



# Effect of Benzocaine and Propranolol on phospholipid-based bilayers

G. Mangiapia,<sup>\*a</sup> M. Gvaramia,<sup>a,b</sup> L. Kuhrts,<sup>a</sup> J. Teixeira,<sup>c</sup> A. Koutsioubas,<sup>a</sup> O. Soltwedel<sup>d</sup> and H. Frielinghaus<sup>a</sup>

Received 00th January 20xx,  
Accepted 00th January 20xx

DOI: 10.1039/x0xx00000x

[www.rsc.org/](http://www.rsc.org/)

Cell membranes play a fundamental role in protecting the cell from its surroundings, in addition to host many proteins with fundamental biological tasks. A study of drug/lipid interactions is a necessary and important step to fully clarify the role and action mechanism of active ingredients, as well as to shed light on possible complications caused by drug overdose. In this paper, the influence of Benzocaine and Propranolol drugs on the structure of L- $\alpha$ -phosphatidylcholines-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 as well as 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 found results may be rationalized in terms of the molecular structure of drugs and may serve as starting point for explaining the toxic behavior in long-term and overdose scenarios.

## Introduction

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

The main task accomplished by cell membranes is essentially to protect the cell from its surroundings, acting as a selective barrier between the internal cytoplasm and the extracellular environment.<sup>2</sup> Part of their 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 include also the possibility to activate cell signaling, to trigger intracellular signaling pathways, and to interact with external substances, including active ingredients.<sup>3</sup> At this aim, most of the drugs designed or used in pharmacological therapies target membrane proteins,<sup>4</sup> like channel blockers or protein binding inhibitors.<sup>5</sup>

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 system. Concerning this, even in the presence of a specific drug interaction with a particular receptor, a partial drug uptake operated 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. A 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; what it is possible only when the protein is embedded in a lipid bilayer membrane. A possible 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 interaction appears a necessary and important step to fully understand and properly evaluate 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,<sup>6-9</sup> antimycotics, and<sup>10, 11</sup> antipsychotic,<sup>12, 13</sup> as well as anti-cancer agents<sup>14, 15</sup> have been studied from the point of view of their possible interaction with lipids,<sup>16</sup> even through simulation studies.<sup>17</sup> It has often been found that, even in the absence of 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.<sup>18-20</sup>

The high complexity shown by the cell membranes makes the

<sup>a</sup> Forschungszentrum Jülich GmbH, Jülich Centre for Neutron Science Außenstelle am Heinz Maier-Leibnitz Zentrum, Lichtenbergstraße 1, D-85747 Garching, Germany. E-mail: mangiapia.gaetano@alice.it

<sup>b</sup> Ivane Javakhishvili Tbilisi State University, 1 Chavchavadze Avenue, GE-0179 Tbilisi, Georgia.

<sup>c</sup> Laboratoire Léon Brillouin (CEA-CNRS), CEA-Saclay, F-91191 Gif-sur-Yvette CEDEX, France.

<sup>d</sup> Heinz Maier-Leibnitz Zentrum, Technische Universität München, Lichtenbergstraße 1, D-85747 Garching, Germany.

physico-chemical and the biophysical interactions with drugs very challenging to investigate. At this aim, simplified membrane systems, like liposomes,<sup>21</sup> giant unilamellar vesicles<sup>22</sup>, Langmuir monolayers,<sup>23</sup> or solid supported membranes<sup>24</sup> 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 a very good alternative to the natural membranes, since they allow investigating the 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 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, 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 formed<sup>25</sup> and on the bilayer characteristics,<sup>26,27</sup> highlighting eventual variations operated by the 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 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 misleading or incomplete picture of the system. Moreover, it will be noted how the present approach, involving complementary neutron-based techniques, is able to give a valuable set of information about the systems investigated.

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

## Experimental

### Materials

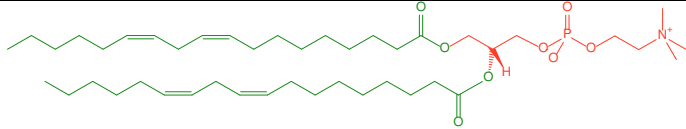
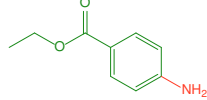
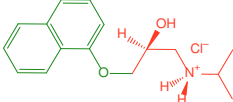
SoyPC (mainly composed by 1,2-dilinoleoyl-*sn*-glycero-3-phosphocholine, purity 95%, averaged molecular weight 775.037 a.m.u.) was obtained from Avanti Polar Lipids Inc. (Alabaster/AL, USA). Benzocaine (purity 98%, molecular weight 165.19 a.m.u.) and Propranolol (in hydrochloride form, purity  $\geq 99\%$ , 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.

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.

**Table 1** – SLDs ( $\rho$ ) 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 from MD simulations carried out on phospholipids<sup>28</sup> and from the theory of group contributions.<sup>29</sup>

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

### 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: at this aim, a selected amount of SoyPC was dissolved in the secondary alcohol, such to have a molar ratio of  $\approx 2$  mol % between the phospholipid and the solvent, and a total volume of  $\approx 15$  ml. The mixed components were stirred, up to have a complete dissolution of the phospholipid in the solvent. Afterwards, always under stirring, the 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: 1.50 cm – 5.0 cm – 2.0 cm) 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 evaporating at room temperature and at a reduced pressure ( $\sim 250$  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  $D_2O$ . 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.<sup>30</sup>

### Preparation of samples for small angle 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 get a concentration of  $\approx 1$  mg ml<sup>-1</sup>. Appropriate amounts of this mother solution were transferred in 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 drug and phospholipid. The dissolution of the components was promoted by a slight warming ( $\approx 40^\circ\text{C}$ ) and a very short sonication treatment ( $\approx 5$  min). Afterwards, a thin film was 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 ( $\approx 30$  min). Subsequently, an aliquot of each suspension was repeatedly extruded through a polycarbonate membrane of 100 nm pore size, for 11 times. The final (molal) concentration of SoyPC was in all cases 5.0 mmol/kg; 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 a 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:<sup>31</sup>

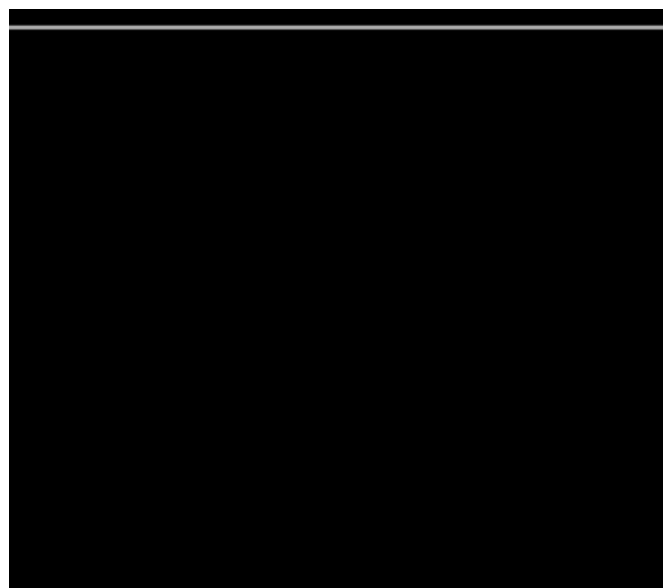
$$R = \frac{16\pi^2}{Q_z^2} \left| \int \rho(z) \exp(iQ_z z) dz \right|^2$$

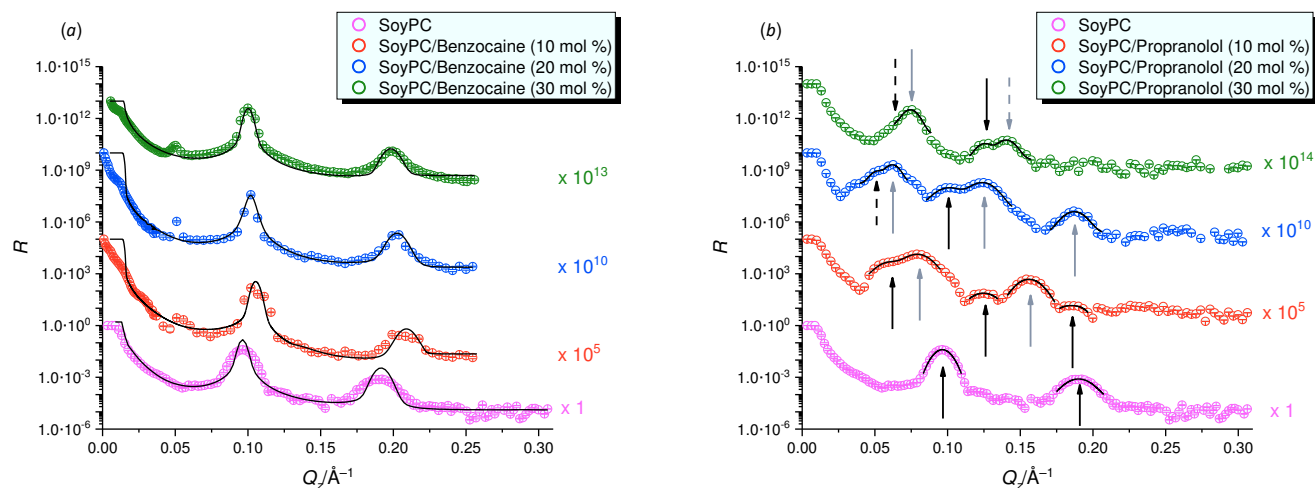
The function  $\rho(z)$ , the scattering length density at the various  $z$  values, is dependent on the local composition and allows obtaining information on the lipid bilayer at the nanoscale.

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.<sup>32</sup>

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.<sup>33</sup>

At MARIA,<sup>34</sup> the wavelength of the neutrons was set to  $\lambda = 10$  Å for NR and to  $\lambda = 5.0$  Å for GISANS investigations, with a wavelength spread of  $\Delta\lambda/\lambda \leq 0.10$  by means of a velocity selector. In NR configuration, an elliptically curved focusing guide along the vertical direction, allowed obtaining a beam size of  $10 \cdot 0.1$  mm<sup>2</sup>; in GISANS setup, a pinhole collimation setting was used, with a beam aperture of  $16 \cdot 2$  mm<sup>2</sup>. 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 range  $3.3 \cdot 10^{-3} < Q_z/\text{\AA}^{-1} < 0.31$  for NR and  $0 < Q_z/\text{\AA}^{-1} < 0.25$ ;  $-0.15 < Q_y/\text{\AA}^{-1} < 0.15$  for GISANS, being  $y$  the





**Fig. 2** – NR data collected for (a) SoyPC/Benzocaine and (b) SoyPC/Propranolol systems in D<sub>2</sub>O, 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 from fitting of 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, have been fitted to the experimental data in proximity to the maxima, and reported as solid curves. Not detected peaks, or weak peaks whose position deviates from that of a lamellar arrangement are indicated with dashed arrows (see also Table 4).

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 measurement ( $\approx 1.5$ –2 h per sample). In GISANS mode, the measurements were mostly performed below the critical angle  $\theta^{(cr)}$  of total internal reflection: in this case an evanescent wave propagating in to the sample occurred, allowing collecting information over the complete surface region down to the depth of this wave.

At NREX,<sup>35</sup> 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  $50 \cdot 1.0 \text{ mm}^2$  was used for defining the incident beam, whereas scattered neutrons were detected by means of a  $^3\text{He}$  detector set at 2.46 m from the sample position. With this configuration, data in the range  $5.6 \cdot 10^{-3} < Q_z/\text{\AA}^{-1} < 0.26$  were collected in NR mode and  $0 < Q_z/\text{\AA}^{-1} < 0.12$ ;  $0 < Q_y/\text{\AA}^{-1} < 0.10$  in GISANS setup.

At the neutron reflectometer TREFF,<sup>36</sup> a wavelength of  $4.7 \text{ \AA}$  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  $0.4 \cdot 60 \text{ mm}^2$ , allowing covering a  $Q_z$  range of  $9.1 \cdot 10^{-4} < Q_z/\text{\AA}^{-1} < 0.25$ .

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 TPA diffractometer, at LLB (Laboratoire Léon Brillouin), Saclay, France.

At KWS-1,<sup>37</sup> neutrons with an average wavelength of  $\lambda = 5.0 \text{ \AA}$  and wavelength spread  $\Delta\lambda/\lambda \leq 0.10$  were used. A two-dimensional  $128 \times 128$  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 dis-

tances in meters) measured neutrons scattered from the samples. These configurations allowed collection of data in a range of the scattering vector modulus  $Q = 4\pi/\lambda \sin(\theta/2)$  ranged between  $0.0028$  and  $0.40 \text{ \AA}^{-1}$ , being  $\theta$  the scattering angle. The investigated samples were contained in a closed quartz cell, in order to prevent the solvent evaporation, and were kept under measurements for a period such 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.<sup>38</sup>

At KWS-3,<sup>39</sup> 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 to 9.6 m. KWS-3 uses a double focusing mirror principle to access to very small angle region; a bidimensional detector with square pixels of  $0.35 \text{ mm}$  in size allowed collecting data ranged between  $6 \cdot 10^{-4}$  and  $2 \cdot 10^{-3} \text{ \AA}^{-1}$ .

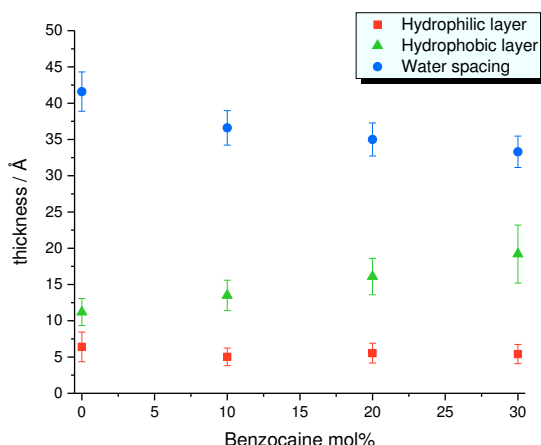
At TPA,<sup>40</sup> neutrons with a wavelength of  $12 \text{ \AA}$  were used, with a wavelength spread  $\Delta\lambda/\lambda \leq 0.11$ . A multibeam design of the collimation system and a high resolution two dimensional image plate detector located at 6.27 m from the sample position, allowed collecting data in the range  $4 \cdot 10^{-4} < Q/\text{\AA}^{-1} < 4.7 \cdot 10^{-3}$ : data corrected for background and empty cell scattering were radially averaged and rescaled on to the small-angle data collected at KWS-1. At this aim, a scale factor was evaluated in order to have the smallest sum of the squares of the difference 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 Fig. 2(b), NR data are reported for SoyPC/Benzocaine and SoyPC/Propranolol systems, respectively. Inspection of data allows recognizing a different behavior shown by the two different drugs.





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

For systems containing Benzocaine, the presence of two Bragg peaks at  $Q_z^{(1)} \cong 0.1 \text{ \AA}^{-1}$  and  $Q_z^{(2)} = 2 Q_z^{(1)}$  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 by 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 or comparable to the minimum angle attained during reflectivity scans ( $Q_z^{(cr,1)} \cong 0.002 \text{ \AA}^{-1}$ ) whereas the second is located around  $Q_z^{(cr,2)} \cong 0.012 \text{ \AA}^{-1}$ . Both these characteristics occur only in the presence of Benzocaine, being absent for the pure SoyPC system.

Data for the above described systems were analyzed by means of the Parratt formalism,<sup>41</sup> describing the scattering length density distribution, along the  $z$  direction, in terms of a stack of distinct layers. In this analysis, Si substrate was modeled as an infinite-thick planar surface covered with a thin layer of oxide ( $\text{SiO}_2$ ).<sup>42</sup> The Parratt approach uses a transition matrix for each layer, able to correlate the amplitude of the neutron wave propagating in a certain layer to the behavior in the previous adjacent layer.<sup>43</sup> A set of matrices minimizing the sum of squares of the differences between the experimental and the calculated reflectivities were found, characterizing thickness, scattering length density, solvation (solvent volume fraction), and interfacial roughness of the different layers. These calculations were implemented in the MOTOFIT application:<sup>44</sup> 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

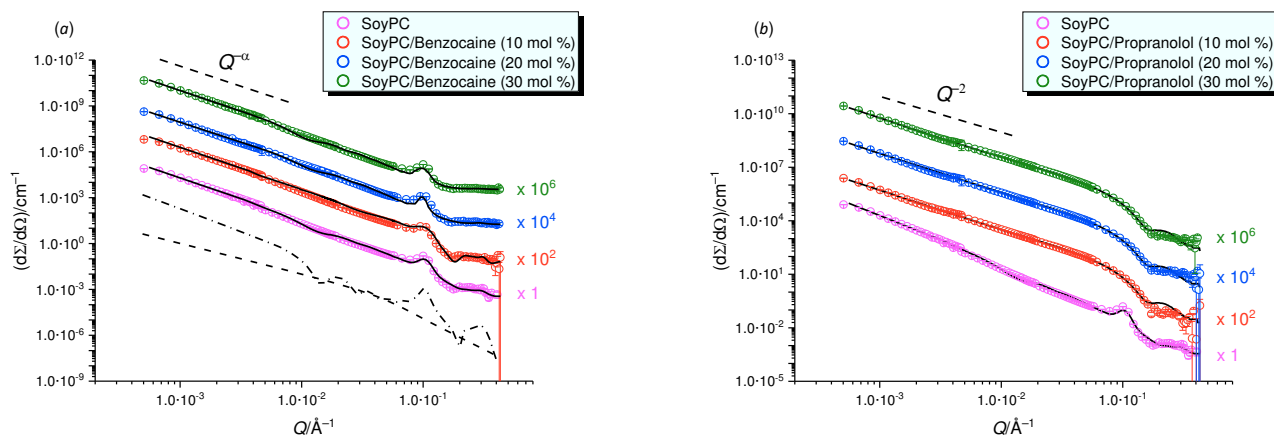
values. For examples, the two thicknesses belonging to the opposite hydrophilic portions of bilayers were constrained to be equal each other. Data obtained from analysis of the systems are reported in Fig. 3.

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. Position of some reflections is, again, typical of a lamellar arrangement, but, differently from what seen for SoyPC/Benzocaine systems, here the peaks appear broad and some of them overlap. This behavior is particularly apparent for the system at 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  $Q_z^{(1b)} \cong 0.0618 \text{ \AA}^{-1}$ , being its shoulder-like shape appearance due to the high signal connected to the Bragg reflection at  $Q_z^{(1a)}$  that partially masks its position. Other peaks most likely correlated to  $Q_z^{(1b)}$ , appear at  $Q_z^{(2b)} \cong 0.124 \text{ \AA}^{-1}$  as well as at  $Q_z^{(3b)} \cong 0.186 \text{ \AA}^{-1}$  (this latter being rather weak). These reflectivity data appear as composed by the overlapping of different structures, and are not susceptible of a simple quantitative analysis.

As mentioned, the study of the interaction of the active ingredients with bilayers was also carried out in the bulk, by means of SANS/USANS. At this aim, Figs. 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 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  $\alpha$  is ranged between 2 and 4. On the other hand, 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 spanned 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 ( $\cong 500 \text{ nm}$ ), 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 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  $\alpha$  of the power law observed at low  $Q$  values is connected to the lamellarity of the aggregates.<sup>45</sup>

Scattering arising from multilamellar liposomes may be de-



**Fig. 4** – Scattering cross sections obtained for (a) SoyPC/Benzocaine and (b) SoyPC/Propranolol systems prepared through 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 from 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 (dash-dotted line, multiplied by 0.01) and unilamellar liposomes (dashed line, multiplied by 0.001) have illustratively been reported for SoyPC aqueous system.

scribed assuming the liposomes as a collection of randomly oriented assembly of lamellar stacks, neglecting their curvature, since the complete missing of the Guinier region. 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:<sup>46</sup>

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

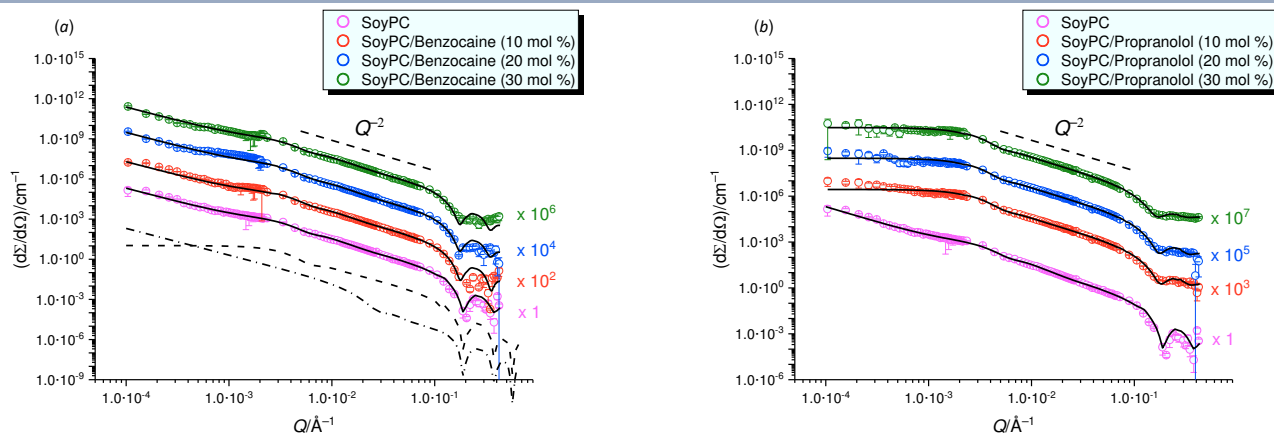
where  $v$  indicates the volume of a single stack,  $\Delta\rho$  is the contrast between the lipid bilayers and the solvent,  $f(Q)$  is the form factor of a single bilayer, *i.e.*  $f(Q) = \sin(Q\tau/2)/(Q\tau/2)$ , whereas  $S(Q)$  represents the structure factor taking into account the interferences 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  $\tau$ , and the distance between the centers of two consecutive layers  $d$ . Both  $\tau$  and  $d$  have been allowed

to be polydisperse: the former with a Schulz–Zimm distribution function related to the polydispersity index  $i_\tau$ , the latter with a Gaussian distribution having a standard deviation  $\sigma_d$ . In particular the ratio  $\sigma_d/d$ , named Hosemann factor, has been estimated. At 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.<sup>46</sup>

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

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

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



**Fig. 5** – Scattering cross sections obtained for (a) SoyPC/Benzocaine and (b) SoyPC/Propranolol systems prepared through 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 from 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 (dash-dotted line, multiplied by 0.001) have illustratively been reported for SoyPC aqueous system.

term  $(d\Sigma/d\Omega)_{\text{incoh}}$  has been added, representing the incoherent contribution to the cross section measured, mainly due to the presence of hydrogen atoms. Data obtained from 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  $Q$  region, *i.e.*  $(d\Sigma/d\Omega) \propto Q^{-2}$  when  $5 \cdot 10^{-3} < Q/\text{\AA}^{-1} < 7 \cdot 10^{-2}$ , similar to what observed for sonicated SoyPC/Propranolol systems. Anyway, differently from that latter case, data at low  $Q$  for SoyPC/Benzocaine suspensions show an excess of scattering that quantitative analysis shows to be imputable 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 to what seen for the corresponding simply sonicated systems, are unquestionably smaller, being the Guinier regime clearly visible when  $Q < 2 \cdot 10^{-3} \text{\AA}^{-1}$ . All data for 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, being the main difference 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 from the presence of the Guinier regime for unilamellar vesicles at very low  $Q$  values, made out 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 are 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 to the drug molecular structure. Generally speaking, the perturbation effects may depend on the hydrophobicity/hydrophilicity degree of the active ingredient, on its molecular shape as well as on possible attractive or repulsive interactions (*e.g.* electrostatic) that can be established: all these sides

contribute in a complex way to (eventually) perturb the phospholipid packing. At this aim, it has been proved to be useful the concept of the packing factor, introduced by Israelachvili, Mitchell, and Ninham,<sup>47</sup> that is able to explain, justify and very often predict the molecular self-assembly behavior in surfactant solutions. The packing factor  $N_s$  is defined as

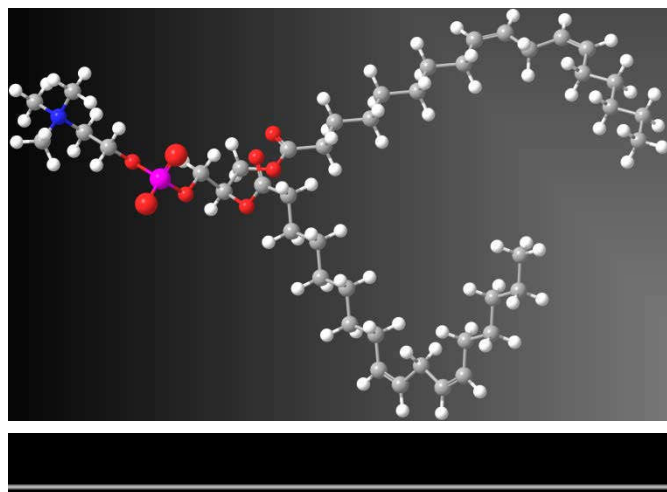
$$N_s = \frac{v}{l \cdot a_0} \quad (3)$$

where  $v$  is the volume of the total surfactant hydrophobic portion,  $l$  represents the tail length, whereas  $a_0$  is commonly defined as the equilibrium area per molecule at the aggregate surface. Nagarajan<sup>48</sup> has stressed that this latter parameter has not to be intended as a simple geometrical variable connected to the size and shape of the polar heads, rather as a thermodynamic quantity coming out from a minimum free energy approach. For the following analysis, it has to be noted that the hydrophobic portion of SoyPC bilayer was assumed to be in the liquid crystalline phase, as the main lipid phase transition (gel-sol,  $P_B \rightarrow L_\alpha$ ) temperature ( $< -40^\circ\text{C}$ ) is well below the room temperature.

Quantitative analysis of membrane characteristics was performed for SoyPC/Benzocaine system. 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 ( $\approx 15 \text{\AA}$ ) water layer was found directly stratified on the  $\text{SiO}_2$  layer ( $\approx 2 \text{ nm}$ ) covering, in turn, the Si block. In all cases, roughness values of the interfaces were ranged between  $2 \text{\AA}$  and  $7 \text{\AA}$ . Extensions of hydrophobic and hydrophilic portions of the bilayers, as well as dimension of the water spacing, have been plotted in Fig. 3 for the drug concentrations investigated. Inspection of data reveals a certain effect operated by the presence of the drug on the bilayer: inclusion of Benzocaine reduces the repetition distance among the bilayers, but not in a monotonic fashion. For pure SoyPC system, this distance results to be  $\approx 2\pi/Q^{(1)} = 65.2 \text{\AA}$ , reaches its minimum value at 10 mol % of the drug, where it is  $59.8 \text{\AA}$ , and slightly increases again for 20% and 30% mol amount ( $61.9 \text{\AA}$  and  $62.7 \text{\AA}$ , respectively). Although these values are independent of any theoretical model, analysis of Fig. 3, based on Parratt algorithm, shows that this trend is the outcome of two different tendencies: (i) the water spacing that slightly decreases with the

**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  $\alpha$  of the power law observed in the range  $4 \cdot 10^{-4} < Q/\text{\AA}^{-1} < 1 \cdot 10^{-2}$ , the number  $N$  of lamellae in multilamellar liposomes, the average thickness  $\tau$  of the bilayer together with its polydispersion index  $i_\tau$ , the distance  $d$  between two consecutive bilayers and the Hosemann factor  $\sigma_d/d$ . In systems composed by both unilamellar and multilamellar liposomes, thicknesses of bilayers have been constrained to be identical each other. The polydispersion 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	$\alpha$	$N$	$\frac{\tau}{\text{\AA}}$	$i_\tau$	$\frac{d}{\text{\AA}}$	$\frac{\sigma_d}{d}$
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	–	–



increase of the drug content (even taking in to account the uncertainty affecting the data), and (ii) the thickness of hydrophobic portion of the bilayer, that is progressively enlarged as soon the drug amount is raised up. Concerning this 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.<sup>49</sup> Although this value refers to the crystallographic distance, and despite the average thickness of the hydrophobic portion of bilayers seldom exceeds 1.8 times the fully extended length of the hydrocarbon chains,<sup>50</sup> it is nevertheless still higher than the experimental thicknesses found, spanning the range from 11.2 to 19.2 Å, symptom of an interdigitation or tilting of the hydrophobic chains. Generally speaking, there are manifold circumstances where interdigitation phenomena are observed. The most important regard 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);<sup>51</sup> the application of high pressures,<sup>52, 53</sup>; and the solubilization of molecules able to promote or increase intermolecular lateral repulsions.<sup>54, 55</sup> 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 argued<sup>56</sup> that the repulsions occurring in the interfacial region of the bilayer initially promote a progressive tilting of the lipid chains, for adapting to the large interfacial molecular area. Increasing of tilting angle increases, in turn, torsional and angular tensions in the lipid

chain: beyond a certain threshold, tilt cannot be further increased, and hydrocarbon interdigitation occurs.

Analysis of the hydrophobic layer thickness shows, indeed, that insertion of Benzocaine reduces the above described phenomena, as an increasing of the thickness is observed at 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 it can be seen in Fig. 6.<sup>57</sup> It can be argued that insertion of Benzocaine is able to promote a better packing of the chains, being capable to “fill” 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 it will be glimpsed in the following. Thickness of the hydrophilic portion is roughly constant in the range of concentration studied: Benzocaine has a small hydrophilic portion represented by the ammine functional group and its insertion does not dramatically modify this part of the bilayer.

Finally, it is interesting to discuss and rationalize the presence of other two 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  $Q_z^{(1/2)} \cong 0.05 \text{ Å}^{-1}$ , both of these features occurring only in the presence of the drug.

There are two distinct sudden changes of slope in data trends, for systems containing Benzocaine: the first is close to the minimum angle reached during investigations, namely  $Q_z^{(cr1)} \cong 0.002 \text{ Å}^{-1}$ , whereas the second is located around  $Q_z^{(cr2)} \cong 0.012 \text{ Å}^{-1}$ . Their presence might be ascribed to the different contrast  $\Delta\rho = \rho_{\text{inf}} - \rho_{\text{Si}}$  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:

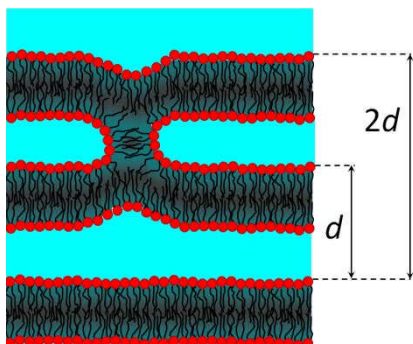
$$Q_z^{(cr)} = \frac{4\pi}{\lambda} \sin\theta^{(cr)} \cong 4\sqrt{\pi} \sqrt{\rho} \quad (4)$$

The presence of macroscopically thick patches of bilayer material with a scarce penetration of D<sub>2</sub>O among the bilayers may rationalize the results observed: these patches, for which D<sub>2</sub>O is

**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  $R_{\text{core}}$  of unilamellar liposomes together with its polydispersion index  $i_{R_{\text{core}}}$ , and the thickness  $\tau$  of the bilayer. The polydispersion 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 coexistence with multilamellar liposomes occurs, the number  $N$  of lamellae, the distance  $d$  between two consecutive bilayers and the Hosemann factor  $\sigma_d/d$  are also reported. In order to reduce the number of fitting parameters, in systems composed by both unilamellar and multilamellar liposomes, thicknesses of bilayers have been constrained to be identical. Furthermore,  $d$  and  $\sigma_d/d$  have been kept fixed and equal to the value averaged for the corresponding sonicated systems.

System	$\frac{R_{\text{core}}}{\text{Å}}$	$i_{R_{\text{core}}}$	$\frac{\tau}{\text{Å}}$	$N$	$\frac{d}{\text{Å}}$	$\frac{\sigma_d}{d}$
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	–	–	–





**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.

not “seen” as final layer (for the purposes of the critical angle determination) coexist with normal hydrated portions of bilayers. It has to be assumed that the size of these “island” of scarcely hydrated material is bigger than the coherence volume of the incident beam, as the two angles would not be visible otherwise. Application of eqn. (4) for  $\rho_{\text{inf}} = \rho_{\text{D}_2\text{O}}$  gives a value ( $0.0146 \text{ \AA}^{-1}$ ) very close to the observed  $Q_z^{(\text{cr},2)}$ . 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, being in qualitative agreement with the experimental observation that  $Q_z^{(\text{cr})} < Q_z^{(\text{cr})}$  (as it 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.<sup>58</sup>

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/Propranol systems, bilayers fluctuate much more, and so they show the typical critical angle of Si/D<sub>2</sub>O interface. This hypothesis may be proven through Grazing Incidence Neutron Spin-Echo Spectroscopy (GINSSES) investigations. At present, the main limitation for such technique is represented by the low counting rates due to the limited scattering volume that only comprises the volume covered by the evanescent wave. However, improvements for dealing with this drawback are currently under development by some of us.<sup>59</sup>

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 coming out 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, that will then result in twice

the typical repetition distance, resulting in turn in the appearance of the peak at half the value of the first Bragg peak (as sketched in Fig. 7). The formation of stalks, that are real crystal-like defects in the lipid arrangement, has to be ascribed to the capability of Benzocaine to induce localized structures with negative curvature of the bilayer membrane: such bending can only be the result of a global increasing of the packing factor  $N_s$  with respect to the value assumed in pure SoyPC bilayers. Actually, it has been seen that addition of Benzocaine provide a packing of hydrophobic chains that tends to approach to the situation existing in the crystal; the experimental collected evidences suggest that this better packing occurs along with reduced repulsive (or even attractive) interactions experienced by the phospholipid heads that approach to shorter distances. 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,<sup>48, 60</sup> shows that  $a_0$  is dependent on a headgroup repulsion parameter  $\beta$ , and on the interfacial free energy  $\sigma$ :  $a_0 = \sqrt{\beta/\sigma}$ . A decrease of  $\beta$  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, that 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.

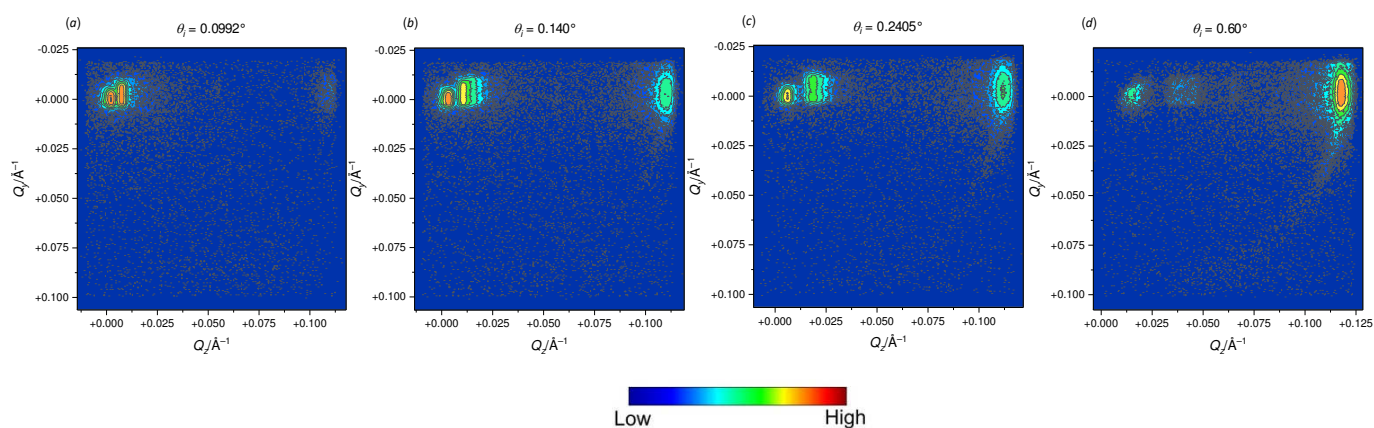
It has often been supposed that the formation of such defects is the prelude to a transition to non-lamellar phases.<sup>61</sup> Indeed, the formation of hexagonal and tetragonal lattices has been found<sup>62</sup> for the investigation of phospholipid-based membranes in the presence of Ibuprofen active ingredient, being coupled to the presence of similar stalks in the double layer arrangement. With the aim to investigate about a possible similar situation in the our system, selected GISANS scans have been performed, most of them below the critical angle of total internal reflection of the Si/D<sub>2</sub>O interface. In this case, eventual lateral correlations are probed inside the bilayers, up to the depth  $\Lambda_{\text{eww}}$  of the evanescent wave given by

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

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  $\theta_i$  ranging from  $0.020^\circ$  to  $0.60^\circ$ , covering the region where the two critical angles were observed in NR scans, with the aim to gain depth resolved results. 2-D detector illustrative images at selected incoming angles  $\theta_i$  of  $0.099^\circ$ ,  $0.14^\circ$ ,  $0.24^\circ$ , and  $0.60^\circ$  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 lamellar arrangement of bilayers appears. Its

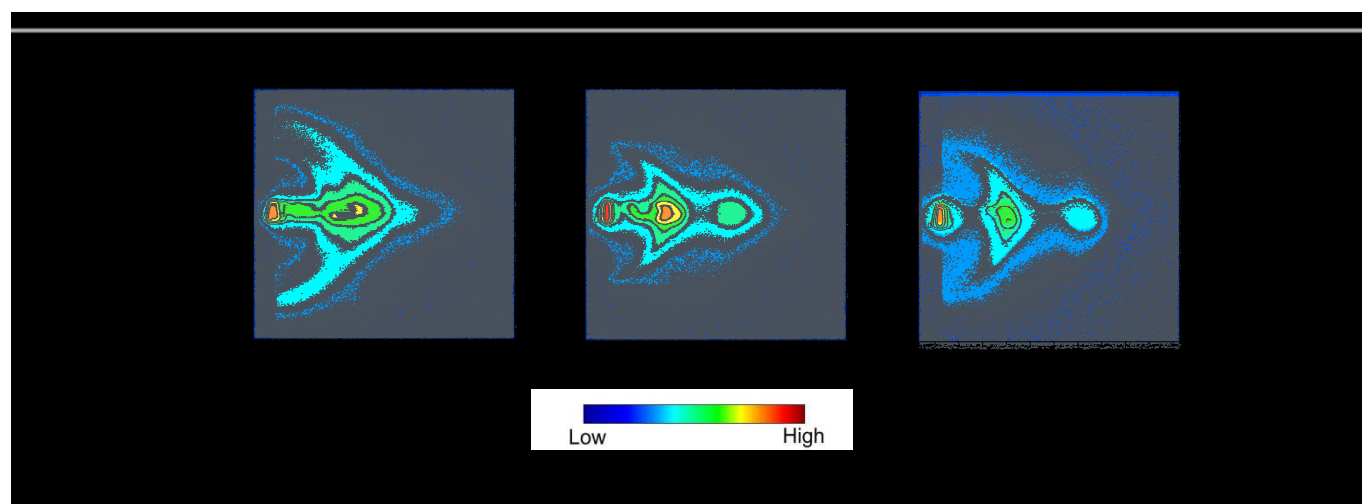


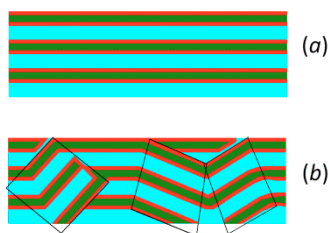


intensity increases as soon  $\theta_i$  increases, as the penetration depth of the evanescent wave is progressively enlarged, so that a higher scattering of the lamellar bilayer structure is observed. Although not appreciable when  $\theta_i < 0.1^\circ$ , a Debye-Scherrer ring starts to appear at  $\theta_i = 0.14^\circ$ , being 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 is far from being complete but rather deviates only to a few degree from a perfect parallel ordering. Interestingly, no reflections arising from crystalline structures inside the double layer are detected, suggesting that the defects corresponding to the formation of isolated stalks, not distributed according to the existence of a lattice. Based on the NR/GISANS results, we may point out that the presence of the stalks inside the bilayer arrangement, the incomplete Si surface coverage with fully hydrated bilayers, and the not complete parallel alignment of bilayer with respect to the Si surface are the main sources for the not perfect description of data by the Parratt model, *i.e.* the not exact fitting of curves in Fig. 2(a).

Much more complicated appear to be the NR data obtained for the system SoyPC/Propranolol. As mentioned in the *Results* section, inclusion of Propranolol drastically changes reflectivity

trends: analysis of data reported in Fig. 2(b) reveals, differently from what was seen for the systems in the presence of Benzo-caine, 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.<sup>62</sup> At this aim, peaks appearing in NR data have been highlighted in Fig. 2(b) by regrouping their  $Q_z$  values with arrows having the same color; positions found have been 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 shows, indeed, a quite complex trend. For 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 by 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. Concerning this lowest amount, five maxima are clearly visible: analysis of their positions shows that they are correlated in the  $Q$ -ratio 1:2 (or 1:2:3), being the second set (the one located at higher  $Q$  and highlighted with grey arrows) clearly visible, whereas the primary reflection of the first set (indicated with a





**Fig. 10** – Sketch of the structural evolution induced by 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 the display of the drug to induce high-curvature of the bilayers. In the drawing, disordered lamellae are highlighted by black lines, for making easier their visualization.

black arrow) appears as a shoulder because of the high signal connected to the primary peak of the second pair. Trends get 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, this latter peak is even not visible (being its expected position indicated with the black dashed arrow). Analysis of the maxima positions 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 lamellae 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. It can be argued a smaller or a nearly null capability of Propranolol to act as “filler”, that is unable to efficiently occupy the space created in consequence of the reduction of tilting/interdigitation of SoyPC hydrophobic chains. Furthermore, differently to what 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 the

lamellar powders, as schematically displayed in Fig. 10.

Summarizing, in the presence of Benzocaine only a very limited local disorder of lamellar arrangement (along with formation of negatively curved structures inside the lamellar phase) occurred, so that the NR data were still describable in 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.<sup>63</sup> 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 evidences are consistent with the results found in the presence of SoyPC: a stronger disorder caused by Propranolol presence 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 found also for other drugs, although with different characteristics.<sup>64</sup>

The results obtained from NR/GISANS analysis of SoyPC bilayers in the presence of Benzocaine and Propranolol drugs allows foreseeing what should be the characteristics of such bilayers in bulk conditions, *i.e.* in water dispersions: in 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) size and characteristics of a population of liposomes are the result of a complex balance between entropy of mixing<sup>65</sup> and bilayer curvature elasticity.<sup>66-68</sup> In the limit of large liposomes, with a dimension much larger than the surfactant size, the curvature energy per unit area of bilayer  $f_c$  is given by<sup>65</sup>

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

being  $K$  and  $\bar{K}$  the bending moduli, depending, in turn, on the

**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 peaks. Most of third reflections are very weak and their positions, even if mentioned in the text, are not susceptible of an accurate determination and have not been reported in the Table. Not detected peaks, or weak peaks whose positions deviates from that of a lamellar arrangement are followed by a question mark and correspond to dashed arrows in Fig. 2(b). All the found values are reported with three significant digits, according to the accuracy of the determination.

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

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 case of a bilayer with relatively high curvature, as appears to be for SoyPC/Propranolol system, the energy requested 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 to a single liposome may be observed. In the most extreme cases, unilamellar liposomes are exhibited. What observed from NR and GISANS data, could suggest the formation of oligolamellar or even unilamellar liposomes for aqueous suspension 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 system SoyPC/Propranolol, independently of the preparation method adopted (simple sonication or extrusion), whereas multilamellar liposomes are observed for SoyPC/Benzocaine sonicated suspensions. Results obtained through the fitting of theoretical models described in the *Results* section are reported in Table 2 and Table 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.<sup>69,70</sup> The essential bilayer characteristics obtained by analysis of NR data are retained for liposomes studied in aqueous environment by means of SANS investigations. Values of bilayer thickness  $\tau$  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. More, 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 if 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 falling in the  $Q$  range spanned 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 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, characteristics of SoyPC-based lip-

osomes 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 extruded 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 liposome spheres to ellipsoid occur during the cross of the pores.<sup>71</sup>

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 rearrange giving rise to 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,<sup>72</sup> even if in some cases, because of rearrangement phenomena, it has also been found to produce multilamellar aggregates,<sup>73</sup> or even more complex morphologies.<sup>74</sup> The general trend is here fulfilled, even if the small fraction of multilamellar aggregates suggests that a higher amount of energy, *i.e.* higher number of extrusion steps, has to be provided for a further or total reduction of the multilamellar aggregates.

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

## Conclusions

In this paper, the influence of Benzocaine and Propranolol active ingredients on SoyPC-based bilayers has been investigated. Despite of their small and simple chemical structure, it has been found a marked and different effect of these two drugs: on one side, Benzocaine induces a local high 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 the presence of charged heads (occurring in 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 of dosage in any medical posology. Nonetheless, it may be argued a correlation between the effects observed at expenses of the bilayers and the complications due to drug high 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.<sup>75</sup>

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 here adopted 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.

## Acknowledgements

Jülich Centre for Neutron Science, Maier-Leibnitz Center, Max Planck Institute for Solid State Research and Laboratoire Léon Brillouin are fully acknowledged for the excellent support obtained, as well as for provision of beam time. One of us (MG) thanks the Georgian Shota Rustaveli National Science foundation for financial support

## Notes and references

- G. M. Cooper and R. E. Hausman, *The Cell: A Molecular Approach*, Sinauer Associates 2016.
- 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.
- W. Lim, B. Mayer and T. Pawson, *Cell Signaling: principles and mechanisms*, Taylor & Francis Group 2014.
- M. A. Yildirim, K. I. Goh, M. E. Cusick, A. L. Barabasi and M. Vidal, *Nat Biotechnol*, 2007, **25**, 1119-1126.
- P. Cohen, *Nat Rev Drug Discov*, 2002, **1**, 309-315.
- 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.
- T. Guinan, C. Godefroy, N. Lautredou, S. Pace, P. E. Milhiet, N. Voelcker and F. Cunin, *Langmuir*, 2013, **29**, 10279-10286.
- K. Hoc-Wydro, J. Kapusta, A. Jagoda, P. Wydro and P. Dynarowicz-Latka, *Chem Phys Lipids*, 2007, **150**, 125-135.
- Z. Sajjadiyan, N. Cheraghi, S. Mohammadinejad and L. Hassani, *J Biol Phys*, 2017, **43**, 127-137.
- 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.
- S. Goncalves, A. Teixeira, J. Abade, L. N. de Medeiros, E. Kurtenbach and N. C. Santos, *Bba-Biomembranes*, 2012, **1818**, 1420-1426.
- Y.-W. Jiang, G. Gao, Z. Chen and F.-G. Wu, *New Journal of Chemistry*, 2017, **41**, 4048-4057.
- A. V. Agasosler, L. M. Tungodden, D. Cejka, E. Bakstad, L. K. Sydnese and H. Holmsen, *Biochem Pharmacol*, 2001, **61**, 817-825.
- S. S. Feng, K. Gong and J. Chew, *Langmuir*, 2002, **18**, 4061-4070.
- A. Preetha, N. Huilgol and R. Banerjee, *Colloid Surface B*, 2006, **53**, 179-186.
- C. Peetla, A. Stine and V. Labhasetwar, *Mol Pharmaceut*, 2009, **6**, 1264-1276.
- A. Yousefpour, H. Modarress, F. Goharpey and S. Amjad-Iranagh, *J. Mol. Model.*, 2017, **23**, 1-18.
- J. A. McLure, J. O. Miners and D. J. Birkett, *Brit J Clin Pharmacol*, 2000, **49**, 453-461.
- S. Nagar and K. Korzekwa, *Drug Metab Dispos*, 2012, **40**, 1649-1652.
- D. A. Smith, L. Di and E. H. Kerns, *Nat Rev Drug Discov*, 2010, **9**, 929-939.
- A. Doisy, J. E. Proust, T. Ivanova, I. Panaiotov and J. L. Dubois, *Langmuir*, 1996, **12**, 6098-6103.
- S. Smeazzetto, F. Tadini-Buoninsegni, G. Thiel, D. Berti and C. Montis, *Phys Chem Chem Phys*, 2016, **18**, 1629-1636.
- G. Brezesinski and H. Mohwald, *Adv Colloid Interfac*, 2003, **100**, 563-584.
- E. T. Castellana and P. S. Cremer, *Surf Sci Rep*, 2006, **61**, 429-444.
- W. T. Heller and K. C. Littrell, *Methods in Molecular Biology (Totowa, NJ, United States)*, 2009, **544**, 293-305.
- C. F. Majkrzak, N. F. Berk, S. Krueger and U. A. Perez-Salas, *Neutron Scattering in Biology*, 2006, 225-263.
- M. Loesche, *Synth. Microstruct. Biol. Res., [Proc. Int. Conf.]*, 3rd, 1992, 91-108.
- R. S. Armen, O. D. Uitto and S. E. Feller, *Biophys J*, 1998, **75**, 734-744.
- P. Gianni and L. Lepori, *Journal of Solution Chemistry*, 1996, **25**, 1-42.
- W. Kern and D. A. Puotinen, *RCA review; a technical journal*, 1970, **31**, 187-206.
- J. Daillant and A. Gibaud, *X-ray and neutron reflectivity : principles and applications*, Springer, Berlin, 2009.
- 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.*, eds. T. Brückel, G. Heger, D. Richter and R. Zorn, Forschungszentrum Jülich GmbH, Jülich 2007, vol. 38, ch. 5.
- S. K. Sinha, E. B. Sirota, S. Garoff and H. B. Stanley, *Phys Rev B*, 1988, **38**, 2297-2311.
- S. Matlack, A. Koutsoubas and S. Pütter, *Journal of large-scale research facilities JLSRF*, 2015, **1**, A8.
- Y. Khaydukov, O. Soltwedel and T. Keller, *Journal of large-scale research facilities JLSRF*, 2015, **1**, A38.
- Neutron Optics - Instrument Service - Science & Projects - MLZ - Heinz Maier-Leibnitz Zentrum, <http://www.mlz-garching.de/neutron-optics>.
- H. Frielinghaus, A. Feoktystov, I. Berts and G. Mangiapia, *Journal of large-scale research facilities JLSRF*, 2015, **1**, A28.
- G. D. Wignall and F. S. Bates, *J Appl Crystallogr*, 1987, **20**, 28-40.
- V. Pipich and Z. Fu, *Journal of large-scale research facilities JLSRF*, 2015, **1**, A31.
- S. Desert, V. Thevenot, J. Oberdisse and A. Brulet, *J Appl Crystallogr*, 2007, **40**, S471-S473.
- L. G. Parratt, *Physical Review*, 1954, **95**, 359-369.
- G. Fragneto-Cusani, *J Phys-Condens Mat*, 2001, **13**, 4973-4989.
- R. J. Roe, *Methods of X-ray and neutron scattering in polymer science*, Oxford University Press, New York, 2000.
- A. Nelson, *J Appl Crystallogr*, 2006, **39**, 273-276.



45. H. Frielinghaus, *Phys Rev E*, 2007, **76**.
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, *Journal of the Chemical Society, Faraday Transactions 2: Molecular and Chemical Physics*, 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-Us*, 1984, **23**, 5570-5577.
52. L. F. Braganza and D. L. Worcester, *Biochemistry-Us*, 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, *Biochimica et Biophysica Acta (BBA) - Biomembranes*, 1983, **731**, 97-108.
55. M. L. Manca, I. Castangia, P. Matricardi, S. Lampis, X. Fernandez-Busquets, A. M. Fadda and M. Manconi, *Colloid Surface B*, 2014, **117**, 360-367.
56. N. E. Nagel, G. Cevc and S. Kirchner, *Biochimica et Biophysica Acta (BBA) - Biomembranes*, 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. Koutsiousbas, S. Mattauch, M. Ohl, M. Monkenbusch and O. Holderer, *Nuclear Instruments & Methods in Physics Research, Section A: Accelerators, Spectrometers, Detectors, and Associated Equipment*, 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*, eds. R. Lipowsky and E. Sackmann, North-Holland 1995, vol. Volume 1, ch. 3, pp. 97-160.
62. S. Jaksch, F. Lipfert, A. Koutsiousbas, S. Mattauch, O. Holderer, O. Ivanova, H. Frielinghaus, S. Hertrich, S. F. Fischer and B. Nickel, *Phys Rev E*, 2015, **91**.
63. C. Pereira-Leite, C. Carneiro, J. X. Soares, C. Afonso, C. Nunes, M. Lúcio and S. Reis, *European Journal of Pharmaceutics and Biopharmaceutics*, 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*, 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*, 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 Pharmaceut*, 2013, **10**, 4408-4417.
71. S. Lesieur, C. Grabielle-Madelmont, M.-T. Paternostre and M. Ollivon, *Analytical Biochemistry*, 1991, **192**, 334-343.
72. B. Mui, L. Chow and M. J. Hope, *Method 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.