

Electrostatic interactions in phospholipid membranes revealed by coherent 2D IR spectroscopy

V. V. Volkov^{*†}, R. Chelli^{*‡}, W. Zhuang[§], F. Nuti[¶], Y. Takaoka^{||}, A. M. Papini[¶], S. Mukamel[§], and R. Righini^{*‡}

^{*}European Laboratory for Nonlinear Spectroscopy, Via Nello Carrara 1, I-50019 Sesto Fiorentino, Italy; [‡]Dipartimento di Chimica, Università di Firenze, Via della Lastruccia 3, I-50019 Sesto Fiorentino, Italy; [§]Department of Chemistry, University of California, Irvine, CA 92697; [¶]Laboratory of Peptide and Protein Chemistry and Biology, Dipartimento di Chimica Organica "Ugo Schiff," Università di Firenze, Via della Lastruccia 13, I-50019 Sesto Fiorentino, Italy; and ^{||}Molecular Simulation Group, Taisho Pharmaceutical Company, 1-403 Yoshino-cho Kita-ku Saitama-shi, 331-9530 Saitama, Japan

Communicated by Peter M. Rentzepis, University of California, Irvine, CA, July 11, 2007 (received for review February 20, 2007)

The inter- and intramolecular interactions of the carbonyl moieties at the polar interface of a phospholipid membrane are probed by using nonlinear femtosecond infrared spectroscopy. Two-dimensional IR correlation spectra separate homogeneous and inhomogeneous broadenings and show a distinct cross-peak pattern controlled by electrostatic interactions. The inter- and intramolecular electrostatic interactions determine the inhomogeneous character of the optical response. Using molecular dynamics simulation and the nonlinear exciton equations approach, we extract from the spectra short-range structural correlations between carbonyls at the interface.

carbonyl | interface | intermolecular | nonlinear exciton equations | molecular dynamics

As a constituent organelle in a cell, the membrane sets the information and energy gradients necessary for life. Carbonyl, phosphate, and choline are the common structural moieties in the polar surface of cellular membranes (1), mediating molecular recognition and signal transduction (1–4). Unfortunately, our knowledge on their arrangement and dynamics is rather limited due to experimental difficulties. In a lipid bilayer, lateral irregularity smears the diffraction pattern in neutron scattering measurements, and NMR resonances are broad because of the restricted motions that result in incomplete motional narrowing. IR spectroscopy, on the other hand, is known to be helpful in application to such systems. Since 1972, the IR resonance of carbonyl moieties in phospholipid membranes has attracted considerable attention (5–9). The absorption band shows a clear inhomogeneous character and can be described as a superposition of several substates (5–7). The carbonyl stretching line-shapes in membranes could yield direct information about molecular architecture and fluctuations in the membrane interface (10, 11), provided that the origin of the spectral inhomogeneity of the carbonyl IR response in phospholipid membranes is understood. The inhomogeneity was attributed to differences in the local environment of the sn-1 and sn-2 carbonyl moieties stemming from the packing arrangements (6, 8), the local chain conformations (8, 9, 12–14), the relative positions of the two C=O groups with respect to the interface (9, 15), and the degree of hydration (16, 17). In an elegant work, Blume *et al.* (16) ruled out all of the scenarios involving local structural differences except hydrogen bonding. In a recent study, we eliminated the variance in hydration as a possible source of inhomogeneity (18).

Here we employ 2D IR spectroscopy (19, 20) to explore whether and to what extent the inhomogeneities of the carbonyl absorption can be attributed to electric field fluctuations. 2D IR techniques are femtosecond optical analogues of 2D NMR that generate 2D correlation plots that can separate homogeneous and inhomogeneous broadenings along the diagonal and anti-diagonal axis and provide a rich cross-peak pattern (21). Electrostatic interactions strongly contribute to the optical response of molecular crystals (22), polypeptides (23, 24), and neat liquids (25). In a previous study the Coulomb coupling between sn-1 and

sn-2 carbonyls in a phospholipid molecule was considered negligible (8). In this study, we use the 2D IR response, which measures the electrostatic interactions, whether of inter- or of intramolecular origin, to determine the role played by such interactions at the polar interface of a phospholipid bilayer.

Our investigation focuses on membrane fragments of dimyristoyl-phosphatidylcholine (DMPC), where the sn-2 chain contains a ¹³C-labeled carbonyl (see molecular structure and schematic representation of the DMPC bilayer in Fig. 1). The details on the phospholipid synthesis are provided in ref. 18. To prepare membrane fragments, first we dissolved DMPC in chloroform and dried it on a glass plate. Second, after addition of deuterium oxide, the phospholipid suspension underwent mechanical mixing until the sample (10- μ m film between two CaF₂ windows) demonstrated proper optical quality. Lipid:H₂O molar ratio of 1:6 for the prepared sample suggests a moderate level of hydration. We performed experiments at 38°C; at this temperature the membrane fragments are in the lamellar liquid phase (26, 27).

The spectral degeneracy of the sn-1 and sn-2 carbonyls was lifted, giving two broad vibrational bands at 1,740 and 1,697 cm⁻¹, respectively (see linear IR spectrum in Fig. 2A) for nonlinear spectral probing. The 2D IR experiment uses 1.2-ps pump pulses and 0.1-ps probe pulses with 16 and 200 cm⁻¹ full bandwidth, respectively. This hole-burning technique uses the narrow band pump to select a subensemble of vibrational modes within an inhomogeneous distribution. The detected 2D correlation spectra were obtained by scanning the pump frequency and spectrally resolving the probe transmission. The 2D IR response was analyzed by using the theoretical approach described in refs. 28–30 combined with molecular dynamics (MD) simulations.

Fig. 2B and C shows the 2D IR spectra at early delay time under parallel and perpendicular polarization geometry for the pump and probe pulses. We noticed that the diagonal resonances in the 2D IR spectra do not show time evolution on the time scale of vibrational relaxation, which means that frequency-fluctuation correlation function decays much slower than the vibrational lifetime. The spectra reveal two strong diagonal resonances corresponding to absorption of the two carbonyls (see also the linear IR spectrum in Fig. 2A). The elongation of these resonances along the diagonal direction of the 2D IR spectrum confirms the inhomogeneous character of the absorption bands of the carbonyl moieties (10, 11, 16). Each resonance has a negative (blue) contribution due to ground-state bleaching and to stimulated emission from the first excited state and a

Author contributions: V.V.V. and R.R. designed research; V.V.V., R.C., S.M., and R.R. performed research; F.N. and A.M.P. contributed new reagents/analytic tools; V.V.V., R.C., W.Z., and Y.T. analyzed data; and V.V.V., R.C., W.Z., S.M., and R.R. wrote the paper.

The authors declare no conflict of interest.

Abbreviations: MD, molecular dynamics; DMPC, dimyristoyl-phosphatidylcholine.

[†]To whom correspondence should be addressed. E-mail: vvolkov@lens.unifi.it.

© 2007 by The National Academy of Sciences of the USA

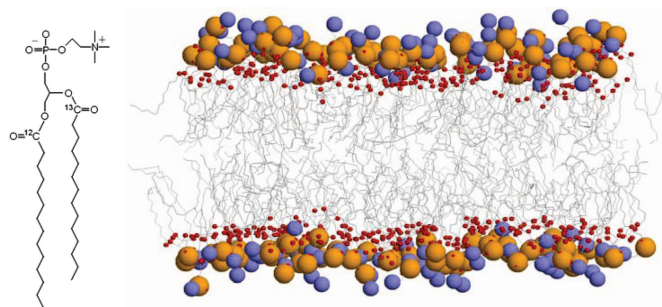


Fig. 1. Chemical structure of the DMPC phospholipid and a snapshot of the DMPC bilayer taken from the MD simulation (for reasons of clarity the water molecules are not shown). The orange, blue, and red spheres represent phosphorus, nitrogen, and oxygen atoms, respectively. The hydrophobic tails are shown with sticks.

positive (red) contribution due to first excited-state absorption. The red shift of the excited-state absorption band reflects the anharmonicity of the carbonyl stretching mode. The 2D IR spectra also show off-diagonal features, known as cross-peaks, that are due to the coupling of the resonantly excited states with the states of different frequency within the same band (intra-band cross-peaks) and with the states which belong to the other resonance (inter-band cross-peaks). In the perpendicular polarization configuration (Fig. 2C), the cross-peaks are more pronounced. In Fig. 3B we present selected horizontal slices of Fig. 2B and C recorded with the pump frequency at 1,675 and 1,752 cm^{-1} (see arrows). The cross-peaks provide a direct measure of vibrational coupling between carbonyl moieties. As in the case of neat liquids (25) and peptides (19), the interaction between transition dipole moments (23, 24) of carbonyl groups is responsible for such coupling, giving rise to delocalized vibrational states, i.e., to vibrational excitons.

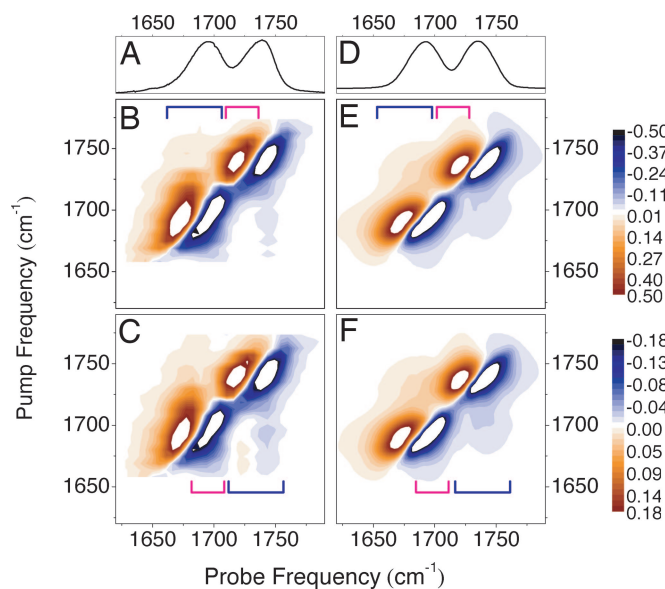


Fig. 2. Experimental and calculated spectra. (A) FTIR optical absorption of the carbonyls in phospholipid membrane fragments. (B and C) Experimental 2D IR spectra recorded under parallel (B) and perpendicular (C) polarization conditions for pump and probe pulses. (D) Calculated linear optical absorption. (E and F) Calculated 2D IR spectra for parallel (E) and perpendicular (F) polarization conditions. Blue and magenta brackets show the spectral regions of the inter- and the intra-band cross-peaks, respectively. Color scale bars show the experimental signal amplitudes in mOD (milli-optical density).

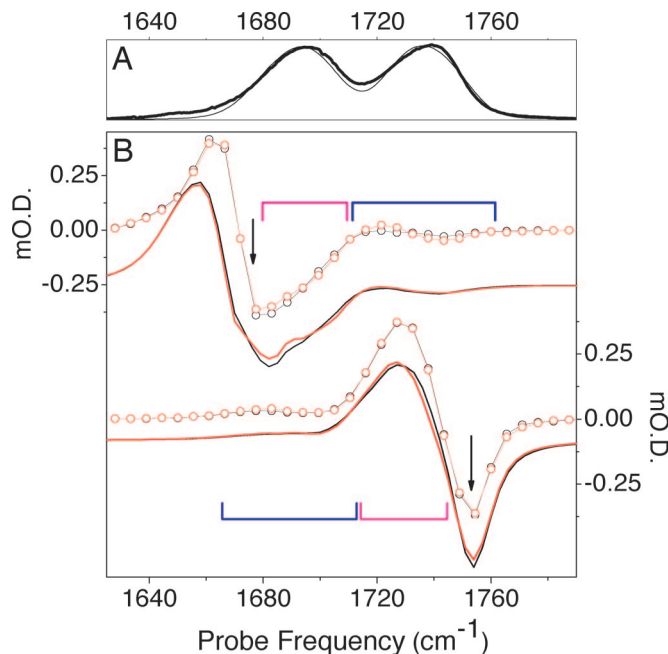


Fig. 3. Experimental and calculated spectra. (A) FTIR optical absorption (thick line) and calculated linear optical absorption (thin line). (B) Experimental (open circle lines) and calculated (solid lines) hole-burning spectra under pump excitation at 1,675 and 1,752 cm^{-1} (see arrows). Black and red colors indicate spectra under parallel and perpendicular polarization conditions, respectively. The spectra under perpendicular polarization are scaled by a factor of three. Blue and magenta brackets specify the spectral regions of the inter- and intra-band cross-peaks, respectively (see also Fig. 2).

The interface of a phospholipid membrane is a highly disordered system (31). To model its linear and nonlinear optical response, we employ an MD simulation of a bilayer system consisting of 112 DMPC molecules (18, 32) to obtain the atomic trajectories. Definition of vibrational states is accomplished by using an $N \times N$ vibrational Hamiltonian matrix, where $N = 112$ is the number of coupled C=O groups in one leaflet of the membrane. The diagonal elements are the zero-order local frequencies corrected by the effect of the electric field acting on each oscillator (33, 34). The Stark frequency shift is $\Delta\omega = k E_{\text{proj}}$, where E_{proj} is the projection of the electric field along the C=O bond. The electric field on each C=O oscillator is calculated in the middle point of the bond following Coulomb's law, i.e., by using the positions and the charges of the atoms known from the MD simulation. The off-diagonal elements of the Hamiltonian matrix are obtained by using the transition dipole coupling mechanism (23, 24, 34), which is based on the dipole-dipole approximation for the electrostatic interaction among the transition dipole moments:

$$\beta_{kl} = \frac{0.1 \boldsymbol{\mu}_k \cdot \boldsymbol{\mu}_l - 3(\boldsymbol{\mu}_k \cdot \mathbf{e}_{kl})(\boldsymbol{\mu}_l \cdot \mathbf{e}_{kl})}{\epsilon R_{kl}^3}, \quad [1]$$

where \mathbf{e}_{kl} is the unit vector connecting the transition dipole moments $\boldsymbol{\mu}_k$ and $\boldsymbol{\mu}_l$ of the carbonyl oscillators k and l at distance R_{kl} and ϵ is the dielectric constant (in our case, set to 1). Diagonalization of the Hamiltonian matrix yields the eigenvectors (excitons) in the carbonyl basis and the corresponding eigenvalues, i.e., the excitonic frequencies. Once the excitons and their frequencies are known, the linear absorption can be calculated as follows:

$$I(\omega) = \left\langle \sum_{m=1}^N \frac{\gamma \left| \sum_{i=1}^N \Phi_{im} \boldsymbol{\mu}_i \right|^2}{(\omega - \omega_m)^2 + \gamma^2} \right\rangle. \quad [2]$$

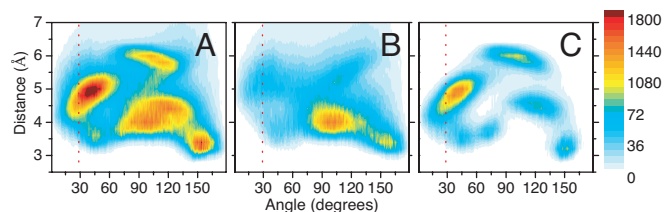


Fig. 5. β' -weighted radial-angular distribution functions (Eq. 5) calculated considering all ^{12}CO - ^{13}CO pairs (A), ^{12}CO - ^{13}CO intermolecular pairs (B), and ^{12}CO - ^{13}CO intramolecular pairs (C). The red dotted lines indicate the angular values obtained from experimental spectral anisotropy. The chromatic bar shows the range of the statistical distribution according to Eq. 5.

The anisotropy recovered from the 2D IR spectra suggests the existence of preferential mutual arrangements of the transition dipole moments and, hence, of the carbonyl moieties. In this respect, the atomistic detail of the MD simulation provides the opportunity of determining the pairing geometry of two transition dipole moments, expressed in terms of mutual distance R and angle θ , that mostly contributes to the cross-peak intensity. The contribution of a given pair arrangement (R, θ) is proportional to the number of transition dipole moment pairs in that arrangement, each pair being weighted by a coupling factor calculated in the context of the transition dipole coupling mechanism. In particular we define a weighted radial-angular pair distribution function as follows

$$h(R, \theta) = \left\langle \sum_{i \in ^{12}\text{CO}}^{N/2} \sum_{j \in ^{13}\text{CO}}^{N/2} \beta'_{ij} \delta(R - R_{ij}) \delta(\theta - \theta_{ij}) \right\rangle, \quad [5]$$

where the sums over i and j run over the ^{12}CO and ^{13}CO carbonyls, respectively. The weight factor β'_{ij} is a pair coupling factor correlated to the parameter β_{ij} (see Eq. 1) and to the difference of the diagonal frequencies, $\omega_i^0 - \omega_j^0$. It is defined as follows:

$$\beta'_{ij} = |\omega_i - \omega_j| - |\omega_i^0 - \omega_j^0|, \quad [6]$$

where ω_i and ω_j are the vibrational frequencies of the two carbonyls obtained considering their coupling in the excitonic fashion, i.e., by diagonalizing the reduced Hamiltonian matrix,

$$H_{\text{red}} = \begin{pmatrix} \omega_i^0 & \beta_{ij} \\ \beta_{ij} & \omega_j^0 \end{pmatrix}. \quad [7]$$

In Fig. 5, we report the $h(R, \theta)$ function calculated considering all ^{12}CO - ^{13}CO pairs, only the ^{12}CO - ^{13}CO intermolecular pairs and only the intramolecular pairs, respectively. In Fig. 5A we note

that for distances of $>6.5 \text{ \AA}$, the $h(R, \theta)$ function vanishes, implying that only neighboring carbonyls give a significant contribution to the cross-peaks, which is in general agreement with the results obtained for the participation ratio reported above. A further interesting feature of the distribution function is its structure. In the total $h(R, \theta)$ (Fig. 5A), we observe the presence of several structural families whose origin (intermolecular or intramolecular) can be easily understood by comparison with Fig. 5B and C. It is relevant to remark that the intermolecular $h(R, \theta)$ function is far from showing random orientation even if it is broader than the intramolecular counterpart. The sharp peak at θ of $\approx 40^\circ$ and $R = 5 \text{ \AA}$ in Fig. 5C corresponds well with the angle between the transition dipole moments obtained from the experimental anisotropy, suggesting that it is mainly originated by the intramolecular pairs. However, we notice that, for this angular value, the intermolecular carbonyl pairs also contribute significantly (up to $26 \pm 5\%$) to the total $h(R, \theta)$ function (see Fig. 5B).

The present experimental and MD simulation study of linear and nonlinear IR response of carbonyl moieties in a phospholipid bilayer reveals the importance of electrostatic interactions at the polar interface. Both the transition dipole moment coupling and the variance of the electric field contribute to the shape and to the bandwidth of the linear IR spectrum. The two contributions, which cannot be distinguished in the linear response, can be clearly separated in the diagonal and in the off-diagonal parts of the 2D correlation plots. The cross-peak intensity is a direct measure of the contribution of coupling to the overall line shape. The diagonal spectral distribution is due to both the frequency dispersion of the excitonic states and the broadening originating from the local electric field. The 2D line shapes allow the unambiguous interpretation of the IR response and specific modeling of the vibrational excitations. The increased localization of the excitonic states in the wings of the IR response allows us to assess local structural properties for the nearest chromophores. 2D IR spectroscopy, in conjunction with the nonlinear exciton equations approach, provides an opportunity to model effectively the structural relations in the phospholipid membranes. This strategy is promising for structural studies through intermolecular coupling in composite phospholipid bilayers, host-guest lipid-protein complexes, lipid systems of reduced dimensionalities, and polymers.

We thank Prof. Akihiro Kusumi for having made available the MD simulation trajectories and Dr. Jason Palmer for proofreading the manuscript. Y.T. thanks Dr. Kunihiro Kitamura and Dr. Hiroh Miyagawa at Taisho Pharmaceutical. This work was supported by Marie Curie Fellowship Contract MTKD-CT2004-509761, European Union Contract RII3-CT-2003-506350, and the Consorzio Interuniversitario Nazionale per la Scienza e Tecnologia dei Materiali (Firenze, Italy). S.M. was supported by National Science Foundation Grant CHE-0446555 and National Institutes of Health Grant 2R01GM59230-05.

- Albert B, Johnson A, Lewis J, Raff M, Roberts K, Walter P (1994) in *Molecular Biology of the Cell* (Garland Science, New York), p 583.
- Simons K, Ikonen E (1977) *Nature* 387:569-572.
- Brown DA, London E (1998) *Annu Rev Cell Dev Bio* 14:111-136.
- Mukherjee S, Maxfield FR (2004) *Annu Rev Cell Dev Bio* 20:839-866.
- Mendelsohn R (1972) *Biochim Biophys Acta* 290:15-21.
- Bunow MR, Levin IW (1977) *Biochim Biophys Acta* 489:191-206.
- Bush SF, Levin H, Levin IW (1980) *Chem Phys Lipids* 27:101-111.
- Bicknell-Brown E, Brown KG, Person WB (1980) *J Am Chem Soc* 102:5486-5491.
- Mushayakarara E, Levin IW (1982) *J Phys Chem* 86:2324-2327.
- Hübner W, Mantsch HH (1991) *Biophys J* 59:1261-1272.
- Lewis RN, McElhaney RN, Pohle W, Mantsch HH (1994) *Biophys J* 67:2367-2375.
- Hitchcock PB, Mason R, Thomas KM, Shipley GG (1974) *Proc Natl Acad Sci USA* 71:3036-3040.
- Pearson RH, Pascher I (1979) *Nature* 281:499-501.
- Hauser H, Pascher I, Pearson RH, Sundell S (1981) *Biochim Biophys Acta* 650:21-51.
- Mushayakarara E, Albon N, Levin IW (1982) *Biochim Biophys Acta* 686:153-159.
- Blume A, Huebner W, Messner G (1988) *Biochem* 27:8239-8249.
- Lewis RN, McElhaney RN (1992) *Biophys J* 61:63-77.
- Volkov V, Nuti F, Takaoka Y, Chelli R, Papini AM, Righini R (2006) *J Am Chem Soc* 128:9466-9470.
- Hamm P, Lim M, Hochstrasser RM (1998) *J Phys Chem B* 102:6123-6138.
- Hamm P, Lim M, Degrado WF, Hochstrasser RM (1999) *Proc Natl Acad Sci USA* 96:2036-2041.
- Scheurer AC, Mukamel S (2002) *J Chem Phys* 116:6803-6816.
- Califano S, Schettino V, Neto N (1981) in *Lattice Dynamics of Molecular Crystals* (Springer, Berlin).
- Krimm S, Abe Y (1972) *Proc Natl Acad Sci USA* 69:2788-2792.

24. Moore WH, Krimm S (1975) *Proc Natl Acad Sci USA* 72:4933–4935.
25. Torii H, Tasumi M (1998) *J Phys Chem B* 102:315–321.
26. Smith G, Sirota E, Safinya C, Piano R, Clark N (1989) *J Chem Phys* 92:4519–4529.
27. Faure C, Bonakdar L, Doufourc E (1997) *FEBS Lett* 405:263–266.
28. Spano FC, Mukamel S (1991) *Phys Rev Lett* 66:1197–1200.
29. Mukamel S, Abramavicius D (2004) *Chem Rev* 104:2073–2098.
30. Zhuang W, Abramavicius D, Hayashi T, Mukamel S (2006) *J Phys Chem B* 110:3362–3374.
31. Sun W, Suter R, Knewton M, Worthington R, Tristram-Nagle S, Zhang R, Nagle J (1994) *Phys Rev E* 49:4665–4676.
32. Takaoka Y, Pasenkiewicz-Gierula M, Miyagawa H, Kitamura K, Tamura Y, Kusumi A (2000) *Biophys J* 79:3118–3138.
33. Park ES, Boxer SG (2002) *J Phys Chem B* 106:5800–5806.
34. Torii H (2004) *J Phys Chem A* 108:7272–7280.
35. Torii H, Tasumi M (1998) *J Raman Spectr* 29:81–86.
36. Volkov V, Palmer DJ, Righini R (2007) *J Phys Chem B* 111:1377–1383.
37. Volkov V, Hamm P (2004) *Biophys J* 87:4213–4225.
38. Mukherjee P, Kass I, Arkin I, Zanni MT (2006) *Proc Natl Acad Sci USA* 103:3528–3533.
39. Torii H (2004) *J Phys Chem A* 108:2103–2107.
40. Thouless DJ (1974) *Phys Rep* 13:93–142.