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## Nanoparticle self-assembly in mixtures of phospholipids with styrene/maleic acid copolymers or fluorinated surfactants

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Self-assembling nanostructures in aqueous mixtures of bilayer-forming lipids and micelle-forming surfactants are relevant to in vitro studies on biological and synthetic membranes and membrane proteins. Considerable efforts are currently underway to replace conventional detergents by milder alternatives such as styrene/maleic acid (SMA) copolymers and fluorinated surfactants. However, these compounds and their nanosized assemblies remain poorly understood as regards their interactions with lipid membranes, particularly, the thermodynamics of membrane partitioning and solubilisation. Using  $^{19}\text{F}$  and  $^{31}\text{P}$  nuclear magnetic resonance spectroscopy, static and dynamic light scattering, and isothermal titration calorimetry, we have systematically investigated the aggregational state of a zwitterionic bilayer-forming phospholipid upon exposure to an SMA polymer with a styrene/maleic acid ratio of 3:1 or to a fluorinated octyl phosphocholine derivative called  $\text{F}_6\text{OPC}$ . The lipid interactions of SMA(3:1) and  $\text{F}_6\text{OPC}$  can be thermodynamically conceptualised within the framework of a three-stage model that treats bilayer vesicles, discoidal or micellar nanostructures, and the aqueous solution as distinct pseudophases. The exceptional solubilising power of SMA(3:1) is reflected in very low membrane saturating and solubilising polymer/lipid molar ratios of 0.10 and 0.15, respectively. Although  $\text{F}_6\text{OPC}$  saturates bilayers at an even lower molar ratio of 0.031, this nondetergent does not solubilise lipids even at >1000-fold molar excess, thus highlighting fundamental differences between these two types of mild membrane-mimetic systems. We rationalise these findings in terms of a new classification of surfactants based on bilayer-to-micelle transfer free energies and discuss the practical implications for membrane-protein research.

### Introduction

Aqueous mixtures of amphiphiles display a rich repertoire of colloidal self-assembly phenomena and give rise to diverse supramolecular structures.<sup>1</sup> In particular, the aggregational state at the nano- and mesoscale in mixtures of bilayer-forming and micelle-forming amphiphiles may greatly vary with composition, temperature, pressure, hydration, ionic strength, and other parameters.<sup>2</sup> Among the most prominent and important examples are combinations of a bilayer-forming polar lipid and a micelle-forming surfactant,<sup>3,4</sup> which have found widespread use for analytical and preparative purposes in the study of biological membranes and membrane proteins.<sup>5,6</sup> From a thermodynamic viewpoint, the supramolecular structures observed in such mixtures can, usually to a very good approximation, be treated as pseudophases.<sup>7</sup>

All other conditions remaining constant, the aggregational state at full hydration then may be captured in a two-dimensional pseudophase diagram, where the concentrations of lipid and surfactant serve as abscissa and ordinate,

respectively (Fig. 1a). In the simplest case, a coexistence range lies between two other ranges hosting either only bilayers (l) or only micelles (m), which is conceptually and formally equivalent to the appearance of a miscibility gap in mixtures of two liquids of limited mutual solubility. Oftentimes, the lipid and surfactant concentrations delimiting the coexistence range are described by straight lines referred to as saturation (SAT) and solubilisation (SOL) boundaries.<sup>4,7</sup> The slopes of these boundaries correspond to the compositions of bilayers (b) saturated with surfactant (S) and micelles (m) saturated with lipid (L), which are expressed in terms of the molar ratios  $R_S^{b,\text{SAT}} \equiv c_S^{\text{SAT}}/c_L$  and  $R_S^{m,\text{SOL}} \equiv c_S^{\text{SOL}}/c_L$ , respectively (cf. eqs 1 and 2 in ESI). These ratios allow calculation of the bilayer-to-micelle partition coefficients,  $K_L^{b \rightarrow m} = (1 + R_S^{b,\text{SAT}})/(1 + R_S^{m,\text{SOL}}) < 1$  and  $K_S^{b \rightarrow m} = K_L^{b \rightarrow m} R_S^{m,\text{SOL}}/R_S^{b,\text{SAT}} > 1$ , as well as the corresponding standard values of the bilayer-to-micelle transfer free energies,  $\Delta G_L^{b \rightarrow m,0} = -RT \ln K_L^{b \rightarrow m} > 0$  and  $\Delta G_S^{b \rightarrow m,0} = -RT \ln K_S^{b \rightarrow m} < 0$ , for the lipid and the surfactant respectively (eqs 3–6). Contrariwise, the thermodynamics of solubilisation determine the experimentally observable saturating and solubilising molar ratios according to  $R_S^{b,\text{SAT}} = (1 - K_L^{b \rightarrow m})/(K_S^{b \rightarrow m} - 1)$  and  $R_S^{m,\text{SOL}} = R_S^{b,\text{SAT}} K_S^{b \rightarrow m}/K_L^{b \rightarrow m}$ , respectively (eqs 7 and 8).

Consequently, phase diagrams of amphiphile mixtures offer quantitative insights into and, conversely, are governed by the thermodynamics of composition-dependent self-assembly.

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typical phospholipid/detergent combinations, all three of the above ranges are easily accessible (Fig. 1a), giving rise to the famous three-stage model of membrane solubilisation.<sup>3</sup> By contrast, membranophobic surfactants with a strong preference for the micellar over the bilayer phase<sup>8</sup> saturate membranes at very low concentrations, thus limiting the abundance of the purely vesicular range (Fig. 1b). At the other extreme, lipids with highly endergonic bilayer-to-micelle transfer free energies are responsible for the phenomenon of solubilisation resistance (Fig. 1c), which is particularly relevant to liquid-ordered domains, so-called “detergent-resistant membranes”, and gel-phase bilayers.<sup>9,10</sup> Unfavourable lipid/detergent interactions in both bilayer and micellar phases result in a large mixing gap, that is, a broad coexistence range (Fig. 1d). Finally, phase diagrams have numerous practical applications, since they predict the aggregational state as a function of sample composition, which can be exploited, for instance, in guiding membrane-protein reconstitution.<sup>11–13</sup>

Owing to its simple yet sound theoretical foundation and predictive power, the pseudophase concept has become popular for the thermodynamic analysis and interpretation of nano- and mesoscale self-assembly phenomena in mixtures of lipids and detergents, a term we use here to refer to any “water-soluble amphiphile with the capacity to solubilise lipids or other hydrophobic molecules”.<sup>14</sup> Recently, however, there has been growing interest in surfactants that fundamentally differ from conventional detergents with regard to chemical composition, lipid interactions, and self-assembly behaviour and, hence, appear to elude the pseudophase model.<sup>15</sup> On the one hand, these include amphiphilic styrene/maleic acid (SMA) copolymers, which solubilise lipids and lipid-embedded proteins to generate SMA-bounded lipid discs known as SMA/lipid particles (SMALPs), lipodisks, or native nanodiscs.<sup>15–18</sup> These colloidal aggregates provide a native-like bilayer environment to membrane proteins but remain amenable to nuclear magnetic resonance (NMR) and optical spectroscopies and other methods requiring small particle sizes.<sup>17–25</sup> On the other hand, fluorinated surfactants<sup>8,26,27</sup> possess very poor detergency, as they do not solubilise lipid membranes and are even believed not to interact with them at all, although there are notable exceptions.<sup>8,28</sup> Therefore, these compounds are employed for solubilising hydrophobic peptides and proteins in the presence of lipid bilayers<sup>29,30</sup> and in similar situations requiring a nanostructured environment that does not compromise membrane integrity or does so in a mild, tightly controllable manner.<sup>8</sup> Taken together, SMA copolymers and fluorinated surfactants currently receive widespread attention as promising membrane-mimetic systems since they enable gentle solubilisation and stabilisation of membrane proteins, but the thermodynamics underlying their peculiar modes of action has remained obscure.

Herein, we show (i) how mixtures of phospholipids with either SMA or a fluorinated nondetergent surfactant, which exhibit extremely strong or poor detergency, respectively, can be made amenable to experimentation and quantitative analysis; (ii) that the pseudophase-diagram approach is a valuable conceptual tool for rationalising membrane solubilisation and

nanoparticle formation by such unconventional surfactants; and (iii) how a broad range of chemically diverse surfactants may be classified in a unifying framework based on simple thermodynamic criteria.

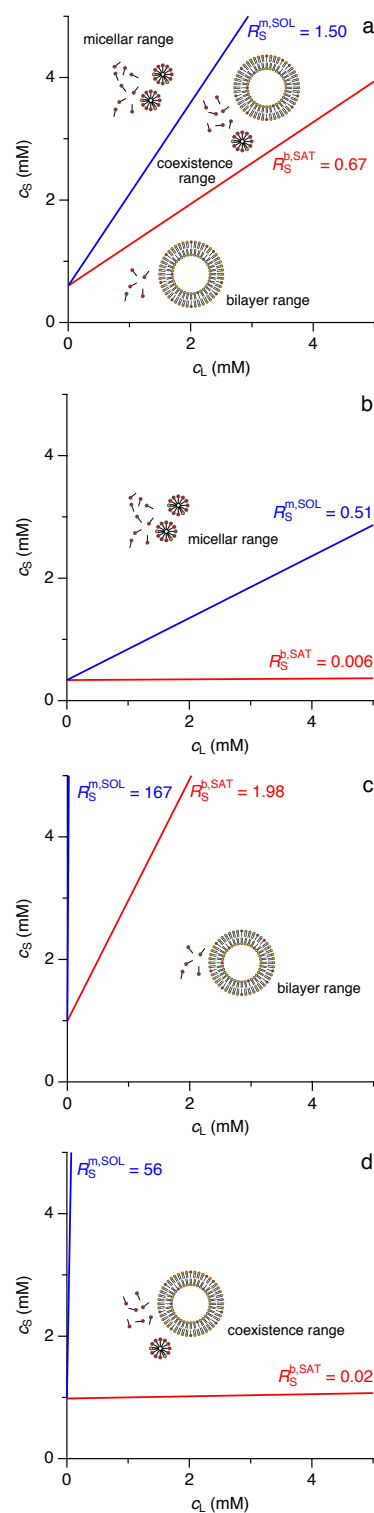


Fig. 1 Phase diagrams of aqueous mixtures of a bilayer-forming lipid and a micelle-forming surfactant with a CMC of 1 mM as predicted for various bilayer-to-micelle transfer free energies. (a) Typical phospholipid and detergent:  $\Delta G_L^{b \rightarrow m,0} = 1.0$  kJ/mol,  $\Delta G_S^{b \rightarrow m,0} = -1.0$  kJ/mol. (b) Membranophobic surfactant:  $\Delta G_L^{b \rightarrow m,0} = 1.0$  kJ/mol,  $\Delta G_S^{b \rightarrow m,0} = -10$  kJ/mol. (c) Solubilisation-resistant lipid:  $\Delta G_L^{b \rightarrow m,0} = 10$  kJ/mol,  $\Delta G_S^{b \rightarrow m,0} = -1.0$  kJ/mol. (d) Mutual insolubility of lipid and surfactant:  $\Delta G_L^{b \rightarrow m,0} = 10$  kJ/mol,  $\Delta G_S^{b \rightarrow m,0} = -10$  kJ/mol. The slopes of the phase boundaries

correspond to the surfactant/lipid molar ratios required for bilayer saturation and complete solubilisation,  $R_S^{b,SAT}$  and  $R_S^{m,SOL}$ , respectively. The predominant range is graphically highlighted in each panel.

## Results and discussion

### Solubilisation by styrene/maleic acid (3:1) copolymer

Random copolymers containing styrene and maleic acid units at various ratios are capable of extracting lipids and proteins directly from artificial<sup>16–20</sup> or native<sup>21–25</sup> bilayer membranes. This peculiarity sets them apart from other compounds that form disc-shaped colloidal particles developed for membrane-protein research, such as bicelles<sup>31</sup> and protein-bounded nanodiscs.<sup>32</sup> It has been suggested<sup>33</sup> that SMA-mediated disruption of phospholipid vesicles is initiated by penetration of its hydrophobic styrene moieties into the hydrocarbon core of the membrane and then proceeds through wrapping of the polymer chain around a lipid bilayer disc. Although mechanistic details remain to be elucidated, it seems that the excision of bilayer patches sets in at very low SMA contents in the membrane.<sup>33</sup> Thus, within the phase-diagram concept, phospholipid/SMA mixtures are expected to display low  $R_S^{b,SAT}$  values, above which further SMA addition shifts the equilibrium from polymer-saturated bilayers to SMA-bounded discs called SMALPs until all lipid is solubilised once  $R_S^{m,SOL}$  is reached.

To corroborate this solubilisation scenario and determine  $R_S^{b,SAT}$  and  $R_S^{m,SOL}$  values, we examined the formation of SMALPs by exposing large unilamellar vesicles (LUVs) formed from the zwitterionic, singly unsaturated phospholipid 1-palmitoyl-2-oleoyl-*sn*-glycero-3-phosphocholine (POPC) to increasing concentrations of SMA(3:1). For this purpose, we found <sup>31</sup>P NMR spectroscopy to be superior to more common techniques such as turbidimetry and light scattering (see below). Under our experimental conditions, the <sup>31</sup>P NMR signal from phosphorus nuclei residing in LUV bilayers is broadened beyond detection.<sup>8,34–36</sup> Hence, any peak in the spectrum is expected to be entirely due to—and, thus, proportional to the amount of—solubilised phospholipid. At a given POPC concentration, an isotropic, Lorentzian-shaped <sup>31</sup>P NMR signal appeared once the SMA(3:1) concentration exceeded a certain threshold, increased sharply upon further addition of polymer, and finally levelled off in overall intensity when a second breakpoint was reached (Fig. 2a). Both breakpoints observed in such titrations were proportional to the concentration of POPC (Fig. 2b), indicating that they reflect the onset and completion of solubilisation.<sup>3,4,37</sup> Accordingly, addition of SMA(3:1) to POPC LUVs initially leads to adsorption of the polymer to the bilayer membrane until the latter is saturated at  $R_S^{b,SAT}$ . A further rise in SMA(3:1) concentration results in solubilisation, which manifests as a linear increase in the area of the <sup>31</sup>P NMR peak before the latter plateaus once all lipid is solubilised at  $R_S^{m,SOL}$ . Plotting the SMA(3:1) concentrations at the breakpoints versus the corresponding POPC concentrations produced a phase diagram (Fig. 2c) characterised by saturating and solubilising SMA(3:1)/POPC molar ratios of  $R_S^{b,SAT} = 0.099 \pm 0.004$  and

$R_S^{m,SOL} = 0.147 \pm 0.005$ , respectively. The vesicle-to-SMALP transfer free energies derived therefrom amount to  $\Delta G_L^{b \rightarrow m,0} = (0.11 \pm 0.01)$  kJ/mol and  $\Delta G_S^{b \rightarrow m,0} = -(0.87 \pm 0.09)$  kJ/mol for POPC and SMA(3:1), respectively. The vanishing ordinate intercept of  $c_S^{aq,0} = 0$  indicates that the concentration of free polymer is negligible.

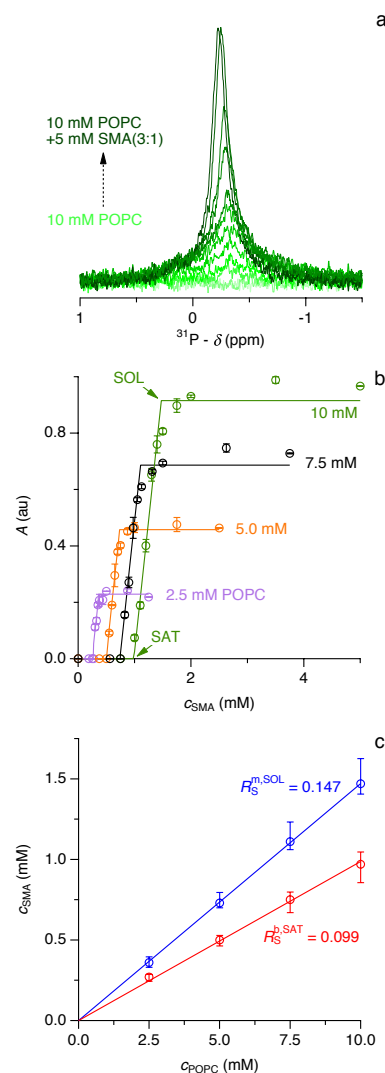


Fig. 2 Solubilisation of POPC vesicles by SMA(3:1) at 25°C as monitored by <sup>31</sup>P NMR. (a) NMR spectra of 10 mM POPC initially present as LUVs upon exposure to increasing concentrations of SMA(3:1). (b) Peak areas, A, at four POPC concentrations as function of SMA(3:1) concentration. Experimental data (circles) derived from titrations such as that in (a) and fits (lines) according to eqs 17–19. Error bars indicate standard deviations from three experiments. (c) Phase diagram of POPC/SMA(3:1). Experimental data (circles) obtained from breakpoints in (b) and linear fits (lines) according to eqs 17 and 18 marking the onset (SAT; red) and completion (SOL; blue) of solubilisation, respectively. Error bars indicate 95% confidence intervals of fits based on eqs 17–19.

In contrast with the clear-cut picture emerging from <sup>31</sup>P NMR spectroscopy, interpretation of dynamic light scattering (DLS) data proved more challenging. Particle size distributions obtained either well below  $R_S^{b,SAT}$  or above  $R_S^{m,SOL}$  were unimodal and relatively narrow, with mean values and standard deviations of  $(160 \pm 10)$  nm and  $(12 \pm 1)$  nm being in excellent agreement with the hydrodynamic diameters expected for, respectively, LUVs (Fig. 3a) and SMALPs (Fig. 3b). By contrast, intermediate SMA(3:1) concentrations gave rise to

multimodal size distributions indicative not only of coexisting LUVs and SMALPs but also of larger aggregates (Fig. 3a). The latter could represent agglomerated vesicles interconnected by polymer chains or even (hemi)fusion products. Their transient appearance and disappearance during solubilisation became more evident when the total light scattering intensity at an angle of 90°, the z-average diameter, and the polydispersity index (PDI) of a POPC suspension were monitored as functions of SMA(3:1) concentration (Fig. 3c). Both the z-average diameter and the PDI increased abruptly below  $R_S^{b,SAT}$ , that is, within the vesicular range. At the same subsolubilising SMA(3:1) concentration, a turbid slurry started to sediment in the sample vial, indicating that the steady decrease in light scattering intensity observed already below  $R_S^{b,SAT}$  was due to sedimentation of particles out of the scattering volume rather than a reduction in particle size prior to solubilisation. Formation of SMALPs above  $R_S^{b,SAT}$ , that is, within the coexistence range, gradually dissolved these large aggregates, thus reducing the z-average diameter and narrowing down the PDI. Beyond  $R_S^{m,SOL}$ , the dispersion became optically clear, and all three of the above quantities decreased monotonically with increasing SMA(3:1) concentration, reflecting a continuous reduction in SMALP size.<sup>38</sup>

#### Lipid interactions of the fluorinated surfactant F<sub>6</sub>OPC

Fluorinated surfactants are used in situations requiring colloidal amphiphile properties without compromising the integrity of lipid membranes.<sup>8,29,30</sup> The fluorinated octyl phosphocholine derivative 3,3,4,4,5,5,6,6,7,7,8,8,8-tridecafluoro-*n*-octylphosphocholine (F<sub>6</sub>OPC) self-assembles into spheroidal micelles above its CMC of 2.9 mM at 25°C but behaves like a nondetergent, as it does not solubilise POPC vesicles even at high concentrations, at elevated temperature, and during prolonged incubation.<sup>8</sup> In particular, no changes in light scattering intensity are observed on exposure of 100 μM POPC in the form of LUVs to 20 mM F<sub>6</sub>OPC, that is, a 200-fold molar excess of fluorinated surfactant over lipid.<sup>8</sup> Making the conservative assumption that such a measurement would pick up any reduction in vesicle number >10%, we conclude that ~17 mM micellar F<sub>6</sub>OPC (i.e., total concentration minus CMC) can solubilise no more than 10 μM POPC. Thus, complete solubilisation would require an F<sub>6</sub>OPC/POPC molar ratio of  $R_S^{m,SOL} > 1000$ , attesting to the nondetergency of this fluorinated surfactant. From a thermodynamic perspective (eq 6), this sets a lower limit of  $\Delta G_L^{b \rightarrow m,o} > 17$  kJ/mol on the transfer free energy of the lipid.

Still, its extremely poor solubilising power does not mean that F<sub>6</sub>OPC does not partition into lipid bilayers, although such interactions must be so weak as to elude standard protocols. In ITC membrane-partitioning experiments, the heats measured during titration of monomeric F<sub>6</sub>OPC with POPC are hardly discernible from dilution effects, and time-resolved fluorescence anisotropy reveals no changes in the nanosecond dynamics of the fluorescent bilayer probe 1,6-diphenyl-1,3,5-hexatriene (DPH) upon addition of F<sub>6</sub>OPC to POPC vesicles.<sup>8</sup> Nevertheless, the presence of lipid had a slight but

significant and systematic influence on the concentration at which F<sub>6</sub>OPC started to self-assemble into micelles, as borne out by ITC (Fig. 4) and <sup>19</sup>F NMR spectroscopy (Fig. 5). In the absence of lipid, both methods determined a CMC of 2.9 ± 0.1 mM at 25°C, in agreement with published data.<sup>8</sup> The presence of POPC raised the apparent CMC, as manifested in a shift of the sigmoidal transition observed in ITC demicellisation experiments to higher F<sub>6</sub>OPC concentrations (Fig. 4a). This shift was proportional to the concentration of lipid, furnishing a phase diagram (Fig. 4b) with a saturation threshold of  $R_S^{b,SAT} = 0.031 \pm 0.004$  and a bilayer-to-micelle transfer free energy of  $\Delta G_S^{b \rightarrow m,o} = -(8.7 \pm 0.3)$  kJ/mol (eq 16). Since the water-to-micelle transfer free energy is given as  $\Delta G_S^{aq \rightarrow m,o} = RT \ln(\text{CMC}/55.5 \text{ M}) = -(24.4 \pm 0.1)$  kJ/mol, the water-to-bilayer transfer free energy amounts to  $\Delta G_S^{aq \rightarrow b,o} = \Delta G_S^{aq \rightarrow m,o} - \Delta G_S^{b \rightarrow m,o} = -(15.7 \pm 0.4)$  kJ/mol.<sup>39</sup>

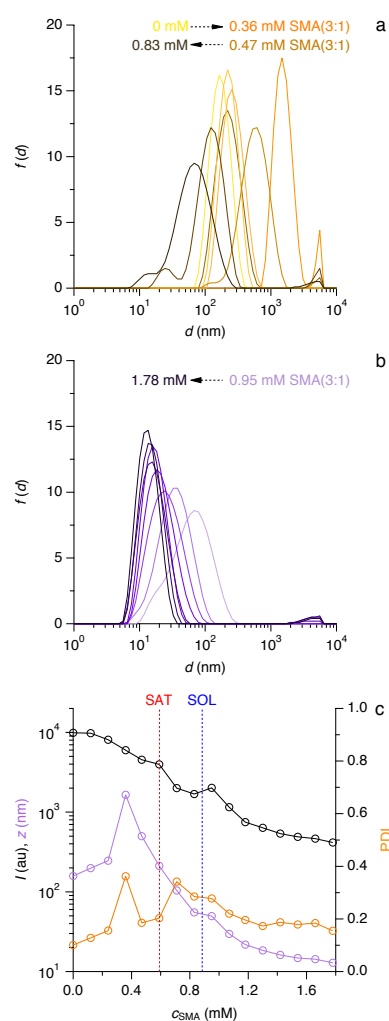


Fig. 3 Solubilisation of POPC vesicles by SMA(3:1) at 25°C as monitored by DLS. (a,b) Intensity-weighted particle size distribution functions,  $f(d)$ , versus hydrodynamic diameter,  $d$ , of 6 mM POPC initially present as LUVs upon exposure to increasing concentrations of SMA(3:1). (c) Total light scattering intensity at 90°,  $I$ , z-average diameter,  $z$ , and PDI as obtained from data in (a) and (b). Also indicated are the SAT and SOL boundaries (dashed lines) derived from <sup>31</sup>P NMR (Fig. 2).



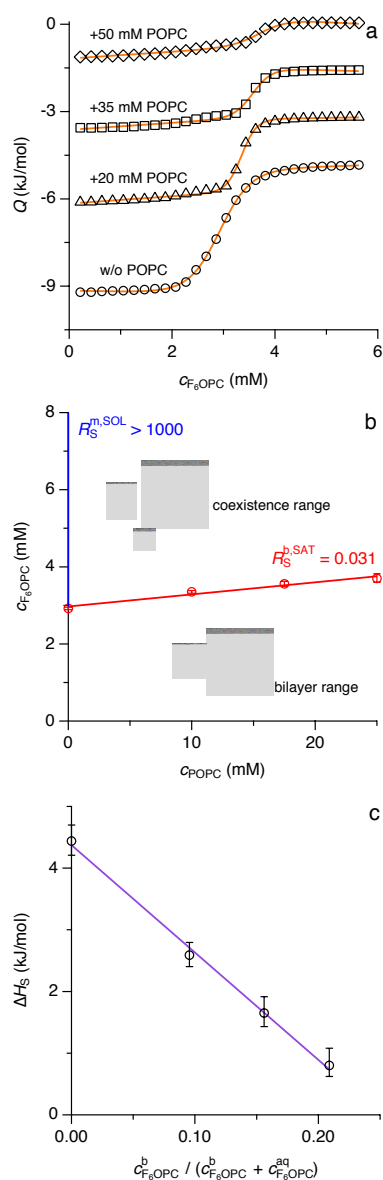


Fig. 4 Weak interactions of  $F_6$ OPC with POPC LUVs at 25°C as quantified by ITC. (a) Isotherms depicting heats of reaction,  $Q$ , (open symbols) obtained upon injection of 30 mM  $F_6$ OPC into buffer containing various POPC concentrations. Isotherms were vertically offset by  $-1.5$  kJ each. Fits (solid lines) were done according to eq 20. (b) Phase diagram of POPC/ $F_6$ OPC. The SAT boundary (red line) was obtained from linear regression of experimental values retrieved from eq 20 (red circles with 95% confidence intervals). The SOL boundary (blue line) was estimated as described in the text. (c) Molar enthalpy change,  $\Delta H_s$ , obtained from eq 20 (circles with 95% confidence intervals) and linear fit according to eq 23 (line) plotted as functions of the fraction of  $F_6$ OPC that partitioned into the bilayer as calculated from eq 22.

Closer inspection of the ITC data allows parsing the exergonic  $\Delta G_S^{b \rightarrow m, o}$  value into enthalpic and entropic contributions. In the absence of lipid, demicellisation of  $F_6$ OPC was accompanied by an enthalpy change of  $-(3.3 \pm 0.2)$  kJ/mol, which translates into a molar micellisation enthalpy of  $\Delta H_S^{aq \rightarrow m} = (3.7 \pm 0.2)$  kJ/mol upon correction for the fraction of monomeric surfactant in the syringe.<sup>40,41</sup> With increasing POPC concentration, the heats measured upon injection of micellar  $F_6$ OPC into the calorimeter cell below the CMC decreased in magnitude (Fig. 4a) because exothermic demicellisation was accompanied by an endothermic process ascribed to membrane partitioning. Plotting the molar heat of reaction

against the fraction of  $F_6$ OPC that partitioned into the bilayer (eq 22) revealed a linear relationship (Fig. 4c), the slope of which returned a value of  $\Delta H_S^{b \rightarrow m} = -(17.4 \pm 0.8)$  kJ/mol for the bilayer-to-micelle transfer enthalpy. Hence, the free-energy gain accompanying the bilayer-to-micelle transfer of  $F_6$ OPC is entirely due to a markedly favourable enthalpy change, and might be expected for the removal of a fluorocarbon chain from a hydrocarbon matrix.

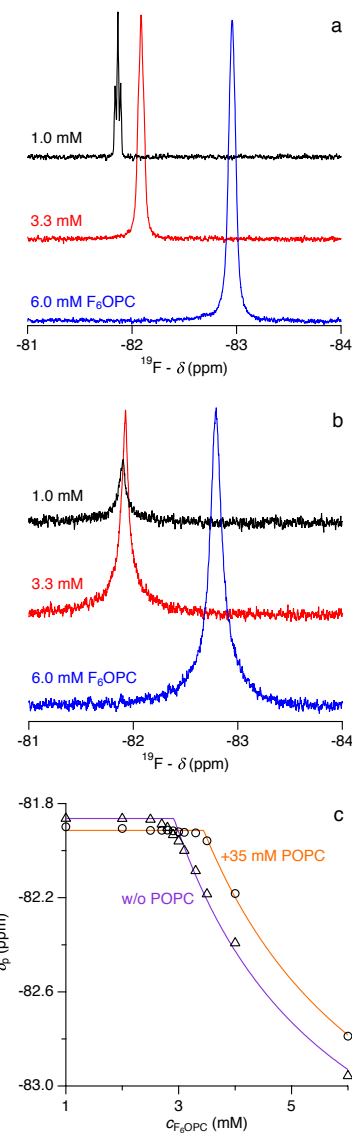


Fig. 5 Weak interactions of  $F_6$ OPC with POPC bilayers at 25°C as revealed by  $^{19}\text{F}$  NMR. (a) Representative spectra showing the  $-\text{CF}_3$  resonance in the absence of POPC at different  $F_6$ OPC concentrations. (b) Spectra in the presence of 35 mM POPC under otherwise identical conditions. At 3.3 mM  $F_6$ OPC, micelles were observed in the absence of POPC but not in its presence. Peak broadening in the presence of LUVs was due to increased viscosity and fast exchange with the membrane-bound state. (c)  $^{19}\text{F}$  NMR peak chemical shift,  $\delta_p$ , versus  $F_6$ OPC concentration in the absence (triangles) and presence (circles) of 35 mM POPC. Fits (lines) were done according to eq 24.

In the above considerations, we assumed the shift in the onset of micelle formation to arise from membrane partitioning of some  $F_6$ OPC, such that the aqueous surfactant concentration is slightly less than the total concentration.  $^{19}\text{F}$  NMR measurements corroborated this interpretation, as they revealed that the lowest  $F_6$ OPC concentration at which

micelles appear (Fig. 5a) increased in the presence of POPC (Fig. 5b). Micellisation was reflected in a sudden change in the environment and, consequently, the chemical shift of fluorine nuclei when the concentration of F<sub>6</sub>OPC exceeded the apparent CMC. With  $R_S^{b,SAT} = 0.031$  derived from ITC (Fig. 4b), we expect a total POPC concentration of 35 mM, only half of which is accessible from the outside of the vesicles, to raise the onset of micellisation by  $0.031 \times 0.5 \times 35 \text{ mM} = 0.54 \text{ mM}$ , which was indeed confirmed experimentally (Fig. 5c).

### Thermodynamics of membrane solubilisation

Considerable efforts are being made towards replacing conventional detergents by milder nanostructured alternatives that should better preserve the native conformations and functions of membrane proteins. Among these, SMA copolymers and fluorinated surfactants appear particularly promising, but a better understanding of their interactions with phospholipid bilayer membranes is mandatory if one is to exploit their full potential. In particular, the powerful arsenal of experimental approaches and thermodynamic concepts that have proven invaluable for studying membrane partitioning and solubilisation by detergents<sup>4,7</sup> has remained largely unexplored for these surfactants. In the following, after setting the stage by reviewing the thermodynamic signatures of established detergents, we discuss the peculiarities of SMA(3:1) and F<sub>6</sub>OPC in the framework of the pseudophase concept and propose a new classification of surfactants based on bilayer-to-micelle transfer free energies.

### Synthetic head-and-tail detergents

The bilayer-to-micelle transfer free energies for many combinations of phospholipids and detergents consisting of a polar headgroup and a hydrocarbon tail are characterised by averages and standard deviations of  $\Delta G_L^{b \rightarrow m,0} = (1.2 \pm 0.5) \text{ kJ/mol}$  and  $\Delta G_S^{b \rightarrow m,0} = -(0.7 \pm 0.4) \text{ kJ/mol}$  (Table S1). The latter value reflects a modest preference of the detergent for the micellar phase, while the former expresses a somewhat more pronounced preference of the lipid for the bilayer phase. Together, these thermodynamic hallmarks result in phase diagrams typically revealing saturation and solubilisation slopes of  $R_S^{b,SAT} = 1.4 \pm 0.6$  and  $R_S^{m,SOL} = 2.9 \pm 1.1$ , respectively.

Gradual addition of detergent to a lipid bilayer initially results in partitioning of the detergent between the aqueous phase and the membrane, thus raising the chemical potentials of the detergent in both phases.<sup>7</sup> Micelle formation begins once the chemical potentials in these two phases reach that in lipid-saturated micelles, just as micellisation in the absence of lipid would set in when the chemical potential of detergent monomers in the aqueous phase reaches that of pure detergent micelles. In both the bilayer and the micellar phases, the interactions of most head-and-tail detergents with lipids are adequately described by models assuming ideal or near-ideal mixing, such as regular solution theory.<sup>7</sup> Consequently, these detergents tend to distribute homogeneously within each phase without forming stable domains or clusters. Membrane insertion of such curvophilic, cone-shaped

molecules generates monolayer curvature stress and, if the detergent does not rapidly flip-flop across the membrane, additional bilayer curvature stress caused by asymmetric expansion of only one leaflet. The bilayer relaxes part of this stress by acyl chain disordering, which, in turn, results in membrane thinning and concomitant lateral expansion, as demonstrated by <sup>2</sup>H solid-state NMR spectroscopy<sup>42</sup> and time-resolved DPH fluorescence anisotropy.<sup>43</sup> The latter method also suggests that this relaxation mechanism is about equally effective for chemically diverse detergents as long as they distribute themselves evenly within the bilayer. The membrane's capacity to alleviate monolayer curvature stress by chain fluidisation appears to be exhausted once the hydrophobic thickness of one leaflet is reduced to  $\sim 9 \text{ \AA}$ ,<sup>43</sup> at which point the degree of acyl chain disorder reaches a critical threshold.<sup>44</sup>

### Bile salts, derivatives, and lipopeptides

Surfactants whose polar group is not clearly segregated from the hydrophobic moiety can be powerful detergents but appear to be more difficult to accommodate in a bilayer environment. This is exemplified by the naturally occurring bile salts cholate and deoxycholate and their synthetic derivatives 3-[(3-cholamidopropyl)dimethylammonio]-1-propanesulphonate (CHAPS) and 3-[(3-cholamidopropyl)dimethylammonio]-2-hydroxy-1-propanesulphonate (CHAPSO). By comparison with alkyl chain surfactants,  $\Delta G_L^{b \rightarrow m,0} = (0.6 \pm 0.3) \text{ kJ/mol}$  is less endergonic and  $\Delta G_S^{b \rightarrow m,0} = -(1.6 \pm 0.8) \text{ kJ/mol}$  is more exergonic for this class of detergents (Table S2). These free energies translate into low saturation and solubilisation ratios of  $R_S^{b,SAT} = 0.2 \pm 0.1$  and  $R_S^{m,SOL} = 0.6 \pm 0.2$ , respectively. Even these measures, the strong detergency of these compounds results from their more unfavourable bilayer incorporation and the ease with which their micelles accommodate phospholipids. These properties can be traced back to the rigid, sterol-based scaffold, which does not impede micelle formation but is less compatible with incorporation into a bilayer environment. Therefore, and in contrast with head-and-tail surfactants, such detergents tend not to mix homogeneously but rather form clusters within the bilayer.<sup>45</sup> This renders it more difficult for the bilayer to counteract curvature stress, as inferred from the observation that membrane insertion of such compounds does not significantly increase acyl chain disorder,<sup>43</sup> and results in low saturation and solubilisation thresholds.

The same is true for surfactin (Table S3), a *Bacillus subtilis* lipopeptide that interacts with and solubilises POPC membranes with transfer free energies of  $\Delta G_L^{b \rightarrow m,0} = 0.40 \text{ kJ/mol}$  and  $\Delta G_S^{b \rightarrow m,0} = -1.4 \text{ kJ/mol}$  and threshold ratios of  $R_S^{b,SAT} = 0.22$  and of  $R_S^{m,SOL} = 0.46$ .<sup>46</sup> A similar case can be made for P2A2, a dipalmitoylated peptide originally designed for the delivery of biological probes and drugs across the blood-brain barrier.<sup>47</sup> Depending on the mixing ratio with phospholipids, P2A2 forms defined nanodisks or liposomal assemblies and therefore can deliver both hydrophilic and hydrophobic cargos.<sup>48</sup> With a bilayer-to-micelle transfer free energy of  $\Delta G_S^{b \rightarrow m,0} = -5.6 \text{ kJ/mol}$ , P2A2

even more membranophobic than bile salts and their derivatives. More importantly, P2A2 provides a favourable micellar environment to POPC, whose bilayer-to-micelle transfer free energy amounts to only  $\Delta G_L^{b \rightarrow m,0} = 0.3$  kJ/mol. Together, these unusual thermodynamic signatures give rise to extremely low critical ratios of  $R_S^{b,SAT} = 0.012$  and  $R_S^{m,SOL} = 0.13$ . Accordingly, POPC membranes become saturated with lipopeptide when the latter accounts for only ~1% of the molecules in the bilayer, and P2A2 can solubilise an ~7-fold molar excess of POPC into mixed micelles. This enormous solubilising power is thought to arise from a disc-shaped micellar morphology, which can accommodate large amounts of lipid in a bilayer patch surrounded by P2A2 and, additionally, offers the advantage of preserving the biologically active, binding-competent conformation of the peptide moiety of P2A2.<sup>48</sup>

#### Detergent activity of styrene/maleic acid copolymers

The transfer free energies describing the formation of SMALPs from POPC bilayers upon exposure to SMA(3:1) amount to  $\Delta G_L^{b \rightarrow m,0} = 0.11$  kJ/mol and  $\Delta G_S^{b \rightarrow m,0} = -0.87$  kJ/mol. Accordingly, transfer of POPC from bilayer vesicles into SMALPs incurs only a minute free-energy cost and is much less unfavourable than solubilisation by any of the above detergents, including P2A2. In keeping with the definition of a detergent as being a “water-soluble amphiphile with the capacity to solubilise lipids or other hydrophobic molecules”,<sup>14</sup> SMA(3:1) appears as an extremely powerful detergent. Hence, extraction of membrane proteins and lipids with the aid of SMA can be regarded as “detergent-free”<sup>15–25</sup> only insofar as no conventional, micelle-forming detergent is involved in this process. This is, of course, not in contradiction with the demonstrated usefulness of SMA as a mild, structure- and activity-preserving solubilising agent for membrane proteins and membrane-protein complexes.<sup>15–25</sup> In fact, the above values quantitatively demonstrate that, whereas SMA(3:1) is preferentially located in SMALPs rather than vesicular membranes, the lipid is almost indifferent, thus reflecting, from a thermodynamic perspective, the native-like properties of the bilayer disc forming the SMALP core.

Two words of caution are in place here: First, for most examples considered so far, the lipid and surfactant components are roughly similar in size, which justifies the use of thermodynamic formalisms based on mole fractions, thereby ignoring size differences between the two mixing components. This approximation is less good for mixtures of phospholipids with polymers such as SMA(3:1), whose number-average molar mass exceeds that of POPC by a factor of ~5. Thus, low critical molar ratios of  $R_S^{b,SAT} = 0.099$  and  $R_S^{m,SOL} = 0.147$  translate into SMA(3:1)/POPC mass ratios of 0.52 g/g and 0.77 g/g, respectively. These values are comparable to those of typical detergents such as *n*-octyl- $\beta$ -D-glucopyranoside,<sup>49</sup> with  $R_S^{b,SAT} = 1.22$  and  $R_S^{m,SOL} = 2.38$  corresponding to detergent/POPC mass ratios of 0.47 g/g and 0.91 g/g, respectively. Accounting for such size dissimilarities by more sophisticated solution theories would require more detailed knowledge of the molar-mass distribution of the

polymer and the lipid/polymer interactions in vesicles and SMALPs. In the absence of such information, a quick calculation reveals that any such correction would have only a minor impact, though: for instance, multiplication of both  $R_S^{b,SAT}$  and  $R_S^{m,SOL}$  by the above factor of 5 yields modestly altered transfer free energies of  $\Delta G_L^{b \rightarrow m,0} = 0.37$  kJ/mol and  $\Delta G_S^{b \rightarrow m,0} = -0.62$  kJ/mol. Second, another complication arises from the fact that lipid and polymer molecules do not mix homogeneously, as would be required for treating SMALPs as a pseudophase, but rather segregate into distinct domains; the same limitation applies to detergents that form clusters with lipid bilayers, as discussed in the preceding section.<sup>43,45</sup> In these cases, free-energy values should not be taken as thermodynamically rigorous measures but as phenomenological yet predictive descriptors.

#### Fluorinated surfactants: detergency vs. lipophobicity

By analogy to the poor miscibility of liquid hydrocarbons and fluorocarbons, fluorinated surfactants are usually assumed to be both hydrophobic and lipophobic, that is, reluctant to interact not only with water but also with lipids. We have recently shown,<sup>8</sup> however, that fluorination and detergency are not mutually exclusive, as the fluorinated surfactant 3,3,4,4,5,5,6,6,7,7,8,8,8-tridecafluoro-*n*-octyl- $\beta$ -D-maltopyranoside ( $F_6$ OM) readily partitions into POPC bilayers at low surfactant concentrations, with  $\Delta G_S^{aq \rightarrow b,0} = -25.3$  kJ/mol, and solubilises them at higher concentrations. With transfer free energies of  $\Delta G_L^{b \rightarrow m,0} = 1.7$  kJ/mol and  $\Delta G_S^{b \rightarrow m,0} = -1.0$  kJ/mol and critical molar ratios of  $R_S^{b,SAT} = 1.0$  and  $R_S^{m,SOL} = 3.0$ ,  $F_6$ OM is indistinguishable from alkyl chain detergents as regards the thermodynamics of solubilisation (Table S3). Nevertheless, this fluorinated detergent does not induce acyl chain disordering and, thus, appears to form clusters rather than being distributed homogeneously within the membrane.<sup>8</sup> The present data show that  $F_6$ OPC, which carries the same fluorinated chain as  $F_6$ OM but a zwitterionic phosphocholine instead of a neutral maltose headgroup, interacts with hydrogenated phospholipids weakly but measurably. However, a distinction has to be made between interactions in the bilayer membrane and those within the micellar phase. Comparably small absolute values of the water-to-bilayer transfer free energy of  $\Delta G_S^{aq \rightarrow b,0} = -15.8$  kJ/mol and the bilayer saturation threshold of  $R_S^{b,SAT} = 0.032$  explain why earlier attempts failed in quantifying the weak membrane affinity of  $F_6$ OPC.<sup>8</sup> Still, this interaction is significant enough to be quantified by dedicated calorimetric (Fig. 4) and NMR (Fig. 5) protocols. By contrast, incorporation of hydrogenated phospholipids into  $F_6$ OPC micelles is so unfavourable that no signs of solubilisation could be observed even under very harsh conditions excluding kinetic barriers,<sup>8</sup> thus explaining why  $F_6$ OPC but not  $F_6$ OM can be used together with free-standing lipid bilayers.<sup>29</sup>

#### Thermodynamic classification of surfactants

Surfactants are usually classified with respect to chemical properties such as headgroup ionisation and hydrophilic-lipophilic balance<sup>50</sup> or practical aspects such as their suitability



for membrane-protein applications.<sup>5,6</sup> Here, we propose a classification based on thermodynamic grounds by considering the two critical mole ratios,  $R_S^{b,SAT}$  and  $R_S^{m,SOL}$ , or—equivalently but more revealingly—the bilayer-to-micelle transfer free energies,  $\Delta G_L^{b \rightarrow m,0}$  and  $\Delta G_S^{b \rightarrow m,0}$ . Since bile salts, their derivatives, lipopeptides, and SMA(3:1) display comparably low saturation and solubilisation thresholds, these strong detergents cluster in the lower left corner in a plot of  $R_S^{m,SOL}$  versus  $R_S^{b,SAT}$  (Fig. 6a). A thermodynamic explanation for this empirical observation comes from the corresponding transfer free energies (Fig. 6b). At given  $\Delta G_S^{b \rightarrow m,0}$ , the transfer of phospholipids from bilayer membranes into mixed micelles is less unfavourable for these detergents, that is,  $\Delta G_L^{b \rightarrow m,0}$  is less endergonic. Conversely, at given  $\Delta G_L^{b \rightarrow m,0}$ , the bilayer-to-micelle transfer of the detergents themselves is more favourable, that is,  $\Delta G_S^{b \rightarrow m,0}$  is more exergonic.

Qualitatively, it takes two properties for a detergent to be an efficient solubiliser. First, the detergent itself must display a clear preference for micellisation over membrane partitioning, as reflected in a markedly exergonic  $\Delta G_S^{b \rightarrow m,0}$ . Second, these micelles have to provide a suitable environment for the lipid in question, as manifested in an only moderately endergonic  $\Delta G_L^{b \rightarrow m,0}$ . Because bile salts, peptides, and polymers are less compatible with a bilayer structure than head-and-tail detergents are, one intuitively expects them to better fulfil the first prerequisite. Moreover, upon solubilisation, these strong detergents tend to give rise to discoidal nanostructures such as CHAPS(O)-containing bicelles,<sup>31</sup> P2A2-bounded nanodiscs,<sup>48</sup> or SMALPS,<sup>15–18</sup> all of which provide a more membrane-like environment than spherical, ellipsoidal, or cylindrical micelles typically formed by head-and-tail detergents. More quantitatively, we can define a simple and unique thermodynamic criterion to distinguish two classes of surfactants: without exception,  $\Delta G_L^{b \rightarrow m,0} < -0.65 \Delta G_S^{b \rightarrow m,0}$  for strong detergents such as bile salts, their derivatives, lipopeptides, and SMA(3:1), whereas relatively weak, synthetic head-and-tail detergents obey the relationship  $\Delta G_L^{b \rightarrow m,0} > -0.65 \Delta G_S^{b \rightarrow m,0}$  (Fig. 6b; Tables S1–3). By contrast, the ratio  $R_S^{m,SOL}/R_S^{b,SAT} = K_S^{b \rightarrow m}/K_L^{b \rightarrow m}$  and, thus, the difference  $\Delta G_L^{b \rightarrow m,0} - \Delta G_S^{b \rightarrow m,0}$  (eq 9), which determine the relative width of the coexistence range, are not significantly different between the two classes of surfactants (Tables S1 and S2).

Heerklotz and Seelig<sup>51</sup> have proposed a thermodynamic classification of detergents by correlating the CMC with the water-to-bilayer partition coefficient.<sup>52</sup> The product of these two parameters decreases with increasing detergent strength, as it reflects a more pronounced preference for the micellar over the bilayer phase. Although relying exclusively on the partitioning behaviour of the detergent,  $\Delta G_S^{b \rightarrow m,0}$ , without considering that of the lipid,  $\Delta G_L^{b \rightarrow m,0}$ , this furnishes a sensible measure of detergency within the class of head-and-tail surfactants. This is because, in these cases, the bilayer-to-micelle transition is primarily governed by the intrinsic curvatures of both lipid and detergent, so that the product of the CMC and the partition coefficient provides a good predictor of  $R_S^{b,SAT}$ .<sup>51</sup> However, this relationship breaks down

when factors other than curvature come into play, as is the case with CHAPS, which is a much more potent detergent than would be expected from its high CMC and strong membrane affinity.<sup>51</sup> At the other extreme is F<sub>6</sub>OPC, which one would expect to be a powerful detergent because of its rather low CMC in conjunction with extremely low membrane affinity. Indeed, F<sub>6</sub>OPC displays an unusually low  $R_S^{b,SAT}$  value. However, the micelles formed upon further addition of surfactant are virtually devoid of lipid because its transfer from a hydrocarbon into a fluorocarbon environment is very endergonic, thus resulting in an exceptionally high  $R_S^{m,SOL}$  value. Such effects, which are related to nonideal lipid/surfactant interactions rather than intrinsic curvature, must evade any approach based on a single transfer free energy but are readily captured by the present classification, making use of both  $\Delta G_L^{b \rightarrow m,0}$  and  $\Delta G_S^{b \rightarrow m,0}$ .

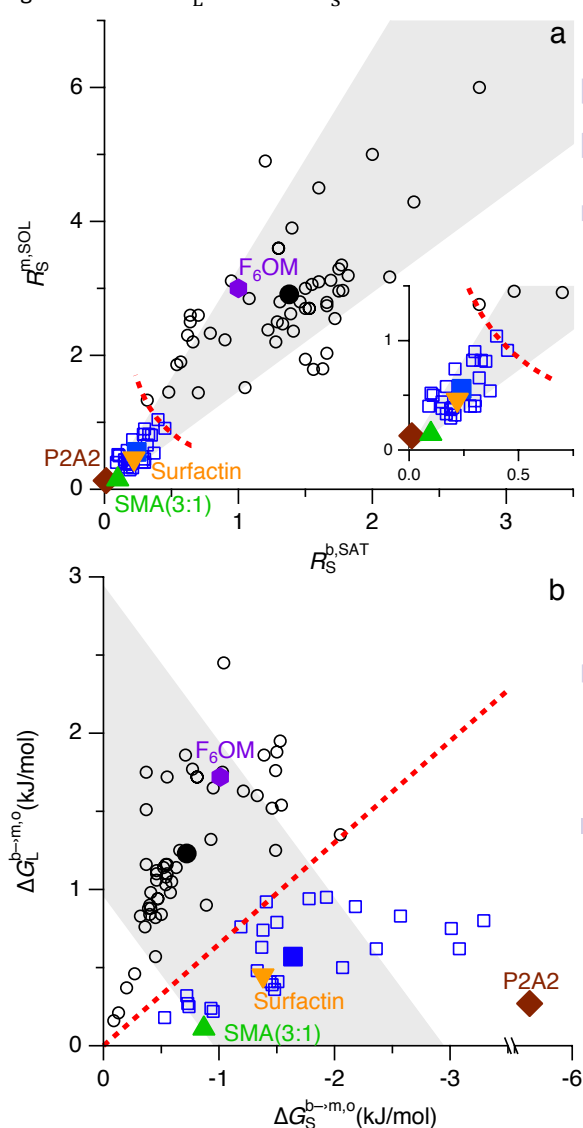


Fig. 6 Thermodynamic classification of surfactants. (a) Surfactant/lipid ratio at the onset and completion of solubilisation,  $R_S^{b,SAT}$  and  $R_S^{m,SOL}$ , respectively. (b) Surfactant and lipid bilayer-to-micelle transfer free energies,  $\Delta G_S^{b \rightarrow m,0}$  and  $\Delta G_L^{b \rightarrow m,0}$ , respectively. Synthetic head-and-tail detergents (open black circles; average: full black circle) on the one hand

and bile salts and their derivatives (open blue squares; average: full blue square), the lipopeptide surfactin (orange down triangle), the dipalmitoylated peptide P2A2 (brown diamond), and SMA(3:1) (green up triangle) on the other hand populate distinct ranges separated by a threshold (red dashed line) defined by  $\Delta G_L^{b \rightarrow m, o} = -0.65 \Delta G_S^{b \rightarrow m, o}$ . F<sub>6</sub>OM (purple hexagon) behaves like a conventional detergent, whereas F<sub>6</sub>OPC (not shown) is a nondetergent with  $R^{sol} > 1000$  and  $\Delta G_L^{b \rightarrow m, o} > 17$  kJ/mol. Also indicated in both panels are the ranges where  $R_S^{m, sol} = (2.37 \pm 0.90) R_S^{b, SAT}$  (gray areas), as obtained from the average and standard deviation for all detergents. Numerical values and references are listed in Tables S1–S3.

## Conclusions

Borrowing from concepts developed for conventional detergents not only furnishes insights into the peculiar properties of self-assembling amphiphilic copolymers and fluorinated surfactants but also highlights quantitative and qualitative differences.

The interactions of SMA(3:1) with lipid membranes follow the three-stage solubilisation scenario observed for many detergents. Polymer/lipid phase diagrams derived with the aid of <sup>31</sup>P NMR spectroscopy pave the way for quantifying the effects of lipid composition and other conditions on solubilisation and define the subsolubilising concentration regime where membrane adsorption of the polymer can be scrutinised at equilibrium. Moreover, these phase diagrams predict the minimum polymer concentration required for complete solubilisation at a given lipid concentration, which is of practical importance for the use of SMA copolymers in membrane-protein research.

The fluorinated nondetergent surfactant F<sub>6</sub>OPC partitions weakly but measurably into POPC bilayers. Once the membrane is saturated, F<sub>6</sub>OPC forms micelles, which, however, do not solubilise lipid and do not compromise membrane integrity. Thus, F<sub>6</sub>OPC qualifies as an amphiphile forming nanoparticles that are compatible with, for example, freestanding lipid bilayers, whereas the related fluorinated surfactant F<sub>6</sub>OM is preferable in situations requiring mild detergent activity. The degree of detergency of new fluorinated surfactants can be tuned by judicious adjustment of headgroup properties and can be quantified with the aid of the ITC and <sup>19</sup>F NMR protocols presented here.

Finally, we proposed a thermodynamic classification of surfactants based on the bilayer-to-micelle transfer free energies of both the lipid and the surfactant of interest. Synthetic head-and-tail detergents, including F<sub>6</sub>OM, partition readily into membranes but form micelles that provide a rather poor environment to phospholipids, thus necessitating relatively high detergent concentrations for complete solubilisation. By contrast, bile salt detergents and their derivatives, natural and designed lipopeptides, as well as SMA(3:1) are more difficult to accommodate in lipid bilayers but fare much better in solubilising phospholipids at low detergent concentrations.

## Experimental

### Materials

POPC was from Lipoid (Ludwigshafen, Germany), F<sub>6</sub>OPC from Anatrace (Maumee, USA), D<sub>2</sub>O from Deutero (Kastellaun, Germany), and NaCl from VWR (Darmstadt, Germany). Sodium trifluoroacetate and 85% H<sub>3</sub>PO<sub>4</sub> in D<sub>2</sub>O were from Sigma-Aldrich (Steinheim, Germany) and 4-(2-hydroxyethyl)-1-piperazineethanesulphonic acid (HEPES) and Tris(hydroxymethyl)aminomethane (Tris) from Carl Roth (Karlsruhe, Germany). SMA(3:1) copolymer solution (trade name Xiran SL25010 S25) was a kind gift from Polyscope (Geleen, Netherlands). All chemicals were obtained in the highest available purity.

### Vesicle preparation

LUVs were prepared by suspension of POPC powder in buffer and 35-fold extrusion through two stacked 100-nm polycarbonate filters with a LiposoFast extruder (Avesco, Ottawa, Canada). Extruded vesicles revealed a unimodal size distribution with a mean diameter and standard deviation of  $(160 \pm 10)$  nm in intensity-weighted size distributions obtained from DLS.

### Preparation of SMA(3:1) stock solutions

The SMA copolymer used in this study had a styrene/maleic acid molar ratio of 3:1, a mass-average molar mass of  $M_w = 10$  kg/mol, and a number-average molar mass of  $M_n = 4$  kg/mol. SMA(3:1) stocks were prepared by dialysis of commercial Xiran SL25010 S25 solutions against buffer (50 mM Tris, 200 mM NaCl, pH 7.4) by use of a 5-mL QuixSep dialyser (Membrane Filtration Products, Seguin, USA) and a Spectra/Por 3 dialysis membrane (Spectrum Laboratories, Rancho Dominguez, USA) with a molar-mass cutoff of 3.5 kg/mol. Dialysis was performed for 24 h at room temperature with gentle stirring and with membrane and buffer exchange after 16 h. Final SMA(3:1) concentrations were determined by refractometry on an Abbatmat 500 (Anton Paar, Graz, Austria) using a molar refractive index increment of  $dn/dc = 1.1178$  M<sup>-1</sup>. This value was obtained from an SMA(3:1) dilution series prepared from a nondialysed Xiran SL25010 S25 solution of known concentration. Molar concentrations were calculated from mass concentrations on the basis of the above number-average molar mass. Samples were aliquoted, stored at  $-20^\circ\text{C}$ , and thawed prior to use.

### <sup>31</sup>P NMR spectroscopy

Stock solutions of 40 mM POPC in the form of LUVs and 17.5 mM SMA(3:1) in buffer (50 mM Tris, 200 mM NaCl, pH 7.4) were used to prepare samples containing 2.5, 5.0, 7.5, or 10.0 mM POPC, 0–5 mM SMA(3:1), and 10% D<sub>2</sub>O for locking. After mixing, all samples were incubated at 25°C for at least 24 h, which was found sufficient for equilibration, as evidenced by the fact that neither NMR nor DLS signals revealed further changes over a period of several weeks. Measurements were performed at 25°C on an Avance 400 spectrometer (Bruker Biospin, Rheinstetten, Germany).

operating at a  $^{31}\text{P}$  resonance frequency of 162 MHz using a 5-mm broadband inverse probe. 256 scans were acquired with an inverse-gated  $^1\text{H}$  decoupling sequence using an acquisition time of 1.6 s, a sweep width of 9746 Hz, and a relaxation delay of 6 s. Data were multiplied by an exponential function with a line-broadening factor of 1.0 Hz before Fourier transformation. Chemical shifts were referenced to 85%  $\text{H}_3\text{PO}_4$  in  $\text{D}_2\text{O}$  as external standard at  $\delta = 0$  ppm.

#### Dynamic light scattering

DLS experiments were carried out on a Zetasizer Nano S90 (Malvern Instruments, Worcestershire, UK) equipped with a 633-nm He–Ne laser and operating at a detection angle of  $90^\circ$ . Samples were thermostatted for 2 min before measurements were carried out at  $25^\circ\text{C}$  in a 45- $\mu\text{L}$  quartz glass cuvette with a cross-section of 3 mm  $\times$  3 mm (Hellma Analytics, Müllheim, Germany). Each sample was measured twice, once with the attenuator position automatically optimised for determination of size distributions and a second time with maximum open attenuator position to ensure comparability of total scattering intensities. The influence of all buffer components on viscosity and refractive index were accounted for during data analysis, which was performed by fitting experimentally determined autocorrelation functions with a non-negatively constrained least-squares function<sup>53</sup> to yield intensity-weighted particle size distributions and by cumulant analysis<sup>54</sup> to obtain z-average particle diameters and associated PDIs.

#### Isothermal titration calorimetry

Demicellisation experiments were done at  $25^\circ\text{C}$  on a VP-ITC (Malvern Instruments) by titrating 30 mM  $\text{F}_6\text{OPC}$  from the injection syringe into the sample cell containing buffer (10 mM HEPES, 150 mM NaCl, pH 7.4). The influence of lipid on the onset of micellisation was assessed by performing the same kind of titration in the presence of 20, 35, or 50 mM POPC in the form of LUVs in both the syringe and the sample cell. Experimental settings included injection volumes of 10  $\mu\text{L}$ , a reference power of 71  $\mu\text{J s}^{-1}$ , a filter period of 2 s, and time spacings of 7 min to allow the signal to reach the baseline before the next injection. Automated baseline adjustment and peak integration were done with NITPIC,<sup>55</sup> and the first injection was always excluded from further analysis.

#### $^{19}\text{F}$ NMR spectroscopy

Stock solutions of 60 mM POPC in the form of LUVs and 20 mM  $\text{F}_6\text{OPC}$  in buffer (10 mM HEPES, 150 mM NaCl, pH 7.4) were used to prepare NMR samples containing 1–6 mM  $\text{F}_6\text{OPC}$  in the absence or presence of 35 mM POPC. All samples contained 10%  $\text{D}_2\text{O}$  for locking. Spectra were recorded at  $25^\circ\text{C}$  on an Avance 400 spectrometer (Bruker Biospin) operating at a  $^{19}\text{F}$  resonance frequency of 376 MHz using a broadband observe probe. 32 scans were acquired with proton decoupling using an acquisition time of 2.2 s, a sweep width of 29762 Hz, and a relaxation delay of 1 s. Data were multiplied by an exponential function with a line-broadening factor of 1.0 Hz before Fourier transformation. Chemical shifts were

referenced to sodium trifluoroacetate in 10%  $\text{D}_2\text{O}$  as external standard at  $\delta = -76.55$  ppm.

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# Nanoparticle self-assembly in mixtures of phospholipids with styrene/maleic acid copolymers or fluorinated surfactants

## Electronic Supplementary Information (ESI)

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## Theoretical Background

### Pseudophases in lipid/surfactant mixtures

The pseudophase concept considers lipid (L) and surfactant (S) molecules in bilayer (b) and micellar (m) phases as well as surfactant monomers in the aqueous (aq) phase.<sup>1</sup> The concentrations of lipid and surfactant,  $c_L$  and  $c_S$ , respectively, together determine the presence and abundance of each of these phases. At given  $c_L$ , increasing  $c_S$  drives a transition from the bilayer range, which contains lipid and surfactant molecules in bilayers as well as surfactant monomers in the aqueous phase, to the coexistence range, where mixed micelles are found in addition to mixed bilayers and surfactant monomers. The first micelles appear at a threshold designated as the saturation (SAT) boundary. A further increase in  $c_S$  results in another transition from the coexistence range to the micellar range containing only micelles and surfactant monomers but no bilayer structures. This transition occurs at the so-called solubilisation (SOL) boundary, where bilayer solubilisation is complete.

Plotting the surfactant concentrations at the SAT and SOL boundaries,  $c_S^{\text{SAT}}$  and  $c_S^{\text{SOL}}$ , respectively, against  $c_L$  gives rise to two straight lines described by:<sup>1,2</sup>

$$c_S^{\text{SAT}} = c_s^{\text{aq},0} + R_S^{\text{b,SAT}} c_L \quad (1)$$

$$c_S^{\text{SOL}} = c_s^{\text{aq},0} + R_S^{\text{m,SOL}} c_L \quad (2)$$

The slopes  $R_S^{\text{b,SAT}}$  and  $R_S^{\text{m,SOL}}$  denote the surfactant/lipid molar ratios in bilayers and micelles at which the membrane becomes saturated with surfactant and at which solubilisation is complete, respectively. Ideally, both lines meet at a common ordinate intercept,  $c_s^{\text{aq},0} = \text{CMC} \times R_S^{\text{m,SOL}} / (1 + R_S^{\text{m,SOL}})$ , which corresponds to the concentration of free surfactant monomers in the aqueous phase in equilibrium with surfactant-saturated bilayers and lipid-saturated micelles within the coexistence range. Note that  $c_s^{\text{aq},0} < \text{CMC}$ , because the latter refers to surfactant monomers in equilibrium with pure surfactant micelles in the absence of lipid.<sup>3,4</sup> The saturating and solubilising mole fractions of surfactant in bilayer and micellar phases,  $X_S^{\text{b,SAT}}$  and  $X_S^{\text{m,SOL}}$ , respectively, thus amount to:

$$X_S^{\text{b,SAT}} = \frac{R_S^{\text{b,SAT}}}{1 + R_S^{\text{b,SAT}}} \quad (3)$$

$$X_S^{\text{m,SOL}} = \frac{R_S^{\text{m,SOL}}}{1 + R_S^{\text{m,SOL}}} \quad (4)$$

The partition coefficients characterising the transfer of lipid and surfactant from the bilayer phase into the micellar phase,  $K_L^{\text{b} \rightarrow \text{m}}$  and  $K_S^{\text{b} \rightarrow \text{m}}$ , then are given by:

$$K_L^{\text{b} \rightarrow \text{m}} \equiv \frac{X_L^{\text{m,SOL}}}{X_L^{\text{b,SAT}}} = \frac{1 - X_S^{\text{m,SOL}}}{1 - X_S^{\text{b,SAT}}} = \frac{1 + R_S^{\text{b,SAT}}}{1 + R_S^{\text{m,SOL}}} < 1 \quad (5)$$

$$K_S^{\text{b} \rightarrow \text{m}} \equiv \frac{X_S^{\text{m,SOL}}}{X_S^{\text{b,SAT}}} = \frac{R_S^{\text{m,SOL}}(1 + R_S^{\text{b,SAT}})}{R_S^{\text{b,SAT}}(1 + R_S^{\text{m,SOL}})} = \frac{R_S^{\text{m,SOL}}}{R_S^{\text{b,SAT}}} K_L^{\text{b} \rightarrow \text{m}} > 1 \quad (6)$$

Note that the above inequalities follow from  $R_S^{\text{m,SOL}} > R_S^{\text{b,SAT}}$ . With the partition coefficients at hand, the corresponding standard molar Gibbs free energies of transfer from bilayers into micelles are obtained as  $\Delta G_L^{\text{b} \rightarrow \text{m},0} = -RT \ln K_L^{\text{b} \rightarrow \text{m}} > 0$  and  $\Delta G_S^{\text{b} \rightarrow \text{m},0} = -RT \ln K_S^{\text{b} \rightarrow \text{m}} < 0$  for lipid and surfactant, respectively.

Conversely, solving the system of eqs 5 and 6 for the saturating and solubilising molar ratios yields:

$$R_S^{b,SAT} = \frac{1 - K_L^{b \rightarrow m}}{K_S^{b \rightarrow m} - 1} \quad (7)$$

$$R_S^{m,SOL} = \frac{K_S^{b \rightarrow m}(1 - K_L^{b \rightarrow m})}{K_L^{b \rightarrow m}(K_S^{b \rightarrow m} - 1)} = \frac{K_S^{b \rightarrow m}}{K_L^{b \rightarrow m}} R_S^{b,SAT} \quad (8)$$

Comparison of eqs 7 and 8 shows that the relative width of the coexistence range is given as:

$$\frac{R_S^{m,SOL}}{R_S^{b,SAT}} = \frac{K_S^{b \rightarrow m}}{K_L^{b \rightarrow m}} = \exp\left(\frac{\Delta G_L^{b \rightarrow m, o} - \Delta G_S^{b \rightarrow m, o}}{RT}\right) \quad (9)$$

In the coexistence range, the concentration of surfactant monomers in the aqueous phase is:

$$c_s^{aq, o} = CMC \frac{R_S^{m,SOL}}{1 + R_S^{m,SOL}} = CMC \frac{K_S^{b \rightarrow m}(1 - K_L^{b \rightarrow m})}{K_S^{b \rightarrow m} - K_L^{b \rightarrow m}} \quad (10)$$

### Membranophobicity

If the surfactant has a pronounced preference for the micellar over the bilayer phase, as expressed by  $K_S^{b \rightarrow m} \gg 1$ , eqs 7 and 8 reduce to:

$$R_S^{b,SAT} \rightarrow \frac{1 - K_L^{b \rightarrow m}}{K_S^{b \rightarrow m}} \ll 1 \quad (11)$$

$$R_S^{m,SOL} \rightarrow \frac{1 - K_L^{b \rightarrow m}}{K_L^{b \rightarrow m}} \quad (12)$$

Thus, even if reliable determination of very low  $R_S^{b,SAT}$  values may be difficult for experimental reasons (Fig. 1b),  $K_L^{b \rightarrow m}$  can still be obtained from  $R_S^{m,SOL}$  alone:

$$K_L^{b \rightarrow m} \approx \frac{1}{1 + R_S^{m,SOL}} \quad (13)$$

### Poor detergency

If the lipid is very reluctant to partition from the bilayer into the micellar phase, as reflected in  $K_L^{b \rightarrow m} \ll 1$ , eqs 7 and 8 simplify to:

$$R_S^{b,SAT} \rightarrow \frac{1}{K_S^{b \rightarrow m} - 1} \quad (14)$$

$$R_S^{m,SOL} \rightarrow \frac{K_S^{b \rightarrow m}}{K_L^{b \rightarrow m}(K_S^{b \rightarrow m} - 1)} \gg 1 \quad (15)$$

Now, it is  $R_S^{m,SOL}$  that is difficult to quantify (Fig. 1c), but  $R_S^{b,SAT}$  still furnishes  $K_S^{b \rightarrow m}$  as:

$$K_S^{b \rightarrow m} \approx \frac{1 + R_S^{b,SAT}}{R_S^{b,SAT}} \quad (16)$$

## Data Analysis

Best-fit parameter values and 95% confidence intervals were derived by nonlinear least-squares fitting in Excel spreadsheets, as detailed elsewhere.<sup>5</sup>

### <sup>31</sup>P NMR spectroscopy

According to the pseudophase model, all lipid molecules reside in bilayer membranes as long as the surfactant concentration is lower than  $c_S^{\text{SAT}}$  (eq 1). In solution-state NMR experiments employing relatively narrow sweep widths, the NMR signal from <sup>31</sup>P nuclei residing in vesicular bilayers is broadened beyond detection.<sup>6–9</sup> Thus, the area under the <sup>31</sup>P NMR peak,  $A$ , vanishes in the absence of solubilised lipid:

$$A(c_S \leq c_S^{\text{SAT}}) = 0 \quad (17)$$

Once the surfactant concentration exceeds  $c_S^{\text{SOL}}$  (eq 2), all lipid molecules are solubilised and produce a sharp, isotropic NMR signal. Hence, the peak area amounts to:

$$A(c_S^{\text{SOL}} \leq c_S) = f c_L \quad (18)$$

where the proportionality factor,  $f$ , depends on experimental conditions but is constant for a given NMR spectrometer operated using identical instrument settings and acquisition parameters. Within the coexistence range, the peak area is proportional to the extent of solubilisation:

$$A(c_S^{\text{SAT}} \leq c_S \leq c_S^{\text{SOL}}) = f c_L \frac{c_S - c_S^{\text{b,SAT}}}{c_S^{\text{m,SOL}} - c_S^{\text{b,SAT}}} \quad (19)$$

Here, the last term on the right-hand side reflects the fraction of solubilised lipid as given by the lever rule.<sup>1,10</sup> Together, eqs 17–19 provide best-fit values of  $c_S^{\text{SAT}}$  and  $c_S^{\text{SOL}}$  at given  $c_L$ .

### Isothermal titration calorimetry

By analogy to calorimetric demicellisation titrations without lipid,<sup>11,12</sup> isotherms obtained by titration of surfactant micelles into buffer in the presence of lipid vesicles can be fitted using a sigmoidal function of the form:<sup>12</sup>

$$Q(c_S) = \frac{(m_1 - m_2)(c_S - c_S^{\text{SAT}}) - \Delta H_S \frac{c_{S,\text{syr}} - c_S^{\text{SAT}}}{c_{S,\text{syr}}}}{1 + \exp\left(\frac{c_S - c_S^{\text{SAT}}}{\Delta c}\right)} + b_2 + m_2 c_S \quad (20)$$

with  $c_S$  denoting the surfactant concentration in the calorimeter cell,  $c_{S,\text{syr}}$  the surfactant concentration in the syringe, and  $c_S^{\text{SAT}}$  the surfactant concentration at which the bilayer is saturated. Here,  $\Delta H_S$  denotes the experimentally determined net change in enthalpy per mole of surfactant, which stems from both (de)micellisation and membrane partitioning, and  $\Delta c$  is a generic parameter modulating the width of the transition region. The coefficients  $m_1$  and  $m_2$  are the slopes of straight pre- and post-transition baselines, respectively, and  $b_2$  is the ordinate intercept of the post-transition baseline.

In the absence of micelles, the water-to-bilayer partition coefficient of the surfactant,  $K_S^{\text{aq} \rightarrow \text{b}}$ , is given as:

$$K_S^{\text{aq} \rightarrow \text{b}} \equiv \frac{X_S^{\text{b}}}{X_S^{\text{aq}}} = \frac{c_S^{\text{b}}/(c_S^{\text{b}} + \gamma c_L)}{c_S^{\text{aq}}/(c_S^{\text{aq}} + c_W)} \approx \frac{c_S^{\text{b}} c_W}{c_S^{\text{aq}} \gamma c_L} \quad (21)$$

with  $c_W = 55.5$  M denoting the molar concentration of water and  $\gamma$  the fraction of lipid that is accessible to the surfactant;  $\gamma = 0.5$  if the surfactant cannot translocate across the bilayer, while  $\gamma = 1.0$  if it has access to both bilayer leaflets on the experimental timescale.<sup>13,14</sup> Thus, the fraction of surfactant in the bilayer phase is:

$$\frac{c_S^b}{c_S^b + c_S^{aq}} = \frac{K_S^{aq \rightarrow b} \gamma c_L}{K_S^{aq \rightarrow b} \gamma c_L + c_W} \quad (22)$$

With this expression, the net enthalpy change retrieved from a nonlinear least-squares fit according to eq 20 can be parsed into contributions from (de)micellisation and bilayer partitioning as:

$$\Delta H_S = \Delta H_S^{aq \rightarrow m} - \frac{K_S^{aq \rightarrow b} \gamma c_L}{K_S^{aq \rightarrow b} \gamma c_L + c_W} \Delta H_S^{aq \rightarrow b} \quad (23)$$

where  $\Delta H_S^{aq \rightarrow m}$  and  $\Delta H_S^{aq \rightarrow b}$  are the molar enthalpy changes accompanying micelle formation and bilayer partitioning, respectively. According to eq 23, a series of demicellisation isotherms recorded at various lipid concentrations provides these two enthalpies as the ordinate intercept and the negative slope, respectively, of a plot of  $\Delta H_S$  versus the fraction of membrane-bound surfactant as given by eq 22.

#### <sup>19</sup>F NMR spectroscopy

In the presence of micelles, the peak chemical shift,  $\delta_p$ , observed by <sup>19</sup>F NMR spectroscopy is a linear combination of the chemical shift of the free monomer in the aqueous phase as measured in the absence of micelles,  $\delta_{mon}$ , and the chemical shift in the micellar state,  $\delta_{mic}$ . The saturating surfactant concentration,  $c_S^{SAT}$ , and  $\delta_{mic}$  were treated as fitting parameters in the equation.<sup>9,15</sup>

$$\delta_p(c_S) = \frac{c_S^{SAT}}{c_S} \delta_{mon} + \left(1 - \frac{c_S^{SAT}}{c_S}\right) \delta_{mic} \quad (24)$$

In the absence of lipid,  $c_S^{SAT}$  is replaced by the CMC in eq 24.

## Literature Data

**Table S1.** Changes in standard molar Gibbs free energy,  $\Delta G^{b \rightarrow m, o}$ , accompanying the transfer of synthetic head-and-tail surfactants (S) and lipids (L) from bilayers (b) into micelles (m).  $\Delta G^{b \rightarrow m, o}$  values were calculated from  $R_S^{b, SAT}$  and  $R_S^{m, SOL}$  with the aid of eqs 3–6.

Surfactant	Lipid	Comment	<i>T</i> (°C)	$R_S^{b, SAT}$	$R_S^{m, SOL}$	$\frac{R_S^{m, SOL}}{R_S^{b, SAT}}$	$\Delta G_S^{b \rightarrow m, o}$ (kJ mol <sup>-1</sup> )	$\Delta G_L^{b \rightarrow m, o}$ (kJ mol <sup>-1</sup> )		Ref.
OG	<i>E. coli</i> lipid		8	0.947	3.11	3.28	-1.03	1.75	-1.69	16
OG	POPC		8	1.22	2.38	1.95	-0.58	0.98	-1.70	16
OG	POPC		25	1.28	2.2	1.72	-0.50	0.84	-1.67	17
OG	EPC		25	1.3	3.6	2.77	-0.81	1.72	-2.13	18
OG	EPC/EPA		25	1.66	2.74	1.65	-0.40	0.84	-2.12	19
OG	SPC		27	1.55	3.06	1.97	-0.54	1.16	-2.16	20
OG	SPC		70	1.72	2.55	1.48	-0.36	0.76	-2.09	20
OG	DMPC		27	1.56	1.79	1.15	-0.13	0.21	-1.67	20
OG	DMPC		70	1.66	2.03	1.22	-0.20	0.37	-1.83	20
OG	DPPC		70	1.5	1.94	1.29	-0.27	0.46	-1.70	20
OG	DSPC		70	1.33	2.47	1.86	-0.63	1.14	-1.80	20
OG	EPC		25	1.3	3.6	2.77	-0.81	1.72	-2.13	21
OG	EPC		28	1.6	3.1	1.94	-0.52	1.14	-2.21	22
OG	POPC		28	1.31	2.8	2.14	-0.66	1.25	-1.90	23
OG	EPC		24	1.5	3	2.00	-0.55	1.16	-2.11	6
OG	EPC	0 M NaCl	25	1.41	2.36	1.67	-0.45	0.82	-1.82	24
OG	EPC	0.02 M NaCl	25	1.75	2.96	1.69	-0.40	0.90	-2.27	24
OG	EPC	0.1 M NaCl	25	1.69	3.12	1.85	-0.46	1.06	-2.28	24
OG	EPC	0.5 M NaCl	25	1.82	3.19	1.75	-0.41	0.98	-2.40	24
OG	EPC	1 M NaCl	25	1.75	3.29	1.88	-0.46	1.10	-2.38	24
OG	EPC	1.5 M NaCl	25	1.77	3.35	1.89	-0.46	1.12	-2.42	24
OG	EPC	0 M sucrose	25	1.66	2.79	1.68	-0.41	0.88	-2.14	24
OG	EPC	0.01 M sucrose	25	1.78	2.97	1.67	-0.39	0.88	-2.29	24
OG	EPC	0.5 M sucrose	25	1.53	2.7	1.76	-0.47	0.94	-2.02	24
OG	EPC	2 M sucrose	25	2.31	4.29	1.86	-0.37	1.16	-3.12	24
OG	EPC	0.01 M urea	25	1.53	2.7	1.76	-0.47	0.94	-2.02	24
OG	EPC	0.5 M urea	25	1.46	2.8	1.92	-0.54	1.08	-2.01	24
OG	EPC	2 M urea	25	1.29	2.5	1.94	-0.59	1.05	-1.79	24
OG	EPC	4 M urea	25	1.63	1.8	1.10	-0.09	0.16	-1.71	24
OG	EPC		25	1.3	3.6	2.77	-0.81	1.72	-2.13	25
NG	EPC		25	1.4	3.9	2.79	-0.77	1.77	-2.30	25
DG	EPC		25	1.6	4.5	2.81	-0.71	1.86	-2.63	25
UG	EPC		25	2	5.0	2.50	-0.55	1.72	-3.11	25
DDG	EPC		25	2.8	6.0	2.14	-0.37	1.51	-4.04	25
NM	POPC		25	0.48	1.45	3.02	-1.49	1.25	-0.84	17
DDM	POPC		25	0.9	2.23	2.48	-0.93	1.32	-1.41	26
DDM	POPC		25	0.7	1.44	2.06	-0.89	0.90	-1.00	17
C <sub>12</sub> EO <sub>5</sub>	POPC		25	3.1	7.3	2.35	-0.37	1.75	-4.66	27
C <sub>12</sub> EO <sub>6</sub>	POPC		25	1.2	4.9	4.08	-1.04	2.45	-2.35	27
C <sub>12</sub> EO <sub>7</sub>	POPC		25	0.7	2.6	3.71	-1.39	1.86	-1.34	27
C <sub>12</sub> EO <sub>8</sub>	POPC		25	0.57	1.9	3.33	-1.46	1.52	-1.04	27
C <sub>12</sub> EO <sub>8</sub>	POPC		11	0.32	1.33	4.20	-2.05	1.35	-0.66	27
C <sub>12</sub> EO <sub>8</sub>	POPC		25	0.54	1.86	3.45	-1.54	1.54	-1.00	27
C <sub>12</sub> EO <sub>8</sub>	POPC		40	0.79	2.33	2.97	-1.21	1.63	-1.34	27
C <sub>12</sub> EO <sub>8</sub>	POPC		50	1.08	2.85	2.63	-0.95	1.65	-1.74	27
C <sub>12</sub> EO <sub>8</sub>	POPC		75	2.13	3.17	1.49	-0.32	0.83	-2.59	27
C <sub>12</sub> EO <sub>8</sub>	EPC		20	0.66	2.2	3.33	-1.33	1.60	-1.20	8
C <sub>12</sub> EO <sub>8</sub>	EPC		25	0.62	2.3	3.71	-1.49	1.76	-1.19	28
LDAO	POPC		25	1.39	2.62	1.88	-0.54	1.03	-1.90	3
SDS	DMPC	0.1 M NaCl	60	1.05	1.52	1.45	-0.45	0.57	-1.26	29
SDS	POPC		65	1.5	2.7	1.80	-0.55	1.10	-2.00	30
Triton	EPC		25	0.64	2.5	3.91	-1.50	1.88	-1.25	18
Triton	EPC		25	0.64	2.6	4.06	-1.53	1.95	-1.28	31
<b>Average</b>				<b>1.38</b>	<b>2.91</b>	<b>2.31</b>	<b>-0.72</b>	<b>1.23</b>	<b>-1.96</b>	
<b>SD</b>				<b>0.55</b>	<b>1.11</b>	<b>0.83</b>	<b>0.44</b>	<b>0.48</b>	<b>0.71</b>	

*Surfactants:* OG, NG, DG, UG, DDG, *n*-octyl-, *n*-nonyl-, *n*-decyl-, *n*-undecyl-, *n*-dodecyl-β-D-glucopyranoside, respectively; NM, DDM, *n*-nonyl-, *n*-dodecyl-β-D-maltopyranoside, respectively; C<sub>12</sub>EO<sub>*n*</sub>, with *n* = 5, 6, 7, 8, penta-, hexa-, hepta-, octa(ethylene oxide) dodecyl ether; LDAO, lauryldimethylamine *N*-oxide; SDS, sodium dodecyl sulphate; Triton, Triton X-100. *Lipids:* DMPC, 1,2-dimyristoyl-*sn*-glycero-3-phosphocholine; DPPC, 1,2-dipalmitoyl-*sn*-glycero-3-phosphocholine; DSPC, 1,2-distearoyl-*sn*-glycero-3-phosphocholine; *E. coli* lipid, *E. coli* polar lipid extract (67.0% phosphatidylethanolamine, 23.2% phosphatidylglycerol, 9.8% diphosphatidylglycerol); EPA, egg phosphatidic acid; EPC, egg phosphatidylcholine; POPC, 1-palmitoyl-2-oleoyl-*sn*-glycero-3-phosphocholine; SPC, soybean phosphatidylcholine.



**Table S2.** Changes in standard molar Gibbs free energy,  $\Delta G^{b \rightarrow m, o}$ , accompanying the transfer of sterol-based surfactants (S) and lipids (L) from bilayers (b) into micelles (m).  $\Delta G^{b \rightarrow m, o}$  values were calculated from  $R_S^{b, SAT}$  and  $R_S^{m, SOL}$  with the aid of eqs 3–6.

Surfactant	Lipid	Comment	T (°C)	$R_S^{b, SAT}$	$R_S^{m, SOL}$	$\frac{R_S^{m, SOL}}{R_S^{b, SAT}}$	$\Delta G_S^{b \rightarrow m, o}$ (kJ mol <sup>-1</sup> )	$\Delta G_L^{b \rightarrow m, o}$ (kJ mol <sup>-1</sup> )		Ref.
CHAPS	POPC		25	0.11	0.5	4.55	-3.01	0.75	-0.25	17
CHAPS	POPC		25	0.09	0.4	4.44	-3.08	0.62	-0.20	1
CHAPS	EPC		25	0.1*	0.52	5.20	-3.29	0.80	-0.24	32
CHAPS	EPC		20	0.4	1.04	2.60	-1.41	0.92	-0.65	33
CHAPSO	EPC		20	0.21	0.74	3.52	-2.18	0.89	-0.41	33
NaC	EPC		25	0.3	0.4	1.33	-0.53	0.18	-0.35	34
NaC	EPC		25	0.3	0.9	3.00	-1.78	0.94	-0.53	18
NaC	POPC		25	0.3	0.45	1.50	-0.73	0.27	-0.37	35
NaC	POPC		60	0.37	0.54	1.46	-0.72	0.32	-0.45	35
NaC	SPC		30	0.33	0.82	2.48	-1.50	0.79	-0.53	35
NaC	SPC		60	0.45	0.91	2.02	-1.19	0.76	-0.64	35
NaC	DPPC		60	0.19	0.29	1.53	-0.95	0.22	-0.24	36
NaC	DPPC	0.1 M NaCl	60	0.21	0.32	1.52	-0.93	0.24	-0.26	37
NaC	DPPG	0.1 M NaCl	60	0.28	0.4	1.43	-0.74	0.25	-0.34	37
NaC	DPPC/PG 3:1	0.1 M NaCl	60	0.15	0.38	2.53	-2.07	0.50	-0.24	37
NaC	DPPC/PG 1:1	0.1 M NaCl	60	0.19	0.37	1.95	-1.46	0.39	-0.27	37
NaDC	DPPC	0.1 M NaCl	60	0.19	0.38	2.00	-1.51	0.41	-0.27	37
NaDC	DPPG	0.1 M NaCl	60	0.17	0.33	1.94	-1.48	0.36	-0.24	37
NaDC	DPPC/PG 3:1	0.1 M NaCl	60	0.17	0.58	3.41	-2.57	0.83	-0.32	37
NaDC	DPPC/PG 1:1	0.1 M NaCl	60	0.15	0.44	2.93	-2.36	0.62	-0.26	37
NaDC	POPC		25	0.25	0.52	2.08	-1.33	0.48	-0.36	35
NaDC	POPC		60	0.32	0.66	2.06	-1.37	0.63	-0.46	35
NaDC	SPC		30	0.35	0.81	2.31	-1.38	0.74	-0.54	35
NaDC	SPC		60	0.29	0.82	2.83	-1.93	0.95	-0.50	35
NaDC	DPPC		60	0.2	0.39	1.95	-1.44	0.41	-0.28	36
<b>Average</b>				<b>0.24</b>	<b>0.56</b>	<b>2.50</b>	<b>-1.64</b>	<b>0.57</b>	<b>-0.37</b>	
<b>SD</b>				<b>0.10</b>	<b>0.22</b>	<b>1.04</b>	<b>0.76</b>	<b>0.25</b>	<b>0.13</b>	

*Surfactants:* CHAPS, 3-[(3-cholamidopropyl)dimethylammonio]-1-propanesulphonate; CHAPSO, 3-[(3-cholamidopropyl)dimethylammonio]-2-hydroxy-1-propanesulphonate; NaC, sodium cholate; NaDC, sodium deoxycholate. *Lipids:* DPPC, 1,2-dipalmitoyl-*sn*-glycero-3-phosphocholine; DPPG, 1,2-dipalmitoyl-*sn*-glycero-3-[phospho-*rac*-(1-glycerol)]; EPC, egg phosphatidylcholine; POPC, 1-palmitoyl-2-oleoyl-*sn*-glycero-3-phosphocholine; SPC, soybean phosphatidylcholine. Included in the table are only datasets for which the saturation and solubilisation boundaries (eqs 1 and 2, respectively) have identical or, at least, similar ordinate intercepts,  $c_s^{aq, o}$ . This is usually not the case for many of the above lipid/surfactant systems at low ionic strength,<sup>35–37</sup> necessitating more sophisticated thermodynamic analysis.<sup>38</sup> \*See ref. 17 for a discussion of this value.

**Table S3.** Changes in standard molar Gibbs free energy,  $\Delta G^{b \rightarrow m, o}$ , accompanying the transfer of unconventional surfactants (S) and lipids (L) from bilayers (b) into micelles (m).  $\Delta G^{b \rightarrow m, o}$  values were calculated from  $R_S^{b, SAT}$  and  $R_S^{m, SOL}$  with the aid of eqs 3–6.

Surfactant	Lipid	Comment	T (°C)	$R_S^{b, SAT}$	$R_S^{m, SOL}$	$\frac{R_S^{m, SOL}}{R_S^{b, SAT}}$	$\Delta G_S^{b \rightarrow m, o}$ (kJ mol <sup>-1</sup> )	$\Delta G_L^{b \rightarrow m, o}$ (kJ mol <sup>-1</sup> )		Ref.
Surfactin	POPC	lipopeptide	25	0.22	0.46	2.09	-1.38	0.45	-0.32	39
P2A2	POPC	lipopeptide	25	0.012	0.13	10.8	-5.63	0.27	-0.05	40
SMA(3:1)	POPC	copolymer	25	0.099	0.147	1.49	-0.87	0.11	-0.05	*
F <sub>6</sub> OM	POPC	fluorinated	25	1.0	3.0	3.00	-1.01	1.72	-1.71	9
F <sub>6</sub> OPC	POPC	fluorinated	25	0.031	>1000	>30000	-8.68	>17	<-1.96	*

*Surfactants:* P2A2, dipalmitoylated tandem peptide derived from apolipoprotein E; SMA(3:1), styrene/maleic acid copolymer with styrene/maleic acid molar ratio of 3:1; F<sub>6</sub>OM, 3,3,4,4,5,5,6,6,7,7,8,8,8-tridecafluoro-*n*-octyl-β-D-maltopyranoside; F<sub>6</sub>OPC, 3,3,4,4,5,5,6,6,7,7,8,8,8-tridecafluoro-*n*-octylphosphocholine. *Lipid:* POPC, 1-palmitoyl-2-oleoyl-*sn*-glycero-3-phosphocholine. Colour code is matched to Fig. 6 in main text. \*This work.

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