

## Comment to the Editor

### Lipid Diffusion, Free Area, and Molecular Dynamics Simulations

In a recent article in *Biophysical Journal*, Falck et al. (2004) present a molecular dynamics (MD) study of phosphatidylcholine (PC)/cholesterol (Chol) bilayers, focusing on the lipid packing and its relation to free area and lateral diffusion of lipids. A significant comparison is made between their MD results and our experimental diffusion measurements using fluorescence recovery after photobleaching (FRAP) and the analysis of those experiments using the free area model (Almeida et al., 1992). In contrast to our conclusion, Falck et al. state that the free area model does not quantitatively represent lipid diffusion. Furthermore, their simulations predict a much stronger effect of cholesterol on diffusion than found experimentally. In our opinion, as written, some of the statements of Falck et al. are prone to misinterpretation. The criticism of the free area model based on the MD simulations is flawed because comparison is made between different systems. If the proper systems are compared, the free area model actually predicts the correct result for lipid diffusion, whereas the MD simulations do not. In this Comment to the Editor, we present our views on the problem and clarify several of the issues. Finally, we attempt to resolve the apparent quantitative disagreement between the MD simulations and our experiments on lipid diffusion.

#### Lipid diffusion in phospholipid/cholesterol systems

A brief summary of experimental diffusion measurements and the phase diagram of phospholipid/cholesterol mixtures is useful to follow this discussion. Above the main phase transition temperature ( $T_m$ ) of the phospholipid, its mixtures with cholesterol result in a phase diagram with three different regions: if the mol fraction of Chol ( $\chi_{cho}$ ) is  $< \sim 0.1$  (depending on temperature), the system is in an liquid-disordered ( $\ell_d$ ) phase; above  $\chi_{cho} \approx 0.30$ , the system is in a liquid-ordered ( $\ell_o$ ) phase (which may also be called a phospholipid/cholesterol condensed complex region; McConnell and Radhakrishnan, 2003); and, in between those Chol concentrations,  $\ell_d$  and  $\ell_o$  phases coexist (Shimshick and McConnell, 1973; Ipsen et al., 1987; Sankaram and Thompson, 1990b; Vist and Davis, 1990; Almeida et al., 1992).

In our work (Almeida et al., 1992), the lateral diffusion coefficient of the phospholipid ( $D_L$ ) in dimyristoylphosphatidylcholine (DMPC)/Chol mixtures drops by a factor of 2.2

from the  $\ell_d$  phase DMPC, in the absence of cholesterol (Vaz et al., 1985), to the  $\ell_o$  phase with at least  $\chi_{cho} = 0.30$  (Almeida et al., 1992). This is in agreement with measurements by different investigators, over two and half decades, using different techniques, as shown in Table 1, which emphasizes DMPC/Chol because this is the system we examined. For the phospholipid/cholesterol systems listed here, the ratio of  $D_L$  in  $\ell_d$ -phase phospholipid to  $D_L$  in  $\ell_o$ -phase phospholipid/cholesterol is always between 2 and 4, with an average value of  $2.7 \pm 0.7$ . Koriach et al. (1999) also report a measurement at  $\chi_{cho} = 0.60$  in dilauroylphosphatidylcholine (DLPC)/Chol. This measurement differs from the data reported by other investigators on other PC/Chol systems in that it is the only one that shows a significant decrease in  $D_L$  in the  $\ell_o$  phase, upon increase in the cholesterol content beyond  $\chi_{cho} = 0.30$ .

In our measurements,  $D_L$  does not decrease with Chol content in the  $\ell_d$  phase; a significant decrease is only observed when the  $\chi_{cho}$  is high enough for the system to enter the  $\ell_d$ - $\ell_o$  coexistence region (Almeida et al., 1992). That lack of dependence of  $D_L$  on  $\chi_{cho}$  in the  $\ell_d$  phase is also apparent in the data from McConnell's group (Rubenstein et al., 1979) for DMPC/Chol and egg PC/Chol, and in the data of Filippov et al. (2003) for chicken egg sphingomyelin (SM)/Chol. The onset of the decrease of  $D_L$  with  $\chi_{cho}$  agrees well with the two-phase boundary of the phase diagram for DMPC/Chol (Almeida et al., 1992) and SM/Chol (Filippov et al., 2003). Data of Filippov et al. (2003) on DMPC/Chol and dioleoylphosphatidylcholine (DOPC)/Chol, on the other hand, show a monotonic decrease of  $D_L$  with Chol concentration even in the  $\ell_d$  phase. These data of Filippov et al. (2003) appear to be also in conflict with earlier measurements by the same group in DOPC/Chol (Lindblom et al., 1981). Filippov et al. (2003) discuss these apparent discrepancies.

#### Use of free area theory to analyze lipid diffusion

Falck et al. (2004) conclude their introduction by stating that free area theories correctly predict a reduction in diffusion caused by the addition of cholesterol to a PC membrane, "but are not applicable to quantitatively describing lateral diffusion in lipid bilayers." Without qualification, this statement appears to apply both to MD simulations and experimental data. Further, in their discussion they add: "In our opinion, one should at least not expect free area theory to yield quantitative results." These statements are contrary to our conclusions. First, Vaz et al. (1985) showed that the free area model can indeed fit the experimental diffusion data quantitatively, when a change in free volume is caused by

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**TABLE 1** Comparison of diffusion coefficients in the  $\ell_d$  and  $\ell_o$  phases for a few phospholipid/cholesterol systems

System	$X_{\text{cho}}$	$T$ (°C)	Phase	$D$ ( $\text{cm}^2\text{s}^{-1}$ )	Ratio	Method	Reference
DMPC	0	35	$\ell_d$	$7.5 \times 10^{-8}$	—	FRAP	Rubenstein et al. (1979)
DMPC/Chol	$\geq 0.30$	35	$\ell_o$	$3.0 \times 10^{-8}$	2.5	—	—
DMPC	0	26	$\ell_d$	$6.0 \times 10^{-8}$	—	FRAP	Alecio et al. (1982)
DMPC/Chol	$\geq 0.30$	26	$\ell_o$	$1.8 \times 10^{-8}$	3.3	—	—
DMPC	0	35	$\ell_d$	$7.6 \times 10^{-8}$	—	FRAP	Vaz et al. (1985)
DMPC/Chol	$\geq 0.30$	34	$\ell_o$	$3.5 \times 10^{-8}$	2.2	—	Almeida et al. (1992)
DLPC	0	25	$\ell_d$	$3 \times 10^{-8}$	—	FCS	Korlach et al. (1999)
DLPC/Chol	0.30	25	$\ell_o$	$1 \times 10^{-8}$	3	—	—
DMPC	0	35	$\ell_d$	$11 \times 10^{-8}$	—	pfg-NMR	Filippov et al. (2003)
DMPC/Chol	0.33	35	$\ell_o$	$3 \times 10^{-8}$	4	—	—
SM	0	55	$\ell_d$	$8 \times 10^{-8}$	—	pfg-NMR	Filippov et al. (2003)
SM/Chol	0.30–0.425	55	$\ell_o$	$3.5 \times 10^{-8}$	2.3	—	—
DOPC	0	30	$\ell_d$	$10 \times 10^{-8}$	—	pfg-NMR	Filippov et al. (2003)
DOPC/Chol	0.33	30	$\ell_o(?)$	$5 \times 10^{-8}$	2	—	—

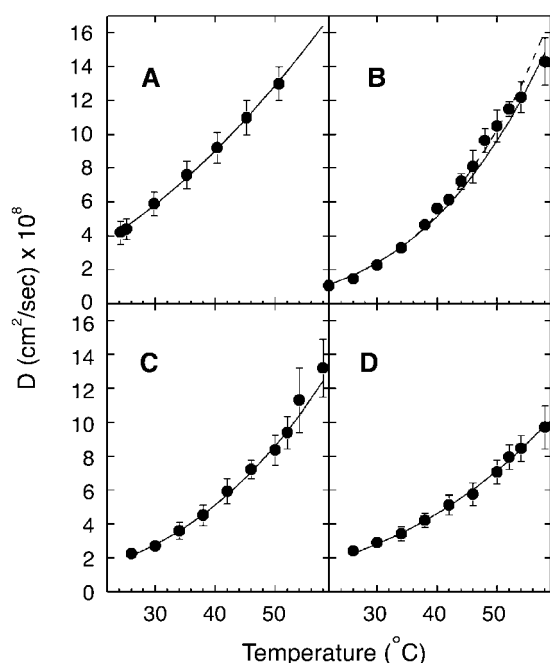
a change in temperature. Second, Almeida et al. (1992), extended this study to the effect of cholesterol, as an obvious way of changing the free volume, using a version of the Macedo-Litovitz equation (Eq. 1) (Macedo and Litovitz, 1965) for the free volume theory (Cohen and Turnbull, 1959; Turnbull and Cohen, 1961, 1970). In fact, the agreement between free area theory and experiment is not only qualitative but also quantitative, as shown in Fig. 1, which presents data and theoretical curves for diffusion in pure DMPC and DMPC/Chol ( $\chi_{\text{cho}} = 0.30, 0.40$ , and  $0.50$ ) as a

function of temperature (Almeida et al., 1992). The theoretical curves were obtained by fitting the equation

$$D_L = A(a^{1/2}) e^{-a_0/a_f} e^{-E_a/kT}, \quad (1)$$

to the experimental data using a simple least-squares analysis. The only free parameters in this procedure were the average cross-sectional, hard-core area (equivalent to the van der Waals volume) of cholesterol ( $a_0^{\text{cho}}$ ), which was subtracted from the total area to obtain the free area ( $a_f$ ), and the activation energy ( $E_a$ ). The preexponential factor depends on the square root of the area over which diffusion occurs as described in detail by Almeida et al. (1992). We found that a value for  $a_0^{\text{cho}} = 26.6 \text{ \AA}^2$ , for all DMPC/Chol compositions examined ( $\chi_{\text{cho}} = 0.30, 0.40$ , and  $0.50$ ), yielded very good agreement with the experimental data; and that  $E_a = 2.7$  kcal/mol for pure DMPC, and 1.9, 2.1, and 2.5 kcal/mol for DMPC/Chol 70:30, 60:40, and 50:50, respectively, gave the best fits. It is interesting that the value of  $26.6 \text{ \AA}^2$  is exactly the same as determined recently by MD simulations, which yielded  $27 \pm 1 \text{ \AA}^2$  (Hofsaß et al., 2003; Khelashvili and Scott, 2004). Such nearly perfect agreement is certainly fortuitous, but it indicates that the value we arrived at is entirely reasonable.

However, Falck et al. (2004) write, “it seems reasonable to expect that  $E_a$  should increase with cholesterol content. Experimental results (Almeida et al., 1992) do support this idea but are partly contradictory. This is, however, probably due to the fitting procedure used.” First, we cannot discern any contradiction in our experimental results: there are no two measurements of the same observable that yield different values. If we understand Falck et al. correctly, they are not questioning the fitting method itself (least squares) but the possible effects of the preexponential factor and the use of constant (temperature-independent) values for the hard-core, cross-sectional areas of DMPC ( $a_0^{\text{pc}}$ ) and cholesterol ( $a_0^{\text{cho}}$ ). In our work,  $a_0^{\text{pc}}$  was taken as the close-packed area per lipid in gel state DMPC ( $45 \text{ \AA}^2$ ) and  $a_0^{\text{cho}}$  was an adjustable parameter in the fits. Slightly different choices for the



**FIGURE 1** Diffusion coefficients ( $D_L$ ) in DMPC (A), and DMPC/Chol 70:30 (B), 60:40 (C), and 50:50 (D) measured by FRAP. The lines are fits of the equations derived from free area theory (Eq. 1) to the data. Reprinted with permission from Almeida et al. (1992). Copyright 1992 American Chemical Society.

preexponential factor have a very minor effect because of its weak (square root) dependence on the area. Reasonable variations in this term lead to no more than  $\sim 0.1$  kcal/mol changes in the values obtained for  $E_a$  for the different systems that we examined and are insufficient to alter the ranking of  $E_a$  values in these four mixtures. Indeed, we noted that  $E_a$  for DMPC and DMPC/Chol 50:50 are essentially the same (Almeida et al., 1992). With the same  $E_a$ , it was then suggested that the effect of cholesterol content in the  $\ell_o$  phase (50:50 mixture) should be equivalent to a shift in temperature, which was shown to be the case (Fig. 2). The meaning of a minimum in  $E_a$  for a mixture with  $\chi_{\text{cho}} = 0.3$  deserves some discussion, which is postponed until the next section. Before concluding this summary we should note that the data of Filippov et al. (2003) yield a different temperature dependence as a function of cholesterol, with a monotonically increasing  $E_a$  with cholesterol content in DMPC/Chol. However, their  $E_a$  values were obtained with simple Arrhenius plots, not with Eq. 1; therefore, in this form, they cannot be directly compared with ours.

### MD simulations, free area, and the timescale of diffusion

The MD simulations of Falck et al. (2004) and their analysis by “slicing” the bilayer at various levels along the normal have made clear an important difficulty of free area theory when applied to diffusion in lipid bilayers: the fact that the free volume is not constant across the bilayer height. Therefore, the assignment of a value to  $a_o^{\text{pc}}$  and  $a_o^{\text{cho}}$  is

conceptually complicated. Free area theory defines an average cross-sectional area for a lipid and treats diffusion in the bilayer as two-dimensional, which is not strictly correct. Movement of the lipids into free volumes at different levels along the bilayer normal would reasonably be expected to contribute to diffusion, probably softening the barriers to displacements along the bilayer plane. Essentially, the free area model ignores all these complications and  $a_o^{\text{pc}}$  and  $a_o^{\text{cho}}$  are then, to some extent, operational parameters. Apart from that conceptual difficulty with free area theory (which we share), Falck et al. appear to have two major problems when comparing their MD simulations with our experiments and their interpretation using free area theory: 1), the magnitude of the decrease in the diffusion coefficient when Chol content is increased from  $\chi_{\text{cho}} \approx 0$  to  $\chi_{\text{cho}} \approx 0.30$ ; and 2), our observation of a minimum in  $E_a$  for DMPC/Chol 70:30, compared to pure DMPC and DMPC/Chol 60:40 and 50:50.

With regard to the first problem, Falck et al. base their conclusions on the fact that their MD simulations show a reduction of a factor of 10 in the lipid diffusion coefficient ( $D_L$ ) when  $\chi_{\text{cho}}$  is increased from 0.047 to 0.297 (Falck et al., 2004). However, if free area theory were correct, they calculate that  $D_L$  should be reduced by a factor of 3 at most. As shown above, in experimental measurements, the effect of cholesterol content on  $D_L$  is consistently a reduction by a factor of 2–3, up to  $\chi_{\text{cho}} \approx 0.50$ , for all measurements including those of Korlach et al. (1999) for  $\chi_{\text{cho}} = 0.30$ . Therefore, experimentally, the ratio of  $D_L$  in the  $\ell_d$  to the  $\ell_o$  phase agrees very well with the prediction of free area theory, as estimated by Falck et al. (2004). In support of their MD results, Falck et al. cite a measurement by Korlach et al. (1999) in DLPC/Chol using fluorescence correlation spectroscopy (FCS), which gives a reduction of 10 upon addition of cholesterol to a final  $\chi_{\text{cho}} = 0.60$ . This decrease is a feature of the data of Korlach et al. (1999) for  $\chi_{\text{cho}} = 0.60$ , but has not been observed by other investigators. To the best of our knowledge, all other studies have shown that above  $\chi_{\text{cho}} = 0.30$  the lipid diffusion coefficient does not vary much. In any case, the comparison made by Falck et al. (2004) was with MD simulations of dipalmitoylphosphatidylcholine (DPPC)/Chol containing  $\chi_{\text{cho}} = 0.297$ , so the relevant experimental data are those for  $\chi_{\text{cho}} = 0.30$ , not 0.60; and at  $\chi_{\text{cho}} = 0.30$   $D_L$  is reduced by a factor of 3 (as predicted by the free area model), not 10 (as predicted by the MD simulations).

Why then do the MD simulations show a stronger effect of Chol on  $D_L$ ? One possibility is that this is due to the timescale of the simulations, which was 100 ns (Falck et al., 2004). This is a long time for an MD simulation, but is still short for diffusion. With a typical  $D_L = 5 \times 10^{-8} \text{ cm}^2 \text{ s}^{-1}$ , the area explored by a lipid molecule in 100 ns is  $200 \text{ \AA}^2$ , which is only three times the average cross-sectional area per phospholipid in a fluid bilayer,  $\sim 65 \text{ \AA}^2$ . Another way of looking at the problem is that this timescale allows only for

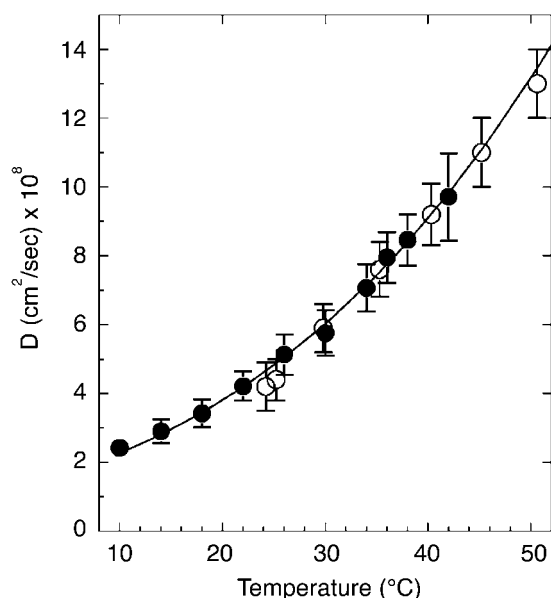


FIGURE 2 Diffusion coefficients ( $D_L$ ) in DMPC (○) and DMPC/Chol 50:50 (●). The values of  $D_L$  for the equimolar mixture were shifted by  $-16^\circ\text{C}$ . The curve is the same as in Fig. 1  $D$ , but also shifted by  $-16^\circ\text{C}$  and extrapolated to higher temperatures. Reprinted with permission from Almeida et al. (1992). Copyright 1992 American Chemical Society.

three “jumps” if lipid diffusion is viewed as a random walk on a lattice. This is certainly not long-range diffusion. In our opinion, this is probably the main reason for the apparent discrepancy between the MD simulations and the experimental data obtained with techniques that measure long-range diffusion, such as FRAP, FCS, and pulsed field gradient (pfg)-NMR. The differences in measurements of long- and short-range diffusion have been addressed previously (Vaz and Almeida, 1991). In addition, could it be that the force fields currently available for MD simulations do not correctly model the water-membrane interfacial region? Falck et al. (2004) point out that their simulations, as well as any other united-atom MD simulations, cannot reproduce the behavior of the experimental  $^2\text{H}$ -NMR order parameter for the deuterons on the second carbon of the *sn*-2 chain (Sankaram and Thompson, 1990a; Seelig and Seelig, 1975), which are near the interface. A third possibility is that the experimental, long-range  $D_L$  in PC/Chol systems is affected by phospholipid–Chol complex formation, which could occur on timescales beyond the reach of the current MD simulations.

With regard to the second problem, the minimum in  $E_a$  for DMPC/Chol 70:30, which results from the analysis of our diffusion data using free area theory (Eq. 1), Falck et al. (2004) find this result unexpected and attribute it to an incompleteness of free area theory. This is possible. Perhaps what appears as an activation energy in that analysis has contributions that the theory does not treat or does not treat adequately. As a type of mean-field theory, its treatment of fluctuations, which are critical for diffusion, is certainly not complete. Nevertheless, with all its imperfections, free area theory has successfully described lipid diffusion in a quantitative way in the experimental systems that we have examined (Vaz et al., 1985; Almeida et al., 1992). The theory is certainly simple; however, simplicity in a theory is not necessarily a weakness. The real test of a theory is its ability to describe experimental data. As stated by Feynman (1963), “The principle of science, the definition, almost, is the following: the test of all knowledge is experiment. Experiment is the sole judge of scientific ‘truth.’”

Yet, another possibility is that the minimum in  $E_a$  for DMPC/Chol 70:30 is real and reflects some important property of the system. If phospholipid/cholesterol systems are understood on the basis of a phase diagram,  $\chi_{\text{cho}} = 0.30$  essentially corresponds to the composition of the  $\ell_o$  phase in equilibrium with the  $\ell_d$  phase in the two-phase region. This may not be a coincidence and may reflect some special property of 2:1 PC/Chol mixtures. If an interpretation of the behavior of phospholipid/cholesterol systems in terms of complex formation is preferred, as proposed by McConnell and collaborators in recent work (see, for a review, McConnell and Radhakrishnan, 2003) this composition corresponds to a pressure cusp (minimum) in the phase diagrams of phospholipid/Chol systems, which has been interpreted by them as indicative of the formation of a

phospholipid/cholesterol 2:1 condensed complex. An interesting observation by Chong (1994), on the basis on our calculated mean areas per DMPC as a function of cholesterol content (Almeida et al., 1992), is that the average value of a DMPC cross-sectional area per chain at 35°C is reduced from 29.5 Å<sup>2</sup> in pure DMPC to 26.7 Å<sup>2</sup> in 70:30 DMPC/Chol. This is exactly the same as the value found for  $a_o^{\text{cho}}$ , 26.6 Å<sup>2</sup>, which, because it corresponds to a rigid molecule (cholesterol) is not expected to change with temperature (Chong, 1994; McConnell and Radhakrishnan, 2003). Therefore, packing may be especially good and the exchange between PC chains and cholesterol may be especially easy at  $\chi_{\text{cho}} = 0.30$ . This could be reflected in a smaller apparent  $E_a$ . Finally, as we have suggested (Almeida et al., 1992), these variations in  $E_a$  may reflect changes in hydration of the bilayer, which may not be monotonic with  $\chi_{\text{cho}}$  when comparing one phase with the other ( $\ell_d$  and  $\ell_o$ ), although they would be expected to be monotonic within each phase when the cholesterol content is changed, as is observed.

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