## **Electronic Supplementary Information**

## Chirality-Dependent Interaction of d- and l-Menthol with Biomembrane Models

Pooja Gusain <sup>1</sup>, Shinya Ohki <sup>1</sup>, Kunihide Hoshino <sup>1,2</sup>, Yoshio Tsujino <sup>1,2</sup>, Naofumi Shimokawa <sup>1</sup>, and Masahiro Takagi <sup>\*1</sup>

<sup>1</sup> School of Materials Science, Japan Advanced Institute of Science and Technology, 1-1 Asahidai, Nomi, Ishikawa 923-1292, Japan.

<sup>2</sup> Takasago International Corporation, 5-37-1, Kamata, Ota-ku, Tokyo 144-8721, Japan.

## NMR Spectroscopy

The <sup>13</sup>C-NMR spectrum of DOPC titrated with d- or l-menthol at 15 °C is shown in Figure S3. As the concentration of d-menthol increased, the peak at 70.4 ppm, corresponding to the carbon at the g2 position of DOPC, remained unchanged. On the other hand, this peak became less intense as the l-menthol concentration was increased. In addition, the peak at 71.57 ppm, which corresponds to C1 of menthol, intensified compared to when d-menthol was added. Since the carbon at the g2 position is located in the hydrophilic region of DOPC, there is a hydrophilic interaction between l-menthol and DOPC.

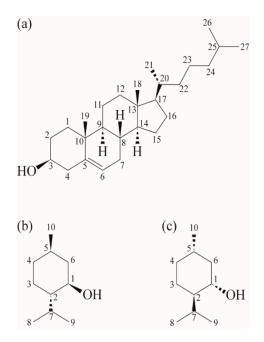


Figure S1. Systematic numbering of the chemical structures of (a) Chol, (b) l-menthol, and (c) d-menthol.

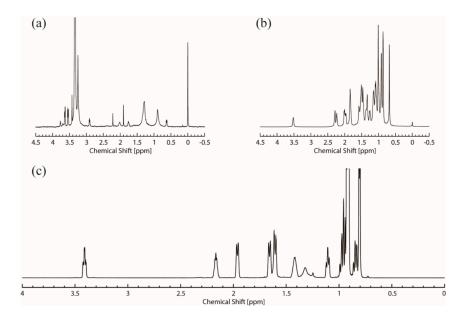


Figure S2. <sup>1</sup>H-NMR spectra of (a) DOPC, (b) Chol, and (c) menthol.

Component	<sup>1</sup> H-site	Chemical Shift (ppm)
DOPC	C18	0.895
	α	3.42
	C9 and C10	1.90
	(CH <sub>2</sub> ) <sub>n</sub>	1.30
	C1	2.01
Chol	C26-27	0.88
	C25	1.56
Menthol	C1	3.41
	C6	1.96
	C7	2.16
	C8	0.93
	С9	0.809
	C10	0.90
	C5	1.41

**Table S1**. <sup>1</sup>H chemical shift values for DOPC, Chol, and menthol at 15 °C. Since there are no significant differences in chemical shifts between d- and l-menthol, only the results for l-menthol are shown.

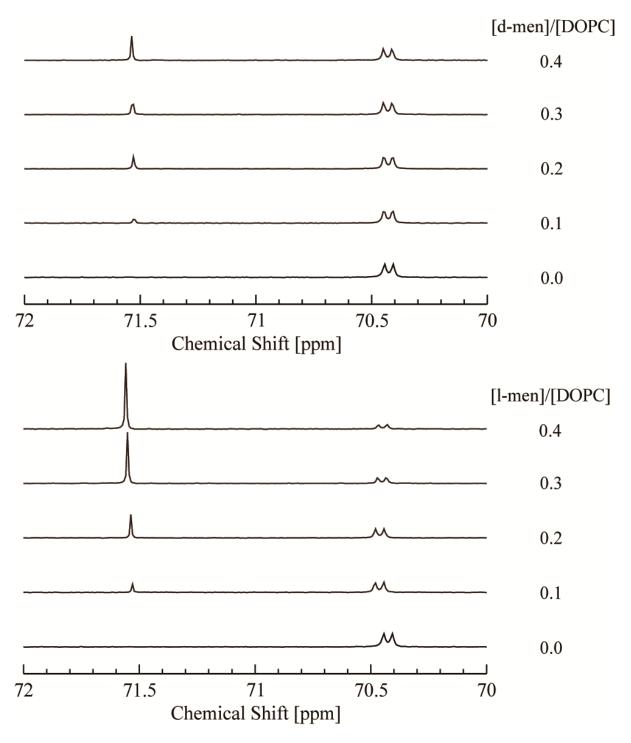
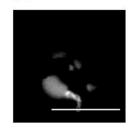


Figure S3. <sup>13</sup>C-NMR spectra of DOPC titrated with d- or l-menthol at 15 °C.

## Phase-separated Structures at Room Temperature



**Figure S4.** Typical microscopic image of reverse domain formation. Scale bar =  $10 \mu m$ .