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Phospholipid binding and transfer by the nonspecific lipid-transfer protein (sterol carrier protein 2)

A kinetic model

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The nonspecific lipid-transfer protein (nsL-TP) from bovine liver was studied by measuring the binding and transfer of the fluorescent phospholipid 1-palmitoyl-2-[6-(1-pyrenyl)-hexanoyl]-sn-glycero-3-phosphocholine (PamPyrGroPCho). A kinetic model is presented involving three steps: (a) interaction of nsL-TP with a membrane surface; (b) equilibration of PamPyrGroPCho monomers between the membrane and nsL-TP; and (c) dissociation of the nsL-TP/PamPyrGroPCho complex from the membrane surface. Steady-state analysis of the model yielded theoretical equations describing both binding and transfer kinetics. Computer analysis, using these equations, showed good fits with the experimental results and several kinetic constants could be calculated. From these constants it was inferred that incorporation of acidic phospholipids into vesicles enhanced the interaction of nsL-TP with the membrane interface (step a), without affecting the equilibrium binding of phospholipid monomers to nsL-TP (step b). As a result, the rate of nsL-TP-mediated PamPyrGroPCho transfer from donor to acceptor vesicles was greatly affected. Under the conditions of incubation, incorporation of the acidic lipids in the donor membrane vesicles stimulated transfer, whereas incorporation of these lipids in the acceptor membranes could lead to a virtually complete inhibition of transfer. From the results it is concluded that the formation of a soluble lipid-nsL-TP complex is the key step in nsL-TP-mediated phospholipid transfer.

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Abbreviations. nsL-TP, nonspecific lipid-transfer protein; PtdCho-TP, phosphatidylcholine-transfer protein; PtdIns-TP, phosphatidylinositol-transfer protein; PtdCho, phosphatidylcholine; PtdOH, phosphatidic acid; PtdIns, phosphatidylinositol; PtdInsP, phosphatidylinositol 4-phosphate; PtdInsP2, phosphatidylinositol 4,5-bisphosphate; Tnp-PtdEtn, N-[2,4,6-trinitrophenyl]-phosphatidylethanolamine; PamPyrGroPCho, 1-palmitoyl-2-[6-(1-pyrenyl)-hexanoyl]-sn-glycero-3-phosphocholine; PamDphGroPCho, 1-palmitoyl-2-[3-(p-diphenylhexatrienyl)-propionyl]-sn-glycero-3-phosphocholine; Nbd, 7-nitro-2,1,3-benzoxadiazol-4-yl.

EXPERIMENTAL PROCEDURES

Materials

Egg-yolk phosphatidylcholine (PtdCho) and phosphatidic acid (PtdOH) were obtained from Sigma. 1-Palmitoyl-2-[6-(1-pyrenyl)-hexanoyl]-sn-glycero-3-phosphocholine (Pam-PyrGroPCho) was synthesized and purified as described [20]. N-[2,4,6-Trinitrophenyl]-phosphatidylethanolamine (Tnp-PtdEtn) was synthesized from egg-yolk phosphatidylethanolamine and trinitrobenzenesulfonic acid [21]. nsL-TP was purified from bovine liver according to Van Amerongen et
Acceptor vesicle preparation

N, atmosphere at 0°C. The Tris buffer was saturated with buffer (0.1-0.5 mM phospholipid) and subjected to ultrasonication for 1 min with a Branson probe sonifier under a N, atmosphere at 0°C. The Tris buffer was saturated with argon prior to resuspension. This procedure did not alter the pyrene moiety as the fluorescence spectrum of the donor vesicle dissolved in ethanol was identical to that of a standard PamPyrGroPCho solution.

Donor vesicle preparation

Solutions of PamPyrGroPCho, egg PtdCho, Tnp-PtdEtn and PtdOH in chloroform, were mixed and dried down by a stream of N2. The dried lipids were resuspended in the Tris buffer (0.1-0.5 mM phospholipid) and subjected to ultrasonication for 1 min with a Branson probe sonifier under a N2 atmosphere at 0°C. The Tris buffer was saturated with argon prior to resuspension. This procedure did not alter the pyrene moiety as the fluorescence spectrum of the donor vesicle preparation dissolved in ethanol was identical to that of a standard PamPyrGroPCho solution.

Acceptor vesicle preparation

Acceptor vesicles consisting of PtdCho and varying amounts of PtdOH were prepared by drying down the lipid mixture from chloroform, followed by suspension of the lipids in the Tris buffer (1-5 mM phospholipid) and ultrasonication under a N2 atmosphere for 5 min with a Branson sonifier. Prior to suspension the Tris buffer was saturated with argon. Vesicles were kept at 0°C and were always used within one day. Vesicle (micellar) suspensions consisting of PtdOH, phosphatidylinositol (PtdIns), phosphatidylinositol 4-phosphate (PtdInsP) or phosphatidylinositol 4,5-bisphosphate (PtdInsP,) were prepared by ultrasonication of a lipid emulsion (50 mM) for 3 min (5 s on and 10 s off) as described above. Interruption of sonication was necessary to prevent degradation of PtdInsP and PtdInsP2 [23].

Phospholipid binding assay

Binding of PamPyrGroPCho to nsL-TP was assayed by the method described for the phosphatidylcholine-transfer protein (PtdCho-TP) [24] or for the phosphatidylinositol-transfer protein (PtdIns-TP) [25]. Briefly, aliquots of nsL-TP (0.09-0.71 µM final concentration) were added to donor vesicles consisting of 50% (by mol) PamPyrGroPCho, 10% (by mol) Tnp-PtdEtn, egg PtdCho and PtdOH (0.25-10 µM total phospholipid, final volume of 2 ml). The increase in pyrene-monomer fluorescence was recorded as a function of time. The slope of the fluorescence increase (initial rate of binding) and the increase at equilibrium (which was usually reached after 10-15 min) were determined. The fluorescence signal was calibrated as described [9], so that the binding could be expressed as the molar amount of PamPyrGroPCho bound to nsL-TP.

Phospholipid transfer assay

This assay is similar to the one described for PtdIns-TP [26] and is based on the pyrene-monomer fluorescence increase as a result of translocation of a PamPyrGroPCho molecule from a quenched donor to an unquenched acceptor vesicle. Briefly, acceptor vesicles (2.5-200 µM) were mixed with donor vesicles (0.25-10 µM) present in the cuvette containing the Tris buffer. The spontaneous transfer of PamPyrGroPCho from donor to acceptor vesicles was followed by recording the increase in pyrene-monomer fluorescence with time. Subsequently, nsL-TP was added (0.09-1.36 µM) to initiate the protein-mediated phospholipid transfer reaction (final volume of 2 ml). Transfer rates were corrected for the spontaneous transfer. The pyrene-monomer fluorescence was calibrated with a standard vesicle preparation which, apart from the 0.05% (by mol) PamPyrGroPCho, was identical to the acceptor vesicles used. Rates of transfer were expressed as pmol PamPyrGroPCho transferred/min.

Kinetic treatment

Kinetic analysis of transfer between donor and acceptor vesicles

The transfer process is described by the following set of reactions:

\[
P + D \xrightarrow{k_1} P \cdot D \xrightleftharpoons{\frac{k_2}{k_1}} PL \cdot D \xrightarrow{k_3} PL + D \quad (A)
\]

\[
PL + A \xrightarrow{k_4} PL \cdot A \xrightleftharpoons{\frac{k_5}{k_4}} P \cdot A \xrightarrow{k_6} P + A \quad (B)
\]

where P, D and A represent nsL-TP, donor membrane and acceptor membrane, respectively. PL designates nsL-TP with a bound PamPyrGroPCho molecule. P · D and P · A are complexes between nsL-TP void of bound lipid, with the donor and the acceptor membrane, respectively. Similarly, PL · D and PL · A represent complexes between nsL-TP that contains PamPyrGroPCho, with the donor membrane and the acceptor membrane. The binding reaction (reaction A) was divided into three steps: two steps concern the association/dissociation reactions between nsL-TP and the donor membrane \((k_a, k_d)\), and one step concerns the equilibration of lipid monomers between nsL-TP and the donor vesicle (reactions \(k_1, k_2\)). It is assumed that the association/dissociation reactions \((k_a, k_d)\) are described by similar rate constants for both P or PL.

In this model the initial rate of transfer is given by the velocity at which PamPyrGroPCho is released in the acceptor membrane. This can be expressed as:

\[
\frac{d[P]}{dt} = -k_4[P][D] + k_3[P \cdot D] - k_5[P][A] + k_2[P][A] = 0
\]

If steady-state conditions are assumed the following equations can be derived:
\[
\begin{align*}
\frac{d[PL]}{dt} &= -k_d[PL][D] + k_{-d}[PL \cdot D] \\
&\quad - k_i[PL][A] + k_{-i}[PL \cdot A] = 0 \\
\frac{d[P \cdot D]}{dt} &= k_i[P][D] - k_{-i}[P \cdot D] \\
&\quad - k_d[P \cdot D] + k_{-d}[PL \cdot D] = 0 \\
\frac{d[PL \cdot D]}{dt} &= k_i[PL][D] - k_{-i}[PL \cdot D] \\
&\quad + k_d[P \cdot D] - k_{-d}[PL \cdot D] = 0 \\
\frac{d[P \cdot A]}{dt} &= k_i[P][A] - k_{-i}[P \cdot A] \\
&\quad - k_d[P \cdot A] + k_{-d}[PL \cdot A] = 0 \\
\frac{d[PL \cdot A]}{dt} &= k_i[PL][A] - k_{-i}[PL \cdot A] \\
&\quad + k_d[P \cdot A] - k_{-d}[PL \cdot A] = 0.
\end{align*}
\]

The total protein concentration \([P_{tot}]\) is given by:
\[
[P_{tot}] = [P] + [PL] + [P \cdot D] + [PL \cdot D] + [P \cdot A] + [PL \cdot A].
\]

From rewriting and substitution of Eqns (4) and (5) and from Eqns (6) and (7), Eqns (9) and (10), respectively, are obtained:
\[
\begin{align*}
[P \cdot D] &= \frac{a_d}{k_{-i}}[D][PL] + \left(\frac{a_d}{k_{-i}} + \frac{a_d}{k_{-d}}\right)[D][P]; \\
\alpha_d &= \frac{k_{-d}k_i}{k_{-i} + k_d + k_{-d}} \\
\end{align*}
\]

\[
\begin{align*}
[P \cdot A] &= \frac{a_d}{k_{-i}}[A][PL] + \left(\frac{a_d}{k_{-i}} + \frac{a_d}{k_{-d}}\right)[A][P]; \\
\alpha_d &= \frac{k_{-d}k_i}{k_{-i} + k_d + k_{-d}}.
\end{align*}
\]

Substitution of Eqns (9) and (10) in Eqn (2) yields:
\[
\begin{align*}
[P] &= \frac{a_d[D] + a_d[A]}{K_d a_d[D] + K_d a_d[A]}; \\
[PL] &= \frac{K_d a_d[D] + K_d a_d[A]}{K_d a_d[D] + K_d a_d[A]}; \\
K_d &= k_d/k_{-d}; \quad K_a = k_a/k_{-a}.
\end{align*}
\]

Addition of Eqns (4) and (5) and of Eqns (6) and (7) yield Eqn (12) and Eqn (13) respectively:
\[
\begin{align*}
[P \cdot D] + [PL \cdot D] &= \{[P] + [PL]\}K_i[D]; \quad K_i = k_i/k_{-i} \\
[P \cdot A] + [PL \cdot A] &= \{[P] + [PL]\}K_i[A]; \quad K_i = k_i/k_{-i}.
\end{align*}
\]

Substitution of Eqns (12) and (13) into Eqn (8) results in Eqn (14):
\[
\begin{align*}
[P_{tot}] &= \{[P] + [PL]\} \{1 + K_i[D] + K_i[A]\}.
\end{align*}
\]

Substitution of Eqn (11) into Eqn (14) yields:
\[
\begin{align*}
[PL] &= \frac{[P_{tot}]}{1 + K_i[D] + K_i[A]} \left\{ \frac{a_d[D] + a_d[A]}{K_d a_d[D] + K_d a_d[A]} + 1 \right\}
\end{align*}
\]

By subsequently using Eqns (6) and (14), Eqn (1) can be rewritten as:
\[
\begin{align*}
\dot{v}_0 &= k_i[P \cdot A] - k_i[P][A] = a_d[A][PL] - K_i[P].
\end{align*}
\]

Finally substitution of Eqns (11) and (15) into Eqn (16) yields:
\[
\begin{align*}
\dot{v}_0 &= \frac{a_d a_d(K_d - K_i)[D][A][P_{tot}]}{[a_d(1 + K_i)[D] + a_d(1 + K_i)[A] \{1 + K_i[D] + K_i[A]\}}.
\end{align*}
\]

The reaction constants \(k_a\) and \(k_d\) are dependent on the mole fraction of PamPyrGroPCho in the membrane [9]. As on time zero the mole fraction of PamPyrGroPCho in the acceptor vesicles is zero, \(k_a\) and hence also \(K_a\) become zero. Under these experimental conditions, Eqn (17) can be simplified to:
\[
\begin{align*}
\dot{v}_0 &= \frac{a_d a_d(K_d - K_i)[D][A][P_{tot}]}{[a_d(1 + K_i)[D] + a_d(1 + K_i)[A] \{1 + K_i[D] + K_i[A]\}} \\
\alpha_d &= \frac{k_d k_i}{k_{-i} + k_{-d}}.
\end{align*}
\]

**Kinetic treatment of equilibrium binding**

The binding process is described by reactions A. The equilibrium binding, therefore, can be directly derived from Eqn (15) by, substituting \([A] = 0\):
\[
\begin{align*}
[PL] &= \frac{K_d}{1 + K_d [1 + K_i[D]]}.
\end{align*}
\]

This equation predicts that equilibrium binding is independent of the donor concentration at conditions where \(K_i[D] \ll 1\).

**RESULTS**

**Exchangeability of donor lipids**

In view of the net-transfer capability of nsL-TP, it is of interest to check whether the protein has access to all lipids in the donor membrane, or only to the lipids in the outer leaflet. At low pyrene-phospholipid concentration (0-5%) in a PtdCho bilayer, the pyrene excimer fluorescence is linear with the pyrene surface concentration [27]. The exchangeability of PamPyrGroPCho from the donor vesicle can therefore be estimated by monitoring the pyrene-exciton fluorescence decrease in the presence of a large excess of unlabeled acceptor vesicles and nsL-TP. Since PamPyrGroPCho in the acceptor vesicles is greatly diluted, the excimer fluorescence in the acceptor vesicles is negligible. As shown in Fig. 1, 60% of PamPyrGroPCho in the donor membrane is transferred by nsL-TP, independent on the amount of PtdOH present in the acceptor membrane. In the absence of nsL-TP an identical amount of PamPyrGroPCho is transferred, though at a much slower rate. This experiment is taken to indicate that nsL-TP, like PtdCho-TP [28] and PtdIns-TP [29] has only access to phospholipids in the outer leaflet of the donor vesicle in agreement with previous studies [30, 31]. The donor vesicles used in the other experiments (see below) contained 10% (by mol) Tnp-PtdEtn, and hence the asymmetry of donor lipid accessibility to nsL-TP could not be determined for these vesicles. Although it is not critical for the proposed kinetic model, we assume that, for the donor vesicles described below, this accessibility is unchanged.
Fig. 1. Exchangeability of PamPyrGroPCho in the donor vesicle. Lipid transfer was measured between donor vesicles (10 μM) consisting of PamPyrGroPCho, PtdOH and PtdCho, molar ratio 5:10:85, and acceptor vesicles (250 μM) consisting of PtdOH and PtdCho, molar ratio 5:95 (●, ○) or of 10:90 (□, □) which were incubated in the presence (●, □) or absence (○, △) of 1.7 μM nsL-TP (final volume 2 ml). The pyrene excimer fluorescence was obtained by exciting at 346 nm (bandpass 2 nm) and measuring the fluorescence at 470 nm (bandpass 10 nm).

Fig. 2. Fluorescence measurement of PamPyrGroPCho binding to nsL-TP. Arrow A indicates the addition of donor vesicles (1 μM) consisting PamPyrGroPCho, Tnp-PtdEtn and PtdCho (molar ratio 50:10:40). At arrow B the binding reaction was initiated by the addition of nsL-TP (0.34 μM, final volume 2 ml). The dashed line indicates the extrapolated initial rate of binding. The fluorescence was recorded at excitation 346 nm (bandpass 2 nm) and emission 377 nm (bandpass 10 nm).

Effect of nsL-TP concentration

Fig. 3. Effect of nsL-TP concentration on PamPyrGroPCho binding to nsL-TP. The experiments were performed as indicated in Fig. 2. The donor vesicles (1 μM) consisted of PamPyrGroPCho, Tnp-PtdEtn and PtdCho (molar ratio 50:10:40). In (A), the symbols represent the equilibrium concentration of the nsL-TP/PamPyrGroPCho complex. In (B), the symbols represent the initial rate of binding.

Effect of donor vesicle concentration

As seen in Fig. 4A, the amount of PamPyrGroPCho binding to nsL-TP increases with increasing donor vesicle phospholipid concentration up to 1 μM, whereas above 1 μM a zero-order dependency between donor vesicle concentration and the PamPyrGroPCho binding is apparent. The fact that, at [D] > 1 μM, binding is independent of the donor vesicle concentration is in agreement with Eqn (19) and lends strong support for the proposed model. The decrease in lipid binding at low [D] contradicts Eqn (19). It cannot be explained by exhaustion of the donor PamPyrGroPCho content, since, even at the lowest [D], only 6% of the PamPyrGroPCho is bound to nsL-TP (8 nM equilibrium concentration cf. 125 nM total PamPyrGroPCho concentration). We propose an additional reaction where nsL-TP releases its bound lipid into the buffer (see Appendix). Under these conditions, the so-called equilibrium binding no longer reflects a true equilibrium, but a steady-state process. In this extended model, the release of bound lipid into the buffer is assumed to be a very slow process (minute timescale) only occurring to a significant extent at very low vesicle (lipid) concentration. Given these assumptions, we could optimize a $K_s$ at 0.046, $k_{cat}/k_{on} = 0.24$ μM, and $K_i = 7$ mM$^{-1}$ (Eqn A9, see legend to Fig. 4). This gives rise to a β value of 0.24 μM representing the donor vesicle concentration at which the release reaction of the bound lipid to the donor membrane (reaction A) and the release reaction to the buffer (reaction C) occur at identical rates. Therefore at [D] > 1 μM, Eqn (A9) can be approximated by Eqn (19).
Fig. 4. Effect of donor vesicle concentration on PamPyrGroPCho binding by nsL-TP. The experiments were performed as described in Fig. 2. Donor vesicles consisted of PamPyrGroPCho, Tnp-PtdEtn and PtdCho (molar ratio 50:10:40). The nsL-TP concentration was 0.34 μM. In (A), the symbols represent the equilibrium concentrations of the nsL-TP/PamPyrGroPCho complex. Nonlinear least-squares analysis according to Eqn (A9) (see Appendix) yielded \( K_d = 0.046 \pm 0.004 \), \( K = 7 \pm 19 \) mM\(^{-1} \) and \( k_+/k_- = 0.28 \pm 0.07 \) μM. The curve as indicated was drawn based on these constants and showed a correlation coefficient \( r = 0.979 \). In (B), the symbols represent the initial rate of binding.

The initial binding velocity is strikingly linear with the donor vesicle concentration (see Fig. 4B). This indicates that the rate-limiting step in the binding process occurs in a first-order dependency with the donor vesicle concentration, and thereby lends further support for a model involving reaction A (see Methods).

Effect of donor vesicle surface charge

In Fig. 5A it is shown that the equilibrium binding at three different nsL-TP concentrations is hardly dependent on the PtdOH surface concentration (or surface charge). Since the association of nsL-TP with these vesicles can be considered as minimal at these low [D], (i.e. \( K_D(D) \approx 1 \), see also below), it is inferred that \( K_c \) (Eqn 19) is independent of the membrane surface charge. In other words the equilibration of lipid monomers on and off the protein at the lipid-protein interface (second step in reaction A) is not dependent on the charge of the membrane. This confirms the experiments described in another study [9]. The initial rate of binding as depicted in Fig. 5B, however, does increase with increasing membrane surface charge. Apparently the first step in reaction A (the association of nsL-TP with membranes) is accelerated by increasingly negatively charged membranes, resulting in a faster equilibration (see also discussion below).

Transfer kinetics

Then nsL-TP mediated transfer of PamPyrGroPCho from donor to acceptor vesicles was monitored as shown in Fig. 6. After addition of unlabeled acceptor vesicles (arrow B) to quenched donor vesicles containing PamPyrGroPCho, the spontaneous rate of transfer from donor to acceptor vesicles is measured \( (V_b) \). After addition of nsL-TP (arrow C) the rate of transfer is increased to a large extent. This nsL-TP-mediat-
ated transfer was always corrected for the spontaneous transfer. The kinetic model was tested by varying the concentration and/or composition of the three components.

**Effect of nsL-TP concentration**

The dependency of PamPyrGroPCho transfer on the nsL-TP concentration was studied by performing transfer experiments at variable nsL-TP concentrations. At a fixed donor and acceptor vesicle concentration, this dependency was found to be linear in a wide nsL-TP concentration range (0–1.5 μM, see Fig. 7), which lends support for the presented kinetic model (see Eqn 18).

**Effect of acceptor vesicle concentration and surface charge**

At a fixed donor vesicle (5 μM) and nsL-TP concentration (0.34 μM), a variation of the acceptor vesicle concentration and composition clearly influenced nsL-TP-mediated PamPyrGroPCho transfer (see Fig. 8). As seen from the curve corresponding to the acceptor vesicles containing 5 mol PtdOH/100 mol, the rate of transfer increased at acceptor vesicle concentrations up to 50 μM whereas above 50 μM the rate of transfer was almost constant. A curve analysis was performed with Eqn (18), using $K_a = 0.046$ and $K_i = 7$ mM$^{-1}$. From the kinetic constants it is clear that both $K_a$ and $a'$ increase with increasing negative surface charge in the acceptor vesicles.

**Effect of donor vesicle concentration and surface charge**

The effect of the donor vesicles on nsL-TP-mediated transfer was tested at a fixed acceptor vesicle concentration (50 μM, 5%, by mol, PtdOH) and nsL-TP concentration (0.34 μM), see Fig. 9. When PtdOH was included in the donor vesicles, the rate of transfer increased significantly. This is in agreement with previous studies [9, 19]. Also the deviation from linearity became more pronounced when the PtdOH concentration was increased in the donor vesicle (see Fig. 9). Kinetic analysis was performed with Eqn (18) using $a' = 259$ pmol $\cdot$ min$^{-1}$ μM$^{-2}$, $K_a = 1.4$ mM$^{-1}$, and $K_i = 59.9$ ± 0.7 mM$^{-1}$ (r = 0.9998) were determined respectively. The curves drawn in the figure are calculated from these constants.

**Inhibition experiments**

To further explore the affinity of nsL-TP for negatively charged surfaces, inhibition experiments were performed em-
DISCUSSION

Validation of the kinetic model

The main reason for proposing a kinetic model is to obtain a better understanding of the mode of action of nsL-Tp-mediated transfer of PtdCho. In this model both binding and transfer characteristics are combined. The validation of reactions A and B (see Experimental Procedures), is given by the observation that nsL-TP is able to extract a phospholipid monomer from a membrane and accommodate it into an endogenous lipid binding site [6-9] after which the phospholipid follows the rotation mobility of a free dissolved nsL-TP monomer [8]. In the model, the binding reaction is divided into three steps: an association of nsL-TP with a donor membrane; a step where one phospholipid monomer translocates from the membrane to nsL-TP; and another association (dissociation) reaction of the nsL-TP/phospholipid complex with the membrane. The translocation step is very likely dependent on the membrane lipid class and its surface concentration, whereas the association steps are more dependent on physical parameters like membrane charge and size. This division allowed us to determine kinetic constants describing these individual steps.

Two assumptions are made in the model. The first is that the association constants $K_1$ and $K_2$ are independent of whether the protein contains an endogenous phospholipid molecule or not. This assumption is based on the facts that (a) upon binding of PtdCho the charge on nsL-TP will not change; (b) the mass of nsL-TP will increase only 5% after PtdCho binding; (c) without this assumption a very complex equation is obtained for $v_0$ (not shown) with which it is practically impossible to determine the various constants; and (d) the correctness of a model is likely to increase with the simplicity of the model [32]. The second assumption pertaining to the steady-state conditions is based on the near linearity of the fluorescence increase after addition of nsL-TP to a mixed donor/acceptor vesicle solution, extending over a long time range (see Fig. 6). This last assumption has also been made for the analysis of the transfer kinetics of PtdCho-TP and PtdIns-TP [26, 33, 34].

An alternative model invoked by the correlation between spontaneous and nsL-TP-mediated transfer [9, 17, 18] which involves pathways where lipid monomers dissociate spontaneously from the membrane into the aqueous phase after which they eventually may bind to nsL-TP, is excluded since such a mechanism would lead to nonlinear relationships between binding and transfer with [nsL-TP] which we did not observe (see Figs 3 and 7). Furthermore such a model cannot explain a stimulation of transfer by nsL-TP because the rate-limiting step in spontaneous transfer is the dissociation of a lipid monomer from the membrane interface [16]. Models in which nsL-TP binds to the membrane and increases the off-rate of lipid monomers or in which nsL-TP forms a ternary complex between donor and acceptor vesicles [11-16] are also excluded. These models can only explain the fluorescence increase observed in the binding experiments in terms of a complex between nsL-TP, the donor membrane and the fluorescent lipid (like [PL - D] in reaction A). First it is expected that such a complex will be accessible to quenching by the trinitrophenyl moieties of the Tnp-PtdEtn present in the donor membrane (so we do not expect a substantial increase in fluorescence intensity upon formation of such a complex). Secondly, these models would predict an increase of equilibrium binding with increasing association of nsL-TP with the donor vesicle which is not observed. Consequently,

ploying donor and acceptor vesicles and a third vesicle/micelle population consisting of pure negatively charged phospholipids. As seen from Fig. 10, relatively low concentrations of these inhibitory vesicles/micelles may have a severe effect on nsL-Tp-mediated transfer. An addition of as little as 5 μM of PtdInsP₂ micelles completely inhibited nsL-TP-mediated PamPyrGroPCho transfer (see Fig. 10). The inhibition is explained by an association of nsL-TP with the PtdInsP₂ micelle after which the protein is incapable of transferring donor lipids to the acceptor membranes.

![Fig. 9. Effect of donor vesicle concentration and charge on the nsL-TP-mediated transfer of PamPyrGroPCho. Transfer was measured as described in Fig. 6. The incubation medium (2 ml) contained acceptor vesicles (50 μM) consisting of PtdCho and PtdOH (molar ratios 95:5), nsL-TP (0.34 μM) and donor vesicles at various concentrations containing PamPyrGroPCho, PtdOH, Tnp-PtdEtn, and PtdCho at molar ratios of 20:0:10:70 (△), 20:5:10:65 (●), 20:0:10:60 (▲), or 20:20:10:50 (■). The experimental data were fitted to Eqn (18) by using $a'_0 = 259 \text{ pmol} \cdot \text{min}^{-1} \cdot \text{μM}^{-1}$, $K_1 = 1.4 \text{ mM}^{-1}$ and $K_2 = 0.046$. For the data corresponding to 0% (△), 5% (●), 10% (▲), and 20% (■) PtdOH, $a'_0 = 378 \pm 8 \text{ pmol} \cdot \text{min}^{-1} \cdot \text{μM}^{-1}$ and $K_1 = 23 \pm 6 \text{ mM}^{-1}$ ($r = 0.997$); $a'_0 = 924 \pm 57 \text{ pmol} \cdot \text{min}^{-1} \cdot \text{μM}^{-2}$ and $K_1 = 35.9 \pm 1.8 \text{ mM}^{-1}$ ($r = 0.9997$); $a'_0 = 2450 \pm 55 \text{ pmol} \cdot \text{min}^{-1} \cdot \text{μM}^{-2}$ and $K_1 = 52 \pm 2 \text{ mM}^{-1}$ ($r = 0.997$) were determined respectively. The curves drawn in the figure were calculated from these constants.

![Fig. 10. Inhibition of nsL-TP-mediated transfer by acidic phospholipid vesicles/micelles. The transfer of PamPyrGroPCho was measured as described in Fig. 6 with donor vesicles (2 μM) containing PamPyrGroPCho, Tnp-PtdEtn, and PtdCho (molar ratio 10:10:80), acceptor vesicles (50 μM) containing PtdCho and PtdOH (molar ratio 95:5) and nsL-TP (0.34 μM). The sonicated acidic phospholipids PtdIns (△), PtdOH (○), PtdInsP (▲) and PtdInsP₂ (■) were added prior to the initiation of transfer by nsL-TP.](image-url)
the zero order dependency with [D] in Fig. 4A and the results in Fig. 5A can only be explained with these alternative models if we assume that 100% of nsL-TP is bound to the donor membrane (at [D] > 1 μM) which would give 100-fold higher association constants than the ones observed in Fig. 8.

**Binding experiments**

One additional pathway ($k_a$) was needed to explain the binding data (see Appendix). This reaction was found to be of minor importance at [D] > 1 μM. Therefore we excluded this pathway from the analysis of the transfer experiments. From combining $\gamma_0 = 372$ pmol · min$^{-1}$ · μM$^{-2}$ (or 0.19 min$^{-1}$ · μM$^{-1}$), (see Figs 8 and 9) with $k_3/\gamma_0 = 0.24$ μM (Fig. 4A) it is inferred that $k_3$ equals 0.05 min$^{-1}$. This means that only 5% of the nsL-TP/phospholipid complex dissociates into nsL-TP and phospholipid/min (regardless of donor, or acceptor, concentration), which supports the notion that at high [D] and [A] this pathway is insignificant. It is of note that this pathway explains the failure to isolate the nsL-TP/phospholipid complex [6, 8, 10] since, upon separation of donor membranes from nsL-TP, [D] ≪ 1 μM after which nsL-TP releases its bound phospholipid into the medium.

In a previous study, Nichols also investigated lipid-binding kinetics by nsL-TP [7]. In that study the initial binding kinetics were studied as opposed to the equilibrium (or steady-state) binding described here. Based on the experiments with 1-palmitoyl-2-[12-(7-nitro-2,1,3-benzoxadiazol-4-yl)amino]-dodecanoyl-sn-glycero-3-phosphocholine (PamNbdLauPyrGroPCho), Nichols concluded that lipid binding and release was a complex mechanism involving not only a collisional pathway (reaction A) but also a monomer-diffusion step. In this study, we could adequately describe our experimental data without including a monomer-diffusion step. Also the linear relationships between lipid-vesicle concentration and nsL-TP concentration with initial binding rates (Figs 3B and 4B) are in agreement with a collisional step determining the initial binding process. In a previous study, in which we described the acyl-chain dependency of equilibrium binding and steady-state transfer, we proposed that nsL-TP lowers the energy barrier for lipids to leave the membrane, but the relative tendency for lipids to leave the membrane is not altered by nsL-TP: short acyl-chain lipids bind faster and to a higher extent to nsL-TP. Perhaps the difference between our results and those of Nichols is explained by the fact that, in contrast to the pyrene moiety of PamPyrGroPCho, the Nbd moiety of Pam(NbdLau)PyrGroPCho, the Nbd moiety of Pam(NbdLau)GroDicho is located near the membrane interface [35] and hence it is expected that Pam(NbdLau)GroPCho dissociates from the membrane with a different mechanism.

A $K_a$ = 0.046 was optimized from the binding experiments. In a previous study, it was shown that this parameter is linearly dependent on the PamPyrGroPCho mole fraction in the donor membrane [9]. The linear relationship of $K_a$ with the mole fraction of various pyrenyl-acyl-labeled PtdCho species as shown in [9] is indicative of the absence of competition of unlabeled PtdCho for the lipid-binding site on nsL-TP: Eqn (2) in [9] can by directly derived from our Eqn (11) by substitution of [A] = 0. This means that at 0% PamPyrGroPCho surface concentration, $K_a = 0$ (or $K_a = 0$), a condition which was used to convert Eqn (17) into Eqn (18). Interestingly, the equilibrium binding (or $K_a$) did not change upon increasing the negative surface charge in the donor membrane, whereas the initial binding velocity was markedly stimulated. This invariance of $K_a$ with negative surface charge confirms a previous study [9] and was taken into account in the fit of the transfer data to Eqn (18). The inverse correlation of $K_a$ with acyl chain length and the positive correlation with phospholipid headgroup polarity [9] strongly suggest that the equilibration of P · D into PL · D (reaction A) is passive in nature. In that study we proposed that by lowering the energy barrier for lipid monomers to dissociate from the membrane surface, nsL-TP only accelerates the natural tendency of lipids to leave the membrane.

**Transfer kinetics**

Eqn (17), which can be considered as a general equation describing phospholipid transfer, has some interesting features. First of all it resembles the equation derived by van den Besselaar et al. which applies to PtdCho-TP- and PtdInsTP-mediated transfer of PtdCho [26, 33]. Secondly, $\gamma_0$ reflects mass transfer instead of exchange (as for PtdCho-TP and PtdIns-TP). It can be seen from Eqn (17) that the sign of $\gamma_0$ is positive (i.e. net transfer from donor to acceptor) when $K_a > K_d$ whereas $\gamma_0$ is negative when $K_a > K_d$ (i.e. net transfer from ‘acceptor’ to ‘donor’). Since a previous study revealed that $K_d$ is linear with the surface concentration [9], it is inferred that nsL-TP will only catalyze transfer of phospholipids down a surface concentration gradient irrespective of the charge or association constant with the membranes.

From the transfer experiments in Figs 8 and 9, the association constants $K_1$ and $K_2$ and the kinetic quantities $\alpha_0$ and $\alpha_d$ were determined as a function of membrane surface charge. We plotted these constants as a function of surface charge as depicted in Fig. 11 after considering the charge of −1 for Tnp-PtdEtN and of −1.2 for PtdOH [36]. Here it is clearly seen that all four constants increase markedly with increasing negative charge and that there is a high correlation between the constants derived for the acceptor and donor membrane. This is taken as a strong support for Eqn (18) and (17) which are symmetric in nature with respect to [A] and [D]. The
association constants \( K_1 \) and \( K_2 \) are much higher than those found for the interaction of PtdCho-TP or PtdIns-TP with negatively charged membranes [26, 33, 34]. This is probably due to the fact that nsL-TP is positively charged and PtdCho-TP and PtdIns-TP are negatively charged at physiological pH. The association constant \( K_2 \) is responsible for the decreased rate of transfer at higher acceptor vesicle concentrations. In fact, it may be concluded that in the experiment with acceptor vesicles containing 20% (by mol) PtdOH at a lipid concentration of 100 \( \mu \)M, about 85% of nsL-TP was associated with the acceptor vesicles and hence was no longer available for transfer. It is of interest that, in a time-resolved fluorescence study [8], it was shown, that in the presence of 100 \( \mu \)M phosphatidylcholine vesicles containing 20% (by mol) PtdOH, the rotational motion of nsL-TP was retarded (1.5-fold). From this it was inferred that the interaction of nsL-TP with membranes containing acidic phospholipids results in a partial rather than in a complete immobilization of the protein [8]. The inhibition experiments as depicted in Fig. 10 further emphasize the marked association of nsL-TP with negatively charged membrane surfaces. The inhibition increases in the order PtdIns < PtdOH < PtdInsP < PtdInsP2 in analogy with the increasing negative charge of these lipids [36, 37]. It is of note that at a PtdInsP2/nsL-TP molar ratio of 7:1, the nsL-TP-mediated transfer is already inhibited more than 50%. A similar study with PtdInsP2nsL-TP yielded comparable results [26].

It is difficult to ascribe one process to the quantities \( a_s' \) and \( a_s \) since they are composed of many rate constants. It is however interesting to consider two extremes. First where \( k_+ \gg k_- \) and \( k_s \gg k_{-s} \), then \( a_s' = k_2 \) and \( a_s = k_1 \) (since \( K_d \ll 1 \)). This would imply that the association rate of nsL-TP with membranes increases with increasing membrane negative charge (see Fig. 11) whereas the dissociation from these membranes would be hardly affected (since \( K_1 \) and \( K_2 \) increase proportionally to \( a_s' \) and \( a_s \)). This would be exactly opposed to the results obtained for PtdIns-TP where \( k_2 \) remained constant and \( k_{-2} \) decreased with increasing membrane negative charge [26]. The second situation to consider is \( k_+ \ll k_- \) and \( k_s \ll k_{-s} \). then \( \alpha_s' = K_d \alpha_s \) and \( \alpha_s = K_1 k_s \). This would imply that \( k_{-s} \) and \( k_+ \) hardly change upon variation of the negative charge of the membrane, and would be consistent with the observation that \( K_d \) is invariant with membrane surface charge. We tentatively favor this last possibility.

In conclusion, we have shown that the kinetic model of PamPyrGroPCho binding and transfer by nsL-TP is consistent with a mode of action of nsL-TP involving the formation of a freely dissolved lipid/protein complex (reaction A). From the binding experiments it became evident that the equilibrium binding of PamPyrGroPCho to nsL-TP (second step in reaction A) is independent of membrane surface charge. The non-specific character of nsL-TP must find its origin in this second step. We propose a passive role of nsL-TP, lowering the energy barrier for lipids to leave the membrane interface [9]. This also explains why the access of nsL-TP to the lipids in the membrane is restricted to the outer leaflet (Fig. 1) [30, 31]. An increase in membrane surface charge significantly influences the binding kinetics as well as the nsL-TP-mediated transfer of PamPyrGroPCho between vesicles (Figs 5B, 7 and 8). These effects can be explained by a preferential interaction of nsL-TP with vesicles containing negative charge, which probably reflects rather aspecific electrostatic interactions between the positively charged protein [38] with the negatively charged vesicle surface. In view of the high membrane concentrations in cells (about 40 mM [7]) it may well be that in cells the majority of the nsL-TP molecules is associated with membranes. From the phospholipid binding experiments it was inferred that at very low membrane concentrations the protein may release its bound phospholipid in the aqueous phase. In view of the high lipid concentrations found in cells, this pathway is of minor importance. In other words, in the cell nsL-TP acts most likely as a phospholipid carrier, in agreement with the conclusions by Nichols [7]. This release pathway, however, may be more significant for sterols as the formation of a nsL-TP sterol complex could not be demonstrated in similar binding experiments [9]. In fact, the assumption that, for sterols, pathways C and D (see Appendix) are predominant (i.e. \( \alpha_s > a_s' \)) may explain why, despite the lack of sterol binding, sterols are transferred by nsL-TP. This statement requires further study but, if true, would have major implications for the targeting of sterols in cells.

**APPENDIX**

**Binding kinetics with irreversible dissociation of lipid monomer from nsL-TP**

A decrease of lipid binding with decreasing donor concentration can be explained by inclusion of an irreversible dissociation reaction of the complex between nsL-TP and PamPyrGroPCho (PL) into P and L (a lipid monomer in the aqueous phase) and a subsequent reassociation of this lipid monomer with the donor membrane. The binding process is then described by reactions A, C and D:

\[
P + D \xrightarrow{k_{a,+}} P \cdot D \xrightarrow{k_{a,-}} PL + D \quad (A)
\]

\[
PL \xrightarrow{k_s} P + L \quad (C)
\]

\[
L + D \xrightarrow{k_{a,-}} D. \quad (D)
\]

By inclusion of this irreversible step, binding will no longer be an equilibrium process but a steady-state reaction. It is of note that any binding model that assumes equilibrium conditions and includes reaction A will always yield the expression for \([PL]\) as given in Eqn (19). By assuming steady-state conditions, this new model involving reactions A, C and D can give an expression of the steady-state concentration of \([PL]\). The steady-state assumption yields Eqs (A1 – A5):

\[
\frac{d[P]}{dt} = -k_1[P][D] + k_{-1}[P \cdot D] + k_s[PL] = 0 \quad (A1)
\]

\[
\frac{d[PL]}{dt} = -k_s[PL][D] + k_{-s}[PL \cdot D] - k_1[PL] = 0 \quad (A2)
\]

\[
\frac{d[P \cdot D]}{dt} = k_1[P][D] - k_{-1}[P \cdot D] - k_d[P \cdot D] + k_{-d}[PL \cdot D] = 0 \quad (A3)
\]

\[
\frac{d[PL \cdot D]}{dt} = k_s[PL][D] - k_{-s}[PL \cdot D] + k_d[P \cdot D] - k_{-d}[PL \cdot D] = 0 \quad (A4)
\]

\[
\frac{d[L]}{dt} = k_3[PL] - k_d[L][D] + k_{-d}[D] = 0. \quad (A5)
\]
It is of note that Eqns (A3) and (A4) are identical to Eqns (4) and (5). The total protein concentration is given by:

$$[P_{\text{total}}] = [P] + [PL] + [P \cdot D] + [PL \cdot D].$$

(A5)

Addition of Eqns (A3) and (A4) and substitution into Eqn (A5) yields:

$$[P_{\text{total}}] = [[P] + [PL] + [P \cdot D] + [PL \cdot D] (1 + K_1)].$$  

(A6)

From Eqns (A3) and (A4), Eqn (A7) (which is identical to Eqn 9) can be generated by elimination of $[PL \cdot D]$:

$$[P \cdot D] = \frac{a_d}{k_1} [D][PL] + \left(\frac{a_s}{k_3} + \frac{a_d}{k_3 - k_{-3}}\right) [D][P];$$

$$a_d = \frac{k_{-3}k_1}{k_{-3} + k_3 + k_{-3}}.$$  

(A7)

Substitution of Eqn (A7) into Eqn (A1) yields:

$$[P] = \frac{a_d[D]}{K_{d,0.5}[D]}.$$  

(A8)

Finally substitution of Eqn (A8) into Eqn (A6) yields:

$$[PL] = \frac{K_{d,0.5}[D]}{1 + K_1 [1 + K_1[D]] [D] + \beta};$$

$$\beta = \frac{k_3}{(1 + K_1)a_d}.$$  

(A9)

At $[D] \gg \beta$, Eqn (A9) becomes equal to Eqn (19).

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REFERENCES