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## Regulation of CD4-p56<sup>lck</sup>-associated phosphatidylinositol 3-kinase (PI 3-kinase) and phosphatidylinositol 4-kinase (PI 4-kinase)

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#### SUMMARY

CD4 serves as a receptor for MHC class II antigens and as a receptor for the human immunodeficiency virus (HIV-1) viral coat protein gp120. It is coupled to the protein-tyrosine kinase p56lck, an interaction necessary for an optimal response of certain T cells to antigen. Although anti-CD4 crosslinking may increase lck activity, the effects of HIV-1 gp120 have been controversial. Activated protein-tyrosine kinases are known to associate with certain intracellular proteins possessing src-homology regions (SH-2 domains) such as phosphatidylinositol 3-kinase (PI 3-kinase). In this paper, we demonstrate that the CD4:p56<sup>lck</sup> complex associates with significant amounts of phosphatidylinositol (PI) kinase activity. High pressure liquid chromatographic (HPLC) analysis of the reaction products demonstrated the presence of phosphatidylinositol 3-phosphate (PI 3-P) and phosphatidylinositol 4-phosphate (PI 4-P), thus indicating that PI 3 and PI 4 kinases associate with CD4-p56lck. The p85 subunit of PI 3-kinase was also detected in anti-CD4 immunoprecipitates by immunoblotting with anti-p85 antiserum. Significantly, p56lck binding to CD4 appears to be necessary for the detection of lipid kinase activity associated with p56lck. Also, anti-HIV gp120 and anti-CD4 crosslinking induced a 10-15-fold increase in levels of both PI 3- and PI 4-kinase activity in anti-CD4 precipitates. Stimulation of CD4-p56lck-linked PI kinases by crosslinked HIV-1 gp120 may play a role in HIV-1-induced immune defects.

#### 1. INTRODUCTION

CD4 serves as a receptor for MHC class II antigens and as a receptor for the human immunodeficiency virus (HIV-1) viral coat protein gp120 (Reinherz et al. 1983; Rudd et al. 1989; Littman 1990; Dalgleish et al. 1984; Bedinger et al. 1988). Both CD4 and CD8 are coupled to the src-related protein-tyrosine kinase p56lck (Rudd et al. 1988; Barber et al. 1989; Veillette et al. 1988). The association is required for an optimal response of certain T cells to antigen (Sleckman et al. 1988; Glaichenhaus et al. 1991; Abraham et al. 1991). CD4-p56<sup>lck</sup> complex can also physically associate with the TcR\(\zeta/\text{CD3 complex (Burgess et al. 1991), and as such synergize with the TcRζ/CD3 complex in augmenting T cell proliferation (Emmerich et al. 1986; Eichmann et al. 1987; Anderson et al. 1987). Although anti-CD4 crosslinking may increase lck activity (Veillette et al. 1989; Luo & Sefton 1990), the effects of HIV-1 gp120 have been controversial (Horak et al. 1990; Juszczak et al. 1991; Kaufmann et al. 1992). p56lck may be responsible for the tyrosine phosphorylation of substrates induced by CD4 or TcR/CD3 ligation (Veillette et al. 1989; June et al. 1990; Deans et al. 1992). Potential targets include phospholipase Cy (PLCγ) (Weber et al. 1992), MAP-2 kinase (Ettehadieh et al. 1992); CD5 (Burgess et al. 1992) and p70<sup>zap</sup> (Chan et al. 1992). Identification of downstream targets of the CD4-p56lck complex will be key to unravelling components of the tyrosine phosphorylation in T cell activation.

Phosphatidylinositol 3-kinase (PI 3-kinase) is a potential component of the signalling cascade initiated by CD4-TcRζ/CD3 ligation. PI 3-kinase phosphorylates the D-3 position of the inositol ring of phosphatidylinositol, phosphatidylinositol 4-phosphate and phosphatidylinositol 4,5 bisphosphate (Cantley et al. 1991; Carpenter & Cantley 1990). It is a heterodimeric protein, comprised of a regulatory subunit (p85) and a catalytic subunit (p110) (Carpenter et al. 1990, Escobedo et al. 1991; Skolinik et al. 1991; Otsu et al. 1991; Hiles et al. 1992). p85 is comprised of two Src-homology 2 domains (SH2 domains) which mediate binding to phosphotyrosine residues within

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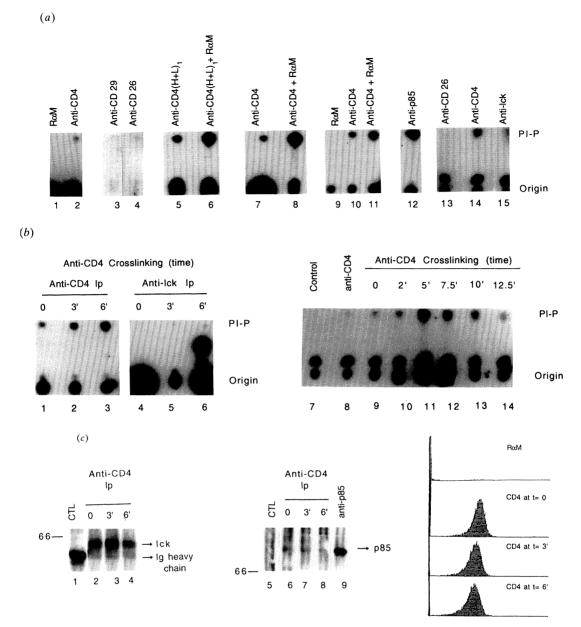


Figure 1. PI kinase activity associated with CD4 is enhanced by antibody induced CD4 crosslinking. TLC analysis of the PI-P formed in anti-CD4 immunoprecipitates is presented. (a) Untreated HPB-ALL cells were lysed in a NP40/digitonin lysis buffer (0.5% each) and subjected to immunoprecipitation with rabbit anti-mouse (lane 1), anti-CD4 (lane 2 and 14), anti CD26 (lane 13) and anti-lck (lane 15). Alternatively, cells were pre-treated with anti-CD29 (lane 3), anti-CD26 (lane 4), univalent anti-CD4 (H+L)<sub>1</sub> (lane 5), univalent anti-CD4 (H+L)<sub>1</sub> plus rabbit anti-mouse (lane 6), bivalent anti-CD4 (lane 7 and 10) or bivalent anti-CD4 plus rabbit anti-mouse (lanes 8 and 11) prior to precipitation, as indicated in the Methods. Addition of anti-CD4 (lanes 7 and 10) for 6 min at 37°C greatly enhanced the levels of precipitable PI kinase activity. Crosslinking of either univalent anti-CD4 (lane 6) or bivalent anti-CD4 (lane 8 and 11) with rabbit anti-mouse increased levels of precipitable activity. Anti-p85 precipitates from cell lysates served as a positive control (lane 12; shorter term exposure of film; see below). (b) Left panel: timecourse of effect of rabbit anti-mouse induced anti-CD4 crosslinking at 37°C on precipitable CD4 associated PI kinase activity. Anti-CD4 precipitates (lanes 1-3); anti-p56lck precipitates from CD4 depleted cell lysates (lanes 4-6). Times of incubation: lanes 1 and 4, 0 min; lanes 2 and 5, 3 min; lanes 3 and 6, 6 min. The intermediate spot in lane 6 is a contaminating hot spot unrelated to the experiment. Right panel: kinetic analysis of the regulation of CD4:p56lck associated PI kinase activity. Lane 7 represents immune complexes obtained from HPB-ALL cells treated with anti-CD29 and crosslinked with rabbit anti-mouse (0 min at 37°C). Cells treated with anti-CD4 alone (0 min at 37°C) (lane 8). Anti-CD4/RaM crosslinked samples correspond to: 0 min (lane 9), 2 min (lane 10), 5 min (lane 11), 7.5 min (lane 12), 10 min (lane 13) and 12.5 min (lane 14). (c) Left panel: anti-lck immunoblotting of CD4 precipitates during the timecourse of CD4 crosslinking with anti-CD4 and rabbit antimouse. Lane 1, rabbit anti-mouse control. Anti-CD4 and rabbit anti-mouse (lanes 2-4). Times of incubation: lane 2, 0 min; lane 3, 3 min; lane 4, 6 min. Middle panel: anti-p85 immunoblotting of CD4 immunoprecipitates during the timecourse of incubation, as in left panel (lanes 5-9). Lane 5, rabbit anti-mouse control. Anti-CD4 plus RaM (lanes 6-8). Times of incubation: lane 6, 0 min; lane 7, 3 min; lane 8, 6 min. Anti-p85 (lane 9). Right panel: flow cytometric analysis of the expression of CD4 receptors after anti-CD4 crosslinking is also shown.

the cytoplasmic tail of receptor tyrosine kinases such as the platelet derived growth factor (PDGF-R) (31–33). p85 also binds to activated pp60src (Fukui & Hanafusa 1989), associated middle T antigen (Talamage et al. 1989) and the insulin receptor substrate IRS-1 (Yonezawa et al. 1992). By contrast, the p110 subunit is related to the yeast protein Vps34 that plays an essential role in protein sorting and transport (Robinson et al. 1988; Herman & Emr 1990). The importance of PI 3-kinase has been shown by its requirement in signalling by PDGF-R (Fantl et al. 1992)

In this study, we demonstrate that the CD4:p56<sup>lck</sup> complex associates with significant amounts PI 3- and PI 4-kinase activity. High pressure liquid chromatographic (HPLC) analysis of the reaction products demonstrated the presence of phosphatidylinositol 3-phosphate (PI 3-P) and phosphatidylinositol 4-phosphate (PI 4-P). Both anti-HIV gp120 and anti-CD4 crosslinking induced a 10–15-fold increase in levels of both PI 3- and PI 4-kinase activity in anti-CD4 precipitates. CD4-p56<sup>lck</sup>–PI 3-kinase binding is likely to constitute a key interaction in HIV-1 gp120 and CD4-TcRζ/CD3 induced T cell function.

#### 2. MATERIALS AND METHODS

#### (a) Antibodies and immunoprecipitation analysis

HPB-ALL cells  $(20 \times 10^6 \text{ ml}^{-1})$  were solubilized in NP-40/Digitonin (0.5% each by volume) in 20 mm Tris-HCL, pH 8.3 containing 150 mm NaCl, 1 mm PMSF (phenyl methylsulphonyl fluoride) and immunoprecipitated with an excess of anti-CD4 antibody (19Thy5D7, 20 μg ml<sup>-1</sup>), as described (Rudd et al. 1988; Barber et al. 1989). Alternatively, the same number of cells in cold RPMI containing 2% (by volume) fetal calf serum (FCS) were incubated with an excess of anti-CD4 antibody, or anti-CD26 (1F7,  $10 \,\mu g \,ml^{-1}$ ) or anti-CD29 (4B4,  $10 \,\mu g \,ml^{-1}$ ) for 30 min at 4°C. Cells were either lysed or further incubated with rabbit anti-mouse antibody (RaM, 6 µg ml<sup>-1</sup>) for 30 min at the same cell density. After washing twice with ice cold media, cells were resuspended in warm RPMI/2% FCS and incubated for various times at 37°C. Activation through receptor crosslinking was arrested by diluting the cells with ice cold RPMI. Cells were centrifuged, washed twice with ice cold RPMI and solubilized in NP-40/Digitonin lysis buffer, as described (Rudd et al. 1988; Barber et al. 1989). The cleared lysate was then incubated with Protein A Sepharose beads for 2 h at 4°C under constant rotation. For a positive control, immunoprecipitations from cell lysates were carried out using anti-p85 rabbit antisera raised against the 85 kDa subunit of PI 3-kinase. Immune complexes were washed three times with PBS (1% NP-40), three times with 100 mm Tris, pH 7.5 with 0.5 m LiCl and twice with TNE (10 mm Tris-Hcl, pH 7.5, 150 mm NaCl and 1 mm EGTA). The lipid kinase reaction was carried out on the beads using phosphatidyl inositol and [32Pγ]-ATP (20 μCi), as described (Whitman et al. 1985; Auger et al. 1989). Lipids were then extracted

using chloroform and methanol (1:1) and separated by thin layer chromatography on a silica gel plate precoated with potassium oxalate using a basic system (chloroform, methanol, water, ammonium hydroxide (60: 47, 11.3, 2) (Whitman *et al.* 1985; Auger *et al.* 1989). The plates were then removed dried, wrapped in a plastic wrap and exposed to a X-ray film, overnight at  $-70^{\circ}$ C, except for lane 12 which is a half-hour exposure.

Cell lysates were further depleted of CD4 by sequential precipitation (three times) using anti-CD4 antibody, followed by two preclearing steps using Protein A Sepharose. Depleted lysates were then subjected to precipitation using an anti-lck sera (figure 1b, left panel, lanes 4-6). The antisera was raised in rabbits against an amino acid terminal peptide (residues 39-64) coupled to key hole limpet haemocyanin (KLH) (Prasad & Rudd 1992). After transfer to nitrocellulose, membranes were blocked with gelatin (2% by mass) in Tris-buffered saline (10 mm Tris-Hcl, pH 8.0 with 150 mm NaCl) and probed with the anti-lck rabbit antisera (1:1000 dilution) (figure 1c, left panel). Immunoblots were visualized using goat anti-rabbit alkaline phosphatase system (Promega). p85 of PI-3 kinase was visualized by immunoblotting with anti-p85 rabbit sera (1:4500) and enhanced by chemiluminescence (ECL) Western blotting (Amersham) (figure 1c, middle panel). Aliquots of the cells at the different time points were pelleted, incubated with goat anti-rabbit-FITC (Fisher) and assayed for CD4 surface expression by flow cytometry using an EPICS cell sorter (Coulter Immunology, Florida) (figure 1c, right panel).

HPB-ALL cells at a density similar to that described for anti-CD4 crosslinking experiment (figure 1) were exposed to a mixture of anti-CD4 antibody (20 μg ml<sup>-1</sup>; 19thy5D7, IgG2A) and rat anti-mouse (0.2 μg ml<sup>-1</sup>, Zymed, California, IgG2a) was added to the cells at 37°C for various times. In contrast to the experiment described in figure 1, cells were shifted to 37°C at the same time as the addition of anti-CD4 and rabbit anti-mouse. In case of control, comparable amounts of anti-CD29 (4B4) and rat anti-mouse were added. Cells were lysed at the indicated times and analysed for CD4-associated PI kinase activity by TLC as described in figure 1. The corresponding PI-P spots were extracted deacylated and subjected to HPLC analysis (Whitman *et al.* 1985; Auger *et al.* 1989).

#### (b) HIV-1 gp120 crosslinking

Conditions of HIV gp120 binding were established as previously described (Kaufmann et al. 1992). gp120 was radiolabelled with  $^{125}\text{I-Bolton}$  Hunter reagent and incubated with  $2\times10^5$  cells for 2 h. Free and bound ligand were separated by centrifugation of cells through silicon oil (specific density: 1.011 g ml¹). The bottom of the 300  $\mu l$  vials (Sarstedt) was cut off for measurement of bound radioligand. Non-specific binding was determined by performing the experiment in the presence of 100 nm soluble CD4. The antigp120 sera blocks the binding of soluble gp120, and binds, but does not dissociate CD4 bound gp120.

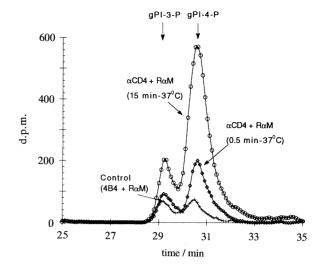
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Computation of the binding parameters (equilibrium dissociation constant  $(K_d)$ , etc.) were determined as described (Kaufmann et al. 1992). For lipid kinase assays, HPB-ALL cells were harvested and suspended at a density of  $20 \times 10^6$  cells ml<sup>-1</sup> in ice cold RPMI (2% by volume FCS) and rotated with recombinant HIV-1 gp120 (ABT, Cambridge) derived from baculovirus Sf9 cell system at 4°C for 2 h (concentration of native pure protein,  $1.0 \times 10^{-8} \,\mathrm{m}$ ;  $K_{\rm d}$ ,  $1.06 \times 10^{-8} \,\mathrm{m}$ ). Based on the measurement of the association kinetics, this procedure results in gp120 binding to 10% of surface CD4 molecules (data not shown). The cells were washed twice with ice cold RPMI (2% FCS) and treated with anti-gp120 rabbit sera (1:100, ABT, Cambridge) for 1 h at 4°C. A 1/100 dilution of rabbit anti-gp120 results in antibody binding to 45-50% of gp120-CD4 complexes on the cell surface (figure 3a). The cells were washed and treated with saturating amounts of goat anti-rabbit antibody (1:200, Sigma) for 0.5 h at 4°C. Cells were then incubated at 37°C for the indicated times. Following cell lysis immunoprecipitations were carried out and subjected to a lipid kinase reaction as described (figure 1). The reaction products were deacylated and analysed by HPLC as described.

#### 3. RESULTS

To determine whether the CD4-p56lck complex can associate with PI kinase, anti-CD4 precipitates from the leukemic T cell line HPB-ALL (figure 1a) were assayed for the ability to generate monophosphorylated phosphoinositides (PIP) from exogenously added lipids. A 5 min incubation of intact HPB-ALL cells with either univalent (H+L)<sub>1</sub> or bivalent anti-CD4 antibodies followed by immunoprecipitation resulted in the detection of significant amounts of PI kinase activity (lanes 5-11). Rabbit anti-mouse (RaM) (lanes 1 and 9), anti-CD29 (4B4) (lane 3) anti-CD26 (1F7) (lanes 4 and 13) failed to co-precipitate enzymic activity, while anti-85 precipitable PI 3-kinase activity served as a positive control (lane 12). Anti-CD4 precipitation from cells that had not been preincubated with anti-CD4 sufficed to precipitate PI kinase activity (lanes 2 and 14). Anti-p56lck serum precipitated small levels of activity (lane 15), although consistently less than that precipitated by anti-CD4 (lane 14). Crosslinking of either univalent anti-CD4 (lane 6) or bivalent anti-CD4 (lanes 8, 11) with RaM increased levels of precipitable activity. Similar results were obtained using other T cell lines and peripheral T cells (data not shown).

We next attempted to determine changes in PI kinase activity over the timecourse of anti-CD4 binding. Anti-CD4 crosslinking has previously been reported to increase the level of p56<sup>lck</sup> phosphorylation within T cells (Veillette *et al.*, 1989; Luo & Sefton 1990). Anti-CD4 crosslinking resulted in a time-dependent increase in precipitable lipid kinase activity (figure 1b, lanes 1–3 and 7–14). Further kinetic analysis showed a transient increase in activity, with maximal precipitable kinase activity at about 5 min, followed by a gradual decrease (lanes 9–14); under



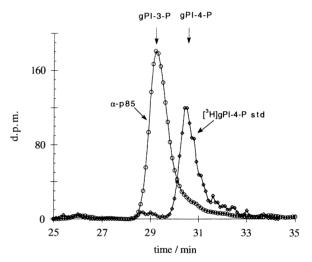
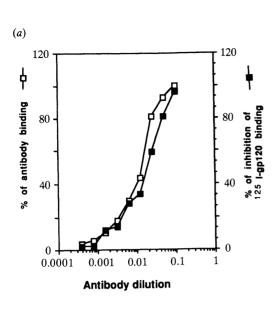
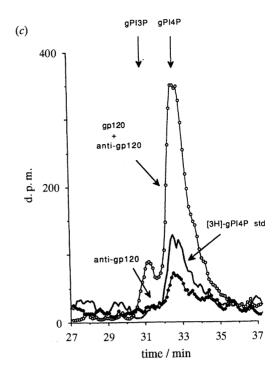


Figure 2. Both PI 3- and PI 4-kinase activities associate with the CD4:p56<sup>lck</sup> complex and are up-regulated upon anti-CD4 antibody crosslinking. An HPLC analysis of the deacy-lated lipids. Upper panel: control (crosses); anti-CD4 and RaM, 0.5 min (diamonds) and 15 min (circles). Lower panel: PI-P analysis corresponding to anti-p85 showed only the presence of PI-3P (circles); the position of <sup>3</sup>H-gPI-4P standard is indicated (diamonds).

some conditions, elevated activity was still detected at 15 min. Under the same conditions, surface CD4 underwent a slight decrease in expression (figure 1c, right panel). Similarly, co-precipitated lck underwent no detectable change as monitored by anti-lck immunoblotting (figure 1c, left panel, lanes 1–4). Anti-p85 PI 3-kinase immunoblotting positively identified co-precipitated p85 associated with CD4-p56<sup>lck</sup> (figure 1c, middle panel, lanes 5–9). The amount of p85 appeared to undergo little change over the time-course; however, given the low amount of detectable material, a conclusive evaluation of this matter was not possible.

The small amount of activity precipitated by the anti-p56<sup>lck</sup> serum (figure 1*a*, lane 15) was readily lost by depletion of lysates using anti-CD4 antibody (figure 1*b*, lanes 4–6). These data indicate that PI kinase preferentially binds to the CD4-p56<sup>lck</sup> complex.





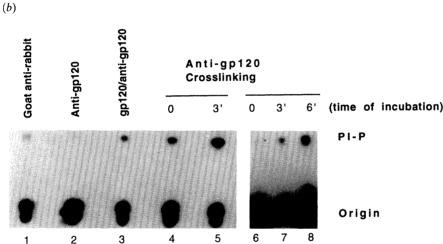


Figure 3. CD4 associated PI kinase activity is also enhanced by HIV-1 anti-gp120-mediated crosslinking of CD4 receptors. (a) Binding analysis of HIV-1 gp120 as detected by anti-gp120. Competition analysis of soluble HIV-gp120 by anti-gp120 antisera. (b) TLC analysis of CD4-precipitable PI kinase activity after gp120 mediated crosslinking. Cells were treated for various periods of time with the following antibodies: goat anti-rabbit (lane 1); rabbit anti-gp120 (lane 2); HIV-1 gp120 and rabbit anti-gp120 (lane 3); HIV-1 gp120, rabbit anti-gp120 and goat anti-rabbit (lanes 4–8). Time of incubation: 0 min (lanes 4 and 6); 3 min (lanes 5 and 7); 6 min (lane 8). Lanes 6–8 are from a separate experiment. (c) HPLC analysis of the deacylated reaction products. PI-P produced by immunocomplexes from cells treated with anti-gp120 alone (filled circles), or HIV-1 gp120, rabbit anti-gp120 plus goat anti-rabbit (open circles) were deacylated and analysed by HPLC; [<sup>3</sup>H]-yPI 4P standard is indicated (dots).

HPLC analysis of the deacylated products of the anti-CD4 associated PI kinase confirmed the presence of PI 3-P and, surprisingly, significant amounts of PI 4-P (figure 2, upper panel). The levels of the two phospholipids varied slightly from experiment to experiment; however, PI 4-P formation exceeded PI 3-P by some 2- to 5-fold. Anti-p85 precipitates showed exclusive labelling of the PI 3-P product (figure 2, lower panel). Crosslinking of CD4-p56lck complexes resulted in time-dependent increase in both PI 3- and PI 4-kinase activities (figure 2, upper panel). Both kinases showed a two- to threefold increase from 0.5 to 15 mm of crosslinking (see legend figure 2).

Of potential importance was the finding that the crosslinking of CD4-p56<sup>lck</sup> by HIV gp120 also increased associated PI kinase activity. Purity of HIV-1 gp120 from Baculovirus Sf9 cells (>80%) and percent native protein (15.7%) were assessed as previously described (Kaufmann et al. 1992). Displacement studies showed gp120 binding to a single binding site with  $K_{\rm d} = 1.06 \times 10^{-8}\,\rm M$  (Kaufmann et al. 1992). Studies were conducted using concentrations of gp120 designed to bind to a relatively low percent (approximately 10%) of CD4 surface receptors. This was followed by exposure of cells to a 1/100 dilution of rabbit anti-gp120 designed to bind 50% of gp120-

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CD4 complexes (figure 3a). The combined exposure to gp120 and rabbit anti-gp120 precipitated moderate levels of PI kinase activity (figure 3b, lane 3). Further crosslinking of the immune complexes with a saturating concentration of goat anti-rabbit serum (1/200 dilution) demonstrated that HIV-1 gp120 aggregation induced a time-dependent increase in precipitable PI kinase activity (figure 3b, lanes 4-8). As controls, neither goat anti-rabbit, nor rabbit anti-gp120 alone precipitated activity (lanes 1 and 2). HPLC analysis of the products revealed the presence of increased levels of PI 3-P and PI 4-P compared with controls (figure 3c). Although previous studies have raised controversy about the effects of HIV-1 gp120-mediated CD4 crosslinking of p56lck activity (Horak et al. 1990; Juszczak et al. 1991; Kaufmann et al. 1992), our data from four experiments indicated that the crosslinking of HIV-1 gp120 has a marked modulatory effect on the activity of PI 3- and PI 4-kinases associated with the receptor.

#### 4. DISCUSSION

CD4-p56 $^{lck}$  associates and synergizes with the TcR $\zeta$ / CD3 complex in the optimal response of T cells to antigen (Rudd 1992). A key issue has been to uncover the downstream events mediated by ligation of the CD4 complex. T cell activation has been reported to generate PI(3,4)P2 and PI(3,4,5)P3, thereby implicating PI 3- and PI 4-kinases in the activation cascade (Ward et al. 1991). In this study, we demonstrate that the CD4:p56<sup>lck</sup> complex associates with significant amounts PI 3- and PI 4-kinase activity, an interaction modulated by anti-HIV gp120 and anti-CD4 crosslinking. Confirmation of the interaction was further made by the detection of p85 by immunoblotting. A previous report by Thompson and co-workers (1992) detected CD4-p56lck associated PI 3-kinase activity; however, others failed to detect activity associated with p56<sup>lck</sup> (Augustine et al. 1991). The basis of this discrepancy may be related to the fact that the PI 3kinase was detected primarily when present in a complex with CD4. Anti-lck precipitates derived from cell lysates depletion of CD4 failed to show detectable PI 3-kinase activity (Thompson et al. 1992). Our studies have also shown that PI 3-kinase binds to the SH3 domain of lck (data not shown). Therefore, it appears that the association of p56<sup>lck</sup> with the CD4 cytoplasmic tail may play a role in facilitating the ability of the SH3 domains to interact with PI 3kinase. Whether this would involve the unfolding of p56<sup>lck</sup> from a restrained C-terminal-SH2 binding remains to be determined.

Surprisingly, unlike previous reports (Augustine et al. 1991; Thompson et al. 1992), PI 4-kinase was also found associated with CD4-p56<sup>lck</sup> at high levels and was regulated by receptor ligation (figures 1 and 2). Although structurally less well characterized, it is a key component in the classical PI pathway, replenishing PI 4,5-P<sub>2</sub> for PI turnover and PIP<sub>3</sub> production. PI 4-kinase associates with the epidermal growth factor receptor (EGF-R) independent of receptor activation and with a region distinct from the kinase domain

(Cochet et al. 1991). If PI 3-kinase associates with internalized vesicles (Kelly et al. 1992), recruitment of PI 4-kinase onto the same vesicles may be necessary for the sustained production of PI 3,4-P<sub>2</sub> and/or PI 3,4,5-P<sub>3</sub>. PI 4-kinase failed to bind the SH2/SH3 domains, and may instead bind to other regions of the kinase, or the CD4 receptor.

The CD4-p56lck-PI 3/PI 4-kinase interaction may influence the pathogenesis of HIV-1 infectivity. Infection is characterized by the impaired function and loss of CD4+ T cells, an event possibly mediated by apoptosis or syncytia (Fauci 1988; Weinhold et al. 1989; Groux et al. 1992). The yeast homolog of the p110 subunit of PI 3-kinase, Vps 34 kinase regulates membrane sorting and morphogenesis (Robinson et al. 1988; Herman & Emr 1990; Hiles et al. 1992). In contrast to CD4-p56lck activity (Horak et al. 1990; Kaufmann et al. 1992) crosslinking by HIV-1 gp120 of a small percent of CD4 molecules on the cell surface (about 5%) was sufficient to allow the detection of a marked increase in associated PI 3- and PI 4-kinase activity. By altering lipid metabolism, these enzymes may facilitate viral entry and inappropriately activate pathways involved in syncytium formation, or be detrimental to T cell signaling. Early activation events have been reported to render T cells susceptible to HIV-1-induced syncytia (Mohagheghpour et al. 1992). Similarly, CD4-p56lck has been reported to inhibit T cell growth (Haughn et al. 1992).

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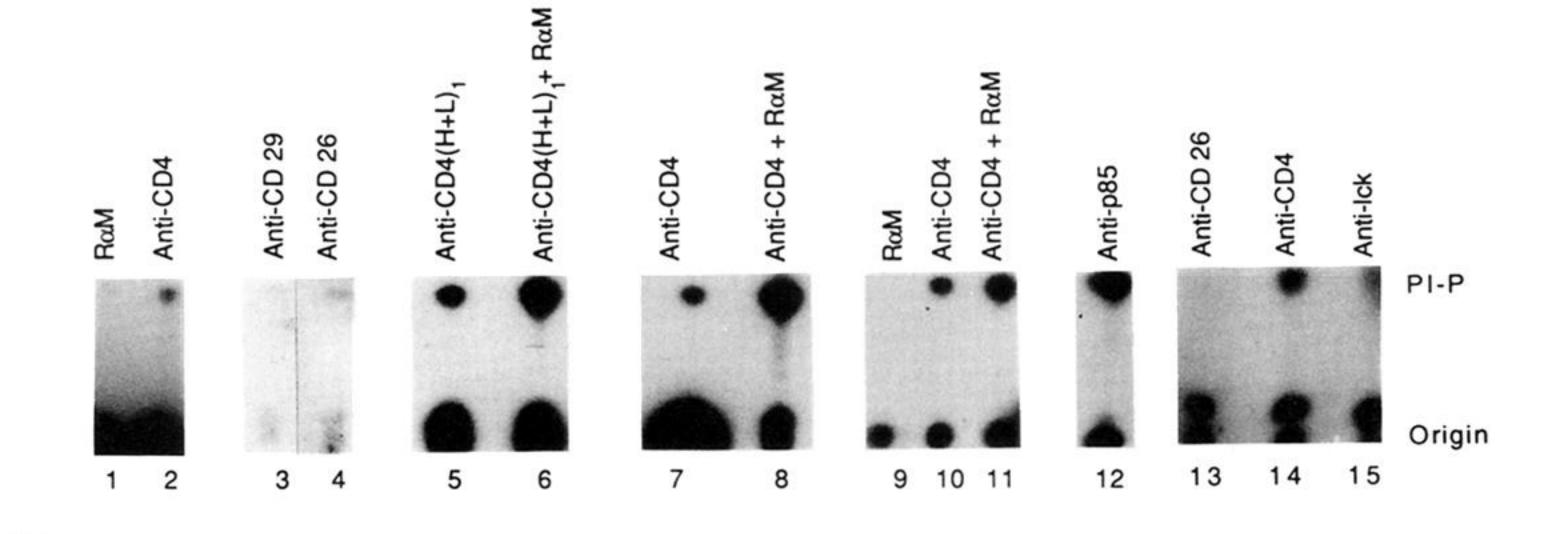
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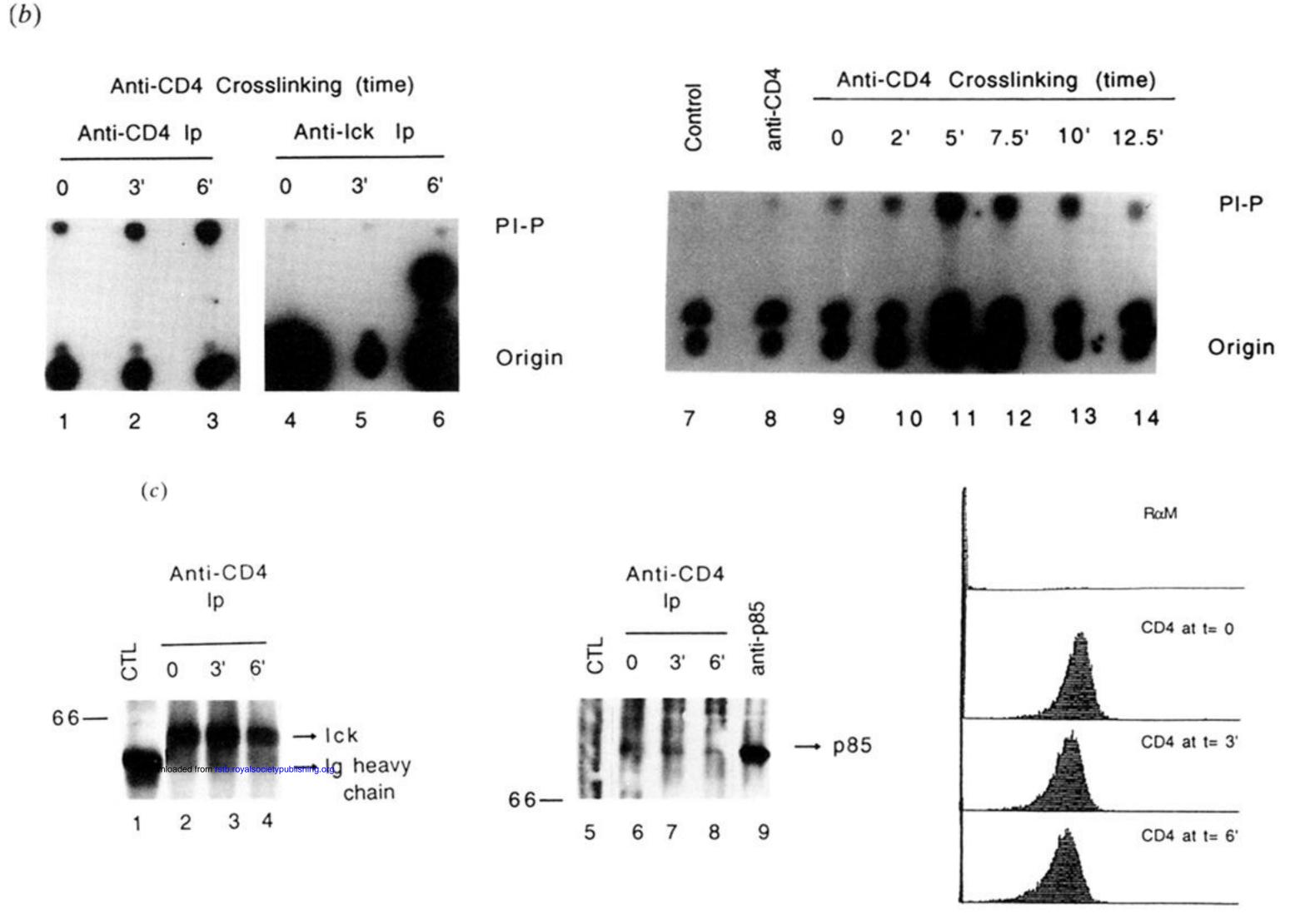
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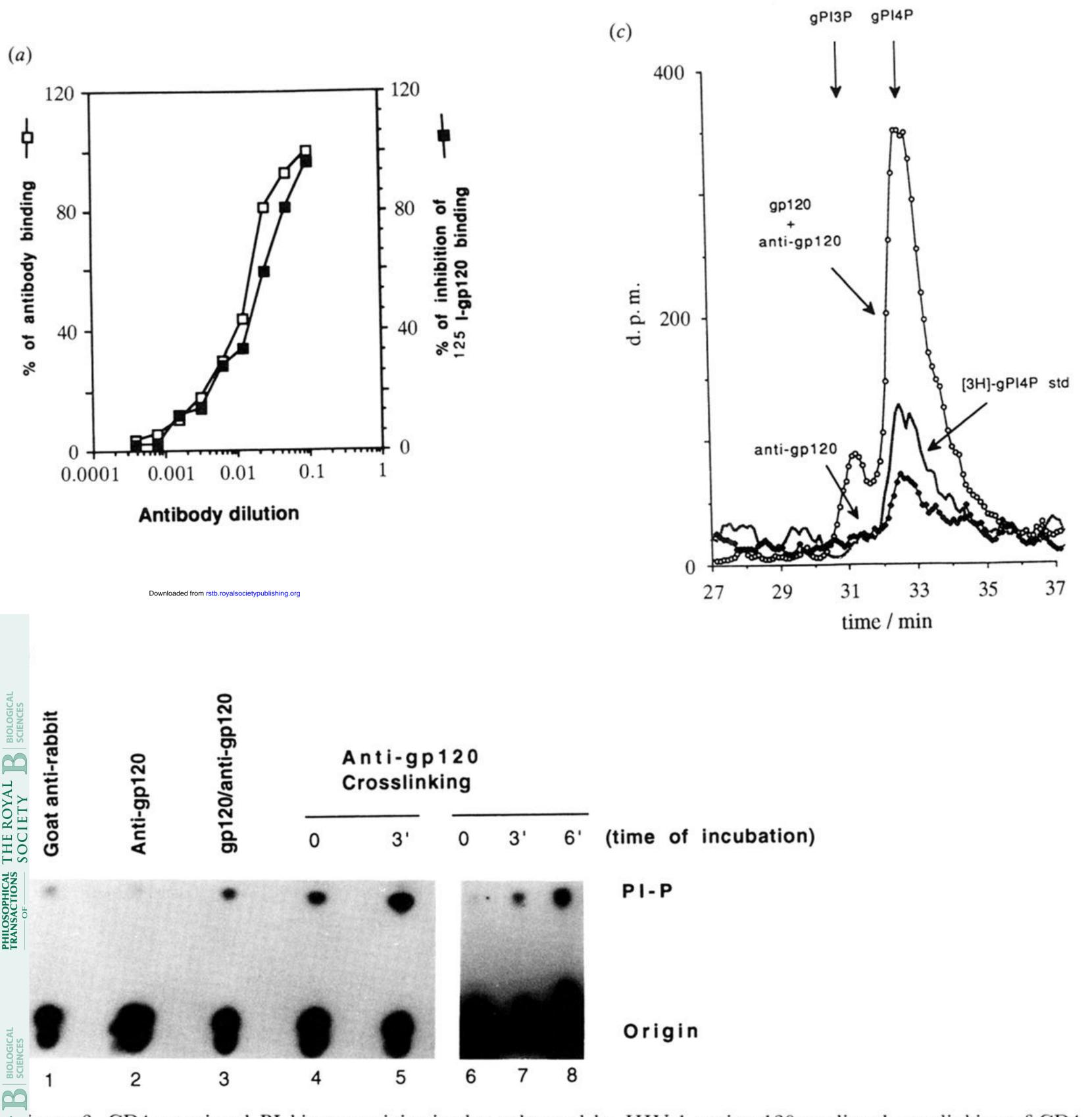
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igure 1. PI kinase activity associated with CD4 is enhanced by antibody induced CD4 crosslinking. TLC analysis f the PI-P formed in anti-CD4 immunoprecipitates is presented. (a) Untreated HPB-ALL cells were lysed in a P40/digitonin lysis buffer (0.5% each) and subjected to immunoprecipitation with rabbit anti-mouse (lane 1), Enti-CD4 (lane 2 and 14), anti CD26 (lane 13) and anti-lck (lane 15). Alternatively, cells were pre-treated with nti-CD29 (lane 3), anti-CD26 (lane 4), univalent anti-CD4 (H+L)<sub>1</sub> (lane 5), univalent anti-CD4 (H+L)<sub>1</sub> plus bbit anti-mouse (lane 6), bivalent anti-CD4 (lane 7 and 10) or bivalent anti-CD4 plus rabbit anti-mouse (lanes 8 nd 11) prior to precipitation, as indicated in the Methods. Addition of anti-CD4 (lanes 7 and 10) for 6 min at 37°C reatly enhanced the levels of precipitable PI kinase activity. Crosslinking of either univalent anti-CD4 (lane 6) or ivalent anti-CD4 (lane 8 and 11) with rabbit anti-mouse increased levels of precipitable activity. Anti-p85 recipitates from cell lysates served as a positive control (lane 12; shorter term exposure of film; see below). (b) Left anel: timecourse of effect of rabbit anti-mouse induced anti-CD4 crosslinking at 37°C on precipitable CD4 ssociated PI kinase activity. Anti-CD4 precipitates (lanes 1-3); anti-p56lck precipitates from CD4 depleted cell rsates (lanes 4-6). Times of incubation: lanes 1 and 4, 0 min; lanes 2 and 5, 3 min; lanes 3 and 6, 6 min. The termediate spot in lane 6 is a contaminating hot spot unrelated to the experiment. Right panel: kinetic analysis of ne regulation of CD4:p56lck associated PI kinase activity. Lane 7 represents immune complexes obtained from IPB-ALL cells treated with anti-CD29 and crosslinked with rabbit anti-mouse (0 min at 37°C). Cells treated with inti-CD4 alone (0 min at 37°C) (lane 8). Anti-CD4/RaM crosslinked samples correspond to: 0 min (lane 9), 2 min ane 10), 5 min (lane 11), 7.5 min (lane 12), 10 min (lane 13) and 12.5 min (lane 14). (c) Left panel: anti-lck munoblotting of CD4 precipitates during the timecourse of CD4 crosslinking with anti-CD4 and rabbit antibouse. Lane 1, rabbit anti-mouse control. Anti-CD4 and rabbit anti-mouse (lanes 2-4). Times of incubation: lane 2, min; lane 3, 3 min; lane 4, 6 min. Middle panel: anti-p85 immunoblotting of CD4 immunoprecipitates during the mecourse of incubation, as in left panel (lanes 5-9). Lane 5, rabbit anti-mouse control. Anti-CD4 plus RaM (lanes -8). Times of incubation: lane 6, 0 min; lane 7, 3 min; lane 8, 6 min. Anti-p85 (lane 9). Right panel: flow ytometric analysis of the expression of CD4 receptors after anti-CD4 crosslinking is also shown.



igure 3. CD4 associated PI kinase activity is also enhanced by HIV-1 anti-gp120-mediated crosslinking of CD4 ceptors. (a) Binding analysis of HIV-1 gp120 as detected by anti-gp120. Competition analysis of soluble HIV-120 by anti-gp120 antisera. (b) TLC analysis of CD4-precipitable PI kinase activity after gp120 mediated cosslinking. Cells were treated for various periods of time with the following antibodies: goat anti-rabbit (lane 1); ibbit anti-gp120 (lane 2); HIV-1 gp120 and rabbit anti-gp120 (lane 3); HIV-1 gp120, rabbit anti-gp120 and goat conti-rabbit (lanes 4–8). Time of incubation: 0 min (lanes 4 and 6); 3 min (lanes 5 and 7); 6 min (lane 8). Lanes 6–8 re from a separate experiment. (c) HPLC analysis of the deacylated reaction products. PI-P produced by nmunocomplexes from cells treated with anti-gp120 alone (filled circles), or HIV-1 gp120, rabbit anti-gp120 plus pat anti-rabbit (open circles) were deacylated and analysed by hplc; [³H]-γPI 4P standard is indicated (dots).