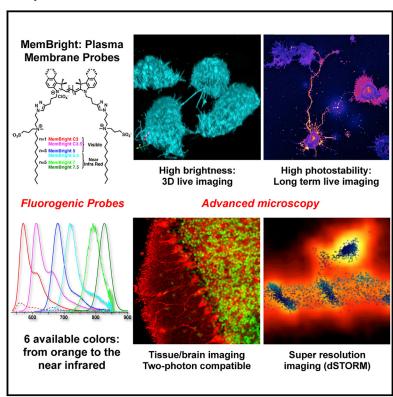
# **Cell Chemical Biology**

# MemBright: A Family of Fluorescent Membrane Probes for Advanced Cellular Imaging and Neuroscience

## **Graphical Abstract**



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#### In Brief

Collot et al. developed MemBright: a family of fluorescent molecules that stain in different colors the cell plasma membrane in a bright and efficient manner. These probes are compatible with various microscopy techniques including live videomicroscopy, multiphoton, and super-resolution. MemBright probes are particularly suitable in neuroscience and cell biology imaging.

#### **Highlights**

- MemBright comprises 6 fluorescent probes emitting from orange to near infrared
- Turn-on probes with fast, homogeneous, and backgroundfree staining
- Compatible with live/fixed cells and with immunostaining
- Compatible with long-term, 2-photon, tissue and superresolution imaging





## MemBright: A Family of Fluorescent Membrane Probes for Advanced Cellular Imaging and Neuroscience

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#### SUMMARY

The proper staining of the plasma membrane (PM) is critical in bioimaging as it delimits the cell. Herein, we developed MemBright, a family of six cyaninebased fluorescent turn-on PM probes that emit from orange to near infrared when reaching the PM, and enable homogeneous and selective PM staining with excellent contrast in mono- and two-photon microscopy. These probes are compatible with long-term live-cell imaging and immunostaining. Moreover, MemBright label neurons in a brighter manner than surrounding cells, allowing identification of neurons in acute brain tissue sections and neuromuscular junctions without any use of transfection or transgenic animals. In addition, MemBright probes were used in super-resolution imaging to unravel the neck of dendritic spines. 3D multicolor dSTORM in combination with immunostaining revealed en-passant synapse displaying endogenous glutamate receptors clustered at the axonal-dendritic contact site. MemBright probes thus constitute a universal toolkit for cell biology and neuroscience biomembrane imaging with a variety of microscopy techniques.

#### INTRODUCTION

Plasma membrane (PM), in addition to its basic function of cell barrier, is a key player in crucial biological processes including cellular uptake, neural communication, muscle contraction, and cell trafficking and signaling (Lingwood and Simons, 2010). In the field of bioimaging, visualizing the PM is of prior importance as it delimits the cell surface. In addition, the shape of the PM directly provides information regarding the cell status such as cell division or cell death processes (Rello et al., 2005). Visualizing PM is particularly important in neuroscience because it enables visualization of neuronal organization and membrane

trafficking involved in the synaptic transmission (Matsuzaki et al., 2004). Recent years have seen a tremendous expansion of fluorescence imaging techniques and tools for cellular research. In addition to genetically encoded fluorescent proteins, a number of molecular probes for monitoring cellular events have been developed (Lavis and Raines, 2008; Fernández-Suárez and Ting, 2008). The key challenge in cellular imaging is to stain cell compartments with high specificity and persistence. Although numerous efficient molecular probes have been designed to selectively stain specific cellular structures including mitochondria (Neto et al., 2013), lipid droplets (Collot et al., 2018), ER (Arai et al., 2014), nucleus (Thermo Fisher), and lysosomes (Despras et al., 2015; Chen et al., 2015), there is still a demand for efficient and bright fluorescent PM markers (Klymchenko and Kreder, 2014). Fluorescently labeled lectins, notably wheat germ agglutinin (WGA) and concanavalin A, are popular fluorescent membrane probes. Despite their efficiency and ease of use, these probes are expensive and much larger than molecular probes. The small size of the latter, comparable with lipids, allows their precise location in the lipid bilayer, which is of key importance for studies of the lateral lipid organization of biomembranes (lipid rafts; Klymchenko and Kreder, 2014; Owen et al., 2012), fluorescence resonance energy transfer with membrane proteins (Ziomkiewicz et al., 2013) and superresolution imaging (Solanko et al., 2013; Honigmann et al., 2014). Therefore, molecular probes are an interesting alternative to protein-based membrane markers. Although highly hydrophobic fluorophores are efficient markers of membrane models such as liposomes, they generally fail in staining the cell PM of live cells as they tend to precipitate before reaching the membrane and quickly cross the PM to stain inner membranes (Klymchenko and Kreder, 2014; Niko et al., 2016). Recent efforts have been made to develop specific and efficient PM probes for cell imaging with various fluorophores including chromone (Kreder et al., 2015; Shynkar et al., 2007), Nile red (Kucherak et al., 2010), styryl pyridinium (Kwiatek et al., 2013), BODIPY (López-Duarte et al., 2014; Zhang et al., 2014), perylene (Heek et al., 2013), oligothiophene (Dal Molin et al., 2015; Fin et al., 2012), conjugated polymers (Cui et al., 2016), or fluorescein isothiocyanate-labeled chitosan (Wang et al., 2015; Jiang et al., 2016). From the rapid



development of multicolor imaging arose a demand for redshifted fluorescent markers. The latter, especially near-infrared ones, are highly popular for bioimaging applications due to their long wavelength (650-900 nm) excitation and emission that reduce photodamage to biological samples, allow deep tissue penetration, and avoid background signal from cell autofluorescence (Yuan et al., 2012; Umezawa et al., 2014). We recently proposed an efficient far-red-emitting PM fluorescent marker called dSQ12S based on a squaraine dye with two zwitterionic amphiphilic anchors, exhibiting superior staining compared with commonly used hydrophobic cyanines, such as DiD and the PKH family (Collot et al., 2015). Indeed, we showed that replacing the octadecyl (C-18) chains of Di-dyes with more polar dodecyl anchors bearing zwitterionic head groups enhance the polarity of the probe, thus facilitating the dissociation of the formed dye aggregates in the presence of membrane.

Application of this unique design concept to cyanine family, highly popular in bioimaging applications (Gonçalves, 2009), could open numerous opportunities. First, cyanines are among the brightest dyes that can be obtained in any desired emission color spanning from the visible (orange) to the near infrared. Second, due to their capacity to form non-fluorescent H-aggregates, they can be transformed into fluorogenic probes (Hou et al., 2015). Third, cyanine dyes have been established as robust tool for super-resolution microscopy (SRM) (van de Linde et al., 2011), which could enable us to propose new membrane probes particularly adapted to SRM. Herein, we thus designed new membrane probes based on zwitterionic anchors and various cyanine fluorophores using click chemistry. These probes provide very selective and bright PM staining and surpass all tested commercial dyes. They constitute a versatile tool as they allow live, fixed, and permeabilized cells/tissue imaging using mono- or two-photon imaging. As a proof of concept we unraveled the nanoscale organization of the axonal enpassant synapse coiling around the dendrite using 3D multicolor SRM.

#### **RESULTS**

#### **Synthesis of MemBright and Spectroscopic Studies**

Spectroscopic properties of cyanines depend on the length of their polymethine chain: the longer the chain, the higher the wavelength it emits. A finer color tuning can be achieved by replacing the indole moieties by benz[e]indole ones. This modification extends the electronic system and therefore gives rise to red-shifted cyanines, named with a ".5" suffix. Herein, to obtain probes of different colors, we first synthesized six dialkyne cyanines (Figure 1A). The latter were reacted with a clickable amphiphilic zwitterion via CuAAC (copper-catalyzed azide alkyne cycloaddition) to provide the MemBright probes family (Figure 1A).

Spectroscopic studies of MemBright were performed in various organic solvents and buffer, as well as in the presence of large unilamellar vesicles (LUVs) composed of dioleoylphosphatidylcholine (DOPC) with and without cholesterol, taken as membrane models. In organic solvents, MemBright probes behaved in the same manner as their parent cyanine fluorophores. Indeed, their absorption spectra were narrow and close to the mirror image of their emission spectra, indicating no sign of

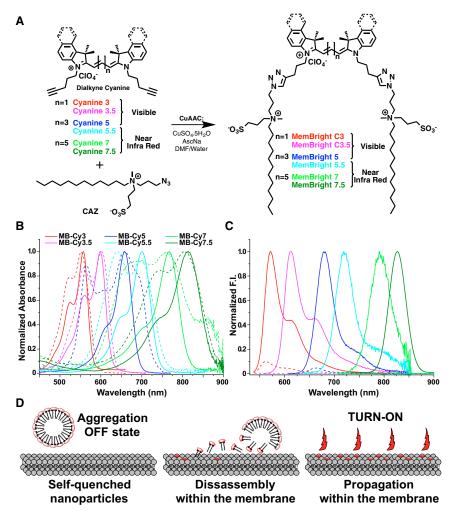
aggregation (Figure S1). Moreover, they displayed good quantum yields (0.12–0.66 in DMSO).

Due to their amphiphilic nature, MemBright markers are expected to form soluble aggregates in aqueous media. Indeed, dynamic light scattering (DLS) measurements in water showed formation of nanoparticles of 22-32 nm (Figure S2E). The critical aggregation concentrations of MemBright were determined by spectroscopy (Figure S2) and were shown to be below 50 nM except for MB-Cy3 and MB-Cy5 (1 µM and 120 nM, respectively). Moreover, their absorption spectra in phosphate buffer displayed a strong short-wavelength shoulder, suggesting formation of non-fluorescent H-type aggregates, known for their blue-shifted absorption (Würthner et al., 2011) (Figure 1B). As expected, MB-Cy3 to MB-Cy5 probes displayed much lower quantum yields in buffer than in organic solvents, while MB-Cy5.5, MB-Cy7, and MB-Cy7.5 were nearly non-fluorescent in water (Table S1). In the presence of LUVs, MemBright probes displayed typical absorption and fluorescence spectra of solubilized cyanine dyes (Figures 1B and 1C) with quantum yield generally higher than those obtained in methanol (Table S1). The fluorescence enhancement between the aqueous media and the DOPC vesicles was 7-, 42-, and 18-fold for MB-Cy3, MB-Cy3.5, and MB-Cy5, respectively, whereas for MB-Cy5.5 to MB-Cy7.5 it was much larger due to their strong quenching in aqueous media. Thus, MemBright aggregated in water disassemble in lipid membranes (Figure 1D), providing a fluorogenic response. This mechanism is in line with the earlier data obtained with membrane probes based on Nile red (Kucherak et al., 2010) and squaraine (Collot et al., 2015).

To determine how fast the aggregates of MemBright disassemble in the presence of LUVs, we added probes to an excess of LUVs and followed their fluorescence emission intensity over time (Figure S2F). Whereas MB-Cy3 and MB-Cy5 rapidly reached their fluorescence intensity plateau in the presence of DOPC vesicles (<3 min), the other probes disassembled in a slower manner. Overall, the disassembly time increased with the polymethine chain length (Cy3 < Cy5 < Cy7) and addition of fused benzene rings (Cy3 < Cy3.5 and Cy5 < Cy5.5). We expect that more hydrophobic MemBright probes tend to form more stable aggregates, which require more time to disassemble in the presence of membranes.

#### **Cellular Imaging**

The ability of MemBright probes to selectively stain the PM was first investigated on live HeLa and KB cells. HeLa cells were chosen as they are widely used human cell lines. KB human cells (derived from HeLa cells) were chosen for their simple shape and appropriate height that facilitates PM imaging. A solution of MemBright probes (20 nM) in a serum-free medium was added to the cells and the latter were directly imaged without washing. WGA-488, a green-emitting membrane marker based on a fluorescently labeled lectin (wheat germ agglutinin), was used to localize the PM. After 5 min, the PM displayed a clean and bright fluorescent staining with a clear co-localization of MemBright probes with WGA-488 (Figure 2A for KB cells, Figure S3 for HeLa cells) and high signal-to-background ratio (up to 18, see Table S2). Vertical projections of the confocal images confirmed perfect membrane staining by MemBright probes with no apparent internalization (Figures 2B and S4). The quality of the



staining also allowed for 3D imaging of live KB cells using reconstruction of z stacks obtained by laser scanning confocal microscopy (Figure 2C and Video S1). These images offered high contrast and clearly revealed intercellular nanotubes, also called tunneling nanotubes (Gousset et al., 2009) as bright thin PM structures connecting two individual cells (Figure 2C).

According to our recent report, in similar imaging conditions the commercial dye DiD at 20 nM concentration failed to stain cellular membranes (Collot et al., 2015). Indeed, due to its highly hydrophobic nature, DiD is poorly soluble in aqueous media, which renders PM staining very inefficient. Another commercial probe, CellVue Claret from the PKH family, is also very hydrophobic and thus requires a specific low-salt medium for PM staining. Moreover, its cell membrane staining is heterogeneous (Collot et al., 2015). Therefore, commercial cyanines, such as DiD (and its analogs Dil and DiO) and the PKH family as well as CellMask membrane stains (Thermo Fisher) require concentrations of around 1-5 µM. Thus, MemBright probes require ~250-fold lower concentration compared with commercial membrane probes. Surprisingly, although MB-Cy7 also possesses a near-infrared excitation wavelength (~760 nm), it was sufficiently excited by the 635-nm laser (corresponding to 10%

Figure 1. MemBright Probes and Their **Properties** 

(A) Synthesis of the MemBright markers. (B and C) Absorption (B) and emission spectra (C) of MemBright probes (200 nM) in the absence (dashed lines) or presence of DOPC vesicles.

(D) Turn-on mechanism of the MemBright probes.

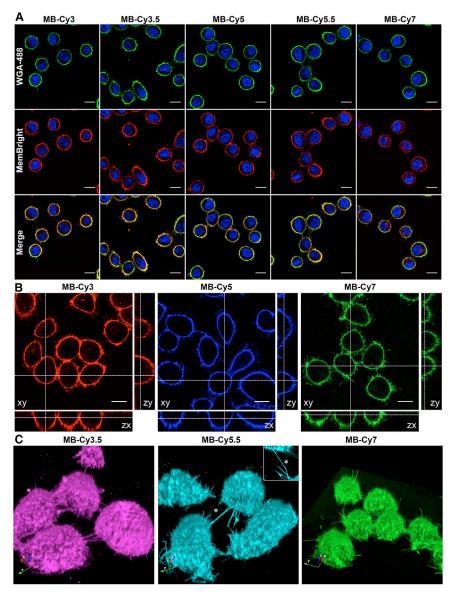
of its peack absorbance) and was bright enough to provide good-quality 2D and 3D images (Figure 2).

In addition, compared with fluorescently labeled lectin WGA-488, MemBright probes displayed a more homogeneous staining of the PM when cells were confluent (Figure S5). In the focal plane, most of the confluent cells' PM is involved in cell-cell contacts. Within these junctions, carbohydrates located on the outer leaflet of the PM are much less accessible for the lectin than those exposed to the external medium. It was measured that in cell junctions, MemBright probes stain the PM >3-fold more efficiently than WGA-488 (Figure S5D). This feature is due to rapid diffusion of MemBright within the whole PM, which ensures a homogeneous staining.

Next, the fluorescent staining of the PM was checked 90 min after addition of MemBright probes. For MB-Cy3 and MB-Cy5, the results showed that although the staining predominantly remains on the PM, internalization partially occurred as some dots appeared in the cytoplasm, revealing the endocytosis

process. In contrast, the red-shifted versions of MB-Cy3 and MB-Cy5, namely MB-Cy3.5 and MB-Cy5.5, conserved a very selective PM staining with less apparent sign of internalization (Figure S6). We expect that the additional fused benzene rings, which significantly increase the lipophilicity of these probes, should either favor their escape from the endocytosis process or stimulate their recycling back to the PM after endocytosis. Consequently, MB-Cy3 and MB-Cy5 will be preferred for short-term PM imaging, whereas MB-Cy3.5 and MB-Cy5.5 will be better for longer-term imaging. At this stage of our work, it was important to evaluate the cytotoxicity of the MemBright probes. MTT assays were performed on KB cells and the results showed no apparent cytotoxicity of the probes at concentrations of up to 1 μM (Figure S7).

Formaldehyde-based cell fixation is an indispensable tool in cellular imaging. However, it leads to partial permeabilization of the PM, so that application of molecular PM probes in fixed cells is challenging. To check the possible application of Mem-Bright for imaging the PM of fixed cells, we first tried to add the dyes on fixed cells (4% paraformaldehyde [PFA]). Although specific staining was obtained (Figure S8), it was noticed that prolonged time of imaging led to diffusion of the dyes within



the fixed cells. However, when images obtained with WGA-488 and MemBright on fixed cells were compared, we noticed that WGA tended to stain the perinuclear envelope as well as the inner vesicles in addition to PM, whereas MemBright probes stained the PM more selectively (Figure S8B). As an alternative attempt, the dyes were added to the cells prior to fixation. This approach provided much cleaner and longer staining of the PM with only a slight amount of internalized dye, demonstrating the compatibility of these probes with fixed cells (Figure S8C).

#### **Two-Photon Excitation Microscopy**

Two-photon excitation (TPE) microscopy imaging has received considerable attention as an advanced optical technique in the field of bioimaging for several reasons (Helmchen and Denk, 2005; Svoboda and Yasuda, 2006). First, TPE microscopy inherently provides 3D sectioning since the excitation occurs only at the focal point of the sample. In addition, due to its

Figure 2. 2D and 3D Confocal Live-Cell Imaging Using MemBright

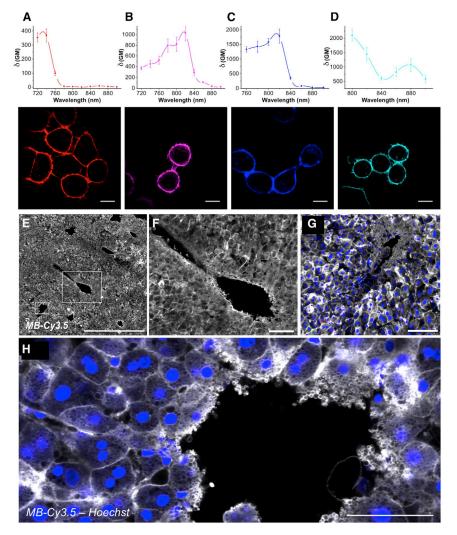
(A) Laser scanning confocal microscopy of live KB cells 5 min after addition of the probes (20 nM of MemBright, except for MB-Cy7, 15 min) without any washing step. WGA-488 was used as a costaining marker (excitation 488 nm). Nuclear staining was done with Hoechst (5 µg/mL). Scale

(B) Orthogonal projection obtained from z stacks (for MB-Cy3.5 and MB-Cy5.5, see Figures S10 and S11). Scale bars, 15 μm.

(C) 3D reconstruction of live KB cells stained with MemBright, Inset in the center panel is the top view of the intercellular nanotube indicated by the asterisk.

localized excitation with a near-infrared laser, it reduces photobleaching, phototoxicity, and cell autofluorescence. It also ensures deeper tissue penetration, which enables experiments on thicker live samples. For these reasons, there is a high demand for bright fluorescent probes with high TPE cross-section values. Ongoing efforts have been made to design such fluorophores (Yao and Belfield, 2012). Although cyanine fluorophores are widely used for biolabeling and bioimaging, only few studies have reported on their efficiency for TPE imaging (Lukomska et al., 2005; Kobat et al., 2009; Albota et al., 1998). Since MemBright markers are localized in PMs in cell experiments, their twophoton fluorescence excitation spectra were measured in an aprotic solvent. DMSO (Figures 3A-3D, top). The twophoton nature of the absorption process was verified from the quadratic dependence of the observed emission intensity versus excitation power (Figure S9). To our surprise, MemBright probes dis-

played high two-photon action cross-section in this solvent with values reaching up to 1,200 Goeppert-Mayer (GM) for MB-Cy5. It was remarkable that fused benzene rings significantly increased TPE cross-section for MB-Cy3.5 and shifted significantly the TPE absorption to the red for both MB-Cy3.5 and MB-Cy5.5 compared with their parent analogs (Figures 3A-3D). To confirm the results obtained for the MemBright probes, we measured the TPE spectrum of the parent dye, DiD (lipophilic cyanine-5), in DMSO and found it to be almost identical to that of MB-Cy5 (Figure S9E). In the light of these unexpected results, MemBright probes were used in twophoton imaging experiments. Live KB cells were stained with MemBright probes without washing and imaged with twophoton excitation, chosen according to their TPE excitation spectra. High signal-to-noise ratio images were obtained for each of the MemBright probes with highly selective staining of the PM (Figures 3A-3D, bottom).



#### Tissue Imaging

Fluorescence tissue imaging with molecular probes is more challenging, as the environment is more complex and presents heterogeneity of components (cells, extracellular matrix, ducts, vessels) and fluids (e.g., plasma, blood, bile) (Kim and Cho, 2015; Guo et al., 2016). Here, applicability of MemBright probes to ex vivo tissue imaging was tested. For this purpose, 1-mm-thick blocks of rat liver were incubated for 3 h in the presence of MB-Cy5. The liver tissues were imaged by two-photon microscopy providing stacks of 50 images up to 50 µm depth. In the case of MB-Cy5 (Figure S10), a clear PM staining was obtained and hepatocytes appeared as polygonal cells (Figure S10G). At 20 µm depth, hepatocytes were well delimited by an intense PM staining, and projections on the yz and xz planes (Figure S10) showed that the PM can be stained on up to 3 layers of cells in the tissue (see Video S2). The lower intensity at higher depths is clearly related to the limited diffusion of the probe through the tissue, which in principle could be improved by optimizing the staining conditions. In sharp contrast, WGA-Alexa647, used in the same conditions, showed unclear staining that appeared mainly in the nuclei (Figures S10D-S10F; Video S3).

#### Figure 3. Two-Photon Properties and Tissue Imaging

(A-D) TPF cross-section spectra (DMSO) and images of KB cells using MB-Cy3 (A), MB-Cy3.5 (B), MB-Cy5 (C), and MB-Cy5.5 (D). The images were obtained without any washing step. Conditions: MB-Cy3 at 40 nM,  $\lambda_{Ex} = 760$  nm (red); MB-Cy3.5 at 40 nM,  $\lambda_{Ex}$  = 810 nm (magenta); MB-Cy5 at 40 nM,  $\lambda_{Ex}$  = 810 nm (blue); MB-Cy5.5 at 120 nM,  $\lambda_{Ex}$  = 860 nm (cyan). The signal was collected using a short-pass filter with 700-nm cutoff wavelength, which explained the use of MB-Cy5.5 at higher concentration, Laser power was set to 5 mW, Each image is the average of five acquisitions. Scale bars, 10 μm.

(E-G) Tile confocal imaging (E) of 1-mm-thick liver slices incubated with MB-Cy3.5. The mosaic is composed of 4 x 4 fields (20x objective). Scale bar, 1 mm. White square corresponds to the magnification in (F) to (H), which show the central vein of the hepatic lobe (G) 3D rendering of the 20× stack (101 z slices, 55 μm thick) (scale bars, 100 µm). MB-Cy3.5 is resistant to continuous illumination and allows detection of cells and tissue arrangement without any other counterstaining.

(H) Confocal 63× zoom around the central vein. MB-Cy3.5 labels the PM but also the intracellular hepatic vesicles of the cells that have been cut during slicing. Scale bar, 50 µm.

Thus, MemBright is clearly advantageous compared with the commercial agent for PM staining in tissue imaging. MemBright Cy3.5 was also very efficient in PM labeling using either TPE microscopy or conventional confocal microscopy. We tested the photoresistance of MemBright by performing either mosaic (Figures 3E and 3F) or 3D imaging (Figure 3G and

Video S4) in large 1-mm-thick liver slices. MB-Cy3.5 was resistant to continuous illumination and mosaic imaging even when acquiring 3D tile picture reconstruction (>100 z slices), indicating a good resistance to photobleaching. At high magnification the high signal-to-noise ratio of MB-Cy3.5 allowed clear discrimination of the PM of hepatocytes and even their intracellular lipoprotein vesicles (Figure 3H).

Encouraged by these results, we investigated the ability of MemBright to stain the PM in brain slices. The study of neuronal arborization has been investigated for years in various types of mutant animals using either Dil labeling thanks to biolistic approaches (Rasia-Filho, 2010) (Dil impregnated gold beads projected over neurons through gene gun) or with the use of transgenic mice reporter (Thy-1-GFP) (Feng et al., 2000; Ricard and Debarbieux, 2014) expressing cytosolic fluorescent protein in neuronal cytoplasm. Although those techniques are robust and accurate, they provide heterogeneous staining. Moreover, they are technically difficult to set up and time consuming. Herein, brain slices were incubated in the presence of MemBright before being imaged by different fluorescence imaging techniques. Taking advantage of its high two-photon absorption cross-section and red-shifted emission, MB-Cy3.5 succeeded in providing good-quality images of pyramidal neurons in the hippocampus of mouse brain slices (Figures 4A–4C). The images clearly revealed the *stratum pyramidale* region (light-blue layer in Figures 4B and 4C), where the soma of the pyramidal neurons were highly stained as well as their dendrites, as thinner structures, localized in the *stratum radiatum*. 3D rendering over z stacks (Figure 4D) allowed visualization of the dendrite deep in the tissue as well as dendritic spines (arrows in Figure 4D). Thus, MemBright enables fast visualization of neuronal cell-layer packing as well as dendrite density.

Moreover, these results suggest that MemBright probes are able to label neuronal cells at higher intensity than other cell types. We then checked whether this characteristic was specific to hippocampal pyramidal cells or whether MemBright could be used in other regions of the brain. Slices (1 mm thick) of cortex (Figures 4E and 4F) and cerebellum (Figure 4G) were efficiently labeled and imaged on either confocal (Figure 4E) or TPE imaging (Figure 4F). Cortical neurons and Purkinje cells can be clearly seen in Figures 4F and 4G, respectively. Altogether, these results indicate that MemBright has the propensity to label some major neuronal cell types.

As MemBright displayed a high affinity toward neurons, we wanted to demonstrate whether it was able to reveal motor neurons in tissue imaging. To this aim, we extracted muscle fibers with nerves from *tibialis anterior* muscles of an adult mouse, and fixed and incubated them with MB-Cy5. Fluorescently labeled α-bungarotoxin, which binds to nicotinic receptors clustered at synapse, was used to label the neuromuscular junctions and served to visualize the whole muscle fiber. As shown in Figure S11, the whole axon and the terminal motor nerve was clearly revealed by MB-Cy5. Both live and fixed samples could be efficiently labeled without any permeabilization; the staining was sufficiently resistant to confocal stack imaging. 3D reconstruction allowed visualization of the nerve terminal in close apposition with the motor plate (Figures S11B-S11D).

#### **Neuronal and Glial Cell Imaging**

The ability of MemBright to label polarized cells organized in networks (neurons) or in confluent layers (pavement of astrocytes) was investigated. In this endeavor we used hippocampal neuronal cells, which can be seeded and fed on an astrocyte feeder layer (Danglot et al., 2003). Primary hippocampal cultures are efficiently labeled with any member of the MemBright family (Figures 5 and S12), revealing neuronal cells lying over astrocytes. This experiment confirmed what was previously noticed in tissue imaging, namely that neurons appeared much brighter than glial cells, suggesting a preferential staining of neurons by MemBright. Although MemBright probes were used at very low concentration (typically, 20 nM directly in the imaging medium), they revealed even very fine protrusions including astrocytic filopodia (Figures 5A3, 5A4, 5E, and 5E1) or fine dendritic processes (Figures 5A-5D). Unlike the use of fluorescent proteins, this method leads to the labeling of all cells in live microscopy without any transfection steps that are poorly efficient and can be deleterious in neuronal cells.

Live cells could be labeled and imaged on confocal or spinning-disc microscopy in combination with other vital staining (Figure S12). Very interestingly, labeled live cells could be kept with the dye during several days without any visible detrimental effects. Indeed, neuronal culture composed of hippocampal primary neurons and glial cells was labeled with MemBright and imaged over several hours under confocal illumination. The brightness of MemBright and its low toxicity allowed us to monitor the cells with only 0.2% power of 561 nm laser line over 13 h without any loss of membrane staining (Video S5). Fine neuronal filopodia and widespread glial lamellipodia could be easily tracked. MemBright is thus able to provide efficient membrane labeling over long periods even at 37°C. We then compared MemBright labeling on live neurons with other classical membrane probes. As expected from HeLa and KB cell labeling, the WGA staining on neurons was very bright and homogeneous from cell to cell (Figure S13A). However, when carefully inspecting the PM, we noticed dashed staining along the PM, impeding fast segmentation of cell shape (Figure S13B). This might be attributed to the non-homogeneous expression of glycosylated protein at the PM. To circumvent this problem, we reasoned that staining by lipidic dyes should be more homogeneous. Consequently, MemBright was compared with DiD (Figures S13C and S13D) over the neuronal network. Although DiD labeling, compared with WGA, was more homogeneous along the PM and allowed perfect visualization of the cell shape, staining intensity from cell to cell was variable, ranging from very faint (white arrowhead in Figures S13C and S13D) to very bright (red arrowhead in Figures S13C and S13D). MemBright was then compared with mCLING, a polylysine-based membrane probe developed for super-resolution imaging (Revelo et al., 2014). For this purpose, we strictly followed the protocol provided by the developers of mCLING (Revelo and Rizzoli, 2001). As shown in Figure S14, after 10 min of incubation mCLING labeling was mostly punctiform and seemed to be mainly internalized in intracellular vesicles as shown previously (Revelo et al., 2014). Conversely, MemBright was still present at the PM and clearly allowed the identification of neuronal cell bodies (Figure S14) as well as axonal or dendritic networks.

#### Monitoring Membrane Trafficking and Cellular Architecture with Endogenous Proteins

We then explored whether MemBright allowed tracking of endocytic vesicles. We incubated MemBright at 37°C with an antibody directed against L1-CAM, an adhesion protein expressed at the cell surface and known to be recycled by trafficking vesicles (Dequidt et al., 2007). MemBright co-localized perfectly with L1-CAM at the cell surface (Figures 6A1–6A3). Live antibody uptake (Figures 6A4–6A6) showed that endocytic vesicles containing L1-CAM were also labeled with MemBright, indicating that a portion of MemBright can be internalized to monitor endocytosis. It is noteworthy that vesicles stained with MemBright are more numerous than vesicles containing L1-CAM, indicating that MemBright can be used to track various endocytic pathways. Additionally, a high amount of MemBright is still present at the PM surface, which can help in assessing membrane morphology and monitoring endocytosis in the same channel.

Using a PM sensor in conjunction with classical immunochemistry would avoid time-consuming transfection of GFP reporters that are usually not efficient and very difficult to handle in primary cells. As discussed previously, PFA fixation followed by permeabilization leads to leakage of the dye from the PM. Herein, after

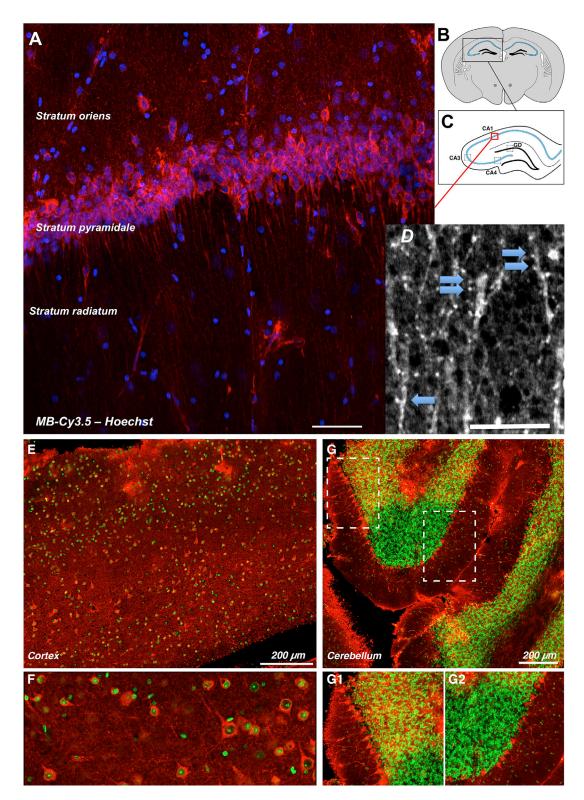


Figure 4. Neuronal Imaging in Brain Slices

(A–D) z projection (A) of two-photon imaging of live 1-mm-thick brain slices incubated with MB-Cy3.5 and Hoechst. Ninety images were acquired every 0.4 μm and were stacked providing a depth of 36 μm. The red box corresponds to the zoomed-in *stratum pyramidale* of CA1 region (C) of the hippocampus (B). Two-photon imaging allows the detection of some dendritic spine in *stratum radiatum* (D, arrows). Scale bars, 60 μm.
(E and F) Tile confocal imaging of 1-mm-thick live cortex incubated with MB-Cy3.5 (E) with strong labeling in cortical neurons (F).
(G) Confocal imaging of the cerebellum with zoom-in on Purkinje cells (G1 and G2).

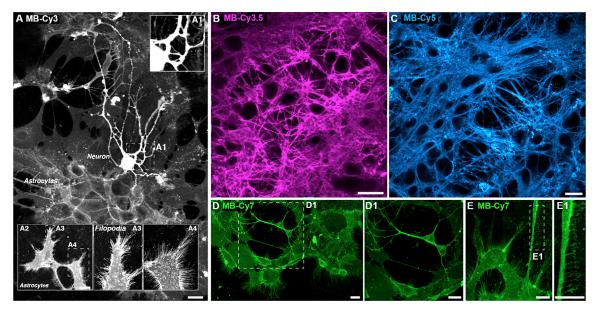


Figure 5. Confocal Microscopy of Hippocampal Primary Neurons and Astrocytes Labeled with MemBright Probes
Images were taken 10–15 min after addition of the probes (20 nM) without any washing step.

(A) Hippocampal neurons appear in a brighter manner than the astrocytes, the latter being either confluent (as shown in A) or isolated (A2). Zooms of the PM show how the MemBright allows detection of very fine membrane protrusion in astrocytes (filopodia in A3 or A4) or in neuronal cells (dendritic processes in A1).

(B and C) Live dense neuronal network is labeled with 20 nM MB-Cy3.5 (B) or 20 nM MB-Cy5 (C) and imaged with spinning-disc microscopy.

(D and E) Example of astrocytic layers and isolated neurons labeled with 200 nM MB-Cy7 excited with a 633-nm laser and imaged with laser confocal scanning microscopy. Even in the near-infrared region, membrane details can be unraveled with the probe both on neuronal (D1) and glial cells (see fine filopodia in E1). Scale bars, 20 μm.

optimizing the permeabilization process we used MemBright labeling in combination with primary VGlut antibody to unravel the glutamatergic synaptic vesicles within axonal presynaptic sites. As shown in Figure 6B, we succeeded in maintaining MemBright labeling after PFA-glutaraldehyde fixation, allowing the detection of VGlut synapses (arrows in Figures 6B1 and 6B2) contacting dendrites labeled with MemBright. Confocal section allowed the visualization of dendritic membrane from a top external view (in Figure 6B2) and from a cross-section revealing the dendritic lumen (Figure 6B1). 3D-view of a z stack led to a clear identification of the neuronal cell body and dendrites (in red) contacted by several glutamatergic synapses (in green).

As dendritic spines are known to be very dynamic tiny structures that change shape during memory (Kasai et al., 2010), we decided to tackle this detection with MemBright. The identification of the spine shape is usually done with phalloidin on fixed samples, with transgenic fluorescent mice, or with transfection on live neurons. Double labeling with fluorescently labeled phalloidin and MemBright on live samples showed that both dendritic spine head and neck can be efficiently labeled by MemBright with preserved staining after fixation (Figure 6C1–6C3).

#### Unraveling Synapse and Dendritic Spine Morphology Using Super-Resolution Imaging

Over the last decade, super-resolution fluorescence microscopy has pushed the diffraction limit in cell imaging (Huang et al., 2010). Among the available techniques, stochastic optical reconstruction microscopy (STORM) (Rust et al., 2006) is based

on successive activation, imaging, and high-precision localization of blinking fluorophores. Herein, MemBright probes were used at 20 nM to stain fixed HeLa cells that were imaged by total internal reflection fluorescence (TIRF) microscopy. Among the developed probes, MB-Cy3.5 was selected as the most efficient for direct STORM (dSTORM) imaging. After an initial bleaching phase, MB-Cy3.5 exhibited a stable blinking. dSTORM image of MB-Cy3.5 labeled PM clearly showed an increase of the resolution compared with the standard TIRF image (Figure S15). Indeed, the intensity plot profile of a cell filopodium evidenced a decrease of the thickness from 430 nm to 120 nm that corresponds to a 3.5-fold increase in the image resolution (Figure S15C).

We then assessed whether MBCy3.5 could reveal dendritic spine morphology on fixed samples in 3D STORM microscopy. First, we showed that MemBright efficiently labeled dendritic spine neck and head, allowing the identification of mushroom spines at either 9 (Figures 7A–7C and Video S6) or 14 days *in vitro* (Figure 7D). 3D imaging with color-coded depth allowed the visualization over several microns from back (Figure 7B) to front views (Figures 7B and 7D). As a comparison, we assessed the ability of our previously developed PM probe dSQ12S to provide dSTORM imaging in similar conditions. Although dSQ12S successfully stained the PM of neurons in wide-field microscopy, it rapidly photobleached (Figure S16) and did not blink sufficiently to provide dSTORM super-resolution images (Figure S17). MemBright was then used as a reference PM probe in multicolor STORM microscopy after fixation and permeabilization. As

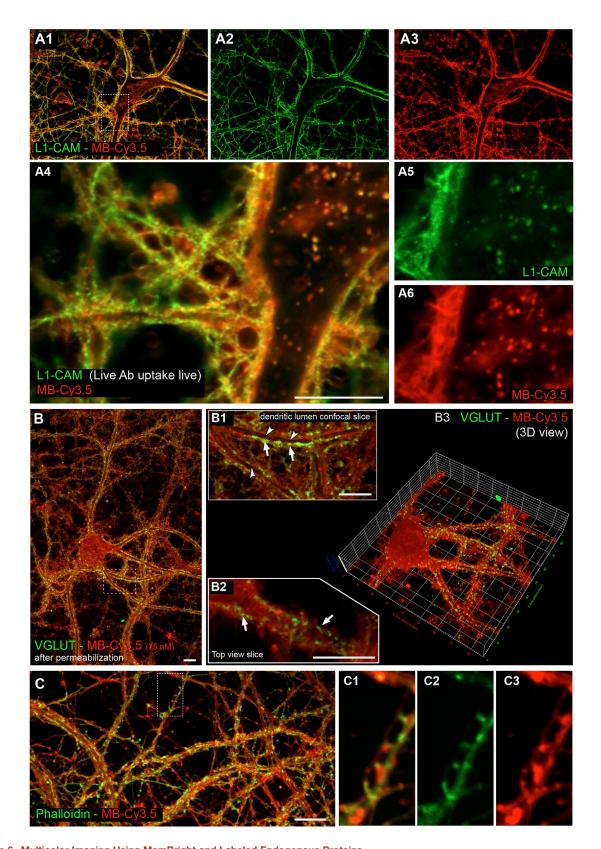


Figure 6. Multicolor Imaging Using MemBright and Labeled Endogenous Proteins (A) Confocal imaging of primary hippocampal neurons incubated with MB-Cy3.5 (red in A1-A3, A4, and A6) and L1-CAM monoclonal antibody (green in A1, A2, A4, and A5) for 10 min at 37°C. MB-Cy3.5 allows the identification of the PM both on the cell body and dendrites as shown with co-localization of the cell surface

shown in Figures 7E–7G, MemBright allowed visualization of both axonal (Figures 7E, 7G, and 7I) and dendritic (Figures 7E, 7F, and 7H) processes. Since MemBright stains axonal processes in a much brighter manner than dendrite, it allowed us to visualize axon coiling around a dendrite in the first thousands of pictures; thereafter, longer acquisition revealed the complete dendritic processes. Additional immunostaining of AMPA ( $\alpha$ -amino-3-hydroxy-5-methyl-4-isoxazole propionate) glutamate receptors (GluR2 subunit) revealed that glutamate receptors were accumulated at the axon-dendritic junctions. Altogether, these experiments showed that MemBright can be used as a reference membrane marker in both conventional and STORM microscopy on either live or fixed samples.

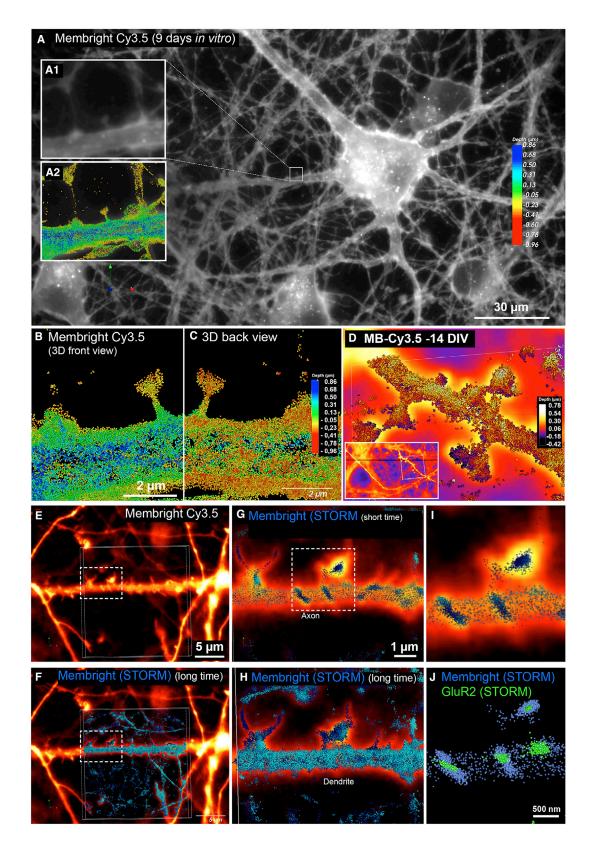
#### **DISCUSSION**

In this work, we developed a family of bright PM probes based on cyanines, so-called MemBright, of six different colors, and demonstrated its versatility for imaging the PM by several imaging techniques. Taking advantage of the symmetric structure of cyanines, they were modified with two amphiphilic zwitterionic anchors ensuring the selective and persistent staining of the PM. In aqueous media, these probes form non-fluorescent dark aggregates, whereas in the presence of lipid membranes these aggregates dissociate into highly fluorescent molecular species solubilized in the bilayer. This turn-on capability of MemBright probes enabled cellular imaging without any washing step with high signal-to-noise ratio in both live and fixed cells. A structure-property relationship was established as the most lipophilic probes (MB-Cy3.5, MB-Cy5.5, and MB-Cy7.5) tend to label the PM in a slower manner (15 min versus 5 min) but, in the meantime, provide longer-term staining of the PM. Owing to the high brightness of their cyanine fluorophore and efficient membrane staining, MemBright probes can be used at concentrations as small as 1 nM, which is >1,000-fold lower than those commonly used for long-chain cyanines, such as DiD, the PKH family, or mCLING. To obtain a sufficient signal, other probes are used at higher concentrations but can then become toxic for cells as shown for mCLING above 2 μM (Revelo and Rizzoli, 2001). Moreover, in confluent cells MemBright probes were found to stain the PM in a more homogeneous manner than the fluorescently labeled WGA, a protein-based membrane marker, because the former are small molecules that can diffuse rapidly within the lipid membrane and unlike WGA are not dependent on the heterogeneous expression of glycan at the cell surface. This property is a real asset for imaging PM in 3D cell cultures, tissues, and small animals, featuring tight packing of the cells. In addition, MemBright can be used to monitor endocytosis, since it can be internalized in endocytic vesicles (Figures 6A and S18). We were able to visualize both PM and endocytic recycling vesicles (Figure S18), which was hardly feasible with conventional FM dyes whose membranous staining does not resist washing steps. MemBright probes were also found to be efficient in TPE imaging, with two-photon cross-section reaching 1,800 GM for Cy5-based probe, and served to image deep PM in liver and brain tissues. In the field of neuroimaging, MemBright probes were successfully applied for imaging live and fixed neuronal networks, showing excellent imaging contrast of the PMs of the primary neurons together with astrocytes. This is an important advantage over the transfection method, which is usually deleterious for the cells and allows detection of only 5%-10% of the neurons within a population. Previously, this has prompted researchers to use transgenic reporter mice, such as THY1-fluorescent transgenic lines (Porrero et al., 2010). However, this method is almost completely reserved for the mouse species since transgenic rat, marmoset, or other mammal animal models are still rare. The MemBright probes described herein are usable whatever the species and eventually on human samples from surgical resections. Moreover, due to the wide range of available colors, MemBright can also be used in combination with green or red transgenic mice using complementary red, deep-red, or near-infrared MemBright. Other alternatives used to label neurons in brain tissue are biocytin injection (Horikawa and Armstrong, 1988) and Dil, DiO, and DiD tissue-labeling pastes or crystals (Honig and Hume, 1986). Whereas the injection needs a fluorescent streptavidin application to reveal the neurons, DiO relies on insoluble dyes in water, which can be problematic for application (Lanciego and Wouterlood, 2011). Indeed those products are usually placed via a needle into the brain area or by using a gene gun. This usually results in a non-homogenous dispersion of the dye, leading to a very bright local fluorescence. MemBright probes are water-dispersible and thus can be used either incubated on cultured cells or directly on slices, which does not require any special skills. Moreover, they can be visualized directly without any additional revealing steps thanks to their intrinsic fluorescence. Finally, MemBright probes combine the capability of the Neurotrace dye (Thermo Fisher) for neuronal soma and Fluoromyeline dye (Thermo Fisher) for labeling myelin in spinal cord (Chevalier and Rosenberger, 2017). Indeed, MemBright stains well the cell soma as well as the dendrites (Figures 5 and 7), axons (Figures 7 and 7), and commissural and association fibers (corpus callosum, fimbria; data not shown). Very interestingly, it appeared that MemBright markers preferentially stain the PM of neurons with a higher intensity compared with surrounding cells present in the brain (e.g., glial cells). This tendency allowed imaging of neurons in various regions of the brain tissue with high signal-to-background ratio. Additionally, MemBright enables high-contrast imaging of motor neurons, which opens perspectives in the important field of nerve-specific fluorescence-guided surgery (Whitney et al., 2011).

L1-CAM adhesion molecule in A1. Internalized vesicles can be tracked with L1-CAM antibody (A5) or with MB-Cy3.5 (A6) for a wider identification of different endocytic pathways. Scale bar: 10 µm.

<sup>(</sup>B) Confocal imaging of primary hippocampal neurons incubated with MB-Cy3.5 (B1-B3) for 10 min at 37°C and then fixed, permeabilized, and incubated with polyclonal antibody to VGlut (green). MB-Cy3.5 allows the identification of the PM both on confocal section passing either through the dendrite (B1) or below the dendrite (B2), which can be visualized in 3D thanks to stack reconstruction (B3). Presynaptic boutons labeled with VGlut (arrows) can be seen in contact on the dendritic membrane. Even the dark lumen of dendrite (arrowheads) can be visualized. Scale bar: 10 μm.

<sup>(</sup>C) Confocal imaging of primary hippocampal neurons incubated with MB-Cy3.5 (red) and phalloidin A488 (green) showing that dendritic spine head/neck can be seen with both dyes. Scale bar: 10 µm.



(legend on next page)

Finally, we showed that MemBright Cy3.5, owing to its capacity to blink, enables STORM-type SRM. To our knowledge, MemBright are the first PM probes operating at nanomolar concentrations in live or fixed samples using conventional confocal, two-photon as well as super-resolution (STORM) microscopy. Since MemBright is compatible with cell fixation and permeabilization, it can be used in combination with immunofluorescence and open a wide avenue to multicolor imaging at nanoscopic resolution. Several teams are currently trying to decipher the nanoarchitecture of molecular complexes with STORM microscopy (Dani et al., 2010; Herrmannsdörfer et al., 2017; Beghin et al., 2017; Lagache et al., 2018; Levet et al., 2015). It is thus crucial to achieve a PM staining that can precisely localize these proteins in their PM environment in order to map their cellular distribution. Additionally, this is of paramount importance in the context of 3D reconstruction. Most of actual studies lack membrane staining as reference and display dotted lines indicating the putative location of cell limits. We provide here a blinking membrane probe that is also resistant enough to permeabilization and to reducing buffer to allow such a use in multicolor STORM immunostaining. We showed the nanoscale organization of both axonal and dendritic compartments in combination with endogenous glutamate receptor clusters in multicolor 3D STORM. Glutamate receptors are clearly aggregated at the contact site between axons and dendrites as well as at extrasynaptic sites. This membrane reference thus permits the fine molecular mapping of endogenous proteins at the nanoscale level within the cellular context that was, up to now, only possible with electron microscopy. In conclusion, due to the availability of six different colors within the spectral range of 550-850 nm, combined with ease of use, brightness, and compatibility with different microscopy techniques, the MemBright family is potentially a powerful toolbox for biomembrane imaging in cell biology and neuroscience.

#### SIGNIFICANCE

Because the plasma membrane (PM) delimits cells and controls many important biological events, visualizing it allows us to study its role in cellular processes. The commonly used PM probes can suffer from limitations such as low selectivity, heterogeneous staining, and fast internalization. We herein proposed a family of selective and fluorogenic PM probes that we called MemBright. These are available in various colors from orange to near infrared, allowing multicolor imaging with complementary fluorescent probes and immunostaining. These probes are compatible with live cells and fixed samples as well as tissues. Additionally, we demonstrated that they are compatible with different fluorescence imaging techniques

including long-term imaging (video microscopy), twophoton excitation, and super-resolution imaging. MemBright was shown to be particularly efficient for neuroscience as it stains neurons in a brighter manner than other cells and thus reveals neurons in brain slices. This feature also permitted us to unravel nerves in muscle tissue samples, which could be a major asset in fluorescence-guided surgery applications. We believe that MemBright is a versatile and efficient tool that could become commonly used in routine bioimaging to image the PM, and could serve in more specialized experiments requiring advanced fluorescence imaging.

#### **STAR**\*METHODS

Detailed methods are provided in the online version of this paper and include the following:

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#### SUPPLEMENTAL INFORMATION

Supplemental Information includes 18 figures, 2 tables, 6 videos, and 1 data file and can be found with this article online at https://doi.org/10.1016/j.chembiol.2019.01.009.

#### Figure 7. STORM Microscopy of Neurons Using MemBright Cy3.5

Imaging of hippocampal neurons labeled with MemBright after 9 (A–C) or 14 (D) days *in vitro*. Spines can be identified in wide-field microscopy (A1) and then super-resolved using STORM (A2, B, and C). 3D STORM microscopy allowed visualization of the spine from the front (B) or back view (C). See Video S6 for animated views. MemBright Cy3.5 allows visualization of the PM through several microns thanks to biplane module; depth is color-coded in (B) to (D) (rainbow look-up table). Dendritic and axonal profiles can be visualized in wide-field microscopy (E). Axonal processes coil around the dendrite (dashed box in E, magnified in G and I) and can be reconstructed in the first 3D-STORM images due to intense blinking. After several hundred pictures, the dendritic shape can be reconstructed (F; magnified in H). Thanks to permeabilization resistance, MemBright (blue) can be used in dual-color immunofluorescence to visualize glutamate AMPA receptor clusters (green) aggregated at the dendrite in front of axonal profile (J) with interleaved 3D STORM stack.

A video abstract is available at https://doi.org/10.1016/j.chembiol.2019.01. 009#mmc8.

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#### **AUTHOR CONTRIBUTIONS**

M.C. and A.S.K. planned the project. M.C. synthesized, purified, and characterized the MemBright and led the spectroscopy experiments with P.A. and A.S.K. P.A. studied the spectroscopic and two-photon properties. Cell imaging was performed by M.C., P.A., H.A. (HeLa dSTORM) and L.D. Two-photon tissue imaging was realized by O.F. and L.D. E.B. led the cytotoxicity evaluation. L.D. led the tissue and neuronal imaging as well as the super-resolution imaging. Y.M., T.G., and A.S.K. contributed materials and analysis tools. M.C., L.D., and A.S.K. wrote the manuscript with some assistance from other co-authors.

#### **DECLARATION OF INTERESTS**

The authors declare no competing interests.

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### **STAR**\***METHODS**

#### **KEY RESOURCES TABLE**

REAGENT or RESOURCE	SOURCE	IDENTIFIER
Antibodies		
L1 CAM	Rabbit antibodies against the L1-CAM extra-cellulardomain(1/2000;	a gift fromF.Rathjen,Max-Delbrueck- CenterforMolecularMedicine,Berlin,Germany
VGLUT	Covalab	Rabbit anti VGLUT1&2 antibody (#pab0047, 200e)
Secondary Ab	Donkey anti rabbit Alexa 488 Highly crossed absorbed	https://www.thermofisher.com/antibody/product/Donkey- anti-Rabbit-IgG-H-L-Highly-Cross-Adsorbed-Secondary- Antibody-Polyclonal/A-21206; RRID: AB_2535792
	Donkey anti mouse Alexa 647 Highly crossed absorbed	https://www.thermofisher.com/antibody/product/Donkey- anti-Mouse-IgG-H-L-Highly-Cross-Adsorbed-Secondary- Antibody-Polyclonal/A-31571; RRID: AB_162542
GluR2	BD Pharmingen 556341	http://www.bdbiosciences.com/eu/reagents/research/ antibodies-buffers/cell-biology-reagents/cell-biology- antibodies/purified-mouse-anti-glutamate-receptor-6c4/p/ 556341; RRID: AB_396373
Biological Samples		
Rat Liver	Janvier Lab	Sprague Dawley Rats
Mouse brain tissue	Janvier Lab	C57BL6/N strain
Mouse liver tissue	Janvier Lab	C57BL6/N strain
Muscle	Janvier Lab	C57BL6/N strain
Chemicals, Peptides, and Recomb	inant Proteins	
WGA-488	ThermoFisher Scientific	http://www.thermofisher.com/order/catalog/ product/W11261
WGA-647	ThermoFisher Scientific	http://www.thermofisher.com/order/catalog/ product/W32466
Phalloïdine marquée	ThermoFisher Scientific	http://www.thermofisher.com/order/catalog/product/ A12379; RRID: AB_2315147
DID	ThermoFisher Scientific	http://www.thermofisher.com/order/catalog/ product/D7757
α-bungarotoxin	ThermoFisher Scientific	http://www.thermofisher.com/order/catalog/ product/B13423
mCLING-ATTO 647N-labeled	Synaptic Systems	#710 006AT1
Hoechst 33342	ThermoFisher Scientific	http://www.thermofisher.com/order/catalog/ product/H1399
Cysteamine hydrochloride	Sigma Aldrich	30078
Glucose Oxidase from Aspergillus niger	Sigma Aldrich	G0543
Catalase from bovine liver	Sigma Aldrich	C3155
Experimental Models: Cell Lines		
HeLa cells		ATCC® CCL-2
KB cells		ATCC® CCL-17
Primary neurons	(From mouse see below)	
Experimental Models: Organisms/S	Strains	
Mouse	Janvier Lab	C57BL6/N strain
Software and Algorithms		
Chemdraw 13.0	PerkinElmer Informatics	http://www.cambridgesoft.com/software/ overview.aspx
Origin 9.1	OriginLab	https://www.originlab.com/

(Continued on next page)

Continued		
REAGENT or RESOURCE	SOURCE	IDENTIFIER
Illustrator CS5	Adobe	https://www.adobe.com
lcy	bioimageanalysis	http://icy.bioimageanalysis.org/
Srx software from Bruker (STORM)	Bruker	https://www.bruker.com/products/fluorescence- microscopes/vutara-super-resolution-microscopy/ overview/srx-software-vutara-super-resolution.html

#### **CONTACT FOR REAGENT AND RESOURCE SHARING**

Further information and requests for resources and reagents should be directed to and will be fulfilled by the Lead Contact, Mayeul Collot (mayeul.collot@unistra.fr).

#### **EXPERIMENTAL MODEL AND SUBJECT DETAILS**

HeLa cells (ATCC® CCL-2, female) were grown in Dulbecco's modified Eagle medium (DMEM, Gibco-Invitrogen), supplemented with 10% fetal bovine serum (FBS, Lonza) and 1% antibiotic solution (penicillin-streptomycin, Gibco-Invitrogen) at 37°C in humidified atmosphere containing 5% CO<sub>2</sub>. KB cells (ATCC® CCL-17, female) were grown in minimum essential medium (MEM, Gibco-Invitrogen) with 10% fetal bovine serum (FBS, Lonza), 1% non-essential amino acids (Gibco-Invitrogen), 1% MEM vitamin solution (Gibco-Invitrogen), 1% L-Glutamine (Sigma Aldrich) and 0.1% antibiotic solution (gentamicin, Sigma-Aldrich) at 37°C in humidified atmosphere containing 5% CO<sub>2</sub>. Cells were seeded onto a chambered coverglass (IBiDi®) at a density of 1×10<sup>5</sup> cells/well 24h before the microscopy measurement. All experiments on mice were performed in accordance with European Community guidelines legislation and reviewed by the local ethical committee of the Paris Diderot University. Animals were maintained on a 12 hour light-dark cycle with ad libitum access to food and water. Hippocampal cultures from 18-day-old male and female Sprague-Dawley rat embryos were pooled and prepared as described previously (Danglot et al., 2003) (Banker and Cowan, 1977).

#### **METHOD DETAILS**

#### **Chemical Synthesis**

All starting materials for synthesis were purchased from Alfa Aesar, Sigma Aldrich or TCI Europe and used as received unless stated otherwise. NMR spectra were recorded on a Bruker Avance III 400 MHz spectrometer. Mass spectra were obtained using an Agilent Q-TOF 6520 mass spectrometer. Synthesis and characterization of DAF is described in Supplemental Information. NMR and mass spectra are available in the Supplemental Information (Data S1). The synthesis of the clickable amphiphilic zwitterion (CAZ) was described in ref (Collot et al, 2015).

#### General Procedure for the Synthesis of Cy3- diAlkyne- and Cy3.5- diAlkyne

The synthesis of Indoleninium 1 and benzoindoleninium 2 were previously described. (Gerowska et al., 2012) To a solution of indoleninium 1 or benzoindoleninium 2 (1 mmol) in pyridine (5 mL) was added triethyl orthoformate (1.5 mmol, 1.5 eq). The solution was allowed to stir at 100°C for 3 hours. The solvents were evaporated and the product was extracted with DCM, washed with an aqueous solution of HClO<sub>4</sub> (1M) before being dried over MgSO<sub>4</sub>. The solution was filtered and evaporated. The crude was purified by column chromatography on silica gel (DCM/MeOH: 99/1 to 95/5).

#### Cy-3-diAlkyne

255 mg (91%), dark pink solid. Rf=0.42 (DCM/MeOH: 95/5).  $^{1}$ H-NMR (400 MHz, CDCl<sub>3</sub>):  $\delta$  8.43 (t, J = 13.5 Hz, 1H, CH), 7.45-7.38 (m, 4H, H Ar), 7.31-7.26 (m, 4H, H Ar), 6.91 (d, J = 13.5 Hz, 2H, 2 CH), 4.32 (t, J = 7.8 Hz, 4H, CH<sub>2</sub>N), 2.53 (dt, J = 6.5, 3.3 Hz, 4H, 2 CH<sub>2</sub>), 2.12-2.08 (m, 6H, 2 CH<sub>2</sub> 2CH alkyne), 1.75 (s, 12H, 4 CH<sub>3</sub>).  $^{13}$ C NMR (126 MHz; CDCl<sub>3</sub>):  $\delta$  174.0 (C=N), 151.0, 142.2, 140.5, 129.0, 125.4, 122.0, 111.1, 104.4, 83.6 (C alkyne), 69.6 (CH alkyne), 49.0 (2 CH<sub>2</sub>), 43.3 (Cq), 28.2 (4 CH<sub>3</sub>), 26.0 (2 CH<sub>2</sub>), 15.8 (2 CH<sub>2</sub>). HRMS (ES<sup>+</sup>), calcd for C<sub>33</sub>H<sub>37</sub>N<sub>2</sub> [M]<sup>+</sup> 461.2951, found 461.2950.

#### Cy-3.5-diAlkyne

238 mg (72%), dark violet solid. Rf=0.32 (DCM/MeOH: 95/5).  $^{1}$ H-NMR (400 MHz, CDCl3):  $\delta$  8.67 (t, J = 13.5 Hz, 1H, CH), 8.14 (d, J = 8.5 Hz, 2H, H Ar), 8.00-7.98 (m, 4H, H Ar), 7.66 (m, J = 8.3, 7.0, 1.1 Hz, 2H, H Ar), 7.58-7.50 (m, 4H, H Ar), 6.97 (d, J = 13.6 Hz, 2H, CH), 4.45 (t, J = 7.8 Hz, 4H, CH<sub>2</sub>N), 2.58 (td, J = 6.5, 2.5 Hz, 4H, 2 CH<sub>2</sub>), 2.19-2.13 (m, 6H, 2 CH<sub>2</sub> 2CH alkyne), 2.09 (s, 12H, 4 CH<sub>3</sub>).  $^{13}$ C NMR (126 MHz; CDCl3):  $\delta$  175.4 (C=N), 149.5, 139.7, 133.3, 132.0, 130.9, 130.3, 128.0, 127.8, 125.1, 121.8, 110.9, 103.8, 83.7 (C alkyne), 69.7 (CH alkyne), 50.7 (2 CH<sub>2</sub>), 43.4 (Cq), 27.8 (4 CH<sub>3</sub>), 26.3 (2 CH<sub>2</sub>), 15.8 (2 CH<sub>2</sub>). HRMS (ES<sup>+</sup>), calcd for C<sub>41</sub>H<sub>41</sub>N<sub>2</sub> [M]<sup>+</sup> 561.3264, found 561.3275.

#### General Procedure for the Synthesis of Cy5- diAlkyne- and Cy5.5- diAlkyne

To a solution of indoleninium **1** or benzoindoleninium **2** (1 mmol, 2 eq) in AcOH (7 mL) was added malonaldehyde dianilide hydrochloride (0.5 mmol, 1 eq) followed by AcONa. The solution was allowed to stir at 50°C for 3 hours. The solvents were evaporated and the product was extracted with DCM, washed with an aqueous solution of HClO<sub>4</sub> (1M) before being dried over MgSO<sub>4</sub>. The solution was filtered and evaporated. The crude was purified by column chromatography on silica gel (DCM/MeOH: 99/1 to 95/5).

#### Cy-5-diAlkyne

254 mg (86%), blue solid. Rf=0.46 (DCM/MeOH: 95/5).  $^1$ H-NMR (400 MHz, CDCl<sub>3</sub>):  $\delta$  8.12 (t, J = 12.8 Hz, 2H, 2 CH), 7.38 (d, J = 6.8 Hz, 4H, H Ar), 7.30-7.19 (m, 4H, H Ar), 7.07 (t, J = 11.9 Hz, 1H, CH), 6.63 (d, J = 13.3 Hz, 2H, 2 CH), 4.30 (s, 4H, CH<sub>2</sub>N), 2.47 (s, 4H, 2 CH<sub>2</sub>), 2.29 (s, 2H, C≡CH), 2.07 (s, 4H, 2 CH<sub>2</sub>), 1.78 (s, 12H, 4 CH<sub>3</sub>).  $^{13}$ C NMR (101 MHz; CDCl<sub>3</sub>):  $\delta$  173.0 (C=N), 153.8, 142.1, 141.2, 128.6, 127.0, 125.1, 122.2, 110.5, 104.3, 82.7 (C alkyne), 70.6 (CH alkyne), 49.3 (2 CH<sub>2</sub>), 43.1 (Cq), 28.2 (4 CH<sub>3</sub>), 26.0 (2 CH<sub>2</sub>), 16.2 (2 CH<sub>2</sub>). HRMS (ES<sup>+</sup>), calcd for C<sub>35</sub>H<sub>39</sub>N<sub>2</sub> [M]<sup>+</sup> 487.3108, found 487.3116.

#### Cy-5.5-diAlkyne

125 mg (36%), Bordeaux red solid. Rf=0.43 (DCM/MeOH: 95/5).  $^{1}$ H-NMR (400 MHz, CDCl<sub>3</sub>):  $\delta$  8.26-8.18 (m, 4H, H Ar), 7.92 (q, J = 6.2 Hz, 4H, H Ar), 7.63-7.59 (m, 2H, 2 CH), 7.46 (m, 4H, H Ar), 6.82-6.76 (m, 1H, CH), 6.42-6.39 (m, 2H, 2 CH), 4.33-4.29 (m, 4H, CH<sub>2</sub>N), 2.44-2.43 (m, 4H, 2 CH<sub>2</sub>), 2.27 (s, 2H, C≡CH), 2.07 (s, 16H, 4 CH<sub>3</sub>, 2 CH<sub>2</sub>).  $^{13}$ C NMR (101 MHz; CDCl<sub>3</sub>):  $\delta$  174.6 (C=N), 153.2, 139.3, 134.2, 131.8, 130.6, 129.9, 128.2, 127.8, 126.4, 125.1, 122.4, 110.4, 82.8 (C alkyne), 70.6 (CH alkyne), 51.2 (2 CH<sub>2</sub>), 42.9 (Cq), 27.7 (4 CH<sub>3</sub>), 26.3 (2 CH<sub>2</sub>), 16.0 (2 CH<sub>2</sub>). HRMS (ES<sup>+</sup>), calcd for C<sub>43</sub>H<sub>43</sub>N<sub>2</sub> [M]<sup>+</sup> 587.3421, found 587.3435.

#### General Procedure for the Synthesis of Cy7- diAlkyne- and Cy7.5- diAlkyne

To a solution of indoleninium 1 or benzoindoleninium 2 (1 mmol, 2 eq) in AcOH (7 mL) was added Glutaconaldehydedianil Hydrochloride (0.5 mmol, 1 eq) followed by AcONa. The solution was allowed to stir at 50°C for 3 hours. The solvents were evaporated and the product was extracted with DCM, washed with an aqueous solution of HClO<sub>4</sub> (1M) before being dried over MgSO<sub>4</sub>. The solution was filtered and evaporated. The crude was purified by column chromatography on silica gel (DCM/MeOH: 99/1 to 95/5).

#### Cy-7-diAlkyne

230 mg (75%), Shiny red solid. Rf=0.47 (DCM/MeOH: 95/5). <sup>1</sup>H-NMR (500 MHz, CDCl<sub>3</sub>): δ 7.92-7.89 (m, 2H, 2 CH), 7.70 (s, 1H, CH), 7.39-7.35 (m, 4H, H Ar), 7.22 (t, J = 7.4 Hz, 2H, 2 CH), 7.17 (d, J = 7.9 Hz, 2H, H Ar), 6.65-6.60 (m, 2H, 2 CH), 6.28-6.25 (m, 2H, 2 CH), 4.17-4.14 (m, 4H,  $CH_2N$ ), 2.41 (dt, J = 6.3, 3.2 Hz, 4H, 2  $CH_2$ ), 2.18 (s, 2H,  $C \equiv CH$ ), 2.05 (t, J = 7.1 Hz, 4H, 2  $CH_2$ ), 1.71 (s, 12H, 4H, CH<sub>3</sub>). <sup>13</sup>C NMR (126 MHz; CDCl<sub>3</sub>): δ 171.7, 152.2, 142.2, 141.1, 128.6, 126.7, 124.8, 122.3, 110.2, 103.8, 82.7 (C alkyne), 70.2 (CH alkyne), 49.1 (2 CH<sub>2</sub>), 42.8 (Cq), 27.9 (4 CH<sub>3</sub>), 25.9 (2 CH<sub>2</sub>), 16.0 (2 CH<sub>2</sub>). HRMS (ES<sup>+</sup>), calcd for C<sub>37</sub>H<sub>41</sub>N<sub>2</sub> [M]<sup>+</sup> 513.3264.3421, found 513.3261.

#### Cy-7.5-diAlkyne

94 mg (26%) dark green solid. Rf=0.29 (DCM/MeOH: 95/5). <sup>1</sup>H-NMR (400 MHz, CDCl<sub>3</sub>): δ 8.12 (d, J = 8.5 Hz, 2H), 8.01 (dd, J = 12.7, 0.4 Hz, 2H, 2 CH), 7.91 (t, J = 9.0 Hz, 4H), 7.81 (broad s, 1H, CH), 7.59-7.55 (m, 2H), 7.48-7.41 (m, 4H), 6.64 (dd, J = 12.0, 0.5 Hz, 2H, 2H), 2H, 3H, 4H, CH), 6.28-6.25 (m, 2H, 2 CH), 4.27-4.24 (m, 4H, CH<sub>2</sub>N), 2.42 (dt, J = 6.2, 3.2 Hz, 4H, 2CH<sub>2</sub>), 2.19 (t, J = 2.4 Hz, 2H, C  $\equiv$  CH), 2.07 (t, J = 2.6.9 Hz, 4H, 2CH<sub>2</sub>), 1.99 (s, 12H, 4 CH<sub>3</sub>). <sup>13</sup>C NMR (101 MHz; CDCl3): δ 173.1 (C=N), 151.30, 139.5, 133.7, 131.7, 130.6, 130.01, 129.97, 129.82, 128.7, 128.2, 127.7, 126.52, 126.50, 126.48, 124.9, 122.3, 110.5, 103.28, 82.9 (C alkyne), 70.4 (CH alkyne), 51.0 (2 CH<sub>2</sub>), 42.9 (Cq), 27.5 (4 CH<sub>3</sub>), 26.3 (2 CH<sub>2</sub>), 16.0 (2 CH<sub>2</sub>). HRMS failed in providing the correct mass.

#### Typical Procedure for the Synthesis of the MemBright Dyes (Click Coupling)

For scheme see Figure 1. To a solution of dialkyne cyanine (18 mg) and CAZ (2.4 eq) in DMF (3 mL) was added an heterogeneous aqueous solution (300 μL) of CuSO<sub>4</sub>.5H<sub>2</sub>O (5 mg) and sodium ascorbate (5 mg). The solution was allowed to stir at 60°C for 1 hour. The solvents were evaporated. The crude was dissolved in a DCM/MeOH (1/1) mixture before being filtered through celite. The product was purified by exclusion phase column chromatography. The pure product was then dissolved in water/dioxane mixture and was lyophilised.

Due to their amphiphilic nature and the low amount of material, NMR spectra of MemBright® markers could not be accurately described. <sup>1</sup>H NMR spectra are available in the Supplemental Information.

38 mg (86%). <sup>1</sup>H-NMR (400 MHz, MeOD):  $\delta$  8.41-8.40 (m), 8.21 (broad s), 7.43-7.34 (m), 7.28-7.25 (m), 6.53 (d), 4.56 (broad s), 4.19 (broad s), 3.95 (broad s), 3.55 (broad s), 3.33 (broad s), 3.19 (broad s), 2.96 (broad d), 2.45 (broad s), 2.37 (broad), 2.09 (broad s), 1.70 (s, 12H, 4 CH<sub>3</sub>), 1.21 (s, 36H, CH<sub>2</sub> C12), 0.84 (s, 6H, 2 CH<sub>3</sub> C12). HRMS (ES<sup>+</sup>), calcd for C<sub>71</sub>H<sub>117</sub>N<sub>10</sub>O<sub>6</sub>S<sub>2</sub><sup>+</sup> [M]<sup>+</sup> 1269.8593, found 1269.8611.

#### MB-Cy3.5

26 mg (65%). Broad NMR signals, see NMR spectrum. HRMS (ES<sup>+</sup>), calcd for C<sub>79</sub>H<sub>121</sub>N<sub>10</sub>O<sub>6</sub>S<sub>2</sub><sup>+</sup> [M]<sup>+</sup> 1369.8906, found 1369.8893.

16 mg (37%). <sup>1</sup>H-NMR (400 MHz, MeOD): δ 8.10, 7.44, 7.29, 6.81, 6.37, 4.62 (broad s), 4.20 (broad s), 3.48 (broad s), 3.28 (broad s), 3.06 (broad s), 2.87 (s, 3H), 2.46 (s, 3H), 2.18 (broad s), 1.72 (s, 12H, 4 CH<sub>3</sub>), 1.28 (m, 32H, CH<sub>2</sub> C12), 0.88 (s, 6H, 2 CH<sub>3</sub> C12). HRMS (ES<sup>+</sup>), calcd for  $C_{73}H_{119}N_{10}O_6S_2^+$  [M]<sup>+</sup> 1295.8750, found 1295.8772.

#### MB-Cy5.5

25 mg (63%).  $^{1}$ H-NMR (400 MHz, MeOD):  $\delta$  8.16, 7.98, 7.64, 7.59-7.55, 7.49, 6.64, 4.38, 3.55, 3.02, 2.49, 2.07 (s, 12H, 4 CH<sub>3</sub>), 1.65, 1.25 (m, 32H, CH<sub>2</sub> C12), 0.85 (s, 6H, 2 CH<sub>3</sub> C12). HRMS (ES<sup>+</sup>), calcd for  $C_{81}H_{123}N_{10}O_6S_2^+$  [M]<sup>+</sup> 1395.9063, found 1395.9059.

#### MR-Cv7

26 mg (62%). H-NMR (400 MHz, MeOD):  $\delta$  7.81, 7.41, 7.24, 6.46, 4.18, 3.54, 3.34, 3.23 (s, 4H), 3.04, 2.92, 2.46 (s, 3H), 2.29 (s, 2H), 2.13 (s, 4H), 1.69 (s, 12H, 4 CH<sub>3</sub>), 1.27 (m, 32H, CH<sub>2</sub> C12), 0.86 (s, 6H, 2 CH<sub>3</sub> C12). HRMS (ES<sup>+</sup>), calcd for  $C_{75}H_{121}N_{10}O_6S_2^+$  [M]<sup>+</sup> 1321.8906, found 1321.8916.

#### MB-Cy7.5

18 mg (47%). Broad signals, see NMR spectrum. HRMS (ES<sup>+</sup>), calcd for C<sub>83</sub>H<sub>125</sub>N<sub>10</sub>O<sub>6</sub>S<sub>2</sub><sup>+</sup> [M]<sup>+</sup> 1421.9219, found 1421.9228.

#### **Lipid Vesicles**

Dioleoylphosphatidylcholine (DOPC) and cholesterol were purchased from Sigma-Aldrich. Large unilamellar vesicles (LUVs) were obtained by the extrusion method as previously described (Hope et al., 1985). Briefly, a suspension of multilamellar vesicles was extruded by using a Lipex Biomembranes extruder (Vancouver, Canada). The size of the filters was first 0.2  $\mu$ m (7 passages) and thereafter 0.1  $\mu$ m (10 passages). This generates monodisperse LUVs with a mean diameter of 0.11  $\mu$ m as measured with a Malvern Zetasizer Nano ZSP (Malvern, U.K.). LUVs were labelled by adding 5  $\mu$ L of probe stock solution in dimethyl sulfoxide to 1-mL solutions of vesicles. A 20 mM phosphate buffer, pH 7.4, was used in these experiments. Molar ratios of probes to lipids were generally 1 to 500-1000.

#### **Spectroscopy**

Absorption spectra were recorded on a Cary 4000 spectrophotometer (Varian) and fluorescence spectra on a Fluoromax 4 (Jobin Yvon, Horiba) spectrofluorometer. Fluorescence emission spectra were systematically recorded at room temperature, unless indicated. All the spectra were corrected from wavelength-dependent response of the detector. The fluorescence and absorption spectra of the corresponding blank suspension of lipid vesicles without the probe was subtracted from these spectra. Concentrations of dyes were 400 nM for MB-Cy3 and MB-Cy3.5, 280 nM for MB-Cy5 and MB-Cy5.5, 220 nM for MB-Cy7 and 600 nM for MB-Cy7.5. Quantum yields were determined by comparison with a reference according to their excitation and emission wavelengths: Rhodamine 101 in EtOH (Rurack and Spieles, 2011) was used as the reference for MB-Cy3, Cresyl Violet in EtOH (Rurack and Spieles, 2011) for MB-Cy3.5, DID in MeOH (Texier et al., 2009) for MB-Cy5, Rhodamine 800 in EtOH (Alessi et al., 2013) for MB-Cy5.5 and Indocyanine green in MeOH(Benson and Kues, 1978) for MB-Cy7 and MB-Cy7.5. The liposomes were used at 200 μM (based on lipid) in phosphate buffer 20 mM, pH 7.4. The full-corrected emission spectra beyond 850 nm were obtained by using the GaussAmp function fit.

#### **DLS Measurements**

The sizes of the particles formed by MemBright in water were measured by Dynamic Light Scattering (DLS) using a Malvern Zetasizer Nano ZSP (Malvern, U.K.). Concentration of the dye was  $5 \mu M$ . The size of particles formed by MB-Cy5 and MB-Cy5.5 could not be measured due to the wavelength of the DLS laser (633 nm).

#### **Cellular Imaging**

For a nuclear staining, the medium was replaced by Hoechst 33342 (Thermo Fisher, 5 μg/mL) in Opti-MEM (Gibco-Invitrogen) and the cells were incubated for 10 minutes at 37°C. For imaging, the medium was removed and the attached cells were washed with HBSS (Gibco-Invitrogen) three times. Then, a freshly prepared solution of MemBright in HBSS (typically 20 nM) was guickly added to the cells without any washing step. Prior to imaging, PM was co-stained by addition of wheat germ agglutinin-Alexa488, WGA-AlexaFluor®488 (Thermo Fisher Scientific, 1 mg/mL in water) at a final concentration of 5 µg/mL. Confocal microscopy experiments were performed by using a Leica TCS SPE-II with HXC PL APO 63x/1.40 OIL CS objective. The microscope settings were: 405 nm laser for excitation of Hoechst 33342, emission was collected between 420 and 470 nm; 488 nm laser for excitation of WGA-AlexaFluor®488, emission was collected between 500 and 550 nm. For excitation of MB-Cy3 and MB-Cy3.5, 561 nm laser was used with 567-750 nm detection range. For excitation of MB-Cy5, MB-Cy5.5 and MB-Cy7, 635 nm laser was used with 640-800 nm detection range. The images were processed with the ImageJ software. In the case of fixed cells 1) fixation followed by staining. The HeLa cells seeded onto the 1 mL Ibidi® Chamber were washed once with D-PBSx1 (Lonza) and fixed in 4% paraformaldehyde (PFA - Electron Microscopy Sciences) at room temperature during 10 min. The fixed cells were washed three times with D-PBS to remove the excess of PFA before usual staining and imaging. For an efficient staining on fixed cells, 2) staining followed by fixation. The cells seeded onto the 1 mL Ibidi® Chamber were stained with 20 nM of MemBright for 10-15 minutes at room temperature, the excess of MemBright was washed with HBSS and the cells were fixed with 4% paraformaldehyde in HBSS for 10 minutes at room temperature before being washed 2 times with HBSS.

#### **Neuromuscular Junction Preparation**

Dissected hindlimb tibialis anterior (TA) muscles from adult C57BL6 mice were fixed (4% PFA in PBS) for 20 minutes at room temperature and further incubated overnight at 4°C with MB-Cy5 (200 nM), Hoechst and alpha-bungarotoxin-Alexa594. After three washed in PBS, mounted under glass in Vectashield (Vector Labs, Burlingame, CA). Incubation of MB-Cy5 for 30 min on non-fixed preparations also gave good results.

#### **Cytotoxicity Assay**

Cytotoxicity assay of the MemBright dyes was quantified by the MTT assay (3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide). A total of 1x10<sup>4</sup> KB cells/well were seeded in a 96-well plate 24 h prior to the cytotoxicity assay in Dulbecco's Modified Eagle Medium (Gibco Lifetechnologies -DMEM) complemented with 10% fetal bovine serum, Penicilin (100 Ul/mL), Streptomycin (100 μg/mL), L-Glutamine (2 mM) and were incubated in a 5% CO<sub>2</sub> incubator at 37°C. After medium removal, an amount of 100 μL DMEM containing 1000 nM, 200 nM or 20 nM of MemBright (MB-Cy3, MB-Cy3.5, MB-Cy5, MB-Cy5.5 and MB-Cy7) was added on the KB cell and incubated during 1 h at 37°C (5% CO<sub>2</sub>). As control, for each 96-well plate, the cells were incubated with DMEM containing the same percentage of DMSO (0,5% v/v) as the solution with the tested dyes or with Triton 1% as a positive control of cytotoxicity. After 1h of dye incubation, the medium was replaced by 100 μL of a mix containing DMEM + MTT solution (diluted in PBS beforehand) and the cells were incubated during 4 h at 37°C. Then, 75 μL of the mix was replaced by 50 μL of DMSO (100%) and gently shaken for 15 min at room temperature in order to dissolve the insoluble purple formazan reduced in living cells. The absorbance at 540 nm was measured (absorbances of the dyes at 540 nm were taken into account). Each concentration of dye was tested in sextuplicate in 3 independant assays. For each concentration, we calculated the percentage of cell viability in reference of the control DMEM+ 0,5% DMSO.

#### **TPE Measurements**

Two-photon absorption cross-section measurements were performed using Rhodamine B in methanol as a calibration standard according to the method of Webb et al. Two-photon excitation was provided by an InSight DS + laser (Spectra Physics) with a pulse duration of 120 fs. The laser was focused with an achromatic lens (f = 2 cm) in a cuvette containing the dye (0.2–1.0 µM in DMSO) and the spectra were recorded with a fibered spectrometer (Avantes) by collecting the fluorescence emission at 90° with a 20 × Olympus objective.

#### **Two-Photon Imaging**

Two-photon fluorescence microscopy imaging was performed by using a home-built two-photon laser scanning setup based on an Olympus IX70 inverted microscope with an Olympus 60x 1.2NA water immersion objective (Clamme et al., 2003). Two-photon excitation was provided by an InSight DS + laser (Spectra Physics), and photons were detected with Avalanche Photodiodes (APD SPCM-AQR-14-FC, Perkin-Elmer) connected to a counter/timer PCI board (PCI6602, National Instrument). Imaging was carried out using two fast galvo mirrors in the descanned fluorescence collection mode. Typical acquisition time was 50 s with an excitation power of 5 mW at the laser output. The images were processed with ImageJ.

#### **Tissue Imaging**

#### Rat Tissue

Fresh samples of rat liver tissue were cut (2×5 mm, 1 mm thickness) and kept in Krebs solution at RT. Before staining, the samples were washed 3 times with Krebs solution at RT. For the staining, the tissue samples were placed 3 h at RT in 1 mL of freshly prepared solution of dye (MB-Cy5: 5 μM in HBSS, WGA-647: 10 μg/mL in HBSS). The tissue samples were washed 3 times with HBSS before being placed in a iBiDi dish containing 3 mL of HBSS, and were then flattened by adding a glass slide on the top of it. The samples were excited at 810 nm with a power of 5 mW for both MB-Cy5 and WGA-647. 50 frames (60 × 60 μm, 512 × 512 pixels) were collected with a depth step of 1 µm, providing stacks of 50 µm depth. Each frame was scanned once with a speed of 4 µs per pixel. The images were processed with ImageJ.

C57BL6/J mice were maintained on a 12 hour light-dark cycle with ad libitum access to food and water. All animal work was conducted following protocol approved by ethical committee. Adult C57BL6 mice were euthanized by CO2 administration. Liver and brain were immediately dissected and washed in PBS at 4°C, before being sliced in 1 mm slices on a Leica Vibratome. Tissue slices were placed either at RT or at 4°C in 1 mL of freshly prepared solution of dye (MB-Cy3.5, Cy5 and Cy5.5: 5 μM in PBS) for a variable period (from 1 to 24 h). The tissue samples were washed 3 times with PBS before being placed in a homemade glass chamber, allowing the imaging on both side of the slides. Briefly a 1 cm hole was breached in a glass slide thanks to a diamond drill bit. The 1 mm tissue slice was inserted in the hole, covered by a coverslip (170 nm, #1.5) on each slide, and sealed with Picodent twinsil. The slices were excited on a LavisionBiotec two-photon Microscope at 800 nm with a TiSa laser (Insight Spectra Physics (690-1300 nm) for both MB-Cy3.5 and Hoechst. Each frame was scanned twice (line average 2) with a speed of 0.4 μs per pixel using a 20x plan Apo Chroma Zeiss (NA:1) water-immersion objective. The fluorescence signal was detected with GaAsP detectors. Confocal microscopy was performed on a Leica SP5 microscope (405 nm (25 mW); Argon Krypton 488 (40 mW), 561 nm (15 mW); HeNe 633 nm (10 mW)). Pictures were acquired with ideal sampling with a PL APO 20x/0.75 or 63x/1.4 CS2 objectives. The images were processed with Icy software.

#### **Neuronal Imaging**

Cells were dissociated by treatment with 0.25% trypsin for 15 min at 37°C and plated on poly-Ornithine (1 mg/mL, Sigma) coated glass coverslips in MEM supplemented with 10% horse serum, 0,6% glucose, 0,2% sodium bicarbonate, 2 mM glutamine, and 10 IU/mL penicillin-streptomycin. After attachment neurons were grown in neurobasal medium supplemented with B27 (Thermo Fisher) conditioned on astroglia. Neurons were then image either on Leica DMI8 spinning disc microscope for live cells or on SP5 confocal microscope and DMRE microscope for fixed samples. Neurons were incubated 10 min with MemBright at 37°C, washed 3 times in Krebs Ringer buffer, and then fixed with 4% PFA, 0.1% glutaraldehyde during 10 min at 37°C. Samples were then permeabilized if mentioned with 0.1% triton for 4 minutes and incubated sequentially with primary and secondary antibodies. Video-imaging of hippocampal neurons (first day *in vitro*) was performed on a Zeiss 880 confocal on gasp detectors, with 561 nm laser line set at 0.2%. Image were acquired every 2 min during 13 hours in Krebs ringer medium containing 200 nM MemBright Cy3.5. The signal in Video S5 was not processed neither for the noise nor the photo-bleaching (raw data).

#### dSTORM on HeLa cells

HeLa cells were cultured as mentioned above. The cells were fixed with 4% PFA during 10 minutes. The PM labelling was performed by incubating the cells with 20 nM MB-Cy3.5 in Opti-MEM during 10 minutes. All non-bound dyes were removed by 3 consecutive washing with PBS. dSTORM imaging was performed on a home built setup based on a Olympus IX-71 inverted microscope with a high-numerical aperture (NA) TIRF objective (Apo TIRF 100  $\times$  , oil, NA 1.49, Olympus) (Kempf et al., 2015). The samples were imaged in a photoswitching buffer containing 100 mM MEA and an oxygen scavenging system (0.5 mg/mL glucose oxidase, 40  $\mu$ g/mL catalase, 10% glucose) in PBS. All chemicals were purchased from Sigma-Aldrich. Samples were illuminated with a 532 nm laser (75 mW, Cobolt) and a 405 nm laser (25 mW, Spectra Physics) was used for the activation of the fluorophores. The signal was recorded on an EM-CCD camera (ImagEM, Hamamatsu) (0.106  $\mu$ m pixel size) with a typical integration time of 8.7 ms. The acquired stacks were processed with ImageJ software(Schneider et al., 2012) and the dSTORM images were reconstructed from a series of 6000 frames using the Thunder STORM plug-in (Ovesný et al., 2014). Acquired images were filtered with a wavelet filter and the approximate fluorophore positions were found by setting the threshold value equal to the standard deviation of the 1st wavelet function. Then, precise subpixel localizations were determined by fitting the spots with the Gaussian PSF by weighted least squares method. High-resolution images were reconstructed from obtained localizations and plotted as a histogram with 20 nm pixel size.

The localization uncertainty of the lateral position of individual fluorophores  $\Delta x$  was calculated according to following formula: (Thompson et al., 2002)

$$\langle (\Delta x)^2 \rangle = \frac{\sigma^2 + a^2/12}{N} + \frac{8\pi\sigma^4 b^2}{a^2 N^2}$$

Where  $\sigma$  is the standard deviation of the fitted Gaussain PSF (nm), a is the pixel size in nm, N is the number of photons detected and b is the background signal level in photons calculated as the standard deviation of the residuals between the raw data and the fitted PSF model (Křížek et al., 2011). In our experiments with typical values  $\sigma \sim$ 150 nm, a=106 nm, N $\sim$ 700 photons and b $\sim$ 15-20photons the resulting localization uncertainty was approximately 17-25 nm.

#### **dSTORM** on Neurons

Stochastic optical reconstruction microscopy (STORM) neuronal imaging was performed on a Vutara microscope (Bruker) with a high-numerical aperture (NA) objective (60x, water, NA 1.2, Olympus). 170  $\mu$ m coverslips (Menzel glaser 18 mm diameter #1.5) were mounted on a glass slide with a 15 mm hole. The hole was filled with imaging buffer (Tris 50 mM, NaCl 10 mM, 10% glucose, 100 mM MEA, 70 U/mL glucose oxidase (Sigma G0543), 20 g/mL catalase) and sealed with Picodent twinsil. Samples were illuminated successively with a 647 and 561 nm laser and a 405 nm laser for the reactivation of the fluorophores. Neurons were isolated with wide field mosaic microscopy (Cool snap camera) and then imaged for STORM for a series of 30,000-100,000 images with a FLASH4 CMOS camera (20 ms,  $20 \times 20$  microns). 3D-STORM imaging was performed using the bi-plane module allowing the localisation in the xyz direction. The Srx software (Bruker) was used to localise particles in 3D. Correction for chromatic aberration in super-resolution microscopy has been done using multispectral (blue/green/orange/dark red) Tetraspeck beads (Thermo Fisher T7279) in Bruker's software.

#### **QUANTIFICATION AND STATISTICAL ANALYSIS**

The statistical details of experiments can be found in the figures' legends or in the Method Details section.