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Effect of benzocaine and propranolol on phospholipid-based bilayers

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Cell membranes play a fundamental role in protecting the cell from its surroundings, in addition to hosting many proteins with fundamental biological tasks. A study of drug/lipid interactions is a necessary and important step in fully clarifying the role and action mechanism of active ingredients, and shedding light on possible complications caused by drug overdose. In this paper, the influence of benzocaine and propranolol drugs on the structure of $\text{L-}\alpha$ -phosphatidylcholine-based membranes has been investigated by means of neutron reflectivity, grazing incidence small angle neutron scattering, and small/ultra-small angle neutron scattering. Investigations allowed discovering a stiffening of the membranes and the formation of stalks, caused by the presence of benzocaine. On the other hand, disordered bilayers (lamellar powders) and highly curved structures were found in the presence of propranolol. The results obtained may be rationalized in terms of the molecular structures of drugs and may serve as a starting point for explaining the toxic behavior in long-term and overdose scenarios.

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Introduction

Cell membranes consist mainly of lipid bilayers as their structural unit.¹ In turn, these bilayers are composed of phospholipids, amphiphilic molecules formed by a functionalized hydrophilic phosphate head esterified with two fatty acid chains.

The main task of a cell membrane is essentially to protect the cell from its surroundings, acting as a selective barrier between the internal cytoplasm and the extracellular environment.² Part of its selectivity is due to the presence of different proteins (known as membrane proteins) that are embedded in the lipid bilayer. The biological functions of these proteins also include the possibilities of activating cell signaling, triggering intracellular signaling pathways, and interacting with external substances, including active ingredients.³ To this aim, most of the drugs designed or used in pharmacological therapies target membrane proteins,⁴ like channel blockers or protein binding inhibitors.⁵

For the above mentioned reasons, most of the action mechanism studies, or pharmacokinetic investigations, are exclusively focused on drug/protein interactions, while a possible drug/phospholipid

interaction is usually neglected. This approach, aside from being not fully justified, neglects some key aspects of the interaction between active ingredients and cellular systems. Concerning this, even in the presence of a specific drug interaction with a particular receptor, a partial drug uptake by the cell membrane may significantly influence the pharmacokinetic properties of the drug, such as its delivery, distribution, and elimination, modulating the overall efficacy of the administration. The second key aspect to consider for a proper and meaningful investigation concerns the structure of the target protein. In fact, action mechanism and pharmacological studies require, for a correct understanding, that the protein has the same conformation assumed inside the cell membrane, which is possible only when the protein is embedded in a lipid bilayer membrane. A scenario in which the protein is extracted from the bilayer, purified, and studied is hardly possible, because of its sensitive nature that often results in a denaturation process once extracted.

For what has been said, a study of drug/lipid interactions appears to be a necessary and important step in fully understanding and properly evaluating the role played by the two different interactions – drug/protein and drug/phospholipid. Such a study is also important, when combined with complementary investigations, for developing and finding new active ingredients with improved pharmacokinetic characteristics. Indeed, in a limited number of cases, different families of drugs such as antibiotics,^{6–9} antimycotics, and^{10,11} antipsychotics,^{12,13} as well as anticancer agents,^{14,15} have been studied from the point of view of their possible interaction with lipids,¹⁶ even through simulation studies.¹⁷ It has often been found that, even in the absence of

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a specific drug/membrane interaction, the binding of the drug to the membrane reduces the availability of the free drug, making the pharmacological effects blander.^{18–20}

The high complexity shown by cell membranes makes their physico-chemical and biophysical interactions with drugs very challenging to investigate. To this aim, simplified membrane systems, like liposomes,²¹ giant unilamellar vesicles,²² Langmuir monolayers,²³ or solid supported membranes,²⁴ have been developed for their capability to mimic closely the natural lipid bilayer.

In this paper we report an investigation of the interaction of benzocaine and propranolol active ingredients with SoyPC phospholipid-based bilayers. Biomimetic model membranes, such as liposomes and lamellar phases, represent very good alternatives to natural membranes, since they allow investigating drug/membrane interactions under very precise and reproducible conditions, without departing from those that are the main features of a cell bilayer.

Benzocaine, chemically the ethyl ester of the 4-aminobenzoic acid, is a local anesthetic normally used as a pain reliever or for treatment of oral ulcers. It is the active ingredient of many non-prescription drugs, including those used for numb teething treatments; it is very commonly administered and it is estimated that around 30 billion doses are annually taken in the U.S.A.

Propranolol is a beta blocker medication. It is used for the treatment of high blood pressure and hyperthyroidism and in some cardiac arrhythmia scenarios.

Investigations at the nanoscale level have been carried out by means of Neutron Reflectivity (NR), Grazing Incidence Small Angle Neutron Scattering (GISANS), and Small/Ultra-Small Angle Neutron Scattering (SANS/USANS). Combination of these techniques is able to provide detailed information both on the supramolecular structures formed²⁵ and on bilayer characteristics,^{26,27} highlighting eventual variations caused by drug addition. In particular, NR is able to probe the nanoscopic characteristics of the bilayers in a very accurate way, providing information about the extensions of hydrophobic and hydrophilic domains inside the bilayers: part

of this accuracy is due to the capability of neutrons to depend on the nuclear composition of the sample, rather than on its electron density, like in the X-ray analogue technique. On the other hand, SANS measures the average structures in the volume probed and is ideal for examining the shape and size of supramolecular aggregates formed.

The results obtained from our investigations will show how the addition of very simple molecules (like those studied here) may modify very strongly the properties of the double layer. These findings strengthen the idea that a simple drug/protein study carried out by neglecting a (possible) interaction with the double layer may lead to a misleading or an incomplete picture of the system. Moreover, it will be noted how the present approach, involving complementary neutron-based techniques, is able to give valuable information about the systems investigated.

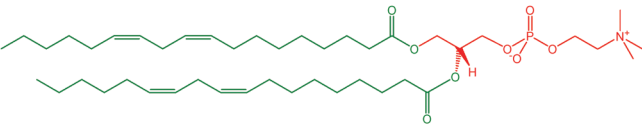
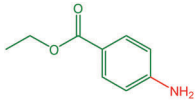
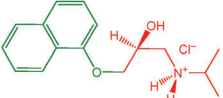
Going into detail, it will be seen that the two drugs exhibit different behaviors: benzocaine tends to stiffen the phospholipid-based double layer and to bend it locally with the formation of stalks, whereas propranolol causes a strong degree of disorder in the double layer, promoting the formation of distorted lamellae and structures with high curvature.

Experimental

Materials

SoyPC (mainly composed of 1,2-dilinoleoyl-*sn*-glycero-3-phosphocholine, purity 95%, averaged molecular weight 775.037 a.m.u.) was obtained from Avanti Polar Lipids Inc. (Alabaster, AL, USA). Benzocaine (purity 98%, molecular weight 165.19 a.m.u.) and propranolol (in the hydrochloride form, purity $\geq 99\%$, molecular weight 295.80 a.m.u.) were purchased from Sigma Aldrich (Taufkirchen, Germany). All the molecular structures, as well as the molecular volumes and the scattering length densities (SLD) of hydrophobic and hydrophilic portions, highlighted in green and red colors, respectively, are reported in Table 1.

Table 1 SLDs (ρ) of the molecules used in the experimental investigations. Values were used for the analysis of some reflectivity data by means of the Parratt algorithm. Molecular volumes have been estimated by using results obtained from MD simulations carried out on phospholipids²⁸ and from the theory of group contributions²⁹

Component	Molecular structure	$10^6 \times \rho/\text{\AA}^{-2}$	
		Hydrophobic portion	Hydrophilic portion
SoyPC		0.312	0.898
Benzocaine		1.72	1.10
Propranolol		2.21	0.508



Heavy water (minimum isotopic enrichment >99.8%, molecular weight 20.0276 a.m.u.) was obtained from ARMAR Chemicals (Leipzig, Germany) and used for the preparation of all the samples.

2-Propanol (molecular weight 60.10 a.m.u.) and trichloromethane (molecular weight 119.38 a.m.u.) pure for HPLC analysis were both purchased from Bachem (Bubendorf, Switzerland).

All chemicals were used as received, without further purification.

Preparation of samples for NR/GISANS experiments

Each sample used for NR and GISANS investigations was prepared starting from a standard solution of SoyPC in 2-propanol: to this aim, a selected amount of SoyPC was dissolved in the secondary alcohol, so as to achieve a molar ratio of $\cong 2$ mol% between the phospholipid and the solvent, and a total volume of $\cong 15$ ml. The mixed components were stirred to ensure complete dissolution of the phospholipid in the solvent. Afterwards, always under stirring, an appropriate amount of a 2-propanol-based stock solution of the drug (benzocaine or propranolol) was slowly added until a clear one-phase system was obtained.

Each of the prepared solutions was deposited on an ultrapolished Si block (r.m.s. roughness <2 Å, dimensions: 15.0–5.0–2.0 cm³) obtained from Andrea Holm GmbH (Tann, Germany): an O-ring in a custom-made scaffold allowed the solution to stay on top of the silicon block. The block was levelled out by means of a spirit level. Subsequently, the solution was allowed to evaporate at room temperature and at a reduced pressure (~ 250 mbar) for 8 h, in order to avoid the formation of bubbles inside the liquid phase due to a drastic decrease of the pressure. After that, the system was kept under a pressure of few millibars ($<10^{-3}$ mbar) for 16 h for removing all the residual isopropanol, leading to a SoyPC/drug layer of macroscopic size (1–2 mm thick). Just after this step, the silicon block was mounted into a custom made sample cell (sketched in Fig. 1), filled with heavy water and studied through NR and GISANS experiments. The cell was equipped with a cover glass, allowing visually inspecting the SoyPC/drug layer and checking for eventual air bubbles formed after injection of D₂O. Ocular inspection of the layer after the NR/GISANS investigations allowed excluding, in all prepared samples, a deterioration of the film coverage during the measurements.

All the silicon block active surfaces used for the experiments were carefully cleaned before their use through the RCA treatment.³⁰

Preparation of samples for SANS/USANS experiments

Samples used for small angle scattering experiments were prepared starting from a stock solution, obtained by dissolving a suitable amount of SoyPC in pure chloroform, in order to achieve a concentration of $\cong 1$ mg ml⁻¹. Appropriate amounts of this mother solution were transferred into different round-bottom glass vials. Subsequently, suitable quantities of chloroform-based stock solutions containing the drug to be investigated were added, in order to have a pre-fixed molar ratio between the drug and phospholipid. The dissolution of the components was promoted by a slight warming ($\cong 40$ °C) and a very short sonication treatment ($\cong 5$ min). Afterwards, a thin film was

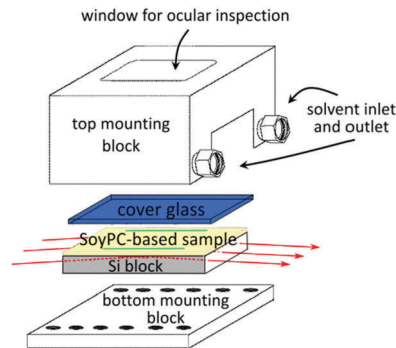


Fig. 1 Drawing of the custom sample cell used for NR and GISANS investigations. Two narrow stripes of cadmium (represented in green in this sketch) prevent the phospholipid-based sample from being touched by the cover glass. The latter allows the sample to be visually inspected at any time. Inlet and outlet plugs allow filling the space between the sample and cover glass with the solvent, whereas a washer (not shown) prevents the solvent leaking from the cell. The path of neutrons is indicated in red.

obtained for every sample, through slow evaporation of the chloroform in a stream of argon, to prevent lipid oxidation. The lipid films were hydrated, and all the resulting suspensions were vortexed and then gently sonicated ($\cong 30$ min). Subsequently, an aliquot of each suspension was repeatedly extruded through a polycarbonate membrane of 100 nm pore size, 11 times. The final (molal) concentration of SoyPC was in all cases 5.0 mmol kg⁻¹; drug concentrations were chosen accordingly to the pre-fixed molar ratio.

NR and GISANS experiments

NR and GISANS measurements were performed at MARIA, N-REX, and TREFF instruments, installed at MLZ (Maier-Leibnitz Zentrum), Garching, Germany. In an NR experiment, the specular reflectivity R is measured as a function of the scattering vector component Q_z along the direction z perpendicular to the interface: under the Born approximation, and in the limit of large Q_z values (*i.e.* low R values), the reflectivity is dependent on the Fourier transform of the scattering length density profile $\rho(z)$ along this direction:³¹

$$R = \frac{16\pi^2}{Q_z^2} \left| \int \rho(z) \exp(iQ_z z) dz \right|^2$$

The function $\rho(z)$, the scattering length density at the various z values, is dependent on the local composition and allows obtaining information on the lipid bilayer at the nanoscale.

Even if in NR experiments the scattering plane is perpendicular to the sample plane, measurements always provide data containing both the specular contribution and the diffuse scattering coming from $Q_x = 0$, where x represents the direction perpendicular both to z and to the detector plane. Anyway, the diffuse contribution is usually orders of magnitude smaller than the specularly reflected signal.³²

On the other hand, in a GISANS investigation, where a non-null component of the scattering vector along a direction perpendicular to z is present, information about eventual lateral correlations inside the sample is easily accessible.³³



At MARIA,³⁴ the wavelength of the neutrons was set at $\lambda = 10 \text{ \AA}$ for NR and at $\lambda = 5.0 \text{ \AA}$ for GISANS investigations, with a wavelength spread of $\Delta\lambda/\lambda \leq 0.10$ by means of a velocity selector. In the NR configuration, an elliptically curved focusing guide along the vertical direction allowed obtaining a beam size of $10 \times 0.1 \text{ mm}^2$; in the GISANS setup, a pinhole collimation setting was used, with a beam aperture of $16 \times 2 \text{ mm}^2$. The sample-to-detector distance was fixed at 1.91 m, whereas a rotating table allowed modulating the incident angle. This configuration allowed collecting data in the $3.3 \times 10^{-3} < Q_z/\text{\AA}^{-1} < 0.31$ range for NR and $0 < Q_z/\text{\AA}^{-1} < 0.25$; $-0.15 < Q_y/\text{\AA}^{-1} < 0.15$ for GISANS, with y being the direction perpendicular to z and parallel to the detector plane. Because of its high-flux and low-background characteristics, MARIA is able to detect reflectivity values down to 10^{-6} with relatively short time measurements ($\cong 1.5\text{--}2 \text{ h}$ per sample). In GISANS mode, the measurements were mostly performed below the critical angle $\theta^{(\text{cr})}$ of total internal reflection: in this case an evanescent wave propagating into the sample occurred, allowing collecting information over the complete surface region down to the depth of this wave.

At NREX,³⁵ a wavelength of $\lambda = 4.28 \text{ \AA}$ with a wavelength spread of $\Delta\lambda/\lambda \leq 0.03$ was used, by means of a HOPG crystal. A slit size of $50 \times 1.0 \text{ mm}^2$ was used for defining the incident beam, whereas scattered neutrons were detected by means of a ³He detector set at 2.46 m from the sample position. With this configuration, data in the $5.6 \times 10^{-3} < Q_z/\text{\AA}^{-1} < 0.26$ range were collected in NR mode and $0 < Q_z/\text{\AA}^{-1} < 0.12$; $0 < Q_y/\text{\AA}^{-1} < 0.10$ in the GISANS setup.

At the neutron reflectometer TREFF,³⁶ a wavelength of 4.7 \AA was used, by means of a graphite monochromator able to provide a wavelength resolution of $\Delta\lambda/\lambda \leq 0.03$. Specular intensities were recorded in $\theta/2\theta$ scan mode, with an incident beam of $0.4 \times 60 \text{ mm}^2$, covering the $9.1 \times 10^{-4} < Q_z/\text{\AA}^{-1} < 0.25$ range.

All the measurements were carried out at room temperature, using the custom made cell, as described in Fig. 1.

SANS/USANS experiments

SANS/USANS measurements were carried out by means of KWS-1 and KWS-3 diffractometers, at MLZ (Maier-Leibnitz Zentrum), Garching, Germany. Additional experiments were also performed with the TPA diffractometer, at LLB (Laboratoire Léon Brillouin), Saclay, France.

At KWS-1,³⁷ neutrons with an average wavelength of $\lambda = 5.0 \text{ \AA}$ and a wavelength spread $\Delta\lambda/\lambda \leq 0.10$ were used. A two-dimensional 128×128 pixel-based array scintillation detector set at three different collimation (C)/sample-to-detector (D) distances (namely C8/D2, C8/D8, and C20/D20, with all distances in meters) measured neutrons scattered from the samples. These configurations allowed collection of data in the scattering vector modulus ($Q = 4\pi/\lambda \sin(\theta/2)$) range between 0.0028 and 0.40 \AA^{-1} , with θ being the scattering angle. The investigated samples were contained in a closed quartz cell, in order to prevent the solvent evaporation, and were kept under measurement for a period so as to have at least 2 million counts of neutrons. The obtained raw data were corrected for background and empty cell scattering, and then radially averaged. Detector efficiency corrections and transformation to

absolute scattering cross sections were executed with a secondary plexiglass standard.³⁸

At KWS-3,³⁹ an incident neutron wavelength $\lambda = 12.8 \text{ \AA}$ (wavelength spread $\Delta\lambda/\lambda \leq 0.2$) was used, while the sample-to-detector distance was fixed at 9.6 m. KWS-3 uses a double focusing mirror principle to access to a very small angle region; a bidimensional detector with square pixels of 0.35 mm in size allowed collecting data ranging between 6×10^{-4} and $2 \times 10^{-3} \text{ \AA}^{-1}$.

At TPA,⁴⁰ neutrons with a wavelength of 12 \AA were used, with a wavelength spread $\Delta\lambda/\lambda \leq 0.11$. A multibeam design of the collimation system and a high resolution two dimensional image plate detector located at 6.27 m from the sample position allowed collecting data in the $4 \times 10^{-4} < Q/\text{\AA}^{-1} < 4.7 \times 10^{-3}$ range: data corrected for background and empty cell scattering were radially averaged and rescaled to the small-angle data collected at KWS-1. To this aim, a scale factor was evaluated in order to have the smallest sum of the squares of the differences of the two datasets in the range in which the overlap between the two instruments occurred.

All the measurements were carried out at room temperature.

Results

In Fig. 2(a) and (b), NR data are reported for SoyPC/benzocaine and SoyPC/propranolol systems, respectively. Inspection of data allows recognizing different behaviors shown by the two different drugs.

For systems containing benzocaine, the presence of two Bragg peaks at $Q_z^{(1)} \cong 0.1 \text{ \AA}^{-1}$ and $Q_z^{(2)} = 2 \cdot Q_z^{(1)}$ is clearly visible. This characteristic is also observed for the system without the drug, although the position of the peaks is shifted to lower Q_z values: according to the method used for lodging the SoyPC/drug mixtures on to the Si-based blocks, all these features may be explained by assuming the formation of a lamellar phase composed of the repetition of several bilayers on the support, separated by water layers. Two additional features are clearly visible by careful perusal of Fig. 2(a): (i) the presence of a weakly pronounced reflection at $Q_z^{(1/2)} \cong 0.05 \text{ \AA}^{-1}$ and (ii) the presence of two different critical angles highlighted by two different slopes shown by data. The first critical angle appears to be very small, *i.e.* smaller than or comparable to the minimum angle attained during reflectivity scans ($Q_z^{(\text{cr},1)} \cong 0.002 \text{ \AA}^{-1}$), whereas the second is located around $Q_z^{(\text{cr},2)} \cong 0.012 \text{ \AA}^{-1}$. Both these characteristics occur only in the presence of benzocaine, while they are absent for the pure SoyPC system.

Data for the above described systems were analyzed by means of the Parratt formalism,⁴¹ describing the scattering length density distribution, along the z direction, in terms of a stack of distinct layers. In this analysis, the Si substrate was modeled as an infinite-thick planar surface covered with a thin layer of oxide (SiO_2).⁴² The Parratt approach uses a transition matrix for each layer, able to correlate the amplitude of the neutron wave propagating in a certain layer with the behavior in the previous adjacent layer.⁴³ A set of matrices minimizing the sum of the squares of the differences between the experimental



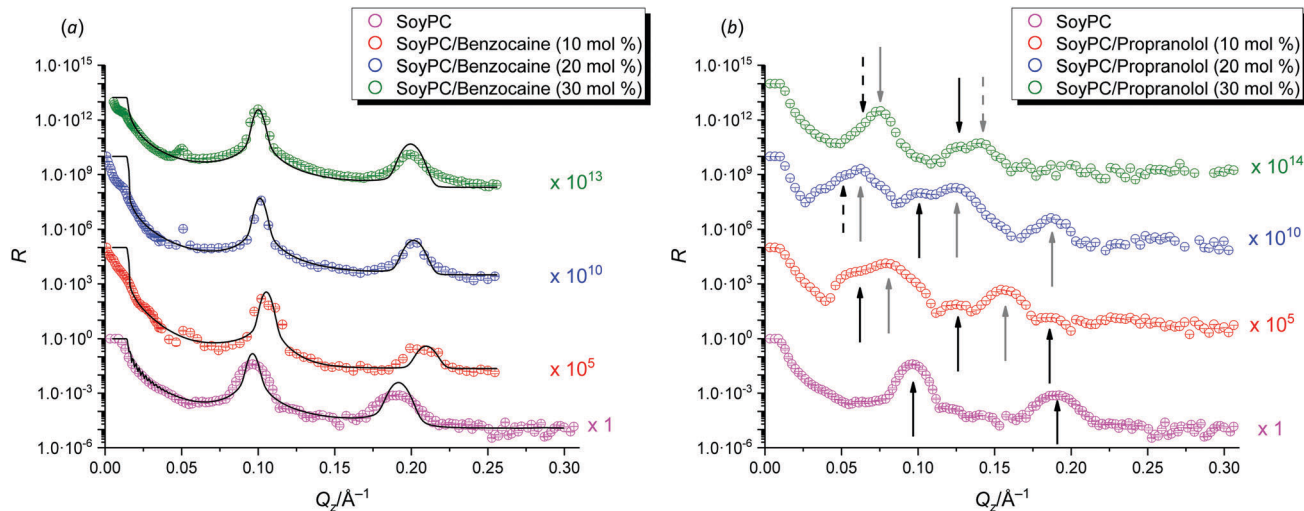


Fig. 2 NR data collected for (a) SoyPC/benzocaine and (b) SoyPC/propranolol systems in D_2O , according to the reported legends. To allow for a better visualization, data have been multiplied by a scale factor, as indicated. In plot (a) solid curves have been obtained by fitting the Parratt algorithm to the experimental data, as described in the text. In plot (b) Bragg reflections have been highlighted with arrows of different colors, according to their Q_z positions and connections (see the Discussion section). For an accurate determination of peak positions, a Gaussian function, or a sum of two Gaussian functions for partially overlapped peaks, has been fitted to the experimental data in proximity to the maxima, and reported as solid curves. Non-detected peaks or weak peaks whose positions deviate from that of a lamellar arrangement are indicated with dashed arrows (see also Table 4).

and the calculated reflectivities were found, characterizing the thickness, scattering length density, solvation (solvent volume fraction), and interfacial roughness of the different layers. These calculations were implemented in the MOTOFIT application:⁴⁴ all the parameters were varied until the optimum fit to the data was found. Although more than one set of parameters can very often be found for a given experimental dataset, the number of these possible sets was significantly reduced by a prior knowledge of the system, which allows imposing a range of values for each parameter to be optimized, as well as a certain number of constraints, avoiding the occurrence of meaningless values. For example, the two thicknesses belonging to the opposite hydrophilic portions of the bilayers were constrained to be equal to each other. Data obtained from the analysis of the systems are reported in Fig. 3.

Reflectivity data collected for SoyPC bilayers in the presence of propranolol show, on the other hand, a more complex behavior: multiple peaks with high broadness are visible. The positions of some reflections are, again, typical of a lamellar arrangement, but, differently from what has been seen for SoyPC/benzocaine systems, here the peaks appear broad and some of them overlap. This behavior is particularly apparent for the system with 10 mol% of propranolol: the most intense reflection is observed at $Q_z^{(1a)} \cong 0.0790 \text{ \AA}^{-1}$, along with two correlated peaks at $Q_z^{(2a)} \cong 0.158 \text{ \AA}^{-1}$ and $Q_z^{(3a)} \cong 0.235 \text{ \AA}^{-1}$. Besides these peaks, additional maxima are clearly present in the spectrum; in particular, a peak is visible at $Q_z^{(1b)} \cong 0.0618 \text{ \AA}^{-1}$, with its shoulder-like shape due to the high signal connected to the Bragg reflection at $Q_z^{(1a)}$ that partially masks its position. Other peaks most likely correlated with $Q_z^{(1b)}$ appear at $Q_z^{(2b)} \cong 0.124 \text{ \AA}^{-1}$ and at $Q_z^{(3b)} \cong 0.186 \text{ \AA}^{-1}$ (the latter being rather weak). These reflectivity data appear as the result of the presence of different structures, and are not susceptible of a simple quantitative analysis.

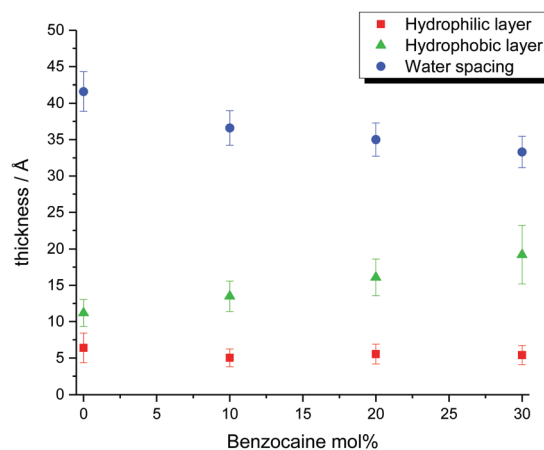


Fig. 3 Thicknesses of SoyPC-based bilayers in the presence of benzocaine, obtained through the Parratt algorithm, according to the reported legend.

As mentioned, the study of the interaction of the active ingredients with the bilayers was also carried out in the bulk, by means of SANS/USANS. To this aim, Fig. 4 and 5 show the scattering cross sections obtained for SoyPC/benzocaine and SoyPC/propranolol systems, including both suspensions subjected to simple sonication and suspensions extruded afterwards. Analogously to what has been seen for reflectivity data, a different trend of cross sections is observed for the two systems. Some differences also occur when extruded and simply sonicated systems are compared.

Sonicated suspensions in the presence of benzocaine, as well as the drug-free SoyPC suspension, show a narrow peak located at $Q \cong 0.1 \text{ \AA}^{-1}$, as also observed for NR data. In the low Q region, cross sections scale with a power law, *i.e.* $(d\Sigma/d\Omega) \propto Q^{-\alpha}$, where the slope α ranges between 2 and 4. On the other hand,



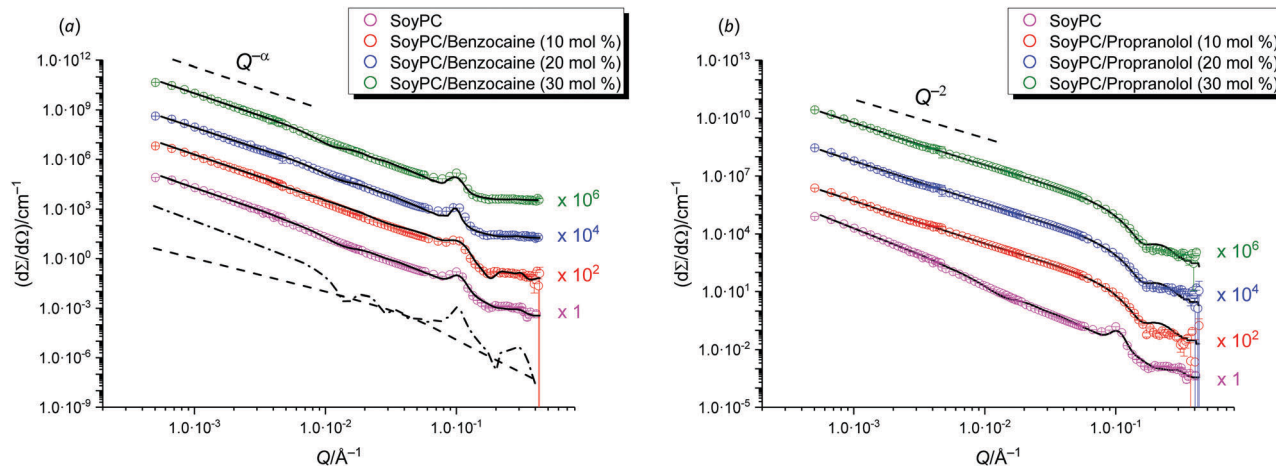


Fig. 4 Scattering cross sections obtained for (a) SoyPC/benzocaine and (b) SoyPC/propranolol systems prepared by sonication of hydrated suspensions in D_2O , according to the reported legends. To allow for a better visualization, data have been multiplied by a scale factor, as indicated. Solid curves have been obtained by fitting of the models described in the text to the experimental data. In plot (a), contributions to the total scattering cross sections coming from multilamellar liposomes (dashed-dotted line, multiplied by 0.01) and unilamellar liposomes (dashed line, multiplied by 0.001) have illustratively been reported for the SoyPC aqueous system.

for sonicated suspensions in the presence of propranolol, the Bragg peak observed for samples containing benzocaine is absent, and data at low Q scale with a power law $(d\Sigma/d\Omega) \propto Q^{-2}$. In all cases no Guinier regime is observed in the Q range covered by SANS/USANS investigations: this behavior is consistent with the opalescent appearance of the simple sonicated suspensions, indicating aggregates whose linear dimensions are larger than the typical wavelength of visible light ($\cong 500$ nm), although samples containing propranolol appeared less opalescent than benzocaine-based suspensions. All these features are typical of the presence of liposomes in all the suspensions: the presence of Bragg peaks and that of a power law with $2 < \alpha < 4$ are typical features of the existence of multilamellar liposomes, whereas for SoyPC/propranolol systems unilamellar liposomes are responsible for the scattering pattern detected. In all cases, the slope α of the power law observed at low Q values is connected to the lamellarity of the aggregates.⁴⁵

Scattering arising from multilamellar liposomes may be described assuming the liposomes as a collection of randomly oriented assemblies of lamellar stacks, neglecting their curvature, since the Guinier region is completely missing. For such structures, the theoretical cross section of a sample containing n_s randomly oriented stacks per unit volume has been obtained by Kotlarchyk and Ritzau:⁴⁶

$$\left. \frac{d\Sigma}{d\Omega} \right|_{MLV} = n_s \cdot N \cdot v^2 \cdot \Delta^2 \rho \times \frac{\langle |f(Q)|^2 \rangle}{Q^2} \left[1 + \frac{|f(Q)|^2}{\langle |f(Q)|^2 \rangle} (S(Q) - 1) \right] \quad (1)$$

where v indicates the volume of a single stack, $\Delta\rho$ is the contrast between the lipid bilayers and the solvent, $f(Q)$ is the form factor of a single bilayer, *i.e.* $f(Q) = \sin(Q\tau/2)/(Q\tau/2)$, and $S(Q)$ represents the structure factor taking into account the interferences occurring

among the bilayers belonging to a single stack. The structure factor depends in a complex way on the geometrical characteristics of the stack, namely the mean layer thickness τ , and the distance between the centers of two consecutive layers d . Both τ and d have been allowed to be polydisperse: the former with a Schulz-Zimm distribution function related to the polydispersity index i_τ , and the latter with a Gaussian distribution having a standard deviation σ_d . In particular, the σ_d/d ratio, named the Hosemann factor, has been estimated. To this aim, the brackets $\langle \rangle$ in eqn (1) refer to averages both over the distribution of layer thicknesses and over that of interlamellar distances.⁴⁶

Scattering coming from unilamellar liposomes may be described by a single bilayer form factor:

$$\left. \frac{d\Sigma}{d\Omega} \right|_{ULV} = n_s \cdot v^2 \cdot \Delta^2 \rho \cdot \frac{\langle |f(Q)|^2 \rangle}{Q^2} \quad (2)$$

Analysis of experimental cross sections collected for SoyPC/benzocaine has shown that the best description has been obtained by modeling the system as a mixture of multilamellar and a small amount of unilamellar vesicles, summing up eqn (1) and (2); on the other hand, for SoyPC/propranolol a description in terms of unilamellar vesicles has been proved to be suitable for the analysis. In all the fitting procedures, a constant term, $(d\Sigma/d\Omega)_{\text{incoh}}$, has been added, which represents the incoherent contribution to the cross section measured, mainly due to the presence of hydrogen atoms. Data obtained from the analysis of the systems are reported in Table 2.

When data for simply sonicated suspensions are compared with the corresponding data obtained for suspensions subjected to extrusion, substantial differences can be detected. For SoyPC/benzocaine, a typical trend ascribable to unilamellar liposomes is observed in the medium and high Q regions, *i.e.* $(d\Sigma/d\Omega) \propto Q^{-2}$ when $5 \times 10^{-3} < Q/\text{\AA}^{-1} < 7 \times 10^{-2}$, similar to what has been observed for sonicated SoyPC/propranolol



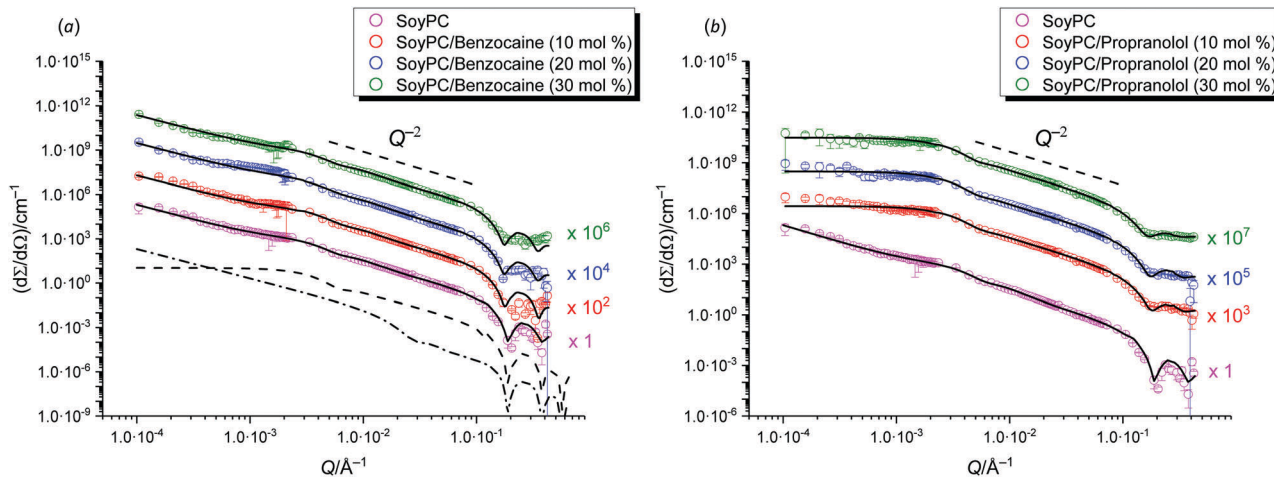


Fig. 5 Scattering cross sections obtained for (a) SoyPC/benzocaine and (b) SoyPC/propranolol systems prepared by extrusion of hydrated suspensions in D_2O , according to the reported legends. To allow for a better visualization, data have been multiplied by a scale factor, as indicated. Solid curves have been obtained by fitting of the models described in the text to the experimental data. In plot (a), contributions to the total scattering cross sections coming from unilamellar liposomes (dashed line, multiplied by 0.01) and multilamellar liposomes (dashed-dotted line, multiplied by 0.001) have illustratively been reported for the SoyPC aqueous system.

Table 2 Structural parameters obtained for sonicated aqueous suspensions of SoyPC/benzocaine and SoyPC/propranolol liposomes, through the analysis of SANS data, as described in the text. The table reports the slope α of the power law observed in the $4 \times 10^{-4} < Q/\text{\AA}^{-1} < 1 \times 10^{-2}$ range, the number N of lamellae in multilamellar liposomes, the average thickness τ of the bilayer together with its polydispersity index i_τ , the distance d between two consecutive bilayers and the Hosemann factor σ_d/d . In systems composed of both unilamellar and multilamellar liposomes, the thicknesses of the bilayers have been constrained to be identical to each other. The polydispersity index is defined as $i_\tau = \langle \tau^2 \rangle / \langle \tau \rangle^2$, where the brackets indicate averages weighted by the Schulz–Zimm distribution function

System	α	N	$\tau/\text{\AA}$	i_τ	$d/\text{\AA}$	$\frac{\sigma_d}{d}$
SoyPC	2.76 ± 0.04	7.3 ± 0.4	31.5 ± 2.5	1.16 ± 0.08	60.9 ± 1.4	0.11 ± 0.04
SoyPC/benzocaine (10 mol%)	2.64 ± 0.03	8.1 ± 0.3	31.8 ± 2.9	1.12 ± 0.07	61.5 ± 1.2	0.10 ± 0.03
SoyPC/benzocaine (20 mol%)	2.76 ± 0.04	7.5 ± 0.3	32.0 ± 2.7	1.22 ± 0.12	63.1 ± 1.5	0.10 ± 0.03
SoyPC/benzocaine (30 mol%)	2.72 ± 0.04	8.4 ± 0.3	32.2 ± 2.8	1.15 ± 0.09	62.9 ± 1.7	0.11 ± 0.03
SoyPC/propranolol (10 mol%)	2.04 ± 0.04	—	41.3 ± 1.2	1.06 ± 0.05	—	—
SoyPC/propranolol (20 mol%)	2.08 ± 0.05	—	40.8 ± 0.6	1.09 ± 0.07	—	—
SoyPC/propranolol (30 mol%)	1.97 ± 0.04	—	40.8 ± 0.4	1.11 ± 0.07	—	—

systems. Anyway, differently from the latter case, data at low Q for SoyPC/benzocaine suspensions reveal an excess of scattering that quantitative analysis shows to be attributable to a residual presence of multilamellar liposomes, although in this case their fraction is low.

Finally, SoyPC/propranolol extruded suspensions show the unique presence of unilamellar liposomes that, differently from what has been seen for the corresponding simply sonicated systems, are unquestionably smaller, with the Guinier regime being clearly visible when $Q < 2 \times 10^{-3} \text{\AA}^{-1}$. All data for the extruded systems are reported in Table 3: quantitative analysis for such systems is similar in many respects to the corresponding description adopted for sonicated suspensions, with the main difference being the model for describing the scattering coming from unilamellar liposomes. In this case a form factor of a hollow sphere with a polydisperse aqueous core has been adopted. This model is justified by the presence of the Guinier regime for unilamellar vesicles at very low Q values, weakly detectable for SoyPC/benzocaine suspensions (because of the scattering coming from multilamellar vesicles) and clearly shown for SoyPC/propranolol systems.

Discussion

In the present work we focused on the study of the perturbation effects produced on the lipid bilayer by molecules used as active ingredients for some treatment, trying to correlate such effects with the drug molecular structure. Generally speaking, the perturbation effects may depend on the hydrophobicity/hydrophilicity degree of the active ingredient, on its molecular shape and on possible attractive or repulsive interactions (e.g. electrostatic) that can be established: all these factors contribute in a complex way to (eventually) perturb the phospholipid packing. To this aim, the concept of the packing factor, introduced by Israelachvili, Mitchell, and Ninham,⁴⁷ has been proved to be useful, because of its capability to explain, justify and very often predict the molecular self-assembly behavior in surfactant solutions. The packing factor N_s is defined as

$$N_s = \frac{v}{l \cdot a_0} \quad (3)$$

where v is the volume of the total surfactant hydrophobic portion, l represents the tail length, and a_0 is commonly defined as the



Table 3 Structural parameters obtained for extruded aqueous suspensions of SoyPC/benzocaine and SoyPC/propranolol liposomes, through the analysis of SANS data, as described in the text. The table reports the aqueous core radius R_{core} of unilamellar liposomes together with its polydispersity index $i_{R_{\text{core}}}$, and the thickness τ of the bilayer. The polydispersity index $i_{R_{\text{core}}}$ is defined as $i_{R_{\text{core}}} = \langle R_{\text{core}}^2 \rangle / \langle R_{\text{core}} \rangle^2$, where the brackets indicate averages weighted by the Schulz–Zimm distribution function. For systems in which the coexistence of unilamellar liposomes with multilamellar liposomes occurs, the number N of lamellae, the distance d between two consecutive bilayers and the Hosemann factor σ_d/d are also reported. In order to reduce the number of fitting parameters, in systems composed of both unilamellar and multilamellar liposomes, the thicknesses of the bilayers have been constrained to be identical. Furthermore, d and σ_d/d have been kept fixed and equal to the value averaged for the corresponding sonicated systems

System	$R_{\text{core}}/\text{\AA}$	$i_{R_{\text{core}}}$	$\tau/\text{\AA}$	N	$d/\text{\AA}$	$\frac{\sigma_d}{d}$
SoyPC	407 ± 6	1.09 ± 0.05	33.0 ± 1.2	6 ± 2	62.1	0.10
SoyPC/benzocaine (10 mol%)	398 ± 6	1.09 ± 0.06	35.9 ± 1.5	4.4 ± 1.5	62.1	0.10
SoyPC/benzocaine (20 mol%)	395 ± 7	1.09 ± 0.05	36.3 ± 1.4	3.5 ± 1.5	62.1	0.10
SoyPC/benzocaine (30 mol%)	408 ± 6	1.09 ± 0.04	36.3 ± 1.5	3.5 ± 1.7	62.1	0.10
SoyPC/propranolol (10 mol%)	471 ± 7	1.12 ± 0.05	36.0 ± 1.8	—	—	—
SoyPC/propranolol (20 mol%)	521 ± 7	1.10 ± 0.05	37.0 ± 1.9	—	—	—
SoyPC/propranolol (30 mol%)	529 ± 7	1.09 ± 0.04	35 ± 2	—	—	—

equilibrium area per molecule at the aggregate surface. Nagarajan⁴⁸ has stressed that the latter parameter has to be considered not as a simple geometrical variable connected to the size and shape of the polar heads, rather as a thermodynamic quantity obtained from a minimum free energy approach. For the following analysis, it has to be noted that the hydrophobic portion of the SoyPC bilayer was assumed to be in the liquid crystalline phase, as the main lipid phase transition (gel–sol, $P_{\beta'}$ → L_{α}) temperature (< -40 °C) is well below the room temperature.

Quantitative analysis of membrane characteristics was performed for SoyPC/benzocaine systems. It showed that SoyPC is organized on the Si substrate as a stack of bilayers, formed by their repetition on the support, separated by layers of water. Besides these repeating layers, a thin ($\cong 15$ Å) water layer was found to be directly stratified on the SiO₂ layer ($\cong 2$ nm), covering, in turn, the Si block. In all cases, the roughness values of the interfaces ranged between 2 Å and 7 Å. Extensions of the hydrophobic and hydrophilic portions of the bilayers, as well as the dimensions of the water spacing, are plotted in Fig. 3 for the drug concentrations investigated. Inspection of data reveals a certain effect on the bilayer caused by the presence of the drug: inclusion of benzocaine reduces the repetition distance among the bilayers, but not in a monotonic fashion. For the pure SoyPC system, this distance is found to be $\cong 2\pi/Q_2^{(1)} = 65.2$ Å, which reaches its minimum value for 10 mol% of the drug (59.8 Å) and slightly increases again for 20 mol% and 30 mol% (61.9 Å and 62.7 Å, respectively). Although these values are independent of any theoretical model, analysis of Fig. 3, based on the Parratt algorithm, shows that this trend is the outcome of two different tendencies: (i) the water spacing that slightly decreases with increase of the drug content (even taking into account the uncertainty affecting the data) and (ii) the thickness of the hydrophobic portion of the bilayer, which is progressively enlarged as soon as the drug amount is increased. Concerning the latter parameter, it is interesting to note that the C1–C18 distance measured in the linoleic acid (the precursor of the main species in SoyPC) is 19.2 Å.⁴⁹ Although this value refers to the crystallographic distance, and although the average thickness of the hydrophobic portion of the bilayers seldom exceeds 1.8 times

the fully extended length of the hydrocarbon chains,⁵⁰ it is nevertheless still higher than the experimental thicknesses found, ranging from 11.2 to 19.2 Å – a symptom of an interdigitation or tilting of the hydrophobic chains. Generally speaking, there are manifold circumstances where interdigitation phenomena are observed. The most important are the asymmetry of the chains forming the phospholipid tail (*i.e.* one chain has more than 5–6 methylene groups with respect to the other one),⁵¹ the application of high pressures,^{52,53} and the solubilization of molecules able to promote or increase intermolecular lateral repulsions.^{54,55} In particular, the last point is what we are concerned about, since solubilized species inside the bilayer may increase (or mitigate) lateral repulsions, enhancing (or reducing) the interdigitation/tilting phenomena. It has been argued⁵⁶ that the repulsions occurring in the interfacial region of the bilayer initially promote a progressive tilting of the lipid chains, for adapting the chains to the large interfacial molecular area. Increasing the tilting angle increases, in turn, the torsional and angular tensions in the lipid chain: beyond a certain threshold, the tilt cannot be further increased, and hydrocarbon interdigitation occurs.

Analysis of the hydrophobic layer thickness shows, indeed, that the insertion of benzocaine reduces the above described phenomena, as an increase of the thickness is observed with increasing content of the drug. The presence of two *cis* double bonds in the hydrocarbon backbone introduces kinks, preventing efficient packing of the hydrophobic chains in the bilayer, as can be seen in Fig. 6.⁵⁷ It can be argued that the insertion of benzocaine is able to promote a better packing of the chains, being capable of “filling” the space that is created in consequence of the reduction of chain tilting/interdigitation. In any case, it has to be taken into account that the final packing is a compromise between the reduction (or increase) of repulsive interactions between the hydrophobic chains and a possible reduction (or increase) of repulsive interactions between the polar heads, as will be glimpsed in the following. The thickness of the hydrophilic portion is roughly constant in the concentration range studied: benzocaine has a small hydrophilic portion represented by the amine functional group and its insertion does not dramatically modify this part of the bilayer.



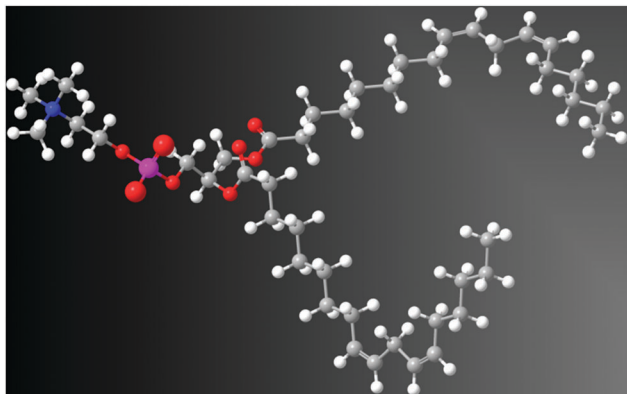


Fig. 6 “Ball and stick” representation of the 1,2-dilinoleoyl-*sn*-glycero-3-phosphocholine molecule (main component of SoyPC) in the staggered conformation.

Finally, it is interesting to discuss and rationalize the presence of two other features appearing in SoyPC/benzocaine reflectivity data, shortly mentioned in the Results section: namely the presence of two possible critical angles and the appearance of weak reflections at $Q_z^{(1/2)} \cong 0.05 \text{ \AA}^{-1}$, both of these features occurring only in the presence of the drug.

There are two distinct sudden changes in slope in data trends, for systems containing benzocaine: the first is close to the minimum angle reached during investigations, namely $Q_z^{(cr,1)} \cong 0.002 \text{ \AA}^{-1}$, whereas the second is located around $Q_z^{(cr,2)} \cong 0.012 \text{ \AA}^{-1}$. Their presence might be ascribed to the different contrast $\Delta\rho = \rho_{inf} - \rho_{Si}$ existing between the “final” (infinite) layer and the “initial” Si layer: the contrast value determines the critical scattering vector modulus $Q_z^{(cr)}$ at which total reflection occurs:

$$Q_z^{(cr)} = \frac{4\pi}{\lambda} \sin \theta^{(cr)} \cong 4\sqrt{\pi \cdot \Delta\rho} \quad (4)$$

The presence of macroscopically thick patches of bilayer material with a scarce penetration of D_2O among the bilayers may rationalize the results observed: these patches, for which D_2O is not “seen” as the final layer (for the purposes of the critical angle determination), coexist with normally hydrated portions of the bilayers. It has to be assumed that the sizes of these “islands” of scarcely hydrated material are bigger than the coherence volume of the incident beam, as the two angles would not be visible otherwise. Application of eqn (4) for $\rho_{inf} = \rho_{D_2O}$ gives a value (0.0146 \AA^{-1}) very close to the observed $Q_z^{(cr,2)}$. On the other hand, a reliable estimation of $\Delta\rho$ for the poorly hydrated phospholipid stacks is hardly possible because of the unknown penetration degree of water; it can only be argued that it must be smaller than the corresponding difference found for water, which is in qualitative agreement with the experimental observation that $Q_1^{(cr)} < Q_2^{(cr)}$ (as can also be seen from Table 1). A similar phenomenon has been described for sodium poly-(styrenesulfonate) based films on silicon, where two critical angles arising from the silicon/film and silicon/air interfaces were observed in X-ray reflectivity investigations.⁵⁸

A possible nanoscopic interpretation for the presence of patches in SoyPC/benzocaine systems might be the partial suppression of bilayer fluctuations. Fluctuations promote the detachment of the bilayers and the subsequent penetration of the solvent among the layers. The presence of benzocaine reduces the strength of these fluctuations, from which we may infer a higher rigidity of the bilayer. On the other hand, in the SoyPC and SoyPC/propranolol systems, bilayers fluctuate much more, showing the typical critical angle of the Si/ D_2O interface. This hypothesis may be proven through Grazing Incidence Neutron Spin-Echo Spectroscopy (GINSES) investigations. At present, the low counting rate is the main limitation for such a technique, because of the limited scattering volume that comprises only the volume covered by the evanescent wave. However, improvements for dealing with this drawback are currently under development.⁵⁹

Most interesting is the appearance, for all the systems in the presence of benzocaine, of a small maximum at $Q_z^{(1/2)} \cong 0.05 \text{ \AA}^{-1}$, besides the first and second order Bragg peaks arising from the multilayer structure. The main point is represented by the position of this reflection that is, for all the concentrations investigated and within the experimental uncertainty, half the value of the first Bragg peak, *i.e.* $Q_z^{(1/2)} = Q_z^{(1)}/2$. A rational justification of such behavior is a local formation, in the bilayers, of stalks in the lamellar arrangement, which will then result in twice the typical repetition distance, resulting in turn in the appearance of a peak at half the value of the first Bragg peak (as sketched in Fig. 7). The formation of stalks, which are real crystal-like defects in the lipid arrangement, has to be ascribed to the capability of benzocaine to induce localized structures with a negative curvature of the bilayer membrane: such bending can only be the result of a global increase of the packing factor N_s with respect to the value assumed in pure SoyPC bilayers. Actually, it has already been seen that addition of benzocaine promotes a packing of hydrophobic chains that progressively resembles that existing in the crystal. The existence of stalks suggests that, at the same time, less repulsive interactions among the phospholipid heads occur. In turn, these interactions affect the global value of a_0 , *i.e.* the molecular area at the aggregate surface. A simple analysis carried out by Tanford^{48,60} shows that

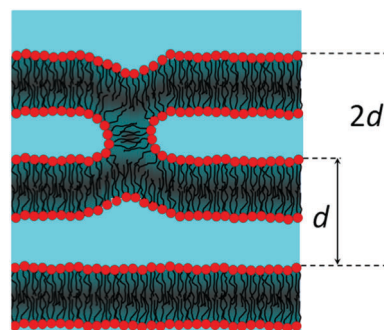


Fig. 7 Sketch of stalk formation induced by the addition of benzocaine (not explicitly shown) to SoyPC bilayers. This phenomenon gives rise to twice the typical repetition distance (e.g. the distance between the water layers), resulting in the appearance of a reflection peak at half the Q_z value of the primary Bragg peak.



a_0 is dependent on a headgroup repulsion parameter (β) and on the interfacial free energy (σ): $a_0 = \sqrt{\beta/\sigma}$. A decrease of β results in an increase of the packing factor N_s , favoring, after all, the formation of morphologies with negative curvatures: the better packing of the hydrophobic chains, which tends to decrease N_s , is overcompensated by the improved interactions among the polar heads, leading to a global increase of N_s . In any case, it has to be highlighted that these structures are formed in a low content, due to the lower intensities of the associated Bragg peaks.

It has often been supposed that the formation of such defects is the prelude to a transition to non-lamellar phases.⁶¹ Indeed, the formation of hexagonal and tetragonal lattices has been found⁶² for the investigation of phospholipid-based membranes in the presence of the ibuprofen active ingredient, which is connected to the presence of similar stalks in the double layer arrangement. With the aim of investigating a possible similar situation in our system, selected GISANS scans have been performed – most of them below the critical angle of total internal reflection of the Si/D₂O interface. In this case, eventual lateral correlations are probed inside the bilayers, up to the depth A_{evw} of the evanescent wave given by

$$A_{\text{evw}} \cong \frac{1}{\text{Re} \sqrt{4\pi\Delta\rho \left[1 - \left(\frac{\theta_i}{\theta(\text{cr})} \right)^2 \right]}}$$

Due to the high integral counting needed for GISANS investigations, long measurement times had to be chosen, so that the analysis was limited to the 30 mol% benzocaine based sample. Such analysis was carried out at several different incident angles θ_i ranging from 0.020° to 0.60° , covering the region where the two critical angles were observed in NR scans, with the aim of gaining depth resolved results. 2-D detector illustrative images at selected incoming angles θ_i of 0.099° , 0.14° , 0.24° , and 0.60° are displayed in Fig. 8(a)–(d), respectively. In this range of incident angles, the biggest change in scattering distribution occurred.

Analysis of images reveals that starting from low angles a spot located at $Q_z \cong 0.1 \text{ \AA}^{-1}$ and corresponding to the Bragg peak arising from the lamellar arrangement of bilayers appears.

Its intensity increases as θ_i increases, since the penetration depth of the evanescent wave is progressively enlarged, so that a higher scattering of the lamellar bilayer structure is observed. Although not appreciable when $\theta_i < 0.1^\circ$, a Debye–Scherrer ring starts to appear at $\theta_i = 0.14^\circ$, which is indicative of an increasing amount of disordered lamellae, *i.e.* bilayers having an increased curviness with respect to the ideal situation corresponding to a parallel alignment of the phospholipid membranes. However, it has to be noted that the Debye–Scherrer ring does not stretch over the whole detector, suggesting that the disordering of the lamellar areas is far from being complete but rather deviates by only a few degrees from a perfect parallel ordering. Interestingly, no reflections arising from crystalline structures inside the double layer are detected, suggesting that the formed stalks are isolated, *i.e.* are not distributed according to the existence of a lattice. Based on the NR/GISANS results, we may point out that the presence of stalks inside the bilayer arrangement, the incomplete Si surface coverage with fully hydrated bilayers, and the imperfect parallel alignment of the bilayer with respect to the Si surface are the main sources for the non-perfect description of data by the Parratt model, *i.e.* the non-exact fitting of the curves in Fig. 2(a).

Much more complicated appear to be the NR data obtained for the SoyPC/propranolol system. As mentioned in the Results section, the inclusion of propranolol drastically changes reflectivity trends: analysis of data reported in Fig. 2(b) reveals, differently from what has been seen for the systems in the presence of benzocaine, the existence of multiple peaks. These peaks are broad and some of them overlap partially, compelling to describe the system in a qualitative fashion. It could be reasonable to analyze the positions of the peaks in terms of the possible existence of lamellae, as already seen for SoyPC/benzocaine systems and as normally observed for systems prepared as described in the Experimental section.⁶² To this aim, peaks appearing in NR data are highlighted in Fig. 2(b) by regrouping their Q_z values with arrows having the same color; positions found are reported in Table 4, according to the grouping operated. In all cases no more than two sets have been found. Analysis of the positions found for the reflections

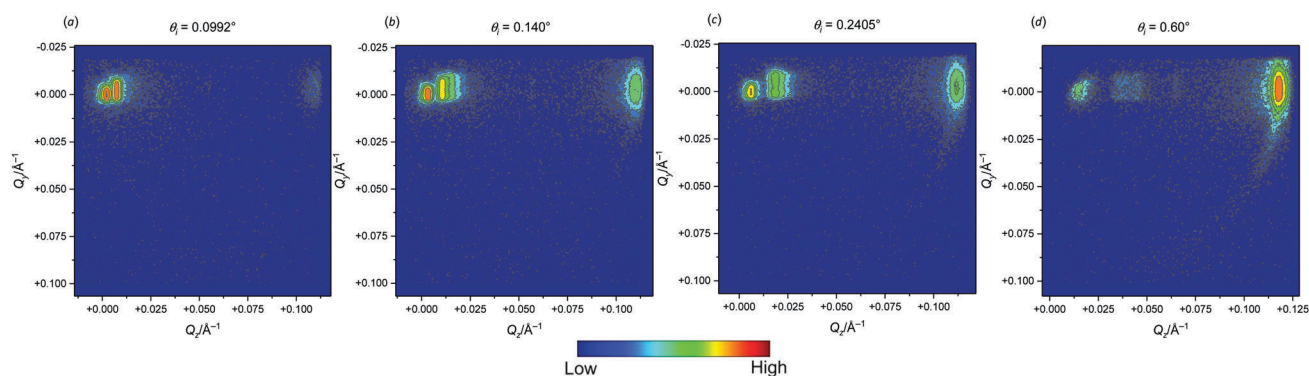


Fig. 8 GISANS images obtained at a wavelength $\lambda = 4.28 \text{ \AA}$ for the SoyPC/benzocaine 30 mol% system at incoming angles (θ_i) of (a) 0.10° , (b) 0.14° , (c) 0.24° , and (d) 0.60° . With increasing angle the lamellar peak at $Q_z = 0.1 \text{ \AA}^{-1}$ appears, and with further increasing angle a Debye–Scherrer ring of a distorted lamellar structure becomes visible. Intensities have been reported on a logarithmic scale, according to the color legend shown.



Table 4 Peak positions determined in NR data for SoyPC/propranolol systems. These peaks are highlighted in Fig. 2(b) with arrows of different colors, according to their possible associations. Some of the peaks appear as a shoulder being partially masked by other reflections: in this case a sum of two Gaussian functions in the region around the maxima has been fitted to the experimental data for evaluating the positions of the peaks. Most of the third reflections are very weak and their positions, although mentioned in the text, cannot be accurately determined and have not been reported in the table. Nondetected peaks or weak peaks whose positions deviate from that of a lamellar arrangement are followed by a question mark and correspond to dashed arrows in Fig. 2(b). All the values found are reported with three significant digits, according to the accuracy of the determination

System	$Q_z^{(1)}/\text{\AA}^{-1}$	$Q_z^{(2)}/\text{\AA}^{-1}$	$Q_z^{(3)}/\text{\AA}^{-1}$	$\frac{Q_z^{(2)}}{Q_z^{(1)}}$	$\frac{Q_z^{(3)}}{Q_z^{(1)}}$	$\frac{d}{\text{\AA}} = \frac{2\pi}{Q_z^{(1)}/\text{\AA}}$
SoyPC	0.0964	0.191		1.98		65.2
SoyPC/propranolol (10 mol%)	0.0790	0.158		2.00		79.5
	0.0618	0.124	0.187	2.01	2.99	102
SoyPC/propranolol (20 mol%)	0.0625	0.125	0.187	2.00	2.99	101
	0.0466 (?)	0.101		2.17		135
SoyPC/propranolol (30 mol%)	0.0752	0.143 (?)		1.90		83.6
	?	0.127				

shows, indeed, quite a complex trend. For the SoyPC system, as already described above, the positions of the Bragg peaks are easily described, both qualitatively and quantitatively, in terms of a (fairly ordered) lamellar phase composed of the repetition of several bilayers on the support. On the other hand, for SoyPC/propranolol systems, even at 10% mol of the drug, the data show a richer behavior. For this lowest amount, five maxima are clearly visible: analysis of their positions shows that they are correlated in the 1:2 Q -ratio (or 1:2:3), with the second set (the one located at higher Q and highlighted with grey arrows) being clearly visible, whereas the primary reflection of the first set (indicated with a black arrow) appears as a shoulder because of the high signal connected to the primary peak of the second pair. Trends become complicated for higher amounts of propranolol. At 20% mol, one peak series is clearly visible (up to the third reflection) having the primary maximum so high and broad to make the detection of the primary Bragg peak at smaller Q hardly detectable. At 30% mol, the latter peak is even not visible (with its expected position being indicated with the black dashed arrow). Analysis of the positions of the maxima indicates two repetition distances for 10% and 20% mol propranolol of 80 Å and 102 Å, and 101 Å and 125 Å, respectively. Furthermore, inspection of GISANS 2-D detector images (Fig. 9) reveals the appearance of diffuse

Debye–Scherrer rings around the reflections, stronger than the corresponding rings observed for systems in the presence of benzocaine.

The indications found can be ascribed to the presence, even at the lowest amount of propranolol investigated, of a powder of lamellar regions in the scattering volume. Likely, the presence of a condensed ring in the molecular structure of propranolol and the contemporaneous charge existing at physiological pH induce strong disorder in the lipid bilayer arrangement: the polycyclic moiety has less conformational freedom than the single aromatic group present in the benzocaine structure. Consequently, propranolol has a smaller or nearly null capability to act as “filler” and is unable to efficiently occupy the space created in consequence of the reduction of tilting/interdigitation of SoyPC hydrophobic chains. Furthermore, differently from what has been seen for SoyPC/benzocaine systems, the charge existing in the hydrophilic region increases the repulsive interactions in the hydrophilic region, promoting the formation of highly curved structures. These two factors may result in the formation of lamellar powders, as schematically displayed in Fig. 10.

In sum, in the presence of benzocaine only a very limited local disorder of lamellar arrangement (along with the formation of negatively curved structures inside the lamellar phase) occurred, so that the NR data were still describable in

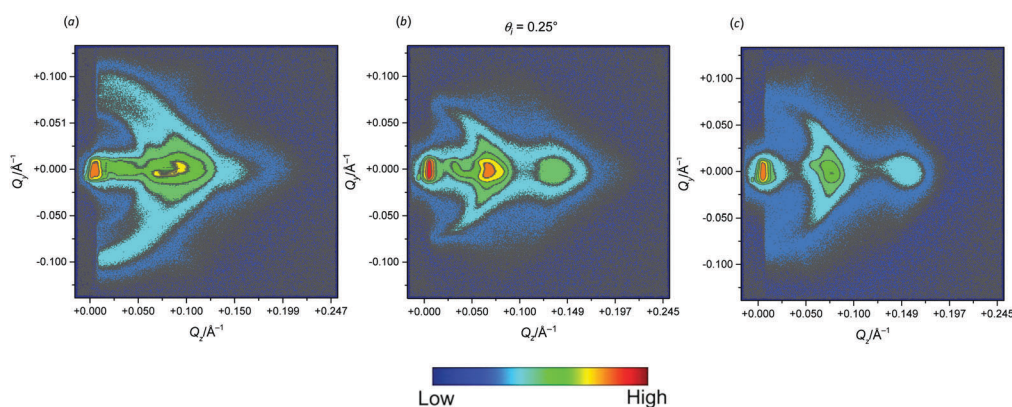


Fig. 9 GISANS images obtained at a wavelength $\lambda = 5.0$ Å for the SoyPC/propranolol system at an incoming angle (θ_i) of 0.25° at (a) 10 mol%, (b) 20 mol%, and (c) 30 mol% of the drug. Intensities have been reported on a logarithmic scale, according to the color legend shown.



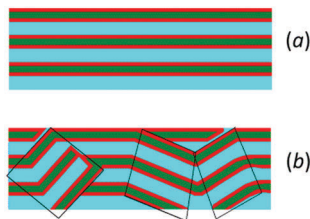


Fig. 10 Sketch of the structural evolution induced by the addition of propranolol. (a) In the absence of the drug, SoyPC exhibits a lamellar structure. (b) The introduction of propranolol induces disordering of lamellar areas, causing the presence of powder scattering of lamellae. These effects are due to the capability of the drug to induce curved bilayers. In the drawing, disordered lamellae are highlighted by black lines, for making their visualization easier.

terms of the Parratt model. On the other hand, the presence of propranolol has a much stronger effect that may not be interpreted as a simple deviation from the perfect parallel ordering, as also highly evidenced by NR data.

It is interesting to compare the strong perturbative effects on the bilayer arrangement due to propranolol addition, with the results found by Pereira-Leite and collaborators.⁶³ The authors investigated the interaction of propranolol with bilayers composed of egg phosphatidylcholine (EPC) and 1,2-dipalmitoyl-*sn*-glycero-3-phosphocholine (DPPC). In particular, they observed a decrease of the gel-sol transition temperature due to the presence of the drug, as well as a reduction of the order parameter associated with the transition. These pieces of evidence are consistent with the results found in the presence of SoyPC: a stronger disorder caused by the presence of propranolol is connected to a less efficient packing of the double layer that results in a gel-sol transition temperature smaller than the one corresponding to the pure phospholipid. The behavior observed for propranolol is also found for other drugs, although with different characteristics.⁶⁴

The results obtained from the NR/GISANS analysis of SoyPC bilayers in the presence of benzocaine and propranolol drugs allow foreseeing what should be the characteristics of such bilayers under bulk conditions, *i.e.* in water dispersions: under these conditions liposomes are formed. Unless a certain amount of mechanical energy is transferred to liposomes during their preparation (*e.g.* in the case of extrusion of aqueous suspensions), the size and characteristics of a population of liposomes are the result of a complex balance between the entropy of mixing⁶⁵ and bilayer curvature elasticity.^{66–68} In the limit of large liposomes, with dimensions much larger than the surfactant size, the curvature energy per unit area of bilayer f_c is given by⁶⁵

$$f_c = 2K \left(\frac{c_1 + c_2}{2} - c_s \right)^2 + \bar{K} c_1 c_2 \quad (5)$$

with K and \bar{K} being the bending moduli, depending, in turn, on the elastic constants determined by the head-head and tail-tail interactions. Finally, c_1 and c_2 indicate the two local curvatures, whereas c_s represents the spontaneous (*i.e.* minimum-energy) radius of curvature. In the case of a bilayer with relatively high curvature, as appears to be the case for SoyPC/propranolol systems, the energy required for hypothetically adding a further

bilayer to the liposome (*i.e.* for increasing the lamellarity) can overcome the attractive interaction between bilayers: in this case, a limited number of lamellae associated with a single liposome may be observed. In the most extreme cases, unilamellar liposomes are exhibited. Observations from NR and GISANS data could suggest the formation of oligolamellar or even unilamellar liposomes for aqueous suspensions prepared in the presence of propranolol, whereas a higher number of lamellae is expected when benzocaine is used.

Analysis of SANS trends shows, indeed, that this is the case: unilamellar liposomes are exclusively observed in the SoyPC/propranolol system, independently of the preparation method adopted (simple sonication or extrusion), whereas multilamellar liposomes are observed for SoyPC/benzocaine sonicated suspensions. The results obtained through the fitting of theoretical models described in the Results section are reported in Tables 2 and 3. For SoyPC/benzocaine systems, multilamellar liposomes with an average lamellarity of $N \cong 7$ are observed: these liposomes coexist with a small fraction of unilamellar structures. Although unilamellar liposomes would be expected to coexist with liposomes having a smaller lamellarity ($2 \leq N \leq 5$), it is important to note that in the adopted model, where no polydispersity for N is considered, the number of layers is generally only approximately estimated; it determines the upturn in scattering at low Q values where the total thickness of the stack is seen.^{69,70} The essential bilayer characteristics obtained from the analysis of NR data are retained for liposomes studied in an aqueous environment by means of SANS investigations. The values of bilayer thickness τ agree, as a rule, to the corresponding data obtained from NR data, although they are slightly overestimated: small discrepancies are normally present depending on the approximation degree of the model adopted for the fitting analysis. In the present case, for example, a homogeneous SLD profile has been assumed for the bilayer. Moreover, the strong Bragg peaks observed give a repetition distance of the bilayer arrangement very close to the values extracted through NR analysis.

The high curvature of bilayers produces, in SoyPC/propranolol systems, only unilamellar liposomes, even without any extrusion process, see Fig. 4(b). Ocular inspections of suspensions have shown a lower turbidity when compared to the corresponding SoyPC/benzocaine samples, suggesting the presence of smaller liposomes. Anyway, their linear size is still too large to make the Guinier region fall in the Q range covered by SANS/USANS measurements. Bilayer thicknesses appear to be comparable to the corresponding values obtained for benzocaine-based dispersions.

Analysis of the systems subjected to the extrusion of the prepared suspensions shows the presence of bilayers having lateral dimensions similar to those observed for simply sonicated systems. On the other hand, the characteristics of SoyPC-based liposomes are, at large scales, mainly influenced by the preparation method adopted: large liposomes (having radius $R \gg 500$ nm) occur when the phospholipid suspensions are only sonicated, whereas relatively small liposomes ($R \cong 400$ nm) are produced if the aqueous suspensions are also subjected to extrusion. Generally speaking, in the presence of extrusion



processes, vesicle dimensions are often (but not always) grossly determined by the pore size of the membrane used (100 nm in the present preparation). Exceptions to this typical rule are observed when elastic deformations of the liposomes occur during their crossing of the pores of the membrane used in the extrusion process.⁷¹

Also the liposome lamellarity and the size polydispersity are influenced by the extrusion process: for the extruded SoyPC/benzocaine systems, the partial tendency of the liposomes to form unilamellar structures has been observed, as proved by SANS data, even if a small residue of the scattering still arises from multilamellar structures. The extrusion process is considered to be a reliable methodology to produce unilamellar vesicles,⁷² even if in some cases, because of rearrangement phenomena, it has also been found to produce multilamellar aggregates,⁷³ or even more complex morphologies.⁷⁴ The general trend is here fulfilled, even if the small fraction of multilamellar aggregates suggests that a higher amount of energy, *i.e.* a higher number of extrusion steps, has to be provided for a further or total reduction of the multilamellar aggregates.

Finally, it is worthy of note that although the prepared liposomes, analogously to most of the natural phospholipids, are found to be metastable, their lifetime (which is of the order of months, at least) does not have any appreciable effect on the timescale covered by the investigations carried out.

Conclusions

In this paper, the influence of benzocaine and propranolol active ingredients on SoyPC-based bilayers has been investigated. Despite their small and simple chemical structures, marked and different effects of these two drugs have been found: on one hand, benzocaine induces a locally highly negative bending of the bilayer membrane, a result of a better packing of phospholipid hydrophobic chains and a closer distance between polar adjacent heads. Over large length scales, only a slight disorder from a perfect parallel ordering is observed.

On the other hand, propranolol, because of its structure and of the presence of charged heads (occurring under physiological conditions), promotes the formation of highly curved bilayers. This has been observed both in NR/GISANS investigations, where a high disorder typical of the formation of lamellar powders exists, and in SANS experiments, where only unilamellar liposomes have been observed.

It is important to recognize that the drug concentrations investigated in the present study are beyond the dosage in any medical posology. Nonetheless, it may be argued that a correlation exists between the effects observed at the expense of the bilayers and the complications due to high drug dosages, where a local excess of active ingredient is in all cases present. A possible structural modification of the bilayer may nevertheless cause a change in the mobility of membrane proteins, reducing or even inhibiting their biological function, giving rise to possible serious cell damages.⁷⁵

A further point to be reaffirmed concerns the necessity, shown by results obtained, to check for eventual interactions between the active ingredients and the phospholipid-based bilayer. The methodology adopted here combining NR, GISANS, and SANS/USANS has been proven to be helpful for characterizing, at the nanoscale, the main features of the bilayer. Integration with GINSES is desirable to integrate the structural information with the bilayer dynamics.

Conflicts of interest

There are no conflicts to declare.

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Notes and references

- 1 G. M. Cooper and R. E. Hausman, *The Cell: A Molecular Approach*, Sinauer Associates, 2016.
- 2 B. Alberts, A. Johnson, J. Lewis, D. Morgan, M. Raff, K. Roberts and P. Walter, *Molecular Biology of the Cell, Sixth Edition*, Taylor & Francis Group, 2014.
- 3 W. Lim, B. Mayer and T. Pawson, *Cell Signaling: principles and mechanisms*, Taylor & Francis Group, 2014.
- 4 M. A. Yildirim, K. I. Goh, M. E. Cusick, A. L. Barabasi and M. Vidal, *Nat. Biotechnol.*, 2007, **25**, 1119–1126.
- 5 P. Cohen, *Nat. Rev. Drug Discovery*, 2002, **1**, 309–315.
- 6 A. Grancelli, A. Morros, M. E. Cabanas, O. Domenech, S. Merino, J. L. Vazquez, M. T. Montero, M. Vinas and J. Hernandez-Borrell, *Langmuir*, 2002, **18**, 9177–9182.
- 7 T. Guinan, C. Godefroy, N. Lautredou, S. Pace, P. E. Milhiet, N. Voelcker and F. Cunin, *Langmuir*, 2013, **29**, 10279–10286.
- 8 K. Hoc-Wydro, J. Kapusta, A. Jagoda, P. Wydro and P. Dynarowicz-Latka, *Chem. Phys. Lipids*, 2007, **150**, 125–135.
- 9 Z. Sajjadiyan, N. Cheraghi, S. Mohammadinejad and L. Hassani, *J. Biol. Phys.*, 2017, **43**, 127–137.
- 10 A. Ambrosini, G. Bossi, S. Dante, B. Dubini, L. Gobbi, L. Leone, M. G. P. Bossi and G. Zolese, *Chem. Phys. Lipids*, 1998, **95**, 37–47.
- 11 S. Goncalves, A. Teixeira, J. Abade, L. N. de Medeiros, E. Kurtenbach and N. C. Santos, *Biochim. Biophys. Acta, Biomembr.*, 2012, **1818**, 1420–1426.
- 12 Y.-W. Jiang, G. Gao, Z. Chen and F.-G. Wu, *New J. Chem.*, 2017, **41**, 4048–4057.
- 13 A. V. Agasosler, L. M. Tungodden, D. Cejka, E. Bakstad, L. K. Sydnes and H. Holmsen, *Biochem. Pharmacol.*, 2001, **61**, 817–825.
- 14 S. S. Feng, K. Gong and J. Chew, *Langmuir*, 2002, **18**, 4061–4070.



- 15 A. Preetha, N. Huilgol and R. Banerjee, *Colloids Surf., B*, 2006, **53**, 179–186.
- 16 C. Peetla, A. Stine and V. Labhasetwar, *Mol. Pharmaceutics*, 2009, **6**, 1264–1276.
- 17 A. Yousefpour, H. Modarress, F. Goharpey and S. Amjad-Iranagh, *J. Mol. Model.*, 2017, **23**, 1–18.
- 18 J. A. McLure, J. O. Miners and D. J. Birkett, *Br. J. Clin. Pharmacol.*, 2000, **49**, 453–461.
- 19 S. Nagar and K. Korzekwa, *Drug Metab. Dispos.*, 2012, **40**, 1649–1652.
- 20 D. A. Smith, L. Di and E. H. Kerns, *Nat. Rev. Drug Discovery*, 2010, **9**, 929–939.
- 21 A. Doisy, J. E. Proust, T. Ivanova, I. Panaiotov and J. L. Dubois, *Langmuir*, 1996, **12**, 6098–6103.
- 22 S. Smeazzetto, F. Tadini-Buoninsegni, G. Thiel, D. Berti and C. Montis, *Phys. Chem. Chem. Phys.*, 2016, **18**, 1629–1636.
- 23 G. Brezesinski and H. Mohwald, *Adv. Colloid Interface Sci.*, 2003, **100**, 563–584.
- 24 E. T. Castellana and P. S. Cremer, *Surf. Sci. Rep.*, 2006, **61**, 429–444.
- 25 W. T. Heller and K. C. Littrell, *Methods Mol. Biol.*, 2009, **544**, 293–305.
- 26 C. F. Majkrzak, N. F. Berk, S. Krueger and U. A. Perez-Salas, *Neutron Scattering Biol.*, 2006, 225–263.
- 27 M. Loesche, *Synth. Microstruct. Biol. Res., [Proc. Int. Conf.]*, 3rd, 1992, 91–108.
- 28 R. S. Armen, O. D. Uitto and S. E. Feller, *Biophys. J.*, 1998, **75**, 734–744.
- 29 P. Gianni and L. Lepori, *J. Solution Chem.*, 1996, **25**, 1–42.
- 30 W. Kern and D. A. Puotinen, *RCA Rev.*, 1970, **31**, 187–206.
- 31 J. Daillant and A. Gibaud, *X-ray and neutron reflectivity: principles and applications*, Springer, Berlin, 2009.
- 32 O. H. Seeck and E. Kentzinger, in *Neutron Scattering. Lectures of the JCNS Laboratory Course held at Forschungszentrum Jülich and the research reactor FRM II of TU Munich.*, ed. T. Brückel, G. Heger, D. Richter and R. Zorn, Forschungszentrum Jülich GmbH, Jülich, 2007, ch. 5, vol. 38.
- 33 S. K. Sinha, E. B. Sirota, S. Garoff and H. B. Stanley, *Phys. Rev. B: Condens. Matter Mater. Phys.*, 1988, **38**, 2297–2311.
- 34 S. Mattauch, A. Koutsoubas and S. Pütter, *J. Large-Scale Res. Facil.*, 2015, **1**, A8.
- 35 Y. Khaydukov, O. Soltwedel and T. Keller, *J. Large-Scale Res. Facil.*, 2015, **1**, A38.
- 36 Neutron Optics – Instrument Service – Science & Projects – MLZ – Heinz Maier-Leibnitz Zentrum, <http://www.mlz-garching.de/neutron-optics>.
- 37 H. Frielinghaus, A. Feoktystov, I. Berts and G. Mangiapia, *J. Large-Scale Res. Facil.*, 2015, **1**, A28.
- 38 G. D. Wignall and F. S. Bates, *J. Appl. Crystallogr.*, 1987, **20**, 28–40.
- 39 V. Pipich and Z. Fu, *J. Large-Scale Res. Facil.*, 2015, **1**, A31.
- 40 S. Desert, V. Thevenot, J. Oberdisse and A. Brulet, *J. Appl. Crystallogr.*, 2007, **40**, S471–S473.
- 41 L. G. Parratt, *Phys. Rev.*, 1954, **95**, 359–369.
- 42 G. Fragneto-Cusani, *J. Phys.: Condens. Matter*, 2001, **13**, 4973–4989.
- 43 R. J. Roe, *Methods of X-ray and neutron scattering in polymer science*, Oxford University Press, New York, 2000.
- 44 A. Nelson, *J. Appl. Crystallogr.*, 2006, **39**, 273–276.
- 45 H. Frielinghaus, *Phys. Rev. E: Stat., Nonlinear, Soft Matter Phys.*, 2007, **76**, 051603.
- 46 M. Kotlarchyk and S. M. Ritzau, *J. Appl. Crystallogr.*, 1991, **24**, 753–758.
- 47 J. N. Israelachvili, D. J. Mitchell and B. W. Ninham, *J. Chem. Soc., Faraday Trans. 2*, 1976, **72**, 1525–1568.
- 48 R. Nagarajan, *Langmuir*, 2002, **18**, 31–38.
- 49 F. D. Gunstone, J. L. Harwood and A. J. Dijkstra, *The lipid handbook with CD-ROM*, CRC Press, Boca Raton, 2007.
- 50 D. F. Evans and H. K. Wennerström, *The colloidal domain: where physics, chemistry, biology, and technology meet*, Wiley-VCH, New York, 1999.
- 51 S. W. Hui, J. T. Mason and C. Huang, *Biochemistry*, 1984, **23**, 5570–5577.
- 52 L. F. Braganza and D. L. Worcester, *Biochemistry*, 1986, **25**, 2591–2596.
- 53 R. Chen, D. Poger and A. E. Mark, *J. Phys. Chem. B*, 2011, **115**, 1038–1044.
- 54 R. V. McDaniel, T. J. McIntosh and S. A. Simon, *Biochim. Biophys. Acta, Biomembr.*, 1983, **731**, 97–108.
- 55 M. L. Manca, I. Castangia, P. Matricardi, S. Lampis, X. Fernandez-Busquets, A. M. Fadda and M. Manconi, *Colloids Surf., B*, 2014, **117**, 360–367.
- 56 N. E. Nagel, G. Cevc and S. Kirchner, *Biochim. Biophys. Acta, Biomembr.*, 1992, **1111**, 263–269.
- 57 H. Martinez-Seara, T. Rog, M. Pasenkiewicz-Gierula, I. Vattulainen, M. Karttunen and R. Reigada, *J. Phys. Chem. B*, 2007, **111**, 11162–11168.
- 58 R. Steitz, W. Jaeger and R. von Klitzing, *Langmuir*, 2001, **17**, 4471–4474.
- 59 H. Frielinghaus, M. Gvaramia, G. Mangiapia, S. Jaksch, M. Ganeva, A. Koutsoubas, S. Mattauch, M. Ohl, M. Monkenbusch and O. Holderer, *Nucl. Instrum. Methods Phys. Res., Sect. A*, 2017, **871**, 72–76.
- 60 C. Tanford, *The hydrophobic effect: formation of micelles and biological membranes*, Wiley, New York, 1973.
- 61 J. M. Seddon and R. H. Templer, in *Handbook of Biological Physics*, ed. R. Lipowsky and E. Sackmann, North-Holland, 1995, ch. 3, vol. 1, pp. 97–160.
- 62 S. Jaksch, F. Lipfert, A. Koutsoubas, S. Mattauch, O. Holderer, O. Ivanova, H. Frielinghaus, S. Hertrich, S. F. Fischer and B. Nickel, *Phys. Rev. E: Stat., Nonlinear, Soft Matter Phys.*, 2015, **91**, 022716.
- 63 C. Pereira-Leite, C. Carneiro, J. X. Soares, C. Afonso, C. Nunes, M. Lúcio and S. Reis, *Eur. J. Pharm. Biopharm.*, 2013, **84**, 183–191.
- 64 C. Nunes, G. Brezesinski, J. L. F. C. Lima, S. Reis and M. Lucio, *Soft Matter*, 2011, **7**, 3002–3010.
- 65 S. A. Safran, P. A. Pincus, D. Andelman and F. C. Mackintosh, *Phys. Rev. A: At., Mol., Opt. Phys.*, 1991, **43**, 1071–1078.
- 66 P. Herve, D. Roux, A. M. Bellocq, F. Nallet and T. Gulikkrzywicki, *J. Phys. II*, 1993, **3**, 1255–1270.
- 67 D. C. Morse and S. T. Milner, *Phys. Rev. E: Stat. Phys., Plasmas, Fluids, Relat. Interdiscip. Top.*, 1995, **52**, 5918–5945.



- 68 B. D. Simons and M. E. Cates, *J. Phys. II*, 1992, **2**, 1439–1451.
- 69 M. Vaccaro, R. Del Litto, G. Mangiapia, A. M. Carnerup, G. D'Errico, F. Ruffo and L. Paduano, *Chem. Commun.*, 2009, 1404–1406.
- 70 R. D. Harvey, N. Ara, R. K. Heenan, D. J. Barlow, P. J. Quinn and M. J. Lawrence, *Mol. Pharmaceutics*, 2013, **10**, 4408–4417.
- 71 S. Lesieur, C. Grabielle-Madelmont, M.-T. Paternostre and M. Ollivon, *Anal. Biochem.*, 1991, **192**, 334–343.
- 72 B. Mui, L. Chow and M. J. Hope, *Methods Enzymol.*, 2003, **367**, 3–14.
- 73 G. D'Errico, A. Silipo, G. Mangiapia, G. Vitiello, A. Radulescu, A. Molinaro, R. Lanzetta and L. Paduano, *Phys. Chem. Chem. Phys.*, 2010, **12**, 13574–13585.
- 74 G. Mangiapia, M. Vaccaro, G. D'Errico, H. Frielinghaus, A. Radulescu, V. Pipich, A. M. Carnerup and L. Paduano, *Soft Matter*, 2011, **7**, 10577–10580.
- 75 M. J. Saxton, *Biophys. J.*, 1987, **52**, 989–997.

