Response of an actin network in vesicles under electric pulses

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We study the role of a biomimetic actin network during the application of electric pulses that induce electroporation or electropermeabilization, using giant unilamellar vesicles (GUVs) as a model system. The actin cortex, a subjacent network of actin filaments, regulates the shape and mechanical properties of the plasma membrane of mammalian cells, and is a major factor influencing the mechanical response of the cell to external physical cues. We demonstrate that the presence of an actin shell inhibits the formation of macropores in the electroporated GUVs. Additionally, experiments on the uptake of dye molecules after electroporation show that the actin network slows down the resealing process of the permeabilized membrane. We further analyze the stability of the actin network inside the GUVs exposed to high electric pulses. We find disruption of the actin layer that is likely due to the electrophoretic forces acting on the actin filaments during the permeabilization of the GUVs. Our findings on the GUVs containing a biomimetic network provide a step towards understanding the discrepancies between the electroporation mechanism of a living cell and its simplified model of the empty GUV.

The plasma membrane is a selective barrier that separates the intracellular environment from the cell exterior and regulates the influx and efflux of differently sized molecules. The membrane consists of a lipid bilayer with numerous anchored and embedded inclusions, providing the membrane with both fluid and elastic properties; additionally, a network of actin filaments, called the actin cortex, lies immediately underneath the plasma membrane. The bi-directional interplay between the membrane and the actin cortex strongly influences the mechanical response of a cell to external stimuli during diverse biological processes, ranging from cell migration and differentiation to cell division.

The actin cortex also plays an essential role in electroporation, also referred to as electropermeabilization, of cells. This is a membrane permeabilization technique used for the delivery of a wide variety of molecules ranging from small molecules, such as drugs, to large molecules, such as DNA, into the cell. Applying direct current (DC) electric pulses to a cell builds up an induced transmembrane voltage, which can permeabilize the cell membrane above a critical transmembrane voltage (~0.2–1 V). The kinetics of electroporation in cells involves five consecutive steps: (i) nucleation of defects in the membrane, (ii) expansion of the defects, (iii) stabilization of the permeabilized state, (iv) resealing of the permeated membrane, (v) “memory effect” including a full structural recovery of the already resealed membrane. It has been shown that cells can remain permeabilized from minutes to hours after pulsation. This long resealing process is considerably slower than that observed for bare lipid bilayers (less than one second). Based on the evidence from electroporation and electrofusion experiments, it has been suggested that the actin cortex is involved in the electroporation mechanism. However, until now, a complete understanding of the role of the actin cortex in the electroporation of a lipid bilayer is lacking.

There have been several studies on the role of the cytoskeleton during the electroporation of living cells. In these studies, the actin filaments were manipulated by drugs that cause chemical disruption, stabilization of actin monomers, or by genetic engineering of the cytoskeleton. Particularly, it has been shown that actin depolymerization in Chinese hamster ovary (CHO) cells and human erythrocytes leads to acceleration of the post-pulse membrane resealing, while it does not considerably affect the initial steps in the permeabilization process. More recently, nanosecond pulses have gained significant attention due to their ability to permeabilize intracellular membranes. Disruption of the actin cortex in CHO cells appears to make the cells more

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susceptible to nanosecond electroporation\textsuperscript{18}. Also, nanosecond electroporation has been found to be more damaging to non-adherent cultures such as Jurkat cells than adherent cell cultures such as HeLa cells, possibly because of the less extensive cytoskeletal network of the former\textsuperscript{26}. Moreover, some biological processes like apoptosis and necrosis can be triggered by nanosecond pulses\textsuperscript{19,21}, which could involve the pulse-mediated disruption of the actin cytoskeleton\textsuperscript{15,22}.

The studies discussed above illustrate how challenging it is to decouple the mechanical interaction between actin filaments and the membrane from other possible biological processes involving actin, such as apoptosis and necrosis, and to manipulate the cytoskeleton of living cells without side effects like cell death. Therefore, simplified cell models like planar lipid bilayer models\textsuperscript{24–26} or giant unilamellar vesicles (GUVs)\textsuperscript{27–29} have been utilized to study the fundamental mechanisms of pore formation during electroporation. Studies on GUVs have revealed the presence of micrometer sized pores (referred to as macro pores) during electroporation\textsuperscript{2,30–33}. The resealing time of these pores is typically in the order of 10 ms\textsuperscript{30}, as opposed to several minutes in living cells. It has been shown that the dynamics of these pores can be modified by adjusting the edge tension of the lipid bilayer, e.g. by adding cholesterol to increase the edge tension\textsuperscript{34,35}. Moreover, gel-phase lipids have been shown to prevent the formation of macropores due to high surface viscosity\textsuperscript{36,37}. From these studies, it can be concluded that the membrane components are crucial for the dynamics of electroporation or permeabilization in cells. However, GUVs and living cells exhibit drastically different permeabilization behaviors. The size of pores formed due to pulse application in GUVs can reach the micrometer range, and the pores reseal within tens of milliseconds. By contrast, the electropores (or permeated membrane) in living cells have been reported to have sizes of in the order of a few nanometers\textsuperscript{38,39} and to express a slow resealing process which takes minutes up to hours\textsuperscript{38,40,41}, suggesting that the permeabilization dynamics are influenced by intracellular structures such as the actin cortex\textsuperscript{10,12}.

In this research, we focus on the role of a biomimetic actin network in the electroporation of the membrane in order to understand the discrepancy between the electroporation of simplified model membranes and the plasma membrane. To isolate the mechanical role of the actin, we incorporated an actin network inside GUVs (see Fig. 1A,B). We utilized a high-speed imaging technique to reveal the pore formation and resealing of GUVs with and without actin in response to electric pulses. Additionally, we assessed the permeability dynamics of both GUVs after electric pulses. Finally, we used a confocal imaging technique to investigate the structural stability of the encapsulated biomimetic actin network after the electric field.

**Materials and Methods**

**Preparation of the GUVs.** The GUVs (with and without actin filaments) were prepared by electroswelling, based on a procedure utilized by Schäfer et al.\textsuperscript{22}. The following membrane composition was used for GUVs in the bright field experiments: 95 mol% 1,2-dioleoyl-sn-glycero-3-phospholine (DOPC) and 5 mol% Ca\textsuperscript{2+}-ionophore (A23187). For the confocal experiments, the membrane was fluorescently labelled using the following membrane composition: 94.5 mol% DOPC, 5 mol% Ca\textsuperscript{2+}-ionophore and 0.5 mol% 1,2-dioleoyl-sn-glycero-3-phosphoethanolamine-N-(lissamine rhodamine B sulfonyl) (ammonium salt) (DOPE-RhoB, ex/em 560 nm/583 nm). The lipids were purchased from Avanti Polar Lipids, Inc. and the Ca\textsuperscript{2+}-ionophore from Sigma Aldrich, and these compounds were stored at −20 °C. All membrane components were dissolved separately in chloroform at a concentration of 1 mg/ml, stored under nitrogen at −20 °C, and mixed prior to the experiments. 10 μl of lipid mixture was deposited on the conductive face of two Indium Tin Oxide (ITO) slides (purchased from Sigma Aldrich) and left under vacuum for more than 2 hours. Afterwards, the ITO slides were inserted in a teflon swelling chamber with the conductive sides facing each other, spaced 1.5 mm apart. 335.6 μl swelling buffer (pH of ~8) was inserted in between the ITO slides, containing 200 mM sucrose, 2 mM Tris–HCl (Thermo Fischer), 0.17 mM Adenosine 5’-triphosphate disodium salt hydrate (Na\textsubscript{2} ATP) and 0.21 mM dithiothreitol (DTT) (both purchased from Sigma Aldrich). In the case of the GUVs with encapsulated filaments, monomeric actin from rabbit muscle (Thermo Fischer), purchased as 1 mg of lyophilised powder, was dissolved in 2 mM Tris–HCl (pH 8.0) to get a stock solution of 2 mg/ml and stored at −80 °C. For visualization of the filaments, fluorescently labeled actin monomers with Alexa Fluor 488 (Thermo Fischer, ex/em 495 nm/519 nm) were purchased as a solution with a concentration of 2 mg/ml and stored at −80 °C. For visualization of the filaments, fluorescently labeled actin monomers with Alexa Fluor 488 (Thermo Fischer, ex/em 495 nm/519 nm) were purchased as a solution with a concentration of 2 mg/ml and stored at −80 °C.
in the range of 3–5 mg/ml in a buffer (5 mM Tris (pH 8.1), 0.2 mM DTT, 0.25 mM CaCl₂, and 0.2 mM ATP) containing 10% (w/v) sucrose. A solution of 7 μM actin monomers, of which 0.5 μM was fluorescently labeled, was added to the swelling buffer. An alternating current (AC) was applied at 70 Hz, 2.4 V for 3 hours at room temperature (around 20 °C), using the Agilent 33220 A 20 MHz Function/Arbitrary Waveform Generator.

Polymerization and binding of actin to the membrane. After the swelling of the GUVs and encapsulation of 7.5 μM actin monomers, actin polymerization at the membrane was initiated. For both the polymerization and the binding of the filaments to the membrane, Mg²⁺ ions were used as shown in Fig. 1B. The GUV solution obtained by electroswelling was diluted about 5 times with 1.3 ml glucose buffer of 200 mM glucose and 2 mM Tris-HCl (pH of ~8) in order to dilute the actin monomer concentration and the sucrose concentration on the outside of the GUVs. 163 μl of 67.3 mM MgCl₂ solution (pH of ~8.5) was then added to increase the MgCl₂ concentration to 6 mM. Subsequently, due to the presence of the Ca²⁺-ionophore in the membrane, the Mg²⁺ ions were transported into the GUVs to initiate polymerization and bind the actin filaments to the membrane as a thin cortical actin shell through electrostatic interactions. Afterwards, the GUVs were left overnight at room temperature at 6 mM MgCl₂ and 1.55 μM actin concentration on the outside of the GUVs. Prior to the electroporation experiments, the GUVs were diluted ten times more with 1 ml of the 200 mM glucose buffer, in order to sediment the GUVs and enhance the optical contrast between the solutions inside and outside the GUVs. Afterwards, the MgCl₂ concentration in the outer solution was increased to 19 mM to ensure the formation of an actin shell at the membrane by the addition of 412.4 μL of 67.3 mM MgCl₂. In addition, this dilution further reduced the concentration of the actin outside the GUVs to 0.04 μM. In our hands, this procedure led to the most consistent results, with a large majority of GUVs containing an actin network (>90%) (see Fig. S.1 in the Supplementary Information). Since the ion concentration has a large impact on the response of the GUVs to electric pulses, altering the deformations of the GUVs and the pore dynamics, we fixed the concentration of MgCl₂ at 19 mM. The stability of the GUVs was not affected by the high ion concentration overnight. The absence of a detectable actin network on the outside of the GUV during the storage overnight and during the electroporation experiments was confirmed by control experiments, where empty GUVs were incubated with 1.55 μM and 0.04 μM actin (see Fig. S.2 in the Supplementary Information). Note that all of our data for electroporation were obtained at a dilute concentration of actin outside of the GUVs (C₀ = 0.04 μM), which is far below the critical concentration to form actin filaments.

Electroporation setup combined with high speed imaging. To image the dynamics of the membrane during an electric pulse, the GUVs were visualized by an inverted microscope (Zeiss Axio Observer.Z1) equipped with a Phantom v9.1 high-speed camera (10,000–15,000 frames per second, Vision Research Inc.) with a 40x oil immersion objective (Ph3, Plan-Neofluar, 40x/1.30), providing a pixel size of 0.3 μm. Prior to the experiments, the imaging chamber was treated with 5 g/l Bovine Serum Albumin (BSA) solution (purchased from Sigma Aldrich) for 20 minutes, to avoid rupture of the GUVs upon contact with the glass at the bottom of the chamber. Custom-made stainless-steel 20 mm-long electrodes with 1 mm distance were submerged in 1 ml sucrose-filled GUVs with glucose solution on the outside assure sufficient optical contrast to image the GUVs in bright field, and indicate permeabilization of the membrane by a contrast loss. Consecutive pulses with increasing amplitudes were applied. The electric pulses induce a transmembrane voltage as follows:

\[ \Psi_m = 1.5 RE(1 - \exp(-t/\tau_e)), \]  

where R, E and t indicate the radius of the GUV, the applied electric field strength and the duration of the pulse, respectively. \( \tau_e \) is the charging time of the membrane given by

\[ \tau_e \approx RC_m(1/\lambda_e + 1/\lambda_i), \]  

where \( C_m \), \( \lambda_e \) and \( \lambda_i \) are the membrane capacitance, and the external and internal solution conductivity, respectively. In our experiments the pulses were chosen such that the induced transmembrane voltage started from about 1 V, with approximately 1 minute interval between the pulses to minimize the effect of the former pulse. The sucrose-filled GUVs with glucose solution on the outside assure sufficient optical contrast to image the GUVs in bright field, and indicate permeabilization of the membrane by a contrast loss. Consecutive pulses with increasing amplitudes were applied until a complete contrast loss of the GUVs was observed.

Electroporation setup for pore dynamics. To assess the resealing of the membrane, the transport of sulforhodamine B through the membrane was monitored. Sulforhodamine B was added to the outer solution at a concentration of 2.5 μM. The same pulse protocol was used as during the high-speed imaging experiments. Dye transport was captured with a Zeiss LSM 710 inverted confocal microscope, using a 100x oil immersion objective (Ph2, Achromplan, 100x/1.25 Oil) and a sampling frequency of 1.5 Hz. To obtain the uptake kinetics of the sulforhodamine B, the intensity of 80% of the inner GUV area was monitored, to exclude the increased intensity of the membrane. In addition, the intensity of the background was captured. The data was normalized as follows:

\[ I_{peak,t} = \frac{I_{peak,t} - I_{dye,0}}{I_{background,t} - I_{dye,0}}, \]  

where \( I_{peak,t} \), \( I_{background,t} \) and \( I_{dye,0} \) represent the normalized intensity inside the GUV at time t, the measured intensity inside the GUV at time t, the initial intensity prior to the pulse and the intensity of the background, respectively.
Electroporation chamber combined with confocal imaging. The response of the actin cortex to the electric pulses was visualized with a Zeiss LSM 710 inverted confocal microscope, using a 100x oil immersion objective (Ph2, Achromplan, 100x/1.25 Oil) to image both the lipid bilayer and the actin cortex. Two different pulsing protocols were used: (i) consecutive gradually increasing pulses, ranging from 10–300 V/mm with steps of approximately 10 V/mm, and (ii) immediate high pulses, ranging from 300–1000 V/mm with steps of approximately 200 V/mm, where only 2 to 4 pulses were applied. In both cases, 500 μs pulses were used, separated by approximately 1 minute. The first pulse protocol (i) was used to test the onset of the breakdown of the actin shell. The second pulse protocol (ii) was used to decouple the breakdown of the actin shell from the cumulative effect of the multiple consecutive pulses. A z-stack of 7 focal planes, with a total range of approximately the diameter of the GUV (the distance between planes is Δz ~ 4 μm), of the fluorescence signal of the membrane and the actin was taken before and after the pulses. The z-stack imaging is a slow technique (approximately 5 to 10 seconds time interval in between the z-stacks), so only the GUV size and the actin network stability before and after the pulse were extracted.

Data analysis of GUVs response. The bright field images of the GUVs were analyzed by a custom written Matlab script (The Mathworks Inc., Natick, MA, USA). The contours of the GUVs were detected with the Canny edge detection method and fitted to an ellipse to extract the equatorial (a) and polar (b, the distance from the center to the poles of a spheroid) radii of the GUVs. For the study of dynamics, these two radii were used to obtain the deformation ratio (a/b) during and after the pulse. Afterwards, the time-dependent deformation was fitted to an exponential decay curve to obtain the relaxation time(s) of the GUVs30 (Fig. 2).

During the confocal experiments, multiple planes of the GUVs were imaged. The radius of the GUVs was determined from the fluorescent confocal membrane images taken at the equatorial plane of the GUV by a similar detection method as described above. These images were used to obtain the area loss of the GUVs as a function of the electric field. In order to remove spurious effects due to motion of the GUV in and out of the imaging plane (which could change their observed radius), the normalized area (Anorm) was calculated at pulse n and corrected for the area loss at the previous pulse (n−1): Anorm = (Ai/Af) × (Af/Af)−1, Here, Ai and Af represent the projected areas before (initial) and after (final) the pulse, respectively. (Ai/Af)−1 and (Ai/Af)n−1 are the normalized areas of the GUV at the pulse n and at the preceding pulse (n−1). By multiplying these two factors, the normalized area was corrected for shrinkage due to earlier pulses. Additionally, the intensity of the fluorescence signal of the actin was obtained from confocal z-stacks by determining the mean and average intensity of all planes. Photobleaching of
the actin shell was removed by correcting the data at scan number \( k \) with a reference of a separate photobleaching experiment of a GUV without applying any pulses. By normalizing the actin fluorescence intensity from the bleaching experiment, the correction factor for the photobleaching per scan is calculated as \( I_{\text{corr}} = I_{\text{ref}}/I_{\text{bleach}} \), where \( I_{\text{ref}} \) and \( I_{\text{bleach}} \) are the intensities of the image at the start of the photobleaching experiment and at the relevant scan number \( k \), respectively. Consequently, the results of the electroporation experiments were corrected as follows: \( I_{\text{corr}} = I_k/I_{\text{bleach}} \). Again, \( I_k \) and \( I_{\text{bleach}} \) represent the intensities of the image at the start of the experiment and at scan number \( k \), respectively, in this case during an electroporation experiment (see Fig. S.3 in the Supplementary Material).

**Results and Discussion**

**The dynamics of the GUVs.** To investigate the influence of the actin network on the electroporation of GUVs, we captured the dynamics of the GUVs by high speed imaging (10,000–15,000 fps) in bright field. The deformations of the GUVs were measured in terms of the deformation ratio \( a/b \) during and after the electric pulse (see the schematic in Fig. 2). The GUVs become electroporated once the accumulated transmembrane voltage exceeds a threshold voltage. Electroporation can be recognized by the observation of a macropore or by the contrast loss due to the exchange of glucose and sucrose molecules between the interior and exterior of the GUV. Below the critical threshold field, where no macropores nor contrast loss were observed, the GUVs exhibited squarelike deformations, as shown in Fig. 2A. These deformations relaxed back exponentially with a characteristic relaxation time \( \tau_1 \) in the order of 100 \( \mu \)s for both the empty and the actin-encapsulated GUVs. Riske and Dimova previously showed that the presence of ions (NaCl, Ca\(^{2+}\) or Mg\(^{2+}\) acetates) flattens the membrane and causes squarelike deformations, which were attributed mainly to the electrohydrodynamic forces of the ions.

The conductivity ratio \( \chi \) is defined as the ratio between the conductivity of the internal \( \lambda_{\text{in}} \) and the external \( \lambda_{\text{out}} \) solution of the vesicle \( \chi = \lambda_{\text{in}}/\lambda_{\text{out}} \). Depending on this conductivity ratio, vesicles are expected to either deform along the field, referred to as a tubelike shape, (deformation ratio \( a/b > 1, \chi > 1 \)) or perpendicular to the field, referred to as a disklike shape, (deformation ratio \( a/b < 1, \chi < 1 \)). We observed both disklike and tubelike deformations during the pulsing experiments of both the empty and the actin-encapsulated GUVs. An ionophore was used to transport Mg\(^{2+}\) into the GUVs to polymerize actin and construct the biomimetic cortex after the GUV formation, as discussed in the methods section. Some variation in the conductivity of the inner solution of the GUVs is possible due to our preparation method, which would explain the observations of both disklike and tubelike deformations. Additionally, permeabilizing pulses cause an exchange of the solutions between inside and outside the vesicles, and consequently can alter their conductivities. As these observations are similar for both the empty and actin-encapsulated GUVs, they can be related to the presence of the ionophore and the ionimbalance of the inner and outer solutions.

The response of the empty and the actin-encapsulated GUVs to an electroporative pulse was strikingly different, as shown in Fig. 2. Macropores were observed for the empty GUVs, as was reported before, whereas the actin-encapsulated GUVs exhibited no visible macropores. Additionally, for the empty GUVs we observed a characteristic relaxation process, where the relaxation of the macropore (with a duration \( \tau_2 \) of ~1 ms) was often followed by another relaxation event (with a duration \( \tau_3 \) of ~10 ms), as illustrated in Fig. 2B and Movie S1. This response agrees with previous studies on empty GUVs composed of egg phosphatidylcholine by Riske and Dimova. They attributed \( \tau_2 \) to the relaxation time of the macropore in a standalone lipid bilayer \( \tau_2 = \eta_p/\lambda_p \), where \( \eta_p, \lambda_p, \gamma \), and \( r \) are the surface viscosity, the line energy per unit length and the pore radius, respectively.

To investigate the influence of the actin network on the electroporation of GUVs, we captured the dynamics of the GUVs by high speed imaging. In a clear contrast to the empty GUVs, the actin-encapsulated GUVs showed only small deformations during the electroporative pulses and did not exhibit macropores (Fig. 2C and Movie S2). Since no macropore was observed for these GUVs, electropermeabilization was determined by the contrast loss of the GUVs. This contrast loss indicates the permeabilization of the membrane, due to the exchange of molecules through the membrane. Due to the limited recording time of the high-speed imaging in the experiments (maximum of 2 seconds), no long-term dynamics of the contrast loss could be captured using this setup.

The relaxation time \( \tau_2 \) characterizes the time-dependent deformation of the actin-encapsulated GUVs in the electroporative regime, is comparable to the relaxation time \( \tau_1 \) of the empty GUVs (Fig. 3A). However, the slow dynamics of the contrast loss after a pulse for the actin-containing GUVs indicates that the membrane remains permeabilized up to minutes, which was not observed for the empty GUVs. Therefore, the pore resealing time of the actin-encapsulated GUVs cannot be deduced from \( \tau_2 \). Previous studies on living cells have proposed that the formed electropores in the membrane cannot expand beyond the mesh size of the actin network. Possibly, a similar mechanism limits the pores in the actin-encapsulated GUVs.

As discussed above, we observed both disklike (\( a/b < 1 \)) and tubelike (\( a/b > 1 \)) deformations for the empty and the actin-encapsulated GUVs. For both types of GUVs, as shown in Fig. 3B, the majority of vesicles underwent a tubelike deformation, indicating a higher conductivity of the interior of the GUVs due to the Mg\(^{2+}\) transport across the membrane by the ionophore. The deformations of the actin-encapsulated GUVs are significantly smaller than those of the empty GUVs, which can be seen from the more narrow distribution of \( (a/b)_{\text{max}} \) (shown in red in Fig. 3B). Experiments on GUVs in alternating current (AC) fields have shown that the extent of the membrane deformation depends on the bending rigidity of the membrane. Additionally, in the DC field experiments, non-electroporative pulses can be used to study the stiffness of the membrane. In our experiments we focused on pulses around the transmembrane voltage of 1 V, therefore we cannot derive the precise mechanical properties. Nevertheless, the smaller deformations of the GUVs with an encapsulated actin shell clearly signify an increase in the bending rigidity of the GUVs, consistent with previous studies employing either AFM-indentation, micropipette aspiration, or flicker spectroscopy. The wide distribution observed for \( (a/b)_{\text{max}} \) of the GUVs is caused...
Dye leakage mainly occurs through macropores. Similarly, we observed macropore formation for the empty GUVs immediately after the pulse is defined as the point where electroporation occurs. To find the amount and characteristic time of dye uptake after the pulse, we use the following fitting equation:

\[ I_\text{uptake} = I_\infty + (I_0 - I_\infty) \exp(-t/\tau_\text{uptake}), \]

where \( I_\infty, I_0 \), and \( \tau_\text{uptake} \) represent the final sulforhodamine B intensity of the GUV, the initial sulforhodamine B intensity of the GUV prior to the pulse and the characteristic time of the uptake of the dye molecules. The two fitting parameters \( I_\infty \) and \( \tau_\text{uptake} \) for the actin-encapsulated GUVs are larger than for the empty GUVs. Employing a high temporal resolution technique, a former study on the dye leakage of GUVs showed that the transport after the pulse could be captured, which is predominantly diffusive. The applied pulses are in the same range where macro-pores were observed for the empty GUVs. An increase in the inner fluorescence of the GUVs immediately after the pulse is defined as the point where electroporation occurs. To find the amount and characteristic time of dye uptake after the pulse, we use the following fitting equation:

\[ I_\text{uptake} = I_\infty + (I_0 - I_\infty) \exp(-t/\tau_\text{uptake}), \]

Resealing of the permeabilized membrane. The permeabilization dynamics of the membranes was determined by the uptake of sulforhodamine B molecules into the GUVs. Due to the relatively slow imaging, only the transport after the pulse could be captured, which is predominantly diffusive. The applied pulses are in the same range where macro-pores were observed for the empty GUVs. An increase in the inner fluorescence of the GUVs immediately after the pulse is defined as the point where electroporation occurs. To find the amount and characteristic time of dye uptake after the pulse, we use the following fitting equation:

\[ I_\text{uptake} = I_\infty + (I_0 - I_\infty) \exp(-t/\tau_\text{uptake}), \]

where \( I_\infty, I_0 \), and \( \tau_\text{uptake} \) represent the final sulforhodamine B intensity of the GUV, the initial sulforhodamine B intensity of the GUV prior to the pulse and the characteristic time of the uptake of the dye molecules. The two fitting parameters \( I_\infty \) and \( \tau_\text{uptake} \) for the actin-encapsulated GUVs are larger than for the empty GUVs. Employing a high temporal resolution technique, a former study on the dye leakage of GUVs showed that the dye leakage mainly occurs through macro pores. Similarly, we observed macro pore formation for the empty GUVs, which was accompanied with ~2% uptake of dye molecules (~0.02I_\infty) after a single pulse and a characteristic uptake time of approximately 12 seconds. Taken together, these observations indicate that most of the dye transport occurs during the pulse and shortly after the pulse. It must be noted that earlier studies have shown resealing of the membrane in 10 ms, which is faster than our temporal resolution. Therefore, the actual characteristic uptake time might be considerably smaller than 12 seconds, although the presence of the ionophore in the membrane may slow down membrane resealing. The actin-encapsulated GUVs, on the other hand, do not show any macro pores and exhibit a longer characteristic uptake time of approximately 146 seconds together with a dye uptake of ~38%. Our observations are reminiscent of prior observations of a high and long-lived permeability for lipid membranes associated with an agarose mesh work that was shown to stabilize the pores formed by electric pulses. The actin network in our experiments appears to affect the membrane stability in a similar fashion, enabling transport through the membrane for a longer duration than for the empty GUVs.

Despite the difference between the structure and organization of our biomimetic cortex and the actin cortex of living cells, similarities can be found between their response to electric pulses. The exponential increase in the dye uptake of actin-encapsulated GUVs appears similar to the exponential uptake of living cells. The presence of an actin network is known to increase the membrane tension, and the enhanced tension likely explains the slow resealing process of permeabilized structures in actin-encapsulated GUVs, compared to empty ones. A slow resealing process of the membrane is consistent with observations in electroporated cells, where the actin cortex influences the resealing of the permeabilized structures.

Stability of the actin network. To assess the response of the actin network inside the GUV to electric pulses, the membrane and actin were concurrently visualized by confocal microscopy. The fluorescence signals of the membrane and actin network were used to detect the projected area of the GUV and the actin intensity, respectively. To determine the coverage of the actin network...
on the inner membrane surface and any possible lateral inhomogeneities of the shell, a confocal z-stack was taken before and after the pulse. Consecutive pulses of increasing voltage were applied to the GUVs, with at least 1 minute in between the pulses to minimize the effect of the former pulse. Similar to earlier studies, the empty GUVs exhibit increasing shrinkage with higher electric pulses (Fig. 5A). Strikingly, a single pulse of 300 V/mm does not induce the same shrinkage as a 300 V/mm pulse preceded by multiple pulses at lower field strength (see Fig. 5A). Thus, multiple pulses exert a cumulative effect on GUV shrinkage. Compared to the empty GUVs, the actin-encapsulated GUVs display shrinkage at higher electric fields, indicating that the actin-supported bilayer has a higher electrical stability (Fig. 5B). This could be caused by the increased surface viscosity of the bilayer due to the presence of the actin layer.

Strikingly, the increasing pulses also result in a decreasing fluorescence signal of the actin at ~150 V/mm, followed by GUV shrinkage at higher electric pulses (~200 V/mm), as illustrated in Fig. 5B and Movie S3. We observe that the decrease in the fluorescence signal of the actin is associated with a breakdown of the actin network. Additionally, the intensity loss of the actin layer due to the electric pulses appears to be slightly more at the poles, where the GUV is facing the electrodes (Movie S3). However, this radial dependence is not always observed clearly, possibly due to GUV’s rotation. After the breakdown of the actin network, we did not observe any increase in the intensity in the center of the GUV (see Section S.5 and Fig. S.4 in the Supplementary Material). By recording the photobleaching of the actin signal in the absence of the pulse and also the response of GUVs with an encapsulated actin network to immediate high pulses, we could confirm that the fluorescence decrease is caused by the breakdown of the network and is not a side effect of bleaching. This notion is further supported by the observation that after the pulse, the actin fluorescence signal progressively decreased for several tens of seconds. This gradual decrease in the signal indicates a slow breakdown of the actin network. In addition, the shrinkage of the actin-encapsulated GUVs at higher electric fields, compared to empty ones, indicates that initially the actin network stabilizes the membrane and that shrinkage only sets in after the breakdown of the network. Our observations are reminiscent of the depolymerization of the actin cortex that has been observed for human cell lines exposed to nanosecond pulsed electric fields (nsPEF). Different mechanisms have been proposed to account for actin depolymerization, including a direct effect of nanosecond pulses or osmotic swelling or biological processes. In the GUV model system, we can exclude biochemical processes and in addition no osmotic swelling was observed in our experiments. The only factors that may cause actin depolymerization are the mechanical forces, originating from the induced electrical membrane stress during the pulse, as well as the electrophoretic forces acting on the actin filaments as soon as the membrane is permeabilized.
We, therefore, estimate the magnitude of both the mechanical and the electrophoretic forces, considering the actin filaments as semiflexible polymers. For simplicity, we ignore the presence of actin bundles (Fig. S.1 in the Supplementary Information). Assuming a uniform distribution of actin filaments over the membrane surface, the total number of filaments underneath the lipid bilayer can be estimated as:

\[ N_f \sim \frac{c_0 \times \frac{4}{3} \pi R^3 \times d}{l} \]  

(4)

where \( c_0 \) and \( d \) are the initial concentration \((4.2 \times 10^{21} \text{ m}^{-3})\) and size of actin monomers, and \( l \) is the average length of the filaments. Assuming the filaments have a length in the order of \( l \sim 4 \mu \text{m} \) (see Fig. S.1 in the Supplementary Material) and with \( R = 10 \mu \text{m} \) and \( d = 2.5 \text{ nm} \), we estimate the total number of filaments per GUV to be \( N_f \sim 10^4 \). The mesh size of the filament network can be approximated as:

\[ \zeta \sim \frac{A}{N_f} \]  

with \( A \) being the surface area of the membrane. The amount of stretch imposed on the length of connections in the actin network reads:

\[ \zeta = \frac{\Delta \ell}{l_0} \]  

As the total enclosed volume of the GUVs is conserved during their deformation into ellipsoidal topologies in the experiments, the equatorial and polar radii become \( b = e^{-1/3} R \) and \( a = e^{2/3} R \), respectively, where \( e = \sqrt{a/b} \) is taken from the example GUV shown in the schematic in Fig. 2E. We choose a value of \( e \sim 1.18 \), corresponding to the maximum deformation experienced by the GUVs. The total surface area of the GUVs increases during the deformation and hence the concomitant mesh size stretches by \( \Delta \zeta \sim 0.8 \text{ nm} \) (if we assume affine deformations for the network). Such an increase in the length of interconnected filaments induces a maximum mechanical force of the order of \( f_{\text{mech}} \sim 34 \text{ pN} \) for a stretching stiffness of \( K \sim 48 \text{ pN/\text{nm}} \) \( ^{59,60} \). This value is markedly smaller than the force needed for either initiating the depolymerization of a filament network \( ^{61} \) or the rupture of single filaments \( ^{62} \), which is in the range of \( 100-400 \text{ pN} \). It is, therefore, unlikely that the mechanical forces generated by in-plane tensions are the only origin for the breakdown of the actin network in our experiments. Other mechanisms, including electrophoretic forces, are hence expected to be involved.

As soon as the membrane is permeabilized by the electric fields, the membrane tension can relax back through the expansion of the pores and the release of the interior fluid. Additionally, upon applying an electric field on any charged molecules in a bulk solution, they experience a driving force. This effective force can drive and direct the motion of a free filament in the bulk fluid \( ^{63} \). In contrast, when entangled and hindered from motion in a network (which is the case for our actin shell), the filaments undergo mechanical forces between their constituent monomers. The force acting on the filaments in the shell due to the electric field is defined as \( ^{63} \):
length of ~4 μm for the actin filaments and considering that the whole filament interacts with the membrane, due to Mg2+–mediated adhesion, the force per unit length can be converted to the electrophoretic force \( f_{\text{electrophoretic}} \). The maximum force experienced by the actin filaments corresponds to a condition in which the filaments are perpendicular to the field. As soon as the GUV is permeabilized and pores are formed, the electric field penetrates into the GUV, with a maximum estimated value of ~0.8E at the poles where the GUV is facing the electrodes (see Fig. S.7 in the Supplementary Material). The fluorescence signal of the actin network drops at approximately 150 V/mm (Fig. 5A). At this electric field and considering a hydrodynamic friction of \( \zeta = 0.034 \text{ N.s/m}^2 \) (for cytoplasmic fluid motion perpendicular to the filament length) and an electrophoretic mobility of \( \mu_E = 10^{-8} \text{ m}^2/(\text{V.s}) \), we predict an electrophoretic force of \( f_{\text{electrophoretic}} \approx 160 \text{ pN} \) acting on a single filament for a vesicle size of 10 μm. Compared to the mechanical forces calculated above, these forces appear most plausible to initiate the disruption of the actin network. Importantly, the generated heat due to Joule heating is estimated to be small (less than 3 K) in our experiments (see Section S9 in the Supplementary Material). Moreover, the disruption of the network mostly occurs above the critical transmembrane voltage (see Fig. S.8 in the Supplementary Material), enabling the electrophoretic forces to affect the actin filaments.

**Conclusion**

To shed light on the role of the actin cortex on the electroporation of cells, we prepared GUVs with a subjacent attached actin network and exposed them to electric pulses. Time-lapse imaging of the membrane revealed that membrane rigidification by the actin network inhibits large deformation of the GUVs during the electric pulses. Additionally, we found that the actin layer prevents the formation of large pores, referred to as macropores. Finally, we observed that the membrane resealing after pulse application takes significantly longer in the presence of the actin shell than for a bare membrane. Interestingly, time-lapse imaging of the actin revealed that, at higher electric fields, the electric pulses cause disruption of the actin shell. Based on the estimation of the relevant forces, we suggest that the actin shell disruption is predominantly triggered by the electrophoretic forces on the actin filaments during the pulses.

Although our biomimetic cortex is still far from mimicking the actual actin cortex of a living cell, the behavior of the actin-encapsulated GUVs already shows the importance of equipping the current membrane models with other components present in the plasma membrane. Our results provide a step towards understanding the major differences between the electroporation of living cells and GUVs such as (lack of) macropore formation and the resealing dynamics of the defects. Consequently, these actin-supported GUVs enable exploring more complex processes, such as the mechanism of electro-gene transfection.

**Data Availability**

The datasets generated and analyzed during the current study are available from the corresponding author on reasonable requests. This manuscript is part of the PhD thesis of DLP65. A digital version of this manuscript is available as a preprint on bioArxiv66.

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**Author Contributions**

P.E.B. and D.L.P. designed the experiments. D.L.P., V.K. and L.S. conducted the experiments. A.V. and L.R. did the theoretical analysis. P.E.B., D.L.P., A.V., M.T.K., Y.M., A.M. and G.H.K. contributed to the analysis and the interpretation of the data. D.L.P. and A.V. wrote the manuscript. P.E.B., D.L.P., A.V., L.R., M.T.K., Y.M. and G.H.K. reviewed the manuscript.

**Additional Information**

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