

Concentration fluctuations and phase transitions in coupled modulated bilayers

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We consider the formation of finite-size domains in lipid bilayers consisting of saturated and hybrid lipids. First, we describe a monolayer model that includes a coupling between a compositional scalar field and a two-dimensional vectorial order parameter. Such a coupling yields an effective two-dimensional microemulsion free energy for the lipid monolayer, and its characteristic length of compositional modulations can be considered as the origin of finite-size domains in biological membranes. Next, we consider a coupled bilayer composed of two modulated monolayers and discuss the static and dynamic properties of concentration fluctuations above the transition temperature. We also investigate the micro-phase separation below the transition temperature and compare the micro-phase separated structures with statics and dynamics of concentration fluctuations above the transition.

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I. INTRODUCTION

Biomembranes are two-dimensional (2D) fluids that separate the inner and outer environment of organelles in biological cells. Naturally occurring biomembranes consist typically of numerous lipid species, sterols, sugars, and membrane proteins. According to the “lipid raft” hypothesis [1], some of the lipid components and/or proteins are incorporated into finite-size domains, which play an important role on cellular functions such as signal transduction processes. Recent experiments suggest that lipid rafts are nothing but dynamical molecular assemblies of 20 nm in size with finite lifetimes in the order of 10–20 ms [2].

Being motivated by the raft hypothesis, a large number of investigations [3–14] have been conducted to reveal the properties of artificial membranes consisting of lipid mixtures and cholesterol. Below the miscibility transition temperature, formation of micron-size domains were observed using fluorescent microscopy [3–6]. In some cases, rather than a macroscopic phase separation, domains with distinct size in the micrometer range have been reported [7–9]. For example, various types of modulated (stripe or hexagonal) patterns have been found for multicomponent lipid and cholesterol mixtures [10]. Above the miscibility transition temperature, even multicomponent membranes do not phase separate, and their concentration fluctuations around the homogeneous state can be investigated [11–13], in particular, close to the critical point, T_c . Furthermore, it is interesting to note that critical concentration fluctuations have been observed in membranes extracted from living cells [14].

One of the main reasons that initiated the notion of lipid rafts in biomembranes is the existence of finite-size domains rather than domains resulting from a macroscopic phase separation [15]. Assuming that membranes are in equilibrium, the same question can be phrased in terms of whether the biomembrane state is above or below the miscibility transition

temperature. In the high-temperature one-phase region, the only relevant length scale is the correlation length associated with concentration fluctuations, and it diverges at the critical point, T_c [16]. Below this temperature, there should be a physical mechanism suppressing domain coarsening in order to explain the existence of finite-size domains in equilibrium.

Yet another characteristic feature of biomembranes is that the lipid composition of the two leaflets (monolayers) constituting the bilayer is not the same [17]. Moreover, such asymmetric monolayers are not independent but are coupled to one another. This was confirmed experimentally by investigating the phase separation of bilayers with different monolayer lipid compositions [18] or seen in simulations [19]. One mechanism that leads to the coupling between the two leaflets is the lipid chain interdigitation occurring at the midplane of the bilayer [20–22], which may affect the domain size in asymmetric bilayers.

One possibility to account for such finite-size domains in lipid mixtures is to consider the special role played by “hybrid lipids” such as POPC or SOPC. These lipids have one saturated hydrocarbon chain and another unsaturated one and are major components of biological membranes. It was suggested that hybrid lipids act as a line-active agent at boundaries of 2D domains of immiscible lipids [23–26]. Hence, a mixture containing hybrid lipids can be regarded as a “2D microemulsion” [13,23–27]. For microemulsions it is known that there is another length scale, in addition to the correlation length mentioned above, related to the size of water and oil microdomains [28].

Using the microemulsion analogy, we suggest that an additional length scale can be responsible for the finite-size structures in biomembranes. We propose a model for bilayers consisting of two coupled monolayers that are both in a 2D microemulsion state, manifesting modulated phases. First, we describe a monolayer model that includes a coupling between the lipid composition and a 2D vectorial order parameter. The model accounts for the line active nature of hybrid lipids. Next, we consider a bilayer in which two such

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monolayers are coupled through an interleaflet interaction. For the coupled bilayer, we discuss the static and dynamic properties of concentration fluctuations above T_c and investigate their micro-phase separation below T_c [29]. Intermediate structures arise when two competing structures have different characteristic length scales. One of our important conclusions is that the micro-phase-separated structures below T_c reflect the static and dynamic properties of concentration fluctuations above T_c .

This paper is organized as follows. In Sec. II, after describing how the microemulsion state is obtained for monolayers, a model for coupled modulated bilayers is presented. In Sec. III, we show the results of static and dynamic structure factors for the coupled bilayers above T_c . In Sec. IV, we describe some results for competing micro-phase separation in coupled bilayers below T_c . Finally, in Sec. V, further discussion and final remarks are presented.

II. MODEL

A. Modulated lipid monolayers

We consider a 2D microemulsion formation in a monolayer consisting of two types of lipids: a saturated lipid (denoted by “S”) and a hybrid one (“H”). As shown schematically in Fig. 1(a), the saturated lipid has two saturated chains, whereas the hybrid lipid has one saturated chain and an unsaturated one. It is well known [3–6] that the resulting liquid-ordered (L_o) and liquid-disordered (L_d) phases are rich in saturated and hybrid lipids, respectively, as is depicted in Fig. 1(c). In the experiments mentioned previously, cholesterol is usually added as a third component and is known to affect the area per headgroup of lipids [30,31]. However, since cholesterol has a strong preference for the L_o phase and affects mainly the saturated lipid, we do not consider cholesterol explicitly in our model and neglect its presence hereafter.

The local area fraction of saturated and hybrid lipids are defined as $\phi_S(\mathbf{r})$ and $\phi_H(\mathbf{r})$, respectively, where $\mathbf{r} = (x, y)$ is a 2D vector. Under the incompressibility condition, $\phi_S + \phi_H = 1$, the only relevant order parameter is the difference between the two lipid compositions, $\phi = \phi_S - \phi_H$. The phenomenological free energy that describes the 2D phase separation between S and H lipids is given by a Ginzburg-Landau expansion in terms of the order parameter ϕ ,

$$F_s[\phi] = \int d^2r \left[\frac{\sigma}{2} (\nabla \phi)^2 + \frac{\tau}{2} \phi^2 + \frac{1}{4} \phi^4 - \mu \phi \right]. \quad (1)$$

Here $\sigma > 0$ is related to the line tension between monolayer domains, $\tau \propto T - T_c$ is the reduced temperature with respect to the critical point T_c , and μ is the chemical potential that regulates the average ϕ value in the monolayer. Without loss of generality, the coefficient of the quartic term is set to be a positive constant by an appropriate rescaling of the position variable, \mathbf{r} .

Next, we discuss the role of the hybrid lipid and its effect on the phase separation. We define a 2D lateral vector \mathbf{b} pointing from the unsaturated tail of the hybrid lipid toward its saturated one, as depicted in Fig. 1(a). We then introduce a coarse-grained 2D vectorial order-parameter field

$\mathbf{m}(\mathbf{r}) = (m_x(\mathbf{r}), m_y(\mathbf{r}))$, as shown in Fig. 1(b). This vector is the spatial average of the \mathbf{b} vectors over areas large as compared with molecular size but still small enough as compared with macroscopic scales. The phenomenological free energy associated with the vectorial field \mathbf{m} (and due to the hybrid lipids) can be written as an expansion up to quadratic order in \mathbf{m} [32],

$$F_h[\mathbf{m}] = \int d^2r \left[\frac{K}{2} (\nabla \cdot \mathbf{m})^2 + \frac{a}{2} \mathbf{m}^2 \right]. \quad (2)$$

The coefficient K is the 2D elastic constant, while a is taken to be positive so that $\mathbf{m} = 0$ is the stable homogeneous state. Although an additional term $(\nabla \times \mathbf{m})^2$ is allowed by symmetry, it is not included in Eq. (2) because this term does not modify our results. The reason why the term $(\nabla \times \mathbf{m})^2$ is irrelevant after the minimization is due mainly to the form of the coupling term discussed in the next.

At the L_o/L_d interface, hybrid lipids orient their saturated and unsaturated chains toward the L_o and L_d phases, respectively, thereby reducing the chain mismatch [25,26] as shown in Fig. 1(c). Within our phenomenological approach, the role of hybrid lipids can be represented by a coupling between

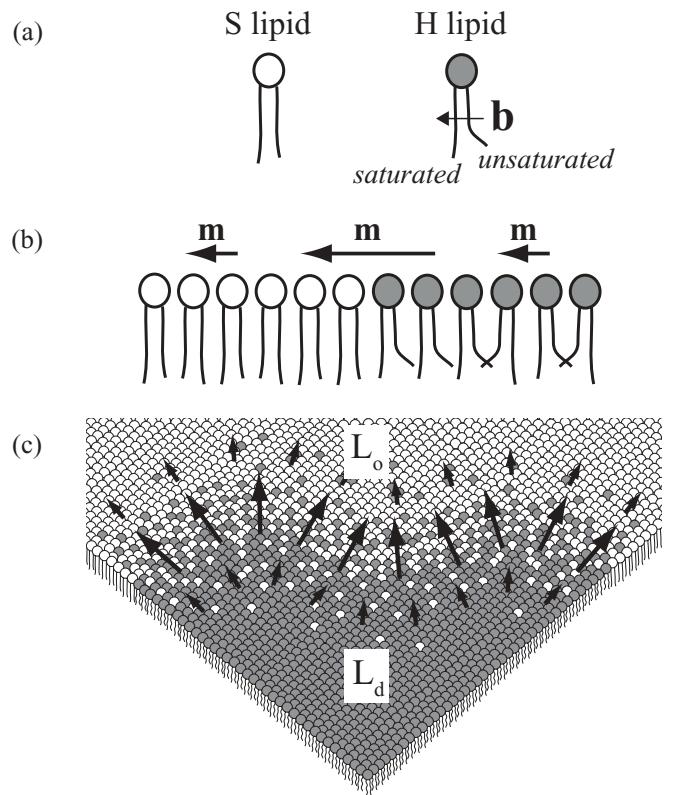


FIG. 1. (a) Saturated lipids (S) with two saturated hydrocarbon chains, and hybrid lipids (H) with one saturated and another unsaturated chain. For each hybrid lipid, we define a 2D vector \mathbf{b} pointing from its unsaturated tail toward the saturated one. (b) The vector \mathbf{m} denotes a coarse-grained 2D vectorial order parameter over the microscopic \mathbf{b} vectors. (c) A binary lipid monolayer consisting of saturated and hybrid lipids. At the interface between the L_o and L_d phases, hybrid lipids orient their saturated chains toward the L_o phase. The distribution of the $\mathbf{m}(\mathbf{r})$ vector field is shown by arrows of variable length.

the lateral variation of ϕ and the vectorial field \mathbf{m} . To lowest orders, the coupling term in the free energy is given by

$$F_c[\phi, \mathbf{m}] = -\Gamma \int d^2r \mathbf{m} \cdot (\nabla \phi), \quad (3)$$

where Γ is a positive coupling constant because the vector \mathbf{m} tends to orient toward the L_o domains. The total monolayer free energy is given here by the sum of the three terms introduced in Eqs. (1)–(3): $F_m = F_s + F_h + F_c$. Notice that our model is valid in the weak segregation limit (close to T_c), because slow spatial variation of the order parameters is intrinsically assumed. We further note that a similar free-energy functional using a vectorial order parameter was proposed for 3D microemulsions [33].

The total free energy can be conveniently expressed using the 2D Fourier transform of ϕ

$$\phi(\mathbf{q}) = \int d^2r \phi(\mathbf{r}) e^{-i\mathbf{q}\cdot\mathbf{r}}, \quad (4)$$

where $\mathbf{q} = (q_x, q_y)$ is a 2D wave vector and, similarly, $\mathbf{m}(\mathbf{q})$ is the Fourier transform of $\mathbf{m}(\mathbf{r})$. Minimizing F_m with respect to \mathbf{m} , we obtain its optimum value as

$$\mathbf{m}(\mathbf{q}) = \frac{i\Gamma\mathbf{q}}{a + Kq^2} \phi(\mathbf{q}), \quad (5)$$

where $q = |\mathbf{q}|$. By substituting back Eq. (5) into the monolayer free energy F_m , the minimized free energy is expressed as

$$F_m[\phi] = \int \frac{d^2q}{(2\pi)^2} \frac{1}{2} \left(\tau + \sigma q^2 - \frac{\Gamma^2 q^2}{a + Kq^2} \right) \phi(\mathbf{q}) \phi(-\mathbf{q}) + \int d^2r \left(\frac{1}{4} \phi^4 - \mu \phi \right), \quad (6)$$

where the Fourier transform has been used only for the second-order terms in ϕ . Expanding the effective binary interaction for small q , we obtain

$$F_m[\phi] \approx \int \frac{d^2q}{(2\pi)^2} \left(2Bq^4 - 2Aq^2 + \frac{\tau}{2} \right) \phi(\mathbf{q}) \phi(-\mathbf{q}) + \int d^2r \left(\frac{1}{4} \phi^4 - \mu \phi \right), \quad (7)$$

where two new parameters are defined

$$B \equiv \frac{K\Gamma^2}{4a^2}, \quad A \equiv \frac{1}{4} \left(\frac{\Gamma^2}{a} - \sigma \right). \quad (8)$$

When the coupling constant Γ is small enough, i.e., $\Gamma^2/a < \sigma$ (or $A < 0$), the minimum of the Gaussian term (first integral) in Eq. (7) occurs at $q^* = 0$ and is a signature of a macroscopic phase separation (for $\tau < 0$). On the other hand, when Γ is large enough, i.e., $\Gamma^2/a > \sigma$ (or $A > 0$), the minimum occurs at a nonzero wave number $q^* \neq 0$ given by

$$q^* = \sqrt{\frac{A}{2B}} = \sqrt{\frac{a(1 - a\sigma/\Gamma^2)}{2K}}, \quad (9)$$

indicating a potential micro-phase separation with $q^* > 0$ modulation.

Taking the inverse Fourier transform, Eq. (7) can be expressed in position space as

$$F_m[\phi] = \int d^2r \left[2B(\nabla^2\phi)^2 - 2A(\nabla\phi)^2 + \frac{\tau}{2}\phi^2 + \frac{1}{4}\phi^4 - \mu\phi \right]. \quad (10)$$

This is the “2D microemulsion” free energy for a monolayer composed of a binary lipid mixture. When $A > 0$, the negative gradient-squared term favors spatial modulations, while the positive Laplacian squared term with $B > 0$ suppresses modulations. As mentioned above, Yamamoto *et al.* [25,26] showed that the effective line tension between domains becomes negative for membranes consisting of saturated and hybrid lipids.

More generally, models based on equations similar to Eq. (10) have been used successfully in the past to describe modulated phases [34] arising in a variety of different biophysical and chemical systems such as Langmuir films [35], lipid membranes [36,37], and diblock copolymers [38].

B. Coupled lipid bilayers

We consider two coupled modulated monolayers forming a bilayer as shown in Fig. 2. Each monolayer is a binary mixture of saturated and hybrid lipids. We define two local order parameters for the two monolayers, $\phi = \phi_S - \phi_H$ and $\psi = \psi_S - \psi_H$, depicted in Fig. 2. The coarse-grained free-energy functional is then written as [29]

$$F_b[\phi, \psi] = \int d^2r \left[2B(\nabla^2\phi)^2 - 2A(\nabla\phi)^2 + \frac{\tau_\phi}{2}\phi^2 + \frac{1}{4}\phi^4 - \mu_\phi\phi + 2D(\nabla^2\psi)^2 - 2C(\nabla\psi)^2 + \frac{\tau_\psi}{2}\psi^2 + \frac{1}{4}\psi^4 - \mu_\psi\psi - \Lambda\phi\psi \right]. \quad (11)$$

The first five terms depend only on ϕ and its derivatives and describe the upper monolayer in Fig. 2 and its possible modulations. These are the same terms as in Eq. (10) for the single monolayer case. Similarly, the latter five ψ terms describe the lower monolayer, where τ_ϕ and τ_ψ are the two reduced temperatures, while μ_ϕ and μ_ψ are the corresponding chemical potentials. The last term, $\Lambda\phi\psi$, represents the coupling between the two leaflets with a coupling constant Λ .

We comment here on the physical origin of the proposed coupling term, $-\Lambda\phi\psi$, in Eq. (11). Note that this quadratic term is invariant under exchange of $\phi \leftrightarrow \psi$. When $\Lambda > 0$,

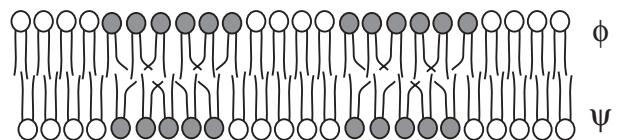


FIG. 2. Schematic illustration of two coupled modulated monolayers forming a bilayer membrane. Each monolayer is composed of a binary S/H lipid mixture, which can have a lateral composition modulation. The relative composition of S and H lipids in the upper and lower leaflets are defined by ϕ and ψ , respectively. The lipid tails interact across the bilayer midplane.

this term can be obtained from a term such as $+(\phi - \psi)^2$, which represents a local energy penalty when the upper and lower monolayers have different compositions [20,21]. For mixed lipid bilayers, such a coupling may result from the conformational confinement of the lipid chains and, hence, would have an entropic origin [20]. By estimating the degree of lipid chain interdigitation, the order of magnitude of the coupling parameter Λ can be evaluated [22]. In general, Λ can also be negative depending on the specific coupling mechanism [39]. However, the sign of Λ does not affect our results in any essential way because the free energy Eq. (11) is invariant under the combined exchange of either $\phi \rightarrow -\phi$ and $\Lambda \rightarrow -\Lambda$ or $\psi \rightarrow -\psi$ and $\Lambda \rightarrow -\Lambda$. Since such an effect only changes the sign of the chemical potential (either μ_ϕ or μ_ψ), it is sufficient to consider the $\Lambda > 0$ case.

The other possible higher-order coupling terms which are allowed by symmetry are $\phi^2\psi^2$, $\phi\psi^3$, and $\phi^3\psi$. However, these terms do not affect the properties of concentration fluctuations in any essential way and will not be considered hereafter.

III. CONCENTRATION FLUCTUATIONS ABOVE T_c

Using the bilayer free energy, Eq. (11), we obtain the static and dynamic structure factors, which describe the properties of concentration fluctuations for two coupled monolayers in their respective one-phase region (above T_c). We shall closely follow the formulation of Ref. [40] in which the coupled macro-phase separation was discussed for bilayers.

A. Static structure factor

The spatially varying $\phi(\mathbf{r})$ and $\psi(\mathbf{r})$ can be written as $\phi(\mathbf{r}) = \phi_0 + \delta\phi(\mathbf{r})$ and $\psi(\mathbf{r}) = \psi_0 + \delta\psi(\mathbf{r})$, respectively, where $\phi_0 = \langle \phi \rangle$ and $\psi_0 = \langle \psi \rangle$ are the spatially averaged monolayer compositions, and $\delta\phi$ and $\delta\psi$ describe the deviations from their average values. In thermal equilibrium, ϕ_0 and ψ_0 satisfy

$$\begin{aligned} \tau_\phi \phi_0 + \phi_0^3 - \mu_\phi - \Lambda \psi_0 &= 0, \\ \tau_\psi \psi_0 + \psi_0^3 - \mu_\psi - \Lambda \phi_0 &= 0. \end{aligned} \quad (12)$$

Expanding the free energy, Eq. (11), and retaining the quadratic order terms in $\delta\phi$ and $\delta\psi$, we obtain the Gaussian free energy

$$\begin{aligned} F_G[\delta\phi, \delta\psi] = \int d^2r \Bigg[&2B(\nabla^2 \delta\phi)^2 - 2A(\nabla \delta\phi)^2 \\ &+ \frac{\epsilon_\phi}{2}(\delta\phi)^2 + 2D(\nabla^2 \delta\psi)^2 - 2C(\nabla \delta\psi)^2 \\ &+ \frac{\epsilon_\psi}{2}(\delta\psi)^2 - \Lambda(\delta\phi)(\delta\psi) \Bigg], \end{aligned} \quad (13)$$

where the notations $\epsilon_\phi = \tau_\phi + 3\phi_0^2$ and $\epsilon_\psi = \tau_\psi + 3\psi_0^2$ as well as Eq. (12) are used.

The static partial structure factor is $S_{\phi\phi}(\mathbf{q}) = \langle \delta\phi(\mathbf{q})\delta\phi(-\mathbf{q}) \rangle$, and because of the radial symmetry, $S_{\phi\phi}(\mathbf{q}) = S_{\phi\phi}(q)$, where $q = |\mathbf{q}|$. It is given by

$$S_{\phi\phi}(q) = \frac{2g_\phi(q)}{4g_\phi(q)g_\psi(q) - \Lambda^2}, \quad (14)$$

and similarly for $S_{\psi\psi}$ and $S_{\phi\psi}$,

$$S_{\psi\psi}(q) = \frac{2g_\psi(q)}{4g_\phi(q)g_\psi(q) - \Lambda^2}, \quad (15)$$

$$S_{\phi\psi}(q) = S_{\psi\phi}(q) = \frac{\Lambda}{4g_\phi(q)g_\psi(q) - \Lambda^2}, \quad (16)$$

and

$$\begin{aligned} g_\phi(q) &= 2Bq^4 - 2Aq^2 + \frac{\epsilon_\phi}{2}, \\ g_\psi(q) &= 2Dq^4 - 2Cq^2 + \frac{\epsilon_\psi}{2}. \end{aligned} \quad (17)$$

Since the structure factors diverge at T_c , we see that the coupling parameter Λ effectively shifts the critical temperature to lower values.

B. Decoupled leaflets ($\Lambda = 0$)

When the two leaflets are decoupled, $\Lambda = 0$, it is sufficient to present results for only one of the two monolayers, say, the ϕ one. From Eq. (14), the decoupled structure factor is simply given by

$$S_{\phi\phi}(q) = \frac{1}{2g_\phi(q)}. \quad (18)$$

The correlation function $G_{\phi\phi}(\mathbf{r}) = \langle \delta\phi(\mathbf{r})\delta\phi(0) \rangle = G_{\phi\phi}(r)$, where $r = |\mathbf{r}|$, is obtained by use of the inverse 2D Fourier transform of Eq. (18) (see Appendix A for the derivation)

$$G_{\phi\phi}(r) = \frac{\xi_\phi \lambda_\phi}{32\pi B} \text{Re} \left[H_0^{(1)} \left(\frac{2\pi r}{\lambda_\phi} + i \frac{r}{\xi_\phi} \right) \right], \quad (19)$$

where $H_0^{(1)}(z)$ is the zeroth-order Hankel function of the first kind and “Re” denotes the real part. This correlation function contains two length scales; the first is the modulation periodicity

$$\frac{\lambda_\phi}{2\pi} = \left(\frac{B}{\epsilon_\phi} \right)^{1/4} \frac{2}{\sqrt{1 - \gamma_\phi}} \quad (20)$$

and the second is the correlation length

$$\xi_\phi = \left(\frac{B}{\epsilon_\phi} \right)^{1/4} \frac{2}{\sqrt{1 + \gamma_\phi}}, \quad (21)$$

where $\gamma_\phi = -A/\sqrt{\epsilon_\phi B}$.

When $-1 < \gamma_\phi < 1$, both λ_ϕ and ξ_ϕ are finite, and the corresponding phase is called the *structured-disordered* phase. In Fig. 3, we plot Eq. (19) for certain parameter values. The correlation function has exponentially decaying oscillations as a function of the distance r , similar to 3D microemulsions [28]. The peak of the structure factor $S_{\phi\phi}$ occurs at $q^* = \sqrt{A/2B} > 0$ for $-1 < \gamma_\phi < 0$, whereas $q^* = 0$ for $0 \leq \gamma_\phi < 1$. In the former case, the peak height is given by

$$S_{\phi\phi}(q_\phi^*) = \frac{1}{\epsilon_\phi - A^2/B} \quad (22)$$

and diverges at $\epsilon_\phi^* = A^2/B$. The line at $\gamma_\phi = 0$ is the *Lifshitz line*. The modulation periodicity λ_ϕ diverges for $\gamma_\phi \rightarrow 1$, and the monolayer transforms into a *disordered* phase for $\gamma_\phi > 1$.

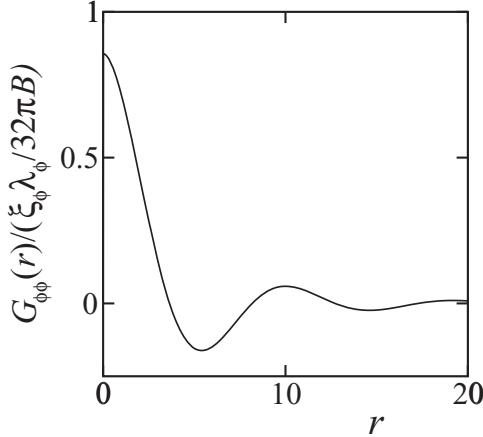


FIG. 3. The rescaled (dimensionless) real-space correlation function, $G_{\phi\phi}$, of a single monolayer in the structured-disordered phase as a function of r . The parameters are $B/\epsilon_\phi = 1$ and $\gamma_\phi = -0.9$, and the two resulting characteristic lengths are $\lambda_\phi/2\pi \simeq 1.45$ and $\xi_\phi \simeq 6.32$.

The line of $\gamma_\phi = 1$ is called the *disorder line* and is not a phase transition line. On the other hand, the correlation length ξ_ϕ diverges for $\gamma_\phi \rightarrow -1$ (hence, this is the critical point), and the *ordered* phase appears for $\gamma_\phi < -1$ [28].

As shown in Fig. 4(a), the phase diagram of a decoupled bilayer is easily obtained by combining the phase sequences in terms of the two independent parameters $\gamma_\phi = -A/\sqrt{\epsilon_\phi B}$ and $\gamma_\psi = -C/\sqrt{\epsilon_\psi D}$. The various phases are expressed by the binary combination of ordered (O), structured-disordered (S), and disordered (D) phases for each of the leaflets. The two disorder lines are shown in the figure by dashed lines, while the Lifshitz lines occur at $\gamma_\phi = 0$ and $\gamma_\psi = 0$ (not shown).

C. Coupled leaflets ($\Lambda \neq 0$)

When the two monolayers are coupled ($\Lambda \neq 0$), the corresponding phase diagram can be obtained by analyzing the poles in the complex plane of Eqs. (14)–(16). One example is shown in Fig. 4(b). The boundary between the OO and SS phases (solid line) is determined by the condition that all the structure factors diverge. In the DD phase, all the poles are pure imaginary, whereas in the SS phase the poles are complex, and at least one real pole exists in the OO phase.

In Fig. 4(b) we see that the asymmetric phases such as DO and SD are suppressed as compared with Fig. 4(a). This is because one of the leaflets induces modulations in the other leaflet due to the interleaflet coupling, Λ . In fact, all the asymmetric phases disappear for any finite value of Λ . In addition, the OO and SS phases are expanded as compared to Fig. 4(a). As mentioned before, increasing the coupling Λ effectively lowers the bilayer temperature. The Lifshitz lines (dotted lines) are obtained by plotting the main peak positions of Eqs. (14) and (15). Due to the coupling effect, they are tilted as compared to the $\Lambda = 0$ case where the lines occur at $\gamma_\phi = 0$ and $\gamma_\psi = 0$.

In Figs. 5(a) and 5(b), we plot the structure factors of the decoupled and coupled bilayers, respectively. As an illustration of the coupling effect with $\Lambda = 0.3$, we consider

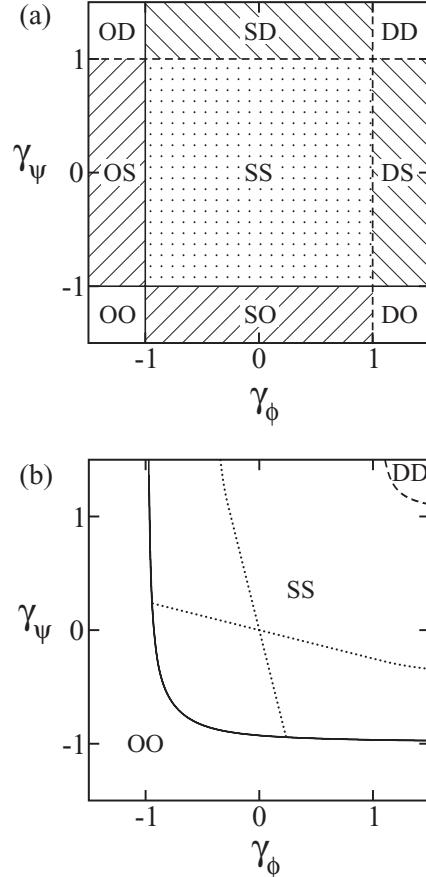


FIG. 4. (a) Phase diagram of a decoupled modulated bilayer ($\Lambda = 0$) plotted in the plane of $\gamma_\phi = -A/\sqrt{\epsilon_\phi B}$ and $\gamma_\psi = -C/\sqrt{\epsilon_\psi D}$. The phases are labeled using a binary combination of two letters representing phases in the ϕ and ψ monolayers. “O,” “S,” and “D” denote the ordered, structured-disordered, and disordered phases, respectively. Solid and dashed lines are the phase transition and disorder lines, respectively. (b) Phase diagram of a coupled modulated bilayer ($\Lambda = 1$) for $\epsilon_\phi = \epsilon_\psi = 2$, $B = D = 0.5$. In addition, the two dotted lines are the Lifshitz lines.

two different characteristic wavelengths, $q_\phi^* \neq q_\psi^*$. The heights of the two peaks are set to be equal by requiring $\epsilon_\phi = \epsilon_\psi$ and $A^2/B = C^2/D = 1$ [see Eq. (22)]. The peak height of $S_{\phi\phi}$ at $q_\phi^* = 1/\sqrt{2}$ is increased [as in Fig. 5(b)] due to the coupling effect, whereas that of $S_{\psi\psi}$ at $q_\psi^* = 3/\sqrt{2}$ is almost unchanged compared with the decoupled case. We also plot $S_{\phi\psi}$ given by Eq. (16) which represents the cross correlation of fluctuations between the two monolayers. This quantity is proportional to the coupling constant Λ and its peak position is essentially determined by that of $S_{\phi\phi}$ at $q_\phi^* = 1/\sqrt{2}$.

The peak position of $S_{\phi\psi}$ denoted as $q_{\phi\psi}^*$ for the coupled case ($\Lambda = 0.3$) is obtained numerically and plotted in Fig. 6(a) as a function of q_ϕ^* and q_ψ^* , which are, respectively, the peak positions of $S_{\phi\phi}$ and $S_{\psi\psi}$ for the decoupled case. We find that for the coupled case, the value of $q_{\phi\psi}^*$ is almost equal to that of the smaller of (q_ϕ^*, q_ψ^*) . In Fig. 6(b), the peak height of $S_{\phi\psi}$ at $q_{\phi\psi}^*$ is plotted as a function of the coupling parameter Λ and the ratio of the two characteristic wave numbers q_ψ^*/q_ϕ^*

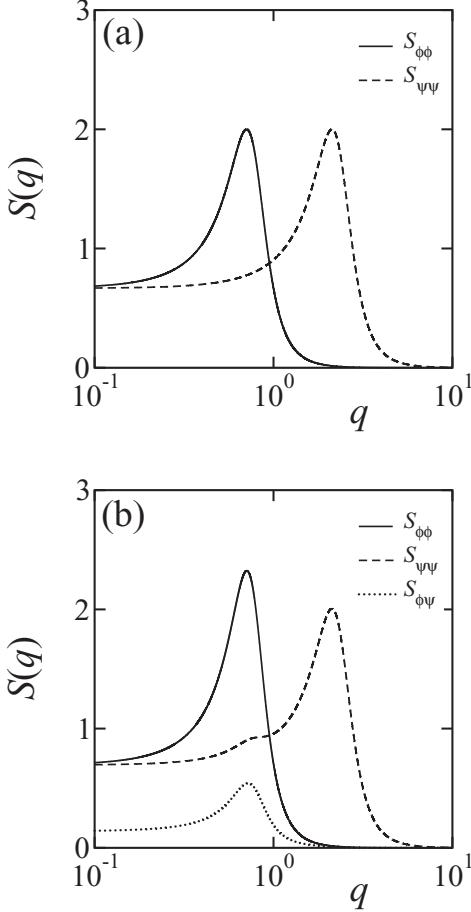


FIG. 5. Bilayer structure factors $S_{\phi\phi}$, $S_{\psi\psi}$, and $S_{\phi\psi}$ as a function of the wave number q for $\epsilon_\phi = \epsilon_\psi = 1.5$, $B = A = 1$, $D = 0.0123$, $C = 0.1111$ ($q_\phi^* = 1/\sqrt{2}$, $q_\psi^* = 3/\sqrt{2}$). (a) The decoupled case, $\Delta = 0$; (b) the coupled case with $\Delta = 0.3$.

for the specific case of $q_\phi^* = 1/\sqrt{2}$, as the parameters are fixed to $B = A = 1$ and $C^2/D = 1$. The peak height increases as Δ is increased because the temperature of the coupled bilayer is effectively lowered. To see this clearly, we have plotted in Fig. 6(b) the phase-transition line (dashed line) separating the structured-disordered and ordered phases. The peak height of $S_{\phi\psi}$ also increases as $q_\psi^*/q_\phi^* = \sqrt{2}q_\psi^*$ approaches unity, which is the case where $S_{\phi\phi}$ and $S_{\psi\psi}$ completely overlap.

D. Dynamic structure factors

The dynamical fluctuations in composition, $\delta\phi(\mathbf{r}, t)$ and $\delta\psi(\mathbf{r}, t)$, depend on time t and are now considered for coupled modulated bilayers. We assume that both $\delta\phi$ and $\delta\psi$ are conserved for each monolayer. This means that the exchange of lipids between the two monolayers is negligible. Such an assumption is justified when the typical time scale for lipid flip-flop motion is considerably large as compared to that for the domain formation (see Sec. V). Hence, the time evolution of $\delta\phi$ and $\delta\psi$ are given by the following diffusive equations

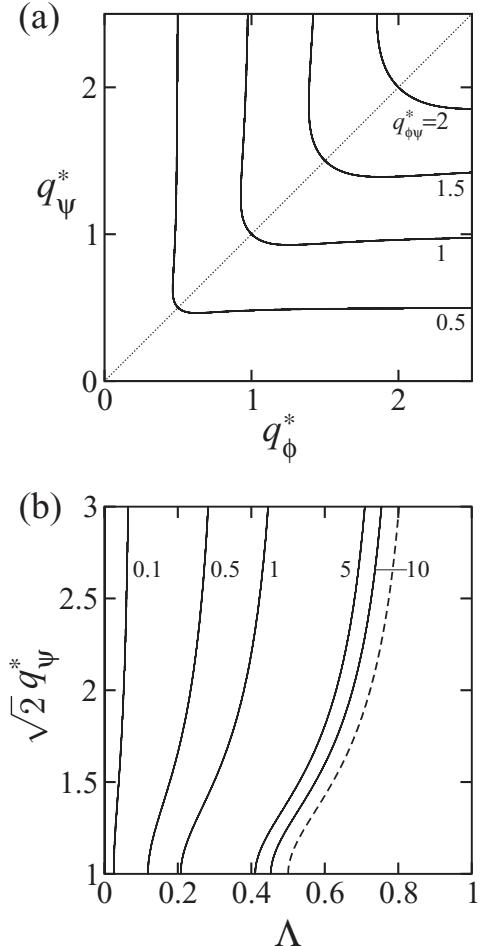


FIG. 6. (a) Contour plot of the peak position $q_{\phi\psi}^*$ of the structure factor $S_{\phi\psi}$ as function of $q_\phi^* = \sqrt{A/2B}$ and $q_\psi^* = \sqrt{C/2D}$. The parameters are $\Delta = 0.3$, $\epsilon_\phi = \epsilon_\psi = 1.5$, and $A^2/B = C^2/D$. The dotted line represents $q_\phi^* = q_\psi^*$. (b) Contour plot of the peak height of the structure factor $S_{\phi\psi}(q_{\phi\psi}^*)$ as a function of Δ and $\sqrt{2}q_\psi^*$. For the choice of parameters $B = A = 1$ and $C^2/D = 1$, $q_\phi^* = 1/\sqrt{2}$, the two characteristic wave numbers satisfy $q_\psi^*/q_\phi^* = \sqrt{2}q_\psi^*$. The dashed line is the phase-transition line.

for each monolayer:

$$\begin{aligned} \frac{\partial \delta\phi(\mathbf{r}, t)}{\partial t} &= L_\phi \nabla^2 \frac{\delta F_G}{\delta(\delta\phi)} + v_\phi(\mathbf{r}, t), \\ \frac{\partial \delta\psi(\mathbf{r}, t)}{\partial t} &= L_\psi \nabla^2 \frac{\delta F_G}{\delta(\delta\psi)} + v_\psi(\mathbf{r}, t). \end{aligned} \quad (23)$$

Here L_ϕ and L_ψ are the kinetic coefficients taken to be constants, and $v_i(\mathbf{r}, t)$ represent Gaussian white noise, satisfying $\langle v_i(\mathbf{r}, t) \rangle = 0$ and

$$\langle v_i(\mathbf{r}, t) v_j(\mathbf{r}', t') \rangle = -\delta_{ij} L_i \nabla^2 \delta(\mathbf{r} - \mathbf{r}') \delta(t - t'), \quad (24)$$

where $i, j = \phi, \psi$ and $\langle \dots \rangle$ now indicates the average over space and time. In the above equations, we have also neglected any hydrodynamic effects. In general, one has to take into account the fluid nature of both the bilayer membrane and the surrounding bulk solvent [41].

TABLE I. Expressions for $\alpha_{ij}(q)$ and $\beta_{ij}(q)$.

| $\alpha_{ij}(q)$ | $\beta_{ij}(q)$ |
|--|--|
| $\alpha_{\phi\phi} = \frac{2L_\phi q^2(\omega_\psi^2 - \omega_+^2)}{\omega_-^2 - \omega_+^2}$ | $\beta_{\phi\phi} = \frac{2L_\phi q^2(\omega_-^2 - \omega_\psi^2)}{\omega_-^2 - \omega_+^2}$ |
| $\alpha_{\psi\psi} = \frac{2L_\psi q^2(\omega_\phi^2 - \omega_+^2)}{\omega_-^2 - \omega_+^2}$ | $\beta_{\psi\psi} = \frac{2L_\psi q^2(\omega_-^2 - \omega_\phi^2)}{\omega_-^2 - \omega_+^2}$ |
| $\alpha_{\phi\psi} = \frac{2L_\phi L_\psi q^4 \Lambda \omega_{\phi\psi}}{\omega_-^2 - \omega_+^2}$ | $\beta_{\phi\psi} = -\frac{2L_\phi L_\psi q^4 \Lambda \omega_{\phi\psi}}{\omega_-^2 - \omega_+^2}$ |

The Fourier transform of fluctuations in both space and time is

$$\delta\phi(\mathbf{q}, \omega) = \int d^2r dt \delta\phi(\mathbf{r}, t) e^{-i(\mathbf{q}\cdot\mathbf{r}-\omega t)}, \quad (25)$$

and a similar Fourier transform is used for $\delta\psi$ and ν_i . The dynamic structure factors such as $S_{\phi\phi}(\mathbf{q}, \omega) = \langle \delta\phi(\mathbf{q}, \omega) \delta\phi(-\mathbf{q}, -\omega) \rangle$ are given by

$$S_{ij}(q, \omega) = \frac{\alpha_{ij}(q)}{\omega^2 + [\omega_+(q)]^2} + \frac{\beta_{ij}(q)}{\omega^2 + [\omega_-(q)]^2}, \quad (26)$$

where the explicit expressions of $\alpha_{ij}(q)$ and $\beta_{ij}(q)$ are given in Table I and the characteristic frequencies $\omega_{\pm}(q)$ are

$$[\omega_{\pm}(q)]^2 = \frac{1}{2} \left[\omega_\phi^2(q) + \omega_\psi^2(q) \right. \\ \left. \mp \sqrt{[\omega_\phi^2(q) - \omega_\psi^2(q)]^2 + 4L_\phi L_\psi q^4 \Lambda^2 \omega_{\phi\psi}^2(q)} \right], \quad (27)$$

$$\omega_\phi^2(q) = 4L_\phi^2 q^4 [g_\phi(q)]^2 + L_\phi L_\psi q^4 \Lambda^2, \quad (28)$$

$$\omega_\psi^2(q) = 4L_\psi^2 q^4 [g_\psi(q)]^2 + L_\phi L_\psi q^4 \Lambda^2, \quad (29)$$

$$\omega_{\phi\psi}(q) = 2q^2 [L_\phi g_\phi(q) + L_\psi g_\psi(q)], \quad (30)$$

and g_ϕ and g_ψ are defined in Eq. (17). In the more special case of $L_\phi = L_\psi = L$, ω_{\pm} reduce to a simpler form,

$$\omega_{\pm}(q) = Lq^2 \left[g_\phi(q) + g_\psi(q) \mp \sqrt{(g_\phi(q) - g_\psi(q))^2 + \Lambda^2} \right]. \quad (31)$$

The intermediate structure factors $S_{ij}(q, t)$ depend explicitly on time and are obtained by taking the inverse Fourier transform in t :

$$S_{ij}(q, t) = \int \frac{d\omega}{2\pi} S_{ij}(q, \omega) e^{-i\omega t}, \quad (32)$$

and from Eq. (26), we obtain

$$S_{ij}(q, t) = \frac{\alpha_{ij}(q)}{2\omega_+(q)} e^{-\omega_+(q)t} + \frac{\beta_{ij}(q)}{2\omega_-(q)} e^{-\omega_-(q)t}. \quad (33)$$

Hence, the decay of concentration fluctuations is described by a sum of two exponentials with two characteristic times

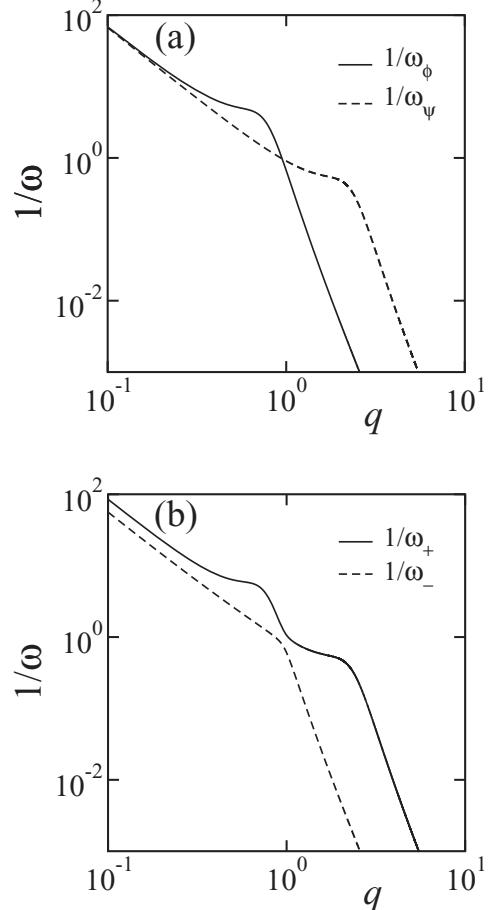


FIG. 7. Decay time of concentration fluctuations as a function of the wave number q for $\epsilon_\phi = \epsilon_\psi = 1.5$, $B = A = 1$, $D = 0.0123$, $C = 0.1111$ (resulting in $q_\phi^* = 1/\sqrt{2}$ and $q_\psi^* = 3/\sqrt{2}$), and $L = 1$. (a) The decoupled case, $\Lambda = 0$; (b) the coupled case with $\Lambda = 0.3$.

$1/\omega_{\pm}(q)$. However, when the two monolayers are decoupled ($\Lambda = 0$), the $S_{\phi\phi}$ and $S_{\psi\psi}$ structure factors decay with a single exponential characterized by a decay rate ω_ϕ and ω_ψ [see Eqs. (28) and (29)], respectively.

For the decoupled case ($\Lambda = 0$), the decay times $1/\omega_\phi$ and $1/\omega_\psi$ are plotted as a function of q in Fig. 7(a), with the same parameters as those in Fig. 5. The plots show a shoulder reflecting the characteristic structure at wave numbers $q_\phi^* = 1/\sqrt{2}$ and $q_\psi^* = 3/\sqrt{2}$. Notice that the larger the initial length scale, the longer the decay time. For the coupled case with $\Lambda = 0.3$, we plot $1/\omega_{\pm}$ in Fig. 7(b). Due to the coupling effect, the two decay times split into a larger and smaller one, $1/\omega_+ > 1/\omega_-$. The larger one, $1/\omega_+$, exhibits two shoulders, while the smaller one, $1/\omega_-$, has a shoulder between the two characteristic wave numbers. The coupling affects the decay time of the structure corresponding to the smaller wave number (larger length), similar to the effect seen in Fig. 5 for the static structure factor.

IV. COUPLED MODULATED PHASES WITH DIFFERENT PERIODICITIES $q_\phi^* \neq q_\psi^*$

In our previous paper [29], we investigated the phase behavior of coupled modulated bilayers by using the free

energy, Eq. (11), below T_c . When the two monolayers have the same preferred periodicity, $q_\phi^* = q_\psi^*$, we obtained the mean-field phase diagram exhibiting various combinations of modulated structures such as stripe (S) and hexagonal (H) phases. In some cases, the periodic structure in one of the monolayers induces a similar modulation in the second monolayer. Moreover, the region of the induced modulated phase expands as the coupling parameter Λ becomes larger.

However, when the preferred periodicities in the two leaflets are different, $q_\phi^* \neq q_\psi^*$, it is difficult to obtain the phase diagram because the free-energy densities cannot be obtained analytically. We then have to rely on numerical simulations [29,42] to solve the time evolution of the two coupled order parameters as explained below.

For the dynamics of coupled modulated bilayers below T_c , we use the following time evolution equations [29]

$$\frac{\partial\phi(\mathbf{r},t)}{\partial t} = L_\phi \nabla^2 \frac{\delta F_b}{\delta\phi}, \quad \frac{\partial\psi(\mathbf{r},t)}{\partial t} = L_\psi \nabla^2 \frac{\delta F_b}{\delta\psi}, \quad (34)$$

where the bilayer free energy, F_b , is given by Eq. (11). As in Eq. (23), we have assumed that both ϕ and ψ are conserved for each monolayer and neglect any hydrodynamic effects. Below T_c the effect of thermal fluctuations is less important and the noise terms have been omitted in the above equations.

Hereafter, the kinetic coefficients L_ϕ and L_ψ are set to unity for simplicity.

We solve numerically the above 2D equations using periodic boundary conditions. Each run starts from a homogeneous state with a small random noise around the average compositions ϕ_0 and ψ_0 . Time is measured in discrete time steps, and $t = 5000$ corresponds to a well equilibrated system. In all our simulations, the reduced temperature parameters are fixed to be $\tau_\phi = \tau_\psi = 0.8$. The characteristic wave number in the ϕ monolayer is fixed as $q_\phi^* = 1/\sqrt{2}$ by setting $B = A = 1$, while the periodicity in the ψ monolayer $q_\psi^* = \sqrt{C/2D}$ is varied and the condition $C^2/D = 1$ is used as before. In the following, we shall consider only the two coupled stripe phases for $\phi_0 = \psi_0 = 0$. However, the combination of stripe and hexagonal phases leads to a rich variety of complex patterns [42].

When the characteristic wave numbers of the two decoupled monolayers differ, $q_\phi^* \neq q_\psi^*$, the two coupled modulated structures cannot apparently match each other. The frustration between the two different periodicities is due to the interleaflet coupling and affects their morphologies. In Fig. 8, we show one example of the time evolution of a coupled micro-phase separation for $q_\psi^*/q_\phi^* = 3$ ($q_\phi^* = 1/\sqrt{2}$ and $q_\psi^* = 3/\sqrt{2}$) and $\Lambda = 0.3$. The spatial patterns of ϕ , ψ , $\phi + \psi$, $\phi - \psi$, and the Fourier transformed pattern, $\psi(\mathbf{q})$, are presented for time steps of $t = 25$, 60 , 250 , and 5000 . Starting from

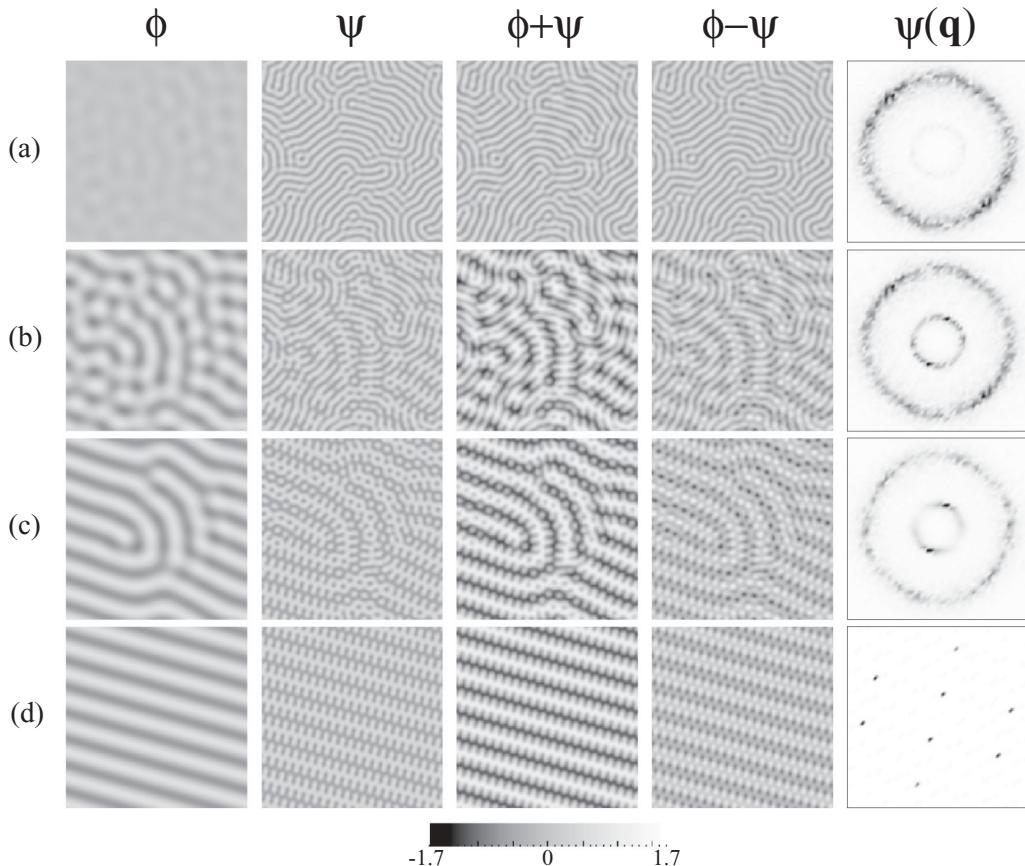


FIG. 8. Time evolution of spatially modulated patterns in a coupled bilayer, which consists of two stripe monolayers with different periodicities, $q_\phi^* \neq q_\psi^*$. Real-space patterns of ϕ , ψ , $\phi + \psi$, $\phi - \psi$, as well as the Fourier transform, $\psi(\mathbf{q})$, are presented. The time steps are (a) $t = 25$, (b) $t = 60$, (c) $t = 250$, and (d) $t = 5000$. The parameters are $\phi_0 = \psi_0 = 0$, $\tau_\phi = \tau_\psi = 0.8$, $B = A = 1$, $D = 0.0123$, $C = 0.1111$, and $\Lambda = 0.3$, yielding $q_\psi^*/q_\phi^* = 3$. In order to emphasize the color contrast, the gray color code of the real-space patterns is chosen to vary between -1.7 and 1.7 .

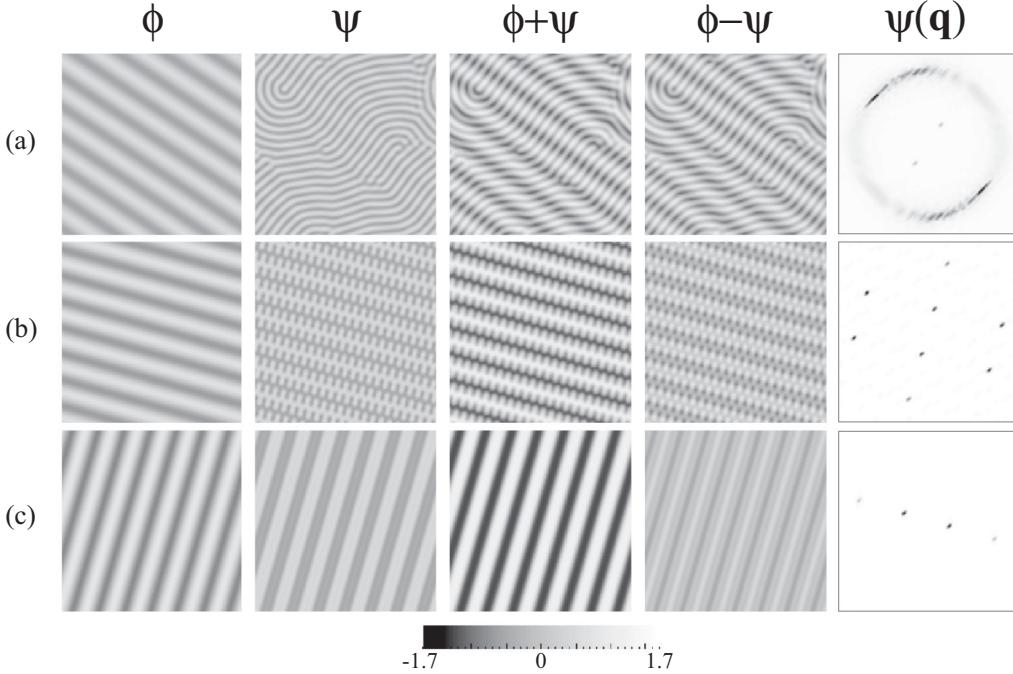


FIG. 9. Equilibrated patterns (time step $t = 5000$) of two coupled bilayers consisting of stripe monolayers with different periodicities. All parameters in addition to Λ are the same as in Fig. 8. The Λ parameter values are (a) $\Lambda = 0.1$; (b) $\Lambda = 0.3$; (c) $\Lambda = 0.5$. In order to emphasize the color contrast, the color code of the real-space patterns is chosen to vary between -1.7 and 1.7 .

the isotropic state, the ψ monolayer forms first stripes (a). Then, as the ϕ monolayer starts to segregate at larger t , it simultaneously chops the ψ stripes into smaller sections (b), and the ψ monolayer transforms into a fingerlike patterns (c). Reconnection of the ϕ stripes takes place after a long time of annealing, and a pattern of alternating fingers in the ψ monolayer is finally obtained (d). From the time evolution of the 2D Fourier patterns of the ψ monolayer, it is apparent that the intermediate structures are characterized by the two length scales of ratio 1:3.

In Fig. 9, we show the spatially modulated patterns at $t = 5000$ of the two coupled ϕ and ψ monolayers with different periodicities ($q_\phi^* = 1/\sqrt{2}$ and $q_\psi^* = 3/\sqrt{2}$ as before) for different values of the coupling parameter $\Lambda = 0.1, 0.3$, and 0.5 . Notice that Fig. 9(b) is the same as Fig. 8(d). In the weak-coupling case ($\Lambda = 0.1$), the two monolayers exhibit two independent stripe morphologies with characteristic periodicities (called the “independent” morphology). Here the two stripes essentially do not affect each other. As the coupling constant is increased ($\Lambda = 0.3$), stripes with a fingerlike structure appear in the ψ monolayer, while the stripe morphology in the ϕ monolayer is almost unaffected (called the “intermediate” morphology). In the Fourier transformed pattern of ψ , we clearly see that the structures with two different characteristic wave numbers are coexisting. This result is also in accord with the properties of the static structure factors, $S_{ij}(q)$, having two different characteristic lengths, as shown in Fig. 5. For a larger coupling parameter ($\Lambda = 0.5$), very similar patterns are obtained for ϕ and ψ and almost coincide with one another (called the “coincident” morphology). It should be noted that the structure with the larger wavelength dominates when the coupling is large

enough, and the sequence of morphological changes shown in Fig. 9, as Λ increases, is rather typical.

In order to quantify the three morphologies (independent, intermediate, and coincident), we calculate the spatial average of the product of the two compositions

$$\langle \phi\psi \rangle = \frac{1}{A} \int d^2r \phi(\mathbf{r}, t)\psi(\mathbf{r}, t), \quad (35)$$

where A is the total system area. In Fig. 10, we plot $\langle \phi\psi \rangle$ at $t = 1000$ (sufficient for equilibration) for various combinations of Λ and $q_\psi^*/q_\phi^* = \sqrt{2}q_\psi^*$ (for the case $q_\phi^* = 1/\sqrt{2}$). The morphology of the obtained patterns is marked on the figure by circles, crosses, and squares. The values of $\langle \phi\psi \rangle$ are small for “independent” (circles) structures, while they become larger for the “intermediate” (crosses) and “coincident” (squares) morphologies. Although the morphological changes are gradual and do not represent a sharp transition, the intermediate patterns appear roughly for $0.05 < \langle \phi\psi \rangle < 0.2$. The region of intermediate structure expands as $q_\psi^*/q_\phi^* = \sqrt{2}q_\psi^*$ increases, and the patterns coincide for $\Lambda = 0.5$. We note that although the morphology cannot be solely determined by the quantity $\langle \phi\psi \rangle$, the behavior of $\langle \phi\psi \rangle$ is similar to that of the peak values of the cross correlation, $S_{\phi\psi}(q_{\phi\psi}^*)$, presented in Fig. 6(b).

For the quantitative argument concerning the micro-phase separation dynamics in each monolayer, we have also calculated the two self-quantities,

$$\langle \phi^2 \rangle = \frac{1}{A} \int d^2r \phi^2(\mathbf{r}, t) \quad (36)$$

and

$$\langle \psi^2 \rangle = \frac{1}{A} \int d^2r \psi^2(\mathbf{r}, t). \quad (37)$$

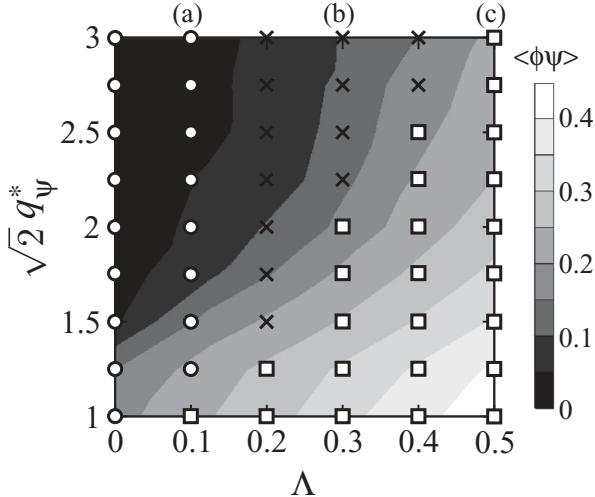


FIG. 10. Contour plot of $\langle \phi\psi \rangle$ at $t = 1000$ as a function of Λ and $\sqrt{2}q_\psi^*$, for the specific case of $q_\phi^* = 1/\sqrt{2}$. (a), (b), and (c) correspond to $\Lambda = 0.1, 0.3$, and 0.5 , respectively, as given in Fig. 9. Circles, crosses, and squares correspond, respectively, to independent, intermediate, and coincident structures (see text).

In Fig. 11, we plot the square root of these quantities as a function of t , with the same parameters as those used in Fig. 9. In all studied cases, the modulation of the ψ monolayer having a larger wave number ($q_\psi^* > q_\phi^*$) grows faster than that of the ϕ monolayer. We also remark that the structure formation in the ϕ monolayer is accelerated for larger Λ , whereas that of the ψ monolayer is almost unchanged. According to the linear stability analysis of Eq. (34), the initial growth rates of the unstable modes are essentially equivalent to the decay rates ω_\pm of the concentration fluctuations given in Eq. (31). This is consistent with Fig. 7, where the characteristic growth time ($1/\omega$) for larger q is smaller than that for smaller q . The growth rates increase with Λ [see Fig. 11(a)] because the coupling effectively reduces the temperature and enhances the phase transition. The faster the decay of concentration fluctuations, the faster the structure formation.

V. DISCUSSION AND FINAL REMARKS

In this paper we present a model for coupled modulated lipid bilayers. We start by considering a monolayer consisting of a mixture of saturated and hybrid lipids and propose a phenomenological model that includes a coupling between the lipid composition and a 2D vectorial field. This coupling arises from the line-active nature of the hybrid lipid, which adjusts its tail orientation in order to reduce the line tension. Minimization of the monolayer free energy with respect to the vectorial field yields a 2D microemulsion exhibiting modulated phases. The characteristic wavelength of modulation is determined by the monolayer coefficients A and B , Eq. (8), reflecting molecular properties of lipid mixture. We then construct a model for lipid bilayers composed of two modulated monolayers that influence each other through an interleaflet coupling.

Based on the model, we study concentration fluctuations of bilayers above T_c and calculate their static structure factors. The calculated phase diagram for coupled bilayers shows

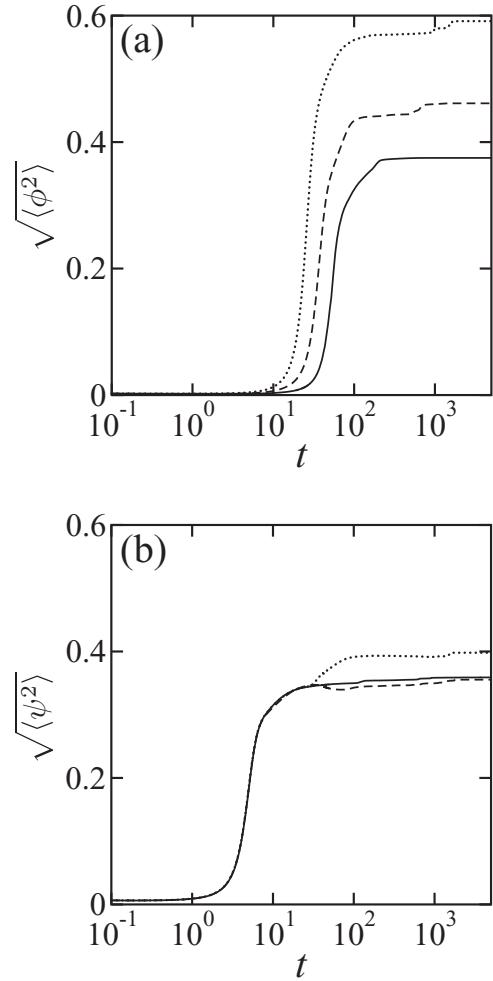


FIG. 11. Time evolution of the modulation amplitudes of the two monolayers: (a) $\sqrt{\langle \phi^2 \rangle}$ and (b) $\sqrt{\langle \psi^2 \rangle}$. The parameters are $\tau_\phi = \tau_\psi = 0.8$, $\phi_0 = \psi_0 = 0$, $B = A = 1$, $D = 0.0123$, $C = 0.1111$ yielding $q_\psi^*/q_\phi^* = 3$. The different lines represent $\Lambda = 0.1$ (solid), $\Lambda = 0.3$ (dashed), and $\Lambda = 0.5$ (dotted).

that the extent of ordered and structured-disordered phases become larger as compared to the decoupled case. When the two monolayers with different preferred wave numbers $q_\phi^* \neq q_\psi^*$ are coupled (say, $q_\phi^* < q_\psi^*$), the peak height of $S_{\phi\phi}$ occurring at smaller q numbers becomes larger as compared to the decoupled case, whereas the peak height of $S_{\psi\psi}$ occurring at larger q numbers almost does not change. Namely the interleaflet coupling strongly affects the compositional modulation in each monolayer. Furthermore, the interleaflet coupling has a clear signature on the cross structure factor, $S_{\phi\psi}$, as well as on the dynamics of concentration fluctuations. By calculating the intermediate structure factor, $S(q, t)$, we show that concentration fluctuations generally exhibit a double exponential decay with two decay rates, ω_\pm . One of the decay times ($1/\omega_+$) exhibits two shoulders at wave numbers describing the monolayer compositional modulations.

For membranes below T_c , we studied the micro-phase separation of a coupled modulated bilayer. When the two

monolayers have different modulations, $q_\phi^* \neq q_\psi^*$, we obtained numerically a variety of complex patterns. The initial growth rates of the unstable modes are identical to the decay rates of the concentration fluctuations.

As mentioned in Sec. III, the special character of concentration fluctuations in our model may explain the finite-size domains (“rafts”) in biological membranes. For ordinary binary mixtures above T_c , the only length scale of the disorder phase is determined by the correlation length, and close to T_c , this length becomes large. But within our 2D microemulsion model, there is another length scale characterizing the modulations as given by Eq. (20). This second length scale may also explain the finite-size domains observed in some experiments as a result of micro-phase separation in the low temperature phase.

We note that our microemulsion picture will not hold if we replace the hybrid lipids with unsaturated lipids (having two unsaturated chains) because unsaturated lipids would not exhibit any line-activity. For mixtures containing unsaturated lipids, we expect to observe macro-phase separation instead of micro-phase separation, in agreement with some experiments [5]. In biological membranes, however, unsaturated lipids are not commonly found and are present only as minor components. The experimental situation using artificial membranes is somewhat less clear because it was shown in some experiments [43,44] that membranes containing a mixture of saturated lipids, hybrid lipids and cholesterol show nanoscopic domains, whereas in other works [6], a macroscopic coarsening of domains was observed for such mixtures.

A related model based on a microemulsion picture was recently proposed by Schick [45], who considered a coupling between curvature and compositional asymmetry between the two leaflets, resulting in a 2D microemulsion. Although Schick’s model as well as ours share the microemulsion viewpoint, the origin of the physical mechanism differs. In Schick’s model the coupling between composition asymmetry and curvature gives rise to domains with different spontaneous curvature. Our model, on the other hand, assumes that for flat monolayers, the line-active nature of hybrid lipids is solely responsible for the microemulsion formation [see Eq. (3)]. Hence, our proposed physical mechanism for the bilayer coupling, as discussed in Sec. II B, differs. In our model, the domains residing on the two leaflets are either correlated for $\Lambda > 0$ or anticorrelated for $\Lambda < 0$, whereas they are always anticorrelated in the model of Ref. [45].

The present work is concerned with the analysis of equilibrium properties and relaxation dynamics toward the equilibrium state. The existence of finite-size domains may also be explained by nonequilibrium lipid transport between the cell interior and the membrane. Such a mechanism was considered [46–49] through a coupling between the membrane and an outer reservoir of lipid or cholesterol. Similarly to our model, these works explain the appearance of the finite-size domains as a result of micro-phase separation.

Asymmetry between lipid composition in the inner and outer leaflets of biomembranes has a very deep significance and is closely related to the cell biological functions. For instance, the breakdown of such compositional asymmetry is related to programmed cell death (apoptosis) [50]. For

living cells, this asymmetry is maintained by an enzyme called “flippase” that actively transports the lipids between the two leaflets [51]. The half-life of lipid composition due to flip-flop motion is measured using time resolved small angle neutron scattering [52] and is estimated to be several hours at physiological conditions.

Although membranal signal transduction is important for various biological functions, its dynamical properties are still not so well understood. As our theory offers an explanation for the size and dynamics of lipid domains, we hope that it and similar models will contribute in the future toward the understanding of functional processes in biomembranes.

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APPENDIX: DERIVATION OF $G_{\phi\phi}(r)$

In this Appendix we present the derivation of $G_{\phi\phi}$ of Eq. (19). The real-space correlation function is given by the 2D inverse Fourier transform of Eq. (18):

$$\begin{aligned} G_{\phi\phi}(r) &= \int \frac{d^2 q}{(2\pi)^2} S_{\phi\phi}(q) e^{i\mathbf{q}\cdot\mathbf{r}} \\ &= \frac{1}{4\pi B} \int_0^\infty dq \frac{q J_0(qr)}{q^4 - (A/B)q^2 + \epsilon_\phi/4B}, \end{aligned} \quad (\text{A1})$$

where $J_0(qr)$ is the zeroth-order Bessel function. We use the relation $J_0(qr) = [H_0^{(1)}(qr) + H_0^{(2)}(qr)]/2$, where $H_0^{(1)}(qr)$ and $H_0^{(2)}(qr)$ are the zeroth-order Hankel functions of the first and second kinds, respectively. The integral in Eq. (A1) then is written as $I = (I_1 + I_2)/2$, where

$$I_i = \int_0^\infty dq \frac{q H_0^{(i)}(qr)}{q^4 - (A/B)q^2 + \epsilon_\phi/4B}, \quad (\text{A2})$$

with $i = 1, 2$.

In order to evaluate the above integral, we performed the integration in the complex plane by replacing q with the complex variable $z = x + iy$. The integrand has four poles at

$$z_j = \frac{(\epsilon_\phi/B)^{1/4}}{2} (\pm\sqrt{1-\gamma_\phi} \pm i\sqrt{1+\gamma_\phi}) = \pm \frac{2\pi}{\lambda_\phi} \pm i \frac{1}{\xi_\phi}, \quad (\text{A3})$$

with $j = 1, 2, 3, 4$ and λ_ϕ and ξ_ϕ have been defined in Eqs. (20) and (21). These poles are located in the quadrants 1, 2, 3, and 4, off the real and imaginary axes. For the integral I_1 , we integrate along a quarter-circle contour in the first quadrant in an counterclockwise direction. The contour radius is taken to infinity (Fig. 12), and, using the residue theorem, we obtain

$$\int_0^\infty dx \frac{x H_0^{(1)}(xr)}{x^4 - (A/B)x^2 + \epsilon_\phi/4B} + \int_{-\infty}^0 dy \frac{i y H_0^{(1)}(iy r)}{y^4 + (A/B)y^2 + \epsilon_\phi/4B} = 2\pi i \text{Res}|_{z_1}, \quad (\text{A4})$$

where “Res” denotes the residue. Similarly, for the integral I_2 , we integrate along the contour of the quarter-circle of infinite radius in the fourth quadrant in the clockwise direction,

$$\int_0^\infty dx \frac{x H_0^{(2)}(xr)}{x^4 - (A/B)x^2 + \epsilon_\phi/4B} + \int_{-\infty}^0 dy \frac{i y H_0^{(2)}(iy r)}{y^4 + (A/B)y^2 + \epsilon_\phi/4B} = -2\pi i \text{Res}|_{z_4}. \quad (\text{A5})$$

Combining Eqs. (A4) and (A5), and further using the relation $H_0^{(1)}(-z) = -H_0^{(2)}(\bar{z})$, we obtain

$$I = \frac{\pi}{2\sqrt{\epsilon_\phi/B}\sqrt{1-\gamma_\phi^2}} [H_0^{(1)}(z_1 r) + H_0^{(2)}(z_4 r)]. \quad (\text{A6})$$

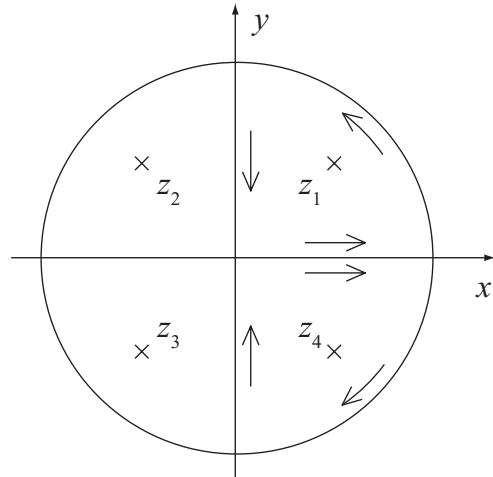


FIG. 12. Four poles in the complex plane at $z = z_j$ ($j = 1, 2, 3$, and 4). The two close integral paths are indicated by the arrows in the first and fourth quadrants.

Finally, using λ_ϕ and ξ_ϕ from Eq. (A3), we obtain

$$G_{\phi\phi}(r) = \frac{\xi_\phi \lambda_\phi}{32\pi B} \text{Re} \left[H_0^{(1)} \left(\frac{2\pi r}{\lambda_\phi} + i \frac{r}{\xi_\phi} \right) \right]. \quad (\text{A7})$$

In the above expressions, we have used the relation $H_0^{(2)}(\bar{z}) = \overline{H_0^{(1)}(z)}$, where \bar{z} is the complex conjugate of z .

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