Effect of dibucaine on phase behavior of ternary liposome

Kazunari Yoshida, Akito Takashima, and Izumi Nishio

College of Science and Engineering, Aoyama Gakuin University,
Sagamihara, Kanagawa, 252-5258, Japan

(Dated: December 1, 2014)

Abstract

Despite many studies, detailed physical mechanism of local anesthetic function is still unclear. In order to clarify the mechanism, the effects of anesthetics on ion channels and the lipids surrounding the ion channels should be evaluated. Thus, we investigated the effect of Dibucaine hydrochloride (DC-HCl), one of the local anesthetics, on phase behavior of ternary liposome composed of dioleoylphosphatidylcholine (DOPC), dipalmitoylphosphatidylcholine (DPPC), and cholesterol (Chol). The large DOPC/DPPC/Chol liposome, that is directly observable by optical microscope, is commonly known to be laterally separated into liquid-ordered (Lo) phase (raft-like domain) and liquid-disordered (Ld) phase under certain conditions and is useful for study of lipid-raft-like domains as a simple model system. In order to confirm the effect of DC-HCl on a miscibility transition temperature, $T_c$, of the ternary liposome, we observed the liposomes with three concentrations, 0, 0.05, and 0.2 mM, of DC-HCl at various temperatures. In addition, we calculated the angle-averaged two-dimensional autocorrelation (2D-AC) functions in order to quantify the phase behavior. The results of these observations and calculations revealed that the DC-HCl molecules induce the reduction of $T_c$ of the ternary liposome. Furthermore, we calculated the circularity of Lo domain in order to confirm the change in the line tension of the Lo/Ld phase boundary and revealed that the insertion of the DC molecules induces the reduction of line tension. In terms of the critical phenomena, we conclude that the insertion of the DC molecules induces the reduction of the $T_c$ of the ternary liposome due to reduction of line tension. This suggests that the DC molecules may disturb function of ion channels via affecting the lipid bilayers which surround ion channels.
INTRODUCTION

Because it is important to understand the physical mechanism of functional expression of anesthetics in the pharmacology, the effect of anesthetic molecules on biological membrane containing ion channels has been studied for many years. For instance, effect of chloroform and other general anesthetics on sodium current in a squid giant axon and interactions of chloroform and/or benzyl alcohol with lipid bilayer have been studied by using experimental and/or simulation methods[1–5].

Recently, local anesthetics have especially been paid attention and studied. A lot of studies have reported that local anesthetics affect the sodium current across the cell membrane[6–12]. In addition, interaction between anesthetics and lipid bilayer which surrounds the ion channels has also been studied because it is suggested that the perturbation of the lipids due to invasion of the anesthetics disturbs the function of ion channels[13–17]. Furthermore, the effect on lipid-raft domains, which are mainly composed of saturated lipids, cholesterol and membrane proteins such as ion channels, and are dispersed in the cell membranes (believed to be a kind of phase separation)[18] should be investigated in order to clarify the detailed mechanism of local anesthetic function because the lipid rafts are believed to play important roles in the various cell functions[18]. Therefore, the phase-separated liposome with the raft-like domains has been used as a simple model system of cell membrane for studies of interactions between local anesthetics and lipid bilayers[19, 20]. However, the detailed physicochemical mechanism of functional expression of local anesthesia is still unclear.

Large phase-separated liposome, that is observable by optical microscope, is useful for the direct investigations of the raft-like domains. The multi-component liposome is commonly known to be separated into two or three phases due to difference of thermodynamical properties of their lipids, and the phase-separated domains of large liposome are observable by fluorescence microscopy with a dye which is only incorporated into a certain phase[21–26]. Indeed, large phase-separated liposome has been used for many studies of the lipid membranes, and the physical properties of the lipid membrane have been revealed[21, 23–29].

In the similar cases to the present study, the large liposome and cell-derived giant plasma membrane vesicle (GPMV) which is similar to the multi-component liposome are also used as a model system. For example, it has been reported that vitamin E, Triton-X 100, and benzyl alcohol affect the phase morphology of large multi-component liposomes[30]. Gray et
al. have also reported that the general anesthetics including alcohols reduce the miscibility transition temperature, $T_c$, of GPMVs. However, a direct observation of the influence of local anesthetics on the phase behavior of large liposome has not been reported.

In the present study, we investigated the influence of dibucaine hydrochloride (DC·HCl), one of the commonly used local anesthetics, on phase behavior of liposome composed of dioleoylphosphatidylcholine (DOPC), dipalmitoylphosphatidylcholine (DPPC), and cholesterol (Chol). The DOPC/DPPC/Chol liposome is one of the famous systems and is known to be laterally separated into liquid-ordered (Lo) and liquid-disordered (Ld) phases under certain conditions. The Lo phase, mainly composed of DPPC and Chol, represents the raft-like domain, and the Ld phase is DOPC-rich region. We calculated angle-averaged two-dimensional autocorrelation (2D-AC) function of images of liposomes with three concentrations of DC·HCl at various temperatures in order to estimate the change in $T_c$ due to the effect of DC. The results indicate that the DC molecules reduce the $T_c$ of ternary liposome.

The 2D-AC and/or angle-averaged 2D-AC functions have been widely used in the various fields such as biology and physics. In addition, the angle-averaged 2D-AC analysis was performed in order to clarify the correlation length of GPMV surface in the similar study to this article. However, to our knowledge, this is the first report of quantitatively demonstrating the change in $T_c$ of ternary liposome due to addition of local anesthetics by calculating the angle-averaged 2D-AC function of membrane domain patterns.

MATERIALS AND METHODS

Materials

Dibucaine hydrochloride (DC·HCl), DOPC (chain melting temperature, $T_m \approx -18^\circ C$), DPPC ($T_m \approx 41^\circ C$), Chol and chloroform were purchased from Wako Pure Chemical Industries, Ltd. (Japan). The chemical structure of DC is described in Figure 1. Methanol was purchased from Showa Chemical Industry Co., Ltd. (Japan). Rhodamine B 1,2-dihexadecanoyl-sn-glycero-3-phosphoethanolamine (rhodamine DHPE, $\lambda_{ex} = 560$ nm, $\lambda_{em} = 580$ nm), fluorescent phospholipid, was obtained from Invitrogen (U.S.A.). Ultra-pure water was obtained using a WT101UV AUTOPURE (Yamato Scientific Co., Ltd., Japan).
Preparation of Ternary Liposome

We prepared liposomes using the natural swelling methods\cite{38,39}. The lipids of DOPC/DPPC/Chol = 50/25/25 (mol%) with 0.5 mol% of rhodamine DHPE were dissolved in mixture solvent of chloroform and methanol with a volume ratio of 2:1. Then, we removed the organic solvents through an air flow and placing the sample in an aspirated desiccator for more than 8 hours in order to make a dry lipid film. Finally, liposomes formed through hydration of the lipid film with the ultra-pure water at 37 °C for more than 24 hours. The lipid concentration was 0.2 mM.

Observation

40 µL of liposome suspension and 40 µL of DC·HCl aqueous solution were gently mixed, and the solution was incubated at room temperature (RT, 21.7 ± 0.4 °C) for more than 10 minutes in order to wait for sufficient dispersion of DC·HCl molecules into the lipid membrane. In this study, we used 0, 0.1, and 0.4 mM of DC·HCl aqueous solutions. The final concentration of lipids was 0.1 mM with 0, 0.05, or 0.2 mM of DC·HCl. Then, 5 µL of the mixed solution was placed between two cover glasses and was sealed with vacuum grease. The sample was placed on a copper plate, and the temperature of the plate was controlled using a peltier device (deviation < 0.1 °C) with monitoring the plate temperature using a thermistor probe placed in the copper plate (under the sample). We put a thin layer of thermal grease between the sample cell and the copper plate in order to keep the high thermal conduction from the plate to the sample. The sample placed on the plate was incubated for more than 10 minutes in order to archive the thermal equilibrium state before observation and was observed using a fluorescence microscope BX40 (Olympus, Japan) through a objective lens of 40× at various temperature. We recorded the microscopic images using a digital camera DP73 (Olympus, Japan). Excitation-light irradiation (λ\text{max} = 546 nm) was applied using a Hg lamp through a WIG filter set (Olympus, Japan).

Analysis of microscopic images

We calculated the angle-averaged 2D-AC using a macro program of the ImageJ\cite{40}. First, fluorescence microscopic images of liposome surface were cut into 64 × 64 pixels and were
converted into gray scale (16 bit). Then, we calculated two-dimensional fast Fourier transform (2D-FFT) using brightnesses of each pixel of images and inverse transform of their power spectrum in order to derive 2D-AC function. Further, the 2D-AC function was averaged for all angles. Finally, we calculated the FFT of \( G(r)/G(0) \) curves, where \( G(r) \) is a angle-averaged 2D-AC as a function of radial position, \( r \), in the real space. Typical results of the analysis of one- and two-phase images are described in the Supplemental Material (SM).

**Calculation of Lo-domain circularity**

The circularity was also calculated using ImageJ\[40\]. We transformed the microscopic images to binary images in order to obtain the phase boundary of Lo/Ld. The circularity, dimensionless parameter, of the Lo domains was defined as

\[
\text{(circularity)} = \frac{4\pi A}{l^2},
\]

where the \( \pi \) is the ratio of the circumference of a circle to its diameter, \( A \) is a area of Lo domain, and \( l \) is a perimeter of Lo domain. The circularity of 1 corresponds to perfect circle. We calculated the circularity in order to estimate the change in line tension of phase boundary in the case of \( T = 20 \) °C.

**RESULTS AND DISCUSSION**

We observed the DOPC/DPPC/Chol liposomes with 0, 0.05, and 0.2 mM of DC·HCl at \( T = 20, 25, 30, 35, \) and 40 °C in order to confirm the influence of DC·HCl on phase behavior of the ternary liposomes. Figure 2 shows the typical microscopic images at each condition. Since rhodamine DHPE is mainly localized in Ld phase of the phase-separated systems\[25, 41\], the dark region corresponds to DPPC- and Chol-rich domain (Lo, raft-like domain) while the bright region corresponds to DOPC-rich domain (Ld). Image 1 described in the Figure 2 shows the phase-separated liposome with scattered Lo domains. The number of phase-separated liposome was reduced with increasing temperature (images 1-5). In the case of 0 mM and at 35 °C, we observed many one-phase liposomes. There is no phase-separated liposome at 40 °C, corresponding to image 5. Next, we also observed the ternary
liposomes at various temperatures in the case of 0.05 mM DC·HCl. We observed the phase separation in the most of liposomes with 0.05 mM DC·HCl at 20 °C, corresponding to image 6. There are several phase-separated liposomes with smaller Lo domains as shown in the image 7 and a few liposomes with large domains as shown in Figure S2 described in SM at 25 °C. We observed a few phase-separated liposomes with smaller Lo domains at 30 °C (image 8), and most of liposomes dose not have the Lo domains at 35 °C (image 9). Further, in the case of 0.2 mM, the number of phase-separated liposomes was also reduced with increasing temperature. At 20 °C, phase-separated liposomes are majority, however, there are several liposomes which have unclear phase boundaries. At 25 °C and above, most of liposomes have one phase without raft-like domains, corresponding to images 11 and 12. We especially focus on the images at 25 °C with the three DC·HCl concentrations, corresponding to images 2, 7, and 11. The Lo domains in the liposome surfaces became smaller with increasing DC·HCl concentration. These results indicate that DC molecule have an ability to change the phase behavior of DOPC/DPPC/Chol liposomes. It is considered that the effect of HCl is ignorable because the lipids used in this study is electrically neutral (not charged lipids)[25, 26].

In order to quantify the phase behavior, we calculated the 2D-AC functions of domain pattern in the ternary liposomes. Figures 3(a, c, and e) show mean values of the normalized angle-averaged 2D-AC, \( G(r)/G(0) \), as a function of radial position, \( r \), at 0, 0.05, and 0.2 mM, respectively, and Figures 3(b, d, and f) show the enlarged view of the short distance region of the angle-averaged 2D-AC functions, corresponding to Figures 3(a, c, and e), respectively. The numbers of the observed liposomes are not less than 10 at each condition. Value of the short distance region (especially \( r \approx 0.25 \mu m \)) of the \( G(r)/G(0) \) decreases with increasing temperature. It is considered that the correlation of the short distance region dependents on the ratio of the phase-separated liposomes. The correlation of the phase-separated liposome tends to have higher value due to Lo domains than that of the one-phase liposome as shown in Figure S1. The results indicate that the ratio of the phase-separated liposomes is reduced with increasing temperature. In the case of the 0.2 mM, dispersion of the values at \( r \approx 0.25 \mu m \) is smaller than that of two other concentrations because the differences of ratio of phase-separated liposomes between three temperatures seem to be smaller than that of 0 and 0.05 mM.

Since we need further informations in order to quantitatively confirm the effect of DC·HCl
molecules on miscibility transition temperature, $T_c$, FFT of the $G(r)/G(0)$ curves of individual liposomes was calculated. Figures 4(a, b, and c) show the Fourier spectrum of the $G(r)/G(0)$ curves at 0, 0.05, and 0.2 mM, respectively ($n \geq 10$). Fourier amplitude of phase-separated liposome at wavenumber $k \approx 0.3$ and 0.6 $\mu$m$^{-1}$ tends to have larger value due to Lo domains than that of one-phase one as shown in Figure S1, where the wavenumber $k \approx 0.3$ and 0.6 $\mu$m$^{-1}$ correspond to wavelength $\lambda \approx 3.3$ and 1.7 $\mu$m, respectively. Therefore, the values of the Fourier amplitudes depend on the rate of phase-separated liposomes. The amplitudes at $k \approx 0.3$ and 0.6 $\mu$m$^{-1}$ are reduced with increasing temperature at all concentrations. In addition, largest distance point of Fourier amplitudes corresponds from 30 to 35 °C at 0 mM (Figure 4(a)), while the points correspond from 20 to 25 °C at 0.05 and 0.2 mM (Figures 4(b and c)). This indicates that the DC·HCl molecules reduce the $T_c$ because it is considered that such largest distance point roughly corresponds to $T_c$.

Furthermore, relation between Fourier amplitude and temperature was shown in Figure 5. The result indicates that the $T_c$ is reduced with increasing DC·HCl concentration. The $T_c$ downwardly shifts more than 5 °C from 0 mM to 0.05 and 0.2 mM. The difference of Fourier amplitude between 0.05 and 0.2 mM at 25 °C is larger at $k \approx 0.3$ $\mu$m$^{-1}$ than at $k \approx 0.6$ $\mu$m$^{-1}$. This indicates that the amount of the liposomes with Lo domains corresponded to $k \approx 0.3$ $\mu$m$^{-1}$ is larger at 0.05 mM than at 0.2 mM, and that the amount of the liposomes with Lo domains corresponded to $k \approx 0.6$ $\mu$m$^{-1}$ at 0.05 mM is similar to at 0.2 mM. In other words, the ratio of liposomes having the large domains is higher at 0.05 mM than at 0.2 mM at 25 °C. This suggests that the difference of the DC·HCl concentration influences the phase behavior. To our knowledge, this is the first report of quantitatively showing the change in $T_c$ of DOPC/DPPC/Chol liposome due to effect of local anesthetic molecule by 2D-AC analysis.

In this study, we demonstrated the change in $T_c$ of DOPC/DPPC/Chol liposome due to effect of DC·HCl molecules. We discuss the cause of change in $T_c$ by calculating a circularity of the Lo domains. Figure 6 shows the circularity, $4\pi A/l^2$, at 20 °C. The circularity is reduced with increasing the DC·HCl concentration. This suggests that the line tension of the Lo/Ld phase boundary is reduced with the DC·HCl concentration because the domain circularity is related to line tension of phase boundary. In other words, the DC molecule induces the reduction of line tension of phase boundary. It was previously shown that the DC molecules has an ability to invade the lipid bilayers, which led us to
consider this possibility in the case of DOPC/DPPC/Chol systems, and the effect of HCl is ignorable as described above. In addition, line-tension change induced by incorporation of other molecules was reported by several groups\cite{44, 45}. Therefore, it is considered that the insertion of the DC molecules into bilayers induces the reduction of the line tension.

Previously, line tension of the phase boundary was expressed as

$$\sigma \propto (T_c - T)^\nu,$$

where the $\sigma$ is a line tension of the Lo/Ld phase boundary, and the $\nu$ is a critical exponent\cite{21, 46}. In terms of this equation, the miscibility transition temperature, $T_c$, is reduced with the decrease in the line tension, $\sigma$, when the temperature of the system, $T$, is constant. Therefore, it is considered that the insertion of the DC molecules into bilayers induces the reduction of the $T_c$ (Figure 5) through the reduction of the line tension.

A similar studies to the present work suggested that the general anesthetics disturb the phase separation of liposome and GPMV (reduction of $T_c$) by using the neutron diffraction and/or X-ray diffraction methods\cite{47, 48} and the direct observation\cite{31}. It is believed that the behavior in these previous studies is also induced by insertion of the anesthetic molecules to bilayers. Actually, Hamada et al. have reported that photoisomerization of the azobenzene derivative reversibly switches the phase pattern due to change in lateral line tension\cite{49}.

**CONCLUSION**

In summary, we have investigated the influence of DC·HCl molecules, one of the local anesthetics, on phase behavior of DOPC/DPPC/Chol = 50/25/25 (mol%) liposome. In order to confirm the effect of DC·HCl molecules, the ternary liposomes with 0, 0.05, and 0.2 mM of DC·HCl have been observed at various temperatures. As a result, we have clarified the DC molecules has an ability to change the miscibility transition temperature, $T_c$ of the ternary liposome by calculating the angle-averaged 2D-AC function. Furthermore, we calculated a circularity of the Lo domains at 20 °C in order to estimate the change in line tension of the Lo/Ld phase boundary and revealed that insertion of the DC molecules into bilayers induces the reduction of the $T_c$ via decrease in line tension. This suggests that the DC molecules disturb the function of the ion channels (anesthetic function of DC) through
affecting the lipid bilayers which surround the ion channels. Since these physicochemical findings hint at the mechanism of the anesthetic function, this study may play an important role in the pharmacology.

ACKNOWLEDGEMENTS

The authors thank Dr. Yasuhiro Fujii (Ritsumeikan Univ.), Prof. Tsutomu Hamada (JAIST), Dr. Rina Kagawa (M.D., Univ. Tokyo), and Prof. Miho Yanagisawa (Tokyo Univ. Agr. Tech.) for their significant advises.

* yoshida.k09@mishio-lab.net
† izumi@phys.aoyama.ac.jp

[1] D. A. Haydon and B. W. Urban, J. Physiol. 341, 429 (1983).
[2] L. Ebihara, J. E. Hall, R. C. MacDonald, T. J. McIntosh, and S. A. Simon, Biophys. J. 28, 185 (1979).
[3] J. Reyes and R. Latorre, Biophys. J. 28, 259 (1979).
[4] R. Reigada, J. Phys. Chem. B 115, 2527 (2011).
[5] R. Reigada, PLoS ONE 8, e52631 (2013).
[6] Y. Muroi and B. Chanda, J. Gen. Physiol. 133, 1 (2009), http://jgp.rupress.org/content/133/1/1.full.pdf+html.
[7] D. A. Hanck, E. Nikitina, M. M. McNulty, H. A. Fozzard, G. M. Lipkind, and M. F. Sheets, Circ. Res. 105, 492 (2009), http://circres.ahajournals.org/content/105/5/492.full.pdf+html.
[8] T. Yamagishi, W. Xiong, A. Kondratiev, P. Vélez, A. Méndez-Fitzwilliam, J. R. Balser, E. Marbán, and G. F. Tomaselli, Mol. Pharm. 76, 861 (2009), http://molpharm.aspetjournals.org/content/76/4/861.full.pdf+html.
[9] M. Arcisio-Miranda, Y. Muroi, S. Chowdhury, and B. Chanda, J. Gen. Physiol. 136, 541 (2010), http://jgp.rupress.org/content/136/5/541.full.pdf+html.
[10] M. Sheets, T. Chen, and D. A. Hanck, Eur. J. Physiol. 461, 91 (2011).
[11] S. Lee, S. J. Goodchild, and C. A. Ahern, J. Gen. Physiol. 139, 507 (2012), http://jgp.rupress.org/content/139/6/507.full.pdf+html.
[12] J. S. Bant, T. K. Aman, and I. M. Raman, J. Neuro. 33, 4976 (2013), http://www.jneurosci.org/content/33/11/4976.full.pdf+

[13] T. Hata, H. Matsuki, and S. Kaneshina, Coll. Surf. B: Biointer. 18, 41 (2000).

[14] T. Hata, T. Sakamoto, H. Matsuki, and S. Kaneshina, Coll. Surf. B: Biointer. 22, 77 (2001).

[15] H. Matsuki, T. Hata, M. Yamanaka, and S. Kaneshina, Coll. Surf. B: Biointer. 22, 69 (2001).

[16] K. Takeda, Y. Sano, S. Ichikawa, Y. Hirata, H. Matsuki, and S. Kaneshina, J. Oleo Sci. 58, 369 (2009).

[17] H. Tsuchiya, T. Ueno, M. Mizogami, and K. Takakura, Chem. Biol. Interact. 183, 19 (2010).

[18] K. Simons and E. Ikonen, Nature 387, 569 (1997).

[19] H. Tsuchiya, T. Ueno, M. Mizogami, and K. Takakura, J. Anesth. 24, 639 (2010).

[20] C. Bandeiras, A. P. Serro, K. Luzyanin, A. Fernandes, and B. Saramago, Eur. J. Pharm. Sci. 48, 153 (2013).

[21] T. Baumgart, S. T. Hess, and W. W. Webb, Nature 425, 821 (2003).

[22] S. L. Veatch and S. L. Keller, Biophys. J. 85, 3074 (2003).

[23] A. Diguet, M. Yanagisawa, Y.-J. Liu, E. Brun, S. Abadie, S. Rudiuk, and D. Baigl, J. Am. Chem. Soc. 134, 4898 (2012), http://pubs.acs.org/doi/pdf/10.1021/ja211664f.

[24] M. Morita, T. Hamada, Y. Tendo, T. Hata, M. C. Vestergaard, and M. Takagi, Soft Matter 8, 2816 (2012).

[25] N. Shimokawa, M. Hishida, H. Seto, and K. Yoshikawa, Chem. Phys. Lett. 496, 59 (2010).

[26] H. Himeno, N. Shimokawa, S. Komura, D. Andelman, T. Hamada, and M. Takagi, Soft Matter 10, 7959 (2014).

[27] P. Cicuta, S. L. Keller, and S. L. Veatch, J. Phys. Chem. B 111, 3328 (2007), http://dx.doi.org/10.1021/jp0702088.

[28] Y. Sakuma, T. Taniguchi, T. Kawakatsu, and M. Imai, Biophys. J. 105, 2074 (2013).

[29] D. Chen and M. M. Santore, Langmuir 30, 9484 (2014), http://dx.doi.org/10.1021/la502089t.

[30] H. S. Muddana, H. H. Chiang, and P. J. Butler, Biophys. J. 102, 489 (2012).

[31] E. Gray, J. Karslake, B. Machta, and S. Veatch, Biophys. J. 105, 2751 (2013).

[32] S. L. Veatch and S. L. Keller, Biophys. J. 85, 3074 (2003).

[33] J. Hwang, L. A. Gheber, L. Margolis, and M. Edidin, Biophys J. 74, 2184 (1998).

[34] S. Facsko, T. Dekorsy, C. Koerdt, C. Trappe, H. Kurz, A. Vogt, and H. L. Hartnagel, Science 285, 1551 (1999), http://www.sciencemag.org/content/285/5433/1551.full.pdf
[35] V. Geringer, M. Liebmann, T. Echtermeyer, S. Runte, M. Schmidt, R. Rückamp, M. C. Lemme, and M. Morgenstern, Phys. Rev. Lett. 102, 076102 (2009).

[36] S. L. Veatch, P. Cicuta, P. Sengupta, A. Honerkamp-Smith, D. Holowka, and B. Baird, ACS Chem. Biol. 3, 287 (2008) [http://dx.doi.org/10.1021/cb800012x]

[37] R. Koynova and M. Caffrey, Biochim. Biophys. Acta - Biomembr. 1376, 91 (1998).

[38] D. D. Lasic, Biochem. J. 256, 1 (1988) [http://www.biochemj.org/bj/256/0001/2560001.pdf]

[39] K. Yoshida, Y. Fujii, and I. Nishio, J. Phys. Chem. B 118, 4115 (2014) [http://dx.doi.org/10.1021/jp412710f]

[40] C. A. Schneider, W. S. Rasband, and K. W. Eliceiri, Nat. Methods 9, 671 (2012).

[41] S. Chiantia, N. Kahya, J. Ries, and P. Schwille, Biophys. J. 90, 4500 (2006).

[42] A. J. García-Sáez, S. Chiantia, and P. Schwille, J. Biol. Chem. 282, 33537 (2007).

[43] N. Weizenmann, D. Huster, and H. A. Scheidt, Biochim. Biophys. Acta - Biomembr. 1818, 3010 (2012).

[44] S. Akimov, E. Hlaponin, P. Bashkirov, I. Boldyrev, I. Mikhalyov, W. Telford, and I. Molotkovskaya, Biochemistry (Moscow) Supplement Series A: Membrane and Cell Biology 3, 216 (2009) [http://dx.doi.org/10.1021/la204225a]

[45] J. B. Hutchison, R. M. Weis, and A. D. Dinsmore, Langmuir 28, 5176 (2012).

[46] A. R. Honerkamp-Smith, P. Cicuta, M. D. Collins, S. L. Veatch, M. den Nijs, M. Schick, and S. L. Keller, Biophys. J. 95, 236 (2008).

[47] M. Weinrich, H. Nanda, D. L. Worcester, C. F. Majkrzak, B. B. Maranville, and S. M. Bezrukov, Langmuir 28, 4723 (2012) [http://pubs.acs.org/doi/pdf/10.1021/la204317k]

[48] M. Weinrich and D. L. Worcester, J. Phys. Chem. B 117, 16141 (2013) [http://pubs.acs.org/doi/pdf/10.1021/jp411261g]

[49] T. Hamada, R. Sugimoto, T. Nagasaki, and M. Takagi, Soft Matter 7, 220 (2011).
FIG. 1. Chemical structure of DC.

FIG. 2. Typical microscopic images of DOPC/DPPC/Chol liposomes with 0, 0.05, and 0.2 mM of DC·HCl at 20, 25, 30, 35, and 40 °C. Scale bars represent 20 µm.

FIG. 2. Typical microscopic images of DOPC/DPPC/Chol liposomes with 0, 0.05, and 0.2 mM of DC·HCl at 20, 25, 30, 35, and 40 °C. Scale bars represent 20 µm.
FIG. 3. Mean values of the angle-averaged 2D-AC functions ($n \geq 10$ at each condition, error bars not shown). (a, c, and e) The angle-averaged 2D-AC functions of ternary liposomes with 0, 0.05, and 0.2 mM of DC·HCl, respectively. (b, d, and f) Enlarged view of the short distance region of the angle-averaged 2D-AC function, corresponding to (a, c, and e), respectively.
FIG. 4. (a, b, and c) Mean values of the Fourier amplitudes of the $G(r)/G(0)$ curves as a function of wavenumber, $k$, at 0, 0.05, and 0.2 mM, respectively ($n \geq 10$ at each condition, error bars not shown).

FIG. 5. (a and b) The Fourier amplitude versus temperature at $k \approx 0.3$ and 0.6 $\mu m^{-1}$, respectively. Error bars represent standard errors ($n \geq 10$ at each condition).
FIG. 6. Circularity of the Lo domains versus DC-HCl concentration at 20 °C (< $T_c$). Error bars represent standard errors ($n \geq 27$ Lo domains at each concentration).