Domain Formation in Model Membranes Studied by Pulsed-Field Gradient-NMR: The Role of Lipid Polyunsaturation

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ABSTRACT   The effects of increased unsaturation in the *sn*-2 fatty acyl chain of phosphatidylcholines (PCs) on the lipid lateral diffusion have been investigated by pulsed-field gradient NMR. Macroscopically oriented bilayers containing a monosaturated PC, egg sphingomyelin, and cholesterol (CHOL) have been studied at temperatures between 0°C and 60°C, and the number of double bonds in the PC was one, two, four, or six. For PC bilayers, with and without the incorporation of egg sphingomyelin and CHOL, the lateral diffusion increased with increasing number of double bonds, as a consequence of the increased headgroup area caused by the unsaturation. Addition of CHOL caused a decrease in lipid diffusion due to the condensing effect of CHOL on the headgroup area. Phase separation into large domains of liquid-disordered and liquid-ordered phases were observed in the ternary systems with PCs containing four and six double bonds, as evidenced by the occurrence of two lipid diffusion coefficients. PC bilayers with one or two double bonds appear homogeneous on the length scales probed by the experiment, but the temperature dependence of the diffusion suggests that small domains may be present also in these ternary systems.

INTRODUCTION

Polyunsaturated fatty acids (PUFA) are essential substances for humans and other higher animals. Lipids containing PUFA are found in very high proportions in neural tissue, and they are important for the development and function of the brain and visual system (1). Apart from these very specialized tissues, in which PUFA may reach 50% of the total fatty acyl chains, PUFA lipids are also found in smaller amounts in other tissues. The levels of PUFA in these membranes can be altered by dietary constraints, and an intake of these fatty acids are considered beneficial to the health. The most studied PUFA, docosohexaenoic acid (DHA), is positively linked to the prevention of an enormous variety of human afflictions, including cancer, heart disease, rheumatoid arthritis, asthma, lupus, alcoholism, and many others (2). For one simple molecule to be able to affect so many seemingly unrelated processes, it seems plausible that its function is not specific but rather that it acts at a fundamental level, common to most cells. This fundamental level, most probably, is connected with the physical properties of the cell membranes, where lipids with PUFA are functioning in the body.

It is well known that the packing of lipids, and therefore their shape, is a very important property for a functioning membrane. This property determines the curvature and elasticity of the membrane. It was shown more than two decades ago that the lipid composition of so-called lamellar- and nonlamellar-forming lipids in the membranes of the bacteria *Acholeplasma laidlawii* and *Escherichia coli* was

regulated to keep an optimal packing or spontaneous curvature for the proteins in the cell membrane (3,4). Similarly, as shown by Brown and co-workers, there is a balance between bilayer-forming and nonbilayer-forming lipids in the native retinal rod membranes, and here the lipids are polyunsaturated with a high content of DHA chains (5). These authors showed that the polyunsaturation has a strong influence on the rhodopsin function, and the molecular packing of wedge-shaped, polyunsaturated lipids was elegantly developed into their ''flexible surface model''.

PUFA is primarily found in the *sn*-2 chain in phosphatidylethanolamine (PE), phosphatidylcholine (PC), and phosphatidylserine (PS), with the *sn*-1 chain being mainly a saturated fatty acid, like palmitic or stearic acid (6). A picture of the way PUFA affects the membrane structure and dynamics is emerging through recent studies of such molecules. Experimental and simulation studies have shown that the DHA chain is extremely flexible, characterized by a high degree of molecular disorder and a rapid interconversion among a diverse set of conformational states (7–9). This is due to a reduced energy barrier for rotational isomerization about the single-C-C bonds that separate the unsaturated carbon atoms in PUFA. Because of this flexibility the DHA chain can rearrange its position in the monolayer to a larger extent than a saturated chain, and both experimental and simulated data show that part of the DHA can even reach up into the polar/apolar interface of the membrane (7,10,11). Also the chain ordering of the adjacent *sn*-1 saturated chain is lowered by the presence of the PUFA in the *sn*-2 chain, and the overall properties of membranes, including chain-melting transition temperature, lipid dynamics, phase behavior, elastic compressibility, permeability, fusion, and flip-flop, are influenced by the introduction of PUFA (2). The effects are largest for the

first and second double bond and then rapidly level off for the introduction of three and more double bonds (2,12).

Another interesting property of polyunsaturated lipids is their possible role in membrane domain formation. There have been a few examples of DHA-induced lipid phase separations in lipid bilayers, in which both liquid-solid and liquid-liquid phase equilibria have been proposed (13). Since cholesterol (CHOL) is believed to be important for the formation of the so-called liquid-ordered (l_o) phase (14,15), several studies have been aimed at the interaction of CHOL with lipids containing PUFA. A general conclusion from these studies is that CHOL interacts only weakly with polyunsaturated chains. This leads to a low solubility of CHOL in dipolyunsaturated lipids, e.g., for di-20:4PC and di-22:6PC it is found to be on the order of 10–15 mol %, whereas CHOL solubility in 18:0–20:4PC and 18:0–22:6PC is on the order of 50%–55% mol % (16). Consequently, the effect of CHOL on membrane properties levels off at low amounts of CHOL for diunsaturated lipids as compared to monosaturated or disaturated lipids (17). The CHOL molecule is also characterized by a larger tilt angle and smaller molecular order in the diunsaturated lipids (16).

A combined ^{2}H and ^{1}H magic angle spinning NMR study showed that CHOL interacts more favorably with the saturated chain in 18:0-X lipids in which X denotes chains of variable unsaturation (9). This leads to the fact that CHOL is most able to order the sn-1 chain and also that the ordering of this chain decreases as the unsaturation of the sn-2 chain increases. Concomitantly, a smaller CHOL-induced area condensation was also observed for all the polyunsaturated phospholipids compared to the monounsaturated PCs.

A molecular dynamics simulation study (11) suggests that the effect of CHOL on the phospholipid acyl chains is highly nonuniform in that the saturated chain shows only a negligible response (essentially a slight increase in the projected length along the bilayer normal), whereas the DHA chain undergoes a significant redistribution of its chain segments from the interfacial region to the hydrophobic core of the bilayer. Thus, a DHA chain can adopt a diverse set of chain conformations with similar (low) energies, whereas a stearic acid has a comparatively reduced conformational freedom.

To our knowledge no direct determination of the lateral diffusion of any polyunsaturated lipids with or without CHOL has so far been published. Polyunsaturated membranes have been proposed to be highly fluid (2), implying a fast lipid translational motion. However, other factors such as acyl chain entanglement and lipid free area will also affect the lateral diffusion in the bilayer. The pulsed-field gradient (pfg) NMR method can be used to measure the lipid lateral diffusion coefficients (D_L) in macroscopically oriented bilayers (18) and, in an attempt to systematically study the effect on D_L of lipid packing and domain formation, several systems containing various lipids and CHOL have been investigated (19–24). These studies are now extended to include PCs with an 18:0 chain in the sn-1 position and with an 18:1, 18:2, 20:4, or 22:6 sn-2 chain.

MATERIALS AND METHODS

Materials

The following substances were used for the preparation of macroscopically aligned lipid bilayers (Fig. 1): 1-stearoyl-2-oleoyl-sn-glycero-3-PC (SOPC), 1-stearoyl-2-linoleoyl-sn-glycero-3-PC (SLPC), 1-stearoyl-2-arachidonoyl-sn-glycero-3-PC (SAPC), 1-stearoyl-2-docosahexaenoyl-sn-glycero-3-PC (SDPC), and egg sphingomyelin (eSM) were obtained from Avanti Polar Lipids (Alabaster, AL) as solutions in chloroform. CHOL and deuterated water (^{2}H$_2$O) were purchased from Sigma (St. Louis, MO).

Preparation of macroscopically oriented bilayers

Macroscopically oriented bilayers were prepared after a procedure reported earlier (18). Appropriate amounts of lipids were dissolved in a mixture of methanol and propanol (1:4 vol) at a concentration of 15 mg/ml. The solution was deposited on glass plates, the solvent was evaporated, and the plates were placed under high vacuum overnight to remove traces of solvent.
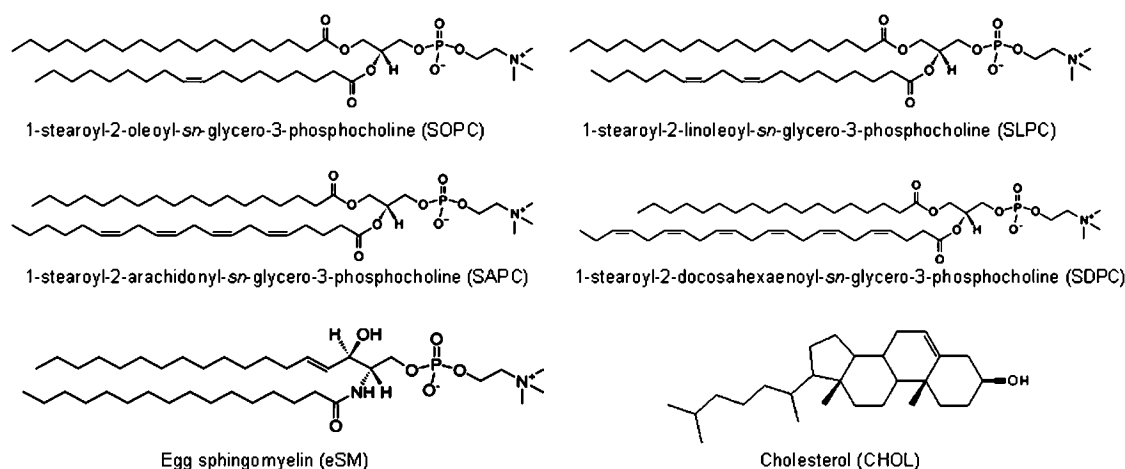

FIGURE 1   Structures for the lipids used in this study.

The plates were then stacked and placed in a glass tube with a square cross section (~30 plates/sample). The sample tube was placed for several days in a humid atmosphere at room temperature. During this time, hydrated and oriented bilayers were formed. After addition of water in excess, the tube was sealed and the sample was left for several hours for final equilibration. Sample orientation was checked with crossed polarizers. Because of the high sensitivity of polyunsaturated lipids to oxidation, all handling of the lipids was performed in a dry nitrogen gas atmosphere. Water used for hydration was depleted of dissolved oxygen by nitrogen gas bubbling.

Pulsed-field gradient NMR technique

The diffusion measurements were performed on a Chemagnetics Infinity (Varian, Fort Collins, CO) NMR spectrometer operating at a proton frequency of 100 MHz and equipped with a specifically designed goniometer probe that enabled macroscopically aligned bilayers to be oriented with the bilayer normal at the magic angle (54.7°) with respect to the main magnetic field. This causes the dipolar interactions to vanish, resulting in a significant reduction of the line width. Details of the pfg-NMR method for measurements of lipid lateral diffusion on macroscopically oriented bilayers can be found elsewhere (18). For all measurements the stimulated spin-echo pulse sequence was used (25). The diffusion decay of the echo amplitude, A, can be described by the equation

$$A = \sum_i \frac{A_i}{2} \exp\left(\frac{-2\tau}{T_{2i}} - \frac{\tau_1}{T_{1i}} - \gamma^2 \delta^2 g^2 D_i \left(\Delta - \frac{\delta}{3}\right)\right), \quad (1)$$

where the summation index goes over all diffusion components present, T_1 and T_2 are the longitudinal and transverse proton NMR relaxation times, γ is the gyromagnetic ratio, Δ is the time interval between two identical gradient pulses (equal to the diffusion time), δ and g are the duration and amplitude of the pfgs, respectively, and D is the self-diffusion coefficient. τ and τ_1 are times in the pulse sequence that govern the T_2 and T_1 relaxation intervals.

In our experiments $g = 1.15 T/m$ and δ was varied in the range 1–9 ms in 10–32 steps and all other variables were kept constant. The ranges of τ and τ_1 were 11 and 50–200 ms, respectively. The resulting D was not dependent on the choice of these parameters. Signal accumulations from 32 to 160 were

made to obtain an acceptable signal/noise level. Experimental data were Fourier transformed into a set of spectra that was analyzed with the component resolved method to extract the diffusion coefficient. This method globally fits all frequency channels and results in spectral shapes and diffusion coefficients for the individual components of the data (26). The lateral diffusion coefficient D_L was calculated as $1.5D$ due to the angle between the pfg and the main magnetic field (18).

The observation of domain formation is based on the difference in D_L and the apparent activation energy for the diffusion process in the l_d and l_o phases (27). If the domains are large enough to make the lifetime within the domains long compared to the diffusion time, two separate D_Ls will be observed. The size of the domains typically needs to be larger than 1 μm for this to occur. If the domains are smaller than this, a weighted average of the D_Ls in the two phases will be observed.

RESULTS AND DISCUSSION

PC and PC/CHOL bilayers

The diffusion decays for the lipid systems containing either PC or PC/CHOL were biexponential; the faster component corresponds to water. This component has not been analyzed further. The slow component, corresponding to lipids, gave NMR spectra with a characteristic appearance of PCs with the most prominent peaks at 0.9 ppm (ω-CH$_3$), 1.1 ppm (chain -CH$_2$-), and 3.1 ppm (choline-CH$_3$). For higher degrees of unsaturation, a signal from the -CH=CH- protons could also be seen at 5.2 ppm. (The chemical shifts are only approximate but are in good agreement with MAS NMR spectra of saturated and polyunsaturated chains (7,28).) Spectra for samples with varying CHOL content are shown in Fig. 2. No signal from CHOL could be observed, because of the fast transverse NMR relaxation of the CHOL protons. Note the larger decrease in signal intensity for the peaks arising from the lipid chains as the CHOL content increases. This is an
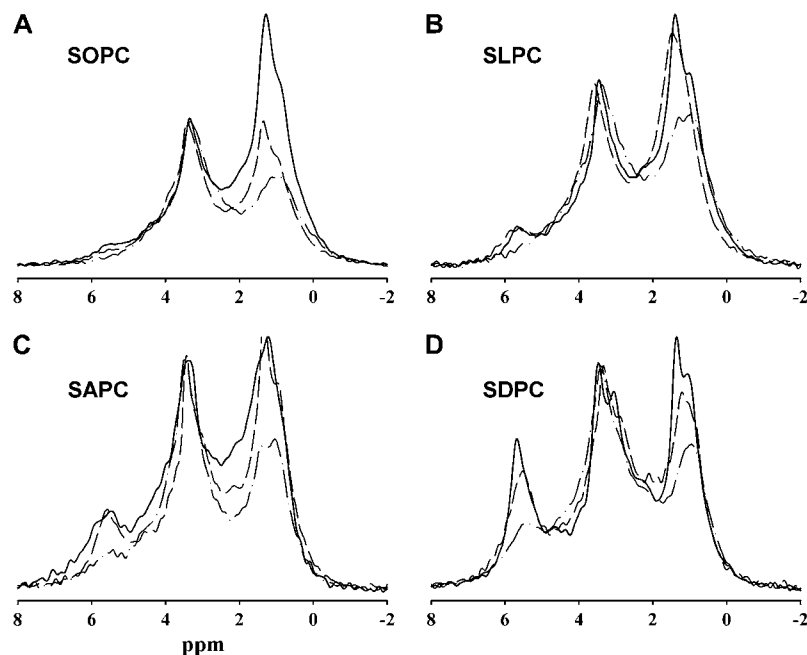

FIGURE 2 ^{1}H-NMR spectra at 21°C from oriented samples of the binary systems PC/CHOL, obtained from the pfg-NMR experiments after water suppression. (*Solid lines*) 0% CHOL, (*dashed lines*) 4% CHOL, (*dash-dotted lines*) 40% CHOL. (*A*) SOPC, (*B*) SLPC, (*C*) SAPC, (*D*) SDPC.

effect of the ordering of the chains caused by CHOL, increasing the T_2 relaxation rate.

The influences of the CHOL concentration and temperature on D_L are shown in Fig. 3. An increase in unsaturation from SOPC to SDPC leads to an increase in D_L at all CHOL concentrations, and an increase in the CHOL content results in a decrease in D_L for all phospholipids. D_L increases with temperature for all the PCs.

The effect of unsaturation can be rationalized if the effect of packing of the lipids on D_L is considered. Vaz et al. (29) and Almeida et al. (30) showed that a free area theory of Cohen and Turnbull (31) could be used to describe the diffusion process in lipid bilayers. This theory considers a particle performing a random walk in two dimensions. Each elementary step in this process is limited by the occurrence of a free volume (or free area a_f) greater than a critical size (a^*) next to the diffusing particle. This leads to the following relation between the lateral diffusion coefficient and the free area (32):

$$D_L = D^* \exp\left(\frac{-\beta a^*}{a_f}\right), \qquad (2)$$

where D^* is a constant, and β is a factor to correct for overlapping free volumes (typically in the range 0.5–1). The free area can be estimated as $a_f = a_{av} - a^*$ where a_{av} is the average area for the molecule in the bilayer. The values of a_{av} for monolayers (12) and bilayers (33) are quite similar, and monolayer data are used here. When data are fit to Eq. 2, values for a^* between 43 and 48 Å^2 are found depending on the chosen value of β. Fig. 4 shows the results using $\beta = 1$. a^* is usually taken to be the van der Waals, area and the value of ~40 Å^2 seems reasonable for the PC molecules. For

example, the area per molecule obtained in the limit of high CHOL content and high lateral pressure in monolayers at 25°C is found to be close to 40 Å^2 for all lipids (12). The good fit to Eq. 2 and the reasonable magnitude for the fitting parameters show that the increase in D_L with increasing unsaturation for our systems is consistent with the increasing area per molecule with increasing unsaturation seen in monolayers (12) and bilayers (33).

It can be inferred from Fig. 3 that the dependence of D_L on the CHOL concentration is approximately linear for SOPC, SLPC, and SAPC. Such a linear dependence has been observed for binary systems with 1-palmitoyl-2-oleoyl-PC and dioleoyl-PC with CHOL (19), which are generally believed to form only homogeneous phases. For SDPC the dependence is more complicated. For CHOL contents smaller than 10% and larger than 25%, D_L varies very little, whereas in the interval 10%–25%, D_L has quite a strong linear dependence on the CHOL content. Such a behavior was also observed in the binary system of eSM/CHOL (19), where it was interpreted as evidence of the formation of microdomains. However, in this system it is difficult to conclude that l_o domains are formed, since the activation energies more closely conform to those usually found in the l_d phase (vide infra). At low CHOL contents the effect is complicated by the fact that there may be two opposing events. In general, the condensing effect of CHOL will decrease the lipid-free volume in the bilayer, causing a decrease in D_L. However, small additions of CHOL can also result in an increase in D_L caused by the reduced entanglement of the lipid chains (34,35). These two effects might balance out to give a constant D_L at low CHOL concentrations. The reason for the constant value of D_L at high CHOL concentrations is not clear. It seems improbable that
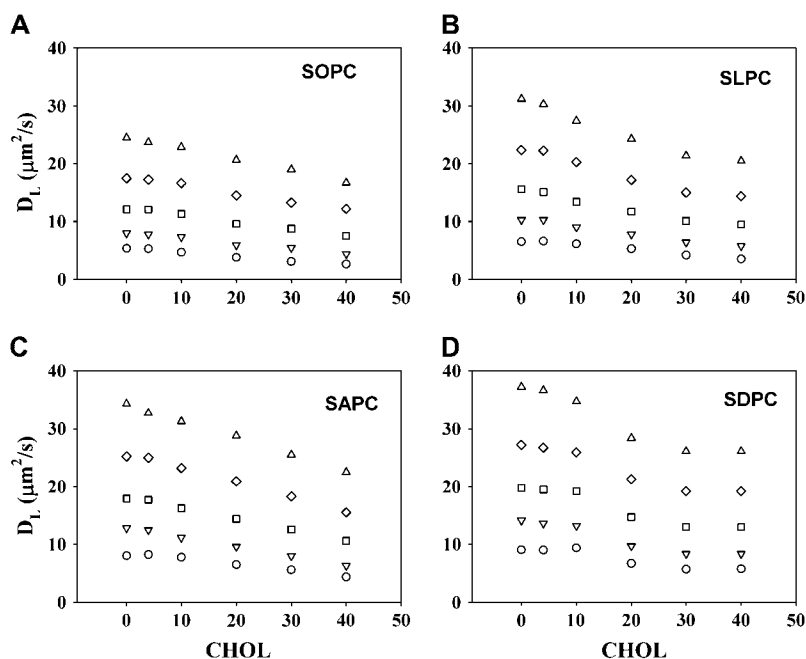

FIGURE 3 Lateral diffusion coefficients as a function of CHOL content for the temperatures 21°C (*circles*), 30°C (*triangles down*), 40°C (*squares*), 50°C (*diamonds*), and 60°C (*triangles up*) for the binary systems of PC/CHOL. (*A*) SOPC, (*B*) SLPC, (*C*) SAPC, (*D*) SDPC.
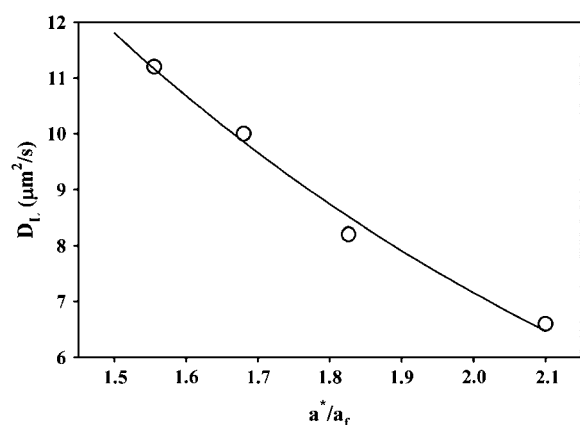
FIGURE 4   Lateral diffusion versus free area for the PC bilayers. The line is the best fit to Eq. 2 with β set to 1. The corresponding values of D^* and a^* are 53 $\mu m^2/s$ and 43 $\mathring{A}^2$, respectively.

the reason is poor solubility of CHOL, since the solubility limit in SDPC bilayers is reported to be 55% (36). More studies are needed to get an understanding of this feature.

The temperature dependence of D_L gives a straight line in an Arrhenius plot, from which an apparent activation energy (E_A) for the diffusion process can be calculated. Fig. 5 shows the influence of the CHOL concentration on E_A. It is found that an increase in lipid unsaturation generally results in a decrease in E_A. This was also observed in an earlier study (19). Moreover, up to a CHOL content of 10 mol % E_A is almost constant, whereas between 10–40 mol % of CHOL E_A increases approximately linearly. However, the increase is rather moderate and the values of E_A are all in the range of what is obtained for other systems in the l_d phase (19,20). Taken together, the data for the binary systems indicate that the bilayers are in a homogeneous l_d phase, although we cannot exclude the existence of small amounts of l_o phase.
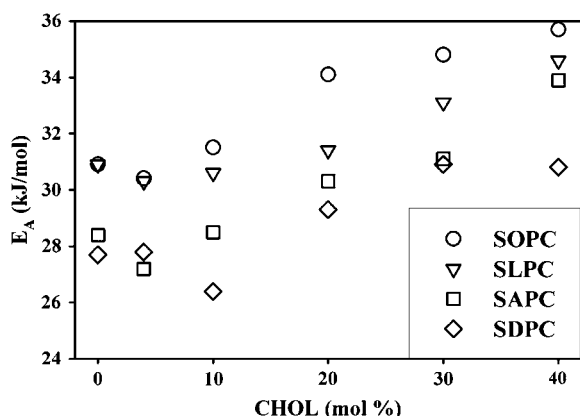

FIGURE 5   Apparent activation energies for the diffusion process in the binary systems of PC/CHOL as a function of the CHOL content.

PC/eSM/CHOL bilayers

The compositions of the bilayers were 37.5/37.5/25 mol % of PC/eSM/CHOL. The temperature dependence of the diffusion coefficients for the ternary systems are shown as circles in Fig. 6 and as Arrhenius plots in Fig. 7. For comparison, data from the binary systems are included in Fig. 6 as solid and dashed lines. Only a single lipid diffusion coefficient is observed in the SOPC and SLPC ternary systems (Figs. 6 and 7, *A* and *B*) with D_Ls intermediate between those observed for the binary systems of eSM/CHOL and PC/CHOL. This "averaging" of the diffusion coefficients in mixed bilayers has been observed previously in other systems (23,37,38) and reflects the fact that the lateral diffusion is governed by the physicochemical properties of the bilayer as a whole and not by the properties of the individual lipids. The bilayers are thus homogeneous in these two systems at all temperatures on the timescale of the diffusion experiment (50–200 ms), i.e., possible domains must be smaller than ~1 μm.

The ternary systems with SAPC and SDPC show a different behavior (Figs. 6 and 7, *C* and *D*), and the diffusion decay for the lipids can only be sufficiently described by a biexponential decay, giving two separate diffusion coefficients. According to observations in similar systems (20,21,24,27), we interpret this finding as a result of a phase separation into the l_d and l_o phases. D_L is ~2–10 times slower in the l_o phase, with the largest difference at low temperatures.

It is difficult to determine the composition of the two phases from the diffusion data, since the signals from the phospholipids overlap and no signal is observed from CHOL. Furthermore, since the relaxation rates differ in the two phases, no direct measure of the relative amounts of the phases can be made (see Eq. 1). However, it is possible to make an estimate of the composition of the two phases from the magnitude of the observed D_Ls and the appearance of the spectra of the two components. We assume that the l_o phase is enriched in the saturated lipid, whereas the l_d phase is enriched in the unsaturated lipid. This is in accordance with phase diagrams in similar systems (23,39,40). Thus, if it is assumed as a first approximation that all eSM goes into the l_o phase and all PC goes into the l_d phase, it is possible to estimate the CHOL content in the phases by comparison with the binary systems. If we compare D_L for the l_d phase with data from the binary PC/CHOL systems, we find that it corresponds to a CHOL content of 30%–40%. This gives a CHOL content in the l_o phase of 20% or less. The corresponding comparison of D_L for the l_o phase to those in the eSM/CHOL system indicates that very little CHOL is located in the l_o phase. This finding is rather surprising since CHOL generally is found to be slightly enriched in the l_o phase.

A plausible explanation is that small amounts of eSM are found in the l_d phase and some of the unsaturated PC goes into the l_o phase. Thus, the diffusion would be slowed down in the l_d phase and speeded up in the l_o phase and a direct comparison with the binary systems will overestimate the
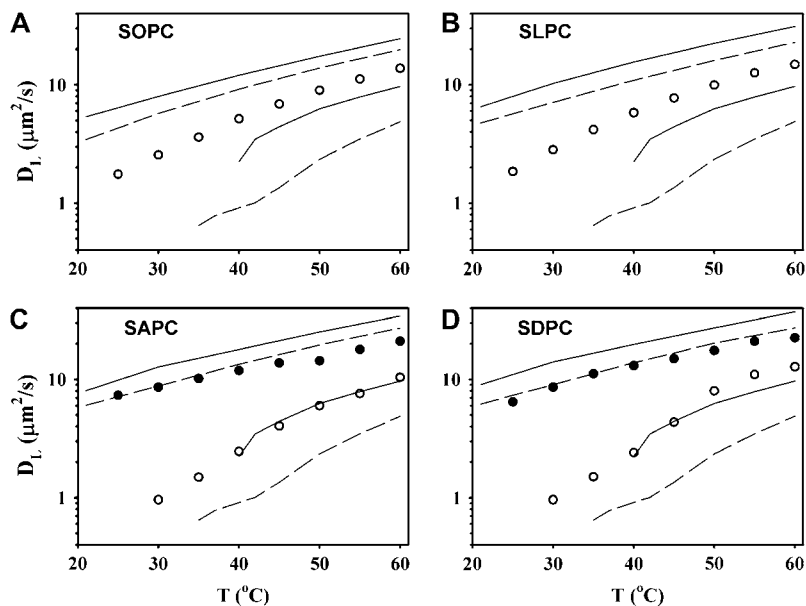
FIGURE 6 Lateral diffusion coefficients as a function of temperature for the ternary systems of PC/eSM/CHOL. The circles indicate diffusion in the ternary systems, and the lines are taken from the binary systems for comparison. (*Solid lines*) PC (*upper*) and eSM (*lower*) diffusion in the pure lipid systems. (*Dashed lines*) PC (*upper*) and eSM (*lower*) diffusion interpolated to a CHOL content of 25% for the binary systems of PC/CHOL and eSM/CHOL, respectively. The values for the eSM systems are taken from Filippov et al. (19).

amount of CHOL in the l_d phase. If we look at the spectra obtained for the l_o phase (Fig. 8, *C* and *D*), no peak from the double bond region at 5.2 ppm is seen. This is an indication that the amount of PC in this phase is low, since this peak is visible in the binary spectra for CHOL concentrations up to 40% at this temperature. Furthermore, a comparison of the spectrum for the l_d phase with those found for the binary PC/CHOL systems reveals that it is similar in appearance as that for 20%–30% CHOL. Thus, the amount of CHOL is substantial in the l_d phase, and CHOL seems to partition fairly equally into the two phases, with only a smaller preference for the l_o phase. This proposal is in agreement with tie-lines found in dioleoyl-PC/dipalmitoyl-PC/CHOL

systems (40), whereas other investigations indicate a more distinct partition of CHOL into the l_o phase (41,42). To further investigate the partitioning issues, we are currently planning measurements with deuterated palmitoyl-SM as well as deuterated CHOL.

Further insight into the nature of the two phases is given by the temperature dependence of D_L. Previous studies have shown that E_A is significantly larger for the l_o phase compared to the l_d phase (19,20), making it possible to discriminate between the two phases. For the SAPC and SDPC systems E_A is found to be 24 and 29 kJ/mol, respectively, for the l_d phase, whereas it is much higher (68 and 78 kJ/mol) for the l_o phase (Fig. 7). For the SOPC and SLPC systems E_A is found to be
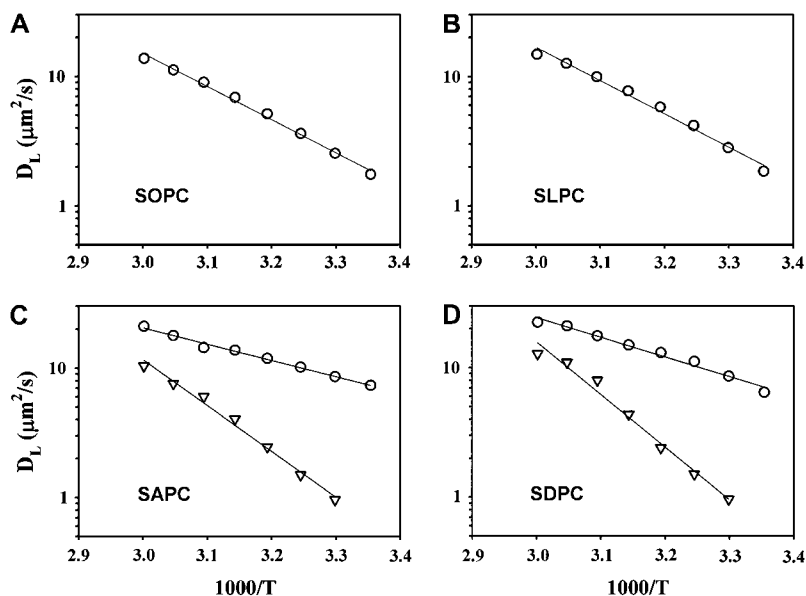

FIGURE 7 Temperature dependence of D_L obtained for the ternary systems of PC/eSM/CHOL. The lines are best fits to the Arrhenius equation and give the following apparent activation energies: (*A*) SOPC: $E_A = 49$ kJ/mol, (*B*) SLPC: $E_A = 49$ kJ/mol, (*C*) SAPC: $E_A = 24$ (l_d) and 68 (l_o) kJ/mol (4), (*D*) SDPC: $E_A = 29$ (l_d) and 78 (l_o) kJ/mol.
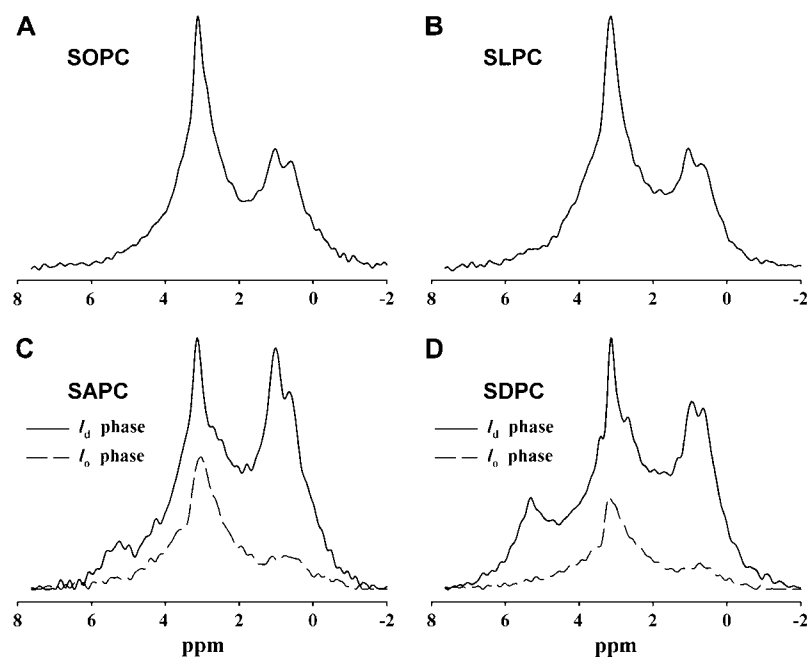
FIGURE 8 Spectra obtained from the component resolved analysis of the pfg-NMR diffusion experiments at 40°C for the ternary systems of PC/eSM/CHOL.

intermediate between these two extremes: 49 kJ/mol for both systems. This may indicate that these systems also form the l_o phase but that the domains formed are small enough to allow for a lipid exchange between the two phases that is fast on the timescale considered for the diffusion experiment. Then, the observed D_L will be an average of the lateral diffusion in the two phases and, provided that the amounts of the two phases do not change significantly with temperature, E_A will also be averaged.

Domain formation in ternary systems

A large number of ternary systems has been investigated with regard to domain formation and they all have common features, i.e., they contain one saturated and one unsaturated lipid, together with a sterol. Systematic pfg-NMR studies have revealed that each of these three components are essential for domain formation and that small changes in the structure of the components can have a large impact on the domain-forming process (20,21,23,24). It has been found that the homogeneity of the SM compound was of crucial importance for the domain formation and that small changes in the structure of the sterol also strongly could affect the process. It was therefore interesting to see how the degree of unsaturation influences the lateral phase separation. Domain formation has been observed for the diunsaturated DOPC with eSM and CHOL, whereas for the monounsaturated POPC it is less clear whether large domains are formed or not.

There are large differences in the two phase diagrams that have been reported (39,41), and pfg-NMR methods have been unable to detect domains in this system (G. Orädd and G.

Lindblom, unpublished results). This discrepancy probably originates from differences in spatial resolution of the methods. Fluorescence microscopy and pfg-NMR rely on large ($>\mu$m) domains, whereas fluorescent anisotropy measurements can detect smaller domains. For the similar systems SOPC/eSM/CHOL and SLPC/eSM/CHOL, we find no evidence for large domains, although the temperature dependence of D_L in these systems could be taken as evidence for the existence of small domains. However, increasing the degree of unsaturation in the lipid chains results in phase separation into the l_d and l_o phases for both the SAPC/eSM/CHOL and the SDPC/eSM/CHOL systems. Therefore, systems that are on the verge of forming larger domains can be triggered into this behavior by an increase in the number of double bonds in monounsaturated PCs.

CONCLUSIONS

We have investigated the effect of increased unsaturation in the fatty acyl chain of PCs on the lateral diffusion in bilayers containing a monosaturated PC, eSM, and CHOL. It can be concluded that an increase in the number of double bonds in the sn-2 chain of the monosaturated PC results in an increase in the free area of the bilayers. For the one-component PC bilayers, this leads to an increase in the lateral diffusion in accordance with theoretical predictions (Eq. 2). It also influences other bilayer properties, such as the lateral compressibility, bending rigidity, and lipid chain localization. Addition of CHOL to such systems will have an ordering effect, primarily on the saturated chain but also on the unsaturated chain. As a consequence of this, the free area in the bilayers

will decrease, resulting in a decreased lateral diffusion. In the ternary systems of PC/eSM/CHOL the ordering by CHOL on eSM will be more effective than on PC, and a lateral phase separation into l_d and l_o phases is observed for the most unsaturated PCs. In these systems the l_o phase contains mostly eSM, whereas the l_d phase is enriched in the unsaturated PC. We propose that the phase separation is driven by the increasing difficulty of incorporating an unsaturated lipid into the highly ordered matrix formed by eSM and CHOL.

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