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Advances in block copolymer-phospholipid hybrid vesicles: from physical–chemical properties to applications

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Hybrid vesicles, made of lipids and amphiphilic block copolymers, have become increasingly popular thanks to their versatile properties that enable the construction of intricate membranes mimicking cellular structures. This tutorial review gives an overview over the different hybrid vesicle designs, and provides a detailed analysis of their properties, including their composition, membrane fluidity, membrane homogeneity, permeability, stability. The review puts emphasis on the application of these hybrid vesicles in bottom-up synthetic biology and aims to offer an overview of design guidelines, particularly focusing on composition, to eventually realize the intended applications of these hybrid vesicles.

1 Introduction

Over the past 20 years since the first reports on hybrid vesicles (HVs),^{1–4} ongoing research has illustrated that the incorporation of amphiphilic block copolymers into lipid bilayers changes the properties of the vesicles, *e.g.*, the stability is enhanced in comparison to liposomes. The combination of block copolymers with lipids not only provides greater design flexibility but also preserves the self-assembly capabilities of the vesicles. Consequently, these HVs are gaining popularity as versatile platforms for drug delivery applications and synthetic biology.⁵ Applications of HVs in synthetic biology have been summarized in several reviews focusing on the obtained artificial systems.^{6–9} In particular, HVs hold great promise as an alternative for assembling plasma membranes imitates, especially when precise control over the membrane properties is essential for the development of synthetic life-like systems.^{10–14} Recently, Reimhult *et al.*¹⁵ reviewed the use of HVs for drug delivery applications comparing selected properties. Further, HVs appeared in many reviews focusing on lipid-based drug delivery vehicles, where they are highlighted as game changing prospects owing to their versatility.^{16–21} However, HVs in a descriptive manner were last time summarized by Schulz *et al.*^{22,23} and Le Meins *et al.*²⁴ almost 10 years ago and only a selection of assemblies was mentioned by Go *et al.*²⁵

The aim of this tutorial review is to provide a detailed overview of the fundamental aspects as well as the recent advances and achievements in the area of HV assembly. We focus on various aspects of the vesicle and membrane properties. We aim to conduct a comprehensive comparison of numerous assemblies, with the intention of deriving conclusions that can

provide inform for design criteria for an optimal synthetic cell membrane. Throughout this tutorial review, HV assemblies are referred to by their composition to highlight the used polymer & lipid.

2 Assembly and composition

To date, various polymers have been combined with phospholipids to assemble HVs and many have demonstrated long-term stability. Table 1 provides a summary of these assemblies including the type and ratio (mol% or w/w%) of their building blocks and the references to the publications. The choice of lipids typically depends on the study's objectives or application. Fully saturated lipid such as 1,2-dipalmitoyl-*sn*-glycero-3-phosphocholine (DPPC) or unsaturated zwitterionic lipids such as 1-palmitoyl-2-oleoyl-*sn*-glycero-3-phosphocholine (POPC), 1,2-dioleoyl-*sn*-glycero-3-phosphocholine (DOPC) or *L*- α -phosphatidylcholine (Soy) (Soy PC) are often the first choices. The most commonly used polymer for the hydrophilic block is poly(ethylene glycol) (PEG) or referred to poly(ethyl oxide) (PEO) in some publications. In contrast, the polymers used as the hydrophobic block are more diverse including poly(1,2-butadiene) (PBd), poly(isobutylene) (PiB), poly(dimethyl siloxane) (PDMS) and poly(cholesterol methacrylate) (PCMA). The chemical structures of the lipids and the polymers are summarized in Scheme 1.

The block copolymer molecular mass typically falls within the range of 1.2 to 20 kDa. Block copolymers with low molecular mass share a more comparable size to the lipid counterpart. They were early on assumed to construct vesicles with a similar curvature, and hence being more suitable to form HVs than high molecular weight block copolymers. Even though it seems that an apparent size mismatch may lead to higher degree of phase separation between the lipids and the block copolymers

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Table 1 Composition of hybrid vesicles^a

Polymer	Lipid	Polymer content	Ref.
PMOXA ₂₁ - <i>b</i> -PDMS ₇₃ - <i>b</i> -PMOXA ₂₁	DPPC	10–60 mol%	1
PDMS ₆₀ - <i>b</i> -PMOXA ₂₁	DMPC	90 mol%	26
PMOXA ₁₅ -PDMS ₆₇ -PMOXA ₁₅ PMOXA ₆ -PDMS ₃₃ -PMOXA ₆	DOPC		27
PDMS- <i>g</i> -PEG ₂	DPPC, DOPC		28
PDMS ₂₂ - <i>g</i> -(PEG ₁₆) ₂	Soy PC	60–100 mol%	29–34
	DOPC		
	DPPC DOTAP		
	Soy PS		
PDMS ₁₆ - <i>g</i> -(PEG ₁₂) ₂	DPPC	0–100 w/w%	35–37
PEG ₈ - <i>b</i> -PDMS ₂₂ - <i>b</i> -PEG ₈	POPC		
PEG ₁₂ - <i>b</i> -PDMS ₄₃ - <i>b</i> -PEG ₁₂			
PEG ₁₇ - <i>b</i> -PDMS ₆₇ - <i>b</i> -PEG ₁₇			
PDMS ₃₆ - <i>b</i> -PEG ₂₃	DOPC	50, 80 w/w%	38
PDMS ₂₃ - <i>b</i> -PEG ₁₃	POPC	75–90 w/w%	39–42
PDMS ₂₇ - <i>b</i> -PEG ₁₇			
PDMS ₃₆ - <i>b</i> -PEG ₂₃			
PEG ₈ - <i>b</i> -PDMS ₂₂ - <i>b</i> -PEG ₈	POPC		41
PEG ₁₂ - <i>b</i> -PDMS ₄₃ - <i>b</i> -PEG ₁₂			
PEG ₃ - <i>b</i> -PDMS ₁₅ - <i>b</i> -PEG ₃	DOPC	0.05–0.18 wt%	43
PEG ₁₄ - <i>b</i> -PDMS ₁₅ - <i>b</i> -PEG ₁₄			
PEG ₂₈ - <i>b</i> -PDMS ₁₅ - <i>b</i> -PEG ₂₈			
PBd ₂₂ - <i>b</i> -PEG ₁₄	HSPC	75, 90 mol%	44
	DOTAP	75 mol%	45
	POPC, EcCl	5–75 mol%	46–53
	Soy PC	50 mol%	54
PBd ₁₂ - <i>b</i> -PEG ₁₁	POPC	25–75 mol%	51–53
PBd ₄₆ - <i>b</i> -PEG ₃₀	POPC	20–90, mol%	55
PBd ₁₂ - <i>b</i> -PEG ₉	DOPS	50%	56
PBd ₃₂ - <i>b</i> -PEG ₂₁	DOPC		
PBd ₃₇ - <i>b</i> -PEG ₂₂	DOPC	10–50 mol%	58
PBd ₂₃ - <i>b</i> -PEG ₁₃	DOTAP, DOPS, DOPC	20–80%	59 and 60
PBd ₃₃ - <i>b</i> -PEG ₂₂			
PBd ₃₇ - <i>b</i> -PEG ₅₅			
PBd ₇₃ - <i>b</i> -PEG ₄₄			
PBd ₂₂ - <i>b</i> -PEG ₂₃	DPPC, POPC	70 w/w%	61 and 62
PBd ₁₁ - <i>b</i> -PEG ₈	DPPC, POPC egg SM		63–65
PBd- <i>b</i> -PEG	POPC	Simulation	66
PBd ₁₂ - <i>b</i> -PEG ₁₀	DOPC, DOPS	50–99 mol%	67
	DOPE		
PBd ₄₃ - <i>b</i> -PEG ₂₀	DPPC	35, 65%	68
PIB ₈₇ - <i>b</i> -PEG ₁₇	DPPC	50 mol%	69–71
PIB ₃₇ - <i>b</i> -PEG ₄₈	DOPC		
PI- <i>b</i> -PEG (877 Da)	DPhPC	90 mol%	72
PI- <i>b</i> -PEG (1530 Da)			
PI- <i>b</i> -PEG (3188 Da)			
PEG ₁₁₃ - <i>b</i> -PCL ₄₆	DPPC	5, 10 mol%	73 and 74
PEG ₁₁₃ -PCL ₅₀			
PEG ₁₁₃ - <i>b</i> -PCL ₂₃			
PCL ₁₂ -PEG ₄₅ -PCL ₁₂	DPPC	0.001–2 mol%	75
PCL ₁₆ -PEG ₁₀₄ -PCL ₁₆			
PEG ₁₂ - <i>b</i> -PCL ₉	POPC	50 mol%	76
PEG ₁₂ - <i>b</i> -PCL ₆			
PEG ₄₅ - <i>b</i> -PCL ₄₄	DPPC	20 mol%	77
PEG ₁₅ - <i>b</i> -PLA ₂₅	POPC	50 mol%	78
PEG ₄₅ - <i>b</i> -PLA ₅₄			
PEG ₁₂ - <i>b</i> -PCL ₂₅			
PEG ₄₅ - <i>b</i> -PCL ₂₅			
PEG ₄₅ - <i>b</i> -PCL ₄₂			
PEG ₁₁₄ - <i>b</i> -PCL ₈₈	DPPC	10 ⁻³ g L ⁻¹ , 0.2 g L ⁻¹	79
PEG ₁₁₄ - <i>b</i> -P(CL ₄₂ - <i>co</i> -LA ₈₅)			
PEG ₁₁₄ - <i>b</i> -P(CL ₁₄ - <i>co</i> -LA ₁₃₈)			
PEG ₁₁₄ - <i>b</i> -PLA ₃₂₈			
P(IPAm- <i>co</i> -DMAAm)- <i>b</i> -PLA	Diyne PC(DC8,9 PC)	27 w/w%	80



Table 1 (Contd.)

Polymer	Lipid	Polymer content	Ref.
POEGA- <i>b</i> -PLauA	HSPC		81
PNIPAM ₁₀₆ - <i>b</i> -PLauA ₂₅	DPPC	0.2–10 mol%	82
PNIPAM ₂₈ - <i>b</i> -PLauA ₁₃			
PDMAEMA ₃₃ - <i>b</i> -PLMA ₁₅	HSPC		83
PDMAEMA ₃₂ - <i>b</i> -PLMA ₂₃			
PDMAEMA ₃₁ - <i>b</i> -PLMA ₈	EPC	10 mol%	84
PDMAEMA ₅₈ - <i>b</i> -PLMA ₂₃			
PEG ₁₁₃ -P(8C-Chol) <i>m m</i> = 1.4, 2.2, 2.8, 3.3, 3.7, 4, 8	DPPC	5 mol%	85
PCMA- <i>b</i> -PDMAEMA	POEPC, POPS, POPE, POPC	90 w/w%	86 and 87
PCMA ₁₁ - <i>b</i> -PEG ₁₁₃	POPC/POPS		88
PCMA ₁₁ - <i>b</i> -P(METMA ₇ - <i>co</i> -CEA ₁₀₄)	DOPC	27 w/w%	89
PCMA ₁₀ - <i>b</i> -PCEA ₁₇₀	DOPC, DOPC, DOPE	10 w/w%	90–92
P(CMA ₉ - <i>co</i> -BuMA ₉)- <i>b</i> -PCEA ₁₈₆	Soy PC	1–90 w/w%	93
P(CMA ₈ - <i>co</i> -HEMA ₃)- <i>b</i> -PCEA ₁₈₆			
PBuMA ₅₀ - <i>b</i> -PCEA ₁₇₅			
PHEMA ₅₈ - <i>b</i> -PCEA ₁₅₀			
PCMA- <i>b</i> -PMPC	DOPS, DOPC, DOPE		94
Oligo(aspartic acid) ₄ - <i>b</i> -PPO ₄₃	DOPC	10 mol%	95
Maltopentaose- <i>b</i> -PPO ₄₃	DOPC	10–90 mol%	96

^a Abbreviations: Lipids: DPPC: 1,2-dipalmitoyl-*sn*-glycero-3-phosphocholine; DMPC: 1,2-dimyristoyl-*sn*-glycero-3-phosphocholine; DOPC: (1,2-dioleoyl-*sn*-glycero-3-phosphocholine); DOTAP: 1,2-dioleoyl-3-trimethylammonium-propane (chloride salt); Soy PC (*L*- α -phosphatidylcholine (Soy)); Soy PS: anionic soy PS; POPC: 1-palmitoyl-2-oleoyl-*sn*-glycero-3-phosphocholine; EcCL: *E. coli* lipid extract and cholesterol; HSPC: *L*- α -phosphatidylcholine, hydrogenated (soy); DPhPC: 1,2-diphytanoyl-*sn*-glycero-3-phosphocholine; DOPS: 1,2-dioleoyl-*sn*-glycero-3-phospho-L-serine (sodium salt); DOPE: 1,2-dioleoyl-*sn*-glycero-3-phosphoethanolamine; Diyne PC(DC8,9PC): 1,2-bis(10,12-tricosadiynoyl)-*sn*-glycero-3-phosphocholine; egg SM: sphingomyelin (egg, chicken); EPC: *L*- α -phosphatidylcholine (egg, chicken) polymer; PMOXA: polymethyloxazoline; PDMS: poly(dimethyl siloxane); PEG: poly(ethylene glycol); Pbd: polybutadiene; PIB: poly(isobutylene); PI: polyisoprene; PCL: poly(ϵ -caprolactone); PLA: poly(lactic acid); APOEGA: poly(oligoethylene glycol acrylate); PLauA: poly(lauryl acrylate); PIPPAm or PNIPAM: poly(*N*-isopropyl amide); PLMA: poly(lauryl methacrylate); PCMA: poly(cholesterol methacrylate); DMAAm or PDMAEMA: poly(dimethyl aminoethyl methacrylate); PCEA: poly(2-carboxyethyl acrylate); PHEMA: poly(2-hydroxyethyl methacrylate); BuMA: poly(*i*-butyl methacrylate); PMPC: poly(2-methacryloyloxyethyl phosphorylcholine); PPO: poly(propylene oxide).

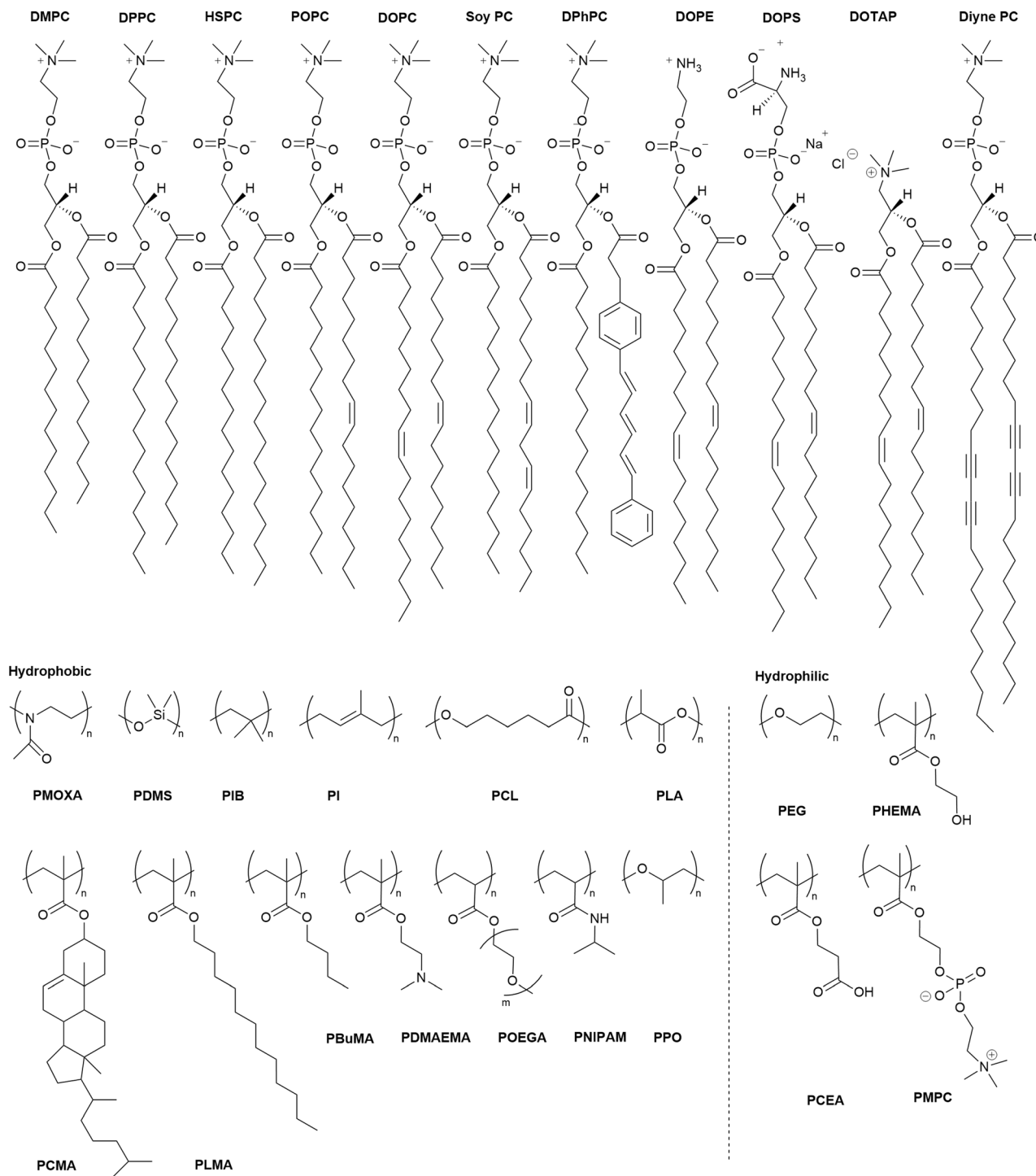
(see later for details), using low molecular weight block copolymer is not a requirement. However, the properties of the HV are most likely to depend on the length of the block copolymer chains and their ability to mix in various ratios (Scheme 2).

Although small HVs and large HVs are spherical assemblies with sizes below 100 nm and 100 nm and below 1 μ m, respectively, this review will refer to both these sizes of vesicles as smaller HV (sHV). Often authors tend to use small and giant as a description to distinguish between the reported vesicle types, that is misleading nomenclature. sHVs are typically assembled using the thin film hydration method, which is well established for liposomes and polymersomes formation.⁵ Briefly, the block copolymer and lipid components are mixed in an organic solvent followed by solvent evaporation either under vacuum on a rotary evaporator or with a stream of nitrogen while rotating the flask, to create a thin lipid-polymer film that is left to dry under vacuum for 4–12 h. The film is rehydrated with aqueous solution (pure water, buffer, cell media *etc.*) that can contain cargo for encapsulation into the void of the vesicles. The assemblies are then subjected to either sonication or extrusion through polycarbonate membranes to form unilamellar sHVs in the range of 50–150 nm with narrow polydispersity. sHVs are often passed through a size exclusion column to remove micelles or non-encapsulated cargo. Complementarily, giant unilamellar HVs (gHVs) (sizes above 1 μ m) are usually assembled *via* electroformation where a thin polymer lipid film is

deposited on the surface of an electrode, usually an ITO coated glass slide. An electric current is applied during the rehydration step to induce the controlled formation of gHVs. Aqueous media with low conductivity (*e.g.*, distilled water) can only be used, therefore, sucrose or glucose solutions are often considered to fill the gHVs in order to allow for their subsequent exposure to physiological buffers without rupturing due to osmotic imbalance between the gHVs void and the environment.⁹⁷ Emulsion-based methods,⁹⁸ which are often used for giant liposome assembly, are not common for gHVs formation. However, this method was used to form asymmetrical vesicles.⁹⁹

The precise process by which block copolymers insert into the lipid bilayer remains debated. It is generally assumed that the components simultaneously detach from the surface during the rehydration phase and form a vesicular structure due to the hydrophobic effect.¹⁰⁰ Nonetheless, evidence suggests that block copolymers can incorporate into pre-existing liposome membranes. For example, Mineart *et al.*⁸⁵ demonstrated that the block copolymer PEG₁₁₃-*b*-P(8C-Chol) could be incorporated into the lipid bilayer post-assembly by adding a micelle solution to pre-made liposomes. Similarly, PCMA-*b*-PCEA-GALA dissolved in DMSO was inserted into pre-made DOPC liposomes.⁹¹ Henderson *et al.*⁵⁷ showed that Pbd-*b*-PEG polymersomes could fuse with DOPC liposomes when the mixture was subjected to high salt concentration and mechanical stimuli.



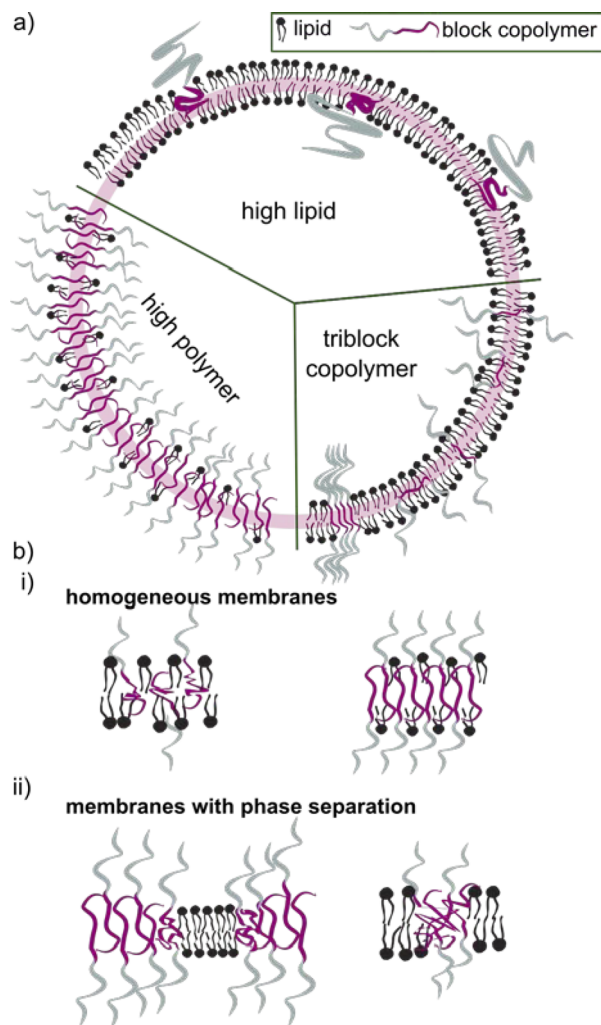


Scheme 1 Chemical structures of the lipids (top row) and the hydrophilic and hydrophobic polymers used in the block copolymers. PDMAEMA is also used as a hydrophilic block. For abbreviations meaning see Table 1.

The successful formation of HVs is typically confirmed through various methods including assessing Förster Resonance Energy Transfer (FRET) effects, measuring changes in zeta potential,^{60,88} determining membrane thickness in recorded cryogenic electron microscopy (cryo-EM) images, utilizing small angle X-ray scattering (SAXS) and monitoring alterations in various membrane properties.

Determining alterations from the theoretical expected membrane composition is important when the effect of the building blocks on the resulting membranes properties is investigated. However, confirming the exact constitution is challenging. The composition of the assemblies after rehydration are commonly reported to be the same as the ratio of the polymer and the lipid in the deposited thin films. However, this





Scheme 2 (a) Schematic representation of hybrid vesicles composed of lipids and block copolymers with various lipid block copolymer ratios. (b) Cartoon of the distribution of lipids and block copolymers in either a homogeneous membrane (i) or a membrane that shows phase separation (ii).

is not necessarily the case as it was pointed out by Dao *et al.* when studying the effect of the polymer's molecular weight on PEG-*b*-PDMS-*b*-PEG & DPPC-based assemblies.³⁶ Cryo-EM images revealed the coexistence of various types of structures such as HVs, micelles, liposomes, and polymersomes. They argued that this finding was the result of the high line tension originating from the differences in thickness of the polymer-rich vs. lipid-rich areas. Similar observations were made for PBD-*b*-PEG & DPPC-based⁶⁸ and poly(oligoethylene glycol acrylate)-*b*-poly(lauryl acrylate) POEGA-*b*-PLauA & hydrogenated 1- α -phosphatidylcholine (HSPC)-based⁸³ assemblies. We also showed recently that assemblies made of P(CMA-*co*-butyl methacrylate (BuMA))-*b*-PCEA or P(CMA-*co*-hydroxyethyl methacrylate (HEMA))-*b*-PCEA & soy PC⁹³ and PCMA-*b*-poly(2-methacryloyloxyethyl phosphorylcholine) (PMPC) & DOPC⁹⁴ had a micellar fraction present when more than ~1 mol% block copolymer was used. However, we could only speculate if the micellar fraction contained lipids or consisted of block

copolymers only. Furthermore, cryo-EM analysis of the vesicles indicated the presence of multiple populations (liposomes, HVs and polymersomes) in several other studies based on membrane thickness analysis.^{30,36,40}

3 Size and composition of smaller HVs (sHVs)

The sizes of sHVs are determined usually by dynamic light scattering (DLS) and reported as hydrodynamic diameter (D_h) and polydispersity (PDI), which ranges, typically, between 40–400 nm with PDI of 0.1–0.3. Several studies have been carried out to determine the influence of the composition on the size of the HVs by changing the molecular weight of the block copolymer, by varying the block copolymer to lipid ratios, or by incorporating charged lipids among others. Determining the size of the sHVs is a commonly performed first characterization effort, which gives an indication of successful HV formation. PDI values above 0.3 are usually considered too polydisperse, suggesting assembly of non spherical vesicles or the presence of multiple populations. The curvature of the liposomes depends on the length of the lipid molecules, *i.e.*, longer chains produce liposomes with larger diameters (D_h). Similarly, longer chain block copolymers form polymersomes with larger diameters. Therefore, the hydrophobic chain length is expected to influence the diameter of the sHVs due to the size mismatch between the lipids and block copolymers.

The amount of the block copolymer content also affects the sHV size. For instance, increasing the PBD₃₇-*b*-PEG₂₂ content in PBD₃₇-*b*-PEG₂₂ & DOPC from 10 to 50 mol% increased the D_h from 150 to 200 nm.⁵⁸ Similarly, the incorporation of higher amounts of poly(ϵ -caprolactone) (PCL)_{*n*}-*b*-PEG_{*m*}-*b*-PCL_{*n*} into DPPC vesicles increased the D_h from 112 to 243 nm.⁷⁵ In addition, the D_h of PB₂₂-PEG₁₄ & POPC-based HVs with 25, 50 or 75% block copolymer content were 60–80% smaller than the respective liposomes and polymersomes.⁴⁶

However, there are also examples where the changes in D_h due to variations in composition were very small. For instance, the D_h slightly decreased with increasing the block copolymer content from 10 to 25 w/w% for PCMA₁₁-*b*-PEG₁₁₃ & POPS-based HVs.⁸⁸ Similarly, D_h of PBD₂₂-*b*-PEG₁₄ & *E. coli* lipid extract-based HVs only changed a few nm when the block copolymer content was increase from 25 to 75 mol%.⁴⁸ In another study, increasing the amount of PMOXA₆-PDMS₃₃-PMOXA₆ from 10 to 20 mol% did not significantly affect the D_h of the resulting HVs. However, increasing the molecular weight of this block copolymer to PMOXA₁₅-*b*-PDMS₆₇-*b*-PMOXA₁₅ led to an increase in D_h from 80 to 120 nm, which further increased by doubling the amount of block copolymer.²⁷ This trend implies that the longer block copolymer chain prefers to form a vesicle with higher curvature. However, a contradictory example reported no change in D_h of approximately 100 nm for sHVs made of POPC and either PEG₁₅-*b*-poly(lactic acid)PLA₂₅ or PEG₄₅-*b*-PLA₅₄.⁷⁸ It should be noted that low block copolymer content might not impact the vesicles curvature. In the same study, HVs made of either PEG₄₅-*b*-PCL₂₅ or PEG₄₅-*b*-PCL₄₂ & POPC had a very broad range of D_h with



a main peak around 20 nm indicating the presence of a large micellar population.

The impact of the HV composition on the size is likely also associated with the interactions occurring between the block copolymer chains and the lipids. For instance, the size of the HVs was affected by the charge of the phospholipids (POPC vs. POPS vs. POEPC) when using PCMA₁₁-*b*-poly(dimethyl aminoethyl methacrylate)₇₇ (PDMAEMA), *i.e.*, HVs containing POPC and POPS were 10–25% smaller than the corresponding liposomes, while POEPC containing HVs were ~16% larger.⁸⁶ This observation was presumably due to the attraction or repulsion between the slightly positively charged PDMAEMA, and the charge of the phospholipids. In contrast, when a block copolymer with a zwitterionic hydrophilic block (PCMA-*b*-PSMA) was used for HV assembly, their size did not change significantly independent whether 1,2-dioleoyl-*sn*-glycero-3-phospho-L-serine (DOPS), DOPC or a mixture of 1,2-dioleoyl-*sn*-glycero-3-phosphoethanolamine (DOPE)/DOPC was employed.⁹⁴ In another example, by employing PCMA and its copolymers with BuMA or HEMA in the hydrophobic block, the size of the resulting HVs together with soy PC decreased from 178 nm to 150 nm upon the introduction of more hydrophilic monomers.⁹³ This reduction reached ~116 nm with an increasing block copolymer content. Here, we observed a micellar population, which might have influenced the measured D_h . Transmission electron microscopy (TEM) and cryo-EM images can also be used to gain information of the size and morphology of the HVs (Fig. 1). Recently, the effect of the polymer's hydrophilic length on the morphology of the vesicles was

studied. Cryo-EM images showed that when the hydrophilic block length increased from 0.12 to 1.2 kDa in PEG-*b*-PDMS-*b*-PEG, the resulting HVs with DOPC became more spherical from distorted sphere. However the PDI also increased.⁴³

4 Lipid-polymer phase separation

The incorporation of block copolymers and lipids into a mixed membrane is typically envisioned *via* the alignment of the hydrophobic tails of the lipids and the block copolymers. The hydrophilic segment of diblock copolymers either faces into the void of the HV or towards the environment. Triblock copolymers are expected to have one hydrophilic blocks facing the external environment, while the other is positioned in the internal void (Scheme 2b). Dissipative particle dynamics simulation showed that the arrangement of block copolymers in the lipid bilayer can be characterized as bridge- and loop-shapes in the membrane, resulting in symmetrical and asymmetrical inner and outer membrane leaflets. The arrangement was highly influenced by the hydrophilic length of the block copolymer as well as repulsive energies between the hydrophobic block and the lipid tails.¹⁰¹ Moreover, the resulting hybrid membrane can have the building blocks either homogeneously distributed or phase separated. Both types of membranes can be beneficial. Homogeneous hybrid membranes have better stability and reproducibility, while (lipid) phase separation and raft formation are essential in membranes of living organisms, and are therefore anticipated to contribute with unique functions to man-made membranes.¹⁰²

Different factors influence if a specific block copolymer and lipid combination results in a homogeneous or phase separated membrane. First, the fluidity of the lipid phase strongly affects the homogeneity of the hybrid membrane, *i.e.* lipids with melting temperature (T_m) below room temperature are more likely to facilitate homogeneous hybrid membranes. Second, the ratio between the block copolymer and lipids is an important parameter, and homogeneous membrane often require very specific ratios. An example in this context are gHVs made of the triblock copolymer PDMS₂₂-*g*-(PEG₁₆)₂ & POPC that had homogeneous membranes with high block copolymer content (75 mol%), while block copolymer amounts below 50 mol% resulted in macrodomain formation that quickly evolved to fission through budding of the vesicle.³³ In contrast, the same block copolymer & DPPC showed stable HVs with macrodomain formation for up to 75 mol% block copolymer content, and homogeneous membranes above 85 mol% block copolymer content. The difference was explained by the curvature energy and the line tension between the polymer-rich and the lipid-rich domains that caused budding of the vesicles. Furthermore, the HVs with DPPC were stable for longer time, due to the lower mobility of the DPPC lipids in the gel phase compared to POPC in the fluid state that allows faster movements of the components. Similarly, gHVs made of maltopentaose-*b*-poly(propylene oxide) (PPO)₄₃ & DOPC had a homogeneous membrane up to 50 mol% block copolymer content, while above domain formation was observed.⁹⁶

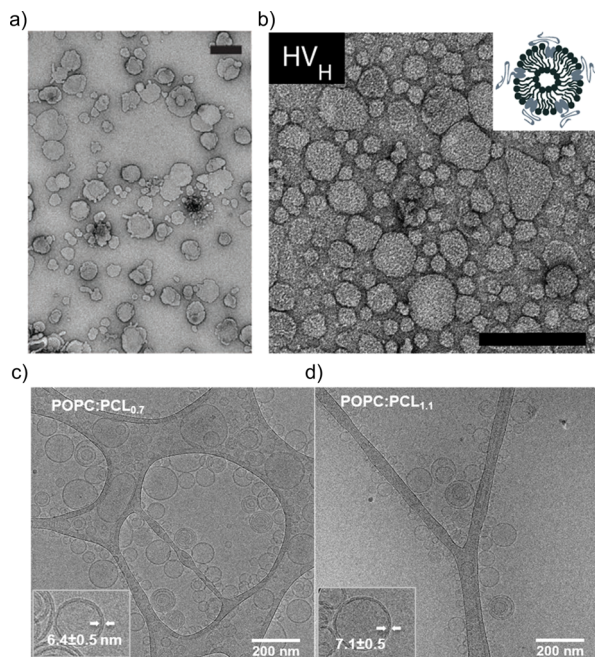


Fig. 1 Typical electron microscopy images of HVs. TEM images of (a) PCMA₁₁-*b*-P(META₇-co-CEA₁₀₄) & DOPC (Reprinted from ref. 89, Copyright 2024, with permission from Wiley) and (b) PCMA₁₁-*b*-PEG₁₁₃ & POPC (Reprinted from ref. 88, Copyright 2024, with permission from Elsevier). Cryo-EM images of (c) PEG₁₂-*b*-PCL₆ & POPC and (d) PEG₁₂-*b*-PCL₉ & POPC.⁷⁶ Used under CC BY 3.0. <https://www.mdpi.com/2073-4360/12/4/979>.



Hu *et al.* used dissipative particle dynamics simulation to model membrane formation with a block copolymer and saturated or non-saturated lipids. They found that phase separation is driven by a weak energy incompatibility between the lipids and the block copolymers. Above a critical lipid concentration, a mixing-demixing transition in the polymer-rich membrane can be observed, until a thermal condition in the hydrophobic layer is satisfied, with low thickness mismatch. This quality depends on the type of lipids and their T_m . Furthermore, significant thickness mismatch also drives phase separation.¹⁰³

FRET measurements are an interesting opportunity to characterize hybrid membranes. The existence of a FRET signal not only gives an indication that the two components are in close proximity to each other, *i.e.* being part of the same assembly,^{1,32,34–36,57,64,89–95} but also the presence of domains can be determined by calculating the FRET efficiency. It is important to note that even though FRET is used very often, it is a qualitative rather than a quantitative measure for domain formation as smaller domains might not be detected, and FRET can always occur at the lipid-polymer interface. FRET measurements supported the existence of lipid domains by using NBD-labelled block copolymers and rhodamine-labelled phospholipids in PDMS-*g*-PEG₂ & DPPC-based or PDMS-*g*-PEG₂ & DOPC-based HVs.³⁴ In a follow up study, the length of the blocks were systematically varied in the PEG_{*n*}-*b*-PDMS_{*m*}-*b*-PEG_{*n*} ($n = 8, 12$ or 17 and $m = 22, 43$ or 67) triblock copolymer, and the effect on the hybrid membrane when assembled into sHV with DPPC was assessed.³⁶ Phase separation was found for all tested compositions illustrated by a combination of small angle neutron scattering (SANS) and FRET measurements. Further, the sample contained liposomes, polymersomes and even worm-like micelles in addition to the sHVs with higher hydrophobic size mismatch of the hydrophobic block and the phospholipids,³⁶ highlighting the impact of the lipid fluidity on the phase separation (Fig. 2a). Membrane homogeneity for gHVs is often determined visually by confocal laser scanning microscopy (CLSM). In contrast, gHVs made of POPC & PEG₈-*b*-PDMS₂₂-*b*-PEG₈ had homogenous hybrid membranes below 10 w/w% lipid content, while gHVs consisting of POPC and higher molecular weight PEG₁₂-*b*-PDMS₄₃-*b*-PEG₁₂ showed homogenous hybrid membranes up to 40 w/w% lipids as observed by CLSM (Fig. 2b).³⁵ The decrease of the overall curvature (sHV *vs.* gHV) allowed for a better mixing of the building block. Macrodomain formation was observed for gHVs assembled from PIB₈₇-*b*-PEG₁₇ & DPPC or PIB₃₇-*b*-PEG₄₈ & DPPC. In the latter case, the DiDC₁₈ dye was incorporated in the hybrid membrane that showed a preference for the less ordered (polymer) phase causing the highly ordered lipid domains to appear dark in the CLSM images. Furthermore, gHVs made of PIB₃₇-*b*-PEG₄₈ & DPPC had an interesting, ragged surface at temperatures below the T_m of DPPC. In contrast, gHVs made of PIB₈₇-*b*-PEG₁₇ & DPPC showed defects and ragged membranes for low block copolymer content (10%), while smooth spherical gHVs were observed at higher block copolymer content.⁷⁰

Domain formation in HVs can be a temperature dependent effect. In particular, if the assembly temperature is raised above the T_m of saturated lipids for HV formation to allow for

sufficient mobility required for assembly, followed by cooling to room temperature. The extent of domain formation was observed to be influenced by the cooling rate after electroformation. Phase-separated gHVs made of PDMS-*g*-PEG₂ and DPPC at 53 °C *via* electroformation had distinct membrane tensions depending on the cooling rates after electroformation, *i.e.*, cooling rates 1 and 5 °C min⁻¹ resulted in low and high membrane tension respectively (Fig. 2c).²⁸ Similarly, applying slower cooling rates after gHVs formation using PBd₄₆-*b*-PEG₃₀ & DPPC resulted in larger domains.¹⁰⁴ In the case of slower cooling rates, the components have increased mobility, allowing them to attain a more energetically favourable arrangement before reaching the bilayer's gelation temperature, at which point the mobility of the components is substantially reduced.

Alternatively, domain formation can be induced post-assembly, *e.g.*, domain formation was induced using the affinity of biotin to neutravidin. The otherwise homogeneous HVs made of PBd₄₆-*b*-PEG₃₀ & POPC with incorporated biotinylated polymer were driven to form macrodomains in gHVs upon crosslinking with neutravidin.⁵⁵ However, when 5 mol% DSPE-PEG-biotin as a lipid was incorporated instead, no domain formation was induced. This difference was attributed to the fact that the hybrid system was not solely driven by the preferential biotin-protein interplay but also by the phospholipid-block copolymer interaction within the hybrid membrane. Furthermore, the addition of cholesterol can induced phase separation in homogeneous HVs,⁸⁶ or it can have a fluidizing effect to overcome nanodomain formation.⁷⁶ Interestingly, the introduction of cholesterol into a hybrid membrane composed of PBd₄₆-*b*-PEG₃₀ & DPPC induced the creation of domains exhibiting quasi-circular shapes, as opposed to the irregular patches observed in the absence of cholesterol.¹⁰⁴ This occurrence is likely attributed to cholesterol's impact on lipid packing and the associated transitions in lipid phases.

In an elegant study, Schulz *et al.* showed the advantage of using PIB₈₇-*b*-PEG₁₇ & DPPC-based HVs in creating biomimetic models to understand receptor-ligand recognition.⁷¹ Ganglioside GM1 modified lipids were incorporated in the HVs, and the lateral diffusion was monitored depending on the amount of block copolymer used during assembly. The binding of cholera toxin B to GM1 caused phase separation in the initially uniform membrane for block copolymer contents up to 30%. However, at block copolymer concentrations exceeding 40%, no binding occurred, presumably due to the steric hindrance caused by the block copolymer chains.

Nonetheless, there are several examples of phospholipid and block copolymer combinations where phase separation was not observed.^{26,29,30,39,44,61,72,76,89} In many of these cases, the study's objective did not focus on the identification of composition boundaries related to phase separation. Consequently, no general conclusions could be drawn regarding whether these components consistently form homogeneous membranes irrespective of the environmental conditions. Often, homogeneity of the hybrid membranes were determined based on the analysis of CLSM images when fluorescently labelled components were used for assembly. A recent observation highlighted that, despite having homogeneous membranes, the quantity of lipid and block copolymer varied among different gHVs. This



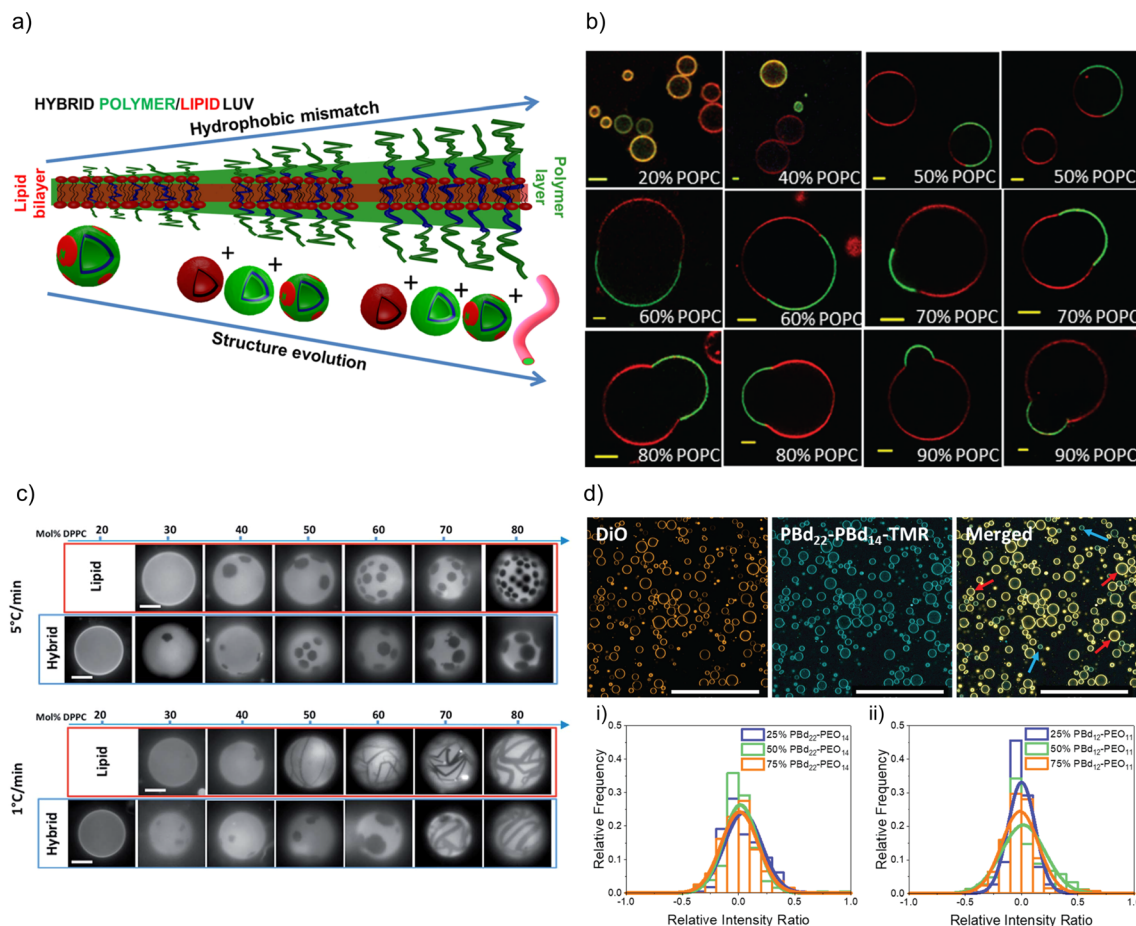


Fig. 2 Lipid and polymer distribution in HVs. (a) Schematics of observed structures with increasing length of polymer including the presence of polymersomes, liposomes, HVs and worm like micelles, using PDMS-*g*-PEG & DPPC.³⁶ Reprinted with permission from ref. 36. Copyright 2024 American Chemical Society. (b) Equatorial slices of hybrid 3K/POPC GHVs labelled with PDMS₂₆-*g*-(PEG₁₂)₂-FITC (green channel) and DOPE-Rhod (red channel) visualized by CLSM at room temperature. Scale bar: 5 μ m. Reprinted from ref. 35. Copyright 2024 the Royal Society of Chemistry. (c) Appearance of PDMS-*g*-PEG & DPPC ("hybrid") vesicles compared with DOPC/DPPC ("lipid") vesicles for different compositions and two cooling rates from the one-phase region. Scale bars: 10 μ m. DOPE-Rhod is employed as the tracer. Reprinted from ref. 28. Copyright 2024 the Royal Society of Chemistry. (d) Images of intensity contributions from DiO and, PBD₂₂-PEG₁₄-TMR dyes individually in 25 mol% PBD₁₂-PEG₁₁-based gHVs, and the merged channels. Red arrows indicate vesicles with greater DiO contribution, while blue arrows indicate vesicles with greater PBD₂₂-PEG₁₄-TMR contribution to the overall fluorescence intensity of the gHV. The histograms show the relative intensity contributions from DiO and PBD₂₂-PEG₁₄-TMR in (a) PBD₂₂-PEG₁₄/POPC and (b) PBD₁₂-PEG₁₁ & POPC-based HVs. An intensity ratio of 0 indicates equal relative fluorescence intensity contributions in HVs from both PBD₂₂-PEG₁₄-TMR and DiO. Scale bars: 200 nm.⁵² Used under CC BY 3.0. <https://pubs.rsc.org/en/content/articlelanding/2022/sm/d1sm01591e>.

conclusion was drawn based on CLSM images captured from a large number of gHVs within the same population. gHVs made of PBD-*b*-PEG & POPC⁵² and either P(CMA-*co*-BuMA)-PCEA or P(CMA-*co*-HEMA)-*b*-PCEA & Soy PC⁹³ showed different levels of fluorescent signal originating from the fluorescently labelled building blocks. This observation indicated that the quality of mixing of the components prior to drying of the lipid-block copolymer film on the electrode surface influenced the composition of the resulting gHVs. Further, there was no (or only very limited) mixing after the gHVs detached from the surface during electroformation (Fig. 2d).

5 Membrane properties

The plasma membrane of living cells has a diversity of properties, including its selective permeability, dynamic fluidity,

integral role in cellular communication, and active participation in maintaining cell structure and homeostasis. This fluidity allows the membrane to undergo dynamic movements, such as lateral diffusion, flexion, and fusion with other membranes. The lateral mobility of the lipid and protein components are essential to maintain the above mentioned cellular processes. Consequently, several important properties need to be considered when assembling artificial membranes that will collectively imitate important aspects of the plasma membrane.

5.1 Lateral mobility

The phospholipids in the bilayer of liposomes, which are often considered a very minimalistic model of the cell membrane, have free lateral diffusion depending on the type of phospholipids and/or the temperature. Consequently, phospholipids in



hybrid membranes also exhibit lateral motion that not only affected by the type of phospholipid and temperature, but also by the amount and type of block copolymer. Lateral mobility can be measured by fluorescence recovery after photobleaching (FRAP), where the reappearance of the fluorescent signal is followed after photobleaching of a selected area. FRAP is typically used for gHVs. Fluorescence correlation spectroscopy (FCS) is typically used for sHVs, where the diffusion of fluorophore containing molecule is detected. The fluorophores can be chemically attached to a lipid or the block copolymer, or inserted into the bilayer, hence its movement is related to the diffusion of the lipids and the block copolymers.¹⁰⁵ For instance, DPPC lipids in liposomes have limited lateral mobility at room temperature since the membrane is in the gel phase. However, when PIB₈₇-*b*-PEG₁₇ was incorporated in a DPPC membrane, the DPPC diffusion coefficient increased with increasing polymer content, indicating that the incorporation of the block copolymer affected the lipid organization, resulting in a more mobile system.⁷¹ In the hybrid membrane with 40 mol% block copolymer DPPC had 25% higher lateral mobility compared to membranes with 18 mol% polymer. In contrast, POPC lipids in liposomes have high lateral mobility at room temperature ($D \sim 9.8 \pm 1.7 \mu\text{m}^2 \text{s}^{-1}$) since the lipid bilayers is in the liquid phase. A $\sim 45\times$ decrease in probe's diffusion coefficient was found when increasing amounts of PBD₄₆-*b*-PEG₃₀ were incorporated, to $0.22 \pm 0.06 \mu\text{m}^2 \text{s}^{-1}$.⁵⁵ Hybrid membranes not only have lateral diffusion of the phospholipids but the incorporated block copolymer also exhibit mobility. For instance, as the block copolymer content decreased from 90 to 50 mol% in membranes composed of PEG₈-*b*-PDMS₂₂-*b*-PEG₈ & POPC, the diffusion coefficient of a polymer-based probe decreased by a factor of 2 (from 3.55 ± 0.79 to $1.70 \pm 0.32 \mu\text{m}^2 \text{s}^{-1}$).³⁷ While when the longer PEG₁₂-*b*-PDMS₄₃-*b*-PEG₁₂ was incorporated with POPC into a hybrid membrane, the diffusion coefficient did not significantly change and was found between 1.41 ± 0.20 and $2.05 \pm 0.41 \mu\text{m}^2 \text{s}^{-1}$. The lipids diffused about $2.5\times$ faster than the block copolymer in the same hybrid membrane.

5.2 Phase transition (melting) temperature (T_m)

The fundamental relevance of the phase transition temperature (T_m) in lipid membranes as well as the glass transition (T_g) and the crystallization temperature (also called melting temperature and referred as T_m) in block copolymers defines the properties of the hybrid membrane, potentially influencing the HVs behavior during storage at 4 °C and during application at room temperature or 37 °C. Therefore, understanding the influence of the block copolymer insertion into the lipid bilayer and its effect on the packing of the lipids is an important question to address. T_m is often determined by measuring the melting enthalpy using differential scanning calorimetry (DSC), but other methods can also be employed (see below in the section about general polarization). For instance, the lipids' main transition peak, occurring at around 40 °C, was diminished and a second peak appeared when considering sHV made of PMOXA₂₁-*b*-PDMS₇₃-*b*-PMOXA₂₁ & DPPC¹ (Fig. 3). This observation was attributed to a change in

the block copolymer's conformation as it was inserted into the membrane. The second peak became more distinct from the actual main transition peak of DPPC when the block copolymer content increased up to 40 mol%, while the temperatures of both peaks remained unaffected by the increase in block copolymer concentration. T_m remained unchanged with respect to the lipid-polymer ratio for gHVs made of PDMS-*g*-PEG & POPC and PDMS-*g*-PEG & DPPC for up to 70 mol% block copolymer content. However, above this threshold, the lipids were unable to meet and nucleate the gel phase, causing the melting transition of the lipids to disappear. Additionally, the melting enthalpy decreased as the block copolymer content increased, from 30 J g^{-1} to 15 J g^{-1} and from 52 J g^{-1} to 20 J g^{-1} for gHVs of PDMS-*g*-PEG & POPC and PDMS-*g*-PEG & DPPC, respectively.³³ The melting temperature of sHVs made of either PMOXA₁₅-PDMS₆₇-PMOXA₁₅ or PMOXA₆-*b*-PDMS₃₃-*b*-PMOXA₆ & DOPC decreased compared to pure DOPC liposomes with increasing amount of block copolymer up to 10 mol% from -17.2 to -19.2 °C.²⁷

Some polymers such as PLA and PCL have a glass transition temperature and a crystallization temperature in the range of interest for biomedical applications. The typical transition behavior of these pure polymers was also observed in HVs made of PEG₁₂-*b*-PCL₁₀ & POPC, indicating that these phase transitions are not significantly affected by the presence of the lipids. The authors argued that the lack of change in T_m is due to phase separation of the components. sHVs with higher molecular weight block copolymers showed subtle features consistent with their melting transition, and the double transition peak maximums were shifted for PEG₄₅-*b*-PCL₄₂ from 36 °C and 46 °C to 32 °C and 44 °C, and for PEG₄₅-*b*-PCL₂₅ from 34 °C to 50 °C. In the same study, the T_m of the block copolymer in PEG_{*n*}-*b*-PLA_{*m*} & POPC was not observed.⁷⁸ Similarly, using PEG_{*m*}-*b*-PCL_{*n*} & DPPC in the sHVs the presence of the polymers led to a significant increase of the main transition temperature (from 41 °C to 54 °C).⁷⁴ The main transition enthalpy decreased significantly by the incorporation of polymer into DPPC liposomes. In sHVs composed of either PCL₁₂-*b*-PEG₄₅-*b*-PCL₁₂ or PCL₁₆-*b*-PEG₁₀₄-*b*-PCL₁₆ & DPPC no change in T_m at low block copolymer content (up to ~ 24 w/w%) was observed. However, T_m was reduced by

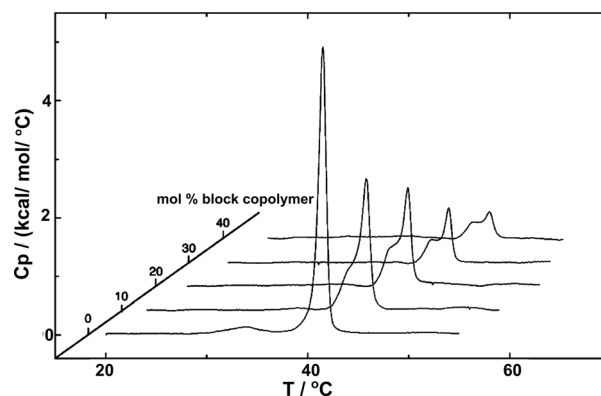


Fig. 3 DSC experiments showing the endotherms of HVs made of DPPC with increasing PMOXA₂₁-*b*-PDMS₇₃-*b*-PMOXA₂₁ content. Reprinted with permission from ref. 1. Copyright 2024 American Chemical Society.



5 °C for a ~38 w/w% block copolymer content in the HVs.⁷⁵ The latter observation indicated that the concentration of the block copolymer was essential to have an effect on the packing of the DPPC lipids. In another case, where HVs made of POEGA-*b*-PLauA & soy (HSPC) (a saturated lipid) were investigated, the pre-transition peak broadened and the transition enthalpy decreased with increased block copolymer concentration.⁸¹ The main peak position did not change significantly.

An interesting type of HVs considered the integration of thermo-responsive PNIPAM, in particular PNIPAM₁₀₆-*b*-PLauA*₂₅/PNIPAM₂₈-*b*-PLauA₁₃ & DPPC.⁸² DSC heating curves revealed differences in the 1st and 2nd heating cycles, with the T_m of the lipid shifting to higher values (above 41 °C), indicating structural rearrangements. However, the contribution of the thermo-responsive phase transition of PNIPAM at 32 °C was unfortunately not detectable during the heat cycles, presumably due to its low concentration.

5.3 Membrane fluidity

The fluidity of cell membranes is crucial for various cellular functions, including signal transduction, membrane trafficking, and the activity of membrane-bound enzymes and receptors. For example, temperature, pH, and the presence of certain ions can influence the membrane fluidity. Laurdan is often used to report on membrane packing since Laurdan's fluorescent emission properties are sensitive to the polarity of its environment (hydration of the membrane), leading to a red shift in the Laurdan emission peak. Fluidity of the membrane can be expressed as the general polarization (GP), which is a relation between Laurdan's emission peaks at 440 nm and 490 nm. GP values close to 1 indicate a highly ordered environment, while values close to -1 are related to a hydrated, disordered environment.¹⁰⁶ Overall, measuring the GP is a valuable tool for understanding the biophysical properties of cell membranes as GP provides a quantitative measure of the fluidity or rigidity of lipid membranes. Changes in GP can be indicative of environmental factors affecting lipid vesicles.

GP values for HVs made of PDMS-*g*-(PEG)₂ & soy PC were found to be -0.43 compared to -0.28 for liposomes, indicating a more fluid membrane in the former case.³⁰ The GP of PDMS-*g*-(PEG)₂ & 1,2-dioleoyl-3-trimethylammonium-propane (DOTAP), PDMS-*g*-(PEG)₂ & anionic soy PS (soy PS) and PDMS-*g*-(PEG)₂ & DOPC also decreased compared to their respective liposomes. Furthermore, the amount of block copolymer in the assemblies influenced the degree of decrease in the measured GP.³²

In contrast, the addition of PBd-*b*-PEG did not change the GP of POPC-based gHV's neither with increasing molecular weight nor with increasing block copolymer content, showing similar hydration and membrane ordering for the different gHV's.⁵² The insertion of cholesterol increases ordering in unsaturated lipid membranes, and the effect of polymerized cholesterol was studied for comparison. No change in the GP was found for HVs made of PCMA-*b*-PMPC and various lipids (DOPC, DOPS or DOPS/DOPE) compared to the corresponding liposomes.⁹⁴ In contrast, PCMA-*b*-PCEA at lower concentration reduced the GP in HVs made with DOPC from ~-0.25 to -0.3, while the GP

increased to -0.10 at higher block copolymer content.⁹⁰ The same effect was found for P(CMA-*co*-BuMA)-*b*-PCEA & soy PC and P(CMA-*co*-HEMA)-*b*-PCEA & soy PC, indicating that the cholesterol units homogeneously distributed in the membrane of the vesicles.⁹³ At the biological relevant temperature, the addition of PEG₁₁₄-*b*-P(CL-*co*-LA)_{*m*} in DPPC with various hydrophobic block lengths slightly increased the GP compared to liposomes.⁷⁹ Similarly, the incorporation of PCL-*b*-PEG-*b*-PCL with DPPC into vesicles had only a small effect on the GP values at low polymer concentrations, but the GP decreased from ~0.55 to 0.45 at a high polymer concentration.⁷⁵ Increasing polymer content in HVs made of PBd₁₁-*b*-PEG₈ & DPPC resulted in decreased GP values from 0.6 to 0.38 (ref. 64) (Fig. 4). As mentioned above, GP values for DPPC lipid containing vesicles are often reported over a range of temperatures and are used for indicating the lipid melting transition temperature. The GP values upon heating reflect the phase transition, and the inflection point of the GP values measured with increasing temperature indicates the T_m . HVs made of PEG₁₁₄-*b*-P(CL-*co*-LA)_{*m*} with various hydrophobic block lengths and DPPC had a 3–4 °C higher T_m compared to DPPC liposomes.⁷⁹ The overall change in GP values of DPPC liposomes was between 0.6 to -0.15 upon heating from 20 to 65 °C with a sharp change at the T_m (41 °C, GP = 0). The addition of increasing amounts of PBd-*b*-PEG decreased the change in GP values upon heating, accompanied by a broader transition temperature.⁶⁴ Furthermore, the T_m decreased more than 20 °C at the highest block copolymer concentrations. Recently, a multipeak analysis of Laurdan emission spectra has been employed to provide deeper insights into the phase behavior of hybrid membranes in several publications. This approach revealed that alterations in both pH and salt concentration induced a shift in the transitional phase of sHV's composed of carboxylated PEG-*b*-PBd & egg sphingomyelin (egg-SM), while variations in molecular weight had no effect.⁶⁵ The application of multipeak analysis to

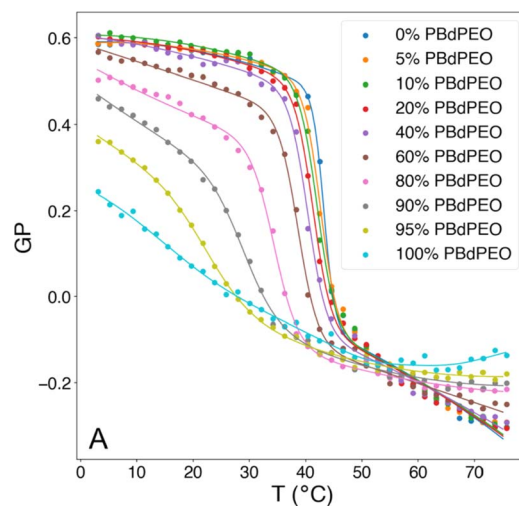


Fig. 4 Generalized polarization (GP) of Laurdan, corresponding to membrane polarity, as a function of temperature and composition for HVs made of Bd₁₁-*b*-PEG₈ & DPPC. Reprinted from ref. 64, Copyright 2024, with permission from Elsevier.



gHVs consisting of the same components (PEG-*b*-PBD & egg-SM and cholesterol) provided insights into the mechanism governing phase separation during the cooling of gHVs.⁶³ The analysis suggested either spinodal decomposition or nucleation and growth as the underlying processes. Furthermore, the analysis indicated that cholesterol was more likely to be partitioned in the block copolymer-rich domains than in the lipid-rich domains.

5.4 Viscoelastic properties

The viscoelastic properties of lipid bilayers determine their ability to change shape and deform in response to various mechanical stresses. This is crucial for the functioning of biological membranes, as cells are constantly subjected to mechanical forces. For example, during cell movement, the lipid bilayer needs to be flexible to allow cell shape changes and migration.

Micropipette aspiration is the most used technique to study the bilayer's mechanical properties. During micropipette aspiration, the membranes of the gHVs are subjected to a mechanical (stretching) force caused by the air suction. The subsequent deformation of the membrane is recorded from where the lysis strain, the lysis stress and the area expansion (stretching) modulus are calculated. For example, PEG₁₄-*b*-PBD₂₂ & HSPC with 75 mol% block copolymer content had an intermediate area expansion modulus of $K_a = 112 \text{ dyn cm}^{-1}$ between polymersomes (72 dyn cm^{-1}) and liposomes (206 dyn cm^{-1}).⁴⁴ However, in the majority of cases, gHVs were reported to have lower lysis strain than their respective liposomes and polymersomes. For example, the area expansion modulus (K_a) in phase separated PDMS-*g*-PEG & DPPC gHVs was calculated to be lower than for the pure vesicles.²⁸ Although these gHVs and the control multi-lipid liposomes (DOPC/DPPC) showed phase separation, the gHVs remained intact under the tested strain conditions, unlike the liposomes that ruptured. In another case, K_a of HVs made of PBD₄₆-*b*-PEG₃₀ & POPC increased up to 17% (from 112 ± 10 to $131 \pm 15 \text{ mN m}^{-1}$) with increasing lipid concentration (up to 30 mol%), while the critical lysis stress and strain decreased by 17% (from 15.3 to 12.7 mN m^{-1}) and 40% (from 0.15 to 0.09), respectively, compared to polymersomes.⁵⁵ The authors argued that despite a reduction in lysis stress and lysis strain, the higher stretching modulus, due to the integration of block copolymers and lipids into the hybrid membrane compensated for the toughness. This compensation was manifested in the greater applied tension needed to obtain a given strain.

In this context, the most studied block copolymer-lipid combination is PEG and PDMS in various architectures such as diblock copolymer, triblock copolymer or grafted copolymer by the Le Mains group. The stretching moduli increased for gHVs made of PEG₁₂-*b*-PDMS₄₃-*b*-PEG₁₂ & POPC with increasing lipid content (0–20 w/w%) without phase separation observed on the macroscale.³⁷ However, a decrease in lysis strain was found even for very low lipid content (5%), which was lower than for POPC liposomes. The authors suggested that, most likely due to the formation of nanodomains, lysis occurred at the borders of

these domains. The elastic modulus exhibited an intermediate value between that of polymersomes and liposomes, but with high dispersity values, ranging from 80 to 155 mN m^{-1} .¹ Interestingly, intermediate mechanical properties were observed for gHVs made of PDMS₂₇-*b*-PEG₁₇ & POPC (with 10% lipid) compared to liposomes and polymersomes. Specifically, the lysis strain was found to be $7 \pm 2\%$, compared to $3 \pm 2\%$ of POPC liposomes and $16 \pm 4\%$ for polymersomes. The lysis stress was the same as for POPC liposomes ($8 \pm 2 \text{ mN m}^{-1}$) while the stretching moduli was comparable to the polymersomes ($123 \pm 12 \text{ mN m}^{-1}$).³⁹ In a follow up report, gHVs made of POPC lipids and PDMS-*b*-PEG with various block lengths were extensively studied to have a better understanding of the change in mechanical properties upon varying the molecular weight of the block copolymer and the block copolymer – lipid composition (up to 20% lipid) (Fig. 5). The area expansion moduli of these gHVs remained almost constant independently of the lipid composition (100 and 130 mN m^{-1}). However, the lysis

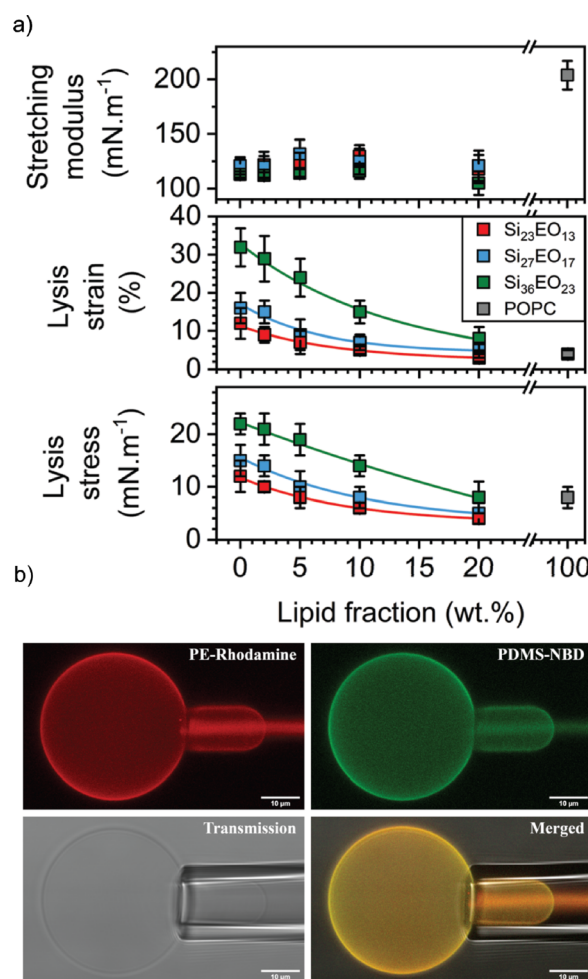


Fig. 5 Membrane properties: (a) mechanical properties determined from the stress–strain curves: stretching modulus (top panel), lysis strain (middle panel) and lysis stress (bottom panel). (b) CLSM images of micropipette experiment in the different observation modes: PE rhodamine channel, PDMS NBD channel, PE rhodamine and PDMS NBD channel merged, and transmission. Scale bars: 10 μm . Reprinted from ref. 41. Copyright 2024 the Royal Society of Chemistry.



strain and stress decreased with the decreasing lipid fraction and increased with the molecular weight of the hydrophobic block. The lysis strain remained higher than that of POPC liposomes, except for gHVs made with the lowest molecular weight block copolymer. The lysis stress was more composition dependent, and lower values than the POPC liposomes were measured in some cases. Consequently, the calculated toughness of the HVs was highly dependent on the molecular weight of the block copolymer. In particular, gHVs assembled with the highest molecular weight PDMS-*b*-PEG had the most resistant hybrid membranes, with a tenacity always higher than POPC liposomes. The authors pointed out that in their previous efforts, gHVs formed with triblock copolymer PEG₁₂-*b*-PDMS₄₃-*b*-PEG₁₂ & POPC had a membrane toughness lower than that of POPC liposomes even at a low lipid fraction (5 wt% POPC).⁴¹ The line tensions between macrodomains of gHVs made of the above mentioned polymers and POPC were calculated to further elaborate on the membrane properties. The compositions of the gHVs were chosen to allow for the assembly of stable budded vesicles with each hemisphere being formed by either the polymer phase or the lipid phase. Similar line tension values (0.32–0.82 pN) were obtained for PDMS₂₃-*b*-PEG₁₃, PDMS₂₇-*b*-PEG₁₇, or PEG₈-*b*-PDMS₂₂-*b*-PEG₈ & POPC with 50% lipid content.⁴² gHVs made of PDMS-*g*-PEG & soy PC exhibited twice lower rigidity than soy PC liposomes.³⁰ These gHVs exhibited lower bending rigidity than liposomes, comparable to that of polymersomes (11.6 ± 2.4 kBT) rather than showing intermediate softening. This result indicated that this property was largely determined by the prevailing membrane component.

In addition to micropipette aspiration, other methods were also utilized to measure the mechanical properties of membranes. For instance, Winzen *et al.* applied atomic force microscopy to characterize the viscoelastic behavior of gHVs.²⁶ They compressed gHVs made of PDMS₆₀-*b*-PMOXA₂₁ & 1,2-dimyristoyl-*sn*-glycero-3-phosphocholine (DMPC) and obtained a Young's modulus of 15 ± 4 MPa that was similar to the corresponding polymersome stiffness. Progressive deformation on PBD-*b*-PEG & DPPC-based gHVs was studied with a microfluidic setup, where the gHVs were passed through a narrow, short channel by controlling the pressure of the flow.⁶⁸ In this case, gHVs with 35% lipid content showed similar longitudinal deformation as the polymersomes despite domain formation. However, gHVs with 65% lipid content might feature both liposome-like transverse flattening and polymersome-like deformation, while slipping through the channel. The variations in stretching techniques employed in pipette aspiration, atomic force microscopy, and microfluidic conditions make it challenging to directly correlate their outcomes. Nevertheless, insights into alterations in membrane properties can be obtained through comparison of HVs to liposomes and/or polymersomes.

5.5 Membrane thickness

The membrane thickness has an effect on the mechanical properties of the membranes as well as on the permeability of the vesicles. Liposomes have a membrane thickness of around

3–6 nm, depending to largely on the length of the lipid tails and environmental factors, such as temperature or ionic strength. The membrane thicknesses can be obtained experimentally by various techniques including SANS, SAXS and cryo-EM. The latter has the advantage that individual vesicle can be observed (to *e.g.*, identify phase separation) and measured and thus, gives information about variations within the population.³⁶

The addition of copolymers has a substantial influence on the membrane thickness due to the higher molecular weight of the copolymer chains compared to the lipids, which therefore occupy a larger volume. For instance, molecular simulations showed that with increasing block copolymer content in PEG-*b*-PBd & DOPC-based HVs, the membrane thickness increased.⁶⁶ Although there is theoretically a variation in membrane thickness based on the lipid-to-polymer ratio, in many instances, this difference may not be significant. For example, the average membrane thickness of HVs made of PDMS₆₀-*b*-PMOXA₂₁ & DMPC did not statistically differ from the thickness of polymersome membranes (17 ± 2 nm) independent on the amount of added lipids (10 or 50%) and were homogenous in all cases.²⁶

Several studies aimed at elucidating the relationship between molecular weight and composition distribution in HVs by analyzing the membrane thickness. For instance, sHVs comprising of PCMA-*b*-pDMAEMA & POPC, POPE or POPS lipids had membrane thicknesses between 5.8 and 6.7 nm, which was thicker than the membrane of a POPC lipid bilayer (5.4 ± 0.5 nm).⁸⁶ Moreover, sHVs made with low molecular weight PCMA-*b*-pDMAEMA had, on average, thinner membranes compared to sHVs assembled with high molecular weight PCMA-*b*-pDMAEMA. Furthermore, the presence of a bilayer structure was observable in all samples, indicating a homogeneous insertion of the block copolymers into the hybrid membrane. Cryo-EM analysis of sHVs assembled from PDMS-*g*-PEG & soy PC indicated that the membrane thickness slightly increased with increasing amount of block copolymer (4.40 ± 0.16 nm for liposomes, 4.86 ± 0.17 nm for sHVs and 5.25 ± 0.17 nm for polymersomes).³⁰ The liposomes had a bilayer structure with sharp outer boundaries, while the block copolymer containing membrane appeared as fuzzy monolayers. Similar assemblies made of PDMS-*g*-PEG & soy PC had an increased membrane thickness when measured in potassium chloride-supplemented buffer compared to sucrose (6.1 ± 0.3 nm compared to 5.3 ± 0.2 nm) due to the ion association in the hydrophilic PEG layer.³¹ Similarly to the potassium chloride containing solution, the presence of the polymer in sHVs made of either PDMS₂₆-*g*-(PEG₁₂)₂ & anionic soy PS or PDMS₂₆-*g*-(PEG₁₂)₂ & cationic DOTAP allowed for strong association of the ions with the vesicle surfaces in tris(hydroxymethyl)amino-methane buffer, *i.e.*, their membrane thickness increased from 5.8 ± 0.4 nm to 6.5 ± 0.4 nm for the anionic sHVs and from 5.8 ± 0.3 nm to 6.4 ± 0.3 nm for the cationic sHVs.³²

Vesicles with inhomogeneous bilayer thicknesses can indicate nanodomain formation. For example, sHVs made of maltopentaose-*b*-PPO & DOPC had a variation within a single sHV between *ca.* 2.5 nm and *ca.* 6.5–7 nm, which corresponded to the average membrane thicknesses in pure vesicles and polymersomes, pointing to lipid and polymer-rich domains.⁹⁶



Cryo-EM is often used as a complementary technique to SANS and SAXS with the aim to visually support the scattering data. Neutron scattering curves can give information about the shape and size of the scattering objects in solution by fitting various types of models to the scattering data. In SANS, scattering of neutron beams depends on the scattering length density (SLD) of the scattering nucleus. There is a difference in SLD between hydrogen and deuterium, thus, by mixing H₂O and D₂O in different proportions, the SLD of the solvent can be systematically varied and matched to the SLD of one of the components of the hybrid vesicle. This technique known as contrast matching, and it is possible to render a phospholipid invisible.¹⁰⁷ For instance, sHVs assembled from PDMS-*g*-PEG₂ & DPPC were measured with or without deuterated DPPC by SANS.³⁴ The obtained shape factor was related only to the copolymer, *i.e.*, the polymer contrast conditions, in the presence of deuterated lipids. In this condition, the obtained curves showed a disk-like shape, indicating polymer domains in the sHVs instead of homogeneous distribution when using the Kratky-Porod fit. In contrast, the assembly had a vesicular shape with an average bilayer thickness of 5.8 nm at 20 °C when non deuterated lipids were used. Cryo-EM images of these sHVs showed either rounded spherical or faceted sHVs. The latter was ascribed to the gel state of the DPPC lipids. In another study, SANS curves obtained from sHV composed of PEG_{*n*}-*b*-PDMS_{*m*} & POPC with increasing overall molecular weight of the block copolymer in polymer contrast and lipid contrast conditions were fitted with the same Kratky-Porod model.⁴⁰ The membrane thickness decreased with increasing lipid fraction when observing the polymer phase, suggesting interactions between the lipid chains and the block copolymer chains in the hybrid membrane. The higher molecular weight of the block copolymer did not influence the membrane thickness significantly (6.4–6.8 nm). In lipid contrast mode, sHVs with shorter block copolymer chains exhibited comparable membrane thicknesses to liposomes, while the sHVs with the longest block copolymer chain displayed a more pronounced deviation. The authors argued that this observation might be explainable by fission. However, in this case, it would be expected that the thickness of the second population should also remain unchanged compared to the polymersome, which was not observed.⁴⁰ Perera *et al.* assembled sHVs made of PEG_{*n*}-*b*-PDMS₁₅-*b*-PEG_{*n*} &

DOPC where PDMS length was kept constant and the PEG was varied systematically.⁴³ They used a polydisperse core-multishell model where the SLDs were taken into account, allowing to estimate the thickness of the water-rich hydrophilic shell and the denser PDMS and lipid-rich core. The SANS data revealed that the hydrophilic mass fraction of an amphiphilic polymer affected the hybrid membrane thickness and the sHV size distribution (Fig. 6a), which was in good agreement with the obtained cryo-EM images.

The scattering profile in SAXS measurements is related to the electron density in the bilayer. Modeled SAXS data can give information about the shape and the size of the vesicle as well as the packing of the components in the bilayer. This technique is often used for studying membrane proteins and their configuration in the bilayer.¹⁰⁸ SAXS was used to assess the effect of the chemical nature of the hydrophobic block on sHVs made of either BuMA-*b*-PCEA & soy PC or (PCMA_{*n*}-*co*-BuMA_{*m*})-*b*-PCEA & soy PC with various molecular weights and polymer to lipid ratios.⁹³ The SAXS data were fitted with a model, which took the presence of micelles in addition to the sHVs in the solution into account. The results showed that the sHVs had a higher membrane thickness between 40.4 ± 0.6 to 44.8 ± 1.2 Å compared to the soy PC liposomes (39 ± 0.2 Å), which slightly increased with increasing block copolymer content. However, the type of block copolymer did not influence the membrane thickness of the sHVs. In a related case, sHVs made of PCMA-*b*-PMPC & DOPC or DOPS had comparable membrane thicknesses to liposomes (~4.6 nm) as determined by SAXS measurements.⁹⁴ However, the scale of the Gaussians describing the head group regions were in general slightly larger for the sHVs than for the liposomes, suggesting the presence of the block copolymer in the membranes. In a recent publication by Senviratne *et al.*, a detailed high resolution cryo-EM and SAXS investigation of PBD₂₂-*b*-PEG₁₄ & POPC-based sHVs with increasing polymer content was presented.⁵⁰ Overall, they found that the membrane thickness increased (between 5–10 nm) with increasing block copolymer fraction. However, two population of vesicles with different membrane thickness were observed at higher polymer content (>50%). The membranes did not show domain formation and atomic resolution cryo-EM images revealed the detailed membrane structure, showing similarly organized bilayer structures with different thickness (Fig. 6b).

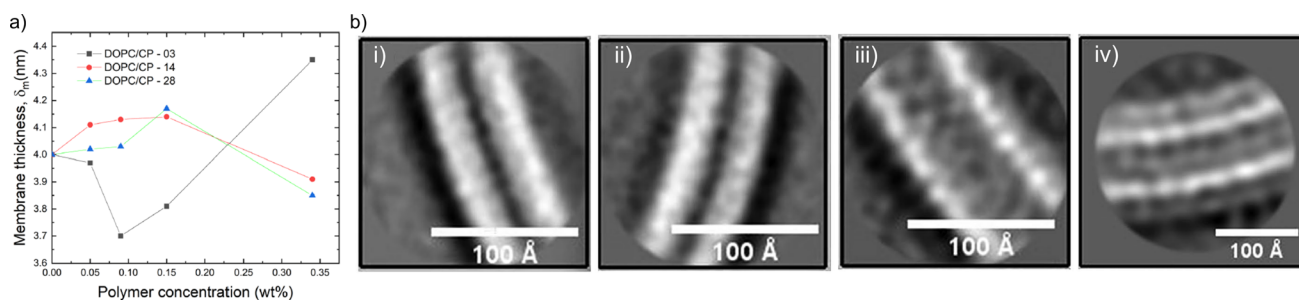


Fig. 6 (a) Change of the total membrane thickness of PEG_{*n*}-*b*-PDMS₁₅-PEG_{*n*} (*n* = 3, 14, 28) & DOPC-based HVs with the increasing polymer concentration. Reprinted with permission from ref. 43. Copyright 2024 American Chemical Society. (b) Cryo-EM images of the membrane of vesicles made of (i) PBD₂₂-*b*-PEG₁₄ (50 mol%) & POPC, (ii) population 1, (iii) population 2 (thin and thick, respectively), and (iv) PBD₂₂-*b*-PEG₁₄ only (polymersomes).⁵⁰ Used under CC BY 4.0. <https://onlinelibrary.wiley.com/doi/full/10.1002/sml.202206267>.

In general, all three techniques give information about the membrane thickness. CryoEM images contain detailed structural information of individual vesicles, while, SAXS and SANS give average structural information in solution. Consequently, using these techniques in a complementary manner provides the most comprehensive understanding.

6 Stability

The limited stability is probably the major shortcoming of liposomes. This issue also includes their rather short shelf life, *i.e.*, the liposomal colloidal stability is affected by membrane fusion within 48 h, at inappropriate storage conditions.⁵ In contrast, HVs often show exceptional stability without detectable changes in D_h and PDI over time. Several examples are reported with stability up to four weeks at 4 °C,^{26,46,78} and in a recent study even over 500 days.⁴⁰ An aspect that affects the HV stability is the length of the polymer chains, *e.g.*, HVs assembled from PEG₁₂-*b*-PCL₁₀ & POPC, PEG₄₅-*b*-PCL₂₅ & POPC or PEG₄₅-*b*-PCL₄₂ & POPC had lower stability for the shorter hydrophobic blocks (4 days vs. 3 weeks) (Fig. 7a).⁷⁸ Conversely, HVs consisting of PEG₁₁₃-P(8C-Chol) & DPPC exhibited higher stability for shorter hydrophobic blocks (up to 245 days), while high molecular hydrophobic segments were insufficiently incorporated leading to precipitation after 2 weeks.⁸⁵ In another example, HVs made of PMOXA₁₅-*b*-PDMS₆₇-*b*-PMOXA₁₅ & DOPC were stable for up to 3 weeks when 10 mol% triblock copolymer was used, while 20 mol% triblock copolymer reduced the stability of the HVs to 5 days.²⁶ HVs assembled from PBD-*b*-PEG & POPC were subjected to different types of sterilization methods that can be required for biological applications such as drug delivery.⁵¹ Not surprisingly, only HVs that were below 400 nm remained intact after filtration through a 0.44 μm filter. Furthermore, freeze-thaw-vortex cycles also showed substantial destruction of the HVs, assessed by the substantial release of their cargo, pointing out that other ways of sterilization methods need to be tested.

Exposure to surfactants such as Triton-X or bile as well as lipases is often used to determine the inherent stability of HVs. Lipases are a group of enzymes that catalyse the hydrolysis of lipids that play a crucial role in the digestion of lipids in the digestive tract. As expected, the stability of HVs towards lipases depends on the building blocks. For instance, sHV s made of PEG₁₂-*b*-PCL₆ & POPC, PEG₁₂-*b*-PCL₉ & POPC or PEG₁₆-*b*-PLA₃₈ & POPC were structurally affected by phospholipase A₂ (PLA₂) as monitored by calcein release and significant changes in their D_h and PDI.⁷⁶ In another study, the hydrophobic block consisted of a copolymers of PCL and PLA with PEG as a hydrophilic segment. The structural integrity of the sHVs containing a rather large polymer PEG₁₁₄-PLA₃₂₈ was best preserved, delaying calcein release up to 5 min after PLA₂ addition.⁷⁹ In contrast, gHVs made of PBD₄₆-*b*-PEG₃₀ & DPPC & cholesterol (50 mol% block copolymer) remained topographically intact and spherical upon PLA₂ addition.¹⁰⁴ The gHVs lost their cholesterol-rich lipid domains when subjected to PLA₂, but they showed self-healing, *i.e.*, recovering the full circular structure after the lipid loss as visualized by CLSM. These gHVs remained

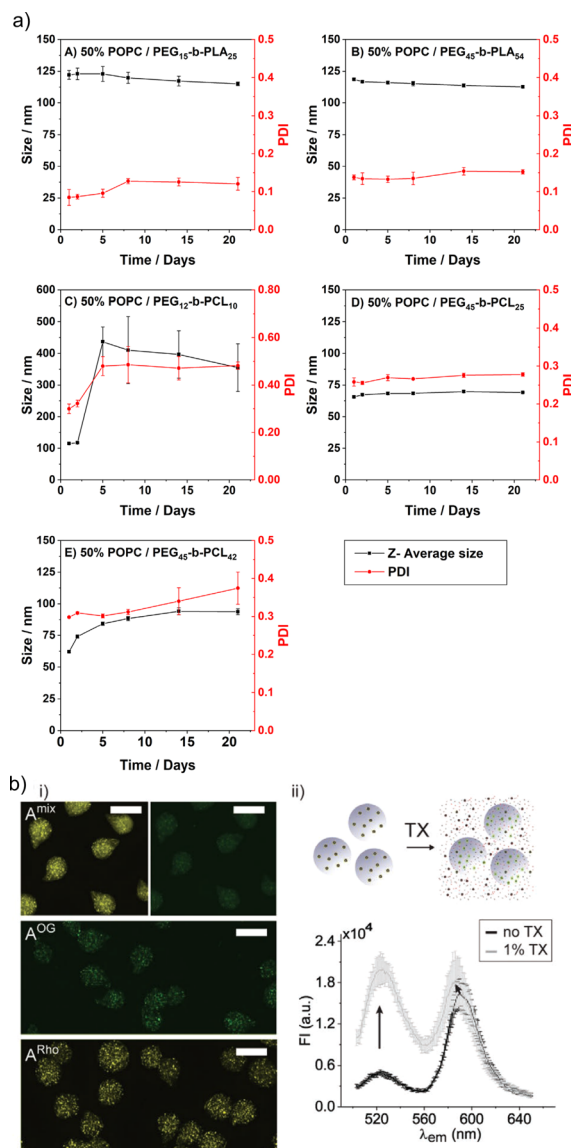


Fig. 7 Stability of HVs. (a) Hybrid vesicle size stability at 4 °C. Samples are monitored by DLS over a period of 3 weeks: (A) PEG₁₅-*b*-PLA₂₅; (B) PEG₄₅-*b*-PLA₅₄; (C) PEG₁₂-*b*-PCL₁₀; (D) PEG₄₅-*b*-PCL₂₅ & POPC; and (E) PEG₄₅-*b*-PCL₄₂ & POPC. Reprinted from ref. 78, Copyright 2024, with permission from Elsevier. (b) (i) Representative fluorescent microscopy images of HVs encapsulated in alginate beads (A) with FRET pair labelled polymer and lipid (A^{mix}) or only one component (A^{OG}, and A^{Rho}). Scale bars: 100 μm. (ii) Illustration of sHVs encapsulated in alginate beads and their disassembly upon addition of Triton-X (top). Fluorescent emission spectra of A^{mix} before (black line) and after (gray line) exposure to 1% Triton-X (bottom). Reprinted from ref. 89, Copyright 2024, with permission from Wiley.

structurally intact despite hole formation that allowed for the release of encapsulated cargo as big as proteinase K (28 kDa). In another study, gHVs made of PBD₁₉-*b*-PEG₂₇ & DPPC, exhibited a burst release of the cargo when exposed to lipases, although the overall structural integrity of the gHVs was preserved.⁶¹ In contrast, gHVs assembled from PBD₁₉-*b*-PEG₂₇ & POPC showed first a continuous release of the encapsulated cargo upon addition of PLA₂ followed by buckling and release of the



remaining cargo. The fact that the lipase could access the lipid molecules showed that the incorporated block copolymer did not provide a protecting stealth layer around the HVs, as it could be expected from PEG. This example illustrated that the HVs, even when containing high amounts of block copolymer, had large lipid domains, even for the non-saturated, more fluid membranes. Destabilization of HVs by surfactants depends on their concentration. Surfactants in low concentrations are often employed for the insertion of membrane proteins (see below). Typically, the HVs undergo complete structural disintegration when the surfactant concentrations reach a certain threshold. A common approach to exploit this fact is to use Triton-X at concentrations ranging from 0.03–1% to release the cargo and evaluate the encapsulation efficiency.^{1,46,51,61,75,76,78,79,93}

The stability of the HVs is an important aspect for their use in drug formulation, *e.g.*, for encapsulation of HVs into hydrogels as potential drug deposits. To this end, sHVs made of PCMA₁₁-*b*-P(METMA₇-*co*-PCEA₁₀₄) & DOPC,⁸⁹ PCMA₁₁-*b*-PEG₁₁₃ & POPC/POPS,⁸⁸ or PCMA-PMPC & DOPS or DOPC⁹⁴ were encapsulated into alginate microbeads. The preservation of the FRET effect due the fluorescently labeled block copolymer and the fluorescently labeled lipids suggested that the building blocks remained closely associated even after the sHVs were trapped in the hydrogel. Yet, the lipids leak out after addition of Triton-X or incubation in simulated gastric fluid, indicating the loss of structural integrity of the sHVs (Fig. 7b).

7 Permeability

The cell membrane's permeability is controlled in various ways and plays a significant role in keeping cellular homeostasis. Being able to influence permeability is a crucial characteristic for achieving cell membrane-like adaptability. Liposomes and especially polymersomes have very low inherent permeability,¹⁰⁹ and recent progresses in this context is discussed in several comprehensive reviews.^{110–112} Often, HVs show improved permeability towards small molecules compared to their polymersome or liposome counterpart. The inherent permeability is usually determined/compared by monitoring the release of encapsulated (small hydrophilic) fluorescent molecules such as calcein or (carboxy)fluorescein. However, the main drawback is the low solubility of these dyes in aqueous media, and to achieve self-quenching concentrations (>20 mM), high pH is required, which can interfere with the assembly process of the vesicles. Further, the measured encapsulation efficiency in vesicles made by the rehydration method is inherently very low, generally below 10% as it is a passive encapsulation process. Nonetheless, this release assay remains the most viable option to address permeability of small vesicles.

Most of the permeability studies are performed on sHVs that are extruded through a 100 nm membrane to obtain unilamellar vesicles. In general, the release of the dyes occurred within 5 days. Multilamellar vesicles likely do not release the encapsulated molecules completely.⁹³ Studies have explored various methods to influence the permeability of HVs, including the lipid to copolymer ratio, molecular weight of the copolymer, and the use of charged lipids or polymer chains. For

example, research has shown that increasing the PEG₁₄-*b*-PB₂₂ to POPC ratio in HVs could result in delayed release of encapsulated carboxyfluorescein at room temperature.⁴⁶ The effect of the block copolymer molecular weight was explored by systematically changing the length of the blocks in PEG_{*n*}-*b*-PLA_{*m*}.⁷⁸ The carboxyfluorescein release rates were influenced by the overall properties of the block copolymer, but there was no clear trend on the dependence of the release profile on the length of the various blocks. However, the release of carboxyfluorescein was slower at 4 °C than at 37 °C. In another example, the molecular weight of the block copolymer was found to affect the permeability of HVs made of PDMS-*b*-PEG & POPC, but not the lipid to block copolymer ratio.⁴⁰ Specifically, assemblies made with the shortest polymer (PDMS₂₃-*b*-PEG₁₃) showed higher permeability, releasing fluorescein more efficiently than less permeable liposomes. HVs made with POPC and the longer block copolymers exhibited slightly lower permeability, but not significantly different from the corresponding polymersomes.

The chemistry of the hydrophobic block, which inserts into the bilayer, can be altered, by synthesizing block copolymers, which, in turn, can modify the permeability of the HVs. The calcein release from sHVs made of POE₁₁₄-*b*-P(CL_{*x*}-*co*-LA_{*y*}) & DPPC, with different compositions of PCL and PLA, increased with increasing PLA content.⁷⁹ When comparing block copolymers of PCMA with BuMA or HEMA as the hydrophobic block, it was observed that increasing the block copolymer content from 1–5 w/w% resulted in a rise in carboxyfluorescein release from the gHVs. gHVs made of soy PC and block copolymers with a PCMA homopolymer hydrophobic block or a copolymer of CMA and BuMA hydrophobic block did not cause any change to the gHVs' permeability. However, when using HEMA instead of BuMA in the copolymer, a significant increase in permeability of RhoX was observed.⁹³ sHVs made of the aforementioned building blocks exhibited a faster release of carboxyfluorescein within 24 h as compared to soy PC liposomes, consistent with the trend observed in gHVs. In another study, when PCMA-*b*-PMPC was used in HVs with DOPC, DOPS or DOPS/DOPE, the encapsulated carboxyfluorescein was released at a similar rate within 24 h, indicating that the lipid type did not influence the permeability.⁹⁴ The molecular weight of the hydrophilic block can also influence the permeability of HVs. Further, HVs made of maltopentaose-*b*-PPO₄₃ & DOPC had lower permeability towards PEG-FITC (550 Da) than HVs made of maltopentaose-*b*-PPO₂₆ & DOPC.⁹⁶

pH-sensitive chemical groups give the possibility to modulate the flow of charged molecules across the membrane. For instance, negatively charged calcein was released within minutes from gHVs made of cationic lipids (CLS) & PDMS₃₆-*b*-PEG₂₃ upon acidification of the environment, *i.e.*, protonation of the lipid heads, while calcein was retained in gHVs containing only zwitterionic POPC and non-chargeable lipids.³⁸ Similarly, gHVs made of oligo(aspartic acid)₄-*b*-PPO & DOPC were permeable to positively charged rhodamine 6G and non-charged PEG-FITC (550 Da) after immersing the assemblies into the fluorophore solution for 4 h. In contrast, negatively charged fluorescein was only able to pass through the hybrid membrane at pH 4 where the polymer side chains were



uncharged.⁹⁵ In addition, these sHVs showed higher permeability towards acetylthiocholine than DOPC liposomes probably due to the attraction forces between the negatively charged polymer chains and the positively charged molecule. The pH responsive nature of PDMAEMA allowed to fully release the encapsulated drug below its pKa (pH 6.5) from PDMAEMA-*b*-poly(lauryl methacrylate) & egg *L*- α -phosphatidylcholine (EPC)-based HVs, while at pH = 7.4 60% of the drug was retained.⁸⁴

Inonophors passively transport specific type of ions across the cell membranes, and they are explored in synthetic cell membranes to establish more acidic or basic conditions in the vesicle's void. The ion channels valinomycin, nigericin and gramicidin A were incorporated into HVs made of PBD₃₇-*b*-PEG₂₂ & POPC to increase the permeability towards potassium ions (Fig. 8).⁵⁸ The H⁺ and OH⁻ permeability can be assessed by encapsulating environmental sensitive dyes such pyranine or trisodium 8-hydroxypyrene-1,3,6-trisulfonate to monitor the pH equalization rates inside of the vesicles. Kleineberg *et al.* showed that the rate of ion permeability was higher for both their HV system compared to liposomes. However HVs made of PBD₂₂-*b*-PEG₁₄ & soy PC showed lower permeability compared to PDMS₁₂-*g*-(PEG₃₂)₂ & soy PC, assessed by valinomycin incorporation. The difference in permeability was probably due to the polymer architecture (linear *vs.* grafted) that likely inserted differently into the hybrid bilayer, forming a denser bilayer when PBD₂₂-*b*-PEG₁₄ was used.⁵⁴

8 Applications in bottom-up synthetic biology

The prospect of constructing artificial vesicular systems through the bottom-up assembly of biological and chemical components presents exciting opportunities for addressing previously unmet needs. By strategically combining modular scaffolds and functional building blocks, it becomes possible to

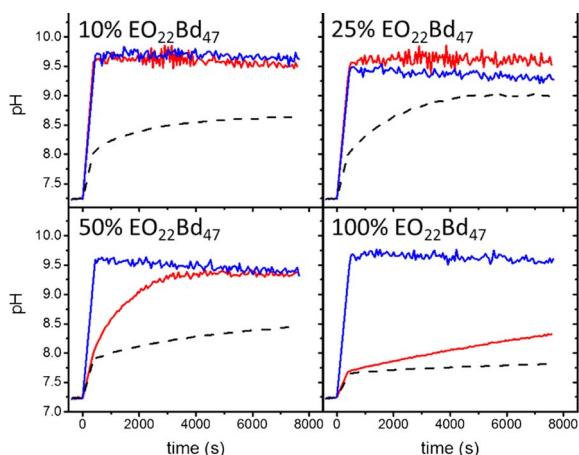


Fig. 8 Permeability of HVs detected *via* monitoring the internal vesicle pH. HVs made of PBD₃₇-*b*-PEG₂₂ & POPC using 10%, 25%, and 50% and 100% block copolymer without any added ionophore (dashes) and after incubating with valinomycin (red tracing) and nigericin (blue tracing). Reprinted from ref. 58, Copyright 2024, with permission from Elsevier.

engineer intricate systems featuring biomimetic or novel functionalities not found in nature.^{113–115}

The majority of the publications discussing HVs focus on physical-chemical properties of the assemblies but increasing numbers of reports push the concept towards an application. HVs were utilized as drug carriers,^{15,84} or mucopenetrating subunits,⁸⁸ but the predominant envisioned purpose of HVs are their use as a scaffold for protein reconstruction, as nanoreactors/artificial organelles or artificial cells. Membrane-bound vesicles in the cellular environment contribute to the complexity, efficiency and specializations of cellular functions.

Membrane fusion is essential as it allows for the merging of membrane-bound organelles and vesicles, facilitating the exchange of materials and enabling cellular processes such as secretion, nutrient uptake, and communication. Controlled membrane fusion was achieved by charge-mediated fusion between positively charged HVs made of PDMS-*g*-(PEG)₂ & DOTAP and negatively charged HVs made of PDMS-*g*-(PEG)₂ & Soy PS for both, sHVs and gHVs. The authors found that the mixing of the semisynthetic membranes were depended on, but did not scale with, the charged lipid content. In gHVs, the fused vesicles even appeared as domains in the newly formed gHV in some cases, which may be a mean to control raft formation in synthetic membranes.³²

8.1 Protein reconstruction

The plasma membrane contains membrane proteins fulfil a variety of crucial functions essential for the cell's survival and proper functioning. These membrane proteins are integral components of the lipid bilayer and contribute to the dynamic nature of the plasma membrane. The improved properties of HVs are anticipated to outweigh their lower resemblance to pristine phospholipid bilayers, *e.g.*, the better hybrid membrane stability and elasticity¹¹⁶ are expected to maintain the functional integrity of the membrane proteins for longer compared to liposomes. The proteins are reconstituted into the hybrid membrane using established detergent-mediated reconstitution protocols used for proteoliposome assembly such as Triton-X,^{29,47,48,53,54} octyl- β -D-glucopyranoside,^{29,56,67} or dodecyl-maltoside.⁴⁸ Various types of enzymes have been reconstructed into HVs, verifying the hypothesis that these vesicles can provide a suitable environment for the proteins as well as that the presence of the polymer did not impact the enzymatic function. Examples include the reconstitution of OmpF into HVs made of polyisopren(PI)-*b*-PEG & 1,2-diphytanoyl-*sn*-glycero-3-phosphocholine (DPHPC),⁷² Cyt b03 into HVs assembled with PBD₂₂-*b*-PEG₁₄ & POPC,^{47,53} or the efflux Pumps NaAtm1 and P-glycoprotein were successfully reconstructed into HVs made of PBD₂₂-*b*-PEG₁₄ & POPC and PBD₂₂-*b*-PEG₁₄ & *E. coli* lipid extract (EcCL), respectively.⁴⁸ The enzyme pair bacteriorhodopsin and F₁F₀ATP synthase were successfully incorporated into HVs made of either PBD₂₂-*b*-PEG₁₄ & POPC or PDMS₁₂-*g*-(PEG₃₂)₂ & Soy PC.^{29,30,54} In the latter case, Marusic *et al.* pointed out that the reason for the successful protein insertion and retained activity of the reconstructed enzyme was due to the suitable choice of copolymer, which resulted in a soft



and fluid hybrid membrane and allowed for unhindered conformational changes and lateral diffusion. In a follow up study, the enzyme pair was inserted into HVs made of PDMS₁₂-*g*-(PEG₃₂)₂ & Soy PS and PDMS₁₂-*g*-(PEG₃₂)₂ & DOTAP. The enzyme activity was reduced in the presence of negatively charged lipids showing its sensitivity towards the environment.³² Recently, Catania *et al.* showed that the Cyt bo3 enzyme could be inserted into HVs made of PBD₂₂-*b*-PEG₁₄ & EcCL HVs without the use of typical surfactants but with the aid of styrene-maleic acid-based copolymer with a 74% efficiency.⁴⁹ The role of surfactants in membrane protein reconstitution is to

destabilize the bilayer by swelling up the bilayer allowing an easy insertion of the protein. This procedure is tedious and requires the removal of the surfactant. Therefore, this finding represents a significant advancement. It not only enhances our comprehension of the bilayer structure of HVs but also provides a more accessible platform for studying membrane-bound enzymes in a more convenient manner.

A recent example illustrated the incorporation of fusiogenic SNARE into sHVs made of PDMS₁₂-*g*-(PEG₃₂)₂ & soy PC with remarkable amount of outward facing proteins and very high incorporation efficiency.³¹ The SNARE-containing sHVs were

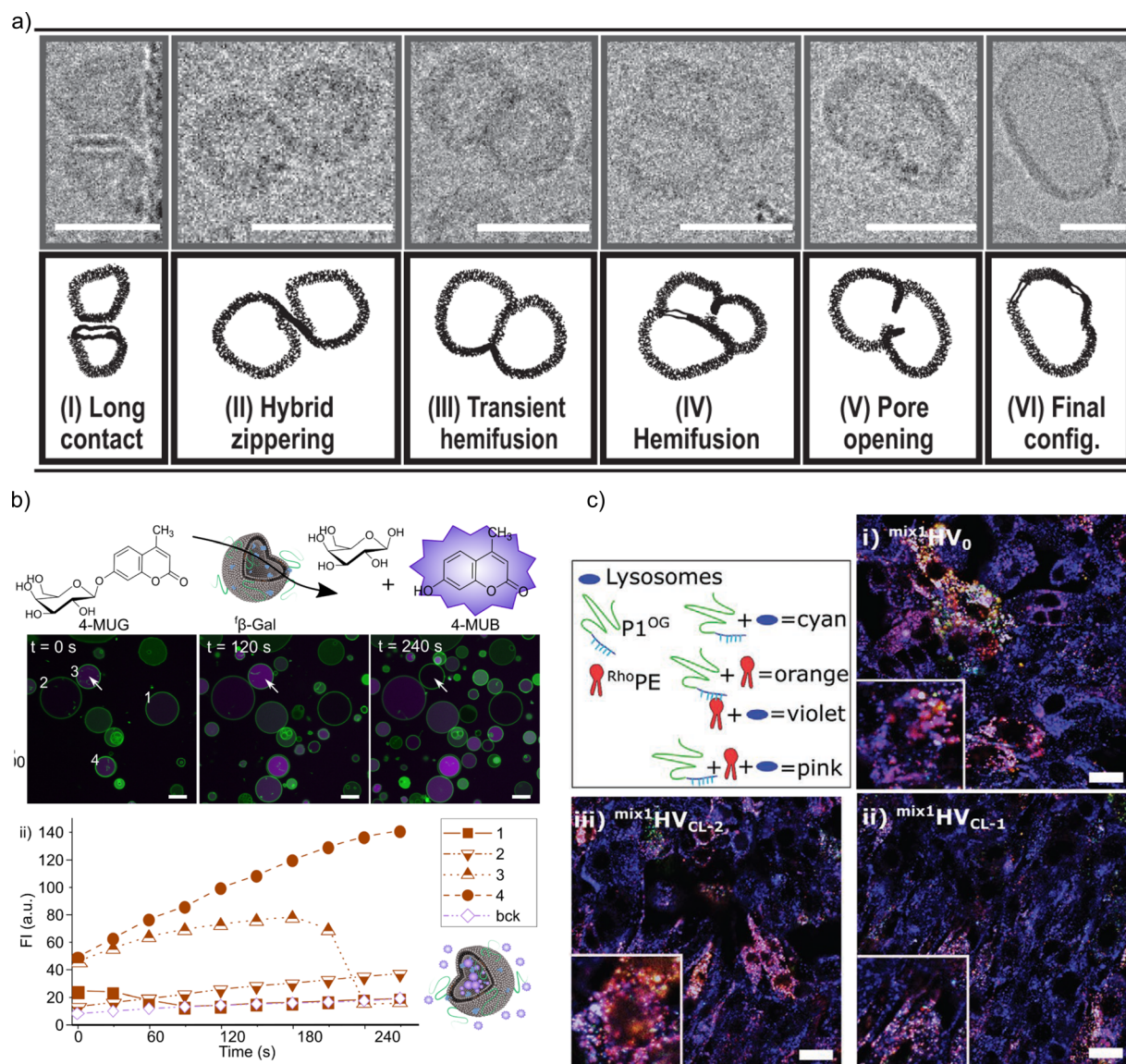


Fig. 9 Applications of HVs as synthetic protocell and as nanoreactors. (a) Proposed fusion intermediates of SNARE-mediated fusion of HVs made of PDMS-*b*-PEG & soy PC. Scale bars: 30 nm.³¹ Used under CC BY 4.0. <https://www.nature.com/articles/s41467-021-25294-z>. (b) (i) Representative time-lapse CLSM images of β -Gal associated with HVs made of P(CMA-co-BuMa)-*b*-PCEA & soy PC immersed into 100 μ M 4-MUG solution. The arrow indicates the sudden change in intensity (green: Oregon green labelled polymer, magenta: 4-MUG). (ii) The increase of the fluorescent intensities of 4-MUG measured on CLSM image over time for selected 4 vesicles in comparison to the background (bck). Scale bars: 10 μ m. Used under CC BY 4.0. <https://www.sciencedirect.com/science/article/pii/S2352940722001846>. (c) Representative CLSM images of cells incubated with HVs made of PCMA-*b*-PCEA & DOPE, PCMA-*b*-PCEA & DOPE with low cross linking, and PCMA-*b*-PCEA & DOPC with high crosslinking for 24 h (blue: LysoTracker deep red stained lysosomes; green: block copolymer; red: RhoPE). Scale bars: 20 μ m.⁹² Used under CC BY 4.0. <https://onlinelibrary.wiley.com/doi/10.1002/adbi.202200209z>.



able to fuse together shown by mixing of their fluorescent content. The bending rigidity and pore edge tension was determined to be the key parameters for fusion. The authors proposed a plausible progression of fusion through cryo-EM snapshots. In addition, such synthetic materials were proposed as an amazing new tool for understanding dynamic membrane phenomenon and for the assembly of synthetic protocells (Fig. 9a).

8.2 Nanoreactors and artificial organelles

Nanoreactors are typically single compartment nanosized carriers that can perform encapsulated biocatalytic reactions due to entrapped (artificial) enzymes as recently reviewed in detail.⁶ These carriers can be based on (porous) silica nanoparticles, polymer capsules, liposomes, polymersomes or HVs among others. An essential property of the vesicular nanoreactors is their selective permeability, *i.e.*, the substrate and product molecules can easily cross the membrane while the encapsulated enzymes are retrained. HVs offer unique opportunities in this context due to their hybrid nature. For instance, the polymer-rich domains in HVs made of oligo(aspartic acid)₄-*b*-PPO_{4.3} & DOPC (10 mol% block copolymer) were shown to be able to serve as synthetic channels, allowing for the transfer of the small substrate molecule acetylthiocholine chloride to be hydrolyzed by the encapsulate enzyme acetylcholine esterase (AChE)s.⁹⁵ The product was then released from the HVs to form a colored compound in a subsequent reaction for detection. Otrin *et al.* utilized HVs made of PDSM-*g*-PEG & soy PC to incorporate the chemical energy driven proton pump through a consecutive reaction of Cyt bo3 oxidase, which was used to drive ATP synthesis through F₁F₀ATP synthase.²⁹ The composition of the hybrid membrane had no effect on the activity of Cyt bo3 oxidase alone. However, the ATP production rates dropped by 50% when HVs were used instead of liposomes. We utilized β -galactosidase to construct simple nanoreactors to evaluate the effect of the chemical changes in the used block copolymer. β -Galactosidase was found to be in the HVs membrane, but nonetheless, the fluorescent methylumbelliferone (4-MUB) production from β -D-galactopyranoside was observed in HVs made of P(CMA₉-*co*-BuMA₉)-*b*-PCEA₁₈₆ & Soy PC as well as in HVs assembled from P(CMA₈-*co*-HEMA₃)-*b*-PCEA₁₈₆. The enzymatic reaction was followed by CLSM imaging using gHVs (Fig. 9b). The produced fluorescent dye built up in the interior of the gHV until it was instantaneously released without any obvious damage of the structural integrity of the gHVs. The various hydrophobic blocks used had a significant influence on the lifetime of the associated enzyme. While using P(CMA₉-*co*-BuMA₉)-*b*-PCEA₁₈₆ the enzymatic reaction was proceeding for up to 7 minutes. However in HVs made of P(CMA₈-*co*-HEMA₃)-*b*-PCEA₁₈₆ & Soy PC the fluorescent content was released after 1 minute.⁹³

Nanoreactors that exhibit catalytic activity inside of mammalian cells are often referred to as artificial organelles. The design of a carrier system that can act as artificial organelles has many challenges including the requirement that they should remain intact upon cellular uptake and placement in the

cytosol. We have observed that HVs made of PCMA₁₁-*b*-P(METMA₇-*co*-PCEA₁₀₄) & DOPC lose their integrity when taken up by HepG2 cells. The block copolymer remained in the cells and the lipids were eliminated after 16 h.⁸⁹ The fusiogenic GALA peptide was covalently attached to the block copolymer for the assembly of PCMA-*b*-PCEA_{GALA} & DOPC-based HVs in addition to HVs made of PCMA-*b*-PCEA & DOPE to enhance their endosomal escape ability. Although both types of HVs could escape the lysosomes, the intracellular retention of the building blocks in RAW 264.7 mouse macrophages was only a few hours.⁹¹ In an attempt to address this issue, the functional groups on the block copolymer and the lipid head were cross-linked in HVs made of PCMA-*b*-PCEA & DOPE to improve their structural integrity. These HVs could withstand low surfactant concentrations. C8-D1A astrocytes showed lower uptake when incubated with the cross-linked HVs compared to the non-cross-linked HVs (Fig. 9c).⁹² In another effort, we demonstrated that HVs consisting of PCMA-*b*-PDMAEMA & POPC were able to produce H₂O₂ and nitric oxide when loaded with glucose oxidase and β -galactosidase, respectively, in RAW264.7 mouse macrophages.⁸⁷ Furthermore, human macrophages differentiated from donor human peripheral blood mononuclear cells had higher intracellular nitric oxide levels when treated with β -galactosidase-containing nanoreactors in the presence of the substrate β -Nonoate compared to the controls.

9 Conclusions & outlook

Since the early days of assembling cell membrane-like systems, phospholipids are a popular choice because of their chemical nature and their ability to self-assemble into vesicles with tuneable sizes. Even though lipid vesicle-based artificial cells are essential in bottom-up synthetic biology, using only lipids is not sufficient to create artificial membranes, as they do not reflect the structural and functional complexity of cell membranes. Consequently, the addition of polymers to these lipid-based membranes is a step towards a more advanced design. The properties of polymers can be easily tuned and adjusted as the monomer of choice largely influences the physical-chemical properties of the resulting polymer chain. HVs therefore combine the best of the two worlds, the self-assembly ability of lipids and the chemical diversity of polymers. The fundamental properties including the homogenous distribution of the building blocks *vs.* their phase separation, the resulting vesicle sizes and membrane properties. In the latter case, different aspects interplay to give the overall hybrid membrane properties including the membrane thickness, the fluidity and viscoelastic properties. The improved stability and better control over the permeability are two core parameters that distinguish HVs from liposomes. Although different fields benefit from HVs, they have proven to be particularly useful in bottom-up synthetic biology to improve protein reconstitution and to offer alternative scaffolds for nanoreactors and eventually artificial organelles and cells.

The next step forward will require larger efforts to implement 'smart' polymers to obtain HVs with responsive properties. In addition, moving away from mostly synthetic polymers (and



monomers) towards natural building blocks (e.g., peptides, sugars, oligonucleotides) is an underexplored opportunity. The biological evaluation of HVs is in the early days, and a better understanding of how cells, tissues and eventually living organisms respond to them is essential for their potential use in biomedicine.

Data availability

No primary research results, software or code have been included and no new data were generated or analysed as part of this review.

Author contributions

Conceptualisation: E. B., B. S.; writing original draft E. B.; writing-review & editing E. B., B. S.; funding acquisition E. B., B. S.

Conflicts of interest

There are no conflicts to declare.

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Notes and references

- 1 T. Ruysschaert, A. F. P. Sonnen, T. Haeefe, W. Meier, M. Winterhalter and D. Fournier, *J. Am. Chem. Soc.*, 2005, **127**, 6242–6247.
- 2 K. Kostarelos, P. F. Luckham and T. F. Tadros, *J. Chem. Soc., Faraday Trans.*, 1998, **94**, 2159–2168.
- 3 K. Kostarelos, P. F. Luckham and T. F. Tadros, *J. Colloid Interface Sci.*, 1997, **191**, 341–348.
- 4 K. Kostarelos, T. F. Tadros and P. F. Luckham, *Langmuir*, 1999, **15**, 369–376.
- 5 E. Rideau, R. Dimova, P. Schwill, F. R. Wurm and K. Landfester, *Chem. Soc. Rev.*, 2018, **47**, 8572–8610.
- 6 X. Qian, I. Nymann Westensee, E. Brodzkij and B. Städler, *Wiley Interdiscip. Rev.: Nanomed. Nanobiotechnol.*, 2021, **13**, e1683.
- 7 C. Guindani, L. C. da Silva, S. Cao, T. Ivanov and K. Landfester, *Angew. Chem., Int. Ed.*, 2022, **61**, e202110855.
- 8 W. Jiang, Z. Wu, Z. Gao, M. Wan, M. Zhou, C. Mao and J. Shen, *ACS Nano*, 2022, **16**, 15705–15733.
- 9 C. G. Palivan, R. Goers, A. Najer, X. Zhang, A. Car and W. Meier, *Chem. Soc. Rev.*, 2016, **45**, 377–411.
- 10 C. Xu, S. Hu and X. Chen, *Mater. Today*, 2016, **19**, 516–532.
- 11 W. M. Smigiel, P. Lefrançois and B. Poolman, *Emerging Top. Life Sci.*, 2019, **3**, 445–458.
- 12 W. Sato, T. Zajkowski, F. Moser and K. P. Adamala, *Wiley Interdiscip. Rev.: Nanomed. Nanobiotechnol.*, 2021, 1–21.
- 13 N. Amy Yewdall, A. F. Mason and J. C. M. Van Hest, *Interface Focus*, 2018, **8**, 20180023.
- 14 J. Shin, B. D. Cole, T. Shan and Y. Jang, *Biomacromolecules*, 2022, **23**, 1505–1518.
- 15 E. Reimhult and M. M. Virk, *J. Biomed. Res.*, 2021, **35**, 301–309.
- 16 M. Mohammadi, S. Taghavi, K. Abnous, S. M. Taghdisi, M. Ramezani and M. Alibolandi, *Adv. Funct. Mater.*, 2018, **28**, 1–18.
- 17 C. E. Meyer, S. L. Abram, I. Craciun and C. G. Palivan, *Phys. Chem. Chem. Phys.*, 2020, **22**, 11197–11218.
- 18 M. G. Gouveia, J. P. Wesseler, J. Ramaekers, C. Weder, P. B. V. Scholten and N. Bruns, *Chem. Soc. Rev.*, 2023, **52**, 728–778.
- 19 V. De Leo, F. Milano, A. Agostiano and L. Catucci, *Polymers*, 2021, **13**, 1027.
- 20 D. Liu, H. Sun, Y. Xiao, S. Chen, E. J. Cornel, Y. Zhu and J. Du, *J. Controlled Release*, 2020, **326**, 365–386.
- 21 R. Tenchov, R. Bird, A. E. Curtze and Q. Zhou, *ACS Nano*, 2021, **15**, 16982–17015.
- 22 M. Schulz and W. H. Binder, *Macromol. Rapid Commun.*, 2015, **36**, 2031–2041.
- 23 M. Schulz, A. Olubummo and W. H. Binder, *Soft Matter*, 2012, **8**, 4849–4864.
- 24 J. F. Le Meins, C. Schatz, S. Lecommandoux and O. Sandre, *Mater. Today*, 2013, **16**, 397–402.
- 25 Y. K. Go and C. Leal, *Chem. Rev.*, 2021, **121**, 13996–14030.
- 26 S. Winzen, M. Bernhardt, D. Schaeffel, A. Koch, M. Kappl, K. Koynov, K. Landfester and A. Kroeger, *Soft Matter*, 2013, **9**, 5883–5890.
- 27 W. Shen, J. Hu and X. Hu, *Chem. Phys. Lett.*, 2014, **600**, 56–61.
- 28 D. Chen and M. M. Santore, *Soft Matter*, 2015, **11**, 2617–2626.
- 29 L. Otrin, N. Marušič, C. Bednarz, T. Vidaković-Koch, I. Lieberwirth, K. Landfester and K. Sundmacher, *Nano Lett.*, 2017, **17**, 6816–6821.
- 30 N. Marušič, L. Otrin, Z. Zhao, R. B. Lira, F. L. Kyrilis, F. Hamdi, P. L. Kastritis, T. Vidaković-Koch, I. Ivanov, K. Sundmacher and R. Dimova, *Proc. Natl. Acad. Sci. U. S. A.*, 2020, **117**, 15006–15017.
- 31 L. Otrin, A. Witkowska, N. Marusic, Z. Zhao, R. B. Lira, F. L. Kyrilis, F. Hamdi, I. Ivanov, R. Lipowsky, P. L. Kastritis, R. Dimova, K. Sundmacher, R. Jahn and T. Vidaković-Koch, *Nat. Commun.*, 2021, **12**, 4972.
- 32 N. Marušič, L. Otrin, J. Rauchhaus, Z. Zhao, F. L. Kyrilis, F. Hamdi, P. L. Kastritis, R. Dimova, I. Ivanov and K. Sundmacher, *Proc. Natl. Acad. Sci. U. S. A.*, 2022, **119**, e2122468119.
- 33 M. Chemin, P. M. Brun, S. Lecommandoux, O. Sandre and J. F. Le Meins, *Soft Matter*, 2012, **8**, 2867–2874.
- 34 T. P. T. Dao, F. Fernandes, M. Er-Rafik, R. Salva, M. Schmutz, A. Brûlet, M. Prieto, O. Sandre and J. F. Le Meins, *ACS Macro Lett.*, 2015, **4**, 182–186.
- 35 T. P. T. Dao, F. Fernandes, E. Ibarboure, K. Ferji, M. Prieto, O. Sandre and J. F. Le Meins, *Soft Matter*, 2017, **13**, 627–637.



- 36 T. P. T. Dao, A. Brûlet, F. Fernandes, M. Er-Rafik, K. Ferji, R. Schweins, J. P. Chapel, A. Fedorov, M. Schmutz, M. Prieto, O. Sandre and J. F. Le Meins, *Langmuir*, 2017, **33**, 1705–1715.
- 37 T. P. T. Dao, F. Fernandes, M. Fauquignon, E. Ibarboure, M. Prieto and J. F. Le Meins, *Soft Matter*, 2018, **14**, 6476–6484.
- 38 V. Passos Gibson, M. Fauquignon, E. Ibarboure, J. Leblond Chain and J.-F. Le Meins, *Polymers*, 2020, **12**, 637.
- 39 M. Fauquignon, E. Ibarboure, S. Carlotti, A. Brûlet, M. Schmutz and J. F. Le Meins, *Polymers*, 2019, **11**, 2013.
- 40 M. Fauquignon, E. Courtecuisse, R. Josselin, A. Mutschler, A. Brûlet, M. Schmutz and J. Le Meins, *J. Colloid Interface Sci.*, 2021, **604**, 575–583.
- 41 M. Fauquignon, E. Ibarboure and J. F. Le Meins, *Soft Matter*, 2021, **17**, 83–89.
- 42 M. Fauquignon, E. Ibarboure and J. F. Le Meins, *Biophys. J.*, 2022, **121**, 61–67.
- 43 R. M. Perera, S. Gupta, T. Li, C. J. Van Leeuwen, M. Bleuel, K. Hong and G. J. Schneider, *ACS Appl. Polym. Mater.*, 2022, **4**, 8858–8868.
- 44 Z. Cheng, D. R. Elias, N. P. Kamat, E. D. Johnston, A. Poloukhtine, V. Popik, D. A. Hammer and A. Tsourkas, *Bioconjugate Chem.*, 2011, **22**, 2021–2029.
- 45 X. Su, S. K. Mohamed Moinuddeen, L. Mori and M. Nallani, *J. Mater. Chem. B*, 2013, **1**, 5751–5755.
- 46 S. K. Lim, H. P. de Hoog, A. N. Parikh, M. Nallani and B. Liedberg, *Polymers*, 2013, **5**, 1102–1114.
- 47 S. Khan, M. Li, S. P. Muench, L. J. C. Jeuken and P. A. Beales, *Chem. Commun.*, 2016, **52**, 11020–11023.
- 48 S. Rottet, S. Iqbal, P. A. Beales, A. Lin, J. Lee, M. Rug, C. Scott and R. Callaghan, *Polymers*, 2020, **12**, 1049.
- 49 R. Catania, J. Machin, M. Rappolt, S. P. Muench, P. A. Beales and L. J. C. Jeuken, *Macromolecules*, 2022, **55**, 3415–3422.
- 50 R. Seneviratne, G. Coates, Z. Xu, C. E. Cornell, R. F. Thompson, A. Sadeghpour, D. P. Maskell, L. J. C. Jeuken, M. Rappolt and P. A. Beales, *Small*, 2023, **22**, 2206267.
- 51 R. Seneviratne, L. J. C. Jeuken, M. Rappolt and P. A. Beales, *Polymers*, 2020, **12**, 914.
- 52 R. Seneviratne, R. Catania, M. Rappolt, L. J. C. Jeuken and P. A. Beales, *Soft Matter*, 2022, **18**, 1294–1301.
- 53 R. Seneviratne, S. Khan, E. Moscrop, M. Rappolt, S. P. Muench, L. J. C. Jeuken and P. A. Beales, *Methods*, 2018, **147**, 142–149.
- 54 C. Kleineberg, C. Wölfer, A. Abbasnia, D. Pischel, C. Bednarz, I. Ivanov, T. Heitkamp, M. Börsch, K. Sundmacher and T. Vidaković-Koch, *ChemBioChem*, 2020, **21**, 2149–2160.
- 55 J. Nam, P. A. Beales and T. K. Vanderlick, *Langmuir*, 2011, **27**, 1–6.
- 56 A. Mech-Doros, N. Bajraktari, C. Hélix-Nielsen, J. Emnéus and A. Heiskanen, *Anal. Bioanal. Chem.*, 2020, **412**, 6307.
- 57 I. M. Henderson and W. F. Paxton, *Angew. Chem., Int. Ed.*, 2014, **53**, 3372–3376.
- 58 W. F. Paxton, P. T. McAninch, K. E. Achyuthan, S. H. R. Shin and H. L. Monteith, *Colloids Surf., B*, 2017, **159**, 268–276.
- 59 K. L. Willes, J. R. Genchev and W. F. Paxton, *Polymers*, 2020, **12**, 745.
- 60 K. L. Willes, S. A. McFarland, T. E. Johnson, D. R. Hart and W. F. Paxton, *ACS Appl. Nano Mater.*, 2022, **5**, 13820–13828.
- 61 M. M. Virk and E. Reimhult, *Langmuir*, 2018, **34**, 395–405.
- 62 M. M. Virk, B. Hofmann and E. Reimhult, *Langmuir*, 2019, **35**, 739–749.
- 63 N. Hamada and M. L. Longo, *Biochim. Biophys. Acta, Biomembr.*, 2022, **1864**, 183887.
- 64 N. Hamada, S. Gakhar and M. L. Longo, *Biochim. Biophys. Acta, Biomembr.*, 2021, **1863**, 183552.
- 65 N. Hamada and M. L. Longo, *Biochim. Biophys. Acta, Biomembr.*, 2022, **1864**, 184026.
- 66 J. Steinkühler, M. L. Jacobs, M. A. Boyd, C. G. Villaseñor, S. M. Loverde and N. P. Kamat, *Biomacromolecules*, 2022, **23**, 4756–4765.
- 67 M. E. Palanco, N. Skovgaard, J. S. Hansen, K. Berg-Sorensen and C. Hélix-Nielsen, *Bioinspiration Biomimetics*, 2018, **13**, 016005.
- 68 C. Magnani, C. Montis, G. Mangiapia, A. F. Mingotaud, C. Mingotaud, C. Roux, P. Joseph, D. Berti and B. Lonetti, *Colloids Surf., B*, 2018, **168**, 18–28.
- 69 A. Olubummo, M. Schulz, B. Lechner, P. Scholtysek, K. Bacia, A. Blume and W. H. Binder, *ACS Nano*, 2012, **6**, 8713–8727.
- 70 M. Schulz, D. Glatte, A. Meister, P. Scholtysek, A. Kerth, A. Blume, K. Bacia and W. H. Binder, *Soft Matter*, 2011, **7**, 8100–8110.
- 71 M. Schulz, S. Werner, K. Bacia and W. H. Binder, *Angew. Chem., Int. Ed.*, 2013, **52**, 1829–1833.
- 72 M. Bieligmeyer, F. Artukovic, S. Nussberger, T. Hirth, T. Schiestel and M. Müller, *Beilstein J. Nanotechnol.*, 2016, **7**, 881–892.
- 73 N. Pippa, E. Kaditi, S. Pispas and C. Demetzos, *Soft Matter*, 2013, **9**, 4073–4082.
- 74 N. Pippa, D. R. Perinelli, S. Pispas, G. Bonacucina, C. Demetzos, A. Forsys and B. Trzebicka, *Colloids Surf., A*, 2018, **555**, 539–547.
- 75 M. A. Palominos, D. Vilches, E. Bossel and M. A. Soto-Arriaza, *Colloids Surf., B*, 2016, **148**, 30–40.
- 76 A. K. Khan, J. C. S. Ho, S. Roy, B. Liedberg and M. Nallani, *Polymers*, 2020, **12**, 979.
- 77 Y. K. Go, N. Kamar and C. Leal, *Polymers*, 2020, **12**, 9–12.
- 78 S. Khan, J. McCabe, K. Hill and P. A. Beales, *J. Colloid Interface Sci.*, 2020, **562**, 418–428.
- 79 K. Flandez, S. Bonardd and M. Soto-Arriaza, *Chem. Phys. Lipids*, 2020, **230**, 104927.
- 80 H. F. Li, C. Wu, M. Xia, H. Zhao, M. X. Zhao, J. Hou, R. Li, L. Wei and L. Zhang, *RSC Adv.*, 2015, **5**, 27630–27639.
- 81 N. Pippa, D. Stellas, A. Skandalis, S. Pispas, C. Demetzos, M. Libera, A. Marcinkowski and B. Trzebicka, *Eur. J. Pharm. Biopharm.*, 2016, **107**, 295–309.
- 82 N. Naziris, A. Skandalis, A. Forsys, B. Trzebicka, S. Pispas and C. Demetzos, *J. Therm. Anal. Calorim.*, 2020, **141**, 751–766.



- 83 N. Naziris, N. Pippa, V. Chrysostomou, S. Pispas, C. Demetzos, M. Libera and B. Trzebicka, *J. Nanopart. Res.*, 2017, **19**, 347.
- 84 N. Naziris, N. Pippa, E. Sereti, V. Chrysostomou, M. Kędzierska, J. Kajdanek, M. Ionov, K. Miłowska, Ł. Balcerzak, S. Garofalo, C. Limatola, S. Pispas, K. Dimas, M. Bryszewska and C. Demetzos, *Int. J. Mol. Sci.*, 2021, **22**, 6271.
- 85 K. P. Mineart, S. Venkataraman, Y. Y. Yang, J. L. Hedrick and V. M. Prabhu, *Macromolecules*, 2018, **51**, 3184–3192.
- 86 W. Zong, B. Thingholm, F. ITEL, P. S. Schattling, E. Brodzkij, D. Mayer, S. Stenger, K. N. Goldie, X. Han and B. Städler, *Langmuir*, 2018, **34**, 6874–6886.
- 87 Y. Zhang, N. Gal, F. ITEL, I. N. Westensee, E. Brodzkij, D. Mayer, S. Stenger, M. Castellote-Borrell, T. Boesen, S. R. Tabaei, F. Höök and B. Städler, *Nanoscale*, 2019, **11**, 11530–11541.
- 88 E. Taipaleenmäki, G. Christensen, E. Brodzkij, S. A. Mouritzen, N. Gal, S. Madsen, M. S. Hedemann, T. A. Knudsen, H. M. Jensen, S. L. Christiansen, F. V. Sparsø and B. Städler, *J. Controlled Release*, 2020, **322**, 470–485.
- 89 E. Brodzkij, I. N. Westensee, M. Bertelsen, N. Gal, T. Boesen and B. Städler, *Small*, 2020, **16**, 1906493.
- 90 P. De Dios Andres, I. N. Westensee, E. Brodzkij, M. A. Ramos-Docampo, N. Gal and B. Städler, *Biomacromolecules*, 2021, **22**, 3860–3872.
- 91 C. Ade, X. Qian, E. Brodzkij, P. De Dios Andres, J. Spanjers, I. N. Westensee and B. Städler, *Biomacromolecules*, 2022, **23**, 1052–1064.
- 92 C. A. Meyer, P. D. D. Andres, E. Brodzkij, I. N. Westensee, J. Lyons, S. H. Vaz and B. Städler, *Adv. Biol.*, 2022, 2200209.
- 93 E. Brodzkij, I. N. Westensee, S. F. Holleufer, C. A. Ade, P. De Dios Andres, J. S. Pedersen, B. Stadler, P. D. D. Andres, J. S. Pedersen and B. Städler, *Applied Mater. Today*, 2022, **29**, 101549.
- 94 J. M. Spanjers, E. Brodzkij, N. Gal and B. Stadler, *Eur. Polym. J.*, 2022, **180**, 111612.
- 95 T. Nishimura, S. Hirose, Y. Sasaki and K. Akiyoshi, *J. Am. Chem. Soc.*, 2020, **142**, 154–161.
- 96 N. Ozawa, S. Kosaka, S. Fujii and T. Nishimura, *Polym. Chem.*, 2023, **14**, 2198–2204.
- 97 P. Walde, K. Cosentino, H. Engel and P. Stano, *ChemBioChem*, 2010, **11**, 848–865.
- 98 P. A. Beales, B. Ciani and A. J. Cleasby, *Phys. Chem. Chem. Phys.*, 2015, **17**, 15489–15507.
- 99 A. Peyret, E. Ibarboure, J. F. Le Meins and S. Lecommandoux, *Adv. Sci.*, 2018, **5**, 1–8.
- 100 D. Chandler, *Nature*, 2005, **437**, 600–647.
- 101 H.-C. Tsai, Y.-L. Yang, Y.-J. Sheng and H.-K. Tsao, *Polymers*, 2020, **12**, 639.
- 102 T. Baumgart, S. T. Hess and W. W. Webb, *Nature*, 2003, **425**, 821–824.
- 103 S.-W. Hu, C.-Y. Huang, H.-K. Tsao and Y.-J. Sheng, *Phys. Rev. E*, 2019, **99**, 1–11.
- 104 J. Nam, T. K. Vanderlick and P. A. Beales, *Soft Matter*, 2012, **8**, 7982–7988.
- 105 E. L. Elson, *Biophys. J.*, 2011, **101**, 2855–2870.
- 106 A. G. Jay and J. A. Hamilton, *J. Fluoresc.*, 2017, **27**, 243–249.
- 107 J. E. Nielsen, T. K. Lind, A. Lone, Y. Gerelli, P. R. Hansen, H. Jenssen, M. Cárdenas and R. Lund, *Biochim. Biophys. Acta, Biomembr.*, 2019, **1861**, 1355–1364.
- 108 J. S. Pedersen, C. Svaneborg, K. Almdal, I. W. Hamley and R. N. Young, *Macromolecules*, 2003, **36**, 416–433.
- 109 D. E. Discher and A. Eisenberg, *Science*, 2002, **297**, 967.
- 110 A. Larrañaga, M. Lomora, J. R. Sarasua, C. G. Palivan and A. Pandit, *Prog. Mater. Sci.*, 2017, **90**, 325–357.
- 111 A. Belluati, I. Craciun, C. E. Meyer, S. Rigo and C. G. Palivan, *Curr. Opin. Biotechnol.*, 2019, **60**, 53–62.
- 112 H. Che and J. C. M. van Hest, *ChemNanoMat*, 2019, **5**, 1092–1109.
- 113 S. Hirschi, T. R. Ward, W. P. Meier, D. J. Muller and D. Fotiadis, *Chem. Rev.*, 2022, **122**, 16294–16328.
- 114 W. Yuan, J. Piao and Y. Dong, *Mater. Chem. Front.*, 2021, **5**, 5233–5246.
- 115 K. A. Podolsky and N. K. Devaraj, *Nat. Rev. Chem*, 2021, **5**, 676–694.
- 116 M. L. Jacobs, M. A. Boyd and N. P. Kamat, *Proc. Natl. Acad. Sci. U. S. A.*, 2019, **116**, 4031–4036.

