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Research Article

AIDS/HIV

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Autophagy is involved in T cell death after binding of HIV-1 envelope proteins to CXCR4

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HIV-1 envelope glycoproteins (Env), expressed at the cell surface, induce apoptosis of uninfected CD4⁺ T cells, contributing to the development of AIDS. Here we demonstrate that, independently of HIV replication, transfected or HIV-infected cells that express Env induced autophagy and accumulation of Beclin 1 in uninfected CD4⁺ T lymphocytes via CXCR4. The same phenomena occurred in a T cell line and in transfected HEK.293 cells that expressed both wild-type CXCR4 and a truncated form of CD4 that is unable to bind the lymphocyte-specific protein kinase Lck. Env-mediated autophagy is required to trigger CD4⁺ T cell apoptosis since blockade of autophagy at different steps, by either drugs (3-methyladenine and bafilomycin A1) or siRNAs specific for *Beclin 1/Atg6* and *Atg7* genes, totally inhibited the apoptotic process. Furthermore, CD4⁺ T cells still underwent Env-mediated cell death with autophagic features when apoptosis was inhibited. These results suggest that HIV-infected cells can induce autophagy in bystander CD4⁺ T lymphocytes through contact of Env with CXCR4, leading to apoptotic cell death, a mechanism most likely contributing to immunodeficiency.

Introduction

The development of AIDS in HIV-infected patients is characterized by a progressive decline in the number of CD4⁺ T cells. Although the multiple mechanisms that lead to this demise of CD4⁺ T cells have yet to be fully elucidated, virus-induced programmed cell death (PCD) of uninfected CD4⁺ T lymphocytes clearly contributes to this phenomenon (1–3). In particular, bystander CD4⁺ T cell death induced by HIV-1 envelope glycoproteins (Env) gp120 and gp141 expressed on infected cells is one of the processes linked to immunodeficiency. These contacts between infected and uninfected cells, which occur in epithelia and lymphoid tissues, are important for HIV transmission (4–7). We and others have demonstrated that cells that express CXCR4-using HIV-1 isolates (X4) Env (either Env-transfected cells; refs. 8, 9; or infected cells; ref. 10) trigger uninfected CD4⁺ T cell apoptosis. The CXCR4/Env interaction plays a central role in this phenomenon by inducing an apoptotic pathway that involves mitochondrial transmembrane depolarization and caspase-3 activation independently of Fas-mediated signaling and G_i protein activation (9). This mechanism may explain the T cell drop that often coincides with the emergence of X4 isolates during the later stages of the disease. As CXCR4 is primarily expressed on cells with a resting naive phenotype, X4 variants were therefore suggested to exercise their harmful effects through enhanced killing of uninfected naive CD4⁺ T cells (11). Massive elimination of naive CXCR4⁺ CD4⁺ T cells has also been observed in highly pathogenic simian/human immunodeficiency virus-infected rhesus monkeys (12).

Although PCD can manifest itself in many forms, 2 major types have been described: type I PCD, or apoptosis, and type II PCD, also called autophagic cell death, which is characterized by the

accumulation of autophagic vacuoles and the absence of caspase activation (13–15). However, cell death – characterized by hallmarks of both type I PCD and type II PCD – is frequently observed, making a clear-cut difference between them difficult (16, 17).

Macroautophagy, herein referred to as autophagy, is a major lysosomal catabolic pathway for cytoplasmic macromolecules and organelles (18). This process is characterized by the formation of double or multiple membrane-bound autophagosomes that engulf cytoplasmic material. After maturation, autophagosomes fuse with lysosomes to digest the sequestered material. Autophagy, first discovered to occur in yeast, is a highly regulated physiological process mediated by 2 conjugation systems related to ubiquitylation. Both systems depend on Atg proteins and their related signaling pathways (19, 20). In the first conjugation system, Atg12 is activated by the E1-like enzyme Atg7, transferred to the E2-like enzyme Atg10, and finally conjugated to Atg5 to form an autophagosomal precursor (21). In the second conjugation system, Atg8 is activated by Atg7, transferred to Atg3, another E2-like enzyme, and finally conjugated to phosphatidyl ethanolamine. Thus, Atg7 is essential for activation of these 2 conjugation systems and, consequently, for the autophagic process (21). Microtubule-associated protein 1 light chain 3 (LC3) is the mammalian ortholog of Atg8 and was the first identified on the autophagosome membrane (22). Class III PI3K is involved in the early stages of autophagic vesicle formation and controls the autophagic pathway through its association with Beclin 1, the ortholog of yeast Atg6 (23, 24).

Autophagy is a cell survival mechanism during nutrient starvation (25, 26) that supplies vital components until conditions improve. Moreover, starvation-induced autophagy protects cells from the induction of apoptosis (27). However, several recent studies have shown that cell death can be blocked after knockdown of autophagy genes (28–30). These results give credence to the role of autophagy as a cell death process (for review see refs. 31–33). Recent data also indicate that a connection exists between pathways that regulate autophagy and apoptosis. Indeed, several molecules such as Bcl-2 family proteins (28, 29,

Nonstandard abbreviations used: Baf A1, bafilomycin A1; Env, HIV-1 envelope glycoproteins; GFP-LC3, LC3 fused to GFP; LC3, light chain 3; 3-MA, 3-methyladenine; PCD, programmed cell death; PI, propidium iodide; Rapa, rapamycin; TEM, transmission electron microscopy; UC, umbilical cord; X4, CXCR4-using HIV-1 isolates; z-VAD, z-Val-Ala-Asp (OMe)-CH₂F.

Conflict of interest: The authors have declared that no conflict of interest exists.

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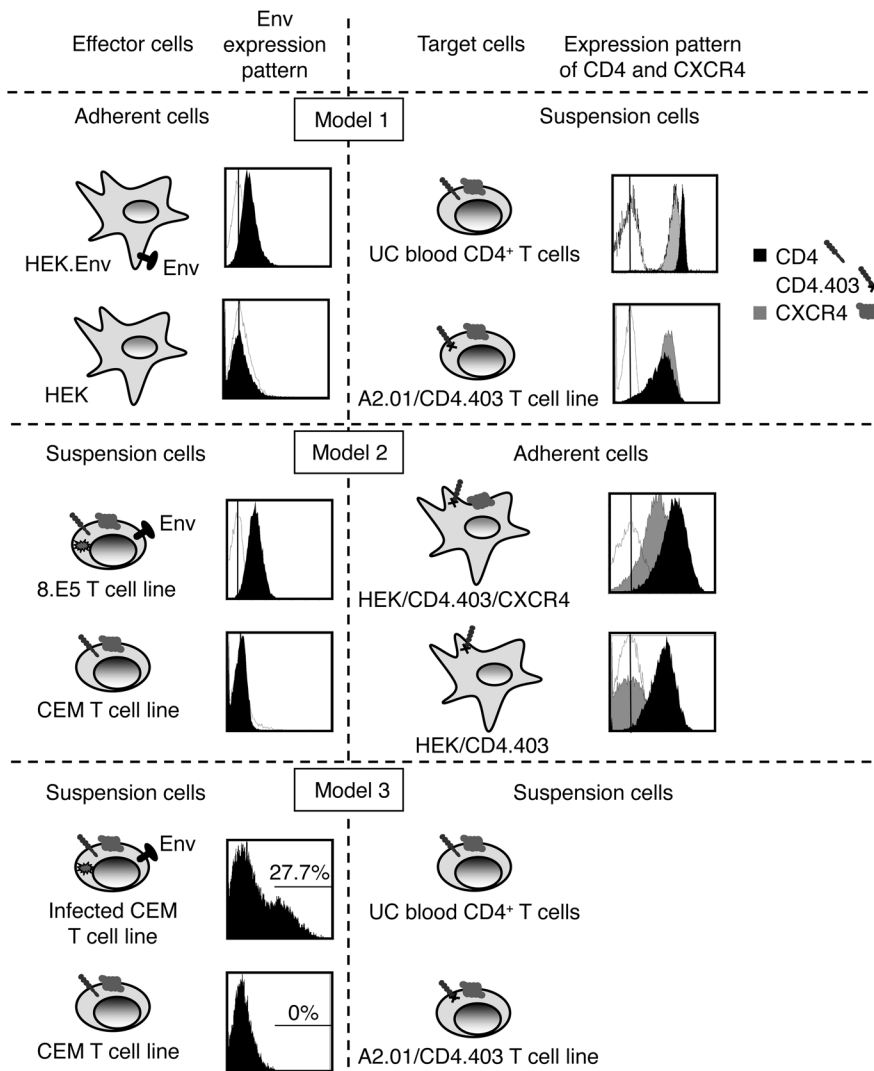


Figure 1

Description of the cellular models. In model 1, target CD4⁺ T cells (UC blood CD4⁺ T cells and a T cell line expressing CXCR4 and CD4.403) are cocultured with effector HEK cells with or without Env expression. In model 2, HEK/CD4.403/CXCR4 and HEK/CD4.403 cells are cocultured with effector T cells with or without Env expression. In model 3, target CD4⁺ T cells are cocultured with HIV_{NL4-3}-infected or uninfected T cells. Expression of gp120 at the surface of HEK, Env, 8.E5, and HIV_{NL4-3}-infected effector cells as well as parental HEK and CEM cells — used as negative controls — were analyzed by flow cytometry after incubation of the cells with PBS (white histograms) or with PBS containing anti-gp120 human polyclonal Ab (black histograms). Bound Ab was detected with a secondary FITC-labeled goat anti-human Ig. Expression analysis of the extracellular domains of CD4 and CXCR4 at the surface of target cells was performed after incubation of the cells with PBS (white histograms) or with anti-CD4 (black histograms) or anti-CXCR4 (gray histograms) mAb at 10 μg/ml. Bound mAb was detected with FITC-labeled goat anti-mouse Ig. The fluorescence intensity was recorded in the log mode on an EPICS XL4 cytofluorometer.

34), the Bcl-2-interacting protein Beclin 1 (28, 35–37), caspase-8 (38) and caspase-9 (35), PI3K class I (39–41), and the sphingolipid ceramide (42, 43) are involved in the control of both apoptosis and autophagy.

Autophagy is triggered in different infections, and several bacteria and viruses have evolved strategies to use components of the autophagic pathway to facilitate their own replication (44–46). Its induction has been observed in response to infection by herpes simplex virus (47), coronavirus (45), and poliovirus (46, 48, 49). Autophagy, in this case named xenophagy, has also been described as a component of the innate immune response that degrades intracellular pathogens (50). This process has also been shown to lead to endogenous MHC class II processing of an Epstein-Barr viral nuclear antigen (51). These data demonstrate that interactions exist between the autophagic pathway and the infectious cycles of several viruses. However, to our knowledge, the occurrence of autophagy has not previously been demonstrated in HIV infection, and there are no descriptions of a putative role of autophagy in cell death induced after viral infection.

On the basis of the hypothesis that HIV-induced bystander naive CD4⁺ T cell killing is central to immunodeficiency, we investigated whether X4 Env is able to trigger autophagy in CD4⁺CXCR4⁺

uninfected cells under the conditions known to induce apoptosis. We demonstrate here that cell-expressed Env binding to CXCR4 induced autophagy of uninfected CD4⁺ T cells and that this was necessary for the induction of apoptosis.

Results

Cellular coculture models. Three models (Figure 1) were used to analyze the capacity of Env expressed on cells to trigger autophagy in the absence of viral replication. All 3 are based on the coculture of effector cells, which express Env, with target cells that express CD4 and CXCR4. These models were created to study the different putative situations in vivo, i.e., the interaction between adherent HIV⁺ epithelial cells and uninfected CD4⁺ T cells, or infected T cells and uninfected CD4⁺ T lymphocytes.

As naive CD4⁺ T cells are the main target of HIV-mediated apoptosis, purified primary umbilical cord (UC) blood CD4⁺ T lymphocytes, which constitute a homogeneous population of cells with a naive phenotype and a high number of CXCR4 molecules, were used in this study. Furthermore, target cells that express a truncated form of CD4 (CD4.403) capable of binding to gp120 but unable to transduce a signal via the lymphocyte-specific protein kinase Lck (52, 53) as well as target cells that express both CD4.403

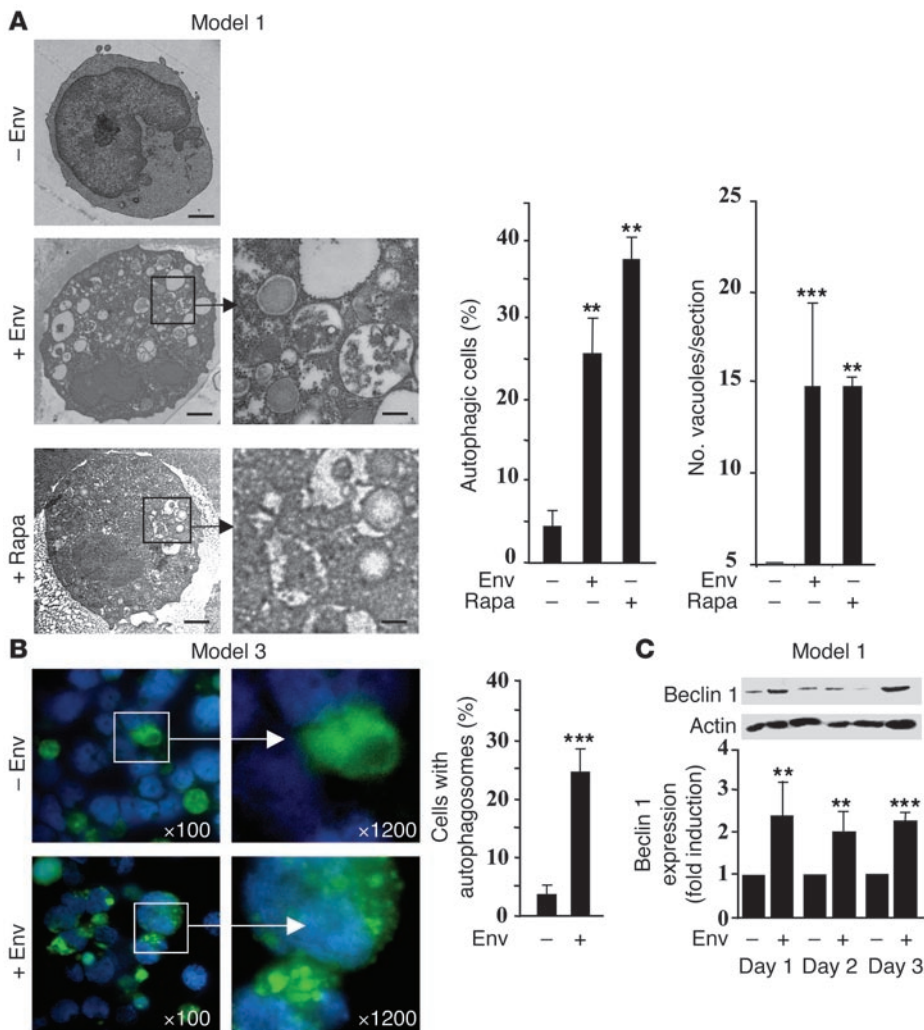


Figure 2

Autophagy is induced by cell-expressed Env, and Beclin 1 is accumulated during this process. (A) UC blood CD4⁺ T lymphocytes were purified by negative selection using the CD4⁺ Rosette separation technique. Cells were cocultured for 3 days with HEK or HEK.Env or treated with 1 μM Rapa and examined by TEM. Autophagic (vacuolated) cells were defined as cells that had 5 or more autophagic vacuoles. The percentage of autophagic cells and the number of vacuoles in the target cell sections were analyzed from at least 100 randomly chosen TEM fields by 2 investigators. Scale bars: 1 μm; 250 nm (indicated enlargements). (B) UC blood CD4⁺ T cells were transfected with GFP-LC3, cocultured with HIV_{NL4-3}-infected or uninfected CEM cells, and examined by epifluorescence. Data are representative of 3 independent experiments; more than 100 cells were counted by 2 investigators. Cells with autophagosomes were defined as cells that had 5 or more LC3 spots in the cytoplasm. Magnification shown. (C) Beclin 1 expression was analyzed in UC blood CD4⁺ T cells cocultured with effector HEK cells with or without Env expression. Fold induction was calculated as the intensity of the Beclin 1 immunoblot obtained in cells cocultured with effector cells expressing Env compared with effector cells that do not express Env. Ratios were normalized to that obtained with anti-actin Ab. Data are representative of at least 3 independent experiments. ***P* < 0.01; ****P* < 0.001.

and wild-type CXCR4 were also analyzed to determine the role of these receptors in Env-mediated autophagy.

In model 1, the target cells (in which autophagy was analyzed) were either CD4⁺ T lymphocytes or the A2.01/CD4.403 T cell line, which expresses wild-type CXCR4 and CD4.403. These target cells were cocultured with effector HEK.293 cells that stably express Env (HEK.Env). The parental HEK cell line was used as a negative control. This model has previously been used to analyze

Env-mediated apoptosis triggered in CD4⁺ T cells (9).

In model 2, the target cells were stably transfected HEK cells that express CD4.403 (HEK/CD4.403) or CD4.403 and CXCR4 receptors (HEK/CD4.403/CXCR4). These target cells were cocultured with a CEM T cell line that harbors a single, defective copy of replication-incompetent HIV-1 proviral DNA and that expresses Env (8.E5 cells). The parental CEM cell line was used as a negative control. This model allowed us to study target cells that do not express wild-type CXCR4. Models 1 and 2 were also created to easily analyze autophagy in the target cells since adherent cells were cocultured with suspension cells.

In model 3, the target cells were uninfected CD4⁺ T lymphocytes and the effector cells were CEM T cells infected with the HIV-1 NL4-3 strain (HIV_{NL4-3}). Uninfected CEM cells were used as a negative control. In this model, both target and effector cells were grown in suspension. In order to distinguish the 2 populations, the effector cells were stained with CMFDA, a nontoxic green fluorescent dye (54). Azidothymidine (AZT; 1 μM) was added to the coculture to inhibit viral replication. This model mimics the contact between infected CD4⁺ T cells and uninfected naive CD4⁺ T cells that mainly occurs in lymph nodes.

Expression patterns of Env on the effector cells and of CD4 and CXCR4 on the target cells are shown in Figure 1.

Env expressed on cells triggers autophagy in uninfected naive CD4⁺ T cells. Autophagy was first analyzed in target UC blood CD4⁺ T lymphocytes after coculture with HEK or HEK.Env (model 1). In the cytoplasm of a significant number of UC blood CD4⁺ T cells cocultured for 3 days with HEK.Env, we observed by transmission electron microscopy (TEM) numerous membrane-bound vacuoles characteristic of autophagosomes that were not present when the CD4⁺ T cells

were cocultured with parental HEK cells (Figure 2A). Rapamycin (Rapa), a classical inducer of autophagy, triggered formation of vacuoles in UC blood CD4⁺ T cells, indicating that these T cells are susceptible to autophagy. Next, to demonstrate that HIV-infected T cells are also able to induce autophagy in naive UC blood CD4⁺ T cells, we analyzed the appearance of autophagosomes after T cell transfection with the reliable autophagosome marker LC3 fused to GFP (GFP-LC3) using model 3. By fluorescence microscopy,

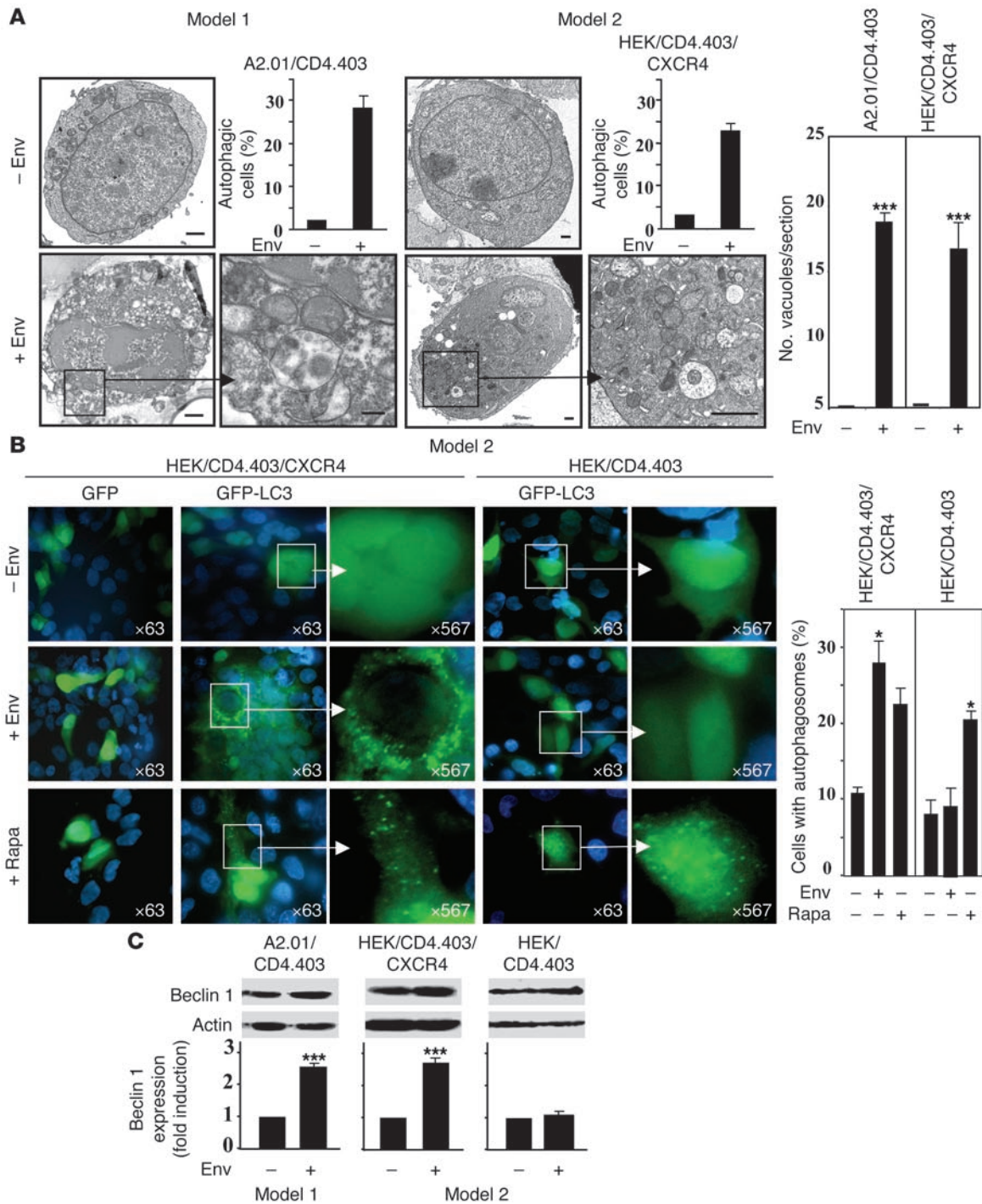


Figure 3

Autophagy is triggered by Env binding to CXCR4. (A) A2.01/CD4.403 and HEK/CD4.403/CXCR4 cells were cocultured for 3 days with cells that do or do not express Env and analyzed by TEM as indicated in Figure 2. Scale bar: 1 μ m; 250 nm (indicated enlargements). (B) Target HEK/CD4.403/CXCR4 and HEK/CD4.403 cells were transfected with GFP or GFP-LC3 using the FuGENE 6 Transfection Reagent and cocultured for 3 days with effector cells with or without Env expression or treated with 1 μ M Rapa. Adherent cells were then examined by epifluorescence. Data are representative of 3 independent experiments and more than 100 cells were counted by 2 investigators. Magnification shown. (C) Immunoblot analysis of Beclin 1 in A2.01/CD4.403, HEK/CD4.403/CXCR4, and HEK/CD4.403 cells cocultured with cells with or without Env expression. * $P < 0.05$; *** $P < 0.001$.

punctate staining was observed in the cytoplasm of GFP-LC3–transfected UC blood CD4⁺ T cells cocultured with HIV_{NL4-3}–infected CEM cells, while only a diffuse pattern was obtained in

GFP-LC3–transfected CD4⁺ T cells cocultured with uninfected CEM cells (Figure 2B). Punctate LC3 staining was also observed in GFP-LC3–transfected UC blood CD4⁺ T cells cocultured with

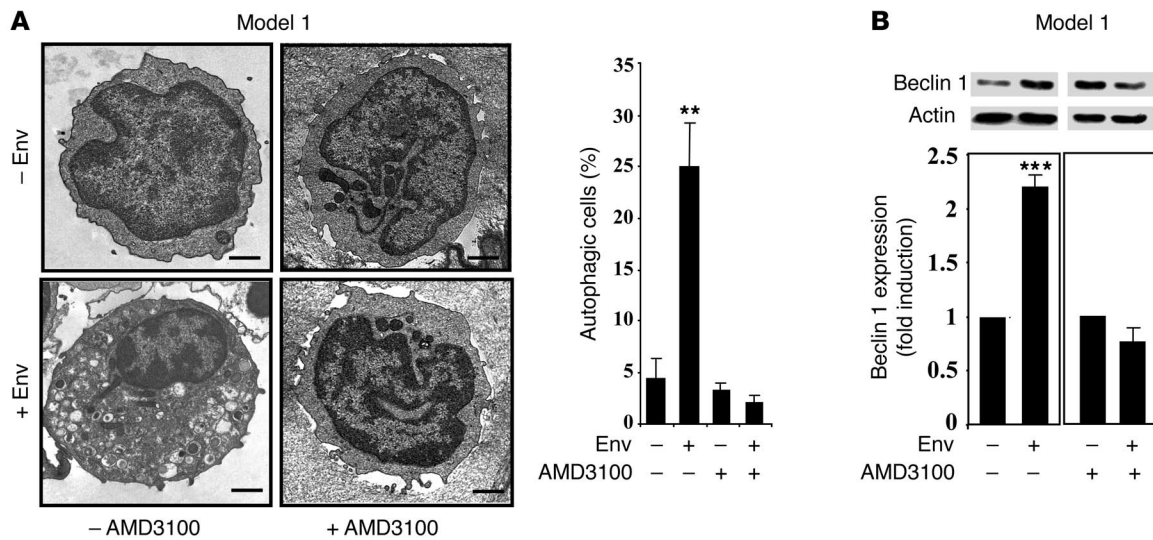


Figure 4

AMD3100 inhibits Env-induced autophagy. (A) UC blood CD4⁺ T lymphocytes were cocultured for 3 days with HEK cells that do or do not stably express Env in the presence or absence of AMD3100 (1 μg/ml) and examined by TEM as previously described. Scale bars: 1 μm. (B) Immunoblot analysis of Beclin 1 in CD4⁺ T cells cocultured for 3 days with cells with or without Env expression in the presence or absence of 1 μg/ml AMD3100. ***P* < 0.01; ****P* < 0.001.

HEK.Env (model 1; data not shown). No punctate staining was observed in UC blood CD4⁺ T cells transfected with GFP-LC3 and cocultured with HEK (model 1; data not shown) or in UC blood CD4⁺ T cells transfected with GFP and cocultured with effector cells in models 1 and 3 (data not shown).

Expression of Beclin 1 in UC blood CD4⁺ T cells during the coculture with cells with or without Env expression (model 1) was also analyzed. Beclin 1 accumulated in UC blood CD4⁺ T cells within 24 hours of coculture with HEK.Env, and its accumulation was maintained during the 3 days of coculture (Figure 2C). On the contrary, Beclin 1 was not accumulated in UC blood CD4⁺ T cells cocultured with HEK. It is worth noting that Rapa at 200 nM or 1 μM did not induce Beclin 1 accumulation in CD4⁺ T cells (data not shown). Thus, contact of X4 Env with HIV receptors, CD4 and CXCR4, on uninfected CD4⁺ T cells, triggers ultrastructural changes associated with autophagy and accumulation of Beclin 1.

Autophagy is triggered by Env binding to CXCR4. Interaction of X4 Env with its receptors at the T cell surface is a 2-step event. First, gp120 binds to CD4, leading to an increased exposure of the gp120 V3 loop and CD4i region which can then interact with CXCR4. Both CD4 and CXCR4 receptors initiate intracellular signaling events after Env binding, including activation and apoptotic signals (9, 55). To determine the role of these receptors in the induction of autophagy, we cocultured effector cells with or without Env expression for 3 days with either the A2.01/CD4.403 T cell line or HEK/CD4.403/CXCR4 cells (models 1 and 2, respectively). It is worth noting that these target cells are susceptible to Env-induced apoptosis (9, 56). By TEM, autophagic vacuoles were observed in A2.01/CD4.403 and HEK/CD4.403/CXCR4 cocultured with effector cells expressing Env (models 1 and 2, respectively), while no vacuoles were present in these target cells cocultured with effector cells that did not express Env (Figure 3A). Target CEM cells that expressed wild-type CXCR4 and CD4, as UC blood CD4⁺ T cells, were still susceptible to Env-mediated autophagy (model 1, data not shown), indicating that CD4 signaling does not prevent the triggering of autophagy.

Punctate staining was observed in the cytoplasm of GFP-LC3-transfected HEK/CD4.403/CXCR4 cocultured with 8.E5 cells, while only a diffuse pattern was obtained in cells cocultured with effector CEM cells and target cells expressing GFP alone. Furthermore, no formation of autophagosomes was observed in HEK/CD4.403 that did not express CXCR4 after coculture with effector cells expressing Env (model 2; Figure 3B), indicating that the presence of CXCR4 is necessary for Env-mediated autophagosome formation. The number of target cells with autophagosomes after contact with effector cells that express Env was similar to those obtained after incubation with 1 μM Rapa (Figure 3B). As observed in UC blood CD4⁺ T cells, Beclin 1 was accumulated in target cells that expressed CXCR4 and CD4.403 after coculture for 3 days with effector cells expressing Env. In the same conditions, the level of Beclin 1 expression was stable in target cells that expressed CD4.403 but not CXCR4 (Figure 3C). Time-course experiments were performed to analyze the kinetics of the autophagic process induced by Env or Rapa in HEK/CD4.403/CXCR4 cells. LC3 punctate staining was significantly induced after 2 days of coculture with 8.E5 cells, and the level of autophagy was maintained after 3 days of coculture with these effector cells. Rapa induced slightly different kinetics, since the number of cells with autophagosomes decreased on the third day of incubation (Supplemental Figure 1A; supplemental material available online with this article; doi:10.1172/JCI26185DS1).

The finding that cell-expressed Env induced formation of autophagic vacuoles in 2 target cell types expressing wild-type CXCR4 and a truncated form of CD4 but not in cells that did not express CXCR4 indicates that Env-mediated autophagy is not cell type dependent and that activation of signaling pathways through CD4 is not needed.

Expression of Beclin 1 was studied in target A2.01/CD4.403, HEK/CD4.403/CXCR4, and HEK/CD4.403 cells after coculture with effector cells with or without Env expression for 1–3 days (models 1 and 2). Accumulation of Beclin 1 was observed in cells

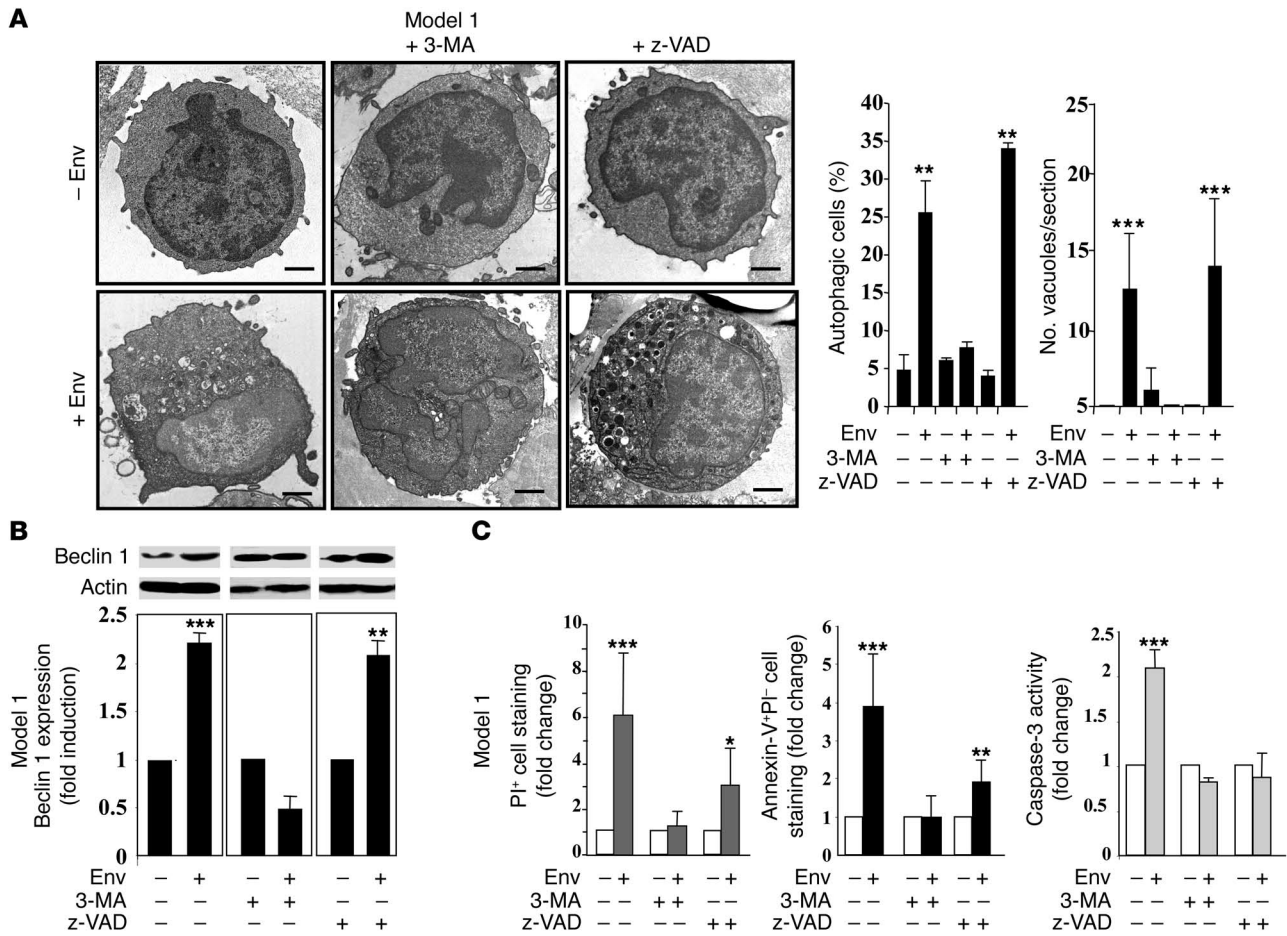


Figure 5

Env expressed on adherent effector cells triggers caspase-dependent and -independent cell death pathways in uninfected CD4⁺ T cells that are blocked by 3-MA. (A) UC blood CD4⁺ T lymphocytes were cocultured for 3 days with HEK or HEK.Env in the presence or absence of 10 mM 3-MA and 50 μM z-VAD. Cells were examined by TEM as previously described. Scale bars: 1 μm. (B) Beclin 1 expression was analyzed in CD4⁺ T lymphocytes cocultured for 3 days with cells with or without Env expression in the presence or absence of 3-MA and z-VAD. (C) Fold increase in Env-mediated CD4⁺ T cell death was analyzed by PI cell staining (global cell death), Annexin-V and PI cell staining (non necrotic cell death) and caspase-3 activity (apoptosis) in the presence or absence of 3-MA and z-VAD. The fold change of cell staining or caspase-3 activity was calculated as the percentage of positive target cells after coculture with cells that express Env versus those that do not express Env. Results are from at least 5 independent experiments. **P* < 0.05; ***P* < 0.01; ****P* < 0.001.

expressing CD4.403 and CXCR4 as early as the first day of coculture with effector cells expressing Env, and this accumulation was maintained during the 3 days of coculture. On the contrary, no accumulation of Beclin 1 occurred in cells that did not express CXCR4 (Supplemental Figure 1B). Thus, Env-induced Beclin 1 accumulation is dependent on the presence of CXCR4 on the target cells and precedes autophagosome formation.

Furthermore, AMD3100, a small-molecule inhibitor of HIV-1 entry via the CXCR4 coreceptor (57) that efficiently blocks Env-induced apoptosis (58), was able to inhibit autophagic vacuole formation (Figure 4A) and Beclin 1 accumulation (model 1; Figure 4B) after contact with Env. It is worth noting that SDF-1, the physiological CXCR4 ligand, was alone unable to trigger autophagy (data not shown). In addition, suramin, which binds to the V3 loop of HIV-1 gp120 (59) and inhibits Env-induced apoptosis (56), blocked autophagosome formation as visualized by fluorescence studies after GFP-LC3 transfection (model 2; Supplemental Figure 2).

This result highlights the role played by gp120 binding on CXCR4 to trigger autophagy. Furthermore, these are the first data to our knowledge to demonstrate that a chemokine receptor engagement is able to trigger autophagy.

Interconnection between Env-induced autophagy and apoptosis. Env expressed on cells is able to activate apoptosis (9, 10, 60) and autophagy (our present results) after binding to CXCR4 on target cells. We then investigated whether these cell death-associated phenomena are linked as well as the role of the autophagic process in Env-mediated T cell death. To answer these questions, we used inhibitors for each pathway. The drug 3-methyladenine (3-MA) is widely used as an inhibitor of autophagy since it was found to prevent autophagosome formation without markedly affecting protein synthesis or ATP levels (61). At the sequestration step, 3-MA interferes with the activity of class III PI3K to interrupt autophagy (40, 62). To inhibit apoptosis, z-Val-Ala-Asp (OMe)-CH₂F (z-VAD), a pan-caspase inhibitor, was used. The percentage of UC blood CD4⁺ T cells accumulating autophagic vacuoles after coculture for

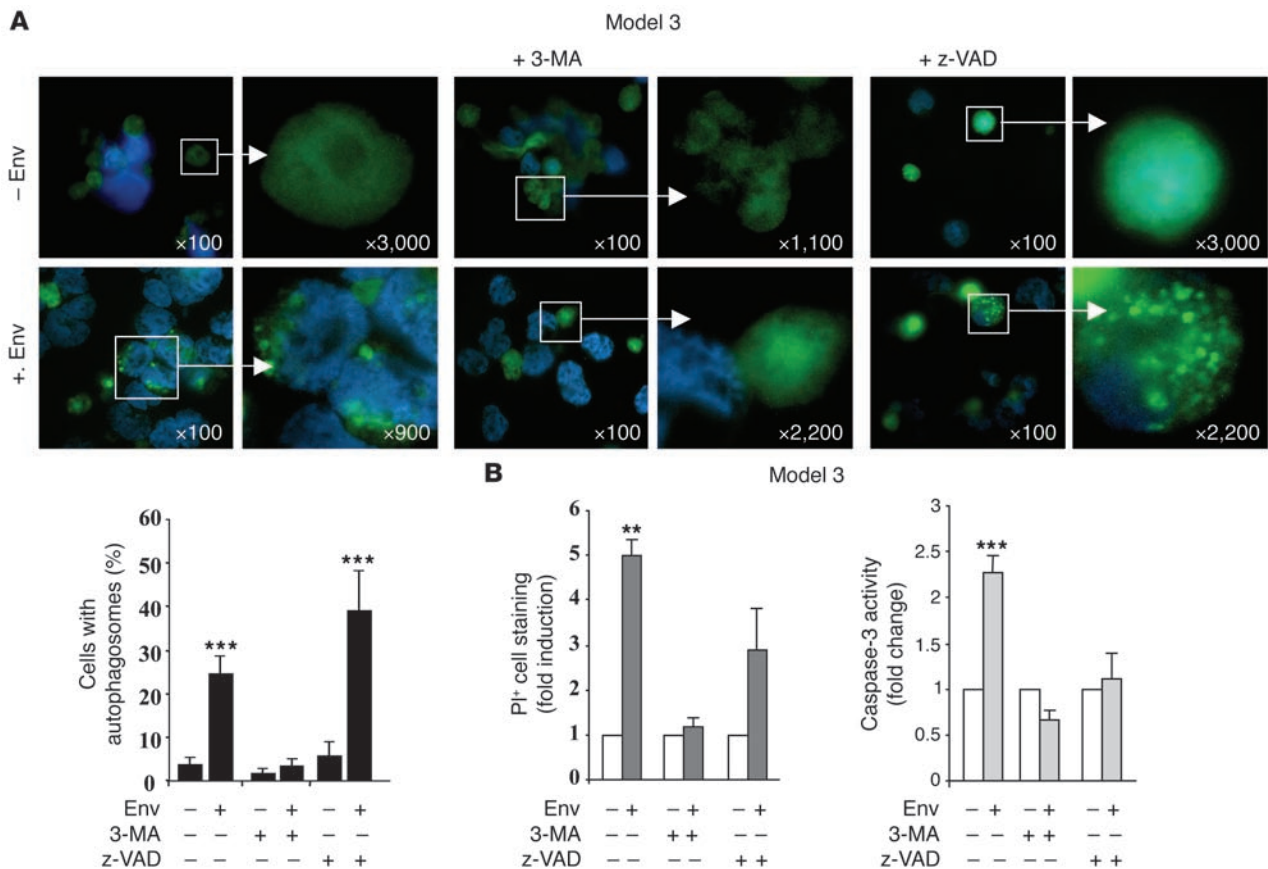


Figure 6

Env expressed on infected effector CD4⁺ T cells triggers caspase-dependent and -independent cell death pathways in uninfected CD4⁺ T cells that are blocked by 3-MA. (A) GFP-LC3–transfected UC blood CD4⁺ T lymphocytes were cocultured for 3 days with HIV_{NL4-3}-infected or uninfected CEM cells in the presence or absence of 10 mM 3-MA or 50 μM z-VAD. Cells were examined by fluorescence as described in Figure 2. Magnification shown. (B) Fold increase in Env-mediated uninfected CD4⁺ T cell death was studied by PI cell staining (global cell death) and caspase-3 activity (apoptosis) in the presence or absence of 3-MA or z-VAD. Results are from at least 3 independent experiments. ***P* < 0.01; ****P* < 0.001.

3 days with effector HEK.Env was strongly reduced in the presence of 3-MA, but no decrease was observed in the presence of z-VAD. Furthermore, z-VAD did not reduce the number of autophagic vacuoles in the autophagic CD4⁺ T cells after coculture with HEK.Env (model 1; Figure 5A). These results demonstrate that 3-MA, but not z-VAD, prevents Env-induced autophagy. Beclin 1 accumulation was also suppressed in CD4⁺ T cells after coculture for 3 days with HEK.Env in the presence of 3-MA but not in the presence of z-VAD (Figure 5B). Next we studied the level of Env-mediated global cell death by propidium iodide (PI) cell staining and the level of Env-mediated apoptosis by analysis of caspase-3 activity. We also analyzed by flow cytometry the percentage of non-necrotic CD4⁺ T cells that exposed phosphatidylserine (PS) at their surface (cells stained with Annexin-V but PI negative) after coculture with cells with or without Env expression. This labeling is currently used to detect apoptosis, but PS exposure was also recently observed in the autophagic process (63). Administration of 3-MA totally inhibited Env-mediated global CD4⁺ T cell death, caspase-3 activity, and PS exteriorization (Figure 5C). On the contrary, z-VAD, which totally inhibited apoptosis, did not abolish Env-mediated global T cell death (Figure 5C). Similar results were obtained using the trypan blue exclusion test (data not shown). This suggests that Env trig-

gers both caspase-dependent and -independent cell death pathways and that Env-mediated autophagy is needed to trigger apoptosis.

We also analyzed the capability of these drugs to inhibit autophagy and cell death triggered in uninfected target CD4⁺ T cells after coculture with effector HIV_{NL4-3}-infected CD4⁺ T cells (model 3). Autophagosome formation in UC blood CD4⁺ T cells after coculture with HIV-infected or uninfected CEM cells was analyzed by fluorescence using LC3 staining in the presence or absence of 3-MA and z-VAD. Punctate LC3 staining was only observed in UC blood CD4⁺ T cells after coculture for 4 days with infected effector cells. Autophagosome formation was still observed in the presence of z-VAD, but was totally blocked by 3-MA (Figure 6A). Next, death of target CD4⁺ T cells was analyzed, after coculture with effector CEM cells infected with HIV_{NL4-3} or uninfected, by flow cytometry after PI staining. In the same conditions, Env-induced apoptosis was studied by analysis of caspase-3 activity. Prior to the coculture, uninfected and HIV_{NL4-3}-infected effector CEM cells were labeled with the nontoxic green fluorescent dye CMFDA to differentiate the target cells. Furthermore, AZT was added to the medium to avoid HIV-1 replication. While 3-MA totally inhibited Env-mediated cell death and caspase-3 activity, z-VAD did not totally block cell death but completely inhibited apoptosis (Figure 6B).

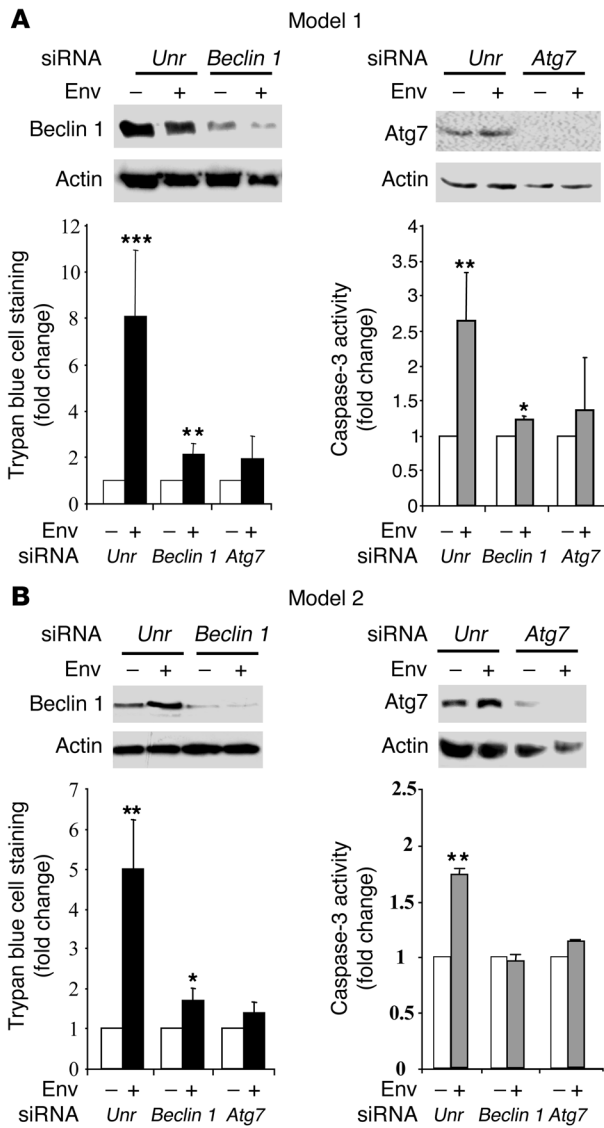


Figure 7

siRNAs against *Beclin 1* and *Atg7* inhibit Env-mediated cell death. (A) CD4⁺ T cells were cocultured with cells with or without Env expression after transfection with *Beclin 1*, *Atg7*, or unrelated (unr) siRNA. Reduction in expression of the corresponding proteins was analyzed by Western blot using the specific Abs. Global cell death — as determined by trypan blue exclusion test and caspase-3 activity, analyzed by colorimetric assay — is shown. Data are representative of at least 5 independent experiments. (B) HEK/CD4.403/CXCR4 cells were cocultured with cells with or without Env expression after transfection with *Beclin 1*, *Atg7*, or unrelated siRNA. Analysis of global cell death and caspase-3 activity was performed as in A. **P* < 0.05; ***P* < 0.01; ****P* < 0.001.

Discussion

The apoptosis of uninfected target CD4⁺ T cells can be induced by the binding of their CXCR4 receptor to Env expressed on effector cells (9, 10, 60, 66). This Env-mediated apoptotic pathway has been proposed as a phenomenon contributing to CD4⁺ T cell loss during HIV infection. We demonstrate here that cells that expressed Env were also capable of inducing autophagy in uninfected CD4⁺ T cells. This was equally true of Env-transfected adherent cells, which mimic infected cells of the epithelia, and infected CD4⁺ T cells. The autophagic process is triggered by Env binding to CXCR4, since target cells that expressed wild-type CXCR4 and a truncated, signal transduction-incompetent form of CD4 still underwent Env-mediated autophagy, while cells that expressed a truncated form of CD4 but did not express CXCR4, did not. In addition, AMD3100, a CXCR4 antagonist that prevents X4 HIV entry and totally inhibits Env-induced apoptosis, also blocked Env-mediated autophagy. Inhibition of Env-mediated autophagy by suramin, a compound that binds the gp120 V3 loop, also strongly suggests that binding of gp120 to CXCR4 is responsible for this phenomenon. It is worth noting that after 3 days of coculture with effector cells that expressed Env, autophagosomes were also present in syncytia: multinucleated, giant cells formed after fusion between the 2 types of cells in culture through interaction of Env with CD4 and CXCR4. This underlines the role of these receptors in the triggering of autophagy.

The autophagic protein Beclin 1 was rapidly accumulated in CD4⁺ T cells cocultured with cells that express Env, and its accumulation was maintained during the 3 days of coculture. In contrast, the formation of autophagosomes was only observed after 2 days of coculture, indicating that Beclin 1 is involved in the very early steps of the Env-induced signaling cascade, preceding autophagic vacuolization. Accumulation of Beclin 1 has also been reported in response to drugs that trigger cell death with autophagic features (29, 35, 43). However, the precise mechanism of Beclin 1 accumulation is unknown and remains to be investigated.

Interestingly, Env-mediated autophagy not only preceded apoptosis but was also required for this process. Indeed, the inhibition of Env-induced autophagy by either administration of general inhibitors of autophagy (3-MA and Baf A1) or inhibition of 2 genes involved in the autophagic pathway (*Beclin 1* and *Atg7*) inhibited caspase-3 cleavage as well as activity, which mainly occurs in CD4⁺ T cells after 3 days of coculture with cells expressing Env (9). These results corroborate several studies that have already demonstrated that autophagy can be required for apoptosis (67–72). It should be pointed out, however, that if the initial phase of autophagy is required, it is not necessarily sufficient for triggering apoptosis (69). In our study, Baf A1, which is known to block the formation

For further confirmation of the involvement of an autophagic process in Env-induced apoptosis, we used gene silencing with siRNA to inhibit expression of 2 genes, *Beclin 1/Atg6* and *Atg7*, related to the autophagic pathway in CD4⁺ T cells (model 1; Figure 7A) and in HEK/CD4.403/CXCR4 cells (model 2; Figure 7B). Reduced expression of these 2 proteins by RNA interference induced a dramatic decrease in Env-induced caspase-3 activity and totally inhibited Env-mediated global cell death (Figure 7).

Finally, to determine the role of vacuolization in Env-mediated cell death, we used bafilomycin A1 (Baf A1), an inhibitor of vacuolar H⁺ ATPase (64, 65) that leads to accumulation of autophagic vacuoles by blocking their fusion with lysosomes. Baf A1 inhibited global cell death and caspase-3 activity in uninfected CD4⁺ T cells cocultured with either Env-transfected (model 1) or infected cells (model 3; Figure 8), indicating that vacuolization is not directly responsible for the Env-mediated cell death related to autophagic genes; rather, the complete autophagic process is needed.

Collectively, these data led us to propose a model in which autophagy is a prerequisite for Env-induced activation of the apoptotic pathway through CXCR4 (Figure 9).

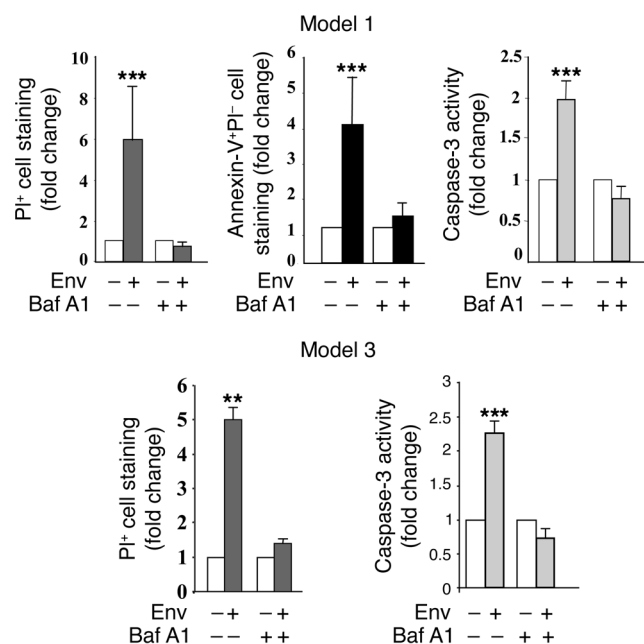


Figure 8

Baf A1 inhibits global CD4⁺ T cell death and apoptosis. (A) Analysis of CD4⁺ T cell death after coculture with HEK and HEK.Env, as described in Figure 5, in the presence or absence of 100 nM Baf A1. (B) Analysis of CD4⁺ T cell death after coculture with HIV_{NL4-3}-infected or uninfected CD4⁺ T cells, as described in Figure 6, in the presence or absence of 100 nM Baf A1. ***P* < 0.01; ****P* < 0.001.

of autophagolysosomes, totally inhibited Env-mediated apoptosis, indicating that the complete autophagic process is needed for this apoptotic pathway. This also suggests that even if autophagic vacuole formation alone does not induce apoptotic cell death, lysosomal degradation of the autophagic vacuole contents does.

In addition, inhibition of autophagy totally blocked Env-induced T cell death, whereas inhibition of Env-mediated apoptosis did not. Even if these data were obtained in nonphysiological conditions, i.e., when apoptosis was blocked, they indicate that Env induces a nonapoptotic cell death with autophagic features. This corroborates recent data indicating that PCD dependent on autophagy genes can directly contribute to cell death (29, 38, 73). In *Dictyostelium*, a nonvacuolar cell death pathway was observed in the presence of conspicuous vacuolization (74). It is worth noting that Baf A1 decreased cell viability independently of Env signaling, indicating that accumulation of vacuoles may lead to a basal level of cell death without triggering the specific cascade of events through interaction of Env with CXCR4.

Together, these results demonstrate that autophagy and apoptosis are responsible for CD4⁺ T cell destruction during HIV infection but also raise unsolved questions about the mechanisms by which these cellular death pathways are triggered after Env binding to CXCR4. One hypothesis may be that Beclin 1, which was first identified as a Bcl-2-interacting protein and accumulates during the Env-mediated autophagic process, could enhance caspase-9 activity by interacting with Bcl-2 and Bcl-x_L antiapoptotic proteins, as recently demonstrated in human gastric cancer cells after *cis*-diamminedichloroplatinum-induced apoptosis (35). This hypothesis certainly needs to be tested, since caspase-9 is cleaved during Env-mediated apoptosis, which was shown here to be activated by the autophagic process. Very recently, Pattingre et al. demonstrated that Bcl-2 antiapoptotic proteins inhibit Beclin 1-dependent autophagy (28). Env-mediated accumulation of Beclin 1 may thus disrupt this equilibrium to trigger the cell toward both autophagy and apoptosis. Indeed, as demonstrated in yeast (75), cell death can also be considered to be a complex cellular response

triggered by mitochondrial dysfunction rather than by one specific cell death pathway. In line with this, Env binding to CXCR4 activates the intrinsic pathway of apoptosis, in which mitochondria play a central role. Moreover, there is now considerable evidence that mitochondria integrate a plethora of cell death signals (76).

We do not currently know whether Env-induced autophagy was first triggered to rescue the CD4⁺ T cells from cell death or if the autophagic process is directly activated to kill the CD4⁺ uninfected T lymphocytes. One hypothesis is that Env-mediated autophagy may reduce the mass of cells prior to apoptosis, facilitating their phagocytosis. However, once activated, the ultimate consequence of autophagy is the triggering of cell death that contributes to pathology. Very recently, in an autosomal-dominant familial neurohypophyseal diabetes insipidus, autophagy-mediated cell death was proposed as a “2-hit” process, in which autophagy is first prosurvival after a cellular stress by removing potentially toxic aggregates, but this ongoing autophagic activity sensitizes cells to the toxic effects of a second insult (77). Exclusive definitions of each PCD may be artificial due to the overlap and shared signaling pathways among these different cell death programs (78), perhaps due to redundant cell death mechanisms. Indeed, Golstein and Kroemer have made the hypothesis that autophagic or apoptotic elements might have been added to a primordial death mechanism, initially improving cell dismantling and later acquiring the ability to act by their own right as death effectors (79).

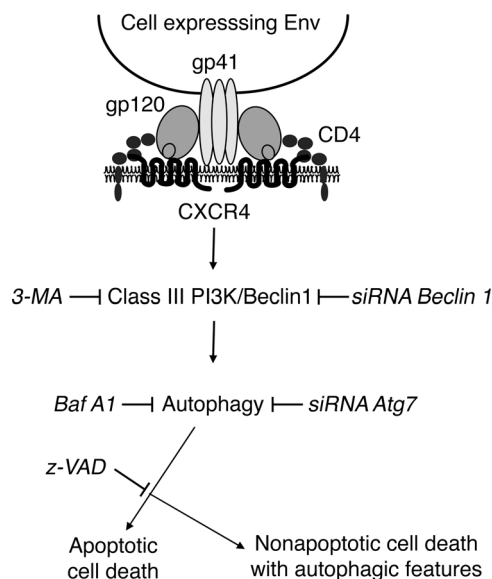


Figure 9

A model for Env-induced signaling cell death cascade. Autophagy is specifically triggered after Env binding to CXCR4, leading to apoptosis. Nonapoptotic cell death with autophagic features is observed when apoptosis is blocked.



Another unsolved question concerns the fact that other signaling pathways coming from molecules that participate in the virological synapse formation may be necessary, in addition to the interaction between Env and CXCR4. Nothing is currently known about autophagy induced in uninfected CD4⁺ T cells by free virions and in HIV-infected CD4⁺ T cells, and there is no direct demonstration of the presence of autophagic T cells *in vivo* in HIV-infected patients. Further investigation will be required to answer these points.

To the best of our knowledge, these findings represent the first example of autophagy triggered through binding of virus envelope proteins to a cellular receptor without viral replication, leading to apoptosis and cell death linked to autophagy.

This study suggests that naive CD4⁺ T cells, which cannot be productively infected by HIV, may take different routes to die after contact with a cell infected by an X4 HIV strain and that sequential, but not exclusive, cell death pathways are triggered by Env binding to CXCR4. Autophagy thus plays a crucial role in Env-mediated CD4⁺ T cell death that leads to the physiopathology of AIDS.

Methods

Cell culture. The original HEK.293 cell line as well as HEK.Env (9), HEK/CD4.403/CXCR4, and HEK/CD4.403 cells were cultured in DMEM supplemented with 1% penicillin/streptomycin antibiotic mixture, 1% Glutamax (Cambrex), and 10% FCS (Perbio, Fisher Scientific).

The CEM T cell line was provided by the ATCC. The CEM-derived A2.01/CD4.403 cell clone, which expresses a mutant form of CD4 truncated at position 403 (80) was provided by D.R. Littman (New York Medical College, New York, New York, USA). The 8.E5 cell line, a CEM-derived T cell line containing a single integrated copy of HIV-1 and unable to produce infectious virions, was provided by F. Barré-Sinoussi (Institut Pasteur, Paris, France). These T cell lines were cultured in RPMI 1640 medium supplemented with 1% penicillin/streptomycin, 1% Glutamax, and 10% FCS. UC blood CD4⁺ T cells from UC blood samples obtained from caesarean deliveries were purified by negative selection using the Human CD4⁺ T Cell Enrichment cocktail (StemCell Technologies). UC blood cells were cultured in complete RPMI 1640 medium containing IL-4 (10 ng/ml). Procedures using human cells were approved by the Human Experimentation and Ethics Committee of the CNRS Institute.

Reagents and antibodies. z-VAD was purchased from Calbiochem (Merck); 3-MA was purchased from MP Biomedicals; and AMD3100, Baf A1, and suramin were purchased from Sigma-Aldrich. The anti-Beclin 1 Ab was purchased from tebu-bio. Anti-cleaved caspase-3 Ab was purchased from Cell Signaling Technology. Rapa was purchased from VWR, and CMFDA was purchased from Invitrogen.

Flow cytometry. Cells (1×10^5) were incubated for 1 hour at 4°C with 50 μ l PBS supplemented – or not – with the appropriate Ab. After washes with PBS, bound Ab was revealed by addition of 50 μ l of a 1:100 dilution of FITC-conjugated secondary Ig. After staining for 30 minutes, cells were washed with PBS, and fluorescence intensity at 543 nm was measured on a COULTER EPICS XL Flow Cytometer (Beckman Coulter).

HIV-1 infection. CEM T cells were infected with the HIV-1 X4 NL4-3 strain. CEM cells (5×10^6) were cocultured with plated HEK.293 cells previously transfected with 5 μ g of HIV-1 NL4-3 DNA construct using FuGENE 6 Transfection Reagent (Roche Diagnostics) according to the manufacturer's instructions. Follow-up of CEM infection was done by p24 and Env expression analysis using flow cytometry.

RNA interference. The sense and antisense oligonucleotides specific for *Beclin 1* and *Atg7* siRNA were synthesized by Eurogentec and annealed according to the manufacturer's instructions. The mRNA sequences to be targeted by *Beclin 1* and *Atg7* siRNA were 5'-CAGUUUGGCA-

CAAUCAAUA-3' and 5'-GGAGUCACAGCUCUCCUU-3', respectively. CD4⁺ T cells were transfected using Nucleofector Technology (amaxes biosystems), according to the manufacturer's instructions, and HEK/CD4.403/CXCR4 cell transfection was performed using the Oligofectamine reagent (Invitrogen). Transfected cells were then cocultured with cells with or without Env expression.

TEM analysis. After 3 days of coculture with cells with or without Env expression, cells were fixed *in situ* with 2.5% glutaraldehyde in cacodylate buffer (pH 7.4) for 60 minutes at 4°C. Cells were then postfixed with 2% osmium tetroxide and washed in cacodylate buffer containing 0.5% tannic acid. After extensive washes in 0.1 M Sorensen phosphate buffer (pH 7.2), cells were included in a fibrin clot as described by Charret and Fauré-Fremiet (81). Cells were then postfixed with 2% osmium tetroxide and 0.5% tannic acid, dehydrated, and embedded in epon (EMbed-812; Electron Microscopy Sciences). Sections were counterstained with uranyl acetate and lead citrate and examined with a Hitachi H7100 TEM.

Western blot analysis. Cells were washed twice in PBS and lysed in buffer containing 50 mM Tris HCl (pH 8), 1% Triton X-100, 100 mM NaCl, 1 mM MgCl₂, 150 μ M PMSF, and complete mini protease inhibitor cocktail (Roche Diagnostics). Cell lysates were electrophoresed in 10% SDS-PAGE and blotted to PVDF membranes. After a blocking step for 1 hour at room temperature, blots were incubated overnight at 4°C with the primary Ab in the blocking buffer. After 3 washes with TBS and 0.05% Tween, the blots were incubated for 1 hour at room temperature with peroxidase-coupled antiserum diluted in TBS, 5% milk, and 0.05% Tween. After further washes, the immune complexes were revealed by ECL (PerkinElmer) and autoradiographed. Quantification of protein expression was performed by densitometry analysis of the autoradiograms using Image J (<http://rsb.info.nih.gov/ij/>). Data were normalized with reference to the density analysis of actin.

Detection of cell death and apoptosis. Cell death of the CD4⁺ T cells was studied after coculture with effector cells with or without Env expression by flow cytometry analysis using PI and Annexin-V-FITC staining. Briefly, suspension cells were washed with PBS, carefully resuspended in 50 μ l binding buffer (100 mM HEPES, pH 7.4; 140 mM NaCl; 5 mM CaCl₂), and a mixture of 4 μ l Annexin-V-FITC and 4 μ l PI was added. After incubation for 20 minutes in the dark at 4°C, cells were analyzed on a COULTER EPICS XL Flow Cytometer (Beckman Coulter). Cell viability was also determined by trypan blue exclusion test. The total number of dead cells was counted and normalized with reference to the total number of cells. Apoptosis was determined using the caspase-3 colorimetric assay from R&D Systems according to the manufacturer's instructions.

LC3 staining. HEK/CD4.403/CXCR4 cells were cultured in 6-well plates and transfected with pEGFP or pEGFP-LC3 plasmid (referred to as GFP and GFP-LC3, respectively) using FuGENE 6 Transfection Reagent. UC blood CD4⁺ T cells were transfected with GFP or GFP-LC3 using the Nucleofector apparatus (amaxes biosystems) according to the manufacturer's instructions. Transfected HEK or CD4⁺ T cells were cocultured with effector cells with or without Env expression for 3 or 4 days, respectively. Target cells were then washed with PBS and fixed in 3.7% paraformaldehyde in PBS for 10 minutes at room temperature (HEK cells) or in an acetone/methanol (vol/vol) mixture for 10 minutes at -20°C (UC blood CD4⁺ T cells) and examined by epifluorescence using a Leica microscope. More than 100 transfected cells were counted by 2 investigators. Only cells with at least 5 spots were counted as positive. Results are from at least 3 independent experiments.

CMFDA staining. Cells were incubated for 30 minutes in prewarmed 0% FCS culture medium containing 20 μ M CMFDA. After centrifugation, cells were resuspended in fresh prewarmed culture medium without FCS for 30 minutes and washed 3 times. Cell fluorescence was controlled by flow cytometry analysis.



Statistics. Analysis of the variance of the results obtained after binding to cells that do not express Env versus cells that express Env was performed after arc sine transformation of the data when percentages were compared (82). Differences were considered significant at * $P < 0.05$, ** $P < 0.01$, and *** $P < 0.001$.

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