met-Tet-ATTO700, and **e** H-Tet-Cy5 in PBS (pH 7.4) without (blue) and with 6 M guanidinium chloride (magenta)

However, the FCS curve recorded from Me-Tet-ATTO655 before addition of TCO*-Lys displays an additional fast decay occurring on the nano- to microsecond time scale together with a change in amplitude (Fig. 4a). Bimolecular experiments with ATTO655 and 25 mM tetrazine reveal a similar FCS decay (Fig. 4b) confirming the idea of diffusion-driven formation of non-fluorescent complexes. By fitting an analytical FCS curve representing two-state fluorescence intermittency to the bimolecular data with a mixture of free ATTO655 and tetrazine we estimated $k_{\text{closing}} = (0.04 \pm 0.01) \times 10^9 \text{ s}^{-1}$ and $k_{\text{opening}} = (0.015 \pm 0.002) \times 10^9 \text{ s}^{-1}$. These values represent an association rate constant of $(1.5 \pm 0.5) \times 10^9 \text{ M}^{-1} \text{ s}^{-1}$ and an association constant of $(100 \pm 23) \text{ M}^{-1}$, close to the values from Stern-Volmer analysis (a sphere-of-action contribution in the Stern-Volmer plots of ATTO655 renders the Stern-Volmer association constant slightly larger)³⁸. Concentration estimates from the amplitude of FCS curves confirm that all fluorophores contribute to the observed bimolecular FCS decays and no population of fluorophores exist, that are quenched over time periods longer than ~ 1 ms (the diffusion-limited observation time) (Fig. 4c).

It is thus reasonable that the nano- to microsecond decay recorded for Me-Tet-ATTO655 reflects fluorescence fluctuations

caused by intramolecular quenching of ATTO655 upon contact with Me-Tet mediated by the conformational flexibility of the linker. The time scale of the fast quenching fluctuations is well separated from that of translational diffusion. The calculated rate constants for opening and closing, k_{opening} of $\sim 9.0 \times 10^6 \text{ s}^{-1}$ and k_{closing} of $\sim 3.3 \times 10^6 \text{ s}^{-1}$, demonstrate that ATTO655 stays on average 27% of the time in its quenched complexed conformation. The fast decay thus accounts for about one fourth of the observed turn-on ratio of 13 (Table 1). The remaining turn-on ratio is indeed reflected in a concentration difference by about a factor of four that is observed from the amplitude of FCS curves (Fig. 4c). This observation of multiple decay components on time scales above and below the FCS diffusion time of ~ 1 ms, which are not detected in intermolecular quenching experiments (Fig. 4b) indicates that the intramolecular linker introduces additional conformational constraints for the process of complex formation.

Whereas Me-Tet-ATTO680 and Me-Tet-ATTO700 show similar behavior, the FCS-curves recorded from Me-Tet-Cy5 and Me-Tet-Cy5B in the absence and presence of TCO*-Lys are identical demonstrating that quenched complexes are not formed. The observed fast decay for Cy5 is due to well-characterized cis/

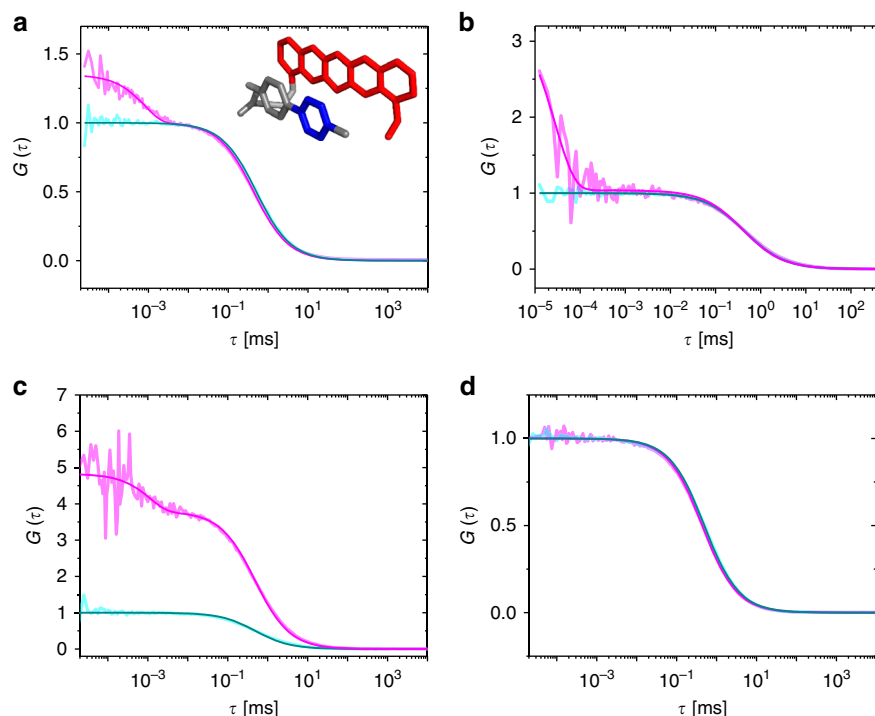


Fig. 4 Fluorescence correlation spectroscopy (FCS) measurements **a** FCS-curve of Me-Tet-ATTO655 recorded in PBS, pH 7.4 in the absence (magenta) and presence (cyan) of 10 μ M TCO*-Lys. Inset: Model for formation of weak or nonfluorescent ground-state complexes between an oxazine dye and tetrazine. Due to the flexible linker the oxazine dye (e.g. ATTO655) (red) can adopt a nearly coplanar stacking conformation with respect to tetrazine (blue). **b** FCS curves of free ATTO655 alone (cyan) and in the presence of 25 mM Me-Tet-amine (magenta). **c** FCS curves of Me-Tet-ATTO655 (magenta) and free ATTO655 (cyan). All data were recorded with dye concentrations of 1 nM and the FCS curves and normalized to the amplitude of the free ATTO655 data. Amplitude differences thus reflect the existence of quenched complexes that are stable on time scales longer than the diffusion time scale. **d** FCS-curve of H-Tet-Cy5B recorded in PBS, pH 7.4 in the absence (magenta) and presence (cyan) of 10 μ M TCO*-Lys. All measurements were performed in PBS, pH 7.4 at 25 $^{\circ}$ C

trans-isomerization (Supplementary Fig. 5)⁴⁶. Accordingly, we observed that Cy5B, which is not capable of isomerization due to the constrained conjugated bond structure, does not show the isomerization decay (Fig. 4d).

Super-resolution microscopy with tetrazine-dyes. In order to enable testing of different tetrazine-dyes for super-resolution microscopy applications in an experimentally easy and comparable way, we followed a recently introduced method³⁷ and synthesized phalloidin-TCO, for labeling of the actin skeleton of fixed cells and subsequent attachment of tetrazine-dyes by click chemistry. Confocal fluorescence images demonstrate that all tetrazine-dyes are well-suited for high-end microscopy (Fig. 5, Supplementary Fig. 6). However, comparative fluorescence labeling and imaging experiments with unmodified phalloidin show that some tetrazine-dyes exhibit the tendency to bind nonspecifically to intracellular structures. Especially some of the yellow to red absorbing ATTO-dyes (e.g. Me-Tet-ATTO550, Me-Tet-ATTO565, and Me-Tet-ATTO590) show substantial nonspecific background signal (Supplementary Fig. 6).

Next, we tested their performance in super-resolution microscopy by re-scan confocal microscopy (RCM)^{47,48} and structured illumination microscopy (SIM)⁴⁹ providing a theoretical resolution improvement factor of 1.4 and 2.0, respectively (Fig. 6, Supplementary Figs. 7 and 8). Most tetrazine-dyes investigated, including the new dyes HMSiR⁵, JF₆₄₆^{6,7}, and Cy5B⁸, exhibit a low tendency to bind nonspecifically to cellular structures and provide super-resolved actin images with high signal-to-noise ratio. Furthermore, fluorogenic tetrazine-dyes such as Me-Tet-ATTO488, Me-Tet-ATTO655, and Me-Tet-ATTO680 with turn-

on ratios ≥ 6 (Table 1) enable wash-free super-resolution fluorescence imaging (Fig. 6b)^{30,31}. In addition, also other tetrazine-dyes such as H-Tet-Cy3 and H-Tet-Cy5 with lower turn-on ratio allow wash-free fluorescence imaging (Supplementary Fig. 9). This finding demonstrates that not only the turn-on ratio but likewise a high water-solubility and low tendency of unspecific binding to cellular components in combination with superior click reactivity can promote wash-free imaging of intracellular structures.

Next, we performed single-molecule localization microscopy experiments to verify that tetrazine-dyes can be used as well for single-molecule sensitive fluorescence imaging (Fig. 7). The obtained super-resolution images demonstrate, that H-Tet-Cy5 coupled to phalloidin-TCO can be reliably photoswitched in the presence of millimolar concentrations of thiols enabling *d*STORM imaging^{14,15} of the actin skeleton with superior spatial resolution (Fig. 7a). In addition, we tested also the bridged carbocyanine dye Cy5B⁸ for PALM-like imaging under reductive imaging conditions (Fig. 7b) and the spontaneously blinking dye HMSiR⁵ in PBS, pH 7.4 without addition of photoswitching buffer (Fig. 7c) and achieved similar image qualities; or to be precise, the dyes exhibit slightly different localization precisions of ~ 12 nm (Cy5), ~ 13 nm (HMSiR), and ~ 11 nm (Cy5B) due to the different localization intensities of 1 (Cy5): 0.42 (HMSiR): 1.8 (Cy5B) recorded under the different experimental conditions.

Visualization of membrane receptors with tetrazine-dyes. To demonstrate the usefulness of click chemistry and tetrazine-dyes for fluorescence imaging of cells, we used GCE for site-specific labeling of extracellular domains of two membrane receptors: the

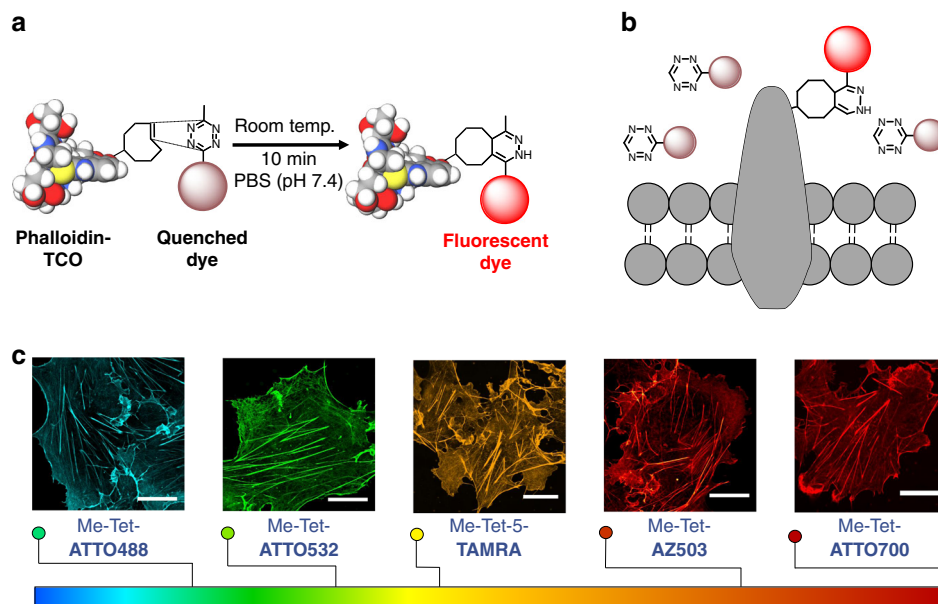


Fig. 5 High-end microscopy of actin with tetrazine-dyes. **a** Cos-7, U2-OS and NIH-3T3 cells were fixed and labeled with phalloidin-TCO followed by click-labeling with different tetrazine-dyes. All tetrazine-dyes investigated show specific labeling of the actin skeleton. **b** Tetrazine-dyes can also be used for site-specific labeling of the extracellular domain of membrane receptors (see also Fig. 8). **c** Various tetrazine-dyes spanning the entire visible part of the electromagnetic spectrum can be used successfully for confocal laser scanning microscopy of intracellular actin structures (see also Fig. 6). Scale bars, 20 μm

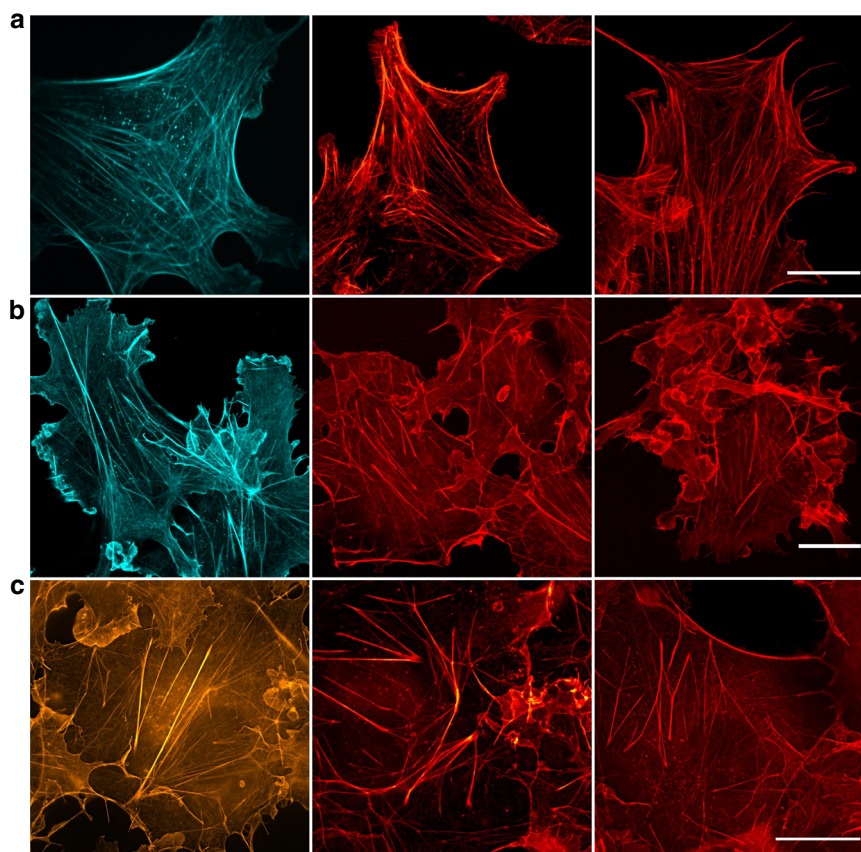


Fig. 6 Super-resolution microscopy with tetrazine-dyes. **a** RCM-images of actin in fixed NIH-3T3 cells labeled with phalloidin-TCO and Me-Tet-ATTO425, large Stokes-shift dyes Me-Tet-AZ503, and H-Tet-SiR are shown from left to right. **b** Wash-free confocal laser scanning microscopy images of fixed Cos-7 cells labeled with Me-Tet-ATTO488, Me-Tet-ATTO655, and Me-Tet-ATTO680 (from left to right). Cells were fixed, 1 h incubated with phalloidin-TCO, rinsed with PBS and then labeled with 3 μM tetrazine-dye and imaged directly after 10 min. without any washing step in the presence of unreacted free tetrazine-dye. **c** SIM-images of actin in fixed Cos-7 cells labeled with phalloidin-TCO and Me-Tet-ATTO565, Me-Tet-ATTO620, and Me-Tet-JF₆₄₆ are shown from left to right. Scale bars, 20 μm

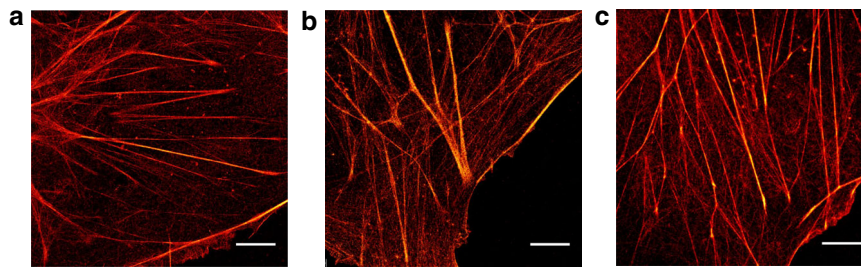


Fig. 7 Single-molecule localization microscopy with tetrazine-dyes. Cos-7 cells were labeled with phalloidin-TCO and different tetrazine-dyes. **a** dSTORM-image of actin labeled with phalloidin-TCO and H-Tet-Cy5 in the presence of photoswitching buffer. **b** PALM-like super-resolution image⁸ of actin labeled with phalloidin-TCO and H-Tet-Cy5B in PBS, pH 7.4. Cy5B was reduced with 0.1% NaBH₄ for 10 min before imaging. **c** Actin was labeled with phalloidin-TCO and H-Tet-HMSiR, a spontaneously blinking silicon rhodamine dye⁵, and imaged in PBS, pH 7.4. Irradiation was performed in all three examples using solely 640 nm laser light at irradiation intensities of ~2 kW/cm². Scale bars, 5 μ m

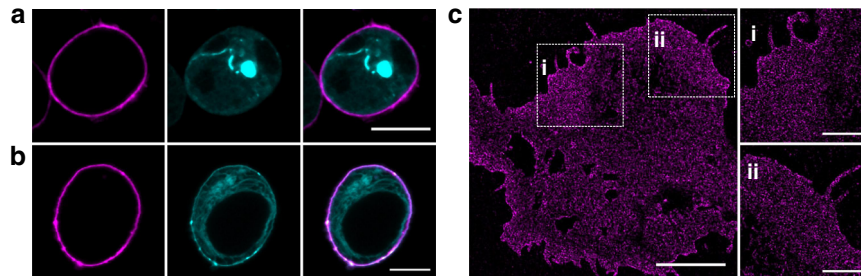


Fig. 8 Visualization of membrane receptors by GCE and bioorthogonal labeling with tetrazine-dyes. **a** Confocal fluorescence image of the clickable TNF receptor 1 (TNFR1^{S42TAG}-tdEOS) and **b** Kainate receptor (GluK2^{S343TAG}-eGFP) in HEK293T cells. Site-specific extracellular labeling was performed with the cell impermeable H-Tet-Cy5 (1.5 μ M) (magenta). The tdEOS and eGFP channels, respectively, are shown in cyan. **c** dSTORM image of membrane receptors GluK2^{S343TAG}-eGFP expressed in HEK293T cells shown a homogeneous distribution of receptors in the plasma membrane. Scale bars, 10 μ m (**a-b**), 5 μ m (**c**) and 2 μ m (expanded images i and ii)

kainate receptor (GluK2) and the tumor necrosis factor receptor 1 (TNFR1). Kainate receptors are ionotropic glutamatergic receptors that mediate fast excitatory neurotransmission. They form homo- and heterotetramers and are localized to the presynaptic and postsynaptic sides of excitatory synapses⁵⁰. Unfortunately, there are no commercial antibodies available that recognize the extracellular domain (N-terminus) of GluK2-tetramers preventing hitherto live-cell imaging of kainate receptors in the plasma membrane with organic dyes.

The TNFR superfamily is a very important class of signal transduction molecules in the immune system⁵¹. TNFRs are single-membrane-spanning proteins that contain an extracellular TNF-binding region and a cytoplasmic tail. There are several antibodies commercially available that recognize the extracellular domain of TNFRs. However, antibody binding often interferes with biological function of TNFRs⁵².

To enable fluorescence imaging of TNFR1 and GluK1 in living cells with organic dyes, we selected five different extracellular positions for ncAAs incorporation into GluK2 and TNFR1. The positions were selected to ensure easy accessibility for bioorthogonal click labeling with tetrazine-dyes based on available structural data (GluK2: S47TAG, S272TAG, K306TAG, S309TAG, S343TAG; TNFR1: S42TAG, Q77TAG, R121TAG, S137TAG, K161TAG). For identification of the best mutant we used the following selection criteria. First, with tdEOS and eGFP, respectively, attached C-terminally after the amber suppression site, the fluorescence signal is only detectable when the ncAA is incorporated properly into the protein of interest. Hence, only those mutants are selected, which exhibit a strong fluorescence intensity (expression level of the fluorescent protein). Second, the fluorescent proteins should be localized at the expected cellular

position (here, the plasma membrane) to ensure selective extracellular labeling with tetrazine-dyes. Third, bioorthogonal labeling with plasma membrane impermeable tetrazine-dyes (H-Tet-Cy5) should show selective membrane labeling. Based on these experiments, we identified GluK2^{S343TAG}-eGFP and TNFR1^{S42TAG}-tdEOS as the best candidates for further experiments.

To improve the incorporation efficiency of ncAAs and reduce the background signal we used a recently optimized pyrrolysine-based CGE system for click chemistry⁵³. By adding a strong nuclear export signal (NES) to the N-terminus of the pyrrolysine tRNA synthase (PylRS^{AF}) sequence (NESPylRS^{AF}) (AF refers to a previously described PylRS mutant that accepts bulky side-chain moieties), unspecific nuclear labeling is reduced⁵³. In combination with cell-impermeable dyes such as H-Tet-Cy5 this approach enables selective and efficient labeling of extracellular domains of correctly incorporated membrane receptors in living cells

First, we exchanged serine at position 42 by TCO*-Lys in the extracellular domain of TNFR1 (TNFR1-tdEOS) in HEK293T cells by amber suppression. We tested different H-Tet-Cy5 concentrations ranging from 10 nM to 5 μ M and found that labeling with 1.5 μ M H-Tet-Cy5 for 10 min provides the best labeling efficiency⁵⁴. Cells were labeled with H-Tet-Cy5 and imaged by confocal laser scanning microscopy. Live-cell fluorescence images clearly show selective labeling of TNFRs in the plasma membrane of cells (Fig. 8a).

Next, we labeled the kainate receptor GluK2 (GluK2^{S343TAG}-eGFP) in HEK293T cells by GCE (amber suppression) with TCO*-Lys and H-Tet-Cy5. Live-cell confocal fluorescence images show again exclusively extracellular fluorescence labeling of GluK2 (Fig. 8b). Corresponding dSTORM images demonstrate

that GluK2 is expressed at high expression rates but homogeneously distributed in the plasma membrane of fixed HEK293T cells (Fig. 8c).

To verify the labeling efficiency, we performed immunofluorescence labeling of kainate receptor GluK2 and TNFR in HEK293T cells. Since commercially available anti-GluK2 antibodies are directed against the intracellular domain cells have to be fixed and permeabilized before labeling. Confocal fluorescence images demonstrate that different antibodies can be used successfully to visualize kainate receptors in fixed cells (Supplementary Fig. 10). Staining of plasma membrane TNFRs with two commercial antibodies showed specific membrane labeling but resulted in negligible or substantially lower labeling efficiency as compared to bioorthogonal click labeling (Supplementary Fig. 11). These results demonstrate that bioorthogonal click chemistry with tetrazine-dyes can be used advantageously for labeling of membrane receptors. Site-specific incorporation of ncAAs into the extracellular domain of plasma membrane receptors enables live-cell visualization of plasma membrane receptors that cannot be labeled by antibodies. In addition, it enables higher labeling densities than standard immunolabeling with antibodies. This suggests that the extracellular labeling sites of plasma membrane receptors is more accessible to small tetrazine-dyes such as H-Tet-Cy5 than to larger IgG antibodies with a molecular size well above 10 nm. Furthermore, antibodies may crosslink to a certain degree during the fixation process and thus induce apparent clustering of receptors in the plasma membrane⁵⁴.

Live-cell bioorthogonal labeling. So far intracellular live-cell bioorthogonal labeling by GCE and fluorescence imaging has been hampered by the high fluorescence background resulting from the excess of unincorporated ncAAs³³. Even water-soluble tetrazine-dyes such as H-Tet-SiR⁴ or tetrazine-dyes with high turn-on ratio do not automatically permit high-end live-cell fluorescence imaging of intracellular target proteins. In general, incorporation of TCO*-Lys into proteins in response to an in-frame amber stop codon does not alter cell viability^{20,25}. Depending on the incorporation site the expression level and cellular location might, however, be influenced. This off-target binding further increases the fluorescence background.

To demonstrate live-cell bioorthogonal labeling and imaging with tetrazine-dyes we selected tubulin as an important cytoskeleton component of cells in our experiments. Microtubules are dynamic structures composed of $\alpha\beta$ -tubulin molecules that are constantly integrated or degraded as the microtubules grow and shorten. Microtubule dynamics can be easily monitored in live cells using fluorescently labeled tubulin and video microscopy. Since overexpression of α -tubulin can potentially influence the dynamics of endogenous microtubules after incorporation and generate a high fluorescent background³⁰, we used two alternative live-cell tubulin labeling strategies. First, we introduced TCO*-Lys into the microtubule-associated protein (MAP) ensconsin (E-MAP-115), which is known to be better tolerated by cells at high expression levels⁵⁵ and second, we synthesized Docetaxel-TCO (see Supplementary Methods), a microtubule-stabilizing antimitotic drug^{56–58} (Fig. 9). It has been shown that cells expressing four to ten times the physiological level of endogenous MAP exhibited microtubule dynamics indistinguishable from those of untransfected cells indicating that E-MAP-115 most likely serves to modulate microtubule functions or interactions with other cytoskeletal elements⁵⁵. We inserted TCO*-Lys into the microtubule binding domain (EMTB) of E-MAP-115, which is C-terminally tagged with three GFPs (EMTB^{K87TAG}-3xGFP)⁵⁹ for live-cell bioorthogonal labeling and fluorescence imaging (Fig. 9). Thus, monitoring of the GFP signal

allowed us to select strongly expressing cells and change to TCO*-Lys free medium one day before bioorthogonal labeling with cell-permeable H-Tet-SiR^{4,30} in order to minimize the non-specific background signal. In addition, we used again the tRNAPyl/NESpYlRS^{AF} pair to improve nuclear export of tRNA-Synthetase⁵³.

Using this optimized approach, we were able to perform fluorescence imaging of microtubule dynamics in living cells by re-scan confocal microscopy (RCM)^{47,48} (Fig. 9d, Supplementary Movies 1 and 2). However, additional single-molecule localization microscopy experiments of fixed EMTB^{K87TAG}-3xGFP cells with H-Tet-HMSiR illustrate that the fluorescence background of unbound EMTB still limits the achievable image quality (Fig. 9c). Next, we tested whether Docetaxel-TCO specifically binds to microtubules and enables live-cell labeling of the cytoskeleton with H-Tet-SiR. Live-cell standard confocal as well as super-resolution microscopy images recorded by RCM and SIM⁴⁹ demonstrate specific labeling of the cytoskeleton by click chemistry (Fig. 9d). Microtubules are polar filaments exhibiting a dynamic instability, i.e. extensive phases of microtubule growth are followed by rapid disassembly and regrowth⁶⁰. To investigate microtubule dynamics, we performed live-cell re-scan confocal time lapse microscopy showing that EMTB^{K87TAG}-3xGFP labeled microtubules exhibit distinct polymerization and degradation events (Supplementary Movies 1 and 2, Supplementary Fig. 12a). Microtubules labeled with 10 μ M Docetaxel-TCO show damped dynamics lacking microtubule polymerization and degradation events (Supplementary Movie 3 and Supplementary Fig. 12b). This demonstrates that the microtubule-stabilizing antimitotic drug Docetaxel-TCO is not suited for studying microtubule dynamics in live cells if used at concentrations > 100 nM⁵⁷.

Discussion

With continuously increasing spatial resolution of super-resolution microscopy the development of small and efficient fluorescent probes generating minimal linkage errors becomes particularly important. Site-specific introduction of a ncAA such as TCO*-Lys into the amino acid chain of the protein of interest followed by bioorthogonal click chemistry with tetrazine-dyes represents a broadly useful possibility to overcome current limitations and enable high-end fluorescence imaging with organic dyes. In addition, a continuously increasing number of organic dyes is commercially available as tetrazine-dyes. Whereas FRET is responsible for fluorescence quenching in blue- and green-absorbing dyes, we discovered PET from the excited dye to tetrazine as the main quenching mechanism in red-absorbing oxazine and Si-rhodamine derivatives. We show that a stacked arrangement of the conjugated π -electron system of the dye and tetrazine is required for strong fluorescence quenching. Upon reaction with dienophiles quenching interactions are reduced resulting in a considerable increase in fluorescence intensity.

The resulting fluorescence increase can be used advantageously for wash-free fluorescence imaging experiments. The two large Stokes-shift dyes Me-Tet-AZ519 and Me-Tet-AZ503 and the red-absorbing dyes H-Tet-Cy5, H-Tet-Cy5B, and H-Tet-HMSiR are particularly interesting for autofluorescence-free super-resolution fluorescence imaging by re-scan confocal microscopy or structured illumination microscopy, and single-molecule localization microscopy, respectively. On the other hand, some tetrazine-dyes tend to bind nonspecifically to intracellular structure highlighting the importance of tetrazine-dye selection and suited control experiments for each specific application. Our data also demonstrate that ncAAs can be incorporated site-specifically into even small extracellular domains of plasma membrane proteins for

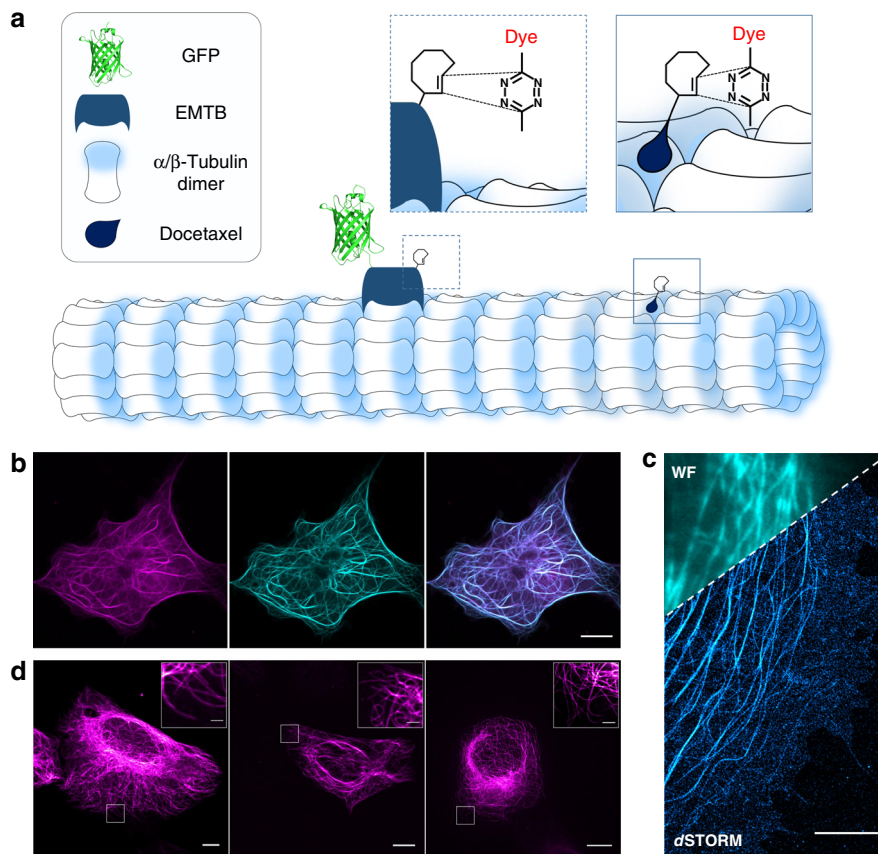


Fig. 9 Intracellular live-cell labeling and imaging of clickable microtubule-associated protein (EMTB) in COS-7 cells. **a** Scheme of incorporation of unnatural amino acid TCO^{*}-Lys into the microtubule-associated protein EMTB^{K87TAG}-3xGFP via GCE and bioorthogonal labeling with a tetrazine-dye. Alternatively, microtubules can be labeled in living cells with Docetaxel-TCO followed by click labeling with a cell permeable tetrazine-dye. **b** Live-cell confocal fluorescence images of the construct EMTB^{K87TAG}-3xGFP labeled with 3 μM of the membrane-permeable H-Tet-SiR for 10 min (GFP: cyan, SiR: magenta, and overlay). **c** Single-molecule localization microscopy image of the same construct. Cells were fixed and labeled with 3 μM H-Tet-HMSiR and then imaged in PBS (pH 7.4). The upper left corner shows an overlay with the corresponding widefield image (WF). **d** Live cell confocal, re-scan confocal, and SIM fluorescence images (from left to right) of U2OS cells treated with 10 μM Docetaxel-TCO for 30 min and labeled with 10 μM H-Tet SiR for 10 min. The insertions in the upper right corner show expanded views of the marked regions demonstrating the improved spatial resolution. Scale bars, 5 μm (**b-d**), 1 μm (expanded views)

which antibodies are not available. Due to the smaller size of tetrazine-dyes they react also with protein sites that are difficult to access by much larger antibodies and thus enable higher labeling efficiencies. Because tetrazine-dyes provide linkage error free stoichiometric labeling of proteins at well-defined extracellular and intracellular positions they are ideally suited for single-molecule fluorescence imaging and tracking as well as super-resolution microscopy experiments in fixed and living cells.

Methods

Tetrazine-dyes/TCO^{*}-Lys. Me-Tet derivatives of ATTO425, ATTO465, ATTO488, ATTO550, ATTO565, ATTO590, ATTO594, ATTO620, ATTO655, ATTO680, ATTO700, AZ503, AZ519, and ATTO647N were provided by ATTO-TEC (Siegen, Germany). Me-Tet-ATTO532, H-Tet-Cy3, Me-Tet-5-TAMRA, H-Tet-Cy5 were purchased from Jena Bioscience (Jena, Germany). HMSiR-NHS was purchased from MoBiTec (Goettingen, Germany), H-Tet-SiR from Spirochrome (Stein am Rhein, Switzerland) and TCO^{*}-Lys from SiChem (Bremen, Germany).

DNA constructs and plasmids. Plasmid amplification was performed via transformation in *E. coli* (XL1-Blue) and DNA isolation via MIDI-prep (NucleoBond[®] Xtra Midi, Macherey & Nagel, #740410). The respective amber stop mutants were generated by introducing a TAG codon through PCR-based site-directed mutagenesis. The plasmid for expression of the TNF-receptor pTNFR1-tdEOS was a gift from Mike Heilemann (Addgene plasmid #98273). The plasmid for the expression of the Kainate receptor (pcDNA3-GluK2) was a gift from Dr. Peter Seeburg (Max-Planck Institute for Medical Research, Heidelberg, Germany). An AgeI restriction site was introduced at the C-terminus of the GluK2 coding sequence in front of the

STOP-codon. Subsequently, GluK2 was cloned into the MCS of the pEGFP-N1 plasmid (Clontech #U55762) using the restriction enzymes EcoRI and AgeI. The plasmid for expression of the microtubule-associated protein Ensconsin (EMTB-3xGFP) was a gift from William Bement (Addgene plasmid #26741). The plasmid for the expression of the tRNA/tRNA-synthetase pair (pCMV tRNA^{Pyl}/NESPylRS^{AF}) was kindly provided by Edward Lemke⁵³.

Transfection. Cells were transfected at 60–80% confluence with a commercial transfection reagent (JetPrime, Polyplus, #114-01) with the suitable plasmid/reagent mixture (according to manufacturer's recommendations). The total DNA amount per well was 500 ng in a 1:1 ratio between the DNA for the respective POI and pCMV-tRNAPyl/NESPylRS^{AF}. The nCAA (TCO^{*}A; SiChem GmbH, Bremen, #SC-8008) was supplemented separately at a final concentration of 250 μM, diluted 1:4 with 1 M HEPES. After 6–8 h the medium was exchanged to fresh cell growth medium. Cells were incubated ~12–36 h before labeling and fixation/imaging.

Bioorthogonal labelling. For bioorthogonal labeling of clickable receptors the transfected HEK293T or COS-7 cells were incubated with 1.5 μM of the respective tetrazine-dye in cell growth medium for 10 min at room temperature or on ice. After incubation the cells were washed twice and fixed at room temperature with a mixture of 4% formaldehyde and 0.25% glutaraldehyde for 10 min. The cells were again washed three times with Hank's Balanced Salt solution (HBSS, Sigma, #55037 C) before imaging. Alternatively, the cells were imaged live without fixation. For labelling of actin filaments via pre-labeling with phalloidin-TCO the cells were first fixed with 4% formaldehyde in cytoskeleton buffer (10 mM MES pH 6.1, 150 mM NaCl, 5 mM EGTA, 5 mM glucose, 5 mM MgCl₂) at room temperature for 10 min. Cells were then washed 2–3 times with HBSS (ThermoFisher, #14025092) and then incubated for at least 1 h with phalloidin-TCO (~0.5 μM) at room temperature or alternatively for overnight at 4 °C. Afterwards the cells were washed

again 2–3 times with PBS and stored until further labeling/imaging. Fluorescence labeling was performed by adding the respective tetrazine-dye (1–3 μM in PBS) for 10 min at room temperature. Cells were then rinsed with 2–3 times PBS and imaged or, for wash-free applications, imaged without removing the tetrazine-dye solution.

dSTORM. The dSTORM images were acquired using an inverted wide-field fluorescence microscope (IX-71; Olympus). For excitation a 641 nm diode laser (Cube 640–100 C, Coherent, Cleanup 640/10, Chroma) was focused onto the back focal plane of the oil-immersion objective (60 \times , NA 1.45; Olympus). Emission light was separated from the illumination light using a dichroic mirror (635rpc, Chroma) and spectrally filtered by a bandpass filter (Em01-R442/514/647–25; Semrock). Images were recorded with an EMCCD (IXON DU897, Andor). Resulting pixel size for data analysis was measured as 128 nm. For each dSTORM measurement, at least 15,000 images with an exposure time of 20 ms and irradiation intensities of $\sim 2 \text{ kW/cm}^2$ were recorded by HILO (highly inclined and laminated optical sheet) illumination. Experiments were performed in PBS-based photoswitching buffer containing 100 mM β -mercaptoethylamine (MEA, Sigma-Aldrich) for Cy5 and an oxygen scavenger system (2% (w/v) glucose, 4 U/ml glucose oxidase and 80 U/ml catalase) or only in PBS for Cy5B and HMSiR, adjusted to pH 7.4. Image reconstruction was performed using rapidSTORM3⁶¹.

Chemical synthesis of H-Tet-HMSiR, H-Tet-Cy5B, Me-Tet-JF₆₄₆, and phalloidin-TCO. H-Tet-HMSiR: 0.034 μmol HMSiR-NHS (MoBiTec, Göttingen, Germany, #A208–01) was added to 0.68 μmol H-Tet-amine (Jena Bioscience, Jena, Germany, #CLK-001–5) in water-free DMSO (Thermo Fisher Scientific, Invitrogen Cat. Nr. D12345) and diisopropylethylamine (Sigma-Aldrich, #D125806, DIPEA, 1.1 μmol) and incubated for 4 h at room temperature.

H-Tet-Cy5B: 0.355 μmol Cy5B-NHS⁸ was added to 3.5 μmol H-Tet-amine (Jena Bioscience, Jena, Germany, #CLK-001–5) in water-free DMSO (Thermo Fisher Scientific, Invitrogen Cat. Nr. D12345) with diisopropylethylamine (Sigma-Aldrich, #D125806, DIPEA, 1.1 μmol) and incubated for 4 h at room temperature.

Me-Tet-JF₆₄₆ was synthesized accordingly using Me-Tet-amine. Phalloidin-TCO: 0.07 μmol Amino-phalloidin (Biomol, Hamburg, Germany, #ABD-5302) was added to 0.7 μmol TCO-PEG4-NHS (Jena Bioscience, Jena, Germany, #CLK-A137–10) in water-free DMSO (Thermo Fisher Scientific, Invitrogen Cat. Nr. D12345) with diisopropylethylamine (DIPEA, 1.1 μmol) and incubated for 4 h at room temperature. Tetrazine-dyes and phalloidin-TCO were purified by reverse phase chromatography on a Phenomenex Kinetex Biphenyl Core-shell LC Columns, (2.6 μm , 150 \times 4.6 mm; water:acetonitrile with 0.1% formic acid and an elution gradient of 0–95% in 45 min).

Measurements of absorbance/emission spectra. Time-dependent fluorescence intensities were measured in quartz glass cuvettes using a FP-6500 spectrofluorimeter (Jasco). Fluorescence was excited at the absorption maxima. The sample temperature was adjusted to 25 $^{\circ}\text{C}$ using a Peltier thermocouple.

Time-correlated single-photon counting (TCSPC). All fluorescence lifetime measurements at 640 nm excitation were performed with a FluoTime200 time resolved spectrometer (Picoquant, Berlin, Germany). Decay curves were analyzed using FluoFit 4.4.0.1 software. Measurements of fluorescence lifetimes of dyes with excitation at 400–560 nm were carried out with a MicroTime200 (Picoquant, Berlin, Germany), equipped with a supercontinuum laser SuperK Extreme EXW12 (NKT Photonics, Germany). The setup is attached to an Olympus IX83 with a 60 \times 1.2 ultra-plan-apochromat water-immersion objective.

Intermolecular fluorescence quenching experiments. To interpret quenching efficiencies of tetrazine and fluorophores, bimolecular quenching experiments with ATTO655, ATTO700, JF₆₄₆, and Cy5 were performed with Me-Tet-amine in phosphate-buffered saline (PBS) at pH 7.4. Bimolecular static and dynamic quenching constants, K_{stat} and K_{dyn} , were determined from time-resolved and steady-state fluorescence quenching experiments using Stern-Volmer analysis:

$$\tau_0/\tau = 1 + K_{\text{dyn}}[Q] = 1 + k_{\text{dyn}}\tau_0[Q] \quad (1)$$

$$F_0/F = (1 + K_{\text{stat}}[Q]) \left(1 + K_{\text{dyn}}[Q] \right) \quad (2)$$

where τ_0 and F_0 are the fluorescence lifetime and intensity in the absence of a quencher, τ and F are the fluorescence lifetime and intensity in the presence of the quencher Q with the concentration $[Q]$, and K_{stat} and K_{dyn} denote the static and dynamic Stern-Volmer constant, respectively. Dynamic quenching is attributed to collisional interactions at a dynamic quenching rate constant of k_{dyn} . Static quenching originates in bimolecular complexes that are very efficiently quenched and K_{stat} can thus be interpreted as an association constant K_a . Experimental estimates are provided as results from linear regression together with standard error from the fitting procedure.

Fluorescence-correlation-microscopy. PET-FCS was performed using a custom-built confocal fluorescence microscope setup as described previously^{39,40} equipped with a 640 nm diode laser (OBIS640, Coherent Europe B.V., Utrecht, Netherlands). Samples were diluted in PBS at pH 7.4 and filtered through a 0.2 μm syringe filter, transferred onto a microscope slide and covered by a cover slip. Samples with concentrations of 1 nM yielded an average of ~ 20 molecules in the detection focus of the microscope setup. Sample temperature was adjusted to 25 $^{\circ}\text{C}$ using a custom-built objective heater. For each sample, at least three individual autocorrelation functions were recorded of at least 5 min measurement time each. PET-FCS data was fitted to an analytical model for diffusion and a two-state equilibrium with off- and on-rate constants k_{closing} and k_{opening} :

$$G_D(\tau) = N^{-1} \left(1 - \frac{\tau}{\tau_D} \right)^{-1} \quad (3)$$

$$G(\tau) = G_D(\tau) \left(1 - A e^{-\tau/\tau_{\text{rel}}} \right) \quad (4)$$

where N is the number of detected molecules; τ_D is the diffusion time; $A = k_{\text{closing}}/k_{\text{opening}}$ is the equilibrium constant and $\tau_{\text{rel}} = (k_{\text{closing}} + k_{\text{opening}})^{-1}$ is the exponential time constant.

Confocal microscopy. Confocal fluorescence images were recorded on a commercial LSM700 (Zeiss; $\times 63/1.4$ oil objective). Single plane images were acquired with a pixel size of 90 nm and suitable settings for the respective dye. Images were adjusted for brightness and contrast (only linear changes).

Rescan confocal microscopy. Rescan confocal microscopy (RCM) imaging was performed using the commercially available RCM-unit (confocal.nl) attached to an inverse Nikon TiE microscope body. The system is equipped with a multiline (405, 488, 561, 640 nm) laser module Skyra (Cobolt) and a sCMOS camera Zylo4.2 P (Andor). Images were acquired with an $\times 100/1.49$ oil objective and a resulting pixel size of 42 nm. Images were corrected for the camera offset and a median filter (kernel: 1px) was applied.

Structured illumination microscopy (SIM). SIM imaging was performed on a commercial ELYRA S.1 microscope (Zeiss AG). The setup is equipped with a Plan-Apochromat $\times 63/1.40$ immersion-oil based objective and four excitation lasers, a 405 nm diode (50 mW), a 488 nm OPSP (100 mW), a 561 nm OPSP (100 mW) and a 642 nm diode laser (150 mW).

Cell culture and sample preparation. HEK293T cells (German Collection of Microorganisms and Cell Cultures, Braunschweig, Germany; #ACC635) were maintained in T25-culture flasks (Thermo Fisher Scientific, Cat. Nr. 156340) with Dulbeccos' Modified Eagle's Medium (DMEM, Sigma Aldrich, #D5796) with 10% FCS (Sigma-Aldrich, #F7524), 2 mM L-Glutamine (already in DMEM suppl.) and 100 U/ml Penicillin + 0.1 mg/ml Streptomycin (Sigma-Aldrich, #P4333) in a 5% CO₂ atmosphere at 37 $^{\circ}\text{C}$.

CO5-7 cells (Cell Lines Service GmbH, Eppelheim, Germany #605470) were maintained in DMEM (Sigma, #D8062) with 10% FCS (Sigma-Aldrich, #F7524), 2 mM L-Glutamine (already in DMEM suppl.) and 100 U/ml Penicillin + 0.1 mg/ml Streptomycin (Sigma-Aldrich, #P4333) in a 5% CO₂ atmosphere at 37 $^{\circ}\text{C}$.

NIH-3T3 (German Collection of Microorganisms and Cell Cultures, Braunschweig, Germany; #ACC59) were maintained in DMEM (Sigma, #D8062) with 10% FCS (Sigma-Aldrich, #F7524), 2 mM L-Glutamine (already in DMEM suppl.) and 100 U/ml Penicillin + 0.1 mg/ml Streptomycin (Sigma-Aldrich, #P4333) in a 5% CO₂ atmosphere at 37 $^{\circ}\text{C}$.

U2-OS cells (Cell Lines Service GmbH, Eppelheim, Germany # 300364) were maintained in DMEM (Sigma, #D8062) with 10% FCS (Sigma-Aldrich, #F7524), 2 mM L-Glutamine (already in DMEM suppl.) and 100 U/ml Penicillin + 0.1 mg/ml Streptomycin (Sigma-Aldrich, #P4333) in a 5% CO₂ atmosphere at 37 $^{\circ}\text{C}$.

Prior to seeding the cells, dishes were coated with 0.01% poly-D-Lysine (Sigma-Aldrich, #P6407) for 1 h at room temperature. For all live and fixed cell images, cells were seeded at least 16 h before transfection on 4-well Lab-Tek II chambered cover glasses (Nunc, cat. no. 155409).

Immunolabeling was performed with commercial anti-GluK2 (monoclonal: Merck-Millipore, #04–921, and polyclonal: Thermo Fisher Scientific, #PA5–32427) and anti-TNFR1 antibodies (monoclonal: abcam, #ab194814, and polyclonal: abcam, #ab19139). Antibodies were fluorescently labeled using the standard protocol for Alexa Fluor 647 NHS. Briefly, 50 μg of the purified antibodies were incubated in a 5 molar excess with NHS-Alexa647 (Thermo Fisher Scientific, #A37566) in 100 mM NaHCO₃ at RT for 3 h in the dark. Antibody conjugates were purified on gel filtration columns (NAP-5, Sephadex G25 DNA Grade, GE Healthcare). Finally, the degree of labeling of the purified antibody was determined by a UV-vis spectrophotometer (Jasco V-650) to DOL-1. Alternatively, primary antibodies were labeled with a secondary antibody (gamAlexa647, Thermo Fisher Scientific, #A-21235 and garAlexa647, Thermo Fisher Scientific, #A-21245, respectively). The conjugated antibodies were stored at 4 $^{\circ}\text{C}$.

Statistics and reproducibility. All experiments were performed at least three times unless otherwise stated. Statistical analysis of spectroscopic data was performed using OriginPro (OriginLab, MA). Experiments with one coherent dataset were performed successively to exclude instrumental variations.

Reporting summary. Further information on research design is available in the Nature Research Reporting Summary linked to this article.

Data availability

All data that support the findings described in this study are available within the manuscript and the related supplementary information, and from the corresponding authors upon reasonable request.

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References

- Sauer, M. & Heilemann, M. Single-molecule localization microscopy in eukaryotes. *Chem. Rev.* **117**, 7478–7509 (2017).
- Toomre, D. & Bewersdorf, J. A new wave of cellular imaging. *Annu. Rev. Cell Dev. Biol.* **26**, 285–314 (2010).
- Shannon, C. E. Communication in the presence of noise. *Proc. IRE* **37**, 10–21 (1949).
- Lukinavicius, G. et al. A near-infrared fluorophore for live-cell superresolution microscopy of cellular proteins. *Nat. Chem.* **5**, 132–139 (2013).
- Uno, S. N. et al. A spontaneously blinking fluorophore based on intramolecular spirocyclization for live-cell super-resolution imaging. *Nat. Chem.* **6**, 681–689 (2014).
- Grimm, J. B. et al. A general Method to improve fluorophores for live-cell and single-molecule microscopy. *Nat. Methods* **12**, 244–250 (2015).
- Grimm, J. B. et al. Bright photoactivatable fluorophores for single-molecule imaging. *Nat. Methods* **13**, 985–988 (2016).
- Michie, M. S. et al. Cyanine conformational restraint in the far-red range. *J. Am. Chem. Soc.* **139**, 12406–12409 (2017).
- Chen, F., Tillberg, P. W. & Boyden, E. S. Optical imaging. *Expans. Microsc. Sci.* **347**, 543–548 (2015).
- Tillberg, P. W. et al. Protein-retention expansion microscopy of cells and tissues labeled using standard fluorescent proteins and antibodies. *Nat. Biotechnol.* **34**, 987–992 (2016).
- Sauer, M. Localization microscopy coming of age: from concepts to biological impact. *J. Cell Sci.* **126**, 3505–3513 (2013).
- Ries, J., Kaplan, C., Platonova, E., Eghlidi, H. & Ewers, H. A simple, versatile Method for GFP-based super-resolution microscopy via nanobodies. *Nat. Methods* **9**, 582–584 (2012).
- Traenkle, B. & Rothbauer, U. Under the microscope: single-domain antibodies for live-cell imaging and super-resolution microscopy. *Front. Immunol.* **8**, 1030 (2017).
- Heilemann, M. et al. Subdiffraction-resolution fluorescence imaging with conventional fluorescent probes. *Angew. Chem. Int. Ed.* **47**, 6172–6176 (2008).
- van de Linde, S. et al. Direct stochastic optical reconstruction microscopy with standard fluorescent probes. *Nat. Protoc.* **6**, 991–1009 (2011).
- Chamma, I. et al. Mapping the dynamics and nanoscale organization of synaptic adhesion proteins using monomeric streptavidin. *Nat. Commun.* **7**, 10773 (2016).
- Virnat, D. et al. A peptide tag-specific nanobody enables high quality labeling for dSTORM imaging. *Nat. Commun.* **9**, 930 (2018).
- Wang, L., Frei, M. S., Salim, A. & Johnsson, K. Small-molecule fluorescent probes for live-cell super-resolution microscopy. *J. Am. Chem. Soc.* **141**, 2770–2781 (2019).
- Liu, C. C. & Schultz, P. G. Adding new chemistries to the genetic code. *Annu. Rev. Biochem.* **79**, 413–444 (2010).
- Chin, J. W. Expanding and reprogramming the genetic code of cells and animals. *Annu. Rev. Biochem.* **83**, 379–408 (2014).
- Lemke, E. A. The exploding genetic code. *ChemBioChem* **15**, 1691–1694 (2014).
- Prescher, J. A. & Bertozzi, C. R. Chemistry in living systems. *Nat. Chem. Biol.* **1**, 13–21 (2005).
- Kozma, E., Demeter, O. & Kele, P. Bio-orthogonal fluorescent labelling of biopolymers through inverse-electron-demand Diels-Alder reactions. *ChemBioChem* **18**, 486–501 (2017).
- Elsaesser, S. J., Ernst, R. J., Walkes, O. S. & Chin, J. W. Genetic code expansion in stable cell lines enables encoded chromatin modification. *Nat. Methods* **13**, 158–164 (2016).
- Ernst, R. J. et al. Genetic code expansion in the mouse brain. *Nat. Chem. Biol.* **12**, 776–778 (2016).
- Nikic, I., Kang, J. H., Girona, G. E., Aramburu, I. V. & Lemke, E. A. Labeling proteins in live mammalian cells using click chemistry. *Nat. Protoc.* **10**, 780–791 (2015).
- Devaraj, N. K., Hilderbrand, S., Upadhyay, R., Mazitschek, R. & Weissleder, R. Bioorthogonal turn-on probes for imaging small molecules inside living cells. *Angew. Chem. Int. Ed.* **122**, 2931–2934 (2010).
- Carlson, J. C. T., MeiMeis, L. G., Hilderbrand, S. A. & Weissleder, R. Bodipy-tetrazine derivatives as superbright bioorthogonal turn-on probes. *Angew. Chem. Int. Ed.* **52**, 6917–6920 (2013).
- Kozma, E., Girona, G. E., Paci, G., Lemke, E. A. & Kele, P. Bioorthogonal double-fluorogenic siliconrhodamine probes for intracellular super-resolution microscopy. *Chem. Commun.* **53**, 6696–6699 (2017).
- Schvartz, T. et al. Direct fluorescent-dye labeling of a-tubulin in mammalian cells for live cell and superresolution imaging. *Mol. Biol. Cell* **28**, 2747–2756 (2017).
- Knorr, G. et al. Bioorthogonally applicable fluorogenic cyanine-tetrazines for no-wash super-resolution imaging. *Bioconjugate Chem.* **29**, 1312–1318 (2018).
- Kozma, E. & Kele, P. Fluorogenic probes for super-resolution microscopy. *Org. Biomol. Chem.* **17**, 215–233 (2019).
- Hoffmann, J. E. et al. Highly stable trans-cyclooctene amino acids for live-cell labeling. *Chemistry* **21**, 12266–12270 (2015).
- Carlson, J. C. T., Mikula, H. & Weissleder, R. Unraveling tetrazine-triggered bioorthogonal elimination enables chemical tools for ultrafast release and universal cleavage. *J. Am. Chem. Soc.* **140**, 3603–3612 (2018).
- Karver, M. R., Weissleder, R. & Hildebrand, S. A. Synthesis and evaluation of a series of 1,2,4,5-tetrazines for bioorthogonal conjugation. *Biocon. Chem.* **22**, 2263–2270 (2011).
- Kormos, A. et al. Bistetrazine-cyanines as double-clicking fluorogenic two-point binder or crosslinker probes. *Chem. Eur. J.* **24**, 8841–8847 (2018).
- Meimets, L. G., Carlson, J. C. T., Giedt, R. J., Kohler, R. H. & Weissleder, R. Ultrafluorogenic coumarin-tetrazine probes for real-time biological imaging. *Angew. Chem. Int. Ed.* **53**, 7531–7534 (2014).
- Doose, S., Neuweiler, H. & Sauer, M. A close look at fluorescence quenching of organic dyes by tryptophan. *ChemPhysChem* **6**, 2277–2285 (2005).
- Doose, S., Neuweiler, H. & Sauer, M. Fluorescence quenching by photoinduced electron transfer: a reporter for conformational dynamics of macromolecules. *ChemPhysChem* **10**, 1389–1398 (2009).
- Troll, Th Reduction potentially of substituted as-triazines and s-tetrazines in acetonitrile. *Electrochim. Acta* **27**, 1311–1314 (1982).
- Gong, Y.-H. et al. Synthesis and physical chemistry of s-tetrazines: which ones are fluorescent and why? *Eur. J. Org. Chem.* **35**, 6121–6128 (2009).
- Magde, D., Elson, E. L. & Webb, W. W. Fluorescence correlation spectroscopy. II. An experimental realization. *Biopolymers* **13**, 29–61 (1974).
- Chattopadhyay, K., Saffarian, S., Elson, E. L. & Frieden, C. Measurement of micro-second dynamic motion in the intestinal fatty acid binding protein by using fluorescence correlation spectroscopy. *Proc. Natl Acad. Sci. USA* **99**, 14171–14176 (2002).
- Neuweiler, H., Doose, S. & Sauer, M. A microscopic view of mini-protein folding: enhanced folding efficiency through formation of an intermediate. *Proc. Natl Acad. Sci. USA* **102**, 16650–16655 (2005).
- Doose, S., Neuweiler, H., Barsch, H. & Sauer, M. Probing polyproline structure and dynamics by photoinduced electron transfer provides evidence for deviations from a regular PPII helix. *Proc. Natl Acad. Sci. USA* **104**, 17400–17405 (2007).
- Widengren, J. & Schwille, P. Characterization of photoinduced isomerization and back-isomerization of the cyanine dye Cy5 by fluorescence correlation spectroscopy. *J. Phys. Chem. A* **104**, 6416–6428 (2000).
- Müller, C. B. & Enderlein, J. Image scanning microscopy. *Phys. Rev. Lett.* **104**, 198101 (2010).
- De Luca, G. M. R. et al. Re-scan confocal microscopy: scanning twice for better resolution. *Biomed. Opt. Express* **4**, 2644–2656 (2013).
- Gustafsson, M. G. Surpassing the lateral resolution limit by a factor of two using structured illumination microscopy. *J. Microsc.* **198**, 82–87 (2000).
- Lerma, J. Roles and rules of kainate receptors in synaptic transmission. *Nat. Rev. Neurosci.* **4**, 481–495 (2003).
- Locksley, R. M., Killeen, N. & Lenardos, J. The TNF and TNF receptor superfamilies: integrating mammalian biology. *Cell* **104**, 487–501 (2001).
- Williams, S. et al. Antibody-mediated inhibition of TNFR1 attenuates disease in a mouse model of multiple sclerosis. *PLoS ONE* **9**, e90117 (2014).
- Nikic, I. et al. Debugging eukaryotic genetic code expansion for site-specific click-PAINT super-resolution microscopy. *Angew. Chem. Int. Ed.* **55**, 16172–16176 (2016).
- Neubert, F. et al. Bioorthogonal click chemistry enables site-specific fluorescence labeling of functional NMDA receptors for super-resolution imaging. *Angew. Chem. Int. Ed.* **57**, 16364–16369 (2018).

55. Faire, K. et al. E-MAP-115 (ensconsin) associates dynamically with microtubules in vivo and is not a physiological modulator of microtubule dynamics. *J. Cell Sci.* **112**, 4243–4255 (1999).
56. Dumontet, C. & Jordan, M. A. Microtubule-binding agents: a dynamic field of cancer therapeutics. *Nat. Rev. Drug Discov.* **9**, 790–803 (2010).
57. Lukinavicius, G. et al. Fluorogenic probes for live-cell imaging of the cytoskeleton. *Nat. Methods* **11**, 731–733 (2014).
58. Lee, Y., Cho, W., Sung, J., Kim, E. & Park, S. B. Monochromophoric design strategy for tetrazine-based colorful bioorthogonal probes with a single fluorescent core skeleton. *J. Am. Chem. Soc.* **140**, 974–983 (2018).
59. Miller, A. L. & Berment, W. M. Regulation of cytokinesis by Rho GTPase flux. *Nat. Cell Biol.* **11**, 71–77 (2009).
60. Mitchison, T. & Kirschner, M. Dynamic instability of microtubule growth. *Nature* **312**, 237–242 (1984).
61. Wolter, S. et al. rapidSTORM: accurate, fast open-source software for localization microscopy. *Nat. Methods* **9**, 1040–1041 (2012).

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Author contributions

G.B. and M.S. conceived and designed the project. M.S. and S.D. supervised the project. G.B., A.J.K., L.B-P., M.M. and A.C.K. performed all labeling and imaging experiments.

Z.-D.S., M.Sch., J.B.G. and L.D.L., provided the dyes Cy5B and JF₆₄₆, and corresponding tetrazine derivatives. N.W. and J.S. provided docetaxel-TCO. G.B., A.J.K. and S.D. performed data analysis. G.B., S.D., and M.S. wrote the paper.

Additional information

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