Modeling the Dynamics of Phospholipids in the Fluid Phase of

Liposomes

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ELECTRONIC SUPPLEMENTARY INFORMATION

Neutron Spin Echo Spectroscopy Basics

Neutron Spin Echo (NSE) spectroscopy detects the sum of coherent and incoherent scattering.

Typically, the coherent part dominates which leads to^{S1}

$$\frac{S(Q,t)}{S(Q)} = \frac{\sigma_{\rm coh}S(Q,t)_{\rm coh} - (1/3)\sigma_{\rm inc}S(Q,t)_{\rm inc}}{\sigma_{\rm coh}S(Q)_{\rm coh} - (1/3)\sigma_{\rm inc}S(Q)_{\rm inc}} \approx \frac{S(Q,t)_{\rm coh}}{S(Q)_{\rm coh}}$$
(S1)

where σ_{coh} and σ_{inc} are the coherent and incoherent scattering cross-sections, respectively. The coherent and the incoherent intermediate scattering functions are represented by $S(Q,t)_{coh}$ and $S(Q,t)_{inc}$, respectively.

The time dependent mean-squared displacement of randomly diffusing, infinitely small and massless particles can be written as $(\langle (r_i(0) - r_i(t))^2 \rangle = 6 D t)$.^{S2} Assuming validity of the Gaussian approximation in the calculation of the time dependent incoherent dynamic structure factor, S(Q, t)^{tr}_{inc}, we arrive at the well-known expression^{S2}

$$S(Q, t)_{\rm inc}^{\rm tr} = N \exp(-Q^2 D_{\rm eff} t)$$
(S2)

Equation S2 implicitly includes *N* particles randomly diffusion independently of each other.

In a next step, we exploit the fact that the pair-correlation function can be separated into inter- and intermolecular contributions. In case of dilute solutions which implies uncorrelated motion, the intermolecular contributions can be neglected.^{S1,S2} Since we assume non-interacting, infinitely small and massless particles the intramolecular interactions can be ignored.^{S1,S2}. In this or in the

more general case, of objects with finite dimensions, center of mass diffusion and a separation ansatz, e.g., to separate rotational motion, can be utilized to derive the coherent dynamic structure factor, $S(Q, t)_{coh}^{tr}$,

$$S(Q, t)_{\rm coh}^{\rm tr} \propto N \exp(-Q^2 D_{\rm eff} t)$$
 (S3)

Eqs. S2 and S3 illustrate the more general principle that the coherent dynamic structure factor contains pair- and self-correlation contributions. In those cases, when objects move independently the self-correlation part in $S(Q, t)_{coh}$ has the same form as in the respective part in $S(Q, t)_{inc}$.



Different Illustrations of the Results

Figure S1: Log-log representations of the normalized intermediate scattering function, S(Q,t)/S(Q), as a function of Fourier time, t, for different Q's, for, (a,d) 5 % lipid mass fraction of protonated DOPC at 20 °C, (b,e) 5 % lipid mass fraction of protonated DMPC at 37 °C and (c,f) the 5 % lipid mass fraction of protonated Soy-PC sample at 30 °C, all in D₂O. The same data sets are analyzed by fits using the (a-c) Zilman-Granek model (ZG) (equation 5) and (d-f) the full model that starts from equation 3 and includes diffusion of liposomes and confined motion of lipid tails (equation 10). The error bars representing one standard deviation.



Figure S2: The $\mathcal{A}(Q)$ for h-DMPC obtained from NSE and QENS studies, over a broad Q-range. The data

is modelled using $\mathcal{A}(Q)$ for a particle diffusion in a cylinder.

References:

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