Stromal cell-derived factor-1α stimulates tyrosine phosphorylation of multiple focal adhesion proteins and induces migration of hematopoietic progenitor cells: roles of phosphoinositide-3 kinase and protein kinase C

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The stromal cell-derived factor-1 (SDF-1) is an alpha chemokine that binds to the CXCR4 receptor. Knock-out studies in mice demonstrate that this ligand-receptor pair is essential in hematopoiesis. One function of SDF-1 appears to be the regulation of migration of hematopoietic progenitor cells. We previously characterized signal transduction pathways induced by SDF-1α in human hematopoietic progenitors and found tyrosine phosphorylation of focal adhesion components, including the related adhesion focal tyrosine kinase (RAFTK), the adaptor molecule p130 Cas, and the cytoskeletal protein paxillin. To better understand the functional role of signaling molecules connecting the CXCR4 receptor to the process of hematopoietic migration, we studied SDF-1α-mediated pathways in a model hematopoietic progenitor cell line (CTS), as well as in primary human bone marrow CD34+ cells. We observed that several other focal adhesion components, including focal adhesion kinase (FAK) and the adaptor molecules Crk and Crk-L, are phosphorylated on SDF-1α stimulation. Using a series of specific small molecule inhibitors, both protein kinase C (PKC) and phosphoinositide-3 kinase (PI-3K) appeared to be required for SDF-1α-mediated phosphorylation of focal adhesion proteins and the migration of both CTS and primary marrow CD34+ cells, whereas the mitogen-activated protein kinases ERK-1 and -2 were not. These studies further delineate the molecular pathways mediating hematopoietic progenitor migration and response to an essential chemokine, SDF-1α. (Blood. 2000;95:2505-2513)

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FAK, Crk, and Crk-L are also phosphorylated on SDF-1α stimulation. With respect to the functional roles of signaling molecules, both phosphoinositide-3 kinase (PI-3) and protein kinase C (PKC), but not p44/42 ERK, appeared to be required for the SDF-1α–induced tyrosine phosphorylation of focal adhesion proteins, and for the migration of CTS cells and of human bone marrow CD34+ progenitor cells.

**Materials and methods**

**Reagents and antibodies**

Recombinant SDF-1α was purchased from R&D Systems (Minneapolis, MN). Rabbit anti-RAFTK antibody (R-4250) was generated as described previously.11,12 Anti-FAK polyclonal antibody was generated from New Zealand white rabbits immunized with a bacterially expressed GST protein containing the C-terminal (750-end aa residues) of FAK cDNA, and was tested to react specifically with FAK. Rabbit anti-PLC-γ, and anti-Crk-L polyclonal antibodies were obtained from Santa Cruz Biotechnology, Inc (Santa Cruz, CA). The mouse antiphosphotyrosine monoclonal antibody (mAb) 4G10 was a generous gift from Dr Brian Druker (University of Oregon, Portland, OR), and rabbit anti-PI-3 kinase polyclonal antibodies were purchased from Upstate Biotechnology Co (St Louis, MO). Anti-Crk, anti-p130 Cas, anti-paxillin, and anti-phosphotyrosine mAb (PY20) were purchased from Transduction Laboratories (Lexington, KY). Normal rabbit serum and purified normal rabbit IgG or mouse anti-PLC-γ polyclonal antibodies were generated from New Zealand rabbits immunized with a bacterially expressed GST-fusion protein containing the C-terminal (750-end aa residues) of FAK cDNA, and 50 µg/mL of aprotinin, leupeptin, and pepstatin, 10 mmol/L sodium orthovanadate, 10 µg/mL aprotinin, leupeptin, and pepstatin, 10 mmol/L sodium orthovanadate) and 1 time with phosphate-buffered saline (PBS). Bound proteins were solubilized in 40 µL of 2 × Laemmli sample buffer and further analyzed by immunoblotting. Samples were separated on 8% to 12% SDS-PAGE and then transferred to nitrocellulose membranes. The membranes were blocked with 5% nonfat milk protein and probed with primary antibody for 2 hours at room temperature (RT) or 4°C overnight. Immunoreactive bands were visualized using horseradish peroxidase (HRP)-conjugated secondary antibody and the enhanced chemiluminescent (ECL) system (Amersham Pharmacia). Immunoreactive bands were quantitated by scanning the blot under a Model GS-700 Imaging Densitometer (Bio-Rad). Control lanes were assigned a value of 1, and the quantitations of the immunoreactive bands were used as multiples of the control, based on the densitometry values. Each experiment was repeated at least 3 times and the presented blots are representative of these experiments.

**Assays of phosphoinositide-3 kinase activity**

SDF-1α-stimulated or –unstimulated cells were lysed in ice-cold lysis buffer containing 137 mmol/L NaCl, 20 mmol/L Tris-HCl (pH 7.4), 1 mmol/L MgCl₂, 1 mmol/L sodium orthovanadate, 10% glycerol, 1% NP-40, and 1 mmol/L PMSF. Immunoprecipitation was performed using antiphosphotyrosine mAb (PY20) (for phosphotyrosine-associated PI-3 kinase activity). Immunoprecipitates were washed 3 times with lysis buffer, 3 times with buffer containing 0.1 mol/L Tris-HCl (pH 7.4), 5 mmol/L LiCl, and 0.1 mmol/L sodium orthovanadate, 2 times with TNE buffer containing 10 mmol/L Tris-HCl (pH 7.4), 150 mmol/L NaCl, 5 mmol/L EDTA, and 0.1 mmol/L sodium orthovanadate. Samples were resuspended in 20 µL TNE buffer, 20 µL phosphatase inhibitor cocktail (10 µg; Avanti Polar Lipids, Alabaster, AL), and 10 µL ATP mix (1 mmol/L HEPES, 10 µmol/L ATP, 1 µmol/L MgCl₂, 10 mmol/L Li₃PO₄), then incubated at 37°C for 10 minutes. The reaction was stopped by adding 40 µL of 3 mol/L HCl and 160 µL chloroform:methanol (1:1 vol/vol). Lipids were separated on oxalate-impregnated silica thin-layer chromatography (TLC) plates with a solvent system of chloroform:methanol:water:ammonium hydroxide (28%) (35:35:3.5:7). TLC plates were dried and subjected to autoradiography at ~80°C.

**Preparation of human bone marrow cells**

Light-density bone marrow mononuclear cells were obtained from normal consenting donors and depleted of adherent cells as previously described.

**Ligand stimulation of cells**

CTS cells were starved in serum-free RPMI-1640 medium for 5 hours. During the last hour of starvation, 0.1 mmol/L of sodium orthovanadate was added. After starvation, cells were washed twice with serum-free RPMI-1640 medium and then resuspended at 15 × 10⁶/mL. Cells were then stimulated in vitro with 20 mmol/L SDF-1α for different time periods at 37°C. After stimulation, cell lysates were prepared by lysis in lysis buffer (50 mmol/L HEPES, pH 7.0, 150 mmol/L NaCl, 10% glycerol, 1% Triton-X 100, 1.5 mmol/L MgCl₂, 1 mmol/L EGTA, 10 mmol/L sodium pyrophosphate, 100 mmol/L NaF, 10 mmol/L dithiothreitol, 1 mmol/L PMSF, 10 µg/mL aprotinin, leupeptin and pepstatin, 10 mmol/L sodium orthovanadate). Total cell lysates (TCL) were clarified by centrifugation at 10 000g for 10 minutes. Protein concentrations were determined by protein assay (Bio-Rad Laboratories). To assess the effects of the PI-3 kinase inhibitor wortmannin, the PKC inhibitor GF109203X, or the MEK inhibitor PD98059 (Bio-Rad Laboratories), samples were resuspended in 40 µL of 2 × Laemmli sample buffer and further analyzed by immunoblotting. Samples were separated on 8% to 12% SDS-PAGE and then transferred to nitrocellulose membranes. The membranes were blocked with 5% nonfat milk protein and probed with primary antibody for 2 hours at room temperature (RT) or 4°C overnight. Immunoreactive bands were visualized using horseradish peroxidase (HRP)-conjugated secondary antibody and the enhanced chemiluminescent (ECL) system (Amersham Pharmacia). Immunoreactive bands were quantitated by scanning the blot under a Model GS-700 Imaging Densitometer (Bio-Rad). Control lanes were assigned a value of 1, and the quantitations of the immunoreactive bands were used as multiples of the control, based on the densitometry values. Each experiment was repeated at least 3 times and the presented blots are representative of these experiments.

**Cell separation or isolation by immunomagnetic beads**

Nonadherent light-density bone marrow cells were resuspended in ice-cold PBS with 1% FCS and 0.5% BSA (isolation medium). CD34+ cells were selected by the Dynal CD34 Progenitor Cell Selection System (Dynal beads® M-450 CD34 and DETACHaBEAD® CD34, Dynal Inc, Lake Success, NY) according to the manufacturer’s instructions. The beads binding to the cells were detached and the cells were washed with and resuspended in a medium for the migration assays (see below). The purity of the CD34+ cells selected by this method was found to be more than 95%.

**Chemotaxis assays**

Chemotaxis assays were performed in triplicate using 5-µm pore filters (Transwell, 24-well cell clusters; Costar, Boston, MA) as described previously.11 Briefly, the filters were rinsed with migration medium (RPMI-1640 with 0.5% BSA for the CTS cells; complete a-medium with 0.5% BSA for the CD34+ bone marrow cells) and the supernatant was aspirated immediately before loading the cells. Either 2 × 10⁵ CTS cells or
1.5 × 10^5 CD34+ cells suspended in 100 µL migration medium were loaded into each Transwell filter. Filters were then carefully transferred to another well containing 650 µL migration medium with 20 nmol/L SDF-1α (R&D Systems). The plates were incubated at 37°C in 5% CO₂ for 3.5 hours. Next, the upper chambers were carefully removed and the cells in the bottom chambers were collected. The cells were washed and resuspended in proper volume and quantitated for viable cells using the trypan blue exclusion method. To assess the effects of wortmannin, GF109203X, or PD98059, the cells were pre-incubated with various concentrations of these inhibitors for 45 minutes, and then the chemotaxic assays were performed, as described above.

Results

Stromal cell-derived factor-1α stimulation induces activation of phosphoinositide-3 kinase and the tyrosine phosphorylation of PLC-γ in CTS cells

Cytokine or chemokine-induced cell migration is a complex process, mediated by multiple signaling mechanisms. PI-3 kinase, PKC, or MAPK pathways have been reported to be involved in this process, mediated by multiple signaling mechanisms. PI-3 kinase, PKC, or MAPK pathways have been reported to be involved in cytokine or chemokine-induced migration in various cell types.34-39 We assessed whether these signaling pathways were involved in cytokine or chemokine-induced cell migration is a complex process, mediated by multiple signaling mechanisms. PI-3 kinase, PKC, or MAPK pathways have been reported to be involved in cytokine or chemokine-induced migration in various cell types.34-39 We assessed whether these signaling pathways were involved in cytokine or chemokine-induced migration in hematopoietic progenitor cells.

Using CXCR4 receptor-transfected L1.2 mouse pre-B cells, we previously showed that SDF-1α stimulation via the CXCR4 receptor selectively activated p44/42 ERK kinase, but not p38 or JNK kinase.36 In CTS cells, we previously observed a similar activation of p44/42 ERK by SDF-1α. Here, we examined the effects of SDF-1α on PI-3 kinase activity and on the tyrosine phosphorylation of PLC-γ in CTS cells.

SDF-1α stimulation of CTS cells induced tyrosine phosphorylation of the p85 PI-3 kinase subunit as detected by immunoblotting of anti-phosphotyrosine (PY) immunoprecipitates with anti-p85 antibody (Figure 1A). To determine whether SDF-1α stimulated PI-3 kinase activity, cell lysates from SDF-1α-stimulated or unstimulated CTS cells were immunoprecipitated with anti-PY antibody. The tyrosine phosphorylation-associated PI-3 kinase activity was measured by an in vitro PI-3 kinase assay. As shown in Figure 1B, PI-3 kinase activity was enhanced by SDF-1α stimulation in a time course that paralleled p85 tyrosine phosphorylation.

PLC-γ has been shown to play an important role in G-protein coupled receptor signaling and is an upstream mediator for PKC activation.40,41 We thus investigated if PLC-γ was tyrosine-phosphorylated after SDF-1α stimulation of CTS cells. Cell lysates from SDF-1α-stimulated or unstimulated CTS cells were immunoprecipitated with anti-PLC-γ antibody and subjected to serial immunoblotting with anti-PY antibody and anti-PLC-γ antibody. We observed that PLC-γ was significantly tyrosine-phosphorylated on SDF-1α stimulation (Figure 2, upper panel). The tyrosine phosphorylation of PLC-γ was rapid and transient, reached a maximum at 1 minute, and returned to a basal level at 5 to 10 minutes after stimulation. Equal amounts of PLC-γ were present in each lane (Figure 2, lower panel). The tyrosine phosphorylation of PLC-γ in response to SDF-1α stimulation suggested that the PKC pathway may be involved in CXCR4 receptor signaling.
as 1 minute after the addition of 20 nmol/L of SDF-1. Thereafter, tyrosine phosphorylation of FAK declined gradually to almost baseline levels after 5 minutes (Figure 3, upper panel). We verified that similar amounts of FAK were recovered from the lysates of cells untreated or treated with SDF-1α for various times (Figure 3, lower panel).

Tyrosine-phosphorylated FAK can form a complex with other focal adhesion proteins. The adaptor protein, p130 Cas, has been shown to associate with FAK and mediate cell migration.26 The cytoskeletal protein, paxillin, is also a substrate of FAK.42 Our previous study showed that, in CTS cells, both p130 Cas and paxillin were tyrosine phosphorylated on SDF-1α stimulation.18 Therefore, we now investigated if FAK associated with either p130 Cas or paxillin. Serum-starved CTS cells were stimulated with SDF-1α for various times. Cell lysates from unstimulated or SDF-1α-stimulated CTS cells were immunoprecipitated with anti-FAK polyclonal antibody. The immunoprecipitates were resolved on SDS-PAGE gels, followed by immunoblotting with anti-p130 Cas antibody (Figure 4A, upper panel) or anti-paxillin antibody (Figure 4B, upper panel). SDF-1α stimulation enhanced the association of FAK both with p130 Cas (Figure 4A, upper panel) and with paxillin (Figure 4B, upper panel). Similar amounts of FAK were recovered from the lysates of cells untreated or treated with SDF-1α (Figure 4A and B, lower panels).

These results indicate that SDF-1α stimulation induces the tyrosine phosphorylation of FAK and its association with p130 Cas and paxillin in hematopoietic progenitor cells.

Stromal cell-derived factor-1α induces the tyrosine phosphorylation of Crk and Crk-L and enhances their association with p130 Cas

Crk and Crk-L are structurally similar, but encoded by separate genes.43 Both Crk and Crk-L are composed of 1 SH2 and 2 SH3 domains. Crk and Crk-L share overall 60% amino acid similarity and their SH2 and SH3 domains are highly conserved.43 Crk and Crk-L have been shown to function as adaptor proteins, linking different proteins in signaling.44 Thus, we sought to determine whether the adaptor protein Crk or Crk-L is tyrosine phosphorylated in CTS cells on SDF-1α stimulation. Cell lysates from unstimulated or SDF-1α-stimulated CTS cells were immunoprecipitated with anti-Crk or anti-Crk-L, followed by immunoblotting with anti-PY antibody. As shown in Figure 5E and F, SDF-1α stimulation resulted in the tyrosine phosphorylation of both Crk (Figure 5E) and Crk-L (Figure 5F). Moreover, we observed that several proteins which communoprecipitated with Crk or Crk-L were also tyrosine phosphorylated. We verified that similar amounts of Crk (Figure 5E, lower panel) or Crk-L (Figure 5F, lower panel) were recovered from the lysates of cells untreated or treated with SDF-1α.

Both Crk and Crk-L have been reported to associate with p130 Cas.25,45,46 Thus, we investigated if the tyrosine-phosphorylated proteins that communoprecipitated with Crk or Crk-L included p130 Cas. Cell lysates from unstimulated or SDF-1α-stimulated CTS cells were immunoprecipitated with anti-Crk (Figure 6A) or anti-Crk-L (Figure 6B), and then immunoblotted with p130 Cas. We observed a constitutive association between p130 Cas and Crk-L but not between p130 Cas and Crk in the unstimulated CTS cells. However, SDF-1α stimulation enhanced the association of both Crk and Crk-L with p130 Cas (Figure 6A and B, upper panels). We verified that similar amounts of Crk (Figure 6A, lower panel) or Crk-L (Figure 6B, lower panel) were recovered from the lysates of cells untreated or treated with SDF-1α.
These results indicate that both Crk and Crk-L are involved in CXCR4 receptor signaling in hematopoietic progenitor cells. SDF-1α stimulated the tyrosine phosphorylation of both Crk and Crk-L and induced their association with p130 Cas.

Effects of signaling inhibitors on stromal cell-derived factor-1α–induced tyrosine phosphorylation of various adhesion proteins in CTS cells

To assess the functional roles of PI-3 kinase, PKC, and p44/42 ERK, before stimulation with SDF-1α, CTS cells were pretreated with 100 nmol/L wortmannin, a PI-3 kinase inhibitor; 3 μmol/L GF109203X, a PKC inhibitor; or 20 μmol/L PD98059, a MEK kinase inhibitor, respectively. Changes in the tyrosine phosphorylation of focal adhesion proteins on inhibitor treatment were investigated by immunoprecipitation with specific antibodies for these proteins and by immunoblotting with anti-PY antibody. As shown in Figure 5, the SDF-1α–induced tyrosine phosphorylation of FAK (A, upper panel), RAFTK (B, upper panel), p130 Cas (C, upper panel), paxillin (D, upper panel), Crk (E, upper panel), or Crk-L (F, upper panel) was significantly reduced by pretreatment with wortmannin or GF109203X but not PD98059. These results indicate that the tyrosine phosphorylation of these focal adhesion molecules was dependent on the activation of PI-3 kinase or PKC but not on that of MEK/p44/42 ERK. When the protein loading controls in each immunoblot were examined by reprobing with the same antibody used for the immunoprecipitation, we observed that the SDF-1α–treated sample showed an apparent decrease in p130 Cas protein level, along with an increase in the tyrosine phosphorylation of this protein (Figure 5C, lower panel). However, similar amounts of protein were recovered in each lane of the other immunoblots (Figure 5A, B, D, E, and F, lower panels).

Inhibition of phosphoinositide-3 kinase or protein kinase C, but not MEK, inhibits stromal cell-derived factor-1α–induced migration of CTS cells or primary bone marrow CD34⁺ progenitor cells. Activation of PI-3 kinase, PKC, and p44/42 ERK after SDF-1α treatment indicated that CXCR4 signaling involves multiple pathways. To determine the functional role of PI-3 kinase, p130 Cas was analyzed after treatment with SDF-1α and various inhibitors. As shown in Figure 6A, the SDF-1α–induced migration of CTS cells was inhibited by pretreatment with wortmannin (Wort.), GF109203X (GF), or PD98059 (PD) but not with PI-3 kinase or PKC inhibitors, indicating that CXCR4 signaling involves multiple pathways. To determine the functional role of PI-3 kinase, p130 Cas was analyzed after treatment with SDF-1α and various inhibitors. As shown in Figure 6A, the SDF-1α–induced migration of CTS cells was inhibited by pretreatment with wortmannin (Wort.), GF109203X (GF), or PD98059 (PD) but not with PI-3 kinase or PKC inhibitors, indicating that CXCR4 signaling involves multiple pathways.
PKC or ERK in SDF-1α–induced migration of hematopoietic progenitor cells, CTS cells were pretreated with different concentrations of wortmannin (Figure 7A), GF109203X (Figure 7B), or PD98059 (Figure 7C). Cell migration in response to SDF-1α was examined by a Transwell migration assay as described in “Materials and methods.” We observed that treatment with wortmannin (Figure 7A) or GF109203X (Figure 7B) significantly inhibited SDF-1α–induced migration in a dose-dependent manner. However, treatment with PD98059 over a concentration range of 1 to 40 µmol/L did not alter cell migration (Figure 7C). Because the specificity of wortmannin, at low doses, was recently questioned,47 we also tested the ability of LY294002,48 a competitive inhibitor of PI-3 kinase, to confirm this observed effect. Similar to the results with wortmannin, LY294002 treatment inhibited cell migration in a dose-dependent manner (data not shown).

We next examined the effects of wortmannin, GF109203X, or PD98059 on the migration of primary bone marrow CD34+ cells in response to SDF-1α. CD34+ cells, isolated from normal human bone marrow, were pretreated with each inhibitor and tested in a Transwell migration assay. As shown in Figure 8, similar to that observed in CTS cells, treatment with wortmannin or GF109203X, but not PD98059, inhibited the SDF-1α–induced migration of CD34+ progenitor cells in response to SDF-1α.

**Discussion**

The molecular mechanisms that regulate hematopoietic progenitor cell migration are not well characterized. On the basis of studies in mice null for either the CXCR4 receptor or its cognate ligand SDF-1, it appears that this chemokine is a major and essential physiological regulator of effective hematopoiesis, because such mice have an absence of both lymphoid and myeloid hematopoiesis.15,16 Moreover, in vitro studies with human progenitors have demonstrated a potent chemotactic effect of SDF-1.10-12 With this background, we sought to characterize the signaling pathways triggered after CXCR4 activation by SDF-1α, first in a model hematopoietic cell line, CTS, and then using primary bone marrow CD34+ cells.

Our studies focused on components of the focal adhesion complex. Focal adhesions are important structures that mediate cell adhesion and migration. They consist of a constellation of signaling molecules and cytoskeletal proteins and are believed to be essential in migration.20,49,50 Our previous studies demonstrated that SDF-1α induced calcium flux in CTS cells, followed by the tyrosine phosphorylation of RAFTK, paxillin, and p130 Cas.51 Calcium flux is the signature initial change in G-protein coupled receptors. RAFTK and the related FAK are well-recognized platform kinases that have multiple binding motifs for a number of different signaling molecules and act as putative bridges to transmit signals to the cytoskeleton.51,52 RAFTK and FAK are highly homologous, sharing an overall 45% amino acid sequence identity, with 60% identity in the catalytic domain. Several tyrosine residues appear to
be conserved between RAFTK and FAK, including an Src family tyrosine kinase SH2-binding site. Both RAFTK and FAK contain proline-rich motifs capable of SH3 domain interaction.32,53 Given the high degree of structural and amino acid sequence similarity, RAFTK and FAK may have some similar or even exchangeable functions. Recently, it has been shown that RAFTK can phosphorylate FAK and functions as an agonist-dependent regulator of FAK.54 FAK appears to play a more important role than RAFTK in regulating cell migration.55 Our studies demonstrated that both FAK and RAFTK were phosphorylated on SDF-1α treatment of CTS cells, and this phosphorylation was followed by their association with 2 other important focal adhesion molecules, p130 Cas and paxillin (Dutt et al18 and our unpublished data). These results indicated that both RAFTK and FAK are involved in CXCR4 signaling induced by SDF-1α in hematopoietic progenitor cells.

With the use of CXCR4-transfected L1.2 cells, our previous study demonstrated that SDF-1α stimulation induced the tyrosine phosphorylation of Crk and its association with paxillin and RAFTK.36 In the current study, we observed that not only Crk but also Crk-L was tyrosine phosphorylated in CTS cells on SDF-1α stimulation. Crk-L has a high homology to Crk and contains a similar domain structure, SH2-SH3-SH3.51,14 Like Crk, Crk-L has been shown to be involved in focal adhesion formation by interacting with several focal adhesion proteins, including p130 Cas, paxillin, Abl, and Bcr-Ab1.44 Crk-L is most abundantly expressed in hematopoietic cells,56 and regulates integrin-mediated cell adhesion.57,58 Although Crk-L has similar in vitro binding characteristics with Crk,43,44 there are a few differences to be noted.45,59 One example of these differences is the observation that Crk-L but not Crk constitutively associates with p130 Cas in Bcr-Ab1-transformed leukemia cells.55 Consistent with this observation, we demonstrated that there was a constitutive association of Crk-L, but not Crk, with p130 Cas in CTS cells. However, SDF-1α stimulation significantly enhanced the association of both Crk and Crk-L with p130 Cas. Our results indicated that both Crk and Crk-L, through their interaction with p130 Cas, were involved in CXCR4 signaling.

p130 Cas was initially identified as a prominent tyrosine-phosphorylated substrate of the oncoproteins v-src and v-ckr.60 p130 Cas contains an SH3 domain and numerous potential SH2 domain docking sites, and also serves as a docking protein for the recruitment of proteins involved in protein-tyrosine kinase-mediated signaling pathways. p130 Cas localizes to focal adhesions and interacts with other focal adhesion components.5 It has been shown that the association of p130 Cas with FAK26 or Crk25 is critical for cell migration. This study, combined with our previous studies, demonstrate that SDF-1α stimulates the tyrosine phosphorylation of p130 Cas and enhances its association with focal adhesion proteins, including RAFTK, FAK, Crk, and Crk-L. These results indicate that p130 Cas may play an important role in CXCR4 signaling and cell migration. The apparent decrease in the protein levels of p130 Cas after SDF-1α stimulation (Figure 5C, lower panel) could be due to p130 Cas redistribution to actin-rich cytoskeletal complexes in the Triton-insoluble fractions on its phosphorylation, as reported previously.60,62,63 However, further study is needed to elucidate the functional significance of p130 Cas redistribution in CXCR4 signaling on SDF-1α stimulation.

Our data indicate that multiple adhesion proteins participated in SDF-1α induced signaling in hematopoietic progenitor cells. The changes in phosphorylation and interaction of these structural components may be important for the cell adhesion and migration mediated by SDF-1α in hematopoietic cells. Thus, we sought to define which upstream signaling components may be critical for these observed changes and the related cell migration induced by SDF-1α.

Accumulating data have implicated that multiple signaling mechanisms exist to regulate cell migration. MAP kinase (p44/42 ERK),38,39 PI-3 kinase,34-36 and PKC37 signaling pathways have been shown to regulate the cell migration induced by chemokines or cytokines. Consistent with our previous observation in CXCR4-transfected L1.2 cells,36 we found that SDF-1α activated PI-3 kinase in CTS cells (Figure 1). Additionally, we observed that SDF-1α significantly induced the tyrosine phosphorylation of PLC-γ (Figure 2). Phosphorylated PLC-γ hydrolyzes phosphatidylinositol diphosphate to the second-messenger molecules IP3 and DAG, which in turn activate PKC.30,41 Our previous study demonstrated that SDF-1α also stimulated p44/42 ERK activity in CTS cells.18 To investigate the functional roles of PI-3 kinase, PKC, and p44/42 ERK in the SDF-1α–induced phosphorylation of focal adhesion components and cell migration, we used a series of compounds that have been used as specific inhibitors for PI-3 kinase (wortmannin or LY294002), for PKC (GF109203X), or for ERK (PD98059, MEK inhibitor). Our results demonstrated that inhibition of PI-3 kinase or PKC significantly decreased the tyrosine phosphorylation of various focal adhesion proteins in response to SDF-1α stimulation, including FAK, RAFTK, p130 Cas, paxillin, Crk, and Crk-L. Inhibition of PI-3 kinase or PKC also decreased cell migration in response to SDF-1α, both in CTS and primary bone marrow CD34+ cells. These results suggest that PI-3 kinase and PKC are both required for the SDF-1α–induced cell migration.

Although activation of the p44/42 ERK signaling pathway has been shown to promote cell motility either by regulating gene expression or by directly activating myosin light chain kinase,39 this was not the case in SDF-1α–induced cell migration. Our studies using the MEK inhibitor PD98059 demonstrated that inhibition of MEK kinase upstream of p44/42 ERK did not inhibit SDF-1α–induced migration in CTS cells or CD34+ bone marrow progenitors. Furthermore, treatment with PD98059 could not interfere with the SDF-1α–induced tyrosine phosphorylation of various focal adhesion proteins (Figure 5). These results imply that the ERK signaling pathway is not involved in the focal adhesion formation and migration induced by SDF-1α in hematopoietic progenitor cells.

In summary, our experiments, first done in the CTS cell line and then confirmed in primary bone marrow CD34+ cells, indicate that both PI-3 kinase and PKC are functionally important in inducing SDF-1α–mediated progenitor cell migration, whereas p44/42 ERK is not. The CXCR4 receptor has also been characterized as the primary coreceptor for certain strains of HIV, and binds the envelope glycoproteins gp120/160 with high affinity. Furthermore, such binding results in the activation of certain downstream signaling molecules, including RAFTK/Pyk2.64,65 With better characterization of these signal transduction pathways triggered by SDF-1 in hematopoietic progenitor cells, we now can study in the context of HIV infection whether such pathways are conserved or whether dysregulated hematopoiesis might occur because of aberrant signaling on gp120/160 binding to CXCR4.

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References


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