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**Influence of Molecular Coherence on Surface Viscosity**

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**ABSTRACT:** Adding small fractions of cholesterol decreases the interfacial viscosity of dipalmitoylphosphatidylcholine (DPPC) monolayers by an order of magnitude per wt %. Grazing incidence X-ray diffraction shows that cholesterol at these small fractions does not mix ideally with DPPC but rather induces nanophase separated structures of an ordered, primarily DPPC phase bordered by a line-active, disordered, mixed DPPC-cholesterol phase. We propose that the free area in the classic Cohen and Turnbull model of viscosity is inversely proportional to the number of molecules in the coherence area, or product of the two coherence lengths. Cholesterol significantly reduces the coherence area of the crystals as well as the interfacial viscosity. Using this free area collapses the surface viscosity data for all surface pressures and cholesterol fractions to a universal logarithmic relation. The extent of molecular coherence appears to be a fundamental factor in determining surface viscosity in ordered monolayers.

**INTRODUCTION**

Despite its ubiquity in cell membranes, there is little quantitative information on the effects of small mole fractions of cholesterol on the phase diagrams, molecular organization, and surface viscosity of mixtures of phospholipids and cholesterol.1,2 Understanding cholesterol’s effects in altering the local molecular organization and rheology of lipid monolayers may give clues to the mechanisms that stabilize nanometer-scale “rafts” in complex lipid-cholesterol mixtures; the raft hypothesis has emerged in recent years as a general organizing principle for the structure of eukaryotic cell membranes.3–6 Rafts are hypothesized to result from nanometer-scale phase separation of ordered and viscous domains in which cholesterol, saturated lipids, and membrane proteins preferentially accumulate, surrounded by a sea of less viscous, unsaturated lipids with greater protein diffusivity.3–6 However, the fundamental physics underlying raft formation is still being developed, especially how interactions between saturated phospholipids and cholesterol determine membrane phase behavior, viscosity and diffusivity.4

The interactions of cholesterol and saturated phospholipids also play an important, but poorly understood, role in the properties of human lung surfactant (LS), a lipid-protein monolayer necessary to reduce the surface tension in the lung alveoli.7 At present, even the existence of cholesterol in native LS is questioned, as the lung lavage required to harvest LS inevitably causes blood and cell debris to be coextracted, potentially contaminating LS with cholesterol. This lack of consensus is reflected in the composition of replacement lung surfactants, which are used to treat neonatal respiratory distress syndrome (NRDS), which occurs in 20 000–30 000 premature infants each year in the U.S.7,8 Survanta and Curosurf, two clinically approved animal extract replacement surfactants for treatment of NRDS, have all cholesterol removed after harvesting. Infasurf, the third clinically approved surfactant, retains 4–5 wt % cholesterol.9 Resolving this controversy is potentially contaminating LS with cholesterol. This lack of consensus is re...
year in the U.S. with a mortality rate of 30–40%. The pathogenesis of ARDS is not fully understood, but in both ALI and ARDS, surfactant is “inactivated” by some primary pathogenesis such as lung inflammation, trauma, pulmonary infection, near-drowning, etc. The cholesterol content of ARDS patients shows an increase in cholesterol content and in vitro, cholesterol levels greater than 10 mol % increase the minimum surface tension at monolayer collapse, which may exacerbate lung damage.

Grazing incidence X-ray diffraction (GIXD) shows that up to 7 mol % added cholesterol does not change the basic alkane packing of DPPC, but does decrease the extent of ordering, or coherence area, suggesting an increased number of lattice defects. We postulate that the free area available for diffuse transport in a two-dimensional analog of the classic Cohen and Turnbull free volume model of viscosity is inversely proportional to the number of molecules in a coherence area. Using this relationship, the surface viscosity data for all surface pressures and cholesterol fractions collapses to a simple logarithmic relation with no adjustable parameters. This suggests that the decreased molecular ordering caused by the incompatibility of cholesterol with the alkane chain lattice is the origin of the orders of magnitude decrease in surface viscosity.

**MATERIALS AND METHODS**

**Isotherms and Fluorescence Imaging.** 1,2-Dipalmitoyl-sn-glycero-3-phosphocholine (DPPC, R-enantiomer) and dihydrocholesterol (Avanti, Alabaster, AL) with 0.1 wt % Texas-Red DHPE (N- (Texas Red sulfonyl)-1,2-dihexadecanoyl-sn-glycero-3-phosphoethanolamine, Invitrogen, Grand Island, NY) were mixed in the appropriate ratios and diluted to ~0.2 mg/mL in HPLC-grade chloroform (Fisher Scientific, St. Louis, MO) to form a spreading solution. Dihydrocholesterol (Chol) was used instead of cholesterol to minimize oxidation but has little impact on phase behavior. Surface pressure—area isotherms were recorded at 25 °C using a Teflon trough (NIMA, Coventry, England) with custom designed stainless steel ribs which reduce film leakage at high surface pressures. A filter paper Wilhelmy plate (Riegler and Kirstein, GmbH; Potsdam, Germany) was used to measure surface pressure. The open area of the trough used was 125 cm² and each complete compression/expansion cycle took about 8 min (0.42 cm²/s). For fluorescence imaging, the Langmuir trough was mounted on a Nikon Optiphot optical microscope with a custom designed stage equipped with long working distance objectives designed for fluorescent light. A dichroic mirror/barrier filter assembly directed the excitation light onto the monolayer films at a normal angle of incidence and filtered the emitted light and the images were detected by a silicon intensified CCD camera. Videos of the monolayer film were recorded during the compression-expansion cycle directly onto the computer using the Pinnacle studio capture software. The continuous, fluid lipid phase appears bright due to the preferential segregation of the Texas Red dye, while the better ordered domains exclude the dye molecules and appear dark.

**Grazing Incidence X-ray Diffraction (GIXD).** Two-dimensional GIXD experiments were carried out at the ChemMatCARS station at beamline 15-ID at the Advanced Photon Source, Argonne National Laboratory. DPCC/Chol at the appropriate ratios were dissolved in chloroform solution at ~1 mg/mL and spread dropwise onto the air/deionized water interface in a custom Langmuir trough, which was temperature-controlled at 22°C. After waiting ~30 min for solvent evaporation, the monolayers were compressed to the desired surface pressure (20, 30, or 40 mN/m) and annealed for an additional 30 min. The trough was enclosed in a helium-filled chamber and the oxygen level was constantly monitored during exposure to the X-ray beam. The analysis of GIXD data for two-dimensional films at the air-water interface follows that of Kaganer et al. It is well established that the Bragg peaks correspond to ordering within the alkane chains of the lipid tailgroups, the organization of the headgroups is inferred from changes in the tailgroup lattice in response to changes in surface pressure and cholesterol fraction.

**Interfacial Microscopy.** Circular ferromagnetic probes (microbuttons) of diameter 20 μm, thickness 1 μm, with “button holes” of diameter 3.5 μm were fabricated by photolithography. The probes were deposited at the air—water interface in a custom-built trough with an attached fluorescence microscope. A uniform magnetic field of magnitude, B, and orientation, θ, was generated by the output of two independent pairs of electromagnets controlled by a custom LabVIEW code to exert a controlled torque, L, on a microbutton of moment m and orientation φ. To measure the frequency-dependent linear viscoelastic response, a sinusoidal magnetic field was applied to generate a time varying applied torque. The driving torque was kept small enough that the response was linear. The microbutton orientation was determined from bright field images of the holes in the microbuttons as a function of applied torque, to determine the rotational resistance. From the rotational resistance, the linear viscoelastic surface moduli were obtained from the solution of the hydrodynamic problem of a rotating cylinder within a viscoelastic monolayer atop a viscous subphase. In terms of measured experimental properties, the surface loss modulus, G″, is the out of phase component of the rotational resistance, the surface storage modulus, G′, is in the in-phase component as in conventional 3-D rheology (Figure 6B). The surface viscosity for the Newtonian response we observed over the frequency range of 0.1–10 Hz is ηn = G″/ω, or for the 1 Hz frequency used in Figure 6, ηn = G″/2π. The same experimental dependence of surface viscosity on surface pressure was obtained using a 100 μm probe showing that continuum values of elasticity and viscosity were being measured. The 100 μm probe is more than an order of magnitude larger than the domain sizes we have seen (Figures 5 and 6). Recent results using macroscopic wire rings (10 cm diameter ring, 0.7 mm diameter wires) showed identical trends with surface pressure and good agreement for similar monolayers. The maximum surface viscosity that we could measure with our rheometer was ~100 μPa·s; the surface viscosity of pure DPPC at 40 mN/m was greater than this and was not measured.

**AFM and Langmuir-Blodgett Deposition.** Freshly-cleaved mica substrates (S&J Trading Inc.; Glen Oaks, NY) connected to a computer-controlled dipping mechanism in a commercial circular NIMA L-B trough (Biolin Scientific, Inc., Linthicum Heights, MD) were pulled through the monolayer at 5 mm/min at a constant surface pressure of 20 mN/m. Transfer ratios were determined by recording the interfacial area change of the trough during transfer and comparing this to the surface area of the mica substrate. A transfer ratio of 1 means that these areas are equal; only films with transfer ratios of ~1 were examined. The mica substrates were glued to stainless steel discs and affixed to the magnetic holder of an MMAF-2 AFM (Digital Instruments; Santa Barbara, CA).
with a cantilever tip (Asylum Research, AC160TS; Santa Barbara, CA) designed for tapping mode operation.

**RESULTS AND DISCUSSION**

Two-dimensional GIXD was carried out at ChemMatCARS, Sector 15-ID at the Advanced Photon Source, Argonne National Laboratory on DPPC monolayers with various mole fractions of dihydrocholesterol (Chol). Figure 1 shows the

![Image](https://example.com/image1.png)

Figure 1. Two-dimensional X-ray maps of (A) pure DPPC at Π = 20 mN/m and (B) 6.4 mol % Chol in DPPC at Π = 40 mN/m. Two Bragg reflections are visible, the degenerate [11] reflection at positive q_y and lower q_x, and the [02] at higher q_y and q_x = 0, indicating nearest neighbor tilt.

GIXD intensity maps for (A) pure DPPC at 20 mN/m and (B) 6.4 mol % Chol/DPPC monolayers at 40 mN/m. Figure 2 shows the q_z-integrated intensity profiles (arbitrary units) for DPPC/Chol monolayers at 20, 30, and 40 mN/m for 0 to 7 mol % Chol (each spectra offset by 1000).

We assign the Bragg peak in Figure 1 at lower q_y and positive q_z as due to the degenerate (11) and (11) reflections of distorted hexagonal packing; the second peak at higher q_y = 0 and q_z is due to the nondegenerate (02) reflection. As the (11) reflection is located at q_x > 0, and the (02) reflection is centered at q_y = 0, the alkane chains are tilted in the nearest neighbor (NN) direction. Regardless of composition or surface pressure, all ordered areas in the monolayers had the same distorted hexagonal packing with NN tilt (Figure 3B, inset).

The q_z-integrated intensity profiles (Figure 2) show that the (11) Bragg peak broadens and moves to lower q_y with increasing cholesterol content (dotted lines) for all surface pressures, while the (02) peak also broadens with increasing cholesterol content but remains at constant q_x. From the q_z values in Table 1, the real space lattice dimensions are d_y = 2π/q_z (Table 2 and Figure 3). The tilt angle, θ, for nearest neighbor tilt is given by tan θ = q_x/[q_11 - (q_02/2)]^{1/2}. The coherence length is determined from the full width at half-maximum (fwhm), of each peak after correction for the instrumental resolution, L_y = (0.9-2π)/(fwhm) (Table 2).

![Image](https://example.com/image2.png)

Figure 2. GIXD intensity integrated over the z-direction. For all surface pressures, the [11] peak migrates to higher q_y with increasing cholesterol. The [02] peak remains at the same location for all cholesterol or surface pressures (dotted lines). The basic motif of the DPPC lattice is unchanged by cholesterol, although the tilt is reduced with increasing cholesterol. Cholesterol broadens both peaks consistent with a decrease in the size of the correlated areas in the monolayer.

![Image](https://example.com/image3.png)

Figure 3. (A) d_11 and (B) d_02 as a function of cholesterol and surface pressure. d_11 decreases with increasing surface pressure and cholesterol fraction. d_02 is invariant with both cholesterol and surface pressure. This is consistent with a decrease in tilt, which decreases d_02, while the alkane chain packing normal to the chain axis, hence d_11, remains the same. Inset: Real space lattice corresponding to the distorted hexagonal lattice parameters. One gray and one white circle are the two alkane chains in a rectangular unit cell of dimensions a and b. Translation of the pair of gray and white circles by a and/or b generates the lattice. The spacing between the dotted lines corresponds to the d-spacings in Table 2 and A, B.
Table 1. Reciprocal Space Lattice Parameters from GIXD for All Samples at 20, 30, and 40 mN/m Surface Pressure As a Function of Cholesterol Mole Fraction (Chol %)\(^a\)

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<th>(q_{11}) (Å(^{-1}))</th>
<th>(q_{\parallel}) (Å(^{-1}))</th>
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<th>(q_{\parallel}) (Å(^{-1}))</th>
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<th>Chol %</th>
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<th>(q_{11}) (Å(^{-1}))</th>
<th>(q_{\parallel}) (Å(^{-1}))</th>
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\(^a\)\(q_{02}\) and \(q_{11}\) are the lattice parameters in Fourier space for the two strong reflections from the \(q\) averaged data. \(q_{\parallel}\) is the \(z\) coordinate of the \((11)\) reflection. All \((02)\) reflections are centered at \(q_{\parallel} = 0\). (See Figure 1).

From the values in Table 2, we determine a rectangular two-molecule unit cell of dimensions, \(a = d_{11} = [d_{11}^{-2} - (2d_{02})^{-2}]^{-1/2}\) and \(b = 2d_{02} = d_{01}\) (Figure 3B Inset), with \(a\) decreasing from 5.8 ± 0.1 at 20 mN/m, to 5.4 ± 0.1 Å at 40 mN/m; \(b\) remains constant at 8.6 ± 0.05 Å. These unit cell dimensions are consistent with a hexagonal packing of the alkane chains,\(^34\) which are tilted to accommodate the mismatch in projected area between the DPPC headgroup and the close-packed chains.\(^25,33\) As is the case for other lipids, accommodation of the larger DPPC headgroup area occurs by dilation of the alkane chain area via an increase in the tilt angle. Tilt costs little favorable alkane chain contact energy, as tilt occurs without changing the distances between the alkane chains.\(^25\) Tilt in the NN direction causes \(a\), which is measured in the plane of the monolayer, to increase. However, \(b\) remains constant as this space does not change with tilt if the alkane chains retain their close-packed configuration. Increasing the surface pressure provides a uniform compression of the DPPC headgroup, which results in a decrease in the headgroup-chain incompatibility, and hence the tilt, without altering the alkane chain packing.\(^25\)

The area per alkane chain perpendicular to the alkane chains, 20.5 ± 0.3 Å\(^2\) \(= (ab \cos \theta)/2\), is constant within the experimental error for all surface pressures and also for all cholesterol fractions. Pure Chol has an untilted hexagonal lattice with a \(d\) spacing of 5.7 Å, and an area per molecule of \(~35\) Å\(^2\),\(^35\) much larger than the 20.5 Å\(^2\) area per alkane chain we measure. The invariance of \(d_{02}\) (or \(b\)), and the linear relationship between \(d_{11}\) and \(\theta\) is consistent with the bulk of the Chol not intercalating uniformly into the DPPC lattice as it does at higher mole fractions.\(^35,36\) but separating primarily into a second phase. Figure 5 shows AFM images of DPPC/Chol monolayers transferred to mica substrates at 20 mN/m showing two distinct morphologies. At 0.8 mol % Chol, dark gray, 10–100 nm circular areas are dispersed in extended light gray domains. The dark gray nanodomains localize preferentially at the boundaries of the light gray domains (arrows).\(^10\) Increasing the Chol fraction causes the circular nanodomains to percolate into linear structures (3.7 mol %), although the width of the linear structures remains \(~10–100\) nm. The linear structures eventually break up the light gray domains (5 mol %). AFM force spectroscopy showed that the nanodomains were more compliant and easier to deform than the DPPC domains,\(^10\) suggesting that the nanodomains are disordered and do not contribute to the GIXD signature of the monolayer.

The alkane chains of DPPC prefer to be untilted to maximize the van der Waals contact between the chains,\(^25,34\) but are frustrated by the conflicting cross-sectional area requirements of the phosphocholine headgroup. This mismatch requires the tailgroup lattice to dilate, which responds by the lower energy tilt deformation in order to fill space efficiently. However, Chol has a complementary shape to DPPC, with a relatively small alcohol headgroup and a relatively large sterol ring tailgroup. Hence, this shape complementarity suggests that Chol can relieve the packing frustration of DPPC by making up some of the mismatch between the headgroups and alkane chains. Palmitic acid (PA) and hexadecanol (HD), which also have complementary shapes with DPPC, also lead to a decreased tilt at a given surface pressure,\(^33\) but do not show nanodomains in AFM images. The saturated C\(_{16}\) alkane chain of PA and HD is the same as the C\(_{16}\) alkane chain of DPPC, which allows PA and HD to be incorporated into the DPPC lattice. PA and HD increase the correlation length of the mixed crystal\(^33\) and the surface viscosity.\(^57\)

Although the shape of cholesterol can relieve the frustration between the area of the headgroup and alkane chains of DPPC, the sterol rings cannot efficiently pack into the alkane chain lattice. This leads to a second type of frustration; the decrease in tilt due to the accommodation of the area mismatch results in disrupting the alkane chain lattice, which is likely higher in energy than decreasing the tilt.\(^25\) This second type of frustration may be accommodated by nanodomain phase separation. The combination of the sterol rings of Chol and more disordered DPPC chains gives the nanodomain phase excess tailgroup area, which may relieve part of the packing frustration of the headgroups in the adjacent ordered DPPC domains, albeit at longer range than if the cholesterol intercalated within the DPPC lattice.

The size of domains in typical phase-separated monolayers is governed by a competition between line tension, \(\lambda\), which leads to larger domains, and entropy and electrostatic interactions, which favor smaller domains.\(^23,38,39\) Fluorescence images
chains in degrees with respect to the water surface, tan nm length scales of the nanodomains, so even with sectional area of the chains in the direction perpendicular to the chains, or electrostatic interactions slowing bulk coalescence, molecular coupling between monolayer curvature (i.e. incompatible area proposed 2-D analogs of 3-D microemulsions, in which indication that we are seeing an example of the recently structure were not stable. This stability may also be an nanodomains with the bulk is of order seconds over the adjacent DPPC domains. The time for nanodomains may be the energy of reducing the tilt in the (Figure 6) at the coexistence surface pressure (9–10 mN/m) show that increasing Chol leads to a dramatic decrease in the width of the domains, which means a large decrease in . This decrease may be sufficient to stabilize the nanodomains against coalescence. An additional factor stabilizing the nanodomains may be the energy of reducing the tilt in the adjacent DPPC domains. The time for diffusive mixing of the nanodomains with the bulk is of order seconds over the ~100 nm length scales of the nanodomains so even with electrostatic interactions slowing bulk coalescence, molecular diffusion should eliminate the nanodomains in minutes if this structure were not stable. This stability may also be an indication that we are seeing an example of the recently proposed 2-D analogs of 3-D microemulsions, in which compositional variations within a single phase are due to coupling between monolayer curvature (i.e. incompatible area requirements of headgroups and tails) and composition. The evolution of the structure between discrete circular nanodomains (analogous to spherical micelles in 3-D) to extended linear structures (rod-like micelles) to an interconnected network (bicontinuous microemulsion) suggests this possibility.

Figure 7A shows that the coherence length, in the untitled direction for pure DPPC is about 70 lattice repeats, or >300 Å, more than five times that in the tilted direction, ~60 Å or about 12 lattice repeats (Table 2). For both 20 and 30 mN/m, decreases monotonically with increasing cholesterol fraction to ~20 lattice repeats, but only decreases to ~10 lattice repeats. At 40 mN/m, does not monotonically decrease with cholesterol fraction, the scatter in is much greater than at lower surface pressures, and is always less than expected from the results for the lower surface pressures. This is likely due to a decrease in film stability caused by a combination of leakage under the trough barriers and slow monolayer collapse during the 3-5 h required for GIXD. The AFM images in Figure 5 show that the average DPPC (light gray) domain size decreases from microns to 100–200 nm with cholesterol. Even with the decreasing domain size, the positional ordering given by the coherence lengths is orders of magnitude smaller than the domain size for a given cholesterol fraction. However, the orientational order extends

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<th>d11 (Å)</th>
<th>a (Å)</th>
<th>b (Å)</th>
<th>θ (°)</th>
<th>area (Å²)</th>
<th>chain area (Å²)</th>
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<th>L11 (Å)</th>
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All d-spacings are referenced to a two molecule rectangular unit cell with . The error in is larger than as the (11) reflection is much broader than the (02) reflection (see Figure 2). The tilt angle of the alkane chains in degrees with respect to the water surface, tan θ = q11/q11 − q02/2|q02|2/2 from Table 1 (see inset to Figure 3B). The chain area is the cross sectional area of the chains in the direction perpendicular to the chains, or ab cos θ/2. L02 and L11 are the coherence lengths in the (02) and (11) directions, L02 ≈ (0.9(Å))/bhw(L02).
cholesterol fraction. This linear relationship between molecular tilt angle measured from the monolayer normal, $\theta$, decreases in the same manner with increasing cholesterol fraction as with increasing surface pressure (over this range of mN/m. With some scatter, the tilt decreases with increasing surface pressure, Open symbols -- 20 mN/m surface pressure, Open symbols -- 30 mN/m and Gray symbols -- 40 mN/m. With some scatter, the tilt decreases with increasing cholesterol fraction. This linear relationship between $d_{11}$ and $\theta$ is the same for changes in cholesterol fraction and surface pressure, which suggests that the local alkane packing of DPPC does not change with added cholesterol, but that both surface pressure and cholesterol act to decrease the mismatch between the lipid headgroup and tailgroup area.

Figure 4. Molecular tilt angle measured from the monolayer normal, $\theta$, decreases in the same manner with increasing cholesterol fraction as with increasing surface pressure (over this range of mN/m. With some scatter, the tilt decreases with increasing surface pressure, Open symbols -- 20 mN/m surface pressure, Open symbols -- 30 mN/m and Gray symbols -- 40 mN/m. With some scatter, the tilt decreases with increasing cholesterol fraction. This linear relationship between $d_{11}$ and $\theta$ is the same for changes in cholesterol fraction and surface pressure, which suggests that the local alkane packing of DPPC does not change with added cholesterol, but that both surface pressure and cholesterol act to decrease the mismatch between the lipid headgroup and tailgroup area in the same way.

for tens of microns as shown by the spiral domain textures in Figure 6.10 DPPC/Chol monolayers have nanometer-range positional order and micron-range orientational order,20,22,27,32,40,45 analogous to the classic free volume model for the viscosity:21,22,27,32,40,45

$$P(V_0) = \exp[-BV_0/V_f]$$

thus giving a diffusivity

$$D = g \frac{\eta}{\mu} \exp[-BV_0/V_f] = D_0 \exp[-BV_0/V_f]$$

in which $g$ is a geometric factor.19 The parameter $B$ in eqs 1 and 2 is to take into account overlaps of free volume, and Cohen and Turnbull suggest a range from $1/2 \leq B \leq 1$.19

In three dimensions, the diffusivity can be related to the bulk viscosity, $\eta$, via a generalized Stokes-Einstein relationship, $D = kT/\eta$. For spherical particles, the friction factor, $f$, is given by the Stokes drag on a sphere of diameter $a$: $f = 3\pi a \eta$. This leads to the free-volume model for the viscosity:

$$\ln \eta = \ln \left[ \frac{kT}{3\pi D_0 a} \right] + [BV_0/V_f] = \ln \eta^0 + [BV_0/V_f]$$

In applications of the model, the free volume is the difference between the measured volume per molecule, $V$, and $V_0$: $V_f = V - V_0$. In applications of the model, $V_0$ is a fitting parameter; theoretically, $V_0$ is related to the volume per molecule at the glass transition where the viscosity diverges.

For a 2-dimensional film of constant thickness, $L$:

Figure 5. Tapping-mode AFM images of DPPC/cholesterol monolayers transferred by Langmuir-Blodgett deposition to mica substrates at 20 mN/m. For 0.8 mol % cholesterol, dispersed, circular 10–100 nanodomains appear (darker gray indicates more compliant relative to the lighter gray background phase), which preferentially locate at the boundaries of the light gray domains (arrows). For 3.7 mol % Chol, the circular nanodomains have condensed into linear features begin to break up the light gray domains, but the light gray domains remain continuous. For 5.0 mol % Chol the nanodomains make up a cocontinuous network separating the light gray domains, while decreasing the size of the light gray domains to 100–200 nm. Figure adapted from ref 10.
In analogy to the free volume, \( A_f(\Pi) = A(\Pi) - A_0 \), in which \( A(\Pi) \) is the area per molecule determined at a surface pressure, \( \Pi \), from a surface pressure-area isotherm and \( A_0 \) is taken to be a fitting parameter. In 2-dimensions, the diffusivity and free area can be related to the surface viscosity, \( \eta_s \), via the Saffmann-Delbrück model\(^{29} \) for a cylinder diffusing within a viscous monolayer, surrounded by a viscous subphase, to give an equation analogous to eq 3 in terms of the free area:\(^{10,21,22,40,45} \)

\[
\ln \eta_s(\Pi) = \ln \eta_s^0(\Pi) + \frac{BA_0}{A_f(\Pi)}
\]  

(5)
From fitting DPPC viscosity data, we found that $A_0 \sim 40 Å^2$, which is roughly equal to $ab \cos \theta$, the molecular area of a DPPC molecule in a close-packed, untitled lattice (Table 2).

However, Cohen and Turnbull did not speculate on the molecular origins of $A_0^{19}$ as they were modeling an unstructured liquid. However, for semicrystalline monolayers, we postulate that the free area is proportional to the number of defects in the lattice, which is inversely proportional to the number of correlated molecules at that composition and surface pressure:

$$A_f(x, \Pi) = \frac{\alpha}{(L_{02}L_{11})/ab}$$  \hspace{1cm} \text{(6)}$$

Lattice defects, such as dislocations, vacancies, and grain boundaries decorrelate the lattice, which creates free area and pathways for diffusion. We define $\alpha$ as the free area (or number of defects times the area per defect$^{20}$) per number of correlated molecules; we take $\alpha$ to be constant, independent of concentration. Combining eqs 5 and 6:

$$\ln \eta_f(x, \Pi) = \ln \eta_0^s(x, \Pi) + \beta \frac{(L_{02}L_{11})}{ab}$$  \hspace{1cm} \text{(7)}$$

in which $\beta = B A_0^{\alpha}/\alpha$. Linear regression to eq 7 (Figure 7C) shows that for 20, 30, and 40 mN/m, the slopes, $\beta = 0.0134 \pm 0.0007, 0.0131 \pm 0.002$ and 0.0196 $\pm 0.007$ are the same within the experimental error, $\eta_0^s(\Pi = 20 \text{ mN/m}) = 0.09 \pm 0.02, \eta_0^s(\Pi = 30 \text{ mN/m}) = 0.37 \pm 0.2$, and $\eta_0^s(\Pi = 40 \text{ mN/m}) = 0.6 \pm 0.5 \mu \text{P-m/s}$, although the physical significance of the surface pressure variation of $\eta_0^s$ is not given by our model. The Pearson correlation coefficient, $r_f$ for the three lines are 0.99, 0.95 and 0.81, respectively, giving a statistically significant fit of eq 7 to the data with $p < 0.001, 0.001$, and 0.05, respectively. To better compare the data at different surface pressures, the surface viscosity and correlated area are normalized relative to a reference composition, $x_{ref}$ at the same $\Pi$:

$$\ln \left[\frac{\eta_f(x, \Pi)}{\eta_f(x_{ref}, \Pi)}\right] = -\beta \left[\frac{(L_{02}L_{11})}{ab}\right]_{ref} - \left[\frac{(L_{02}L_{11})}{ab}\right]$$  \hspace{1cm} \text{(8)}$$

$\eta_f(x_{ref}, \Pi)$ is the surface viscosity and $((L_{02}L_{11})/ab)_{ref}$ is the number of molecules in the coherence area at $x_{ref}$ (taken to be pure DPPC for $\Pi = 20$ and 30 mN/m and 0.4% Chol for $\Pi = 40$ mN/m). Figure 7D shows that eq 8 eliminates the surface pressure dependence. Linear regression to eq 8 gives $\beta = 0.0133 \pm 0.007$, which is the same as the fits to eq 7. The Pearson correlation coefficient is 0.91, giving $p < 0.001$, showing that eq 8 gives a statistically significant representation of the surface viscosity over the almost three orders of magnitude change in surface viscosity (Figure 7B). For $A_0 \sim 40 Å^2$, $\alpha \sim 1500 Å^2$ (for $B = 1/2$) $\sim 3000 Å^2$ (for $B = 1$). For pure DPPC, $(L_{02}L_{11})/ab \sim 450$, giving $A_0 \sim 3-6 Å^2$ from eq 6, which is consistent with the variation in area per molecule, $A(x, \Pi) = A_0 + A(x, \Pi)$, measured from DPPC isotherms.$^{10,25}$

Adding cholesterol decreases $(L_{02}L_{11})/ab$ to $\sim 100$, thereby increasing the free area per molecule to $15-30 Å^2$, resulting in the dramatic decrease in surface viscosity. These values of $A_f$ are in agreement with the assumptions behind the Cohen and Turnbull free volume theory, which postulates that $V_t \ll V_0$, hence $A_t \ll A_0$.

## CONCLUSIONS

In summary, GIXD shows that increasing the Chol fraction at constant surface pressure, or increasing the surface pressure at constant Chol fraction, decreases the tilt of the DPPC lattice, while leaving the alkane chains close-packed with an invariant area per molecule normal to the alkane chain direction. This confirms AFM images that show cholesterol does not intercalate homogeneously into the DPPC lattice but is expelled to disordered nanodomains that break up the DPPC domains. The nanodomains evolve from isolated, circular 10–100 nm diameter domains to $\sim 10–50$ nm wide linear nanodomain aggregates to an interconnected network structure with increasing cholesterol. However, the monolayer is homogeneous at micron-scale optical images; macroscopic phase separation does not occur as at lower surface pressure (Figure 6). Hence, the DPPC/cholesterol monolayer is better described as a nанostructured, single-phase monolayer, rather than a mixture of two distinct phases, similar to recently proposed two-dimensional microemulsions.$^{6,41}$ As such, this system is analogous to 3-D nanostructured surfactant-oil-water bicontinuous microemulsions, or copolymer systems that separate into nanometer-scale regions more enriched in one or the other monomer, which are considered single phase.

Adding cholesterol does not change the local hexagonal DPPC chain packing, but does reduce the molecular tilt, suggesting that cholesterol relieves some of the packing frustration between the DPPC headgroup and tailgroups. Most important to the monolayer dynamics, the extent of order, or the number of correlated molecules, decreases with increasing cholesterol fraction at all surface pressures, suggesting an increased number of lattice defects that create free area for visco-diffusive transport. At all cholesterol fractions, the positional order is much shorter ranged than the orientational order, consistent with an overall hexatic organization.$^{20}$ We show that a simple model that proposes that the “free” area available for visco-diffusive transport$^{19}$ is inversely proportional to the number of correlated molecules, which collapses the surface viscosity data for all surface pressures and cholesterol fractions onto a single universal curve. Molecular defects and the associated decrease in domain size caused by the incompatibility of cholesterol packing into the alkane chain lattice enhance visco-diffusive transport in monolayers. Our model and data show that the extent of molecular correlations is an excellent predictor of the effects of the cholesterol on the surface viscosity of model lung surfactant monolayers. Future work should show if this relationship is universal to other phospholipid monolayers.

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**Notes**

The authors declare no competing financial interest.

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