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# Phase Separation and Nanodomain Formation in Hybrid Polymer/ Lipid Vesicles

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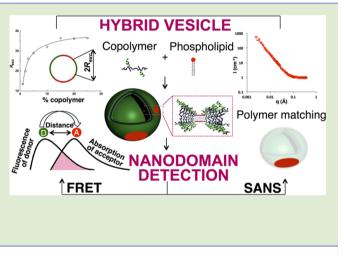
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**Supporting Information** 

**ABSTRACT:** Hybrid polymer/lipid large unilamellar vesicles (LUVs) were studied by small angle neutron scattering (SANS), time-resolved Förster resonance energy transfer (TR-FRET), and cryo-transmission electron microscopy (cryo-TEM). For the first time in hybrid vesicles, evidence for phase separation at the nanoscale was obtained, leading to the formation of stable nanodomains enriched either in lipid or polymer. This stability was allowed by using vesicle-forming copolymer with a membrane thickness close to the lipid bilayer thickness, thereby minimizing the hydrophobic mismatch at the domain periphery. Hybrid giant unilamellar vesicles (GUVs) with the same composition have been previously shown to be unstable and susceptible to fission, suggesting a role of curvature in the stabilization of nanodomains in these structures.

ipid nanodomains in biological membranes are thought to ✓ play a key role in several cell processes,<sup>1,2</sup> but their small size (10-200 nm) implies complex detection and characterization methodologies. Therefore, many theoretical and experimental works<sup>3,4</sup> were devoted to synthetic model membranes in order to unveil the salient features and the parameters governing the formation of such domains and the modulation of their size. Studies performed on multicomponent lipid vesicles showed that the fluidity of the different lipids and lipid-lipid interactions through their hydrophobic tails or polar head play important roles on phase separation and, consequently, on domains formation.<sup>5</sup> So far, the energy per unit length at the domain boundary (line tension) was shown to be of paramount importance. Bilayer thickness mismatch in multicomponent lipid vesicles modulates this line tension and domain size, as nicely evidenced by Heberle et al. using small angle neutron scattering (SANS) with contrast variation.<sup>6</sup> In parallel, polymersomes obtained by selfassembly of amphiphilic block copolymers emerged in the late 90s as an alternative to liposomes, for example, in the field of drug delivery: the improved stability of their membrane compared to liposomes and the ease to modulate their functionality by coupling chemistry offers unprecedented new



possibilities.<sup>7</sup> The formation of domains in polymersome membrane was also investigated, in order to tune their membrane properties.<sup>8,9</sup>

Recently, hybrid polymer/lipid vesicles appeared as an ideal "upgrade"<sup>10</sup> of their forerunners as they can marry in a single membrane the best characteristics of the two different systems (stability, biofunctionality, controlled permeability, and so on). Promising results were observed regarding for instance drug targeting or biomolecular recognition.<sup>11,12</sup> To date, physical and molecular parameters governing the phase separation in these hybrid membranes are not well understood. In addition to the expected chemical incompatibility between polymer block chains and lipids, one has to consider also their dimensional differences. In order to perfectly exploit the potential of such structures, phase separation needs to be tuned, especially the way to formulate lipid nanodomains. As previously pointed out, the characterization of nanodomains in multicomponent vesicles is not straightforward, as they cannot be observed directly by optical microscopy. Therefore, only

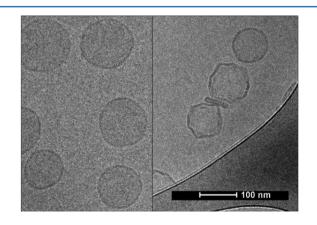
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rather few results are available for lipid systems,<sup>6,13–18</sup> and no evidence has yet been offered for hybrid polymer/lipid small or large vesicles.<sup>10</sup> We seek to clarify the membrane structure of hybrid polymer vesicles and especially the way to obtain nanodomains enriched (or pure) either in lipids or in polymers.

Although biophysical mechanisms that prevent ripening of small lipid domains in a membrane involve several parameters and are not yet completely understood, we decided to minimize line tension, which naturally arises from the thickness mismatch between lipid domain and the surrounding polymer membrane, since this parameter has been shown to play a role in the membrane structuration of giant hybrid vesicles.<sup>19,20</sup> In this way, we selected a close to matching system, using a commercial grafted copolymer with a flexible poly-(dimethylsiloxane) backbone and two poly(ethylene oxide) pendant moieties (PDMS-g-(PEO)<sub>2</sub>) of  $M_w = 3000 \text{ g} \cdot \text{mol}^{-1}$  (see Supporting Information for other details).

This copolymer is well-known to form vesicles by itself with a membrane thickness ( $\sim$ 5 nm) close to that of liposomes ( $\sim$ 3-4 nm).<sup>21,22</sup> It was mixed with 1,2-dipalmitoyl-sn-glycero-3phosphocholine (DPPC) or palmitoyl-oleoyl-sn-glycero-3phosphocholine (POPC) at molar ratios of 50/50 (~80/20 in polymer/lipid weight ratio). For the same composition in giant unilamellar vesicles (GUV), phase-separation with micrometer-sized lipid domains was observed by fluorescence microscopy.<sup>10,20</sup> The present study was performed with lipid either in a fluid state (POPC at room temperature or DPPC at 46 °C) or in the gel state (DPPC at room temperature). The vesicles produced (see Supporting Information, section S2.2) were analyzed by dynamic and static light scattering (DLS/ SLS). They possess a narrow size distribution and a hydrodynamic radius in agreement with the extrusion process (see Figure S-3). The hybrid character of the vesicles was checked by flow cytometry (FACS) and zeta potential measurements. (Supporting Information, Figures S-4, S-5, S-6).

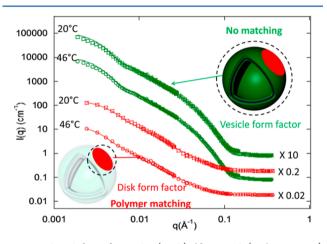
The nanostructures formed were characterized by cryo-TEM which has recently been used to characterize homogeneous hybrid polymer/lipid vesicles obtained from mixtures of another copolymer (PDMS<sub>60</sub>-*b*-PMOXA<sub>21</sub>) with DMPC.<sup>23</sup> Figure 1 is a representative micrograph of the two main morphologies observed: on the one hand, rounded spherical vesicles, on the other hand, faceted vesicles. The faceting is ascribed to the gel state of the DPPC phase. Similar pictures were indeed already obtained with pure DPPC vesicles in previous studies<sup>24</sup> (see also control in Supporting Information,



**Figure 1.** Cryo-TEM pictures of 80/20 weight ratio PDMS-*g*-(PEO)<sub>2</sub>/ DPPC vesicles, quenched from room temperature.

section S3.6). Rounded spherical vesicles are probably hybrid vesicles in which the lipid content is too small to see the faceting effect. It is important to note that the polymer/lipid composition is not perfectly controlled within the vesicle population as vesicles are known to be out-of-equilibrium objects<sup>25-27</sup>

To get more information about the membrane structure of LUV hybrid vesicles, we performed SANS experiments using  $d_{62}$ -DPPC, that is, DPPC with fully deuterated fatty chains. To separate the contribution to the scattering of the lipid and of the copolymer components, two  $D_2O/H_2O$  mixtures were used (Supporting Information, section S3.4). LUV hybrid vesicles were also prepared in  $D_2O$  with classical DPPC. Thus, the solvent mixture matching the polymer enables the observation the lipid phase, while the lipid-matching solvent allows seeing the copolymer. Without matching, we can observe the whole hybrid vesicles. Such results are illustrated in Figure 2.



**Figure 2.** SANS data of PDMS-g-(PEO)<sub>2</sub>/ $d_{62}$ -DPPC (80/20 wt ratio) hybrid vesicles at 20 and 46 °C in polymer matching and no matching conditions. Solid lines: fitting curves.

Interestingly, the curves obtained in polymer matching condition could not be fitted with the vesicle form factor commonly used to model pure lipid or polymer vesicles, but they were well fitted with a polydisperse flat cylinder (disk-like) form factor (Supporting Information, section S3.4). The fitting values of radius ( $R_{cyl}$ ) and height (L) indicated in Table 1 suggest that lipid phase presents disk-like shapes, both at 20

Table 1. Parameters Obtained by Either Fitting the SANS Curves from Hybrid PDMS-g-(PEO)<sub>2</sub>/DPPC (50/50 mol ratio) with Flat Cylinder (Polymer Matching) or Vesicle\* (No Matching) Form Factor, and from Guinier and Kratky– Porod Representations

	$R_{cyl} \pm \sigma_{R}$ (nm) or $R_{ves} \pm \sigma_{R}$ (nm)*	R <sub>G</sub> (nm; Guinier plot)	flat cylinder $L_{\rm cyl} \pm \sigma_{\rm L} (\rm nm)$ or shell thickness $\delta$ (nm)*	membrane thickness (nm; Kratky–Porod plot)
polymer matching 20 °C	54 ± 9	35	3.4 ± 0.7	3.4
polymer matching 46 °C	73 ± 7	43	3.1 ± 0.9	3.3
no match- ing 20 °C	$30 \pm 17^*$	52	$5.1 \pm 1.2^{*}$	5.8
no match- ing 46 °C	$37 \pm 14^{*}$	54	$5.1 \pm 1.2^{*}$	5.6

and 46 °C, whose thicknesses well correspond to the one of a pure lipid bilayer. Radius of gyration  $(R_G)$  and membrane thicknesses ( $\delta$ ) were also estimated using Guinier and Kratky– Porod plots, respectively (see Supporting Information). Disk radii calculated from the independently measured values of  $R_{G}$ and  $\delta$  through the equation  $R_{\rm G}^2 = R_{\rm cyl}^2/2 + \delta^2/12$  are in good agreement with disk radii obtained using the disk-like form factor fit. It is interesting to note that the lipid/polymer volume ratio estimated geometrically is  $\pi R_{cyl}^2 \times L_{cyl}/(4\pi R_G^2 \times \delta) =$ 0.18 at 20 °C and 0.28 at 46 °C, thus, not differing too much from the mass ratio of 0.20 calculated from the composition and the molar masses. In lipid matching condition, data could be precisely fitted with neither a cylinder nor a vesicle form factor either (Supporting Information, section S3.4). The curves would probably be fitted by a holey shell form factor that needs to be computed numerically in further studies.

In pure D<sub>2</sub>O (no matching), the data are very well fitted with a vesicle form factor as shown in Figure 2: the characteristic parameters are indicated in Table 1, as well as estimates of  $R_{\rm G}$ and  $\delta$ . The values are in agreement with the size of the vesicles expected from the extrusion process ( $R \sim 50$  nm) and with membrane thicknesses reported previously for DPPC<sup>28</sup> and for the copolymer.<sup>22,29</sup>

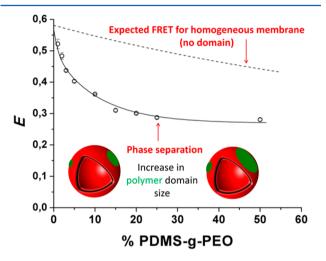
This set of results proves that hybrid vesicles are obtained and that phase separation occurs within the membrane, leading to the formation of lipid domains detected as disks by SANS, floating in the surrounding polymer membrane. It also appears that phase separation is still present above the melting transition of DPPC, although the characteristic values of disk radii seem to be a little too large compared to the measured radii of gyration of the vesicles ( $R_{\rm G} \sim 56$  nm by SLS and  $R_{\rm G} \sim$ 52 nm by SANS in no matching condition), especially at 46 °C. As previously mentioned, polymer/lipid composition is not perfectly controlled from a vesicle to another and that probably leads to higher dispersity in disk sizes and to the existence of a population with lower lipid contents. Also, we insist of the point that the term "disk" is not rigorous for these lipid domains surrounded by polymer membrane as their shape may not be circular, especially in the gel state, and should follow the convex curvature of the vesicle in the fluid state. These aspects are not taken into account in our simple fitting procedure, which however correctly describes the phase-separation phenomenon. The parameter that is measured the most precisely is the membrane thickness, whose values are close to the expected ones. A more precise estimate of the lipid domain size and shape by small angle neutron scattering deserves to be undergone in the future by a comprehensive study on a larger range of compositions in a higher flux neutron reactor.

To get more insight into the membrane structure of these hybrid vesicles, and especially to probe demixing at the nanometer scale between the lipid and polymer phases, we used Förster resonance energy transfer (FRET).<sup>17</sup> This is a powerful method to detect and characterize lateral membrane domains presenting sizes smaller than 50-100 nm. PDMS-g-(PEO)<sub>2</sub> tagged with the NBD molecule and *N*-(lissamine Rhodamine B sulfonyl)-1,2-dioleoyl-*sn*-3-phosphatidylehanolamine (Rhod-PE) were used, respectively, as the donor and acceptor pair (see section S3.5 of Supporting Information for experimental procedure and data analysis). At first, the phase separation within the membrane was proven by measuring the partition coefficient of the probes, PDMS-g-(PEO)<sub>2</sub>-NBD and Rhod-PE in both DPPC and POPC with PDMS-g-(PEO)<sub>2</sub> hybrid

vesicles when the lipid was in its fluid state (Supporting Information, section S3.5.1).

The PDMS-g-(PEO)<sub>2</sub>-NBD probe partitions almost exclusively in the polymer phase, which implicitly demonstrates phase separation between polymer and lipid. We also observed that the labeled lipid incorporates in the polymer phase to some extent (39 and 37 mol %) for equimolar PDMS-g-(PEO)<sub>2</sub>/DPPC and PDMS-g-(PEO)<sub>2</sub>/POPC mixtures, respectively (see Supporting Information, section S3.5.1).

The phase separation was further proven by comparing the experimental FRET efficiency in PDMS-g-(PEO)<sub>2</sub>-NBD/POPC/DOPE-Rho (1/98.5/0.5 mol/mol/mol) mixed vesicles (Supporting Information, section S3.5.3) with the theoretical expectation value (formalisms described in Supporting Information, section S3.5.2). FRET efficiencies were measured at different polymer/lipid ratios (Figure 3). The significant



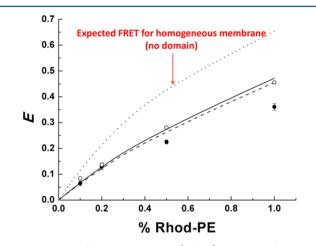
**Figure 3.** FRET efficiency vs content (mol %) PDMS-*g*-(PEO)<sub>2</sub>. Solid line is a guide to the eyes.

decrease in FRET efficiencies observed for increasing molar fractions of polymer is consistent with the increase of polymer domain sizes, and therefore increase of the average distance between labeled PDMS-g-(PEO)<sub>2</sub>-NBD donor and Rhod-PE acceptor. In the case of a homogeneous lipid—polymer membrane, the predicted decrease in FRET efficiency upon increase of PDMS-g-(PEO)<sub>2</sub> molar fraction would be much smaller (dotted line in Figure 3).

In the following, we prove that the plateau observed at PDMS-g-(PEO)<sub>2</sub>/PC proportion higher than 25 mol % is due to the formation of polymer-rich domains larger that 5–10 times  $R_0$ , the Förster distance for this donor–acceptor pair, that is, about 25–50 nm (see Supporting Information, sections S3.5.2 and S3.5.4, for details). The acceptor Rhod-PE lipid is still present in these polymer raft-like domains, as already quantified through its partition coefficient between the lipid and polymer phases and reported in the preceding paragraph.

We measured the FRET efficiency within the PDMS-g-(PEO)<sub>2</sub>-rich phase of the 50/50 PDMS-g-(PEO)<sub>2</sub>/DPPC mixture with different concentrations of the acceptor Rhod-PE. The recovered experimental FRET efficiencies are in perfect agreement with the FRET values calculated with acceptor concentrations obtained from the partition coefficient experiments and well below those expected for a homogeneous membrane. This demonstrates that there exists a negligible percentage of bordering PDMS-g-(PEO)<sub>2</sub>-NBD donor mole-

cules transferring to acceptors in the DPPC-rich phase and provides good evidence for diameters of PDMS-g-(PEO)<sub>2</sub> domains larger than 25–50 nm in these hybrid polymer/lipid vesicles. A more detailed FRET experiment was conducted on a 50/50 POPC/PDMS-g-(PEO)<sub>2</sub> mixture with comparable results (Figure 4).



**Figure 4.** FRET efficiencies vs content (mol %) of acceptor for DPPC at 46 °C ( $\bigcirc$ ) and for POPC at 25 °C ( $\bigcirc$ ). Theoretical values for FRET (Supporting Information) with DPPC (-) or POPC (---).

In conclusion, this work evidenced from neutron scattering and FRET data that a nanoscale phase separation is obtained in hybrid polymer/lipid LUV, leading to the formation of stable nanodomains enriched in either lipid or polymer, coexisting in the same patchwork membrane of the vesicles.

These nanodomains, whose exact shape and size remain to be determined, are observed whatever the fluid or gel state of the lipid. We hypothesize that the use of a flexible copolymer, well-known to form polymersomes with a membrane thickness close to that of liposomes, decreases the thickness mismatch and allows a conformational adaptation of the polymer chains at the boundaries, decreasing the line tension and stabilizing the nanodomains, even in the fluid state. This thickness mismatch is of paramount importance for membrane structuration and control of domain sizes, as shown in the literature for membranes made by mixing lipids of various chain lengths.<sup>6,30</sup>

Interestingly, phase-separation inside these hybrid/polymer lipid vesicles was previously observed (through stable micrometric lipid domains) for giant vesicles obtained from the same polymer/lipid composition with lipid in a gel state.<sup>20</sup> However, with lipid in a fluid state for such composition, budding and fission phenomenon was observed after a few hours, leading to the formation of "pure" liposomes and polymersomes. There is no evidence of such a phenomenon in our experiments for LUVs. This suggests that membrane curvature could play a role also in the stabilization of nanodomains. A complete study in parallel on GUV, LUV, and SUV vesicles, involving several block copolymers with different molar masses and a large compositional range is currently in progress to gain more insight into the parameters governing the phase separation and the formation of nanodomains in hybrid polymer/lipid vesicles.

This study is a first important step to elucidate the properties of these new self-assembled hybrid structures, and to allow their optimization regarding different fields of application, namely the design of drug delivery nanosystems with particulate targeting capability (through the possibility of biofunctionalization of the lipid and/or the polymer nanodomains) or of biomimetic experiments aimed at reproducing for instance focal adhesion points of biological cells.

#### ASSOCIATED CONTENT

#### **S** Supporting Information

Structure of copolymer used, chemical modification procedures, light and neutrons scattering experiments, cryo-TEM, zeta potential, flow cytometry, FRET experiments, and analysis. This material is available free of charge via the Internet at http://pubs.acs.org.

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### Notes

The authors declare no competing financial interest.

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