Directed Growth of Biomimetic Microcompartments

Ivan Ivanov, Rafael B. Lira, T.-Y. Dora Tang, Titus Franzmann, Adam Klosin,
Lucas Caire da Silva, Anthony Hyman, Katharina Landfester, Reinhard Lipowsky,
Kai Sundmacher, and Rumiana Dimova*

Contemporary biological cells are sophisticated and highly
compartmentalized. Compartmentalization is an essential principle of
prebiotic life as well as a key feature in bottom-up synthetic biology research.
In this review, the dynamic growth of compartments as an essential
prerequisite for enabling self-reproduction as a fundamental life process
is discussed. The micrometer-sized compartments are focused on due
to their cellular dimensions. Two types of compartments are considered,
membraneless droplets and membrane-bound microcompartments.
Growth mechanisms of aqueous droplets such as protein (condensates) or
macromolecule-rich droplets (aqueous two phase systems) and coacervates
are discussed, for which growth occurs via Ostwald ripening or coalescence.
For membrane-bound compartments, vesicles are considered, which are
composed of fatty acids, lipids, or polymers, where directed growth can occur
via fusion or uptake of material from the surrounding. The development
of novel approaches for growth of biomimetic microcompartments can
eventually be utilized to construct new synthetic cells.

1. Introduction

Despite the long and ongoing debate about the essence of life,
there is a widespread consensus about some common features
of all living systems. While some phenomena, such as metabolism, remain arguable benchmarks (and place viruses in the
twilight zone), it is widely recognized that
the formation of micro- and nanocompartments is an essential ingredient of
all forms of life. This spatial constraint
serves numerous purposes, including
the segregation and protection from the
environment (to allow for individuality
and maintenance), the establishment of
gradients (to enable and make use of out-of-equilibrium conditions), and the role
of 2D and 3D confinement for self-organization phenomena, all of which serve to
overcome the overall "dilution problem."
In order to fulfill another cornerstone
of life—proliferation—compartments
also need to change with time by fusion,
growth, and division. In particular, the
dynamic growth of compartments is an
essential prerequisite for enabling self-reproduction as a fundamental life process, both in simplistic systems such as
droplets or fatty acid–based vesicles, as
well as for lipid vesicle compartments with membranes that
resemble the biomembranes of today's cells. In this context, we
argue that growth deserves more attention, not only because
growth precedes division but also because of the difficulty to
realize growth compared to division, especially in the case of
lipid vesicles, where budding and division has been observed
in response to various factors. This growth aspect has also been
recognized in basic theoretical models of living systems such
as Ganti's chemoton,[1,2] for which the increase of membrane
area (referred to as membrane formation) was postulated to be
one of three subsystems, characterizing living entities from a
chemical viewpoint. The autopoietic theory was also centered
on the membrane but focused on self-maintenance rather than
on (self-)reproduction.[3,4] However, such a self-maintaining
system could also enter a self-reproduction mode, manifested
by growth, if homeostatic misbalance leads to excess membrane formation as shown in a thought experiment by Luisi.[3]

In this progress report, we review the existing approaches
for growth of compartments in the context of bottom-up synthetic biology and protobiology. We consider mainly micrometer-sized compartments due to their characteristic cellular
dimensions; this feature also ensures that the area and curvature of the interface or the bounding membrane has less influence on the enclosed solution. Microcompartments of various
origins and chemistries have been used as protocell models,
and many studies have addressed growth and division simultaneously in an ambitious effort to mimic self-reproduction.

Dr. I. Ivanov, Prof. K. Sundmacher
Max Planck Institute for Dynamics of Complex Technical Systems
Process Systems Engineering
Sandtorstrasse 1, 39106 Magdeburg, Germany
E-mail: ivanov@mpi-magdeburg.mpg.de
Dr. R. B. Lira, Prof. R. Lipowsky, Dr. R. Dimova
Max Planck Institute of Colloids and Interfaces
Theory and Bio-Systems
Science Park Golm, 14424 Potsdam, Germany
E-mail: dimova@mpikg.mpg.de
Dr. T.-Y. D. Tang, Dr. T. Franzmann, Dr. A. Klosin, Prof. A. Hyman
Max Planck Institute of Molecular Cell Biology and Genetics
Organization of Cytoplasm & Dynamic Protocellular Systems
Pfotenhauerstrasse 108, 01307 Dresden, Germany
Dr. L. Caire da Silva, Prof. K. Landfester
Max Planck Institute for Polymer Research
Physical Chemistry of Polymers
Ackermannweg 10, 55128 Mainz, Germany

Here, we focus on growth and divide the literature examples into two categories: membraneless and membrane-bound microcompartments, whereby we review aqueous droplets (such as protein- or macromolecule-rich droplets and coacervates) on one side, and vesicles, composed of fatty acids, lipids, or polymers, on the other. This is by no means a universal and comprehensive classification of compartment types in nature,[5] but rather an attempt to classify reported examples.

2. Coacervates and Protein-Rich Droplets: Volume and Area Grow Simultaneously

In the 1920s Oparin hypothesized that membrane-free liquid-like droplets formed via complex coacervation could have been precursors to modern cells.[6,7] Extensive research has demonstrated that small molecules and oppositely charged polymers can phase separate to form structures with emerging liquid-like properties. More recently, it was discovered that certain types of proteins undergo intracellular phase separation to form membrane-free/membraneless liquid-like protein-rich droplets. Both of these membraneless microcompartments represent interesting systems to circumvent the coupling between volume and surface area, and their austerity in comparison to membrane-bound microcompartments has qualified them as useful models to demonstrate compartment formation and growth. The relevance of membraneless compartments to bottom-up synthetic biology, regardless of their chemical composition, remains undisputed, since they successfully represent or mimic certain biological aspects. Thus, phase-separated droplets represent useful models of self-organized entities, independent of the natural or man-made origin of the macromolecules. From a protobiology perspective, coacervates made of synthetic polymers are logically excluded as potential ancestors of today's cells; but such coacervates have played an important role in the history of protocells, starting with the first experiments on gum arabic and gelatin.

2.1. Protein-Rich Droplets

Compartmentalization is a hallmark of the eukaryotic cell. It enables cells to spatially separate their complex biochemistry into microreactors. Well-known examples of compartments are the mitochondrion and the nucleus, both of which are surrounded by lipid bilayers. However, in recent years it has become clear that the eukaryotic cytoplasm is further organized by compartments that lack membranes. These compartments have been termed biomolecular condensates, also known as membraneless organelles.[8] Intracellular condensates form by the process of liquid–liquid phase separation.[9–14] Phase separation is a highly cooperative and concentration-dependent process: above a certain saturation concentration, a solution becomes unstable and demixes to form two or more distinct coexisting aqueous phases, a protein-dense and a protein-poor phase that continuously exchange molecules and maintain a steep concentration gradient across their interfaces. Recent work suggests that the formation of biomolecular condensates is primarily driven by RNA and proteins that exhibit a large

Ivan Ivanov obtained his Ph.D. at the Max Planck Institute for Dynamics of Complex Technical Systems (Magdeburg, Germany) in the field of enzymatic fuel cells. Afterward, he spent two years as a postdoc at the Penn State University, working in the area of microbial electrochemical technologies. Currently, he is a team leader at his alma mater Max Planck Institute and is active in the field of bottom-up synthetic biology. His research interests include biological electron transfer and electrocatalysis, biomimetics, multiphase systems, and compartmentalization.

Rumiana Dimova obtained her Ph.D. at Bordeaux University (France) in 1999. Afterward she joined the Max Planck Institute of Colloids and Interfaces (Potsdam, Germany) as a postdoctoral fellow, where she became a group leader in 2000. Since then she has lead the Biophysics Lab in the Department of Theory and BioSystems. Her main research interests are in the field of membrane biophysics and synthetic biology.

degree of intrinsic disorder and/or multivalency. Among the growing list of proteins that undergo liquid-like phase separation are the well-studied P-granule proteins LAF1 and PGL-3,[15,16] the stress granule proteins FUS and hnRNPA1,[17–19] the centrosome protein SPD-5,[20] as well as the nucleolar proteins Fib1 and Npm1 that form immiscible phases to organize the nucleolus.[21] When expressed in a test tube, these proteins readily phase-separate to form two-phase systems at physiological concentrations. These protein-rich droplets grow by taking up material from the environment (Ostwald ripening) or via coalescence of two smaller droplets to form a larger one (see **Figure 1**). The condensates exhibit liquid-like characteristics, such that they can coalesce, wet surfaces, and deform under shear stress. The dynamics of these processes are governed by internal viscosity and surface tension.[9,15,22] Importantly, condensates display selectivity, and client molecules such as interacting proteins or RNA will partition into the condensates.

Recent work has identified that the saturation concentration of the FET family of proteins (FUS-EWSR1-TAF15 protein family) is quantitatively determined by the associative cation–π interactions between segregated tyrosine and arginine sticker residues.[24] Other types of interactions, such as π–π and electrostatics between charged residues, play important roles as
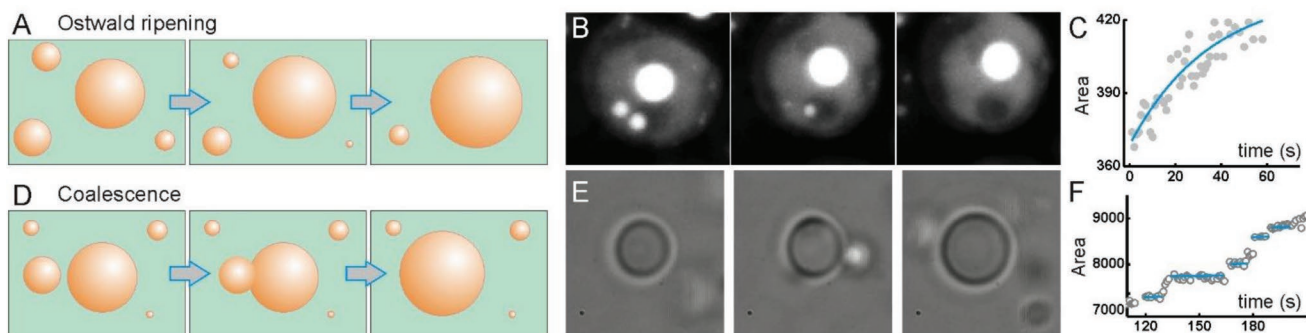
Figure 1. A) Protein-rich droplets can grow by the uptake of material from the surrounding and at the expense of smaller droplets (Ostwald ripening). B) Still fluorescence images from a time-lapse movie of a Hela cell expressing the YFP-labeled disordered region of the RNA helicase DDX4 (unpublished data). The disordered region forms an intracellular liquid-like droplet. With time, the bigger droplet grows, whereas the smaller droplets shrink and eventually disappear. C) Quantification of the bigger droplet area shown in (B) as a function of time (unpublished data). D) Growth of liquid-like protein droplets can also occur via coalescence upon encounter of two or more droplets that merge to form a bigger one. E) Still bright field images from a time-lapse movie of protein droplets formed from the prion protein Sup35.[23] The protein droplet was held with an optical tweezers and brought in the vicinity of other droplets, which coalesce leading to growth with time. F) Quantification of the droplet area shown in (E) as a function of time. The step-wise increase in area is indicative of coalescence.

well.[25] For instance, the disordered region of the condensate protein helicase DDX4 displays a great degree of segregated charges.[26] Importantly, weak multivalent interactions underlie the nucleation process.[12]

In the context of the cell, these systems are driven away from equilibrium, e.g., by chemical reactions. However, in the test tube, droplet growth continues until the thermodynamic equilibrium has been reached. Accordingly, mechanisms must exist that regulate the nucleation as well as the size and the mechanical properties. Liquid-like condensates can also undergo liquid-to-solid transitions.[19,23,27,28] For FET family proteins, the composition of interspacing residues (so-called spacers) that segregates the stickers plays an important role in modulating the interaction strength and thus the liquid-to-solid transition.[24] The liquid-to-solid transition of the centrosome forming protein SPD-5 may be mediated by coiled-coil domains to establish a force-resistant meshwork for microtubule-mediated chromosome segregation.[20] Coupling phase separation to gelation may provide an ideal size-determining mechanism. Moreover, coupling the phase behavior to biochemical reactions that drive the system away from equilibrium establishes control over nucleation, droplet growth and, under certain circumstances, may even provide a mechanism for droplet division.[29,30] Phase separation is exquisitely sensitive to environmental changes, including temperature, ionic strength, or pH, parameters that also fluctuate within cells.[31,32] Cells have harnessed the phenomenon of phase separation to orchestrate complex processes and evolved precise mechanisms to control nucleation of biomolecular condensates. Taken together, it is now possible to predict the saturation concentration, as well as the material constants for minimal proteinaceous systems. Linking synthetic phase separation units with enzyme functions provides an opportunity to organize and polarize biochemical reactions within a synthetic cell.

2.2. Polymer-Rich Droplets

Coacervate microdroplets are generated from associative liquid–liquid phase separation (coacervation) processes

between oppositely charged polymers or small highly charged molecules. They form from synthetic polymers (PDDA, polyethylemine, etc.),[33,34] as well as from biologically relevant molecules including proteins (BSA),[35] polypeptides (polylysine, polyarginine),[36,37] nucleic acids (DNA, RNA),[34,38,39] and cofactors (ATP, FAD, NAD).[40] In comparison, dissociative liquid–liquid phase separation into two aqueous phases can be triggered in solutions of neutral macromolecules such as polyethylene glycol (PEG) and dextran, and more recently, this process was also established in the closed compartment of lipid vesicles.[41,42] It is commonly considered that the general mechanism of phase separation occurs via two steps: an enthalpic contribution to the free energy from the electrostatic interaction between the molecules (in the case of charged polymers), which draws the molecules toward each other, and an entropic driving force from the rearrangement of ions and water leading to a lowering of the Gibbs free energy and the formation of membraneless, chemically enriched microdroplets. It has been proposed that this mechanism is analogous to liquid–liquid phase separation in biology.[43,44] The surface tension of coacervates is low, between 1 μN/m and 1 mN/m,[45] and even lower for aqueous two-phase systems of neutral polymers.[46] The droplets will grow in size over time via coalescence events.[47] The rate of coalescence and, therefore, growth can be tuned by the overall charge ratios of the coacervate components.

Studies have shown that polymer-rich microdroplets will selectively partition a range of different molecules leading to molecular localization and further chemical enrichment.[40,48–50] Moreover, enzymes, which partition into the microdroplets, maintain activity within the highly charged and crowded interior.[51,52] Enzymes capable of phosphorylation and dephosphorylation have led to cycles of growth and degradation of the coacervate droplets.[38,53] In addition, growth and decay of coacervate droplets can be instigated by switching the pH of the solution by bubbling carbon dioxide and ammonia through the dispersion of coacervate droplets.[40] These results indicate that the physical parameters such as molar ratio of polymers or pH and temperature of coacervate droplets can be used to tune and drive droplet growth and disassembly. Additionally, the ability

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of membraneless coacervate droplets to partition and support enzyme reactions could suggest that these systems could be plausible models for predicted growth and division cycles driven by flux of molecules across the interface.[30]

3. Fatty Acid Vesicles as Primitive Protocells: Membrane Formation due to Fast Equilibrium

The vesicular (membrane) systems represent another compartment type with the same structural features as living cells. Worth noting with respect to the field of minimal cells, and growth in particular, are the pioneering works of Luisi and Szostak,[54,55] which mainly relied on fatty acid micelles and vesicles. The latter have been investigated extensively as protocell models because fatty acids have been proposed to be prebiotically relevant due to their structural simplicity compared to phospholipids.[56] Therefore, the spontaneous uptake of fatty acids into preformed vesicles has often been considered as a primitive growth mechanism. As far as structural simplicity is concerned, the emphasis on fatty acids is analogous to the RNA hypothesis (single-/double-stranded vs one or two hydrophobic tails) but the quest for the origin of life is intrinsically speculative and we will refrain from statements about the evolutionary relevance of these compartments. Regardless of this open question, fatty acid vesicles and fatty acids have served as rewarding models for mimicking life processes, including growth.

We also note that the majority of studies with fatty acids employed small (≈30–50 nm) and large (≈100–200 nm) unilamellar vesicles—below the limit of optical resolution—which poses the question whether those are relevant protocell models from another perspective, namely, their small size. A workshop dedicated to defining the minimal compartment size of life, related to possibilities for extraterrestrial life and nanobacteria, dealt with this question 20 years ago (see proceedings).[57] The biologists' view regarding the threshold size of primitive compartments was set to diameters of 250 ± 50 nm, determined by a requirement for the compartment to accommodate minimum amount of necessary proteins, genes, and ribosomes. Others' views were more speculative suggesting ≈50 nm as the minimal size required to sustain potential forms of primordial life. The latter notion was supported by reports on enhanced protein expression in 100 nm liposomes.[58] Considered from a simplistic chemoton perspective, smaller size implies higher surface-to-volume ratio, which is associated with enhanced inward flux (proportional to the surface) of membrane precursors and other metabolites and thus could be considered as beneficial. However, other factors such as packing, curvature, etc., could result in micrometer size (as in modern cells) optimal for a specific function. In this progress report, we focus on biomimetic microcompartments, while we do not aim to set the size limits of protocells. However, we include reports on nanocompartments as well, due to their historical and mechanistic significance. Furthermore, microcompartments such as giant unilamellar vesicles (GUVs)[59–62] have proven to be a practical system, whose properties, growth, and response to external factors can be monitored and manipulated directly under a microscope.

Compared to phospholipids, fatty acids exhibit a very quick exchange between the membrane leaflets as well as faster exchange kinetics between the vesicle and the solution. In addition, stabilizing head group interactions are more important, which endows distinct properties, such as growth and self-reproduction of vesicles.[63–65] Due to their single-chain structure, the concentration of monomers in equilibrium with vesicles is significantly higher than for phospholipids, which enables fast flip-flop and exchange of molecules (**Figure 2**A). These beneficial properties have qualified fatty acids as a prominent system for studying protocells, including the demonstration of membrane growth.[66] A notable phenomenon, observed during the growth of oleate vesicles, was the so-called matrix effect, which exemplified itself when a seed of preformed vesicles was added to the micellar oleate solution, resulting in a narrow size distribution, corresponding to the size of the seed.[66] The growth of oleate vesicles was further investigated in detail, which led to kinetic[67] and molecular dynamics[68] models. The growth process was also demonstrated under flow,[69] whereby filamentous microcompartments were formed, and also employed as a mechanism for ribozyme activation[70] to manifest a form of homeostasis. In addition, the filamentous growth of multilamellar oleate vesicles (≈4 µm) was used to mimic the full proliferation cycle of growth and division,[71] and the chemistry was extended to other surface-active molecules beyond oleate.[72]

While fatty acids and surfactants are efficiently incorporated into existing membranes due to their optimal physicochemical properties, phospholipids—the membrane constituents of modern living cells—form structures, which are stable over a wider range of concentrations (phospholipids exhibit lower critical micelle concentration, CMC, compared to fatty acids). With respect to this distinction, the addition of oleate to preformed phospholipid vesicles was used in other attempts to mimic self-reproduction as an intermediate approach. Growth and division has been shown and investigated in detail in several examples of fatty acid incorporation into smaller liposomes[73–75] and the method has been later on scaled up to GUVs.[76,77] In a similar fashion, the addition of detergents and other water-soluble membrane active molecules at low concentration leads to their intercalation into the bilayer, which in turn could lead to an increase in the membrane area. In GUVs this increase depends on whether the inserting molecules exhibit fast or slow flip-flop. Fast flip-flop induces a large, visible increase in membrane area and fluctuations.[76,78,79] Slow flip-flop molecules induce an increase in the membrane spontaneous curvature due to their asymmetric incorporation into the outer leaflet of the membrane, resulting in tubulation,[80–82] increased membrane tension, and eventually rupture.[76,78] Interestingly, the incorporation of simpler amphiphilic molecules or micelles leads to vesicle growth, often followed by division,[71,83] which is in line with predictions for phospholipids.[84] The limitation of detergent-like molecules is that at high concentrations, they lead to membrane permeabilization and eventual solubilization, depending on the membrane phase state and the CMC of the surfactant.[85,86] In general, although this mixed approach provides a tangible system for studies of liposome growth, there is major conceptual drawback in the context of minimal
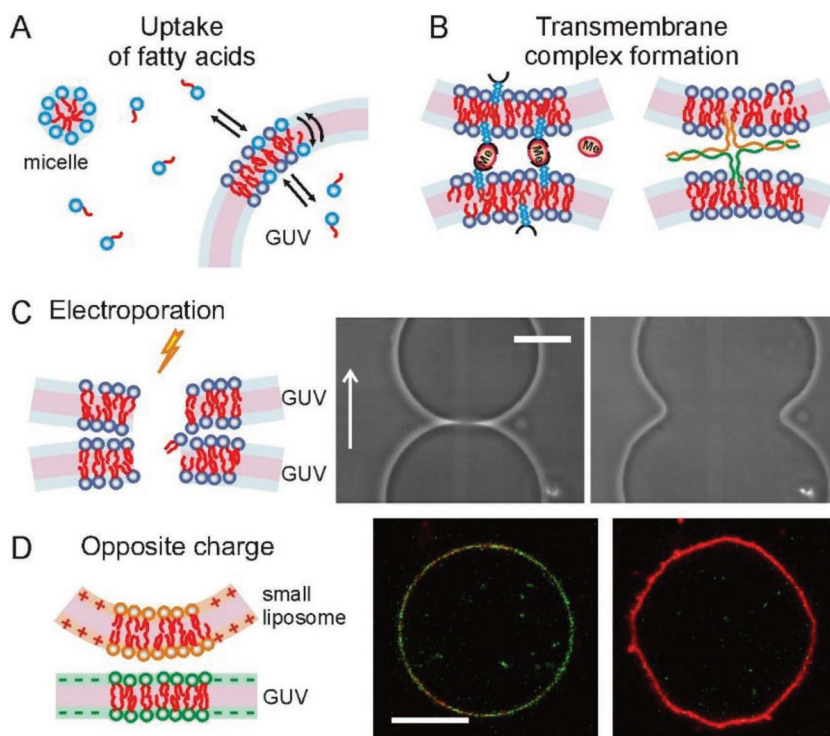
Figure 2. Some approaches for establishing growth of giant vesicles. A) Growth via uptake of fatty acid molecules, which remain in dynamic exchange with the external and internal GUV solution and exhibit fast flip-flop across the membrane. B) Growth via fusion mediated by the formation of transmembrane metal-ligand complex (left) or fusion protein mimetics (right). C) Electrofusion relies on poration of two opposing membranes exposed to strong electric fields. The phase contrast snapshots on the right illustrate the fusion of two GUVs brought in contact and aligned by means of AC field, followed by the application of a DC pulse. Reproduced with permission.[88] Copyright 2006, National Academy of Sciences. The white arrow indicates the direction of the field. (D) Growth of giant vesicles can be established by efficient fusion of small liposomes with the GUV, where the two membrane types are oppositely charged. Fusion is illustrated with the cartoon (left) and the confocal images (right) of one negatively charged GUV (green) exposed to a solution of positively charged small liposomes (red). Upon fusion, Förster resonance energy transfer leads to decrease in the red signal and the GUV area increases significantly.[89] The change occurs within seconds. Reproduced with permission.[89] Copyright 2019, Elsevier. Scale bars in (C,D) represent 20 µm.

cells—such microcompartments would lose their chemical identity over several generations and therefore could not sustain a continuous life cycle. The same drawback applies to another approach, in which blends of phospholipids and fatty acids or cationic surfactants were fused by attractive electrostatic interactions.[87]

4. Growth of Lipid Vesicles: A Difficult Task, Solved by Fusion

The system of highest interest for mimicking the proliferation of living cells is the phospholipid vesicle, which grows as a result of the incorporation of phospholipids in its membrane and eventually divides. The importance of liposomes arises from the chemical analogy with modern cell membranes. The cellular self-reproduction found in nature involves very complex biochemical interactions and machineries. According to the reductionist concept, research in the context of protobiology

and bottom-up synthetic biology aims to establish this phenomenon in a minimal system, based on simpler physicochemical mechanisms in a proof-of-concept for the origin and understanding of life. So far, however, growth of pure lipid vesicles has proved to be a challenge. Due to the difficulty of experimental realization of the membrane formation subsystem, however, theoretical studies have preceded practical examples in the case of phospholipids. Different kinetic[90] and thermodynamic[84,91,92] aspects have been considered and summarized together with other chemoton-like approaches in a recent review.[93] On the practical side, the self-reproduction of a biomembrane, i.e., a phospholipid vesicle building its own membrane, has been addressed several times, but with only modest success so far.

4.1. Growth via Uptake of Synthesized Membrane Components

Efforts for synthesizing membrane components in liposomes date back to the early 90s. In particular, Luisi and co-workers assembled a four-enzyme cascade for phosphatidylcholine (PC) synthesis and deduced its incorporation into the membrane based on geometrical considerations (eventually the liposome size decreased, which was ascribed to the higher spontaneous curvature triggered by the newly synthesized short-chain PC—a partially undesired outcome with respect to growth).[94] The authors did not unequivocally determine whether the enzymes were present in the interior, whereby significant dimensional limitations could arise from the small liposome radius (23–26 nm), but they ascertained the enzyme association with the membrane to maintain an active form. In another study, the synthesis of phosphatidic acid by acyltransferases, expressed via cell-free methods in vesicles, was demonstrated but no growth was observed.[95] The latter observation was ascribed to the low amount of product due to the limited encapsulation, which will be difficult to overcome unless efficient transport mechanisms for the various precursors are established. Recently, this cell-free approach was extended to a more comprehensive pathway, starting from acyl-CoA and glycerol-3-phosphate, and the liposome compartments were equipped with α-hemolysin to facilitate uptake of small polar molecules.[96] Although the authors did not specifically follow the vesicle growth, they identified practical barriers causing the low lipid synthesis rate, speculated about transport mechanisms for acyl-CoA, and discussed relevant crowding and confinement effects. Another example in a similar direction was the biochemical synthesis of palmitate based on a FAS type I enzyme, encapsulated in liposomes,[97] where vesicle growth has indeed been observed

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but the incorporation of foreign membrane components remains a conceptual problem.

To circumvent protein synthesis issues, inherent to cell-free systems, an eight-enzyme cascade for the biosynthesis of phospholipids from fatty acids and glycerol 3-phosphate as building blocks has recently been assembled in vitro.[98] In the latter study, the enzymes and precursors were not encapsulated and synthesis occurred outside the liposomes. Experiments measuring the dequenching of rhodamine dye showed an approximately 30% membrane expansion, which was subsequently limited by the depletion of substrate. The growth was ascribed to the incorporation of oleic acid into the preformed liposomes, while the subsequent biosynthetic conversion to phosphatidic acid did not contribute to further expansion. Despite these limitations, the reconstitution of the biochemical membrane formation machinery is undisputedly a landmark in the area of bottom-up synthetic biology.

The continuous search for more realistic growing protocell models has led to the use of phospholipid-like molecules, supplied from outside, instead of internally synthesized. The Sugawara group demonstrated growth, division, and amplified DNA distribution in daughter vesicles.[99] This was achieved by adding a cationic precursor, which was hydrolyzed into the membrane lipid by an embedded catalyst. Notably, DNA amplification enhanced the efficiency of both growth and division. However, the elegant process was ultimately exhausted because it was limited by the consumption of DNA precursors inside the vesicles and the dilution of membrane catalyst and phospholipids. Similar approaches were undertaken by the replacement of the complex biochemical lipid synthesis pathway with a self-reproducing autocatalyst, which resulted in triazole phospholipids formation and membrane growth.[100] Both cases seem to function with aggregates and multilamellar vesicles, and the exotic chemistry of the membranes might hinder interactions with some biological species (e.g., sensitive membrane proteins), thus potentially limiting other applications in the context of minimal cells.

4.2. Growth via Vesicle Fusion

In the case of authentic phospholipids, the only established and practical approach for achieving growth until now seems to be via vesicle fusion. Membrane fusion is a ubiquitous process in biology and it is fundamental for a number of cellular processes. It involves merging of two otherwise separated membranes, forming a compartment, whose area is the sum of the two fusing bilayers. In other words, increase in membrane area inevitably accompanies fusion, although the former is scarcely reported because the vast majority of fusion assays are insensitive to it. Driving forces of various nature have been employed to trigger fusion, and some approaches are illustrated in Figure 2. These range from simple physicochemical triggers such as electrostatic interactions[101–103] and volume depletion[104] to biochemical approaches, relying on natural protein fusion machinery,[105] as well as combinations thereof.[106] Furthermore, fusion methods have been expanded to biomimetic strategies[88,107] in addition to more exotic approaches, involving the use of light (to heat nanoparticles[108] or isomerize

azocompounds[109]) or electric fields[88,110,111] to perturb and porate the bilayer. In some of the examples fusion has led to spontaneous budding, correlating to predictions made by theoretical models,[91] which has been put in the context of growth and division cycles for self-reproduction.[110] Presumably, the simplest mechanism to induce membrane fusion is based on membrane tension as has been elucidated by molecular simulations.[112]

Membrane fusion of small liposomes has been used for decades[113,114] and only more recently fusion using giant vesicles has been reported.[115] In cells, membrane fusion is mediated by specific interactions between fusion proteins and such fusion reactions have been reconstituted in synthetic systems, in which small liposomes fuse with giant vesicles.[105,116,117] However, possibly due to the low fusion efficiency of protein-reconstituted systems or the low protein density, vesicle growth has not been detected. Fusion mediated by pH-sensitive lipids has also resulted in undetectable area increase.[118] In contrast, when mediated by high charge density, the fusion of positively charged liposomes induces extensive area increase of the negative GUVs[89,119] (see also Figure 2D). Although very efficient, the process is limited by eventual charge neutralization.

The fusion of two GUVs can also be induced and observed in a more controllable way. This can be achieved by bringing a pair of vesicles in contact by manipulation with ultramicroelectrodes,[120] by trapping them within microfluidic devices,[121] by applying an electric field,[88,111] or by direct manipulation using micropipettes[88,122] or optical tweezers.[108,123] After initial contact, fusion can be triggered by the application of a strong DC pulse,[88,124] via ligand-mediated ion interactions,[88] transmembrane domains of fusogenic proteins,[125] or through localized nanoparticle-mediated[108] or laser-mediated[123] heating. The common feature of all these studies is that an increase in membrane area is clearly observable. Although GUV fusion may result in visible area increase, the resulting area has never been quantified and fusion efficiency has been determined by changes in the measured fluorescence rather than the measured volume changes of the vesicle to characterize vesicular growth. Generally, the excess area results in increased membrane fluctuations and formation of membrane folds as a consequence of volume expansion lagging behind the area growth. The GUV–GUV fusion approach in its variations has been used to mix chemical precursors for nanoparticle formation inside the vesicular compartment,[120,121,124] to controllably obtain vesicles with well-defined compositions,[122] and to unravel the formation of the hemifusion diaphragm[125,126] and the rapid nature of the fusion neck expansion.[88]

Importantly, fusion is not limited to pure phospholipid GUVs, but can also be extended to fusion of GUVs and cells[127,128] or GUVs made of synthetic polymers. In contrast to liposomes, fusion of polymersomes is limited by the restricted mobility and high stability of the polymer membranes[129–131] associated with slowed down polymer rearrangement, which is a prerequisite for processes such as fusion. Some strategies have been successful in inducing fusion events in polymer GUVs, including exposure of vesicles to osmotic pressure gradients,[129] ultrasound treatment,[132] membrane perturbation by

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azobenzene photoisomerization,[133] and addition of membrane disruptive agents into the dispersing medium.[134]

In relation to bottom-up synthetic biology and protobiology, although fusion has been traditionally considered from a biophysical and biochemical perspective, spontaneous liposome growth could in principle occur (or could have occurred in the evolutionary history) by the discrete steps of vesicle fusion.[135] In addition, apart from a mechanism for increasing the membrane area, it has been speculated as a scenario, in which two systems combine their properties and thus increase the degree of biocomplexity,[136] for instance considering that vesicular trafficking is a major transport mechanism for proteins in natural living cells.[137] This aspect was used to supply feeding components and trigger gene expression in the case of fusion of oppositely charged vesicles.[138]

5. Synthetic Chemistry Enables New Potential Approaches for Growth

Parallel to the continuing efforts to form and grow new types of membraneless microcompartments, enable efficient fusion via new approaches, and ideally reconstitute the natural machinery for phospholipid synthesis, there is another possible pathway, which brings synthetic chemistry in the foreground of bottom-up synthetic biology. This aspect has been already partially addressed by the demonstration of life-like properties of dichloromethane and nitrobenzene oil droplets, which grow, divide, and partition cargo.[139,140] The versatility of synthetic reactions was also used to augment phospholipid vesicles and make them grow[99,100] as noted above. With respect to this, advances in the preparation of functional amphiphilic block copolymers provide powerful synthetic tools for the engineering of artificial compartments that could grow upon stimulation by light, temperature, or pH. Here the stimuli would induce the build-up of membrane stresses that could relax through fusion events. In this way the chemistry of living systems could be extended far beyond natural building blocks, enabling not only the reverse engineering or re-engineering, but also the de novo engineering of life. Besides a powerful toolbox, which may enable futuristic applications, the latter concept of synthetic (in its conventional semantics of chemical) has resorting implications on the understanding of life as a display of self-organization and will expand our search criteria for other forms of life.

Conflict of Interest

The authors declare no conflict of interest.

[1] T. Ganti, *Biosystems* **1975**, *7*, 15.
[2] T. Ganti, *J. Theor. Biol.* **1997**, *187*, 583.
[3] P. L. Luisi, *Naturwissenschaften* **2003**, *90*, 49.
[4] F. G. Varela, H. R. Maturana, R. Uribe, *Biosystems* **1974**, *5*, 187.
[5] We conceptually include examples of fatty acid vesicles as presumable predecessors, as well as vesicles made of synthetic chemicals as prospective successors of today's cells.
[6] A. I. Oparin, *Origin of Life*, Macmillan, New York **1938**.
[7] A. I. Oparin, K. L. Gladilin, *Biosystems* **1980**, *12*, 133.
[8] S. F. Banani, H. O. Lee, A. A. Hyman, M. K. Rosen, *Nat. Rev. Mol. Cell Biol.* **2017**, *18*, 285.
[9] C. P. Brangwynne, C. R. Eckmann, D. S. Courson, A. Rybarska, C. Hoege, J. Gharakhani, F. Julicher, A. A. Hyman, *Science* **2009**, *324*, 1729.
[10] C. P. Brangwynne, T. J. Mitchison, A. A. Hyman, *Proc. Natl. Acad. Sci. USA* **2011**, *108*, 4334.
[11] A. A. Hyman, C. A. Weber, F. Juelicher, *Annu. Rev. Cell. Dev. Biol.* **2014**, *30*, 39.
[12] P. L. Li, S. Banjade, H. C. Cheng, S. Kim, B. Chen, L. Guo, M. Llaguno, J. V. Hollingsworth, D. S. King, S. F. Banani, P. S. Russo, Q. X. Jiang, B. T. Nixon, M. K. Rosen, *Nature* **2012**, *483*, 336.
[13] Y. Shin, C. P. Brangwynne, *Science* **2017**, *357*, eaaf4382.
[14] X. L. Su, J. A. Ditlev, E. F. Hui, W. M. Xing, S. Banjade, J. Okrut, D. S. King, J. Taunton, M. K. Rosen, R. D. Vale, *Science* **2016**, *352*, 595.
[15] S. Elbaum-Garfinkle, Y. Kim, K. Szczepaniak, C. C. H. Chen, C. R. Eckmann, S. Myong, C. P. Brangwynne, *Proc. Natl. Acad. Sci. USA* **2015**, *112*, 7189.
[16] S. Saha, C. A. Weber, M. Nousch, O. Adame-Arana, C. Hoege, M. Y. Hein, E. Osborne-Nishimura, J. Mahamid, M. Jahnel, L. Jawerth, A. Pozniakovski, C. R. Eckmann, F. Julicher, A. A. Hyman, *Cell* **2016**, *166*, 1572.
[17] K. A. Burke, A. M. Janke, C. L. Rhine, N. L. Fawzi, *Mol. Cell* **2015**, *60*, 231.
[18] T. Murakami, S. Qamar, J. Q. Lin, G. S. K. Schierle, E. Rees, A. Miyashita, A. R. Costa, R. B. Dodd, F. T. S. Chan, C. H. Michel, D. Kronenberg-Versteeg, Y. Li, S. P. Yang, Y. Wakutani, W. Meadows, R. R. Ferry, L. Dong, G. G. Tartaglia, G. Favrin, W. L. Lin, D. W. Dickson, M. Zhen, D. Ron, G. Schmitt-Ulms, P. E. Fraser, N. A. Shneider, C. Holt, M. Vendruscolo, C. F. Kaminski, P. St George-Hyslop, *Neuron* **2015**, *88*, 678.
[19] A. Patel, H. O. Lee, L. Jawerth, S. Maharana, M. Jahnel, M. Y. Hein, S. Stoynov, J. Mahamid, S. Saha, T. M. Franzmann, A. Pozniakovski, I. Poser, N. Maghelli, L. A. Royer, M. Weigert, E. W. Myers, S. Grill, D. Drechsel, A. A. Hyman, S. Alberti, *Cell* **2015**, *162*, 1066.
[20] J. B. Woodruff, B. F. Gomes, P. O. Widlund, J. Mahamid, A. Honigmann, A. A. Hyman, *Cell* **2017**, *169*, 1066.
[21] M. Feric, N. Vaidya, T. S. Harmon, D. M. Mitrea, L. Zhu, T. M. Richardson, R. W. Kriwacki, R. V. Pappu, C. P. Brangwynne, *Cell* **2016**, *165*, 1686.
[22] A. Hubstenberger, S. L. Noble, C. Cameron, T. C. Evans, *Dev. Cell* **2013**, *27*, 161.

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[23] T. M. Franzmann, M. Jahnel, A. Pozniakovsky, J. Mahamid, A. S. Holehouse, E. Nuske, D. Richter, W. Baumeister, S. W. Grill, R. V. Pappu, A. A. Hyman, S. Alberti, *Science* **2018**, *359*, eaao5654.

[24] J. Wang, J. M. Choi, A. S. Holehouse, H. O. Lee, X. J. Zhang, M. Jahnel, S. Maharana, R. Lemaitre, A. Pozniakovsky, D. Drechsel, I. Poser, R. V. Pappu, S. Alberti, A. A. Hyman, *Cell* **2018**, *174*, 688.

[25] S. Boeynaems, S. Alberti, N. L. Fawzi, T. Mittag, M. Polymenidou, F. Rousseau, J. Schymkowitz, J. Shorter, B. Wolozin, L. Van den Bosch, P. Tompa, M. Fuxreiter, *Trends Cell Biol.* **2018**, *28*, 420.

[26] T. J. Nott, E. Petsalaki, P. Farber, D. Jervis, E. Fussner, A. Plochowietz, T. D. Craggs, D. P. Bazett-Jones, T. Pawson, J. D. Forman-Kay, A. J. Baldwin, *Mol. Cell* **2015**, *57*, 936.

[27] T. S. Harmon, A. S. Holehouse, M. K. Rosen, R. V. Pappu, *Elife* **2017**, *6*, 30294.

[28] A. Molliex, J. Temirov, J. Lee, M. Coughlin, A. P. Kanagaraj, H. J. Kim, T. Mittag, J. P. Taylor, *Cell* **2015**, *163*, 123.

[29] D. Zwicker, A. A. Hyman, F. Julicher, *Phys. Rev. E* **2015**, *92*, 012317.

[30] D. Zwicker, R. Seyboldt, C. A. Weber, A. A. Hyman, F. Julicher, *Nat. Phys.* **2017**, *13*, 408.

[31] T. Franzmann, S. Alberti, *J. Biol. Chem.* **2018**, https://doi.org/10.1074/jbc.TM118.001190.

[32] K. M. Ruff, S. Roberts, A. Chilkoti, R. V. Pappu, *J. Mol. Biol.* **2018**, *430*, 4619.

[33] R. Chollakup, W. Smitthipong, C. D. Eisenbach, M. Tirrell, *Macromolecules* **2010**, *43*, 2518.

[34] T. Y. D. Tang, C. R. C. Hak, A. J. Thompson, M. K. Kuimova, D. S. Williams, A. W. Perriman, S. Mann, *Nat. Chem.* **2014**, *6*, 527.

[35] K. A. Black, D. Priftis, S. L. Perry, J. Yip, W. Y. Byun, M. Tirrell, *ACS Macro Lett.* **2014**, *3*, 1088.

[36] D. Priftis, K. Megley, N. Laugel, M. Tirrell, *J. Colloid Interface Sci.* **2013**, *398*, 39.

[37] D. Priftis, M. Tirrell, *Soft Matter* **2012**, *8*, 9396.

[38] W. M. Aumiller, C. D. Keating, *Nat. Chem.* **2016**, *8*, 129.

[39] J. R. Vieregg, M. Lueckheide, A. B. Marciel, L. Leon, A. J. Bologna, J. R. Rivera, M. V. Tirrell, *J. Am. Chem. Soc.* **2018**, *140*, 1632.

[40] S. Koga, D. S. Williams, A. W. Perriman, S. Mann, *Nat. Chem.* **2011**, *3*, 720.

[41] R. Dimova, R. Lipowsky, *Soft Matter* **2012**, *8*, 6409.

[42] C. D. Keating, *Acc. Chem. Res.* **2012**, *45*, 2114.

[43] C. W. Pak, M. Kosno, A. S. Holehouse, S. B. Padrick, A. Mittal, R. Ali, A. A. Yunus, D. R. Liu, R. V. Pappu, M. K. Rosen, *Mol. Cell* **2016**, *63*, 72.

[44] V. N. Uversky, I. M. Kuznetsova, K. K. Turoverov, B. Zaslavsky, *FEBS Lett.* **2015**, *589*, 15.

[45] E. Spruijt, J. Sprakel, M. A. C. Stuart, J. van der Gucht, *Soft Matter* **2010**, *6*, 172.

[46] Y. G. Liu, R. Lipowsky, R. Dimova, *Langmuir* **2012**, *28*, 3831.

[47] D. S. Williams, S. Koga, C. R. C. Hak, A. Majrekar, A. J. Patil, A. W. Perriman, S. Mann, *Soft Matter* **2012**, *8*, 6004.

[48] E. A. Frankel, P. C. Bevilacqua, C. D. Keating, *Langmuir* **2016**, *32*, 2041.

[49] T. Y. D. Tang, M. Antognozzi, J. A. Vicary, A. W. Perriman, S. Mann, *Soft Matter* **2013**, *9*, 7647.

[50] M. S. Long, C. D. Jones, M. R. Helfrich, L. K. Mangeney-Slavin, C. D. Keating, *Proc. Natl. Acad. Sci. USA* **2005**, *102*, 5920.

[51] E. Sokolova, E. Spruijt, M. M. K. Hansen, E. Dubuc, J. Groen, V. Chokkalingam, A. Piruska, H. A. Heus, W. T. S. Huck, *Proc. Natl. Acad. Sci. USA* **2013**, *110*, 11692.

[52] T. Y. D. Tang, D. van Swaay, A. deMello, J. L. R. Anderson, S. Mann, *Chem. Commun.* **2015**, *51*, 11429.

[53] K. K. Nakashima, J. F. Baaij, E. Spruijt, *Soft Matter* **2018**, *14*, 361.

[54] P. Stano, P. L. Luisi, *Chem. Commun.* **2010**, *46*, 3639.

[55] M. M. Hanczyc, S. M. Fujikawa, J. W. Szostak, *Science* **2003**, *302*, 618.

[56] D. Deamer, J. P. Dworkin, S. A. Sandford, M. P. Bernstein, L. J. Allamandola, *Astrobiology* **2002**, *2*, 371.

[57] N. R. C. Steering Group for the Workshop on Size Limits of Very Small Microorganisms, *Size Limits of Very Small Microorganisms: Proceedings of a Workshop*, National Academies Press (US) Copyright 1999 by the National Academy of Sciences, Washington, DC, **1999**.

[58] T. P. de Souza, P. Stano, P. L. Luisi, *ChemBioChem* **2009**, *10*, 1056.

[59] R. Dimova, S. Aranda, N. Bezlyepkina, V. Nikolov, K. A. Riske, *J. Phys. : Condens. Matter* **2006**, *18*, S1151.

[60] R. Dimova, in *Advances in Planar Lipid Bilayers and Liposomes* (Ed: A. Iglič), Academic Press, Burlington **2012**, pp. 1–50.

[61] P. Walde, K. Cosentino, H. Engel, P. Stano, *ChemBioChem* **2010**, *11*, 848.

[62] S. F. Fenz, K. Sengupta, *Integr. Biol.* **2012**, *4*, 982.

[63] K. Morigaki, P. Walde, *Curr. Opin. Colloid Interface Sci.* **2007**, *12*, 75.

[64] I. A. Chen, R. W. Roberts, J. W. Szostak, *Science* **2004**, *305*, 1474.

[65] P. Walde, R. Wick, M. Fresta, A. Mangone, P. L. Luisi, *J. Am. Chem. Soc.* **1994**, *116*, 11649.

[66] E. Blochliger, M. Blocher, P. Walde, P. L. Luisi, *J. Phys. Chem. B* **1998**, *102*, 10383.

[67] I. A. Chen, J. W. Szostak, *Biophys. J.* **2004**, *87*, 988.

[68] A. J. Markvoort, N. Pfleger, R. Staffhorst, P. A. Hilbers, R. A. van Santen, J. A. Killian, B. de Kruijff, *Biophys. J.* **2010**, *99*, 1520.

[69] C. Hentrich, J. W. Szostak, *Langmuir* **2014**, *30*, 14916.

[70] A. E. Engelhart, K. P. Adamala, J. W. Szostak, *Nat. Chem.* **2016**, *8*, 448.

[71] T. F. Zhu, J. W. Szostak, *J. Am. Chem. Soc.* **2009**, *131*, 5705.

[72] A. Shioi, T. A. Hatton, *Langmuir* **2002**, *18*, 7341.

[73] N. Berclaz, M. Muller, P. Walde, P. L. Luisi, *J. Phys. Chem. B* **2001**, *105*, 1056.

[74] T. P. de Souza, M. Holzer, P. Stano, F. Steiniger, S. May, R. Schubert, A. Fahr, P. L. Luisi, *J. Phys. Chem. B* **2015**, *119*, 12212.

[75] M. L. Rogerson, B. H. Robinson, S. Bucak, P. Walde, *Colloids Surf., B* **2006**, *48*, 24.

[76] P. Peterlin, V. Arrigler, K. Kogej, S. Svetina, P. Walde, *Chem. Phys. Lipids* **2009**, *159*, 67.

[77] J. Dervaux, V. Noireaux, A. J. Libchaber, *Eur. Phys. J. Plus* **2017**, *132*, 284.

[78] T. P. Sudbrack, N. L. Archilha, R. Itri, K. A. Riske, *J. Phys. Chem. B* **2011**, *115*, 269.

[79] V. N. Georgiev, A. Grafmüller, D. Bléger, S. Hecht, S. Kunstmann, S. Barbirz, R. Lipowsky, R. Dimova, *Adv. Sci.* **2018**, *5*, 1800432.

[80] R. Dasgupta, M. S. Miettinen, N. Fricke, R. Lipowsky, R. Dimova, *Proc. Natl. Acad. Sci. USA* **2018**, *115*, 5756.

[81] R. Lipowsky, *Faraday Discuss.* **2013**, *161*, 305.

[82] P. Bassereau, R. Jin, T. Baumgart, M. Deserno, R. Dimova, V. A. Frolov, P. V. Bashkirov, H. Grubmüller, R. Jahn, H. J. Risselada, L. Johannes, M. M. Kozlov, R. Lipowsky, T. J. Pucadyil, W. F. Zeno, J. C. Stachowiak, D. Stamou, A. Breuer, L. Lauritsen, C. Simon, C. Sykes, G. A. Voth, T. R. Weikl, *J. Phys. D: Appl. Phys.* **2018**, *51*, 343001.

[83] Y. Inaoka, M. Yamazaki, *Langmuir* **2007**, *23*, 720.

[84] B. Bozic, S. Svetina, *Eur. Biophys. J.* **2004**, *33*, 565.

[85] D. Lichtenberg, H. Ahyayauch, F. M. Goni, *Biophys. J.* **2013**, *105*, 289.

[86] B. Mattei, R. B. Lira, K. R. Perez, K. A. Riske, *Chem. Phys. Lipids* **2017**, *202*, 28.

[87] F. Caschera, P. Stano, P. L. Luisi, *J. Colloid Interface Sci.* **2010**, *345*, 561.

[88] C. K. Haluska, K. A. Riske, V. Marchi-Artzner, J. M. Lehn, R. Lipowsky, R. Dimova, *Proc. Natl. Acad. Sci. USA* **2006**, *103*, 15841.

[89] R. B. Lira, T. Robinson, R. Dimova, K. A. Riske, *Biophys J.* **2019**, *116*, 79.

[90] V. P. Zhdanov, B. Kasemo, *Langmuir* **2000**, *16*, 7352.

[91] B. Bozic, S. Svetina, *Eur. Phys. J. E* **2007**, *24*, 79.

[92] D. Fanelli, A. J. McKane, *Phys. Rev. E* **2008**, *78*, 051406.

[93] F. Mavelli, E. Altamura, L. Cassidei, *Entropy* **2014**, *16*, 2488.

[94] P. K. Schmidli, P. Schurtenberger, P. L. Luisi, *J. Am. Chem. Soc.* **1991**, *113*, 8127.

[95] Y. Kuruma, P. Stano, T. Ueda, P. L. Luisi, *Biochim. Biophys. Acta, Biomembr.* **2009**, *1788*, 567.

[96] A. Scott, M. J. Noga, P. de Graaf, I. Westerlaken, E. Yildirim, C. Danelon, *PLoS One* **2016**, *11*, 0163058.

[97] G. Murtas, *Sys. Synth. Biol.* **2010**, *4*, 85.

[98] M. Exterkate, A. Caforio, M. C. A. Stuart, A. J. M. Driessen, *ACS Synth. Biol.* **2018**, *7*, 153.

[99] K. Kurihara, M. Tamura, K. Shohda, T. Toyota, K. Suzuki, T. Sugawara, *Nat. Chem.* **2011**, *3*, 775.

[100] M. D. Hardy, J. Yang, J. Selimkhanov, C. M. Cole, L. S. Tsimring, N. K. Devaraj, *Proc. Natl. Acad. Sci. USA* **2015**, *112*, 8187.

[101] T. Tanaka, M. Yamazaki, *Langmuir* **2004**, *20*, 5160.

[102] D. P. Pantazatos, R. C. MacDonald, *J. Membr. Biol.* **1999**, *170*, 27.

[103] T. Sunami, F. Caschera, Y. Morita, T. Toyota, K. Nishimura, T. Matsuura, H. Suzuki, M. M. Hanczyc, T. Yomo, *Langmuir* **2010**, *26*, 15098.

[104] B. R. Lentz, *Eur. Biophys. J.* **2007**, *36*, 315.

[105] D. Tareste, J. Shen, T. J. Melia, J. E. Rothman, *Proc. Natl. Acad. Sci. USA* **2008**, *105*, 2380.

[106] M. Kyoung, Y. Zhang, J. Diao, S. Chu, A. T. Brunger, *Nat. Protoc.* **2013**, *8*, 1.

[107] Y. H. M. Chan, B. van Lengerich, S. G. Boxer, *Biointerphases* **2008**, *3*, FA17.

[108] A. Rorvig-Lund, A. Bahadori, S. Semsey, P. M. Bendix, L. B. Oddershede, *Nano Lett.* **2015**, *15*, 4183.

[109] Y. Suzuki, K. H. Nagai, A. Zinchenko, T. Hamada, *Langmuir* **2017**, *33*, 7393.

[110] H. Terasawa, K. Nishimura, H. Suzuki, T. Matsuura, T. Yomo, *Proc. Natl. Acad. Sci. USA* **2012**, *109*, 5942.

[111] K. A. Riske, N. Bezlyepkina, R. Lipowsky, R. Dimova, *Biophys. Rev. Lett.* **2006**, *01*, 387.

[112] J. C. Shillcock, R. Lipowsky, *Nat. Mater.* **2005**, *4*, 225.

[113] D. K. Struck, D. Hoekstra, R. E. Pagano, *Biochemistry* **1981**, *20*, 4093.

[114] J. Wilschut, N. Duzgunes, R. Fraley, D. Papahadjopoulos, *Biochemistry* **1980**, *19*, 6011.

[115] G. H. Lei, R. C. MacDonald, *Biophys. J.* **2003**, *85*, 1585.

[116] A. Witkowska, R. Jahn, *Biophys. J.* **2017**, *113*, 1251.

[117] T.-T. Kliesch, J. Dietz, L. Turco, P. Halder, E. Polo, M. Tarantola, R. Jahn, A. Janshoff, *Sci. Rep.* **2017**, *7*, 12070.

[118] S. Trier, J. R. Henriksen, T. L. Andresen, *Soft Matter* **2011**, *7*, 9027.

[119] J. Solon, J. Pecreaux, P. Girard, M. C. Faure, J. Prost, P. Bassereau, *Phys. Rev. Lett.* **2006**, *97*, 098103.

[120] D. T. Chiu, C. F. Wilson, A. Ryttsen, A. Stromberg, C. Farre, A. Karlsson, S. Nordholm, A. Gaggar, B. P. Modi, A. Moscho, R. A. Garza-Lopez, O. Orwar, R. N. Zare, *Science* **1999**, *283*, 1892.

[121] T. Robinson, P. E. Verboket, K. Eyer, P. S. Dittrich, *Lab Chip* **2014**, *14*, 2852.

[122] N. Bezlyepkina, R. S. Gracia, P. Shchelokovskyy, R. Lipowsky, R. Dimova, *Biophys. J.* **2013**, *104*, 1456.

[123] S. Kulin, R. Kishore, K. Helmerson, L. Locascio, *Langmuir* **2003**, *19*, 8206.

[124] P. Yang, R. Lipowsky, R. Dimova, *Small* **2009**, *5*, 2033.

[125] J. Nikolaus, M. Stöckl, D. Langosch, R. Volkmer, A. Herrmann, *Biophys. J.* **2010**, *98*, 1192.

[126] J. Heuvingh, F. Pincet, S. Cribier, *Eur. Phys. J. E* **2004**, *14*, 269.

[127] A. C. Saito, T. Ogura, K. Fujiwara, S. Murata, S. M. Nomura, *PLoS One* **2014**, *9*.

[128] A. Strömberg, F. Ryttsén, D. T. Chiu, M. Davidson, P. S. Eriksson, C. F. Wilson, O. Orwar, R. N. Zare, *Proc. Natl. Acad. Sci. USA* **2000**, *97*, 7.

[129] B. M. Discher, Y. Y. Won, D. S. Ege, J. C. M. Lee, F. S. Bates, D. E. Discher, D. A. Hammer, *Science* **1999**, *284*, 1143.

[130] J. C. M. Lee, M. Santore, F. S. Bates, D. E. Discher, *Macromolecules* **2002**, *35*, 323.

[131] R. Dimova, U. Seifert, B. Pouligny, S. Forster, H. G. Dobereiner, *Eur. Phys. J. E* **2002**, *7*, 241.

[132] Y. F. Zhou, D. Y. Yan, *J. Am. Chem. Soc.* **2005**, *127*, 10468.

[133] W. Su, Y. H. Luo, Q. Yan, S. Wu, K. Han, Q. J. Zhang, Y. Q. Gu, Y. M. Li, *Macromol. Rapid Commun.* **2007**, *28*, 1251.

[134] T. P. Smart, C. Fernyhough, A. J. Ryan, G. Battaglia, *Macromol. Rapid Commun.* **2008**, *29*, 1855.

[135] J. W. Szostak, D. P. Bartel, P. L. Luisi, *Nature* **2001**, *409*, 387.

[136] P. L. Luisi, T. P. de Souza, P. Stano, *J. Phys. Chem. B* **2008**, *112*, 14655.

[137] J. E. Rothman, F. T. Wieland, *Science* **1996**, *272*, 227.

[138] F. Caschera, T. Sunami, T. Matsuura, H. Suzuki, M. M. Hanczyc, T. Yomo, *Langmuir* **2011**, *27*, 13082.

[139] K. P. Browne, D. A. Walker, K. J. M. Bishop, B. A. Grzybowski, *Angew. Chem., Int. Ed.* **2010**, *49*, 6756.

[140] F. Caschera, S. Rasmussen, M. M. Hanczyc, *ChemPlusChem* **2013**, *78*, 52.