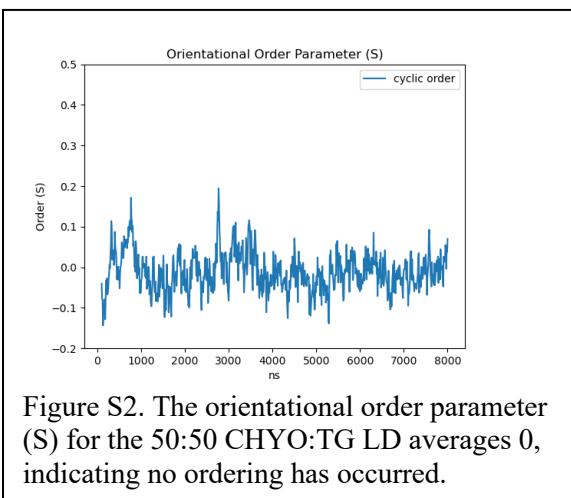
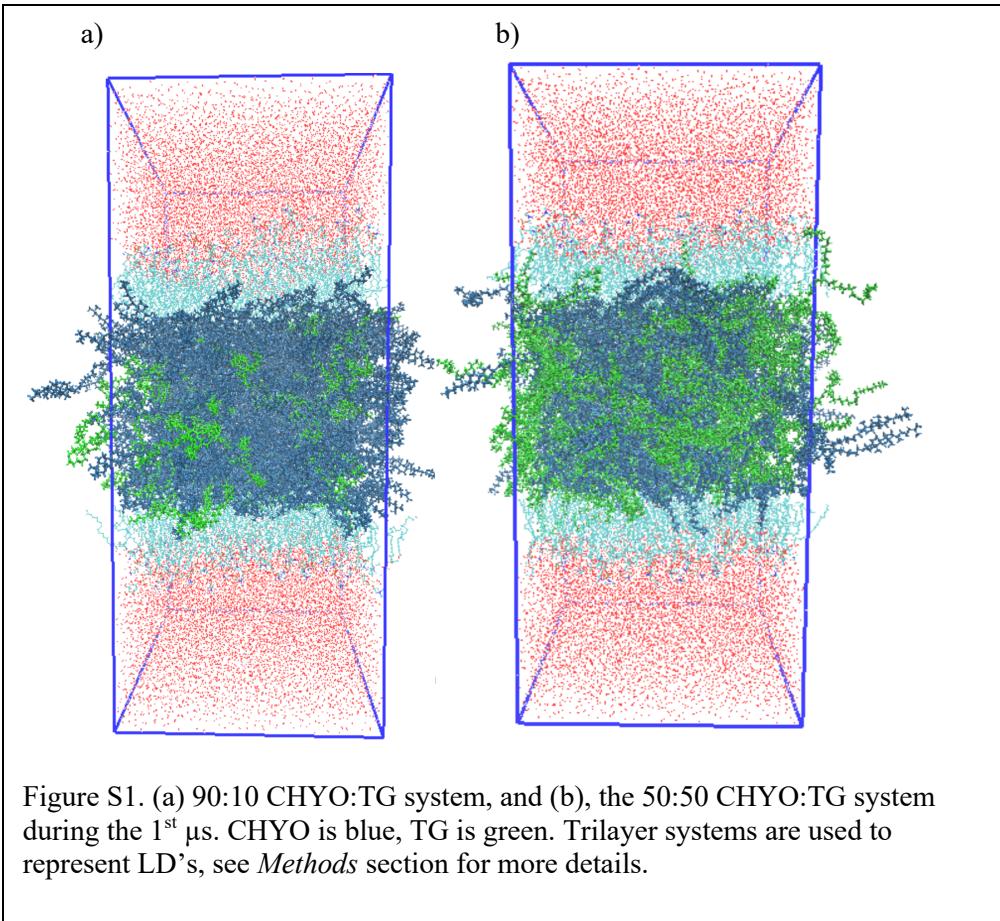


Supplementary Material

Capturing the liquid-crystalline phase transformation: Implications for protein targeting to sterol ester-rich lipid droplets

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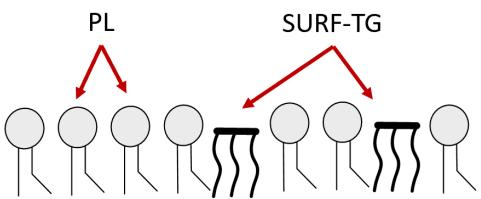


Figure S3. Representation of SURF-TGs in the PL monolayer. The SURF-TGs behave as membrane components

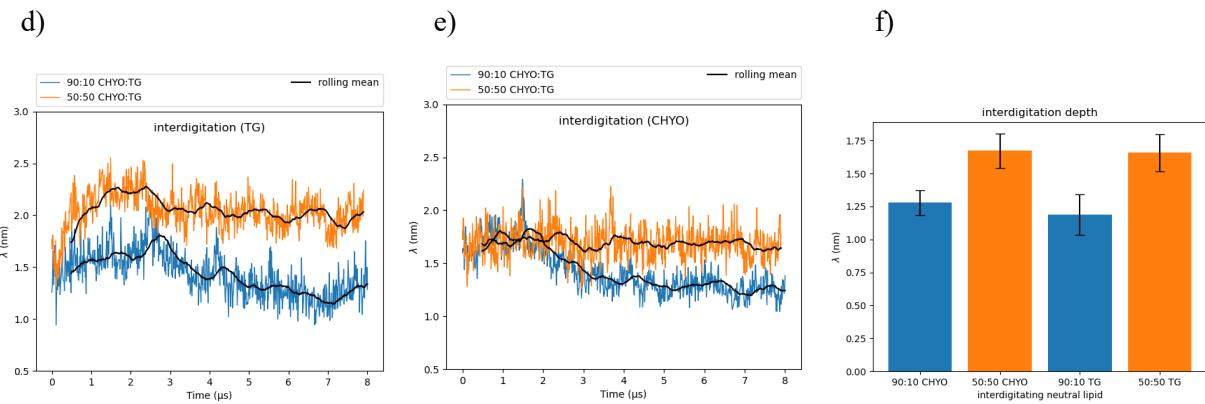
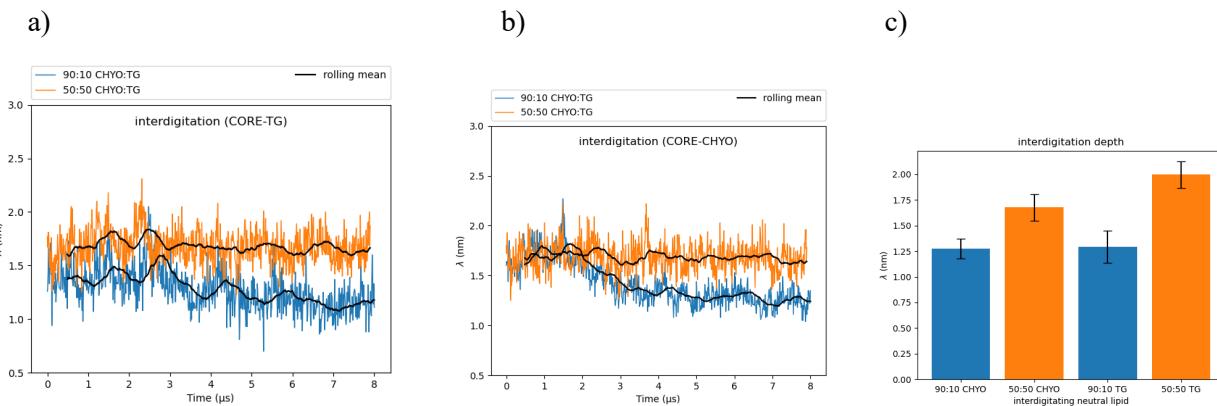


Figure S4. (a) Interdigitation as a function of time for CORE-TG, and (b) CORE-CHYO. (c) Interdigitation depth as a function of time for total-TG, and (d) Average interdigitation after initial equilibration of 3 μ s to allow for CHYO ordering to occur. (e) Interdigitation total-TG and (f) total-CHYO. (f) Average CORE interdigitation depth after initial equilibration of 3 μ s.

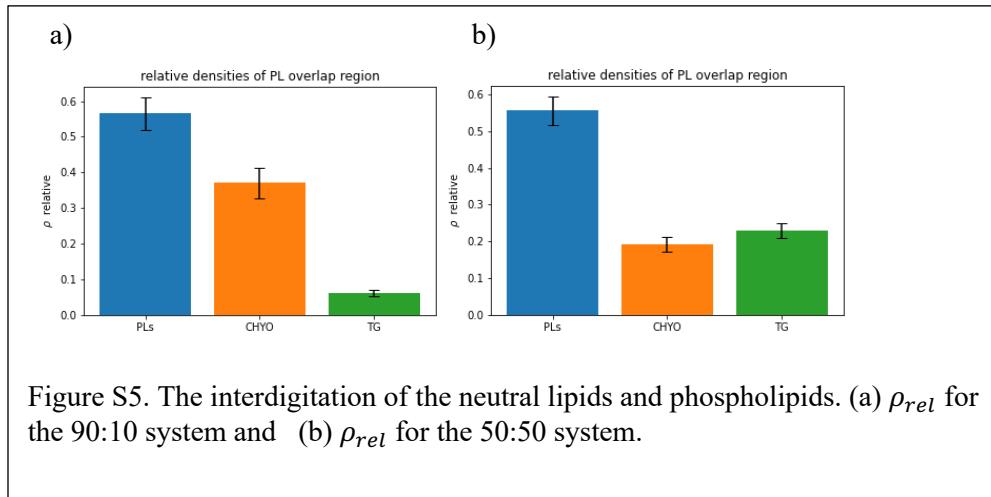
SI table S1. The average magnitude of interdigitation of neutral lipids for multiple LD species. The calculation were taken after 4 μ s to allow for equilibration

LD type	90:10 CHYO:TG LD	50:50 CHYO:TG LD	Pure TG LD
SURF-TG	0.130 \pm 0.014	0.605 \pm 0.039	\sim 1.27
CORE-TG	1.188 \pm 0.155	1.657 \pm 0.139	\sim 1.83
TG total	1.294 \pm 0.157	1.998 \pm 0.0127	\sim 2.42
CHYO	1.278 \pm 0.097	1.678 \pm 0.129	

Standard errors SURF: 90:10 TG: 0.001, 50:50 TG: 0.001

Standard errors CORE: 90:10 CHYO: 0.005, 50:50 CHYO: 0.008, 90:10 TG: 0.008, 50:50 TG: 0.007

Standard errors total: 90:10 CHYO: 0.005, 50:50 CHYO: 0.008, 90:10 TG: 0.008, 50:50 TG: 0.007



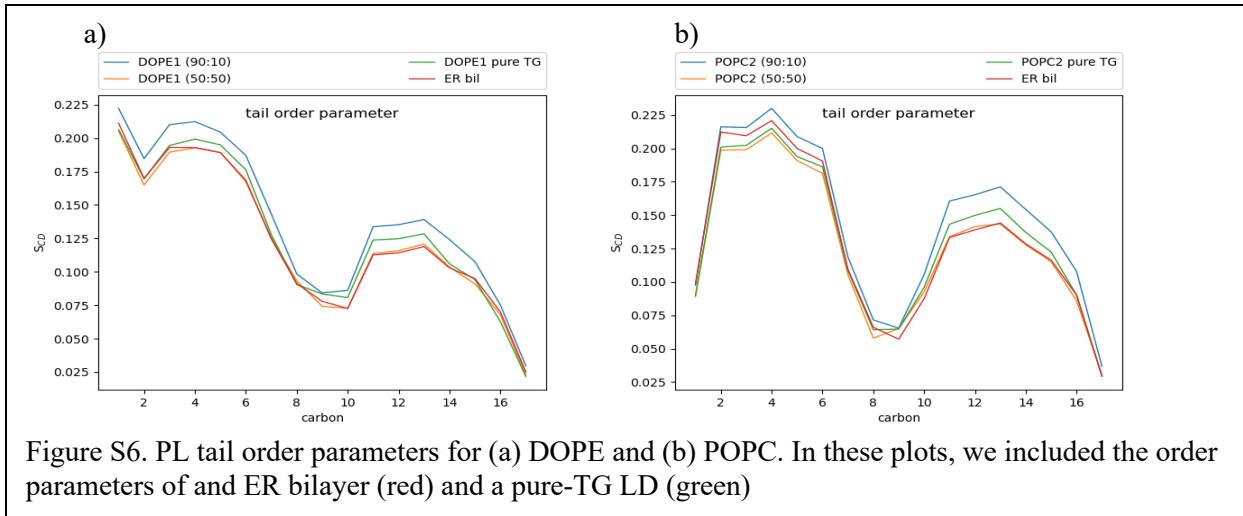


Figure S6. PL tail order parameters for (a) DOPE and (b) POPC. In these plots, we included the order parameters of an ER bilayer (red) and a pure-TG LD (green)

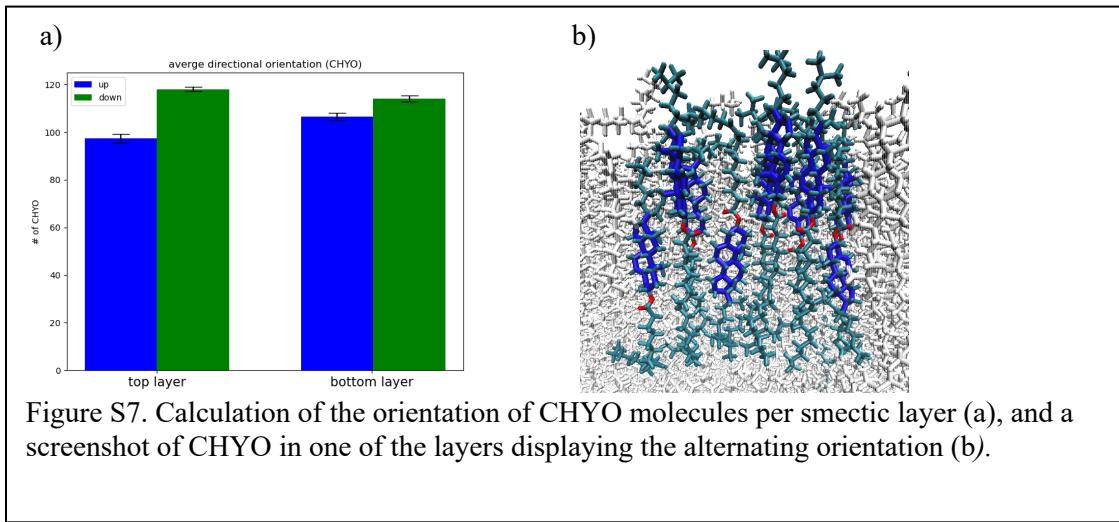


Figure S7. Calculation of the orientation of CHYO molecules per smectic layer (a), and a screenshot of CHYO in one of the layers displaying the alternating orientation (b).