

## Article

## How Cholesterol Could Be Drawn to the Cytoplasmic Leaf of the Plasma Membrane by Phosphatidylethanolamine

Ha Giang<sup>1</sup> and M. Schick<sup>1,\*</sup><sup>1</sup>Department of Physics, University of Washington, Seattle, Washington

**ABSTRACT** In the mammalian plasma membrane, cholesterol can translocate rapidly between the exoplasmic and cytoplasmic leaves, so that its distribution between them should be given by the equality of its chemical potential in the leaves. Due to its favorable interaction with sphingomyelin, which is almost entirely in the outer leaf, one expects the great majority of cholesterol to be there also. Experimental results do not support this, implying that there is some mechanism attracting cholesterol to the inner leaf. We hypothesize that it is drawn there to reduce the bending free energy of the membrane caused by the presence of PE (phosphatidylethanolamine). It does this in two ways: first by simply diluting the amount of PE in the inner leaf, and second by ordering the tails of the PE to reduce its spontaneous curvature. Incorporating this mechanism into a model free energy for the bilayer, we find that between 50 and 60% of the total cholesterol should be in the inner leaf of human erythrocytes.

### INTRODUCTION

The importance of cholesterol in the regulation of the properties of mammalian cells is widely recognized, and it has been the subject of intense research (1,2). Nonetheless, some very basic questions about it remain unanswered. Among these is its distribution between the two leaves of the plasma membrane. It is well known that cholesterol can translocate rapidly between these leaves (3–5). Consequently, its distribution should be determined from the equilibrium requirement that the chemical potential of cholesterol be the same in both leaves. Given this, the well-known preference of cholesterol for SM (sphingomyelin) among phospholipids (6), and the fact that almost all of the SM is in the exoplasmic leaf of the plasma membrane (7), one might expect that the free energy of the system would decrease with an increase in the concentration of cholesterol in the outer leaf, and that the cholesterol would be found predominantly in that leaf. Indeed, molecular-dynamics simulations of some simple models of asymmetric bilayers incorporating SM and cholesterol do find the majority of cholesterol in the outer leaf (8,9), just as early experiments did (10).

However, since 1982, experiments have consistently reported distributions of cholesterol that run contrary to these expectations. There is less agreement on what the actual distribution is. Some estimate that the cholesterol is rather evenly divided between leaves (4,11), others that it is found to a greater extent in the inner, cytoplasmic, leaflet of the plasma membrane of various cells (12–16). Initial observations were made on human erythrocytes

(12,13) with 75–80% of the cholesterol reported to be in the inner leaf. Similar observations were then made in plasma membranes of other cells, such as neurons in mice (14,15), ovaries in Chinese hamsters (16), and the endocytotic recycling compartment of the latter (16). Nonetheless, because the experimental evidence can be characterized as indirect, and a rationale for the results is absent, a 2011 review could describe the situation by stating that “...the transbilayer orientation of the sterols that make up one-third of the lipids in the eukaryotic plasma membrane has still not been resolved satisfactorily” (17).

In this article, we propose two related mechanisms that would counteract the attraction of cholesterol for the SM in the outer, exoplasmic, leaflet and would draw it to the inner, cytoplasmic, one. We begin with the observation that almost all of the PE (phosphatidylethanolamine) is in the cytoplasmic leaf (7). PE has a small headgroup, and thus a relatively large spontaneous curvature (18). Because of this, PE forms inverted hexagonal phases at high temperatures, at which the entropy of its hydrocarbon tails dominates, and forms lamellar phases only at lower temperatures (19). Thus, the free energy of bilayers containing PE in the inner leaf must encompass a significant amount of bending energy. This bending energy is quadratic in the concentration of PE, and therefore acts as a repulsive interaction between PE molecules. Such an interaction is equivalent to an attractive interaction between PE and all other components, and affects their distribution. In particular, the bending energy can be reduced simply by diluting the PE and replacing it with any other component that does not increase the spontaneous curvature of the leaf. We assume that this is true of cholesterol, due to its small size and its

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\*Correspondence: [schick@phys.washington.edu](mailto:schick@phys.washington.edu)

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placement below the headgroups of the phospholipids, as in the umbrella model (20).

In addition to this, the bending energy penalty is also quadratic in the spontaneous curvature of the PE, which increases with the disorder of its tails. But cholesterol is known to decrease the disorder of hydrocarbon tails of phospholipids (21). In particular, in excess of 0.35 mol fraction cholesterol, POPE (palmitoyloleoylphosphatidylethanolamine) bilayers are very well ordered, comparable to those of POPC (palmitoyloleoylphosphatidylcholine) cholesterol bilayers (22). Thus cholesterol will be drawn to the inner leaflet to reduce the bending energy penalty of PE by decreasing its spontaneous curvature. That this bending energy penalty is reduced by a sufficient concentration of cholesterol is in accord with the sterol's effect on the temperature of transition of PE from the high-temperature hexagonal phase to the low-temperature lamellar one. Whereas the initial addition of cholesterol decreases this transition temperature and stabilizes the inverted-hexagonal phase, amounts greater than 0.3 mol fraction increase the transition temperature and stabilize the lamellar phase (23,24). Some lipids, like POPS (palmitoyloleoylphosphatidylserine) and POPC, also stabilize a lamellar phase in mixtures with PE (25), but do so because their own architecture and interactions favor a lamellar phase, and not because they order the chains of PE. Thus, in contrast to cholesterol, they are not expected to relieve the bending energy penalty of incorporating PE into bilayers.

In sum, we suggest that the free energy of the system is decreased if cholesterol is drawn to the cytoplasmic leaf of the plasma membrane because it reduces the bending energy caused by the inclusion of PE. It does this in two ways: first, by simply diluting the PE in the inner leaf; second, by actually reducing the magnitude of the spontaneous curvature of the PE itself.

We incorporate this hypothesis into a model of an asymmetric membrane consisting of PC (phosphatidylcholine), SM, and cholesterol in the outer leaf and PS (phosphatidylserine), PE, and cholesterol in the inner leaf. We take from experiment on erythrocytes the ratios of SM to PC and of PE to PS as well as the ratio of the total amount of cholesterol to the total amount of lipid. Requiring that the chemical potential of cholesterol in the two leaves be the same, we determine the fraction of cholesterol in the inner leaf. For reasonable values of the interaction parameters we find, in the absence of the bending energy penalty, that only about 25% of the cholesterol is in the inner leaf. The majority is in the outer leaf due to the presence of SM there. Including the bending energy, but ignoring any effect of cholesterol on the spontaneous curvature of PE, we determine that about 39% of cholesterol is now in the inner leaf. Finally, taking into account the ordering effect of cholesterol on PE, we find that between 50 and 60% of the cholesterol should be found in the cytoplasmic leaf of the human erythrocyte membrane.

## THEORETICAL MODEL

### Procedure

We consider a bilayer of which the outer, exoplasmic, leaf consists of  $N_{SM}$  molecules of sphingomyelin,  $N_{PC}$  molecules of phosphatidylcholine, and  $N_{Co}$  molecules of cholesterol; and the inner, cytoplasmic leaf, consists of  $N_{PE}$  molecules of phosphatidylethanolamine,  $N_{PS}$  molecules of phosphatidylserine, and  $N_{Ci}$  molecules of cholesterol. We assume that the glycerophospholipids are, for the most part, unsaturated in the sn-2 chain. We denote the total number of molecules in the outer leaf by  $N_o$ , the total number of molecules in the inner leaf by  $N_i$ , and the total number of molecules in the bilayer by  $N_{bi}$ . We assume that each leaf is a liquid with the areas of the outer leaf,  $A_o$ , and of the inner leaf,  $A_i$ , directly related to their molecular compositions. If the area per molecule of the phospholipids be denoted by  $a$  and that of cholesterol by  $r_a a$ , then

$$\begin{aligned} A_o &= [N_{SM} + N_{PC} + r_a N_{Co}]a = [N_o - (1 - r_a)N_{Co}]a, \\ A_i &= [N_{PE} + N_{PS} + r_a N_{Ci}]a = [N_i - (1 - r_a)N_{Ci}]a. \end{aligned} \quad (1)$$

In the absence of lateral pressure, the Helmholtz free energy of the bilayer,  $F_{bi}$ , depends only on the temperature,  $T$ , and the numbers of molecules of each component. Inasmuch as the free energy is an extensive quantity, it can be written in the form

$$\begin{aligned} F_{bi}(N_{SM}, N_{PC}, N_{Co}, N_{PE}, N_{PS}, N_{Ci}, T) \\ = N_{bi} f_{bi}(x_{SM}, x_{PC}, x_{Co}, x_{PE}, x_{PS}, x_{Ci}, T), \end{aligned} \quad (2)$$

where  $N_{bi} = N_o + N_i$  and  $x_{SM} = N_{SM}/N_{bi}$  comprise the mol fraction of SM in the bilayer; this is similar for the other components as well. By definition, the sum of the mol fractions of all components is unity.

It will be more convenient to express quantities in terms of the mol fractions of a component in the inner or outer leaf rather than the mol fraction in the bilayer. Thus, we introduce

$$\begin{aligned} y_{SM} &= \frac{N_{SM}}{N_o} = x_{SM} \frac{N_{bi}}{N_o}, \\ y_{PC} &= x_{PC} \frac{N_{bi}}{N_o}, \\ y_{Co} &= x_{Co} \frac{N_{bi}}{N_o}, \end{aligned} \quad (3)$$

$$\begin{aligned} y_{PE} &= \frac{N_{PE}}{N_i} = x_{PE} \frac{N_{bi}}{N_i}, \\ y_{PS} &= x_{PS} \frac{N_{bi}}{N_i}, \\ y_{Ci} &= x_{Ci} \frac{N_{bi}}{N_i}. \end{aligned} \quad (4)$$

By definition,  $y_{SM} + y_{PC} + y_{Co} = 1$  and  $y_{PE} + y_{PS} + y_{Ci} = 1$ , so that only four of these mol fractions are independent. We take the areas of the two leaves to be equal as the fractional area difference, being of the order of the ratio of the membrane thickness to the cell diameter, is small, of order  $10^{-3}$ . Although this difference can be of importance for the gross morphology of cells (26), it is not of importance here. From the equality of areas, the fractions  $N_{bi}/N_i$  and  $N_{bi}/N_o$  can be obtained, and the mol fractions  $x$  can be expressed in terms of the mol fractions  $y$  and vice versa. In particular, the total mol fraction of cholesterol in the bilayer,  $x_C$ , is given by

$$x_C = \frac{y_{Ci} + y_{Co} - 2(1 - r_a)y_{Ci}y_{Co}}{2 - (1 - r_a)(y_{Ci} + y_{Co})}. \quad (5)$$

The four independent mol fractions, then, are determined by the requirement that the chemical potentials of cholesterol in the two leaves be the same; that the ratios of SM to PC in the outer leaf,  $y_{SM}/y_{PC}$ ; and PS to PE, in the inner leaf,  $y_{PS}/y_{PE}$  be equal to their experimental values; and that the total mol fraction of cholesterol in the bilayer (Eq. 5) be equal to its experimental value. Then the distribution of cholesterol between leaves is determined.

We now turn to three models for the free energy of the bilayer. In the first model, we consider each leaf to be described by a phenomenological, regular solution, free energy (27). The coupling between leaves is provided solely by the equality of the cholesterol chemical potentials in the two leaves. We do this because it is not only unclear what other coupling mechanisms are important (28), but also because none of them so directly affect the cholesterol distribution as the one we do incorporate.

In the second model, we include a bending energy due to the presence of PE in the inner layer. This draws the cholesterol to the cytoplasmic leaf where it reduces the bending energy penalty caused by the presence of PE simply by diluting it.

In the third model, we include the effect on the spontaneous curvature of PE due to cholesterol, which draws additional amounts of it to the inner leaf to further reduce the bending energy caused by the presence of PE.

### Regular solution free energy

We take as the model free energy a sum of the free energies of the two leaves in the form

$$\begin{aligned} F_{bi}(N_{SM}, N_{PE}, N_{Co}, N_{PE}, N_{PS}, N_{Ci}, T) \\ = N_a f_o(y_{SM}, y_{PC}, y_{Co}, T) + N_b f_i(y_{PE}, y_{PS}, y_{Ci}, T), \\ f_i = 6\epsilon_{PS,PE} y_{PS} y_{PE} + 6\epsilon_{PC,C} y_{PS} y_{Ci} + 6\epsilon_{PE,C} y_{PE} y_{Ci} \\ + k_B T (y_{PS} \ln y_{PS} + y_{PE} \ln y_{PE} + y_{Ci} \ln y_{Ci}), \end{aligned} \quad (6)$$

$$\begin{aligned} f_o = 6\epsilon_{SM,PC} y_{SM} y_{PC} + 6\epsilon_{SM,C} y_{SM} y_{Co} + 6\epsilon_{PC,C} y_{PC} y_{Co} \\ + k_B T (y_{SM} \ln y_{SM} + y_{PC} \ln y_{PC} + y_{Co} \ln y_{Co}). \end{aligned} \quad (7)$$

We have assumed an average of six nearest-neighbor interactions per molecule. From this free energy we calculate the chemical potential of the cholesterol in the inner and outer leaves:

$$\begin{aligned} \mu_{Ci} &= \frac{\partial F_{bi}}{\partial N_{Ci}} = \frac{\partial N_a f_i(y_{PE}, y_{PS}, y_{Ci}, T)}{\partial N_{Ci}} \\ &= \frac{\partial f_i}{\partial y_{Ci}} + f_i - \sum_j \frac{\partial f_i}{\partial y_j} y_j, \\ j &= PE, PS, C_i, \end{aligned} \quad (8)$$

$$\begin{aligned} \mu_{Co} &= \frac{\partial F_{bi}}{\partial N_{Co}} = \frac{\partial N_o f_o(y_{SM}, y_{PC}, y_{Co}, T)}{\partial N_{Co}} \\ &= \frac{\partial f_o}{\partial y_{Co}} + f_o - \sum_k \frac{\partial f_o}{\partial y_k} y_k, \\ k &= SM, PC, C_o. \end{aligned} \quad (9)$$

Again, to determine the six mol fractions, we equate these two chemical potentials, utilize the two constraints

$$\begin{aligned} \sum_j y_j &= 1, \\ j &= PE, PS, C_i, \end{aligned} \quad (10)$$

$$\begin{aligned} \sum_k y_k &= 1, \\ k &= SM, PC, C_o, \end{aligned} \quad (11)$$

and set to their experimental values the ratios of SM to PC in the outer leaf,  $y_{SM}/y_{PC}$ , of PS to PE in the inner leaf,  $y_{PS}/y_{PE}$ , and the total mol fraction of cholesterol in the bilayer ( $x_C$ ) (Eq. 5). Once  $y_{Ci}$  and  $y_{Co}$ , the mol fractions of cholesterol in each leaf are obtained, the percentage of the total cholesterol in the inner leaf follows from

$$p = \frac{y_{Ci}}{y_{Ci} + y_{Co} (N_o/N_i)} \times 100,$$

where the ratio of the number of molecules in each leaf is determined by the equality of the areas (Eq. 1).

We must now set the parameters of our model. For the ratio of the area per molecule of cholesterol to the area per molecule of the other phospholipids, we take  $r_a = 0.6$  because the average area per molecule of phospholipids is  $\sim a = 0.7 \text{ nm}^2$  and that of cholesterol is  $\sim 0.4 \text{ nm}^2$  (21,29). For the binary interactions, we choose  $\epsilon_{SM,C} = -0.58 k_B T$ ,  $\epsilon_{PC,C} = 0.2 k_B T$ ,  $\epsilon_{SM,PC} = 0.30 k_B T$ ,  $\epsilon_{PS,C} = -0.06 k_B T$ ,  $\epsilon_{PE,C} = 0.28 k_B T$ , and  $\epsilon_{PS,PE} = 0$ . We discuss the selection of these values in the Appendix.

We must also specify the membrane. We assume that it is at a temperature  $T = 37^\circ\text{C}$ . We take the ratios of the

components to be those of the human erythrocyte as given by Zachowski (30). There we find that the SM accounts for 0.22 of phospholipids in the outer leaf and 0.02 in the inner leaf, whereas PC accounts for 0.20 of the phospholipids in the outer leaf and 0.07 in the inner. For simplicity, we assume that all the SM and PC are in the outer leaf and take the ratio  $y_{SM}/y_{PC} = 0.22:0.20 = 1.1$ . Similarly, PS accounts for 0.13 of phospholipids in the inner leaf and 0.02 in the outer, whereas PE accounts for 0.25 in the inner leaf and 0.08 in the outer. Assuming that all PS and PE are in the inner leaf, we take the ratio  $y_{PS}/y_{PE} = 0.13:0.25 = 0.52$ . Lastly, we set the total mol fraction of cholesterol in the bilayer to be  $x_C = 0.4$  (17).

It is now straightforward to carry out our program, and we find a single solution of our equations. By examining the matrix of second derivatives of the free energy, we have verified that this solution is stable. The inner leaf contains a mol fraction of cholesterol  $y_{Ci} = 0.22$ . This corresponds to only 25% of the total cholesterol being in the inner leaf. This is easy to understand, inasmuch as the energy of cholesterol is reduced if it goes to the outer layer where it can interact favorably with the SM concentrated there. It is easy to understand, but clearly not in accord with the reported results of experiment. What physics is missing?

### Addition of the bending energy

It is our hypothesis that what is missing is that cholesterol can ameliorate the bending energy cost of having PE in the cytoplasmic leaf both by diluting the PE and by increasing the order of its tails, reducing its curvature.

We incorporate this hypothesis into our model by adding to the free energy of the flat bilayer a bending energy. Because absolute values of the spontaneous curvatures of SM, PC, and PS are an order-of-magnitude smaller than that of PE (18,31–33), we consider the curvature only of PE and write the bending energy as

$$F_b = \left( \frac{A_0 + A_i}{2} \right) \frac{\kappa}{2} y_{PE}^2 H_{PE}^2 \quad (12)$$

$$= 1/2 [N_i + N_o - (1 - r_a)(N_{C_o} + N_{C_i})] f_b,$$

$$f_b = \frac{1}{2} a \kappa y_{PE}^2 H_{PE}^2, \quad (13)$$

with  $\kappa$  the bending modulus. We shall take  $\kappa = 44 k_B T$ , which is appropriate for red blood cells (34).

Our model free energy is now

$$F_{bi} = N_a f_o + N_i f_i + 1/2 [N_i + N_o - (1 - r_a)(N_{C_o} + N_{C_i})] f_b, \quad (14)$$

with  $f_i$  and  $f_o$  given by Eqs. 6 and 7 and  $f_b$  by Eq. 13. We calculate the chemical potential of the cholesterol in the

outer and inner layers. We then set the areas of the two leaves, Eq. 1, to be equal, after which we obtain

$$\mu_{C_o} = f_o + \frac{\partial f_o}{\partial y_{C_o}} - \sum_j \frac{\partial f_o}{\partial y_j} y_j + \frac{1}{2} r_a f_b, \quad (15)$$

$$j = \text{SM, PC, } C_o,$$

$$\begin{aligned} \mu_{C_i} = & f_i + \frac{\partial f_i}{\partial y_{C_i}} - \sum_k \frac{\partial f_i}{\partial y_k} y_k + \frac{1}{2} r_a f_b \\ & + [1 - (1 - r_a) y_{C_i}] \left[ \frac{\partial f_b}{\partial y_{C_i}} - \sum_k \frac{\partial f_b}{\partial y_k} y_k \right], \end{aligned} \quad (16)$$

$$k = \text{PE, PS, } C_i.$$

We must now specify the intrinsic curvature of PE. We first take its value to be that in the absence of cholesterol,

$$H_{PE} = H_{PE}^0, \quad (17)$$

one which has been measured to be  $H_{PE}^0 = -0.316 \text{ nm}^{-1}$  (18). Repeating our procedure, we now find a solution corresponding to a stable bilayer in which the inner layer contains a mol fraction  $y_{C_i} = 0.32$ . This corresponds to 38.6% of the total cholesterol now being found in the inner leaf. As stated above, the reason for this increase from the previous 25% is simple. The bending energy penalty due to the presence of the PEs is equivalent to a pairwise repulsion between them, and thus an attraction between them and all other components. Cholesterol in the outer layer responds, decreasing the free energy of the system by going to the inner layer and diluting the effect of the PE.

We now consider the additional effect of cholesterol acting on the tails of PE which, for sufficient mol fraction, will order the tails of PE and further reduce the bending energy cost of having PE in a planar bilayer. We incorporate this effect by having the spontaneous curvature of PE depend upon the cholesterol mol fraction. We choose

$$H_{PE}(y_{C_i}) = H_{PE}^0 - B \frac{y_{C_i}}{y_{\min}} + \frac{B}{\lambda} \left( \frac{y_{C_i}}{y_{\min}} \right)^\lambda \quad (18)$$

with  $B = 0.05$ ,  $y_{\min} = 0.3$ , and  $\lambda = 8$ . This description is dictated by the following considerations:

1. From the behavior of the temperature of transition of PE from the inverted-hexagonal to lamellar phase (23), we know that the addition of cholesterol initially stabilizes the former phase with respect to the latter. This could result from the cholesterol decreasing the free energy of the inverted-hexagonal phase, or increasing the free energy of the lamellar phase, or a combination of the two. We choose the second of these possibilities by having the spontaneous curvature of PE become more negative with the initial addition of cholesterol. This leads to the second term in Eq. 18 above.

- Again from the behavior of the temperature of transition of PE from the inverted-hexagonal to lamellar phase (23), we expect that the magnitude of the spontaneous curvature  $H_{PE}(y_{Ci})$  no longer increases for mol fractions of cholesterol  $> y_{\min} \approx 0.3$ .
- Finally, the observation that, with a mol fraction cholesterol of 0.35, the tails of PE are as well ordered as those of PC with cholesterol (22), dictates a choice of  $\lambda$  such that the magnitude of the spontaneous curvature of PE in the presence of cholesterol decreases rapidly for values of  $y$  somewhat larger than 0.35. The behavior of the spontaneous curvature  $H_{PE}(y)$  is shown in Fig. 1 for values of  $\lambda = 6, 8, 10,$  and  $12$ . We choose  $\lambda = 8$ . The dependence of our results on our choice of  $\lambda$  will be shown below.

With the above cholesterol-dependent PE spontaneous curvature, we now find that the inner leaf contains  $y_{Ci} = 0.45$  mol fraction of cholesterol, which corresponds to 58% of the total cholesterol being in the inner leaf. We note that the mol fraction of cholesterol in the inner leaf is below the maximum solubility of cholesterol in PE, 0.51 (20). With our solution, the mol fractions of the other components in the plasma membrane are  $y_{PS} = 0.19$  and  $y_{PE} = 0.36$  in the inner leaf, and  $y_{Co} = 0.35, y_{SM} = 0.34,$  and  $y_{PC} = 0.31$  in the outer leaf.

Because our hypothesis depends so directly upon the bending energy penalty, our results are clearly a function of the magnitude of the bending modulus. We have taken it to be  $\kappa = 44 k_B T$ , appropriate for mammalian red blood cells (34). The dependence of the fraction of cholesterol in the inner leaf upon the bending modulus is shown in Fig. 2.

We have taken the total cholesterol content of the membrane to be  $x_C = 0.4$ , appropriate for the plasma membrane of erythrocytes (17). As for membranes of other cells, only the plasma membrane and late-stage endosomes are characterized by such large levels of cholesterol (35), whereas other membranes contain much less cholesterol. It is of interest, therefore, to determine the dependence of the per-

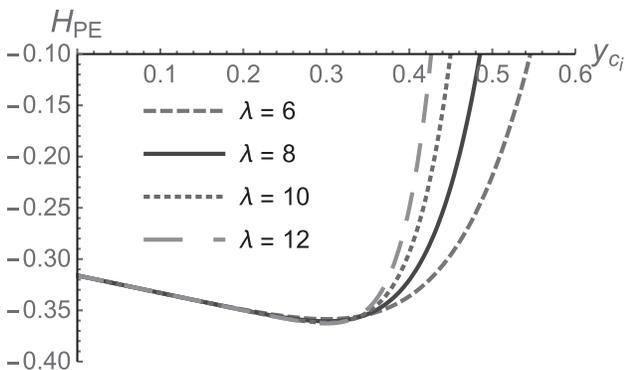


FIGURE 1 Spontaneous curvature of PE, in inverse nanometers, in the presence of cholesterol as modeled by Eq. 18 for four values of  $\lambda$ .

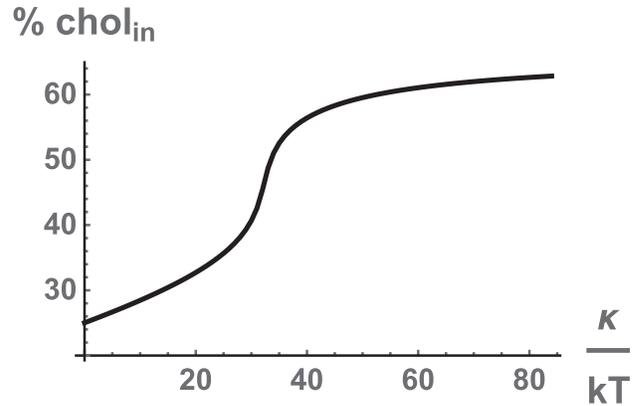


FIGURE 2 Percentage of cholesterol in the inner leaf as a function of the bending modulus  $\kappa$ .

centage of total cholesterol in the inner leaf as a function of total cholesterol content. This is shown in Fig. 3.

There are a few things to note:

- When the total amount of cholesterol is small, the percentage in the inner leaf is also small. Hence, for those membranes with small amounts of cholesterol, we find that most of it will be in the outer leaf.
- The percentage of cholesterol in the inner leaf initially increases linearly. This is because of the reduction of the bending energy penalty due to the dilution of the PE by cholesterol. The sharp increase reflects the reduction by additional cholesterol of the PE spontaneous curvature.
- There is a maximum in the percentage of cholesterol in the inner leaf as a function of total cholesterol content. This follows from the fact that we have found that the percentage in the inner leaf can exceed 50% and that this percentage must approach 50 in the limit in which  $x_C \rightarrow 1$ . It is interesting that the maximum occurs near  $x_C \approx 0.35$ , not very different from the value characterizing erythrocytes.

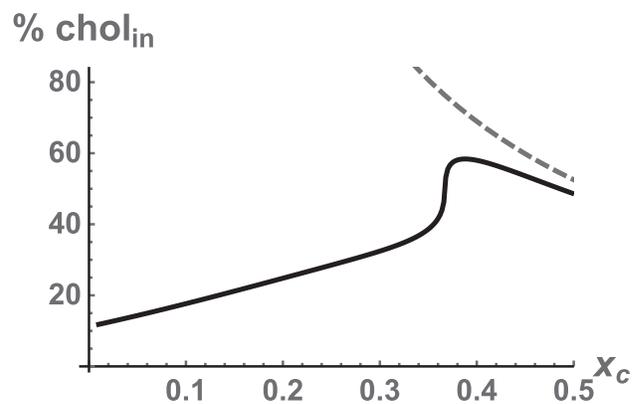


FIGURE 3 Percentage of total cholesterol in the inner leaf as a function of the total cholesterol mol fraction  $x_C$ . (Dashed line) Solubility limit of cholesterol.

## DISCUSSION

We have proposed that cholesterol is drawn to the cytoplasmic leaf of the plasma membrane in the plasma membrane because that is where the phosphatidylethanolamine is; that by going there, it reduces in two ways the bending energy penalty of incorporating PE into a planar bilayer:

1. By simply diluting the PE mol fraction, cholesterol reduces the penalty, which is quadratic in the PE mol fraction; and
2. In sufficient quantity, cholesterol orders the tails of PE, thereby further reducing the penalty of incorporating PE into the bilayer.

By these means, the chemical potential of cholesterol in the inner leaf will be negative, and matches the chemical potential of cholesterol in the outer leaf, which is negative due to the presence of sphingomyelin. We employed a simple model to show that in the absence of bending energy considerations, only ~25% of the cholesterol would be in the cytoplasmic leaf, again due to the presence of sphingomyelin in the exoplasmic leaf. Including the bending energy penalty simply of incorporating the PE, i.e., without any effect of cholesterol on its tails, we found that the fraction of total cholesterol in the inner leaf would increase to 39%. Finally, by including the ordering effect of cholesterol on PE, we obtained a fraction of total cholesterol in the inner leaf of some 58%.

Several comments are in order. The additional increase in the fraction of cholesterol in the inner leaf due to the ordering of the PE tails by cholesterol depends upon our phenomenological modeling of this effect by our choice of the cholesterol-dependent spontaneous curvature of PE, as given in Eq. 18. The effect of varying the parameter  $B$ , reflecting the coupling of the cholesterol to the order of the PE tails, is simple. Were  $B = 0$ , then the spontaneous curvature of PE would be constant, equal to  $H_{PE}^0$ , and the only response of cholesterol to the bending energy would be to dilute the PE. The fraction of the total cholesterol that is in the inner leaf would be 39%. This corresponds to a mol fraction of cholesterol in the inner leaf of 0.32, too little to order the tails of PE. For small values of  $B$ , this remains the case even though the spontaneous curvature does depend on the amount of cholesterol in the inner leaf. For values of  $B > 0.045$ , however, the amount of cholesterol drawn to the inner leaf is 0.45 mol fraction, sufficient to order the tails of the PE and greatly reduce its curvature. The fraction of total cholesterol in the inner leaf is 58%, and is not very sensitive to further increases in  $B$ . As to the parameter  $\lambda$ , we argued that a range of values was permissible. We show in Fig. 4 that the fraction of cholesterol in the inner leaf exceeds 50% for the values of  $\lambda$  in this range. In principle, one should be able to go beyond this phenomenological modeling by utilizing more microscopic, analytic theories that include an explicit description of the lipid tails and can describe

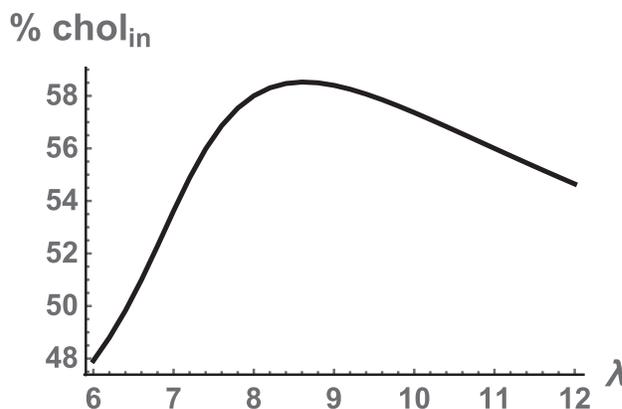


FIGURE 4 Percentage of cholesterol in the inner leaf as a function of the power  $\lambda$  in the modeling of the spontaneous curvature, Eq. 18.

not only the polymorphism of PE (36), but also the effect of cholesterol upon the tails (37). Alternatively simulations similar to, but larger than, those of Mori et al. (38) should evince the effect we have described.

We have checked the dependence of our results on other assumptions that we have made. The effect of reducing the number of nearest-neighbor interactions from six to four is small; the percentage of total cholesterol in the inner leaf increases from 58 to 61%.

Because the cholesterol is driven to the leaf in which the PE is located, the largest effect on our result for the cholesterol distribution comes from including the amount of PE in the outer leaf, which is a fraction 0.08, compared to 0.25 in the inner leaf (30). If we do this, as well as include the 0.07 PC in the inner leaf, compared to 0.20 in the outer leaf, we find that for  $x_C = 0.4$  the percentage of total cholesterol in the inner leaf decreases from 58 to 51%.

Even this modest majority, however, demonstrates our major point: that the long-standing expectation that cholesterol should be found predominantly in the exoplasmic leaf of the plasma membrane, because that is where the sphingomyelin is located, is very likely incorrect. The presence of PE in the cytoplasmic leaf at the cost of a bending energy penalty attracts cholesterol there to reduce that penalty. The specific percentage of cholesterol in the inner leaf depends on several factors that we have tried to illustrate, but values on the order, or in excess, of 50% can be understood.

We close with some observations concerning possible experiments:

1. The effect of total cholesterol content on the percentage of cholesterol in the inner leaf, as shown in Fig. 3, should be measurable in model asymmetric membranes that mimic the plasma membrane (39).
2. As noted earlier, our model is consistent with results on the effect of cholesterol on the inverted-hexagonal to lamellar phase transition temperature in mixtures of cholesterol and POPE (23). There are some correlations

clearly expected between cholesterol content and the specific form of PE in the membrane. Because the *trans* double bond in DEPE (dielaidoylphosphatidylethanolamine) is more easily ordered by cholesterol than is a *cis* double bond (23,24), one expects the asymmetry in the cholesterol distribution to be less. It would be interesting to know how cholesterol affects a lipid with a polyunsaturated tail such as 18:0/20:4 PE, inasmuch as these polyunsaturated tails make up a nonnegligible fraction, perhaps 18%, of the chains of PE in the plasma membrane (40).

3. It is of interest to note that the effects of ergosterol on PE membranes is not the same as the effect of cholesterol on them. Although the chain order of PE keeps increasing with cholesterol content to fractions of ~0.45 mol fraction (22), the order saturates at ergosterol mol fractions of ~0.1 (41). Thus, as Richard Epand (McMaster University, Hamilton, Ontario, Canada) has kindly pointed out to us, one might expect the fraction of cholesterol in the inner leaf of yeast plasma membrane to be rather different from that in mammalian cells. To our knowledge, the distribution of ergosterol between the leaves of the yeast plasma membrane has not been measured. The result of such an experiment would certainly be of interest and relevant to the considerations we have put forth here.

## APPENDIX

The free energy of a binary mixture of molecules of species *A* and *B*, as obtained from regular solution theory, is simply the mean-field approximation to the exact free energy obtained from a lattice-gas Hamiltonian of form

$$H = \sum_{\langle ij \rangle} \left[ E_{A,A} n_i^A n_j^A + E_{B,B} n_i^B n_j^B + E_{A,B} (n_i^A n_j^B + n_i^B n_j^A) \right], \quad (19)$$

where  $n_i^A = 1$  if there is a molecule of species *A* at the site *i* and is zero otherwise, and similarly for  $n_i^B$ . The sum is over all distinct nearest-neighbor pairs of molecules. This Hamiltonian is easily mapped to that of an Ising model

$$H = -J \sum_{\langle ij \rangle} S_i S_j - B \sum_i S_i, \quad (20)$$

$$S_i = \pm 1$$

via the relation  $S_i = 2n_i^A - 1 = 1 - 2n_i^B$ , so that the presence of an *A* molecule is related to an up-spin, and a *B* molecule to a down-spin. With this mapping, the interaction *J* of the Ising model is then  $J = \omega_{A,B}/2$  with  $\omega_{A,B} \equiv E_{A,B} - (E_{A,A} + E_{B,B})/2$ . The exact transition temperature of the two-dimensional Ising on a triangular lattice is known to be  $k_B T_c = 4J/\ln 3 = 2\omega_{A,B}/\ln 3$  (42). Thus, if a physical system is known to undergo a phase separation at a critical temperature *T*, then the interaction parameter in a model including fluctuations should be positive, and taken to be  $\omega_{A,B} = (1/2)\ln 3 k_B T \approx 0.55 k_B T$ .

Regular solution theory, however, does not include fluctuations. Within this theory, the free energy per particle of a binary mixture on a triangular lattice can be written

$$f = 6\epsilon_{A,B} y_A y_B + k_B T (y_A \ln y_A + y_B \ln y_B).$$

It yields a transition  $k_B T_c^{\text{rs}} = 3\epsilon_{A,B}$ . Therefore, to obtain in regular solution theory the same transition temperature as the exact result, one must choose  $\epsilon_{A,B} = (2/3 \ln 3) \omega_{A,B} \approx 0.6 \omega_{A,B}$ .

The values of the interaction parameters,  $\omega_{A,B}$ , for many pairs of lipids can be estimated from experiment and have been collected by Almeida (43). In particular for the interactions between components of the outer leaf at  $T = 37^\circ\text{C}$ ,  $\omega_{PC,C} = 0.34 k_B T$ ,  $\omega_{SM,C} = -0.97 k_B T$ , and  $\omega_{SM,PC} = 0.51 k_B T$ . Because  $\omega_{PC,C}$  is positive, we use in our regular solution theory the estimate  $\epsilon_{PC,C}/k_B T = 0.6$ ,  $\omega_{PC,C}/k_B T = 0.20$ . If we take the ratios of the other interactions parameters to be the same as in the table, i.e.,  $\epsilon_{SM,C}/\epsilon_{PC,C} = \omega_{SM,C}/\omega_{PC,C}$ , then  $\epsilon_{SM,C}/k_B T = -0.58$  and similarly  $\epsilon_{SM,PC}/k_B T = 0.30$ .

The interactions relevant to the inner leaf, those between PE and cholesterol, between PS and cholesterol, and between PE and PS, are not included in the table of Almeida (43). To obtain an estimate for them, we proceed as follows: Cholesterol and PE do not phase-separate at  $T = 37^\circ\text{C}$ , so that we should take  $\epsilon_{PE,C}/k_B T < 1/3$ . We choose  $\epsilon_{PE,C}/k_B T = 0.28$ .

Next, we estimate the interaction between PS and cholesterol. Niu and Litman (6) have measured the differences  $\Delta_{SM} \equiv \omega_{SM,C} - \omega_{PC,C} = -1181 \text{ cal/mol} = -1.92 k_B T$  at  $37^\circ\text{C}$ , and  $\Delta_{PS} \equiv \omega_{PS,C} - \omega_{PC,C} = -0.65 k_B T$ .

We assume that

$$\frac{\epsilon_{PS,C} - \epsilon_{PC,C}}{\epsilon_{SM,C} - \epsilon_{PC,C}} = \frac{\omega_{PS,C} - \omega_{PC,C}}{\omega_{SM,C} - \omega_{PC,C}} = \frac{0.65}{1.92} = 0.34,$$

so that  $\epsilon_{PS,C}/k_B T = \epsilon_{PC,C} + 0.34(\epsilon_{SM,C} - \epsilon_{PC,C}) = -0.06$ .

Lastly, because the tails of PE and PS lipids are similar, we take  $\epsilon_{PE,PS}/k_B T = 0$ . This completes the specification of the interactions.

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