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Effect of dibucaine hydrochloride on raft-like lipid domains in model membrane systems†

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To clarify the biophysical and/or physicochemical mechanism of anaesthesia, we investigated the influence of dibucaine hydrochloride (DC-HCl), a local anaesthetic, on raft-like domains in ternary liposomes composed of dioleoylphosphatidylcholine (DOPC), dipalmitoylphosphatidylcholine (DPPC) and cholesterol (Chol). The large DOPC/DPPC/Chol liposome is directly observable by optical microscopy and is known to be laterally separated into a liquid-ordered (Lo) phase (raft-like domain) and liquid-disordered (Ld) phase under certain conditions. Hence, it is useful for the study of lipid raft domains as a simple model system of cell membranes. We observed ternary liposomes with three concentrations (0, 0.05 and 0.2 mM) of DC-HCl at various temperatures and calculated the angle-averaged two-dimensional autocorrelation functions to confirm the change in miscibility transition temperature (T_c) of ternary liposomes. Furthermore, we calculated the circularity of the Lo domain to confirm the change in line tension of the Lo/Ld phase boundary. The results indicated that the insertion of DC molecules into lipid bilayers induces reduction in T_c of the ternary liposome, accompanied by reduction in line tension of the phase boundary. This suggests that the DC-HCl molecules may disturb ion channel functions by affecting the lipid bilayers surrounding the ion channels.

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1 Introduction

Understanding the biophysical and/or physicochemical mechanisms of the functional expression of anaesthetics in pharmacology is important. As such, the effect of anaesthetic molecules on cell membranes containing ion channels has been extensively studied. For instance, the effect of chloroform and other general anaesthetics on the sodium current in a squid giant axon and interactions of chloroform and/or benzyl alcohol with the lipid bilayer have been studied through experimental and/or simulation methods.^{1–5}

Recently, local anaesthetics have attracted significant attention. Numerous studies have been reported involving the effect of local anaesthetics on the sodium current across the cell membrane.^{6–12} In addition, interactions between anaesthetics and the lipid bilayer, which surrounds the ion channels, have also been studied as it has been suggested that the perturbation of the lipids due to invasion of anaesthetics disturbs the ion channel function.^{13–17}

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† Electronic supplementary information (ESI) available: Typical results of the 2D-AC analyses of liposomes with three types of phase patterns, a typical microscopic image of an ambiguous phase pattern, and result of Fourier analysis. See DOI: 10.1039/c5md00108k

Furthermore, it is still unknown how local anaesthetics affect lipid raft domains. Lipid raft domains are mainly composed of saturated lipids, cholesterol and membrane proteins, such as ion channels, and are dispersed in cell membranes, which are thought to occur *via* phase separation.^{18,19} Since lipid rafts play an important role in various cell functions,¹⁸ the effect of local anaesthetics on raft domains should be investigated to clarify their detailed mechanism of action. Phase-separated liposomes with raft-like domains have been used as simple model systems of the cell membrane to study the interactions between local anaesthetics and lipid bilayers.^{20,21} However, the detailed physicochemical mechanism of the functional expression of local anaesthesia is still unclear.

Large phase-separated liposomes that are observable by optical microscopy are useful for the direct investigation of raft-like domains. Multicomponent liposomes are known to be separated into two or three phases due to the difference in thermodynamic properties of their lipids. In addition, the phase-separated domains of large liposomes are observable by fluorescence microscopy with a dye incorporated in a certain phase.^{22–27} Indeed, large phase-separated liposomes have been used as models for many studies, revealing the properties of the lipid membrane.^{22,24–30}

In cases similar to the present study, the large liposome and cell-derived giant plasma membrane vesicle (GPMV), which is similar to the multicomponent liposome, are also used as model systems. For example, it has been reported that vitamin E, Triton-X 100 and benzyl alcohol affect the



phase morphology of large multicomponent liposomes.³¹ Gray *et al.* reported that general anaesthetics, including alcohols, reduce the miscibility transition temperature (T_c) of GPMVs.³² They estimated the percentage of phase-separated vesicles and determined T_c as the temperature where 50% of vesicles have two phases. However, it is difficult to distinguish one- or two-phase liposomes near the critical point using this method as the phase boundary becomes unclear. In other words, determination of a threshold between one- and two-phase liposomes is very difficult near the critical point.

In the present study, we investigated the influence of dibucaine hydrochloride (DC·HCl), a commonly used local anaesthetic, on the phase behaviour of liposomes composed of dioleoylphosphatidylcholine (DOPC), dipalmitoylphosphatidylcholine (DPPC) and cholesterol (Chol). The DOPC/DPPC/Chol liposome is a system known to be laterally separated into liquid-ordered (Lo) and liquid-disordered (Ld) phases under certain conditions.²³ The Lo phase, mainly composed of DPPC and Chol, represents the raft-like domain, while the Ld phase represents the DOPC-rich region.²³ We calculated angle-averaged two-dimensional autocorrelation (2D-AC) functions of images of liposomes with three concentrations of DC·HCl at various temperatures without determining a threshold of one- and two-phase liposomes to quantify the phase pattern of liposomes and estimate the change in T_c . The results indicated that DC molecules reduce the T_c of ternary liposomes.

The 2D-AC and angle-averaged 2D-AC functions have been widely used in various fields, such as biology³³ and physics,^{34,35} to quantify the image pattern, such as dotted patterns.^{34,35} In addition, angle-averaged 2D-AC analysis was performed to clarify the correlation length of the GPMV surface in a similar study.³⁶ To the best of our knowledge, this is the first report demonstrating the change in T_c of ternary liposomes with addition of local anaesthetics by calculating the angle-averaged 2D-AC function of membrane domain patterns.

2 Materials and methods

2.1 Materials

DC·HCl, DOPC (chain melting temperature, $T_m \approx -18$ °C),³⁷ DPPC ($T_m \approx -41$ °C),³⁷ Chol and chloroform were purchased

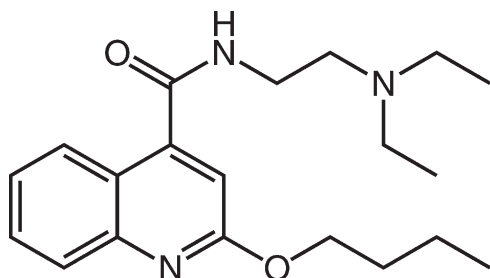


Fig. 1 Chemical structure of dibucaine (DC).

from Wako Pure Chemical Industries, Ltd. (Japan). The chemical structure of DC is shown in Fig. 1. Methanol was purchased from Showa Chemical Industry Co., Ltd. (Japan). Rhodamine B 1,2-dihexadecanoyl-*sn*-glycero-3-phosphoethanolamine, triethylammonium salt (rhodamine DHPE, $\lambda_{ex} = 560$ nm, $\lambda_{em} = 580$ nm), a fluorescent phospholipid, was obtained from Invitrogen (USA). Ultra-pure water was obtained using a WT101UV Autopure system (Yamato Scientific Co., Ltd., Japan).

2.2 Preparation of ternary liposomes

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

2.3 Observations

The liposome suspension (40 μ L) and aqueous DC·HCl solution (40 μ L) were gently mixed, and the solution was incubated at room temperature (RT, 21.7 ± 0.4 °C) for more than 10 min to achieve sufficient dispersion of DC·HCl molecules into the lipid membrane. In this study, we used 0, 0.1 and 0.4 mM of aqueous DC·HCl solutions. The final DOPC/DPPC/Chol concentration was 0.1 mM with 0, 0.05 or 0.2 mM DC·HCl and 0.5 μ M rhodamine DHPE. The mixed solution (5 μ L) was then placed between two cover glasses and 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) and monitored using a thermistor probe placed in the copper plate (under the sample). A thin layer of thermal grease was placed between the sample cell and copper plate to maintain high thermal conduction from the plate to the sample. The sample placed on the plate was incubated for more than 10 min to achieve thermal equilibrium prior to observation. The sample was then observed using a fluorescence microscope BX40 (Olympus, Japan) through a 40 \times objective lens at various temperatures. The microscopic images were recorded using a DP73 digital camera (Olympus, Japan). Excitation-light irradiation ($\lambda_{max} = 546$ nm) was applied using a Hg lamp through a WIG filter set (Olympus, Japan). The observed liposomes were unilamellar vesicles with a diameter range of approximately 15–40 μ m. The diameter of the majority of observed liposomes was 15–25 μ m, and that of a few liposomes was nearly 40 μ m.

2.4. Analysis of microscopic images

We calculated the angle-averaged 2D-AC using a macro program of ImageJ.⁴⁰ First, fluorescence microscopic images of the liposome surface were cut into 64 \times 64 pixels and converted to a greyscale (16 bit). This pixel size is appropriate



for calculating angle-averaged 2D-AC of liposomes with a diameter greater than 15 μm . The influence of pixel size on 2D-AC values is believed to be very small as most of the observed liposomes are 15–25 μm . We then calculated the two-dimensional fast Fourier transform (2D-FFT) using the brightness of each pixel in the images and inverse transformed their power spectra to derive the 2D-AC function as

$$g(r, \theta) = \mathcal{F}^{-1}[\mathcal{F}[I] \cdot \mathcal{F}^*[I]], \quad (1)$$

where $g(r, \theta)$ is the 2D-AC as a function of real space position (r, θ) , \mathcal{F} represents 2D-FFT, \mathcal{F}^* represents complex conjugation of the Fourier component, \mathcal{F}^{-1} represents the inverse 2D-FFT and I is the brightness of each pixel in the microscopic image. $\mathcal{F}[I]$ represents the 2D-FFT of I . The 2D-AC function was averaged for all angles as

$$G(r) = \langle g(r, \theta) \rangle_{\theta}, \quad (2)$$

where $G(r)$ is the angle-averaged 2D-AC as a function of radial position (r) in real space. Finally, we calculated the FFT of normalised $G(r)/G(0)$ curves, which represent the correlation between the brightness of a certain point $(r = 0)$ and another one placed within the radial distance (r) . Typical results of the analysis of three domain patterns are described in the ESI† (Fig. S1). It is not necessary to determine the phase pattern (one- or two-phase) of an ambiguous case such as that shown in Fig. S2.†

2.5 Calculation of Lo-domain circularity

Circularity was also calculated using ImageJ.⁴⁰ We transformed the microscopic images to binary images to obtain

the phase boundary of Lo/Ld. The circularity, a dimensionless parameter, of the Lo domains was defined as

$$(\text{circularity}) = \frac{4\pi A}{l^2}, \quad (3)$$

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

3 Results and discussion

We observed the DOPC/DPPC/Chol liposomes with 0, 0.05 and 0.2 mM DC·HCl at $T = 20, 25, 30, 35$ and 40 °C to confirm the influence of DC·HCl on the phase behaviour of ternary liposomes. Fig. 2 shows the typical microscopic images under each set of conditions. Since rhodamine DHPE is mainly localised in the Ld phase of phase-separated systems,^{26,41} the dark regions correspond to DPPC- and Chol-rich domains (Lo, raft-like domain), while the bright regions correspond to DOPC-rich domains (Ld). In addition, it was previously reported that dye lipids influence the phase behaviours of ternary liposomes.⁴² However, rhodamine DHPE was also incorporated into control systems; hence the effect of only local anaesthetics was observed in this study. Image (1) in Fig. 2 shows the phase-separated liposome with scattered Lo domains, which disappeared with increasing temperature (images (1)–(5)). In the case of 0 mM at 35 °C, we observed many one-phase liposomes. No phase-separated liposome exists at 40 °C, as shown in image (5). Veatch *et al.* reported

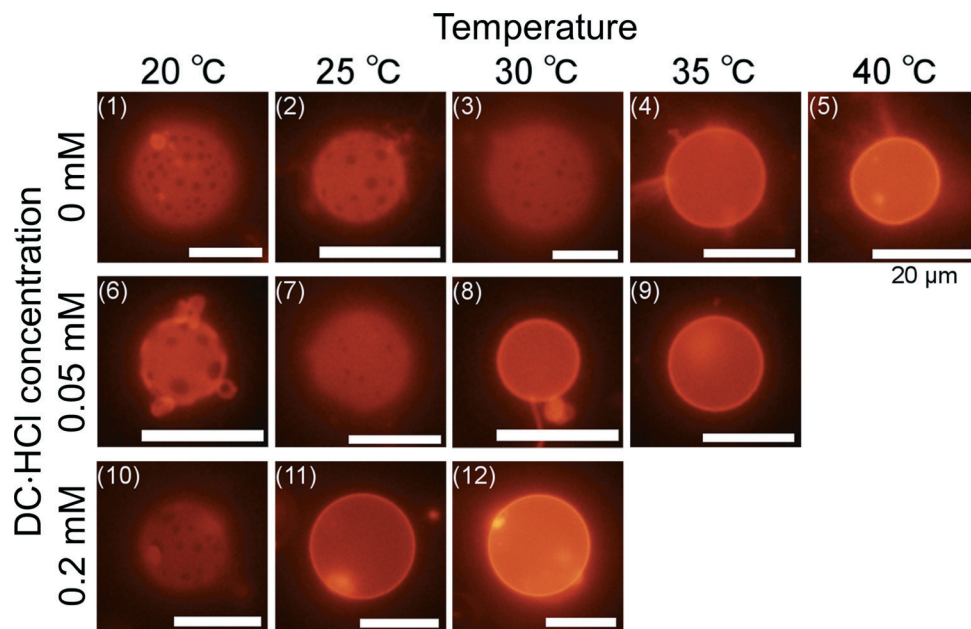


Fig. 2 Typical microscopic images of DOPC/DPPC/Chol liposomes with 0, 0.05 and 0.2 mM DC·HCl at 20, 25, 30, 35 and 40 °C. Scale bars represent 20 μm .



$T_c = 30\text{ }^\circ\text{C}$ in the case of the DOPC/DPPC/Chol = 50/25/25 system,²³ which is in agreement with the behaviour of the present system. Next, we observed the ternary liposomes at various temperatures in the case of 0.05 mM DC-HCl. We observed phase separation in most liposomes with 0.05 mM DC-HCl at 20 °C, as shown in image (6). There were several phase-separated liposomes with smaller Lo domains, as shown in image (7), and a few liposomes with one large domain, as shown in Fig. S1(c),† at 25 °C. We observed many one-phase liposomes at 30 °C (image (8)) and 35 °C (image (9)). In the case of 0.2 mM, the number of phase-separated liposomes decreased as the temperature increased. At 20 °C, the majority of liposomes were phase-separated; however, there were several liposomes which had unclear phase boundaries. At 25 °C and above, most liposomes had one phase without raft-like domains, as shown in images (11) and (12). We focused on the images of all three DC-HCl concentrations at 25 °C, as shown in images (2), (7) and (11). The Lo domains in the liposome become smaller as the DC-HCl concentration increased. These results indicate that DC-HCl molecules have the ability to change the phase behaviour of DOPC/DPPC/Chol liposomes. Moreover, we could not confirm the influence of vesicle size on phase behaviors.

We calculated the 2D-AC functions of the ternary liposome domain pattern to quantify the phase behaviour and averaged the functions. The typical results of this analysis are shown in Fig. S1,† as described above. The near region from the reference point, $r = 0$ ($r \leq 0.5\text{ }\mu\text{m}$), is a high correlation domain. The nearest similar phase domains generally exist at the high correlation region of $2.0 \leq r \leq 3.0$, due to phase separation, as in the case of the two-phase image with scattered Lo domains (Fig. S1(b),†). Therefore, the $G(r)/G(0)$ curve of phase-separated liposomes shows a certain fluctuation due to the scattered domain pattern (Fig. S1(d),†),^{34,35} and the Fourier amplitudes at low wavenumber (especially $k \approx 0.3$ and $0.6\text{ }\mu\text{m}^{-1}$) have higher values (Fig. S1(e),†). Furthermore, in the case of large phase separation, as shown in Fig. S1(c),† the high correlation region is quite long due to its large Lo or Ld domain (Fig. S1(d),†) and the $G(r)/G(0)$ does not show fluctuation within $3.5\text{ }\mu\text{m}$. Hence, the correlation of the $r \leq 0.5\text{ }\mu\text{m}$ region is also very high, while the Fourier amplitude is low (Fig. S1(e),†). In terms of this consideration, we must focus on both the short distance region of $G(r)/G(0)$ and the Fourier amplitudes at low wavenumber. In previous methods, the ambiguity of the phase pattern, as shown in Fig. S2,† was problematic, as there was no circular domain but slight fluctuations in the lipid density. However, this method enabled us to estimate the approximate percentage of phase-separated liposomes without determination of the one- and two-phase threshold of each liposome.

Fig. 3(a, c and e) show the mean values of $G(r)/G(0)$ as a function of radial position (r) at each concentration, while Fig. 3(b, d and f) show the enlarged view of the short distance region of the angle-averaged 2D-AC functions (mean \pm standard error), corresponding to Fig. 3(a, c and e), respectively. The error bars can be seen only in the enlarged view as they

confuse the measured values in Fig. 3(a, c, and e). The number of observed liposomes was greater than 10 under each set of conditions. We focused on the correlation of the short distance region, as described above. The value of the short distance region (especially $r \approx 0.25\text{ }\mu\text{m}$) of $G(r)/G(0)$ decreased as the temperature increased (Fig. 3(a and b)). It was determined that the correlation of the short distance region depends on the ratio of the phase-separated liposomes, because the correlation of the short distance region of the two-phase liposome is different from that of the one-phase liposome, as described above. The results in Fig. 3(b) indicate that the ratio of phase-separated liposomes decreased as the temperature increased. The difference in correlation values of $r \approx 0.25\text{ }\mu\text{m}$ at several temperatures gradually decreased as the DC-HCl concentration increased. This behaviour was especially shown above 30 °C. This indicates that the percentage of phase-separated liposomes becomes similar due to the DC-HCl molecules. In other words, the percentage of phase-separated liposomes at 20 and 25 °C decreased upon the addition of DC-HCl. It was suggested that the DC-HCl molecules disturb the phase separation of ternary liposomes. Further observations are required to clarify the detailed behavior of autocorrelations.

Since we need further information to confirm the effect of DC-HCl molecules on miscibility transition temperature (T_c), the FFT of the $G(r)/G(0)$ curves of individual liposomes was calculated and averaged. As shown in Fig. S1,† FFT calculations clarify the difference between one-phase (without pattern) and scattered small Lo domain patterns. Fig. S3(a–c),† show the Fourier spectrum of the $G(r)/G(0)$ curves at 0, 0.05 and 0.2 mM, respectively ($n \geq 10$). The Fourier amplitude of phase-separated liposomes at wavenumber $k \approx 0.3$ and $0.6\text{ }\mu\text{m}^{-1}$ tends to have larger values than that of one-phase liposomes, due to Lo domains, as shown in the Fig. S1(e),† the wavenumbers $k \approx 0.3$ and $0.6\text{ }\mu\text{m}^{-1}$ correspond to wavelengths $\lambda \approx 3.3$ and $1.7\text{ }\mu\text{m}$, respectively. Therefore, the averaged values of the Fourier amplitudes depend on the ratio of phase-separated liposomes. The amplitudes at $k \approx 0.3$ and $0.6\text{ }\mu\text{m}^{-1}$ decreased as the temperature increased at all concentrations.

The relationship between the Fourier amplitude and the temperature is shown in Fig. 4. The drop of the Fourier amplitudes with increasing temperature between 30 and 35 °C is shown at 0 mM, while the drop between 20 and 25 °C is shown at 0.05 and 0.2 mM. In other words, the dropping temperature region decreased with addition of the DC-HCl molecules. This indicates that the DC-HCl molecules reduce the T_c as the dropping region roughly corresponds to T_c . The T_c shifts downward more than 5 °C from 0 mM to 0.05 and 0.2 mM. The behaviour seen in Fig. 4 differs from the results of the 2D-AC functions. This is because there are large phase-separated liposomes, such as those seen in Fig. S1(c),† in the case of 0.05 mM DC-HCl at 25 °C. The Fourier amplitude was low in this case, while the correlation value of the short distance region was very high (Fig. S1(d and e),†). This shows that both the angle-averaged 2D-AC functions and their Fourier amplitudes are required to clarify the effect of



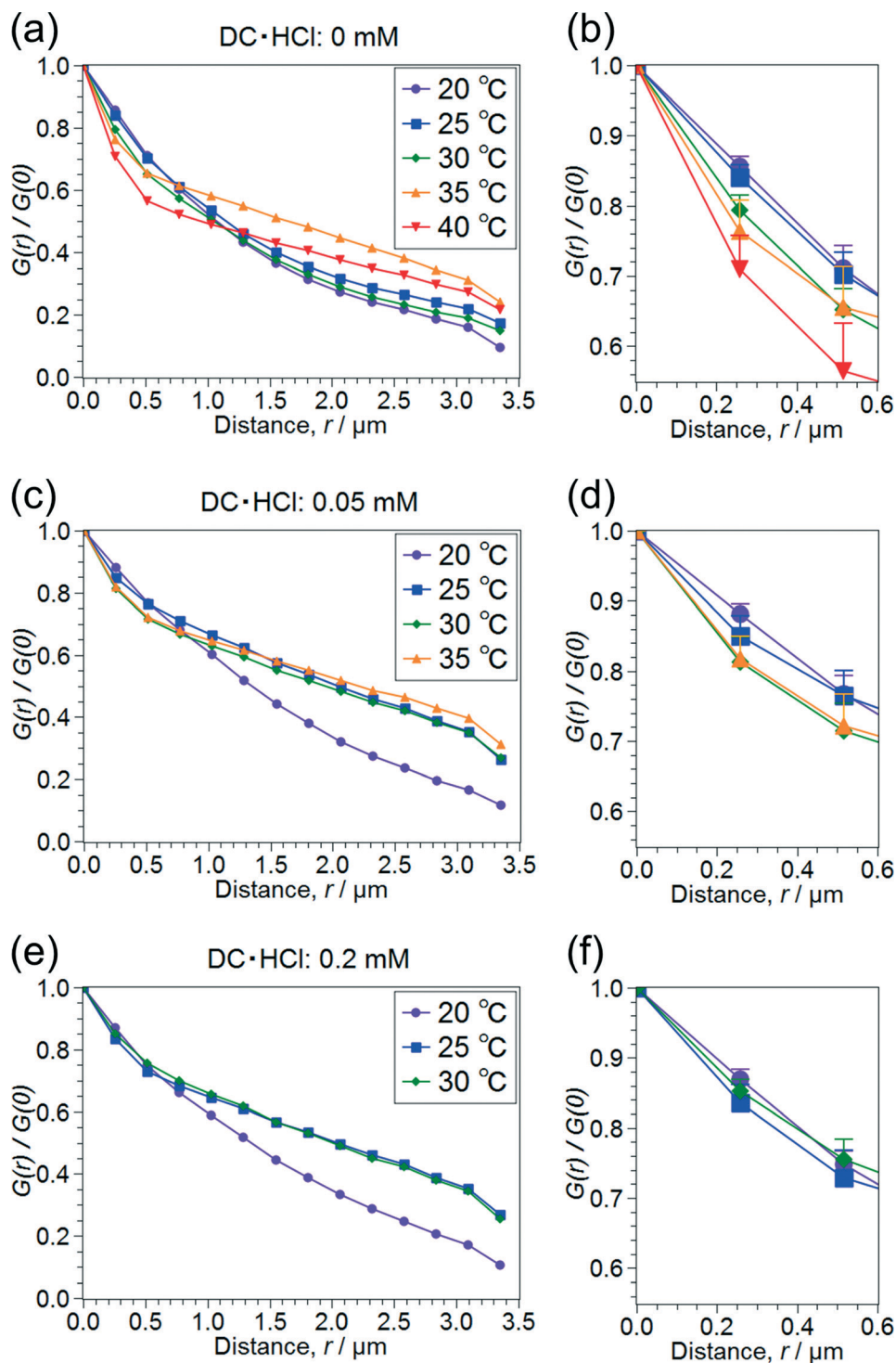


Fig. 3 Mean values of the angle-averaged 2D-AC functions ($n \geq 10$ under each set of conditions). (a, c and e) The angle-averaged 2D-AC functions of ternary liposomes with 0, 0.05 and 0.2 mM DC-HCl, respectively (error bars not shown). (b, d and f) Enlarged view of the short distance region of the angle-averaged 2D-AC function, corresponding to a, c and e, respectively. Error bars in enlarged view represent standard errors.

anaesthetics on the phase behaviour of ternary liposomes. These results suggest that DC-HCl has the ability to decrease the T_c of ternary liposomes.

In this study, we demonstrated that DC-HCl molecules induce a decrease in T_c of DOPC/DPPC/Chol liposomes. Furthermore, we calculated the Lo domain circularity to discuss

the change in the line tension of phase boundary. We were not able to determine the exact value of the line tension of the phase boundaries, as it is difficult to determine the fluctuations in Lo/Ld phase boundaries of small Lo domains. However, we could easily calculate the small Lo domain circularity, showing that the domain circularity is related to the



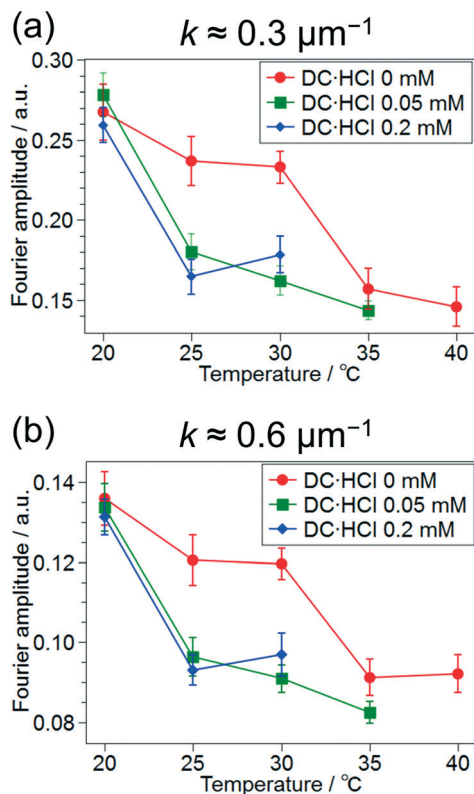


Fig. 4 (a and b) The Fourier amplitude versus temperature at $k \approx 0.3$ and $0.6 \mu\text{m}^{-1}$, respectively. Error bars represent standard errors ($n \geq 10$ under each set of conditions).

line tension of the phase boundary, as higher line tension tends to induce more circular domains and minimises energy.⁴³ Therefore, this is an appropriate method for these conditions, allowing us to estimate the change in line tension. Fig. 3 shows the circularity ($4\pi A/l^2$) at 20 °C. The circularity decreased as the DC-HCl concentration increased, suggesting that the line tension of the Lo/Ld phase boundary decreased with DC-HCl concentration. In other words, the DC-HCl molecules induce reduction in line tension of the phase boundary.

Near the critical point, line tension of the phase boundary was expressed as

$$\sigma \propto (T_c - T)^\nu, \quad (4)$$

where the σ is the line tension of the Lo/Ld phase boundary and ν is the critical exponent.^{22,44} In terms of this equation, the line tension (σ) decreased the miscibility transition temperature (T_c) when the temperature of the system (T) was constant. It was determined that insertion of the DC molecules into bilayers induced the reduction in T_c (Fig. 4), leading to the decrease in line tension. DC molecules have previously been shown to have the ability to invade lipid bilayers,⁴⁵ which led us to consider this possibility in the case of DOPC/DPPC/Chol systems. In addition, the change in line tension induced by the incorporation of other molecules was reported by several groups.^{46,47} Therefore, it was determined

that the insertion of the DC molecules into bilayers induces reduction in the miscibility transition temperature and line tension of the phase boundary.

The drop of the domain circularity with increasing DC-HCl concentration between 0.05 and 0.2 mM is shown in Fig. 5, while the difference of the Fourier amplitude between 0.05 and 0.2 mM (Fig. 4) is very small. This is because the exact line tension of the phase boundary was not obtained, but the domain circularity was calculated. We are not able to quantitatively compare the circularity with the behaviours of the Fourier amplitudes but are able to compare the line tension with T_c using eqn (4). However, the approximate behaviour of the line tension with T_c was confirmed.

Similar studies to the present work suggested that general anaesthetics disturb the phase separation of liposomes and GPMVs (reduction in T_c) using neutron diffraction and/or X-ray diffraction methods^{48,49} and direct observation.³² It is believed that the behaviour shown in these previous studies was also induced by insertion of anaesthetic molecules into bilayers. In addition, Hamada *et al.* have reported that the photoisomerisation of an azobenzene derivative reversibly switches the one-/two-phase pattern due to the change in T_c , accompanied by change in the lateral line tension.⁵⁰ This previous study also suggested that the change in line tension is related to the change in T_c .

Furthermore, compared with previous studies by Veatch *et al.*³⁶ and Honerkamp-Smith *et al.*,⁴⁴ the present results do not provide in-depth information about the critical phenomena, such as discussion of the correlation length. Hence, further detailed studies are required.

Finally, we discuss the biological implications of the present study. We demonstrated that DC-HCl has the ability to decrease the T_c of multicomponent liposomes. Previously, it has been revealed that changes in the lipid state affects the

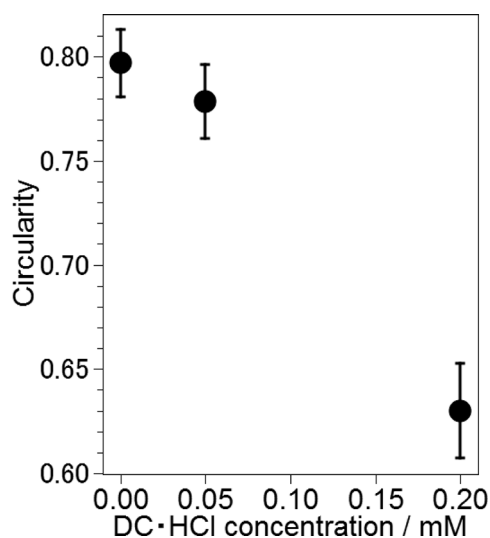


Fig. 5 Circularity of the Lo domains versus DC-HCl concentration at 20 °C ($<T_c$). Error bars represent standard errors ($n \geq 27$ Lo domains under each set of concentrations).



structure and function of the membrane protein.^{51,52} In terms of the previous and present studies, it was determined that the DC-HCl molecules have the ability to influence the function of ion channels by affecting the lipids that surround the ion channels. We expect that these findings will further the fields of pharmacology and medicinal chemistry.

4 Conclusion

In summary, we have investigated the influence of DC-HCl molecules, a local anaesthetic, on the phase behavior of DOPC/DPPC/Chol = 50/25/25 (mol%) liposomes. To confirm these effects, ternary liposomes with 0, 0.05 and 0.2 mM DC-HCl were observed at various temperatures and their phase patterns were quantified by angle-averaged 2D-AC functions. As a result, we clarified that the DC-HCl molecules have the ability to change the miscibility transition temperature (T_c) of ternary liposomes. Furthermore, we calculated the circularity of the Lo domains at 20 °C to estimate the change in line tension of the Lo/Ld phase boundary, revealing that insertion of the DC molecules into bilayers induces reduction in line tension of the phase boundary with decrease in T_c . These results suggest that the DC-HCl molecules disturb the function of ion channels (anaesthetic function of DC-HCl) by affecting the lipid bilayers that surround the ion channels. Since these physicochemical findings suggest the mechanism of anaesthetic function, this study will play an important role in pharmacology.

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References

- D. A. Haydon and B. W. Urban, *J. Physiol.*, 1983, **341**, 429–439.
- L. Ebihara, J. E. Hall, R. C. MacDonald, T. J. McIntosh and S. A. Simon, *Biophys. J.*, 1979, **28**, 185–196.
- J. Reyes and R. Latorre, *Biophys. J.*, 1979, **28**, 259–280.
- R. Reigada, *J. Phys. Chem. B*, 2011, **115**, 2527–2535.
- R. Reigada, *PLoS One*, 2013, **8**, e52631.
- Y. Muroi and B. Chanda, *J. Gen. Physiol.*, 2009, **133**, 1–15.
- D. A. Hanck, E. Nikitina, M. M. McNulty, H. A. Fozzard, G. M. Lipkind and M. F. Sheets, *Circ. Res.*, 2009, **105**, 492–499.
- T. Yamagishi, W. Xiong, A. Kondratiev, P. Vélez, A. Méndez-Fitzwilliam, J. R. Balser, E. Marbán and G. F. Tomaselli, *Mol. Pharmacol.*, 2009, **76**, 861–871.
- M. Arcisio-Miranda, Y. Muroi, S. Chowdhury and B. Chanda, *J. Gen. Physiol.*, 2010, **136**, 541–554.
- M. F. Sheets, T. Chen and D. A. Hanck, *Eur. J. Appl. Physiol.*, 2011, **461**, 91–97.
- S. Lee, S. J. Goodchild and C. A. Ahern, *J. Gen. Physiol.*, 2012, **139**, 507–516.
- J. S. Bant, T. K. Aman and I. M. Raman, *J. Neurosci.*, 2013, **33**, 4976–4987.
- T. Hata, H. Matsuki and S. Kaneshina, *Colloids Surf., B*, 2000, **18**, 41–50.
- T. Hata, T. Sakamoto, H. Matsuki and S. Kaneshina, *Colloids Surf., B*, 2001, **22**, 77–84.
- H. Matsuki, T. Hata, M. Yamanaka and S. Kaneshina, *Colloids Surf., B*, 2001, **22**, 69–76.
- K. Takeda, Y. Sano, S. Ichikawa, Y. Hirata, H. Matsuki and S. Kaneshina, *J. Oleo Sci.*, 2009, **58**, 369–377.
- H. Tsuchiya, T. Ueno, M. Mizogami and K. Takakura, *Chem.-Biol. Interact.*, 2010, **183**, 19–24.
- K. Simons and J. L. Sampaio, *Cold Spring Harbor Perspect. Biol.*, 2011, **3**, a004697.
- R. Lipowsky and R. Dimova, *J. Phys.: Condens. Matter*, 2003, **15**, S31.
- H. Tsuchiya, T. Ueno, M. Mizogami and K. Takakura, *J. Anesth. Clin. Res.*, 2010, **24**, 639–642.
- C. Bandejas, A. P. Serro, K. Luzyanin, A. Fernandes and B. Saramago, *Eur. J. Pharm. Sci.*, 2013, **48**, 153–165.
- T. Baumgart, S. T. Hess and W. W. Webb, *Nature*, 2003, **425**, 821–824.
- S. L. Veatch and S. L. Keller, *Biophys. J.*, 2003, **85**, 3074–3083.
- A. Diguët, M. Yanagisawa, Y.-J. Liu, E. Brun, S. Abadie, S. Rudiuk and D. Baigl, *J. Am. Chem. Soc.*, 2012, **134**, 4898–4904.
- M. Morita, T. Hamada, Y. Tendo, T. Hata, M. C. Vestergaard and M. Takagi, *Soft Matter*, 2012, **8**, 2816–2819.
- N. Shimokawa, M. Hishida, H. Seto and K. Yoshikawa, *Chem. Phys. Lett.*, 2010, **496**, 59–63.
- H. Himeno, N. Shimokawa, S. Komura, D. Andelman, T. Hamada and M. Takagi, *Soft Matter*, 2014, **10**, 7959–7967.
- P. Cicuta, S. L. Keller and S. L. Veatch, *J. Phys. Chem. B*, 2007, **111**, 3328–3331.
- Y. Sakuma, T. Taniguchi, T. Kawakatsu and M. Imai, *Biophys. J.*, 2013, **105**, 2074–2081.
- D. Chen and M. M. Santore, *Langmuir*, 2014, **30**, 9484–9493.
- H. S. Muddana, H. H. Chiang and P. J. Butler, *Biophys. J.*, 2012, **102**, 489–497.
- E. Gray, J. Karstlake, B. B. Machta and S. L. Veatch, *Biophys. J.*, 2013, **105**, 2751–2759.
- J. Hwang, L. A. Gheber, L. Margolis and M. Edidin, *Biophys. J.*, 1998, **74**, 2184–2190.
- S. Facsko, T. Dekorsy, C. Koerdt, C. Trappe, H. Kurz, A. Vogt and H. L. Hartnagel, *Science*, 1999, **285**, 1551–1553.
- V. Geringer, M. Liebmann, T. Echtermeyer, S. Runte, M. Schmidt, R. Rückamp, M. C. Lemme and M. Morgenstern, *Phys. Rev. Lett.*, 2009, **102**, 076102.
- S. L. Veatch, P. Cicuta, P. Sengupta, A. Honerkamp-Smith, D. Holowka and B. Baird, *ACS Chem. Biol.*, 2008, **3**, 287–293.
- R. Koynova and M. Caffrey, *Biochim. Biophys. Acta, Biomembr.*, 1998, **1376**, 91–145.
- D. D. Lasic, *Biochem. J.*, 1988, **256**, 1–11.
- K. Yoshida, Y. Fujii and I. Nishio, *J. Phys. Chem. B*, 2014, **118**, 4115–4121.



- 40 C. A. Schneider, W. S. Rasband and K. W. Eliceiri, *Nat. Methods*, 2012, **9**, 671–675.
- 41 S. Chiantia, N. Kahya, J. Ries and P. Schwille, *Biophys. J.*, 2006, **90**, 4500–4508.
- 42 S. L. Veatch, S. S. W. Leung, R. E. W. Hancock and J. L. Thewalt, *J. Phys. Chem. B*, 2007, **111**, 502–504.
- 43 A. J. García-Sáez, S. Chiantia and P. Schwille, *J. Biol. Chem.*, 2007, **282**, 33537–33544.
- 44 A. R. Honerkamp-Smith, P. Cicuta, M. D. Collins, S. L. Veatch, M. den Nijs, M. Schick and S. L. Keller, *Biophys. J.*, 2008, **95**, 236–246.
- 45 N. Weizenmann, D. Huster and H. A. Scheidt, *Biochim. Biophys. Acta, Biomembr.*, 2012, **1818**, 3010–3018.
- 46 S. A. Akimov, E. A. Hlaponin, P. V. Bashkirov, I. A. Boldyrev, I. I. Mikhalyov, W. G. Telford and I. M. Molotkovskaya, *Biochem. (Mosc.) Suppl. Ser. A: Membr. Cell Biol.*, 2009, **3**, 216–222.
- 47 J. B. Hutchison, R. M. Weis and A. D. Dinsmore, *Langmuir*, 2012, **28**, 5176–5181.
- 48 M. Weinrich, H. Nanda, D. L. Worcester, C. F. Majkrzak, B. B. Maranville and S. M. Bezrukov, *Langmuir*, 2012, **28**, 4723–4728.
- 49 M. Weinrich and D. L. Worcester, *J. Phys. Chem. B*, 2013, **117**, 16141–16147.
- 50 T. Hamada, R. Sugimoto, T. Nagasaki and M. Takagi, *Soft Matter*, 2011, **7**, 220–224.
- 51 M. Ø. Jensen and O. G. Mouritsen, *Biochim. Biophys. Acta, Biomembr.*, 2004, **1666**, 205–226.
- 52 A. Laganowsky, E. Reading, T. M. Allison, M. B. Ulmschneider, M. T. Degiacomi, A. J. Baldwin and C. V. Robinson, *Nature*, 2014, **510**, 172–175.

