Molecular areas of phospholipids as determined by ²H NMR spectroscopy

Comparison of phosphatidylethanolamines and phosphatidylcholines

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ABSTRACT The role of lipid diversity in biomembranes is one of the major unsolved problems in biochemistry. One parameter of possible importance is the mean cross-sectional area occupied per lipid molecule, which may be related to formation of nonbilayer structures and membrane protein function. We have used 2H NMR spectroscopy to compare the properties of 1,2-diperdeuteriopalmitoyl-sn-glycero-3-phosphocholine (DPPC- d_{c2}) and 1,2-diperdeuteriopalmitoyl-sn-glycero-3-phosphocholine (DPPC- d_{c2}) in the L_{α} phase. We find that DPPE has greater segmental order than DPPC, and that this increase in order is related to the smaller area per acyl chain found for DPPE. Values of the mean cross-sectional chain area are calculated using a simple diamond lattice model for the acyl chain configurational statistics, together with dilatometry data. The results obtained for the mean area per molecule are comparable with those from low angle x-ray diffraction studies.

INTRODUCTION

The significance of lipid diversity in biological membranes still remains obscure. A greater understanding of the differences between the principal lipid constituents in eukaryotic systems, such as phosphatidylcholine and phosphatidylethanolamine, could help resolve this problem. One excellent technique to study phospholipids is deuterium nuclear magnetic resonance (NMR)¹ spectroscopy, which has been used widely to study the average configurational properties and molecular dynamics of phospholipid systems (Seelig, 1977; Seelig and Seelig, 1980; Davis, 1983; Salmon et al., 1987; Dodd and Brown, 1989; Brown et al., 1990; Brown and Söderman, 1990). The bulk of the work has been directed toward phosphatidylcholines; however, a few groups have applied ²H NMR spectroscopy to investigate phosphatidylethanolamines. Seelig and Gally (1976) studied the polar head group of 1,2-dipalmitoyl-sn-glycero-3-phosphoethanola-

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¹Abbreviations used in this paper: DLPE, 1,2-dilauryl-sn-glycero-3-phosphoethanolamine; DMPC-d₂₇, 1-myristoyl-2-perdeuteriomyristoyl-sn-glycero-3-phosphoethanolamine; DMPE-d₂₇, 1-myristoyl-2-perdeuteriomyristoyl-sn-glycero-3-phosphoethanolamine; DPPC-d₂₇, 1-myristoyl-2-perdeuteriomyristoyl-sn-glycero-3-phosphoethanolamine; DPPC-d₆₂, 1,2-diperdeuteriopalmitoyl-sn-glycero-3-phosphoetholine; DPPE, 1,2-dipalmitoyl-sn-glycero-3-phosphoethanolamine; DPPE-d₆₂, 1,2-diperdeuteriopalmitoyl-sn-glycero-3-phosphoethanolamine; EDTA, ethylenediaminetetraacetic acid; egg PE, egg yolk phosphatidylethanolamine; ²H NMR, deuterium nuclear magnetic resonance; H_{III}, inverted hexagonal phase; L₆, liquid-crystalline phase; MOPS, morpholinopropane sulfonic acid; S⁽¹⁾_{CD}, carbon-deuterium bond order parameter of ith segment; TLC, thin layer chromatography; T_m, main (gel to liquid-crystalline) phase transition temperature; T_{red}, reduced temperature.

mine (DPPE) using ²H NMR and obtained evidence suggesting that the phosphoethanolamine moiety undergoes rapid transitions between two enantiomeric conformations, each of which lies approximately flat on the surface of the bilayer. Blume et al. (1982a,b), in a series of papers, showed for DPPE deuterated specifically in three acyl positions that the chains are approximately all-trans in the low temperature ordered state with the molecules rotating about their long axes. Finally, Marsh et al. (1983) studied the phase behavior and configurational order of sn-2 chain perdeuterated 1,2-dimyristoylsn-glycero-3-phosphoethanolamine (DMPE-d₂₇) in the L, phase, concluding that the order parameter plateau of the phosphatidylethanolamine was higher compared with the corresponding phosphatidylcholine (DMPCd₂₇).

²H NMR spectroscopy has also been used to estimate the effective acyl chain lengths and average chain cross-sectional areas of phospholipids in the liquidcrystalline state in terms of simple statistical models (Seelig and Seelig, 1974; Schindler and Seelig, 1975; Salmon et al., 1987; DeYoung and Dill, 1988; Ipsen et al., 1990). The cross-sectional chain area is related to the mean area occupied by a lipid molecule at the membrane lipid-water interface. The area per molecule in turn is important because the effective or average shape of the molecules is associated with the geometry of lipid aggregates (Israelachvili et al., 1976; Tanford, 1980). The latter reflect various contributions to the free energies of lipid systems which may play a role in regulating membrane protein function (Kirk et al., 1984; Wieslander et al., 1986; Wiedmann et al., 1988). In this paper we use ²H NMR spectroscopy of acyl chain perdeuterated DPPE and DPPC along with de-Pakeing procedures (cf. Bloom et al., 1981; Sternin et al., 1983) to compare the orientational order of the two systems. We relate the differences in order to differences in average chain cross-sectional area. It is found that the acyl chains of DPPE-d₆₂ have greater orientational order than for DPPC-d₆₂ at the same absolute and reduced temperatures. This difference in order is reflected in a smaller average chain cross-sectional area for DPPE-d₆₂ compared with DPPC-d₆₂ in the liquid-crystalline state.

MATERIALS AND METHODS

Palmitic acid was obtained from Sigma Chemical Co. (St. Louis, MO) and was perdeuterated by passing deuterium gas over it in the presence of a 10% palladium on charcoal catalyst (Aldrich Chemical Co., Milwaukee, WI) as described (Salmon et al., 1987). The 1,2diperdeuteriopalmitoyl-sn-glycero-3-phosphocholine (DPPC-d₆₂) was synthesized by acylating the cadmium chloride adduct of sn-glycero-3phosphocholine with the anhydride of palmitic acid-d₃₁ (Jensen and Pitas, 1976; Mason et al., 1981). The 1,2-diperdeuteriopalmitoyl-snglycero-3-phosphoethanolamine (DPPE-d₆₂) was synthesized by transphosphatidylation of DPPC-d₆₂ using phospholipase D (from Savoy cabbage) in the presence of ethanolamine (Yang et al., 1967; Comfurius and Zwaal, 1977). The DPPE-d₆₂ was purified using silica gel chromatography and yielded a single spot upon TLC analysis. The sample was dried under high vacuum, placed in a 8-mm test tube, and mixed with 50 wt% buffer containing 20 mM MOPS, 1 mM EDTA, in deuterium depleted water, pH = 7.1. It was then vortexed lightly (<1min) and freeze-thawed five times to ensure homogeneity. 2H NMR spectroscopy of the sample as a function of temperature showed a narrow phase transition ($T_m = 55$ °C, range 53–56°C) which indicated relatively high purity.

The ²H NMR spectra were collected using a home-built high-power probe with a horizontal solenoid inductor. The spectrometer used was a General Electric (Fremont, CA) GN-300 spectrometer operating with a magnetic field strength of 7.05 tesla (²H frequency of 46.1 MHz). A phase-cycled quadrupolar echo pulse sequence was employed with a 1.8 µs 90° pulse, recycle time of 0.5 s, and a digitization dwell time of 2 us (Davis et al., 1976; Bloom et al., 1980). Both quadrature channels were used, taking care to initiate the Fourier transformation at the maximum of the quadrupolar echo. The data were transferred to a Digital Equipment Corporation Microvax II computer for off-line processing. All spectra were numerically deconvoluted, that is de-Paked, to obtain subspectra corresponding to the $\theta = 0^{\circ}$ orientation of the bilayer normal relative to the main magnetic field (Bloom et al., 1981; Sternin et al., 1983). The C-2H bond segmental order parameters, $S_{CD}^{(i)}$, of the de-Paked ²H NMR spectra were calculated from the quadrupolar splittings using the relation

$$\Delta \nu_{\rm Q} = \frac{3}{2} \left(\frac{e^2 q Q}{h} \right) P_2 \left(\cos \theta \right) \left| S_{\rm CD}^{(i)} \right|, \tag{1}$$

where $P_2(\cos \theta) = 1$ and $(e^2qQ/h) = 170$ kHz. In Eq. 1 the C-²H bond segmental order parameters are defined by

$$S_{CD}^{(i)} \equiv \frac{1}{2} \langle 3 \cos^2 \beta_i - 1 \rangle, \tag{2}$$

in which β_i is the angle between the $C^{-2}H$ bond direction at any instant, and the brackets indicate an average over all conformations sampled

on the ²H NMR time scale. Based on geometrical considerations the values of $S_{CD}^{(i)}$ are assumed to be negative.

The segmental order parameters can be used to estimate the average length of the acyl chain $\langle L \rangle$ relative to the all-trans state using a simple statistical model developed by Schindler and Seelig (1975) and later modified by Salmon et al. (1987) to yield

$$\langle L \rangle = l_0 \left[\left(\frac{n - m + 1}{2} \right) - \sum_{i=m}^{n-1} S_{CD}^{(i)} - 3S_{CD}^{(n)} \right].$$
 (3)

Eq. 3 takes into account the fact that different order profiles are observed for the sn-1 and sn-2 chains of phospholipid lamellar phases in the liquid-crystalline state (Seelig and Seelig, 1974; Salmon et al., 1987). Here the index i refers to the numbering of the acyl chains beginning with the ester carbon (i = 1) and ending with the methyl carbon (i = n), where n is the number of carbon atoms (n = 16 for DPPC and DPPE). The effective acyl chain length is taken as extending from the C_1 carbon to the methyl carbon in the case of the sn-1 chain, i.e., m = 2. In the case of the sn-2 chain, the contribution from the C_2 carbon is neglected so that m = 3 (cf. Salmon et al., 1987). The length of one carbon-carbon bond projected onto the long axis of the all-trans reference state is $l_0 = 1.25$ Å. From the effective chain length $\langle L \rangle$ an estimate can be made of the average cross-sectional area of the acyl chain $\langle A \rangle$ by the equation

$$\langle A \rangle = V_{\text{chain}}/\langle L \rangle$$
 (4)

Here $V_{\rm chain}$ is the volume of the acyl chain derived from the volume per methylene for each lipid (Nagle and Wilkinson, 1978; Wilkinson and Nagle, 1981; Salmon et al., 1987; Nagle and Wiener, 1988), and is given by

$$V_{\text{chain}} = n' V_{\text{CH}_1} + V_{\text{CH}_2}. \tag{5}$$

In the above expression $V_{\rm CH_3} \cong 2V_{\rm CH_2}$ and n' = n - m is the number of methylene segments used to calculate the projected chain length (n' = 14 and 13 for the sn-1 and sn-2 chains, respectively, of DPPC and DPPE). A value of $V_{\rm CH_2} = 28.0 \text{ Å}^2$ was used in the calculations (cf. Nagle and Wilkinson, 1978).

RESULTS AND DISCUSSION

A representative 2H NMR spectrum of an aqueous dispersion of DPPE- d_{62} in the L_{α} phase is shown in Fig. 1 a, along with the corresponding de-Paked spectrum in Fig. 1 b. The powder-type 2H NMR spectrum (Fig. 1 a) consists of a series of overlapping powder patterns due to the various deuterated chain segments and is very similar to 2H NMR spectra for DPPC- d_{62} (Davis, 1979; Salmon et al., 1987). The de-Paked spectrum (Fig. 1 b) corresponds to the $\theta=0^\circ$ orientation of the bilayer normal with respect to the magnetic field. The improvement in the resolution is due to the fact that the subspectrum is spread out to the maximum extent possible which allows for easier assignment of the C- 2H bond order parameters, $S_{CD}^{(i)}$.

The values of $S_{CD}^{(i)}$ extracted from the de-Paked ²H NMR spectra are shown for DPPE-d₆₂ at two temperatures in Fig. 2 a. The chain segment assignments were

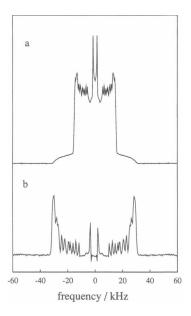


FIGURE 1 Representative ²H NMR spectra of multilamellar dispersion of DPPE-d₆₂ at 69°C ($T_{\rm red} = 0.0419$) in the liquid-crystalline ($L_{\rm e}$) phase. The ²H NMR powder-type spectrum corresponding to a random distribution of the lamellae is indicated in a. The de-Paked ²H NMR spectrum in b represents the $\theta = 0^{\circ}$ orientation of the lamellar normal relative to the magnetic field.

made by integrating the intensities of the de-Paked spectra and by comparison to previous results for DPPC with specifically deuterated and perdeuterated acyl chains (Seelig and Seelig, 1974, 1975; Brown et al., 1979; Blume et al., 1982a; Salmon et al., 1987). Because of the different orientations with respect to the bilayer surface, the two acyl chains of DPPC are inequivalent in the L phase, leading to separate profiles for the sn-1 and sn-2 chains (Seelig and Seelig, 1975). In the case of DPPC-d_s, in the L_{α} phase, the quadrupolar splittings arising from the sn-1 and sn-2 chains were assigned by comparison to results for DPPC-d₃₁, in which the sn-2 chain was perdeuterated (Salmon et al., 1987). An identical pattern of splittings was evident for DPPE-d₆₂ (cf. Fig. 1), which were assigned by analogy to the results for DPPC-d₆₂. The order profiles of both DPPE-d₆₂ and DPPC-d₆₂ show a plateau region where the order parameters for the carbons closest to the water interface are approximately equal (Fig. 2). Beyond the plateau, the order decreases as one moves down the chain. The shape of the profile for DPPE-d₆₂ is the same as seen for DPPC- d_{62} (Fig. 2b), except that the magnitudes of the order parameters are greater at the same absolute temperatures. Because DPPE-d₆₂ has a higher orderdisorder phase transition temperature (55°C vs. 38°C for DPPC-d₆₂) (this work; Trouard et al., 1989) it is helpful to compare the order on the same temperature scale.

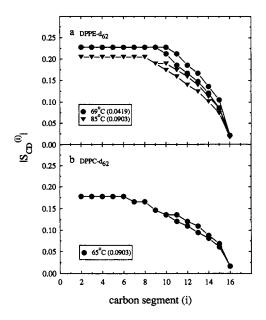


FIGURE 2 Comparison of order profiles of DPPE- d_{62} and DPPC- d_{62} at different absolute and reduced temperatures in the L_a phase. The order parameters $S_{CD}^{(1)}$ are plotted as a function of chain position for (a) DPPE- d_{62} at 69 and 85°C ($T_{\rm red}=0.0419$ and 0.0903) and (b) DPPC- d_{62} at 65°C ($T_{\rm red}=0.0903$). Different order profiles are obtained for the two acyl chains and are connected by solid lines. The smaller and larger values of $S_{CD}^{(1)}$ arise from the sn-1 and sn-2 chains, respectively. Note that the segmental ordering is greater for DPPE- d_{62} at all absolute and reduced temperatures compared to DPPC- d_{62} .

For this the reduced temperature is used where $T_{\text{red}} \equiv (T - T_{\text{m}})/T_{\text{m}}$ and T_{m} is the main transition temperature (Seelig and Browning, 1978; Barry et al., 1990). At the same reduced temperatures the order is also *greater* for DPPE-d₆₂ than for DPPC-d₆₂.

The different order profiles of these two systems are related to differences in the polar headgroup regions because the acyl chains are identical. The most obvious difference is the headgroup size and x-ray diffraction data indicate that phosphatidylethanolamines have a smaller surface area per molecule relative to phosphatidylcholines (Lis et al., 1982; Wilkinson and Nagle, 1981; McIntosh and Simon, 1986; Nagle and Wiener, 1988). Mely et al. (1975) have shown that order profiles for potassium laurate extracted from ²H NMR spectra are related to the mean area per polar headgroup. The theoretical model of Meraldi and Schlitter (1981) stresses the importance of short-range repulsive forces, which are related to the size and shape of the molecules, in governing the orientational order observed by ²H NMR in lipid systems. Therefore the smaller surface area per molecule for DPPE-d₆₂ is most likely associated with the larger order parameters compared with DPPC-d₆₂; cf. also Marsh et al. (1983). The smaller cross-sectional area available to the chains indicates that fewer gauche isomers are needed to maintain the packing near the density of liquid hydrocarbons. This same conclusion was reached by Marsh et al. (1983) when comparing DMPC-d₂₇ and DMPE-d₂₇ at the same reduced temperatures, except they found that only the plateau carbons exhibited an increase in order for DMPE-d₂₇. The difference between what we observe and the observations by Marsh et al. (1983) is due to the increased resolution obtained by de-Paking the spectra.

As described above, a simple diamond lattice model can be used to relate the profiles of the $S_{CD}^{(i)}$ values to the average projected length $\langle L \rangle$ and the mean crosssectional area $\langle A \rangle$ of the acyl chains. For DPPE-d₆₂ at 69°C, the calculated length $\langle L \rangle$ is 12.9 Å for the sn-1 chain and 12.1 Å for the sn-2 chain. When the temperature is increased to 85°C, $\langle L \rangle$ decreases to 12.4 and 11.7 Å for the sn-1 and sn-2 chains, respectively. In Fig. 3 the calculated cross-sectional areas are compared for DPPEd₆₂ and DPPC-d₆₂. As expected, the calculated average areas of the acyl chains are significantly less for DPPEd₆₂ at all absolute and reduced temperatures compared with DPPC-d₆₂. Because the only difference in the two molecules is in the headgroup region, the area occupied per acyl chain must reflect the effective headgroup area at the membrane lipid-water interface. If one adds the areas of the sn-1 and sn-2 chains an estimate of the average cross-sectional area per molecule can be obtained. This technique yields a value of 71.7 Å² for DPPC-d₆₂ at 50°C ($T_{red} \approx 0.04$) compared with 57.6–70.9 Å² from x-ray data (Nagle and Wiener, 1988). The wide range of values from low angle x-ray diffraction studies underscores the ambiguities present when using this method in the L_a phase (Nagle and Wiener, 1988). A value of $\langle A \rangle = 69.5 \,\text{Å}^2$ is obtained for the interfacial area

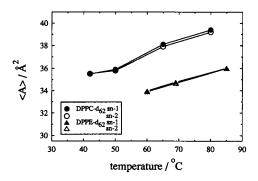


FIGURE 3 Average cross-sectional area $\langle A \rangle$ as a function of temperature for DPPE-d₆₂ (\blacktriangle , \triangle) and DPPC-d₆₂ (\spadesuit , \bigcirc) in the L_a phase. The average area per chain is calculated from the order parameters using a diamond lattice model; cf. text. The cross-sectional areas of the sn-1 and sn-2 chains are smaller for DPPE-d₆₂ than for DPPC-d₆₂ at all temperatures.

occupied per molecule of DPPE- d_{62} at 69°C ($T_{\rm red} \cong 0.04$) in the L_{α} phase, which is less than for DPPC- d_{62} . Although x-ray data are currently unavailable for DPPE, the smaller surface area per molecule relative to phosphatidylcholines is in qualitative agreement with lowangle x-ray diffraction studies of egg PE, DLPE, and DMPE in the L_{α} phase (Lis et al., 1982; Wilkinson and Nagle, 1981; McIntosh and Simon, 1986; Nagle and Wiener, 1988). A reduction of ~4 Ų in the average chain cross-sectional area of DPPE- d_{62} relative to DPPC- d_{62} is evident at the same absolute temperature (cf. Fig. 3), from which a difference of ~8 Ų in the cross-sectional area per molecule (a 10% decrease) is suggested.

The above approach may perhaps overestimate the area per molecule because the values obtained for DPPC-d₆₂ are at the high end of the x-ray data range. An alternate method assumes that only the carbons nearest to the headgroup reflect the molecular area at the lipid/water interface. This would yield lower values for the mean cross-sectional area per molecule. Such an approach may be reasonable because the motions of carbon segments further down the chain are less restricted by the headgroup. On the other hand this assumption could lead to packing problems when one gets to the center of the bilayer. If we use only carbons two through ten (approximately the plateau region) in calculating the area per molecule, the values decrease to 57.4 Å² for DPPC-d₆₂ at 50°C and 55.4 Å² for DPPE-d₆₂ at 69°C. These values still reflect the smaller area per molecule for DPPE-d₆₂ relative to DPPC-d₆₂, and the differences in molecular area for these two lipids are the same as for the previous method ($\sim 10\%$ decrease).

In summary, the differences in configurational properties of the acyl chains of DPPC-d₆₂ and DPPE-d₆₂ manifest differences in the polar headgroup region of the bilayers. These differences cause an increase in the main transition temperature for DPPE-d₆₂ as well as an increase in the segmental order at the same absolute and reduced temperatures compared with DPPC-d₆₉. The ²H NMR data indicate that the average area per acyl chain is significantly smaller for DPPE-d₆₂ than for DPPC-d₆₂. This difference in the chain cross-sectional area is consistent with the molecular areas determined by x-ray studies. 2H NMR thus provides an additional method for comparing the molecular areas of different phospholipids in the L_a phase. The area available for a lipid molecule in membranes is of fundamental importance in understanding their roles in biological systems. This parameter has been related to such phenomena as the formation of non-bilayer phases (including the inverted hexagonal, H_{II}, and cubic phases), the role of lipid diversity in biological systems, lipid-protein interactions, and diffusion/permeability in membranes (Israelachvili

et al., 1976; Kirk et al., 1984; Wiedmann et al., 1988; De Young and Dill, 1988; Lindblom and Rilfors, 1989).

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REFERENCES

- Barry, J. A., T. P. Trouard, A. Salmon, and M. F. Brown. 1990. ²H NMR spectroscopy of the low temperature behavior of bilayers containing docosahexaenoyl phospholipids. *Biophys. J.* 57:273a. (Abstr.)
- Bloom, M., J. H. Davis, and M. I. Valic. 1980. Spectral distortion effects due to finite pulse widths in deuterium NMR. Can. J. Phys. 58:1510-1517.
- Bloom, M., J. H. Davis, and A. L. MacKay. 1981. Direct determination of the oriented sample NMR spectrum from the powder spectrum for systems with local axial symmetry. *Chem. Phys. Lett.* 80:198–202.
- Blume, A., D. M. Rice, R. J. Wittebort, and R. G. Griffin. 1982a. Molecular dynamics and conformation in the gel and liquidcrystalline phases of phosphatidylethanolamine bilayers. *Biochemis*try. 21:6220–6230.
- Blume, A., R. J. Wittebort, S. K. Das Gupta, and R. G. Griffin. 1982b.
 Phase equilibria, molecular conformation, and dynamics in phosphatidylcholine/phosphatidylethanolamine bilayers. *Biochemistry*. 21: 6243–6253.
- Brown, M. F., A. Salmon, U. Henriksson, and O. Söderman. 1990. Frequency dependent ²H N.M.R. relaxation rates of small unilamellar phospholipid vesicles. *Mol. Phys.* 69:379–383.
- Brown, M. F., J. Seelig, and U. Häberlen. 1979. Structural dynamics in phospholipid bilayers from deuterium spin-lattice relaxation time measurements. *J. Chem. Phys.* 70:5045-5053.
- Brown, M. F., and O. Söderman. 1990. Orientational anisotropy of nuclear spin relaxation in phospholipid membranes. Chem. Phys. Lett. 167:158-164.
- Comfurius, P., and R. F. A. Zwaal. 1977. The enzymatic synthesis of phosphatidylserine and purification by CM-cellulose column chromatography. *Biochim. Biophys. Acta.* 488:36–42.
- Davis, J. H. 1979. Deuterium magnetic resonance study of the gel and liquid crystalline phases of dipalmitoyl phosphatidylcholine. Biophys. J. 27:339-359.
- Davis, J. H. 1983. The description of membrane lipid conformation, order and dynamics by ²H-NMR. *Biochim. Biophys. Acta.* 737:117–171.
- Davis, J. H., K. R. Jeffrey, M. Bloom, M. I. Valic, and T. P. Higgs. 1976.Quadrupolar echo deuteron magnetic resonance spectroscopy in ordered hydrocarbon chains. *Chem. Phys. Lett.* 42:390–394.
- DeYoung, L. R., and K. A. Dill. 1988. Solute partitioning into lipid bilayer membranes. *Biochemistry*. 27:5281–5289.

- Dodd, S. W., and M. F. Brown. 1989. Disaturated phosphatidylcholines in the liquid-crystalline state studied by deuterium NMR spectroscopy. *Biophys. J.* 55:102a. (Abstr.)
- Ipsen, J. H., O. G. Mouritsen, and M. Bloom. 1990. Relationships between lipid membrane area, hydrophobic thickness, and acylchain orientational order. *Biophys. J.* 57:405-412.
- Israelachvili, J. N., D. J. Mitchell, and B. W. Ninham. 1976. Theory of self-assembly of hydrocarbon amphiphiles into micelles and bilayers. J. Chem. Soc. Faraday Trans. II. 72:1525-1568.
- Jensen, R. G., and R. E. Pitas. 1976. Synthesis of some acylglycerols and phosphoglycerides. Adv. Lipid Res. 14:213-247.
- Kirk, G. L., S. L. Gruner, and D. L. Stein. 1984. A thermodynamic model of the lamellar to inverse hexagonal phase transition of lipid membrane-water systems. *Biochemistry*. 23:1093-1102.
- Lindblom, G., and L. Rilfors. 1989. Cubic phases and isotropic structures formed by membrane lipids—possible biological relevance. Biochim. Biophys. Acta. 988:221-256.
- Lis, L. J., M. McAlister, N. Fuller, R. P. Rand, and V. A. Parsegian. 1982. Interactions between neutral phospholipid bilayer membranes. *Biophys. J.* 37:657-666.
- Marsh, D., A. Watts, and I. C. P. Smith. 1983. Dynamic structure and phase behavior of dimyristoylphosphatidylethanolamine bilayers studied by deuterium nuclear magnetic resonance. *Biochemistry*. 22:3023-3026.
- Mason, J. T., A. V. Broccoli, and C. Huang. 1981. A method for the synthesis of isomerically pure saturated mixed-chain phosphatidylcholines. *Anal. Biochem.* 113:96–101.
- McIntosh, T. J., and S. A. Simon. 1986. Area per molecule and distribution of water in fully hydrated dilauroylphosphatidylethanolamine bilayers. *Biochemistry*. 25:4948–4952.
- Mely, B., J. Charvolin, and P. Keller. 1975. Disorder of lipid chains as a function of their lateral packing in lyotropic liquid crystals. *Chem. Phys. Lipids.* 15:161–173.
- Meraldi, J.-P., and J. Schlitter. 1981. A statistical mechanical treatment of fatty acyl chain order in phospholipid bilayers and correlation with experimental data A. Theory. *Biochim. Biophys. Acta*. 645:183–192.
- Nagle, J. F., and M. C. Wiener. 1988. Structure of fully hydrated bilayer dispersions. *Biochim. Biophys. Acta.* 942:1-10.
- Nagle, J. F., and D. A. Wilkinson. 1978. Lecithin bilayers. Density measurements and molecular interactions. *Biophys. J.* 23:159–175.
- Salmon, A., S. W. Dodd, G. D. Williams, J. M. Beach, and M. F. Brown. 1987. Configurational statistics of acyl chains in polyunsaturated lipid bilayers from deuterium NMR. J. Am. Chem. Soc. 109:2600-2609.
- Schindler, H., and J. Seelig. 1975. Deuterium order parameters in relation to thermodynamic properties of a phospholipid bilayer. A statistical mechanical interpretation. *Biochemistry*. 14:2283–2287.
- Seelig, J. 1977. Deuterium magnetic resonance: theory and application to lipid membranes. *Q. Rev. Biophys.* 10:353–418.
- Seelig, J., and J. L. Browning. 1978. General features of phospholipid conformation in membranes. FEBS (Fed. Eur. Biochem. Soc.) Lett. 92:41-44.
- Seelig, J., and H.-U. Gally. 1976. Investigation of phosphatidylethanolamine bilayers by deuterium and phosphorus-31 nuclear magnetic resonance. *Biochemistry*. 15:5199-5204.
- Seelig, A., and J. Seelig, 1974. The dynamic structure of fatty acyl chains in a phospholipid bilayer measured by deuterium magnetic resonance. *Biochemistry*. 13:4839–4845.

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- Seelig, A., and J. Seelig. 1975. Bilayers of dipalmitoyl-3-sn-phosphatidylcholine. Conformational differences between the fatty acyl chains. *Biochim. Biophys. Acta.* 406:1-5.
- Seelig, J., and A. Seelig. 1980. Lipid conformation in model membranes and biological membranes. *Q. Rev. Biophys.* 13:19-61.
- Sternin, E., M. Bloom, and A. L. MacKay. 1983. De-Pake-ing of NMR spectra. J. Magn. Reson. 55:274-282.
- Tanford, C. 1980. The Hydrophobic Effect. John Wiley and Sons, Inc., New York.
- Trouard, T. P., R. L. Thurmond, J. A. Barry, S. S. Berr, A. Salmon, S. W. Dodd, S. C. Shekar, and M. F. Brown. 1989. Deuterium NMR spectroscopy of polyunsaturated and disaturated phosphatidylcholine lipid bilayers. *Biophys. J.* 55:103a. (Abstr.)
- Wiedmann, T. S., R. D. Pates, J. M. Beach, A. Salmon, and M. F. Brown. 1988. Lipid-protein interactions mediate the photochemical function of rhodopsin. *Biochemistry*. 27:6469-6474.
- Wieslander, Å., L. Rilfors, and G. Lindblom. 1986. Metabolic changes of membrane composition in *Acholeplasma laidlawii* by hydrocarbons, alcohols, and detergents: arguments for effects on lipid packing. *Biochemistry*. 25:7511-7517.
- Wilkinson, D. A., and J. F. Nagle. 1981. Dilatometry and calorimetry of saturated phosphatidylethanolamine dispersions. *Biochemistry*. 20:187-192.
- Yang, S. F., S. Freer, and A. A. Benson. 1967. Transphosphatidylation by phospholipase D. J. Biol. Chem. 242:477-484.