## **Supporting Information**

## SPICA Force Field for Lipid Membranes: Domain Formation Induced by Cholesterol

Sangjae Seo<sup>1</sup> and Wataru Shinoda<sup>1</sup>\*

<sup>1</sup>Department of Materials Chemistry, Nagoya University, Furo-cho, Chikusa-ku, Nagoya,

464-8603, Japan

\* w.shinoda@chembio.nagoya-u.ac.jp

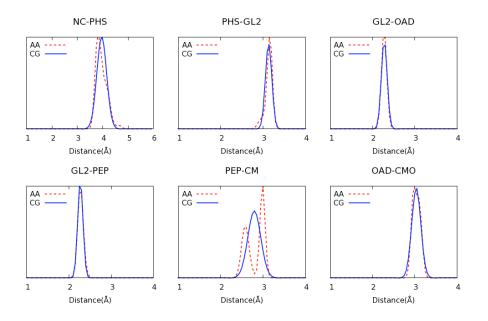


Figure S1. Bond distribution of SM from both AA and CG-MD simulations.

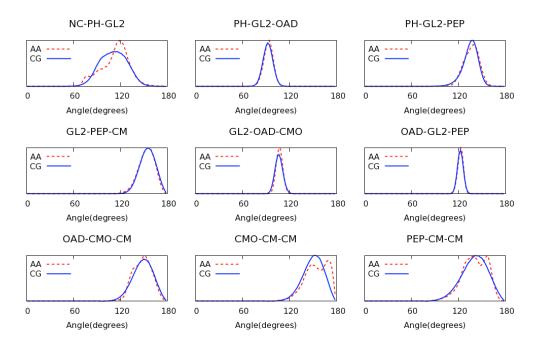
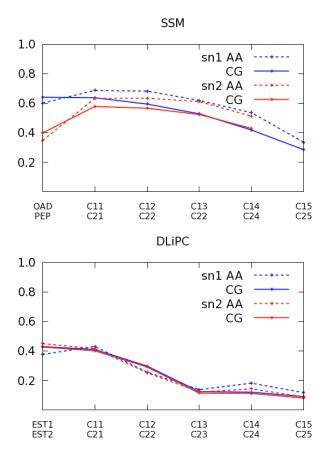
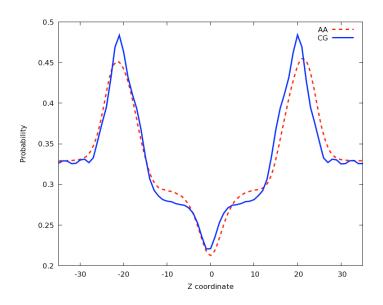


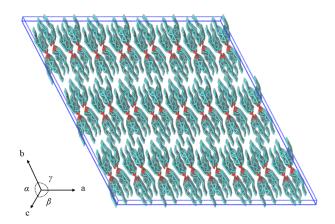
Figure S2. Angular distribution of SM from both AA and CG-MD simulations.



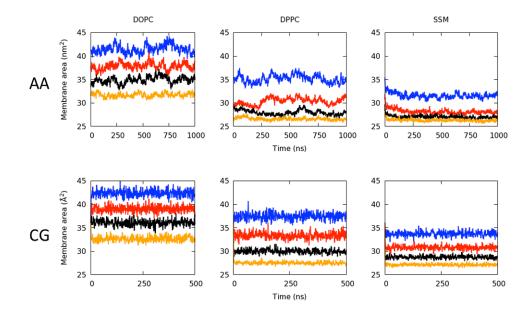
**Figure S3**. CG order parameters of SSM and DLiPC. First line and second line in x-label corresponds to the *sn*-1 and *sn*-2 segments, respectively.



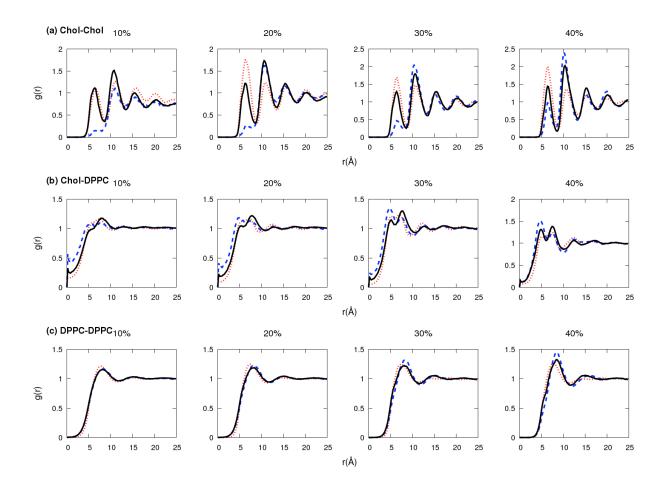
**Figure S4**. Electron density profile of SSM along the bilayer normal. The bilayer center was taken at Z = 0.



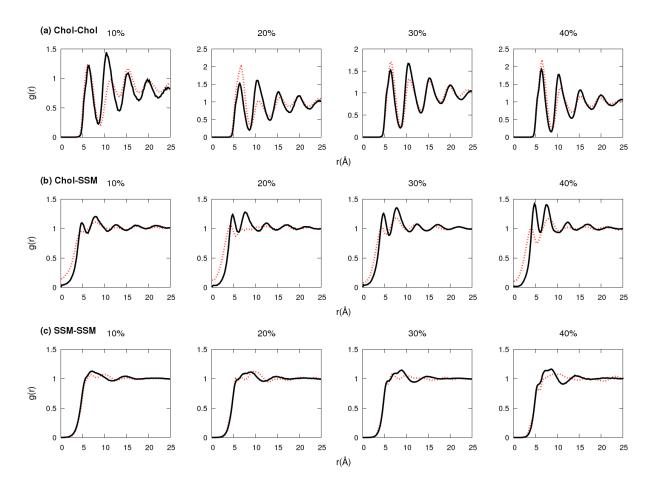
**Figure S5**. The configuration of crystal structure of CHOL. The initial coordinate of the crystal structure was downloaded from The Cambridge Crystallographic Data Centre (https://www.ccdc.cam.ac.uk/).



**Figure S6**. Time evolution of membrane area fluctuation. Color code is as follows; Blue-10% Chol, Red-20% Chol, Black-30% Chol, Orange-40% Chol.



**Figure S7**. 2D RDF of DPPC-CHOL membrane. Red dotted and blue dashed lines are the results obtained from AA-MD and CG-MD with previous SDK force field, respectively. Black solid lines represent the results from CG-MD with the current SPICA force field.



**Figure S8**. 2D RDF of SSM-CHOL membrane. Red dotted lines are the RDF obtained from AA-MD simulation and black solid lines are the results obtained from CG-MD with the SPICA force field.

System	Lipid composition	Chol.	# of	# of lipid Temp. <sup>b</sup>	Temp. <sup>b</sup>	Simulation
		contents	water <sup>a</sup>		· · ·	time
Pure	SSM	-	4304	128	323 K	500 ns <sup>c</sup>
	PSM	-	4678			
	DLiPC	-	5696			
Binary mixture	DOPC/Chol	10%	4200	128	323 K	500 ns °
		20%	4200			
		30%	4200			
		40%	4200			
	DPPC/Chol	10%	4017			
		20%	4017			
		30%	4017			
		40%	4017			
	SSM/Chol	10%	4445			
		20%	4620			
		30%	4758			
		40%	4954			
	DLiPC/Chol	10%	5506			
		20%	5220			
Ternary mixture <sup>d</sup>	DPPC/DOPC/Chol	33%	88767	2700	298 K	4 µs
	DPPC/DLiPC/Chol		100350			
	SSM/DOPC/Chol		81243			
	SSM/POPC/Chol		92376			
<sup>a</sup> The number of CC water hands can calculated by dividing three						

 Table S1. Summary of the simulated bilayer systems.

<sup>a</sup> The number of CG water beads can calculated by dividing three.

<sup>b</sup> The temperature of one component and binary mixture system was set to 323 K, which is above phase transition temperature of DPPC (314 K) and SSM (318 K). For ternary mixture, the temperature selected to follow the experiment condition.

<sup>c</sup> AA simulation for reference data was conducted for 1µs. Refer to the text.

<sup>d</sup> Ternary mixtures consist of [saturated lipid]:[unsaturated lipid]:Chol=1:1:1.