Binding of His-tagged fluorophores to lipid bilayers of giant vesicles†

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His-tagged molecules can be attached to lipid bilayers via certain anchor lipids, a method that has been widely used for the biofunctionalization of membranes and vesicles. To observe the membrane-bound molecules, it is useful to consider His-tagged molecules that are fluorescent as well. Here, we study two such molecules, green fluorescence protein (GFP) and green-fluorescent fluorescein isothiocyanate (FITC), both of which are tagged with a chain of six histidines (6H) that bind to the anchor lipids within the bilayers. The His-tag 6H is much smaller than the GFP molecule but somewhat larger than the FITC dye. The lipid bilayers form giant unilamellar vesicles (GUVs), the behavior of which can be directly observed in the optical microscope. We apply and compare three well-established preparation methods for GUVs: electroformation on platinum wire, polyvinyl alcohol (PVA) hydrogel swelling, and electroformation on indium tin oxide (ITO) glass. Microfluidics is used to expose the GUVs to a constant fluorophore concentration in the exterior solution. The brightness of membrane-bound 6H-GFP exceeds the brightness of membrane-bound 6H-FITC, in contrast to the quantum yields of the two fluorophores in solution. In fact, 6H-FITC is observed to be strongly quenched by the anchor lipids which bind the fluorophores via Ni²⁺ ions. For both 6H-GFP and 6H-FITC, the membrane fluorescence is measured as a function of the fluorophores' molar concentration. The theoretical analysis of these data leads to the equilibrium dissociation constants $K_d = 37.5 \text{nM}$ for 6H-GFP and $K_d = 18.5 \text{nM}$ for 6H-FITC. We also observe a strong pH-dependence of the membrane fluorescence.

1 Introduction

Lipid bilayers, which represent a universal building block for all biomembranes and, thus, for the architecture of the living cell, provide a versatile biomimetic module to understand the mechanical properties of cells and organelles as well as their responses to external stimuli. In aqueous solutions, lipid bilayers form closed vesicles which have a wide range of applications in the field of synthetic biology,¹⁻⁵ intracellular cargo transport and trafficking⁶⁻⁹ and the systematic study of biomembrane properties.¹⁰⁻¹³ From an experimental point of view, giant unilamellar vesicles (GUVs) are particularly useful because they are micrometer-sized, comparable to the size of eukaryotic cells. In addition, GUVs can be directly imaged by optical microscopy and their mechanical responses can be probed by optical tweezers and micropipettes.¹⁴

Two broad categories of membrane proteins can be distinguished. First, integral membrane proteins have one or several transbilayer domains, which span the hydrophobic core of the bilayer membranes. Second, peripheral membrane proteins, which take part in cell signaling and membrane trafficking, can attach to one leaflet of the bilayer membranes by binding to a specific lipid or to a small cluster of several lipids.¹⁵ Likewise, to achieve biofunctionalisation of GUVs, integral membrane proteins are inserted into the membranes using detergents¹⁶ or proteoliposomes¹⁷,¹⁸ whereas peripheral membrane proteins are attached via certain anchor lipids. Here, we will consider a specific lipid anchor as provided by DGS-NTA(Ni) or NTA lipid for short, which binds to poly-histidine tagged molecules via coordinate bonds. Two such molecules will be considered, His-tagged GFP and His-tagged FITC, both of which are green fluorescent, see Fig. 1.

Poly-histidine tags are generally attached to one of the terminals of the proteins for their purification.¹⁹ Hence, this specific interaction can be harnessed to attach proteins to lipid bilayers.¹⁶,²⁰⁻²⁴ The NTA(Ni) in the lipid head group forms an octahedral coordinate complex with poly-histidine chains. Four
Fig. 1 Cartoon depicting the lipid bilayer composed of POPC, cholesterol, and DGS-NTA(Ni) or NTA lipid, which acts as an anchor for a poly-histidine chain. Two types of molecules tagged with a chain of six histidines will be studied. 6H-GFP and the 6H-FITC, both of which are green fluorescent. The barrel height of 6H-GFP is comparable to the membrane thickness (left) whereas the linear extension of 6H-FITC is governed by the length of the histidine chain (right). The optical image displays a GUV with membrane-bound 6H-GFP. Scale bar: 10 μm.

The binding between His-tagged proteins and NTA anchor lipids has been frequently used in previous studies. Examples include protein crowding on vesicles, lipid-coated substrates for drug delivery systems, artificial cell adhesion, and high spontaneous curvature generated by the dilute concentration of surface proteins. In general, the Ni^{2+} ion can be replaced by a Co^{3+} ion in the NTA group. Poly-histidine chains form a stronger and less labile coordinate bond with NTA-Co^{3+} compared to the corres-
ponding bond with NTA-Ni^{2+}.20–22

GUVs can be prepared using a variety of well-established protocols such as swelling lipid layers on a substrate by hydration, electroformation, phase transfer across liquid–liquid interfaces, and droplet stabilized compartments; for a recent review, see ref. 14. All of these techniques have their advantages and disadvantages. The swelling methods involve the spreading of a lipid mixture on a substrate followed by the evaporation of the (organic) solvent. As a result, one obtains a stack of lipid bilayers which can then be swollen by hydration. Spontaneous swelling is achieved by spreading the lipid on a non-reactive surface and gently hydrating it for an appropriate period of time.35 This gentle hydration method is limited to the use of specific lipids and solution conditions.36

The influx of the hydrating solution can be enhanced by hydrogel-assisted swelling.36,37 The lipid film is deposited on the surface of the dried hydrogel. The hydrogel absorbs the hydrating solution and GUVs form rapidly at the hydrogel–aqueous solution interface. The hydrogels typically used for this purpose are agarose,37,38 polyvinyl alcohol (PVA),28,36,39 and polyacrylamide gels.40 Sometimes the hydrogel becomes incorporated into the lumen of the GUV or even within the lipid bilayer, which represents a drawback of this method because it affects the mechanical bilayer properties.38,40 To avoid these unwanted side effects, electroformation can be used to control the hydration of the lipid films by external alternating currents (ACs).32 The substrate for spreading the lipid film must be electrically conductive. Commonly used substrates include platinum wires,43 ITO coated glasses,40,43 stainless steel wires,43 copper electrodes,44 and carbon fiber microelectrodes.48

One implicit assumption that is often made in the preparation of GUVs is that these vesicles have the same composition as the lipid mixture that was initially used to grow them. This assumption should be checked, especially when the GUVs are prepared from a lipid mixture with a certain fraction of DGS-NTA(Ni). Furthermore, the binding of the poly-histidine with the NTA(Ni) lipid is also sensitive to the solution conditions. Here, we will analyze and compare different preparation techniques of GUVs in different pH conditions. As shown in Fig. 1 and 2, we will use two different fluorescent probes, 6H-GFP and 6H-FITC, where 6H stands for a linear chain of six histidines, and study the binding of these fluorophores to NTA lipids embedded in GUV membranes.

The molecular weight of the two fluorophores is quite different: GFP has a molecular weight of 27 kD whereas FITC has a much smaller molecular weight of only 389 D. Therefore, the two fluorophores are also quite different in size. As shown below, one important consequence of this size difference is that the fluorescence of FITC is strongly quenched by the NTA lipids, in contrast to the fluorescence of GFP.

2 Results and discussion

2.1 Brightness of GUV membranes

Three different preparation methods are used to form the GUVs: electroformation on platinum wire, PVA hydrogel
swelling, and electroformation on ITO glass. In all cases, we start from the same initial lipid mixture with 3 mol% of DGS-NTA(Ni) to grow the GUVs, which are then exposed to a certain concentration of the His-tagged fluorophores. After 10 min, the GUVs are directly observed with a confocal microscope, see the images in Fig. 3.

In order to expose the GUVs to a constant concentration of the His-tagged molecules in the exterior solution, we control this concentration by microfluidics and a specific design of the microfluidic chip: we trap individual GUVs in the short side channels of the chip and use its main channel for the fast exchange of the exterior solution. Further details about the microfluidic approach are described in the Material and method and Fig. 10.

Two types of His-tagged molecules are studied: 6H-GFP, corresponding to a chain of six histidines attached to the N terminal of GFP,28 and 6H-FITC consisting of FITC covalently bound to the same histidine chain. The histidine chain binds to the NTA(Ni) head group of the DGS-NTA(Ni) lipid in the vesicle via a coordinate bond (Fig. 2).

As shown in Fig. 3, the observed brightness of the GUV membranes depends both on the His-tagged fluorophore and on the preparation method. For both fluorophores, the GUVs are observed to have the largest brightness when prepared by platinum wire electroformation. As will become clear further below in Fig. 6 and 7, both molar concentrations chosen in Fig. 3 – 320 nM for 6H-GFP and 240 nM for 6H-FITC – belong to the saturation regime for the platinum wire method, in which almost all NTA lipids are occupied by His-tagged molecules.

For a spherical GUV, the membrane fluorescence can be determined quantitatively by measuring the fluorescence intensity profile as a function of the radial coordinate \( r \), which represents the distance from the center of the vesicle. One example for such a profile is shown in Fig. 4a together with the baseline intensity that interpolates smoothly between the background intensities of the interior and exterior solutions. The excess intensity of the fluorescence is equal to the difference between the total fluorescence and baseline intensities, see Fig. 4b, and the membrane fluorescence is obtained by integrating this excess intensity over the radial coordinate \( r \). The excess intensity profiles corresponding to the GUVs in Fig. 3a–f are displayed in Fig. 3g and h.

The GUV images and line profiles show that the brightness of membrane-bound 6H-FITC is reduced compared to the one of membrane-bound 6H-GFP. In contrast, in the absence of lipids, 6H-FITC has a quantum yield that is about 1.2 larger than the quantum yield of 6H-GFP, as obtained from independent measurements, see Material and method section. This different behavior indicates that the fluorescence of 6H-FITC is quenched when this molecule is bound to the membrane or, more precisely, to an NTA anchor lipid.

### 2.2 Fluorescence quenching by NTA lipids

In order to elucidate the quenching effect of the NTA lipids, we added, for each fluorophore, an increasing concentration of NTA lipids to a 120 nM solution of the fluorophore. The resulting fluorescence intensities are displayed in Fig. 5. As shown in Fig. 5a, the fluorescence of 6H-GFP is hardly affected by large concentrations of NTA lipids, whereas the fluorescence of 6H-FITC is strongly reduced by small concentration of these lipids, see Fig. 5b.

The data in Fig. 5a and b were obtained by adding liposomes with the lipid composition POPC : Chol : NTA as given by 6 : 1 : 3. Similar quenching effects were observed when the fluorophores were exposed to nickel sulphate (data not shown). On the other hand, no fluorescence quenching of 6H-FITC was observed when the liposomes contained only POPC and cholesterol but no NTA lipids, as demonstrated in Fig. S1 (ESI).

Therefore, the fluorescence quenching of 6H-FITC is caused by NTA lipids which bind the fluorophore via their Ni\(^{2+}\) ion, see Fig. 2. Fluorescence quenching by Ni\(^{2+}\) ions has been previously reported for a variety of fluorophores. For membrane-bound 6H-GFP, such a quenching effect does not occur because the chromophore of 6H-GFP is located in the middle of the...
protein’s β-barrel and well-separated from the Ni$^{2+}$ ion of the NTA lipid. In contrast, the fluorescein group of membrane-bound 6H-FITC is much closer to this Ni$^{2+}$ ion, see Fig. 2, which explains the strong quenching effect in Fig. 5b.

2.3 Fluorescence of membrane-bound 6H-GFP

Using the Pt wire method, we measured the membrane fluorescence $I_{flu}$ of 6H-GFP as a function of its molar concentration $X$ for the range $0 \leq X \leq 320$ nM. The corresponding normalised intensity $I_{flu}/I_{sat}$ is plotted in Fig. 6a with $I_{sat} = 320.4$ a.u. where a.u. stands for arbitrary (or procedure-defined) units. These data are well fitted by the intensity–concentration relationship as given by eqn (6) in the Material and method. From this fit, we obtain the equilibrium dissociation constant $K_d = (37.5 \pm 7.5)$ nM for 6H-GFP.

Inspection of Fig. 6a shows that the fluorescence intensity becomes saturated for large values of the molar concentration $X$. This saturation regime is characterized by the maximal number of 6H-GFP that can be bound to the NTA anchor lipids. For a lipid bilayer consisting of POPC and cholesterol with a molar ratio of 8:2, the average area per lipid is 0.5 nm$^2$ as follows from previous studies. For such a bilayer doped with 3 mol% anchor lipids, neighboring anchor lipids have an average separation of 4.1 nm. This anchor–anchor separation exceeds the lateral size of membrane-bound 6H-GFP, as explained in the Material and method. Therefore, we conclude that, in the saturation regime, the membrane-bound fluorophores have the same average separation as the anchor lipids, corresponding to an average surface coverage of $6.0 \pm 10^4$ molecules per mm$^2$, see eqn (1). In Fig. 6b, we use this saturation-based calibration to transform the fluorescence intensities into surface coverages for the three preparation methods. The scatter in the data of Fig. 6b reflects the different lipid compositions of individual GUV membranes that deviate from the overall lipid composition used for the preparation of these vesicles. The numerical values of the data in Fig. 6a and b are given in Tables S1 and S2 (ESI†).

2.4 Binding of 6H-FITC to GUV membranes

Next, we studied the His-tagged molecule 6H-FITC, consisting of six histidines attached to the small fluorescent probe FITC. We used the same experimental protocol to determine membrane fluorescence and surface coverage as described in the previous section for 6H-GFP. To obtain the equilibrium...
dissociation constant for 6H-FITC, we varied the molar concentration $X$ of 6H-FITC over the range $0 \leq X \leq 480$ nM and measured the X-dependent fluorescence intensity of the GUV membranes, which were prepared by platinum wire electroformation. The corresponding normalised intensity $I_{\text{flu}}/I_{\text{sat}}$ is plotted in Fig. 7a with $I_{\text{sat}} = 40.9$ a.u. These data are well fitted by the intensity–concentration relationship as given by eqn (6) in the Material and method. From this fit, we obtain the equilibrium dissociation constant $K_d = (18.5 \pm 3.7)$ nM for 6H-FITC.

Comparison of the equilibrium dissociation constant for 6H-FITC with the one for 6H-GFP as given by $K_d = 37.5$ nM reveals that 6H-FITC is more strongly bound to the GUV membranes compared to 6H-GFP. Furthermore, the different values for the saturation intensity $I_{\text{sat}}$ imply that the fluorescence of membrane-bound 6H-FITC is strongly reduced compared to the one of membrane-bound 6H-GFP, in agreement with the results on fluorescence quenching by NTA lipids in Fig. 5.

The GUV membranes in Fig. 7a and 6a contain the same molar fraction of 3 mol% NTA lipids. Therefore, the saturation values of the surface coverage by the two fluorophores are also identical and equal to $6.0 \times 10^4$ molecules per $\mu$m$^2$ as obtained for 6H-GFP by the platinum wire method. In Fig. 7b, we use this saturation-based calibration to transform the fluorescence...
intensities into surface coverages for the three preparation methods. The scatter in the data of Fig. 7b reflects the different lipid compositions of individual GUV membranes that deviate from the overall lipid composition used for the preparation of these vesicles. The numerical values corresponding to the data in Fig. 7a and b are given in Tables S3 and S4 (ESI†).

2.5 Different compositions of lipid bilayers

**Dependence on mole fraction of NTA lipids.** To get additional insight into the weak binding of His-tagged molecules to GUVs that are grown by electroformation on ITO glass, we compare this method with electroformation on platinum wire, using two different lipid mixtures with mole fractions of 3 mol% and 30 mol% DGS-NTA(Ni) lipids. After harvesting the GUVs, they are exposed to 120 nM 6H-FITC in the exterior solution. As displayed in Fig. S2 (ESI†), GUVs prepared by electroformation on ITO glass exhibit a smaller membrane fluorescence than GUVs prepared by electroformation on platinum wire, indicating that only a fraction of the DGS-NTA(Ni) lipid deposited on the ITO glass becomes incorporated into the vesicles. However, for both electroformation methods, the fluorescence increases with increasing mole fraction of the NTA lipids.

A possible explanation for the low membrane fluorescence of GUVs prepared by ITO glass electroformation is as follows. After the deposition of lipid stock on the ITO glass surface, the lipid molecules have orientational freedom to arrange themselves into bilayers.† The surface properties of the ITO glass are likely to affect the properties of the produced vesicles. The ITO film consists of indium(III) ions, which can bind to NTA. Presumably, the NTA lipid prefers to stay close to the ITO surface and forms a weak bond with indium, when the lipid stock dissolved in chloroform is deposited on the ITO surface. When the fraction of NTA is increased in the lipid stock to 30 mol%, the ITO surface becomes saturated and some excess NTA is then incorporated into the vesicle membrane.

**Dependence on mole fraction of cholesterol.** GUVs with two different lipid compositions are prepared by platinum wire electroformation and exposed to 120 nM 6H-FITC. Both lipid compositions contain 3 mol% NTA lipids but differ in their mole fractions of cholesterol. For lipid bilayers with 20 mol% cholesterol, the intensity is about twice as large as for bilayers without cholesterol, see Fig. S3 (ESI†). This change in intensity can arise by two different mechanisms. First, cholesterol may reduce the fluorescence quenching by the NTA lipids; second it may increase the surface coverage of 6H-FITC. For a certain variant of His-tagged GFP, the binding affinity to lipid bilayers was found to increase with the mole fraction of cholesterol.†

2.6 pH dependence of membrane fluorescence

The data on membrane fluorescence and surface coverage as displayed in Fig. 6 and 7 are obtained for pH 7.4. These two quantities depend on the pH-value as can be directly concluded from the brightness of the GUV membranes, which varies with the pH, see Fig. 8. To determine the pH-dependence of the fluorescence intensity in a systematic manner, the vesicles were grown using platinum wire electroformation. The swelling solution consisted of 50 mM sucrose and 2 mM HEPES buffer at pH 7.4. The vesicles were then exposed to an exterior solution containing 20 nM 6H-GFP or 120 nM 6H-FITC as well as 22.5 mM NaCl, 5 mM sucrose and 2 mM HEPES buffer. The pH of these solutions was increased and decreased by adding NaOH and HCl, respectively.

For each pH value, the membrane fluorescence was determined via the method described in Fig. 4. The resulting fluorescence intensities of the GUV membranes are displayed in Fig. 9a and b for 6H-GFP and 6H-FITC, respectively. For 6H-GFP, the membrane fluorescence first increases and then saturates for pH values above pH 8.5. For 6H-FITC, the intensity exhibits a pronounced maximum at about pH 8.5. The numerical values of the data in Fig. 9a and b are given in Tables S5 and S6 (ESI†).

In general, the fluorescence of a dye molecule can vary with the pH value even in the absence of lipids. In order to determine the latter pH dependence, we also measured the background intensities for 6H-GFP and 6H-FITC in the exterior solution far away from the vesicle membranes, see the corresponding data in Fig. 9. Inspection of both panels shows that the background intensity in the exterior solution is negligible over the whole range of pH values studied here. Therefore, the pH dependence of the membrane fluorescence in Fig. 9 is caused by the pH dependence of the membrane-bound fluorophores rather than by the intrinsic pH dependence of these dye molecules in solution.
3 Conclusions and outlook

The binding of His-tagged molecules to GUV membranes has been investigated, using the anchor lipids DGS-NTA(Ni) that form coordinate bonds with the His-tags (Fig. 2). Two different green fluorescent molecules, 6H-GFP and 6H-FITC, have been studied, both of which were tagged to a chain of six histidines. Using three different preparation methods – electroformation on platinum wire, swelling of PVA hydrogels, and electroformation on ITO glass – as well as microfluidics (Fig. 10) to expose the GUVs to a constant bulk concentration of the fluorophores, we obtained the images in Fig. 3 which show that the brightness of the GUV membranes depends both on the type of His-tagged molecule and on the preparation method. To determine the membrane fluorescence in a quantitative manner, we measured the fluorescence intensity profiles, see Fig. 4a, and obtained the membrane fluorescence by integrating the excess intensity profiles as in Fig. 4b over the radial coordinate.

One surprising result of our study is the different behavior of the two fluorophores when exposed to an increasing amount of NTA lipids. Indeed, we observed essentially no change in the fluorescence of 6H-GFP, see Fig. 5a, but a strong quenching effect on 6H-FITC as demonstrated by the data in Fig. 5b. Both for 6H-GFP and 6H-FITC, the highest brightness of the GUV membranes was obtained when we prepared these vesicles via electroformation on platinum wire. We used this latter method to measure the dependence of the fluorescence intensities on the molar concentrations of the His-tagged fluorophores (Fig. 6a and 7a). Both sets of data are well fitted by the intensity–concentration relationship in eqn (6), which depends on only two parameters, the equilibrium dissociation constant $K_d$ and the saturation intensity $I_{sat}$.

The equilibrium dissociation constant turns out to be $K_d = (37.5 \pm 7.5)$ nM for 6H-GFP and $K_d = (18.5 \pm 3.7)$ nM for 6H-FITC which implies that 6H-FITC is more strongly bound to the GUV membranes. On the other hand, the strong quenching of 6H-FITC by the Ni$^{2+}$ ions of the NTA lipids (Fig. 5b) leads to the saturation intensity $I_{sat} = 40.9$ a.u. for 6H-FITC which is much smaller than for 6H-GFP.
smaller than \( I_{\text{sat}} = 320.4 \text{ a.u.} \) for the membrane fluorescence of 6H-GFP.

Our method to determine the equilibrium dissociation constant \( K_d \) is quite general and can be applied to the binding between other His-tagged fluorophores and/or other anchor lipids. First, one measures the fluorescence intensities as a function of the molar concentration \( X \) and then plots the normalised fluorescence intensities \( I_{\text{flu}}/I_{\text{sat}} \) versus \( X \) as in Fig. 6a and 7a. Second, one fits the plotted data by the intensity-concentration relationship in eqn (6), which involves two fit parameters, the equilibrium dissociation constant \( K_d \) and the saturation intensity \( I_{\text{sat}} \). If the data can be well-fitted by this equation, one obtains a reliable estimate for \( K_d \). On the other hand, if the fit turns out to be unacceptable, one can conclude that the model used to derive eqn (6) should be modified. The most important simplifying assumption of this model is that each anchor lipid can bind one fluorophore. Therefore, a poor fit will provide direct evidence that the latter assumption does not apply to the system under consideration. Possible modifications of the model include (i) crowding, i.e., the steric hindrance between the membrane-bound fluorophores, (ii) the binding of several fluorophores to one anchor lipid, and (iii) the recruitment of several anchor lipids by one fluorophore.

Another surprising outcome of our study is that it reveals a pronounced pH dependence of the membrane fluorescence for both 6H-GFP and 6H-FITC, see Fig. 9a and b. For 6H-GFP, the membrane fluorescence first increases and then saturates for pH values above pH 8.5. For 6H-FITC, on the other hand, the membrane fluorescence exhibits a pronounced maximum at about pH 8.5 (Fig. 9b). Furthermore, the observed membrane fluorescence was always much larger than the background intensity, corresponding to those fluorophores in the exterior solution that were not in contact with the lipid membranes. Therefore, the pH dependence displayed in Fig. 9 represents the pH dependence of the membrane fluorescence rather than the intrinsic pH dependence of the fluorescent molecules in solution.

Previous experiments have shown that GUV membranes exposed to nanomolar solutions of 6H-GFP acquire a large spontaneous curvature that can be fine-tuned to divide the GUVs in a controlled manner. As shown here, 6H-FITC has a smaller equilibrium constant and is, thus, more strongly bound to GUV membranes compared to 6H-GFP. Therefore, it will be interesting to study the spontaneous curvature generated by 6H-FITC and the shape transformations of GUVs arising from this curvature generation.

4 Material and method

4.1 Preparation of giant unilamellar vesicles

Chloroform stock solutions of 1-palmitoyl-2-oleoyl-sn-glycero-3-phosphocholine (POPC) and cholesterol were mixed at a molar ratio of 8:2 with a final lipid concentration of 4 mM. The required mol% of DGS-NTA(Ni) was added to the lipid stock solution. This stock was used as it is or diluted depending on the preparation method. All lipids were obtained from Avanti Polar Lipids. His-tagged GFP was obtained from Astrid Krämer of the protein facility at the Max Planck Institute of Molecular Physiology (Dortmund, Germany). His-tagged FITC was purchased from Biomatik Corporation (Ontario, Canada).

Electroformation on platinum wire

Using the lipid mixture as specified in the previous paragraph, we spread 6 \( \mu \text{L} \) of a 0.5 mM solution of this mixture on two platinum wires via a syringe. The wires were kept under vacuum for an hour to remove traces of chloroform. The wires were then dipped in a quartz cuvette, which was filled with 50 mM of sucrose in 2 mM HEPES buffer (pH 7.4). An AC electric field with peak-to-peak voltage of 3 V and frequency of 10 Hz was applied for 2 hours at 35 °C to speed up the vesicle swelling process. Subsequently, the frequency of the AC field was reduced to 3 Hz and the low-frequency field was applied for 5 min, to detach the vesicles from the wires. After cooling to room temperature, the vesicles were harvested using a micropipette with a broad tip.

PVA hydrogel swelling

A PVA solution was prepared by dissolving 40 mg of PVA in 1 mL of water. 40 \( \mu \text{L} \) of this solution was spread on clean cover glasses and dried in an oven. An AC electric field with peak-to-peak voltage of 3 V and frequency of 10 Hz was applied for 2 hours at room temperature (23 °C) for 30 minutes. The vesicles were harvested after gentle tapping on the PVA coated glass.

Electroformation on ITO glasses

Commercially available ITO glass from PGO-GmbH (Iserlohn, Germany) was used without further treatment. In general, the ITO glass surface can have a complex molecular structure, depending on the pretreatment conditions. A 4 \( \mu \text{L} \) of a 4 mM solution of the lipid mixture was spread on two ITO coated glasses using a syringe. The resulting lipid films formed bilayers stacks on the ITO glass surfaces, which were then kept under vacuum for an hour to remove remaining traces of chloroform. The two glasses were then arranged into a closed chamber by using a Teflon spacer as side walls. The chamber was filled with 50 mM of sucrose in 2 mM HEPES buffer (pH 7.4). An AC electric field with peak-to-peak voltage of 2 V and frequency of 10 Hz was applied for 2 hours at room temperature (23 °C). Subsequently, the frequency of the AC field was reduced to 3 Hz and the low-frequency field was applied for 5 min, to detach the vesicles from the glass surfaces. The vesicles were harvested using a micropipette with a broad tip.
4.2 Microfluidic observation chamber

The experiments were performed on polydimethylsiloxane (PDMS) microfluidic chips. The design of the device is such that the vesicles are trapped in the dead-end side channels that are perpendicular to the main channel, as in Fig. 10a. The global design of the microfluidic chip is displayed in Fig. S4 (ESI†). The solution volume that can be loaded in the microfluidic device is 2.5 μL and each of the side channels is 250 μm deep and 100 μm wide. The reservoir in the inlet is filled with vesicles and NaCl solution. The vesicles are loaded on the chip using a syringe connected to a pump at the outlet. Then the chip is oriented vertically in such a way that the vesicles slowly fall into the side channels. After about 20 min, the chip is placed on the microscope. The solution in the reservoir is exchanged by another solution with a certain concentration of His-tagged molecules. Because of the continuous flow from the inlet, the concentration of these molecules remains constant during the whole experiment.

On the other hand, the His-tagged molecules reach the vesicles in the side channels only by diffusion which implies that the vesicles are shielded from the hydrodynamic flow and do not suffer any mechanical perturbations that could otherwise arise from this flow. The trapping in the side channels also helps to monitor the same GUV throughout the experiment. As shown in Movie S1 (ESI†), the complete solution exchange including the side channels takes around 3.5 min. For all experiments performed here, the pump was used to exchange 20 μL of the his-tagged molecules, corresponding to 8 times the solution volume within the microfluidic device, before the vesicles have been imaged.

4.3 Quantum yield of fluorophores in the absence of lipids

To determine the different quantum yields of the two fluorophores, the detector gain was adjusted to give the same intensity for the same molar concentrations of 6H-FITC and 6H-GFP in the absence of lipids. Briefly, 100 nM solutions of 6H-FITC and 6H-GFP were imaged using the HyD detector on the Leica SP8 confocal microscope. Both molecules were excited using a 488 nm Argon laser and emission was collected between 495 and 550 nm. The detector gain was adjusted to obtain the same brightness for both solutions. From this adjustment, we concluded that 6H-FITC is about 1.2 times brighter than 6H-GFP.

4.4 Measurements of fluorescence quenching

Experiments to observe the possible quenching of the fluorescence of the His-tagged fluorophores by the DGS-NTA(Ni) lipids were conducted using the FluorMax-4 fluorimeter. LUVs were prepared by hydrating dried lipid films (POPC:cholesterol (8:2) and 30 mol% DGS-NTA(Ni)) with 2 mM HEPES buffer. After vortexing for 30 min, the suspension was extruded 21 times through a 200 nm polycarbonate filter (Avanti mini extruder). Fluorescence spectra were recorded for 120 nM 6H-GFP and 120 nM 6H-FITC in the presence and in the absence of the LUVs by exciting at 488 nm. Control experiments were also performed for LUVs without the DGS-NTA(Ni) lipids, see Fig. S1 (ESI†).

4.5 Image acquisition

All fluorescent images were recorded using a Leica SP8 confocal microscope. Argon laser with wavelength 488 nm was used to excite both the GFP proteins and the FITC dye. The emission was collected in the range of wavelengths from 495 to 550 nm. The images for quantification of the fluorescence intensity on the membrane were obtained using a 40 × 1.3 oil immersion objective.

4.6 Separation of NTA lipids and lateral size of fluorophores

For a lipid bilayer consisting of POPC and cholesterol with a molar ratio of 8:2, the average area per lipid is 0.5 nm² as follows from previous studies. When such a bilayer is doped with 3 mol% NTA lipids, these anchor lipids have an average spatial separation of 4.08 nm. This separation determines the surface density ρanc of the NTA lipids via

$$\rho_{\text{anc}} = \frac{1}{(4.08 \text{ nm})^2} = \frac{6.0 \times 10^4}{\mu \text{m}^2}$$

(1)

When each NTA anchor lipid is occupied by one His-tagged molecule, corresponding to the saturation regime, the surface density ρanc of the anchor lipids is equal to the surface density or coverage of the His-tagged fluorophores.

In order to estimate whether or not this saturation regime leads to crowding of the membrane-bound fluorophores, we need to compare the average separation of the NTA lipids with the lateral size of the membrane-bound fluorophores. The lateral size of membrane-bound 6H-GFP should be comparable with the diameter of its β-barrel. This diameter is about 3 nm which is smaller than the anchor–anchor separation of 4.08 nm and implies that crowding effects between membrane-bound 6H-GFP molecules can be ignored even when each NTA anchor lipid is occupied by one such molecule. The 6H-FITC peptide, on the other hand, has a linear extension of about 3.2 nm which provides an upper bound on the lateral size of membrane-bound 6H-FITC. Because this upper bound on the lateral size is again smaller than the average separation of the NTA anchor lipids, crowding effects between membrane-bound 6H-FITC molecules can again be ignored even when each NTA lipid binds one such molecule.

4.7 Fluorescence of membrane-bound fluorophores

Radial Profile Angle from ImageJ was used to plot the fluorescence intensity as a function of the radial coordinate r, see Fig. 4a. The peak analyzer of Origin2021b from OriginLab was used to determine the excess intensity in Fig. 4b by subtracting the baseline intensity from the fluorescence intensity. Then the membrane fluorescence was obtained by integrating the excess intensity over the radial coordinate r, using the whole range of r-values.

To measure the membrane fluorescence at different pH values, the vesicles were grown using platinum wire electroformation at pH 7.4. These vesicles were then trapped and...
exposed to the solution of the His-tagged molecules at different pH, ranging from 6.4 to 9.4. The pH of the solution was adjusted using HCl or NaOH. The final pH of the solutions was measured using a micro pH electrode from Mettler Toledo. The results of these measurements are displayed in Fig. 9.

4.8 Binding kinetics and equilibrium dissociation constant

The binding sites for the His-tagged molecules are provided by the NTA anchor lipids. We consider a simple binding model and assume that each anchor lipid can bind at most one His-tagged molecule. The outer leaflet of the GUV membrane contains a total number of $N_{anc}$ anchor lipids which can bind $N_{bH}$ His-tagged molecules with $N_{bH} \leq N_{anc}$. For each anchor lipid, the binding (or association) rate is taken to be proportional to the molar concentration $X$ and, thus, has the form $k_{on}X$ with the association rate constant $k_{on}$, while unbinding (or dissociation) is an activated process with the dissociation rate $\omega_{off}$. The number $N_{bH}$ of bound His-tagged molecules then changes with time $t$ according to the evolution equation

$$dN_{bH}/dt = k_{on}X(N_{anc} - N_{bH}) - \omega_{off}N_{bH}. \tag{2}$$

In equilibrium, the binding flux is equal to the unbinding flux which implies $dN_{bH}/dt = 0$. The equilibrium value $N_{eq}^{bH}$ of the bound His-tagged molecules is then given by

$$N_{eq}^{bH} = N_{anc}X/K_{d} + X \tag{3}$$

which depends on the equilibrium dissociation constant

$$K_d = \omega_{off}/k_{on}. \tag{4}$$

When we divide eqn (3) by the surface area $A$ of the membrane, we obtain the equilibrium surface density or coverage $\Gamma_{eq}^{bH}$ of the His-tagged molecules which depends on the surface density $\rho_{anc}$ of the NTA anchor lipids and on the molar concentration $X$ according to

$$\Gamma_{eq}^{bH} = \frac{N_{eq}^{bH}}{A} = \frac{N_{anc}X}{A} \frac{X}{K_{d} + X} = \frac{\rho_{anc}}{K_{d} + X} \tag{5}$$

For 3 mol% NTA lipids, the surface density $\rho_{anc}$ is equal to 6.0 $\times$ 10$^7$ molecules per $\mu$m$^2$ as in eqn (1).

4.9 Fluorescence intensity of membrane-bound fluorophores

The fluorescence intensity of the membrane-bound fluorophores is proportional to their surface density. Because the fluorescence of these fluorophores can be quenched by the NTA anchor lipids, we introduce a reduction factor $\phi$ with $0 < \phi \leq 1$ for the fluorescence of a fluorophore bound to the NTA lipid. The data in Fig. 5 imply that $\phi = 1$ for 6H-GFP and $\phi < 1$ for 6H-FITC. We then obtain the relationship

$$I_{flu} = I_{sat} \frac{X}{K_d + X} \quad \text{with} \quad I_{sat} = \phi \rho_{anc} \tag{6}$$

between the fluorescence intensity $I_{flu}$ and the molar concentration $X$. As a consequence, the fluorescence intensity $I_{flu}$ increases monotonically with the molar concentration $X$ and attains its saturation value $I_{flu} \approx I_{sat} = \phi \rho_{anc}$ for large $X$ as in Fig. 6a and 7a.

For 6H-GFP and 6H-FITC, the fluorescence intensities attain the saturation values $I_{GFP} = 320.4$ a.u. and $I_{FITC} = 40.9$ a.u. as obtained from the HyD detector on the Leica SP8 confocal microscope. Taking into account that the data in Fig. 6a and 7a were obtained for GUV membranes with the same mole fraction of 3 mol% NTA lipids and thus with the same surface density $\rho_{anc}$, we obtain the ratio

$$\frac{I_{FITC}}{I_{GFP}} = \frac{\phi_{FITC}}{\phi_{GFP}} = \frac{40.9}{320.4} = 0.13 \tag{7}$$

of the two fluorescence reduction factors $\phi_{FITC}$ and $\phi_{GFP}$. It then follows from $\phi_{GFP} = 1$ for 6H-GFP, that the fluorescence of membrane-bound 6H-FITC molecules is reduced by the factor $\phi_{FITC} = 0.13$.

4.10 Calibration of surface coverages via saturation regime

The saturation regime of the His-tagged fluorophores, which corresponds to the limit of large molar concentrations $X$, is used to calibrate the transformation from membrane fluorescence to surface coverage. For large $X$, the coverage $\Gamma_{eq}^{bH}$ of the membrane-bound fluorophores is equal to the surface density $\rho_{anc}$ of the NTA anchor lipids as in eqn (5). A combination of this latter equation with eqn (6) leads to

$$\Gamma_{eq}^{bH} = \rho_{anc} \frac{X}{K_d + X} = \rho_{anc} \frac{I_{flu}}{I_{sat}} \tag{8}$$

which allows us to compute the surface coverage $\Gamma_{eq}^{bH}$ of the His-tagged molecules from (i) the surface density $\rho_{anc}$ of the NTA anchor lipids, see eqn (1), and (ii) the data for the normalized fluorescence intensity $I_{flu}/I_{sat}$ as displayed in Fig. 6a and 7a.

Abbreviations

AC Alternating current
DGS-NTA(Ni) or NTA for short
1,2-Dioleoyl-sn-glycero-3-[(N-(5-aminocarbonyl)pentyl)iminodiacetic acid]succinyl] (nickel salt)
FITC Fluorescein isothiocyanate
GFP Green fluorescent protein
ITO Indium tin oxide
POPC 1-Palmitoyl-2-oleoylsn-glycero-3-phosphocholine
PVA Polyvinyl alcohol
6H-GFP Green fluorescent protein tagged by six histidines
6H-FITC Fluorescein isothiocyanate tagged by six histidines

Conflicts of interest

There are no conflicts to declare.

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References

1. C. Schmitt, A. H. Lippert, N. Bonakdar, V. Sandoghdar and L. M. Voll, *Front. Bioeng. Biotechnol.*, 2016, 4, 19.
2. K. Kamiya, *Micromachines*, 2020, 11, 559.
3. L. Van de Cauter, F. Fanalista, L. van Buren, N. De Franceschi, E. Godino, S. Bouw, C. Danelon, C. Dekker, G. H. Koenderink and K. A. Ganzinger, *ACS Synth. Biol.*, 2021, 10, 1610–1702.
4. J. W. Hindley, R. V. Law and O. Ces, *SN Appl. Sci.*, 2020, 2, 1–10.
5. R. Lipowsky, *Adv. Biol.*, 2022, 6, 2101020.
6. O. Stauffer, S. Antonia, D. Zhang, J. Csatári, M. Schröter, J.-W. Janiesch, S. Fabritz, I. Berger, I. Platzman and J. P. Spatz, *Biomaterials*, 2021, 264, 120203.
7. L. van der Koog, T. B. Gandek and A. Nagelkerke, *Adv. Healthcare Mater.*, 2021, 2100639.
8. Y. Lee and D. Thompson, *Wiley Interdiscip. Rev.: Nanomed. Nanobiotechnol.*, 2017, 9, e1450.
9. J. Lou and M. D. Best, *Chem. Phys. Lipids*, 2020, 232, 104966.
10. M. Karimi, J. Steinkühler, D. Roy, R. Dasgupta, R. Lipowsky and R. Dimova, *Nano Lett.*, 2018, 18, 7816–7821.
11. R. Dimova, *Annu. Rev. Biophys.*, 2019, 48, 93–119.
12. S. Aden, T. Snoj and G. Anderluh, *Methods Enzymol.*, 2021, 649, 219–251.
13. K. Karamdad, R. Law, J. Seddon, N. Brooks and O. Ces, *Lab Chip*, 2015, 15, 557–562.
14. R. Dimova, P. Stano, C. M. Marques and P. Walde, *The Giant Vesicle Book*, CRC Press, Taylor & Francis, 2020, pp. 3–20.
15. R. V. Stahelin, *J. Lipid Res.*, 2009, S299–S304.
16. M. Dezi, A. Di Cicco, P. Bassereau and D. Lévy, *Proc. Natl. Acad. Sci. U. S. A.*, 2013, 110, 7276–7281.
17. I. L. Jorgensen, G. C. Kemmer and T. G. Pomorski, *Eur. Biophys. J.*, 2017, 46, 103–119.
18. M. Garten, D. Lévy and P. Bassereau, *The Giant Vesicle Book*, Taylor & Francis, 2020, ch. 3.
19. J. A. Bornhorst and J. J. Falke, *Methods Enzymol.*, 2000, 326, 245–254.
20. S. Knecht, D. Ricklin, A. N. Eberle and B. Ernst, *J. Mol. Recognit.*, 2009, 22, 270–279.
21. J. A. Nye and J. T. Groves, *Langmuir*, 2008, 24, 4145–4149.
22. M. Weiss, J. P. Frohnmayer, L. T. Benk, B. Haller, J. W. Janiesch, T. Heitkamp, M. Börsch, R. B. Lira, R. Dimova, R. Lipowsky, E. Bodenschutz, J. C. Baret, T. Vidakovic-Koch, K. Sundmacher, I. Platzman and J. P. Spatz, *Nat. Mater.*, 2018, 89–96.
23. D. Deshpande, Y. Caspi, A. E. Meijering and C. Dekker, *Nat. Commun.*, 2016, 1–9.
24. K. Dürre and A. R. Bausch, *Soft Matter*, 2019, 9676–9681.
25. J. C. Stachowiak, E. M. Schmid, C. J. Ryan, H. S. Ann, D. Y. Sasaki, M. B. Sherman, P. L. Geissler, D. A. Fletcher and C. C. Hayden, *Nat. Cell Biol.*, 2012, 14, 944–949.
26. M. Farell, A. Self, C. Guza, H. Song, L. Apollon, E. W. Gomez and M. Kumar, *ACS Appl. Mater. Interfaces*, 2020, 12407–12416.
27. S. Togo, K. Sato, R. Kawamura, N. Kobayashi, M. Noiri, S. Nakabayashi, Y. Teramura and H. Y. Yoshikawa, *APL Bioeng.*, 2020, 016103.
28. J. Steinkühler, R. L. Knorr, Z. Zhao, T. Bhatia, S. M. Bartelt, S. Wegner, R. Dimova and R. Lipowsky, *Nat. Commun.*, 2020, 11, 905.
29. S. V. Wegner, F. C. Schenk and J. P. Spatz, *Chem. – Eur. J.*, 2016, 22, 3156–3162.
30. S. V. Wegner and J. P. Spatz, *Angew. Chem., Int. Ed.*, 2013, 52, 7593–7596.
31. J. Di Russo, J. L. Young, A. Balakrishnan, A. S. Benk and J. P. Spatz, *Biomaterials*, 2019, 192, 171–178.
32. M. I. Angelova and D. S. Dimitrov, *Faraday Discuss.*, *Chem. Soc.*, 1986, 303–311.
33. M. Abkarian, E. Loiseau and G. Massiera, *Soft Matter*, 2011, 7, 4610–4614.
34. M. C. Blosser, B. G. Horst and S. L. Keller, *Soft Matter*, 2016, 12, 7364–7371.
35. J. P. Reeves and R. M. Dowben, *J. Cell. Physiol.*, 1969, 49–60.
36. A. Weinberger, F. C. Tsai, G. H. Koenderink, T. F. Schmidt, R. Itri, W. Meier, T. Schmatko, A. Schröder and C. Marques, *Biophys. J.*, 2013, 154–164.
37. K. S. Horger, D. J. Estes, R. Capone and M. Mayer, *J. Am. Chem. Soc.*, 2009, 1810–1819.
38. R. B. Lira, R. Dimova and K. A. Riske, *Biophys. J.*, 2014, 107, 1609–1619.
39. A. Witkowska, L. Jablonski and R. Jahn, *Sci. Rep.*, 2018, 8, 9422.
40. E. Parigoris, D. L. Dunkelmann, A. Murphy, N. Will, A. Kaeck, C. Dumrese, N. Jimenez-Rojo and U. Silvan, *Sci. Rep.*, 2020, 10, 1–10.
41. E. M. Schmid, D. L. Richmond and D. A. Fletcher, *Methods Cell Biol.*, 2015, 319–338.
42. J. Steinkühler, P. De Tillieux, R. L. Knorr, R. Lipowsky and R. Dimova, *Sci. Rep.*, 2018, 8, 11838.
43. Y. Okumura and Y. Iwata, *Membranes*, 2011, 1, 109–118.
44. Y. Okumura and S. Oana, *Membranes*, 2011, 1, 345–353.
45. V. Pereno, D. Carugo, L. Bau, E. Sezgin, J. Bernardino De La Serna, C. Eggeling and E. Stride, *ACS Omega*, 2017, 994–1002.
46. J. A. Bellon, M. J. Pino and N. Wilke, *HardwareX*, 2018, 4, e00037.
47. H. G. Behuria, B. K. Biswal and S. K. Sahu, *J. Liposome Res.*, 2020, 255–266.
48. Y. Li, J. Jing, X. Zhang, J. Cao, Y. Li and S. Zhan, *Electrochem. Commun.*, 2012, 25, 151–154.
49. S. Lata, A. Reichel, R. Brock, R. Tampé and J. Pielhér, *J. Am. Chem. Soc.*, 2005, 127, 10205–10215.
50. Y. O. Posokhov, A. Kyrychenko and A. S. Ladokhin, *Anal. Biochem.*, 2010, 407, 284–286.
51. S. S. Tan, S. J. Kim and E. T. Kool, *J. Am. Chem. Soc.*, 2011, 133, 2664–2671.
52. R. Y. Tsien, *Annu. Rev. Biochem.*, 1998, 67, 509–544.
53. A. Lefin, T. R. Molugu, C. Job, K. Beyer and M. F. Brown, *Biophys. J.*, 2014, 107, 2274–2286.
54 G. Shahane, W. Ding, M. Palaiokostas and M. Orsi, *J. Mol. Model.*, 2019, 25, 76.
55 C. Herold, G. Chwastek, P. Schwille and E. P. Petrov, *Langmuir*, 2012, 28, 5518–5521.
56 C. V. Chrysikopoulos and P. Kruger, PhD thesis, Stanford University, 1986.
57 M. A. Malyarick, A. B. Ilyuhin and S. P. Petrosyants, *Main Group Met. Chem.*, 1994, 17, 707–718.
58 T. Biver, R. Friani, C. Gattai, F. Secco, M. R. Tine and M. Venturini, *J. Phys. Chem. B*, 2008, 112, 12168–12173.
59 G. Raghunath and R. B. Dyer, *Langmuir*, 2019, 35, 12550–12561.
60 C. Donley, D. Dunphy, D. Paine, C. Carter, K. Nebesny, P. Lee, D. Alloway and N. R. Armstrong, *Langmuir*, 2002, 18, 450–457.
61 J. A. J. Arpino, P. J. Rizkallah and D. D. Jones, *PLoS One*, 2012, 7, e47132.
Electronic Supplementary Information (ESI)

Binding of His-tagged fluorophores to lipid bilayers of giant vesicles

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This Electronic Supplementary Information contains four Supplementary Figures and seven Supplementary Tables as well as the caption of Movie 1:

Supplementary Figures:

\textbf{Figure S1}: No fluorescence quenching of 6H-FITC in the absence of NTA lipids
\textbf{Figure S2}: Dependence of membrane fluorescence on mole fraction of NTA lipid
\textbf{Figure S3}: Dependence of membrane fluorescence on mole fraction of cholesterol
\textbf{Figure S4}: Design of microfluidic chip

Supplementary Tables:

\textbf{Table S1}: Membrane fluorescence versus molar concentration of 6H-GFP (Figure 6a)
\textbf{Table S2}: Surface coverage of GUV membranes by 6H-GFP (Figure 6b)
\textbf{Table S3}: Membrane fluorescence versus molar concentration of 6H-FITC (Figure 7a)
\textbf{Table S4}: Surface coverage of GUV membranes by 6H-FITC (Figure 7b).
\textbf{Table S5}: Membrane fluorescence intensity of 6H-GFP versus pH (Figure 9a)
\textbf{Table S6}: Membrane fluorescence intensity of 6H-FITC versus pH (Figure 9b).
\textbf{Table S7}: Membrane fluorescence for increased mole fractions of NTA (Figure S2).
Figure S1: Control experiments to demonstrate that the fluorescence of 6H-FITC is not quenched by liposomes which contain a mixture of POPC and cholesterol with a molar ratio of 8:2 but no NTA lipids.
Figure S2: Binding of 6H-FITC to GUV membranes with 3 mol% and 30 mol% DGS-NTA(Ni) anchor lipid: GUVs formed by (a,b) Pt wire electroformation and by (c,d) ITO glass electroformation. All GUVs are exposed to a 120 nM solution of 6H-FITC; and (e) Membrane fluorescence for the four preparation conditions in (a-d). The filled squares represent the mean values of the respective data. The numerical values of these data are given in Table S3. All scale bars are 10 µm.
Figure S3: Influence of cholesterol on the fluorescence intensity of GUV membranes: The GUVs are prepared by platinum wire electroformation and exposed to 120 nM 6H-FITC for two different lipid compositions. In the presence of 20 mol% cholesterol, the intensity is about twice as large as in the absence of cholesterol. This change in intensity can arise by two different mechanisms. First, cholesterol may reduce the fluorescence quenching by the NTA lipids; second it may increase the surface coverage by 6H-FITC.
Figure S4: (Top) Design of microfluidic chip: The device has two main channels parallel to each other. A pipette tip is used as the reservoir for the solution at the inlet. A syringe pump connected to the outlet draws the solution from the reservoir to the outlet via the main channels. The side channels trap the GUVs; and (Bottom) Bright field image showing and magnifying the portion of the chip within the broken red frame. The solution within the side channels is exchanged by diffusion from the main channel. The complete solution exchange including the side channels takes about 3.5 mins as shown in Movie 1.
Table S1: Statistics for the normalised fluorescence intensity of GUV membranes exposed to different molar concentrations \(X\) of 6H-GFP, as obtained via the platinum wire method, corresponding to the data in Figure 6a. The membranes contained 3 mol% NTA lipids and were exposed to a 320 nM solution of 6H-GFP at pH 7.45. The number of GUVs used for the statistics is denoted by \#GUVs and displayed in the 2nd column. The numerical values for the mean value and the standard deviation (SD) of the normalised intensity are dimensionless.

| \(X\) [nM] | \#GUVs | Mean intensity | SD  |
|------------|--------|----------------|-----|
| 0          | 4      | 0.002          | 0.001|
| 20         | 4      | 0.296          | 0.047|
| 40         | 4      | 0.567          | 0.048|
| 80         | 4      | 0.674          | 0.055|
| 120        | 4      | 0.771          | 0.061|
| 160        | 4      | 0.813          | 0.056|
| 240        | 4      | 0.851          | 0.064|
| 320        | 4      | 0.894          | 0.062|

Table S2: Statistics for the surface coverage of GUV membranes by 6H-GFP as obtained for three different preparation methods, corresponding to the data in Figure 6b. The membranes contained 3 mol% NTA lipids and were exposed to a 320 nM solution of 6H-GFP at pH 7.45. The number of GUVs used for the statistics is denoted by \#GUVs and displayed in the 2nd column. The numerical values for the mean value and the standard deviation (SD) of the coverage are given in units of membrane-bound molecules per \(\mu m^2\).

| Preparation method                  | \#GUVs | Mean coverage | SD  |
|-------------------------------------|--------|---------------|-----|
| Pt wire electroformation            | 19     | 53 703        | 4 449|
| PVA hydrogel swelling               | 18     | 31 742        | 9 071|
| ITO glass electroformation          | 18     | 7 261         | 4 582|
Table S3: Statistics for the normalised fluorescence intensity of GUV membranes exposed to different molar concentrations $X$ of 6H-FITC, as obtained by the platinum wire method, corresponding to the data in Figure 7a. The membranes contained 3 mol% NTA lipids and were exposed to a 240 nM solution of 6H-FITC at pH 7.45. The number of GUVs used for the statistics is denoted by #GUVs and displayed in the 2nd column. The numerical values for the mean value and the standard deviation (SD) of the normalised intensity are dimensionless.

| $X$ [nM] | #GUVs | Mean intensity | SD   |
|----------|-------|----------------|------|
| 0        | 3     | 0.014          | 0.004|
| 15       | 4     | 0.347          | 0.088|
| 30       | 4     | 0.585          | 0.050|
| 60       | 4     | 0.707          | 0.184|
| 120      | 4     | 0.748          | 0.085|
| 180      | 4     | 0.822          | 0.127|
| 240      | 4     | 0.818          | 0.075|
| 480      | 4     | 0.834          | 0.027|

Table S4: Statistics for the surface coverage of GUV membranes by 6H-FITC for three different preparation methods, corresponding to the data in Figure 7b. The membranes contained 3 mol% NTA lipids and were exposed to a 240 nM solution of 6H-FITC at pH 7.45. The number of GUVs used for the statistics is denoted by #GUVs and displayed in the 2nd column. The numerical values for the mean value and the standard deviation (SD) of the coverage are given in units of membrane-bound molecules per $\mu m^2$.

| Preparation method           | #GUVs | Mean coverage | SD |
|------------------------------|-------|---------------|----|
| Pt wire electroformation     | 29    | 55 718        | 8 751|
| PVA hydrogel swelling        | 20    | 12 411        | 4 376|
| ITO glass electroformation   | 29    | 13 904        | 8 230|
Table S5: Statistics for the fluorescence intensity of GUV membranes arising from membrane-bound 6H-GFP for different pH values in the exterior solution, corresponding to the data in Figure 9a. The membranes were prepared by platinum wire electroformation with 3 mol% NTA lipids and were exposed to a 20 nM solution of 6H-GFP. The pH was varied by the addition of HCl or NaOH solution. The number of GUVs used for the statistics is denoted by #GUVs. The numerical values for the mean intensity and its standard deviation (SD) are given in arbitrary units.

| pH  | #GUVs | Mean Intensity | SD  |
|-----|-------|----------------|-----|
| 6.38| 5     | 14.8           | 5.9 |
| 6.98| 5     | 36.8           | 18.0|
| 7.25| 5     | 70.1           | 16.9|
| 7.43| 5     | 72.4           | 21.5|
| 8.02| 5     | 80.2           | 24.8|
| 8.54| 5     | 110.6          | 27.0|
| 9.06| 5     | 108.3          | 29.7|
| 9.36| 5     | 114.3          | 29.2|

Table S6: Statistics for the fluorescence intensity of GUV membranes arising from membrane-bound 6H-FITC for different pH values in the exterior solution, corresponding to the data in Figure 9b. The membranes were prepared by platinum wire electroformation with 3 mol% NTA lipids and were exposed to a 120 nM solution of 6H-FITC. The pH was decreased and increased by the addition of HCl or NaOH solution. The number of GUVs used for the statistics is denoted by #GUVs. The numerical values for the mean intensity and its standard deviation (SD) are given in arbitrary units.

| pH  | #GUVs | Mean Intensity | SD  |
|-----|-------|----------------|-----|
| 6.38| 5     | 10.3           | 2.0 |
| 6.98| 5     | 14.0           | 4.0 |
| 7.25| 5     | 21.4           | 2.9 |
| 7.43| 5     | 33.7           | 4.1 |
| 8.02| 5     | 56.8           | 6.7 |
| 8.54| 5     | 76.9           | 7.3 |
| 9.06| 5     | 67.4           | 7.5 |
| 9.36| 5     | 20.7           | 3.0 |

Table S7: Statistics for the fluorescence intensity of GUV membranes exposed to 6H-FITC for different electroformation methods and two different mole fractions of NTA lipids, corresponding to the data in Figure S2. The membranes were prepared with 3 mol% or 30 mol% NTA lipids and exposed to a 120 nM solution of 6H-FITC at pH 7.45. The number of GUVs used for the statistics is denoted by #GUVs. The numerical values for the mean value and the standard deviation of the fluorescence intensity are given in arbitrary units.

| Preparation method    | #GUVs | Intensity, 3 mol% | #GUVs | Intensity, 30 mol% |
|-----------------------|-------|-------------------|-------|--------------------|
| Pt wire electroformation| 17    | 34.4 ± 4.9        | 7     | 42.1 ± 16.1        |
| ITO glass electroformation | 12    | 6.1 ± 7.6         | 6     | 17.8 ± 7.3         |
Movie Caption

**Movie 1:** Time-lapse movie showing the exchange of 2 µM carboxyfluorescein solution to the side channels. The images are obtained by an overlay of the brightfield and the fluorescent channels. Initially, the device is filled with water and then the solution in the reservoir is exchanged with carboxyfluorescein. The pump connected to the outlet is drawing the solution at the speed of 150 µL/hr. The device is imaged close to the outlet. It takes around 3.5 mins to observe homogeneous fluorescence in the two main channels as well as in all side channels.