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Bone degradation machinery of osteoclasts: An HIV-1 target that contributes to bone loss

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Bone deficits are frequent in HIV-1-infected patients. We report here that osteoclasts, the cells specialized in bone resorption, are infected by HIV-1 in vivo in humanized mice and ex vivo in human joint biopsies. In vitro, infection of human osteoclasts occurs at different stages of osteoclastogenesis via cell-free viruses and, more efficiently, by transfer from infected T cells. HIV-1 infection markedly enhances adhesion and osteolytic activity of human osteoclasts by modifying the structure and function of the sealing zone, the osteoclast-specific bone degradation machinery. Indeed, the sealing zone is broader due to F-actin enrichment of its basal units (i.e., the podosomes). The viral protein Nef is involved in all HIV-1-induced effects partly through the activation of Src, a regulator of podosomes and of their assembly as a sealing zone. Supporting these results, Nef-transgenic mice exhibit an increased osteoclast density and bone defects, and osteoclasts derived from these animals display high osteolytic activity. Altogether, our study evidences osteoclasts as host cells for HIV-1 and their pathological contribution to bone disorders induced by this virus, in part via Nef.

osteoclast | HIV-1 infection | bone loss | Nef | podosome

Reduced bone mineral density is a frequent complication of HIV-1-infected patients and often progresses to osteoporosis and high prevalence of fractures. A sixfold increased risk of low bone mineral density is observed in HIV-1⁺ individuals compared with the general population (1). The use of highly active antiretroviral therapy (HAART) has significantly improved the lifespan of patients, revealing these long-term effects of the infection and the persistence of latent proviruses in reservoir cells (2). Multiple factors can contribute to bone loss in infected patients. HAART is one of these factors, especially during the first years of therapy. In addition, there is evidence of bone deficit in nontreated patients, showing that the virus alone alters bone homeostasis (3–6).

Bones undergo continual remodeling, which mainly relies on the sequential actions of bone-resorbing osteoclasts (OC) and bone-forming osteoblasts, under the control of osteocytes (7, 8). In the case of aging or HIV-1 infection, this balance can be disrupted in favor of bone loss. HIV-1-induced bone disorders are associated with an increase of blood biomarkers for bone resorption and minor changes in bone formation-specific markers, suggesting a major contribution of OC in this process (6, 9).

OC are multinucleated cells derived from the monocytic lineage, which have the unique ability to resorb bone matrix. They terminally differentiate by fusion from mononucleated precursors, including blood-circulating monocytes and bone-resident precursors (10). This process is regulated by macrophage colony-stimulating factor (M-CSF) and the key osteoclastogenic cytokine, receptor activator of NF- κ B ligand (RANKL), mainly secreted by osteocytes but also by osteoblasts and activated B and T cells (10, 11). Terminally differentiated OC express high levels of the α v β 3 integrin adhesion receptor and enzymes involved in resorption including cathepsin K, matrix metalloproteinase 9 (MMP9), and tartrate-resistant acid phosphatase (TRAP). Bone attachment and resorption are mediated

Significance

Bone deficits are frequent complications observed in HIV-1-infected patients. Our study demonstrates that HIV-1 infects osteoclasts, the cells specialized in bone degradation, using different models including HIV-1-infected humanized mice. We decipher the cellular mechanisms by which HIV-1 contributes to enhanced bone degradation in human osteoclasts, showing that the virus modifies the structure and function of the sealing zone, the bone resorption machinery of osteoclasts. We identify the viral protein Nef as the key factor responsible for such effects. As a proof-of-concept, we correlate bone deficit in transgenic Nef-expressing mice with enhanced osteoclast activity. Therefore, our findings provide formal evidence that osteoclasts constitute HIV-1 host target cells, contributing to bone deficits in vivo.

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by an OC-specific structure called the sealing zone (SZ). It is composed of a dense array of interconnected F-actin structures, the podosomes. The SZ anchors the cells to the bone surface and creates a confined resorption environment where protons and osteolytic enzymes are secreted (11–14). Alteration in SZ formation and dynamics are linked to defective bone resorption, and ultimately to bone disorders, as demonstrated by knocking out regulators or constituents of the SZ (14–17).

To explain the increase in osteolytic activity associated to HIV-1-induced bone loss, only a few mechanisms have been proposed: disruption of the immune system (6, 18), increased production of proinflammatory cytokines (19), and direct infection of OC (20). Regarding the immune system, studies from the HIV-1-transgenic rat model revealed that bone damage results, in part, from an altered production of regulatory factors of osteoclastogenesis secreted by B cells (18). This modified cytokine profile correlates with some bone mineral defects in non-treated HIV-1-infected patients (6). Along with CD4⁺ T lymphocytes, macrophages serve as primary target cells for HIV-1 in vivo (21–23). Because OC share a common myeloid origin with macrophages, the last proposed hypothesis is that OC are targets for HIV-1, and their infection would then contribute to bone loss. Indeed, it has recently been shown that HIV-1 may replicate in vitro in human monocyte-derived OC and enhance their bone resorption activity (20). However, the relevance of this observation has yet to be provided in vivo, along with the corresponding cellular and viral mechanisms involved in the bone resorption process.

Here, we report the occurrence of infected OC in bones of HIV-1-infected humanized mice and in human synovial explants exposed to HIV-1. We further demonstrate that the exacerbated osteolytic activity of infected OC results from modified structure and function of the SZ, correlates with Src activation, and is dependent on the viral protein Nef.

Results

Infected OC Are Found in the Bones of HIV-1-Infected Humanized Mice and in Human Synovial Explants Exposed to HIV-1. We first investigated whether OC are infected in vivo. Because bone marrow/liver/thymus (BLT) humanized mice infected with HIV-1 reproduce most hallmarks of infection in humans (22, 24, 25), we used these mice infected for 14–21 d (2×10^4 – 8×10^4 RNA copies/mL in blood, $n = 4$) to examine the growth plate of femurs and tibias, the OC-enriched zone. For each bone section (head of femur or tibia from four mice), we quantified 85 ± 22 cells that exhibit OC characteristics (multiple nuclei, TRAP activity, and localization at the bone surface). Importantly, for each infected animal, we identified by immunohistochemistry (IHC) one or two OC positive for the viral protein p24, which is used as an indicator of productive viral infection (Fig. 1*A* and Fig. S1). Negative controls were included for each sample by omitting the primary antibody.

We then assessed whether OC can be infected in human tissue using synovial membrane explants, which contain fibroblasts, macrophages, lymphocytes, dendritic cells, and OC, in an abundant extracellular matrix (26). Fresh human synovial tissues were incubated ex vivo with the HIV-1 macrophage R5-tropic ADA strain and maintained in culture with osteoclastogenic cytokines to keep resident OC and OC precursors alive throughout the experiment. Fifteen days postinfection, OC were characterized by multiple nuclei, TRAP-, and cathepsin K-positivity by IHC. Remarkably, we observed that about 10% of these cells were positive for the viral p24 ($n = 5$ synovial explants examined) (Fig. 1*B*).

Altogether, these data show that infection of OC occurs both in vivo in humanized mice and ex vivo in human tissues.

Human OC Are Permissive to HIV-1 Infection by Cell-Free Viruses at Different Stages of Differentiation. To examine the stage of differentiation at which the cells become capable to be infected, we

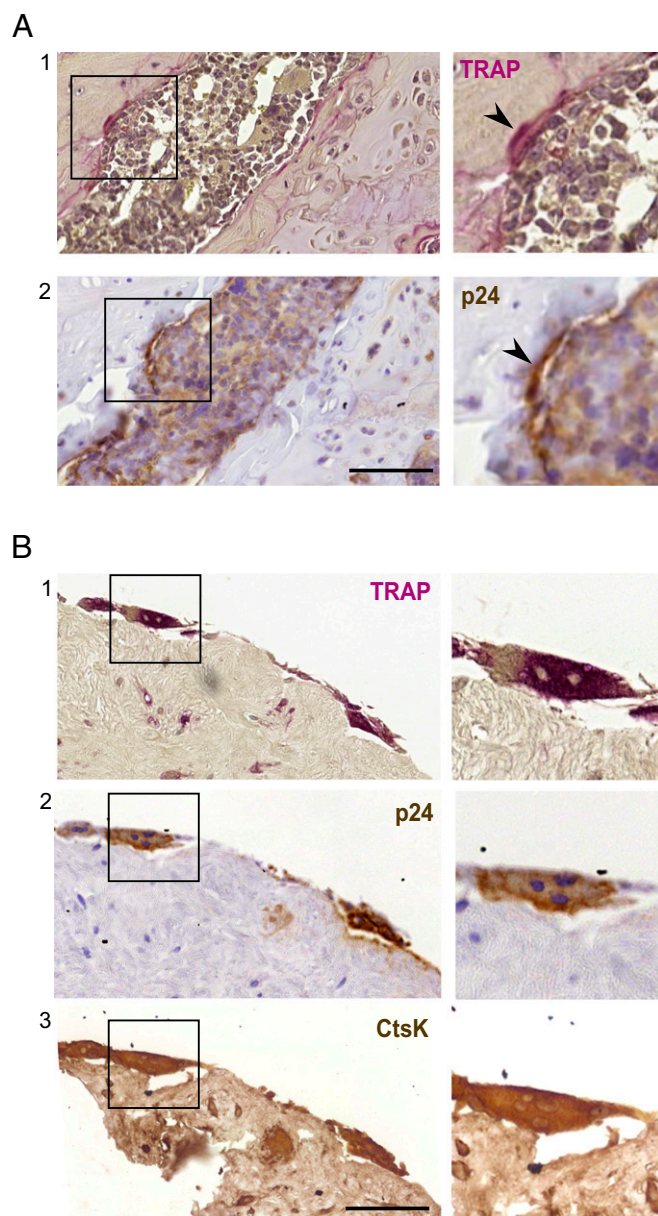


Fig. 1. Tissue OC are infected with HIV-1 in vivo and ex vivo. (A) HIV-1 infects OC in vivo in HIV-1-infected BLT-humanized mice. Two serial sections (3-μm thick, 1 and 2) of the head of a tibia of HIV-1-BLT mice were stained for TRAP activity (in purple, 1) and with a monoclonal antibody directed toward human viral protein p24 (in brown, 2). Representative sections ($n = 4$ mice, two sections per mouse of tibia and femur heads). Arrowheads show an infected OC. (Scale bar, 50 μm.) Enlarged frames, 2x zoom. (B) Human OC are infected with HIV-1 in synovial explants. Pieces of a noninflammatory human synovial tissue were cultured with M-CSF and RANKL and infected with HIV-1 (ADA strain). Two weeks postinfection, histological analyses [TRAP activity in purple (1), p24 (2) and cathepsin K (CtsK) (3) IHC in brown, nuclei in blue] were performed on three serial sections (3-μm thick, 1–3). Representative histological sections ($n = 3$ synovial tissues, four pieces per tissue). (Scale bar, 100 μm.) Enlarged frames, 2.3x zoom.

turned to human OC derived from primary monocytes differentiated in vitro in the presence of M-CSF and RANKL. The OC differentiation process was assessed by measuring OC protein level at different stages. While we observed a rapid increase of TRAP and $\beta 3$ integrin as soon as day 1, we noticed the acquisition of cathepsin K (a late-stage differentiation marker) at day 6; the expression level for all these proteins increased until day

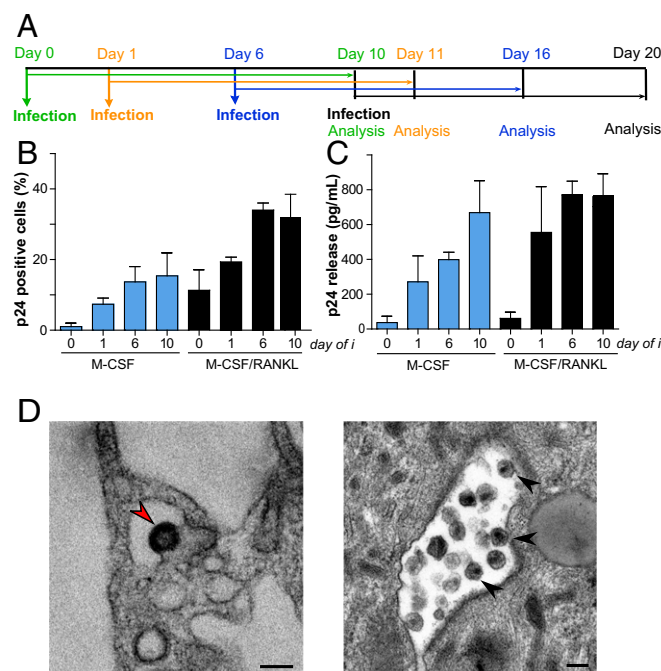


Fig. 2. HIV-1 infects human OC and their precursors in vitro. (A–C) OC are infected by cell-free viruses at different stages of differentiation. (A) Experimental design of infection (ADA strain). (B and C) Kinetics of the percentage of p24⁺ cells (B, determined by IF) and of p24 release in the supernatants (C, determined by ELISA) in cells maintained in M-CSF plus RANKL (OC, black bars) or M-CSF alone (MDM, blue bars) from the same donors are shown. Results are expressed as mean \pm SEM ($n = 3$ donors and 5 donors for day 10). Day of i, day of infection. (D) Transmission electron microscopy images of infected OC. Mature OC were infected with HIV-1 and examined 10 d postinfection. Images of a budding virus (Left, red arrow-head) and viruses contained in a cytoplasmic membrane compartment (Right, black arrowheads show mature viruses). (Scale bar, 100 nm.)

10, when cells presented characteristics of mature OC, including high fusion index, high TRAP and MMP9 activities, and bone degradation activity (Fig. S2 A–E). Of note, monocyte-derived macrophages (MDM) from the same donors differentiated with M-CSF (at day 10) only exhibited undetectable or low levels of OC markers, low fusion index, and lacked osteolytic activity (Fig. S2 B–E). Cells were infected in vitro with the HIV-1 R5 ADA strain at day 0, 1, 6, or 10 of differentiation (Fig. 2A). The extent of HIV-1 infection and replication was evaluated at day 10 postinfection by immunofluorescence (IF) analysis of p24 and quantified by measuring the concentration of p24 released in the supernatant. While monocytes (day 0) were poorly able to sustain infection, cells became increasingly permissive to infection along the differentiation process (Fig. 2B and C, black bars); this correlated with the increased expression of the CCR5 entry coreceptor from day 1 (Fig. S2F). Moreover, we observed that virus production was inhibited by pretreatment of OC with the CCR5 antagonist Maraviroc, the reverse-transcriptase inhibitor AZT, the integrase inhibitor Raltegravir, or the protease inhibitor Ritonavir (Fig. S2G), indicating that the p24 signal corresponds to productively infected cells. We noticed that MDM and OC equally sustained infection (Fig. 2B and C) ($\geq 95\%$ donors supported infection, $n = 29$), which is consistent with similar levels of CD4 and CCR5 receptors expressed during differentiation (Fig. S2F) and at day 10 (20). Moreover, the viral particles produced by infected OC or MDM had comparable infectivity, as assessed using the TZM-bl reporter cell line ($27 \pm 5\%$ of p24⁺ TZM-bl for OC-produced particles vs. $26 \pm 7\%$ for MDM, $n = 5$), indicating that both cell types released infectious

virions. Importantly, HIV-1 did not affect OC viability, as cell density was not altered even at day 20 postinfection ($1,580 \pm 276$ nuclei/mm² for noninfected OC vs. $1,560 \pm 352$ for infected OC, $n = 6$ donors, $\geq 3,000$ cells per condition). Finally, we observed virus budding in OC and both mature and immature virus particles accumulating in membrane-delineated intracellular compartments by electron microscopy (Fig. 2D).

Collectively, these results show that HIV-1 infects and replicates in OC and their precursors, without significant cytotoxic effect.

Human OC Are Preferentially Infected by Transmission from Infected T Cells. HIV-1 spreads by infecting target cells either as cell-free particles or more efficiently via cell-to-cell transmission, both in vitro and in vivo (27–30). We thus examined whether mature OC could be infected by contact with infected CD4⁺ T lymphocytes, first using Jurkat T cells infected with the HIV-1 R5-tropic NLAD8-VSVG strain ($>50\%$ of infected T cells, $n = 8$). Briefly, after 6 h of contact with OC, Jurkat cells were washed out (more than 99% of the T cells were eliminated) (Fig. S3A) and OC were harvested either immediately (day 0) or 5 d later (day 5) and stained for intracellular viral p24 by IF (Fig. 3A and B). We observed that 6-h contact (day 0) was sufficient for T cell-to-OC virus transfer with about 15% of p24⁺ OC, while no detectable infection was observed at this time point when OC were cultured with cell-free virions produced by T cells. The difference was maintained at day 5. At this time point, the high rate of infected OC could result from the initial infection by T cell-to-OC transmission and from enhanced OC cell fusion. Noticeably, virus transfer via infected T cells led to a productive infection of OC as shown by the amount of p24 detected in the supernatant at day 5 (Fig. 3C), which is higher than in the case of infection with cell-free virions. Finally, we confirmed the efficient virus transfer from infected T cells leading to productive infection of

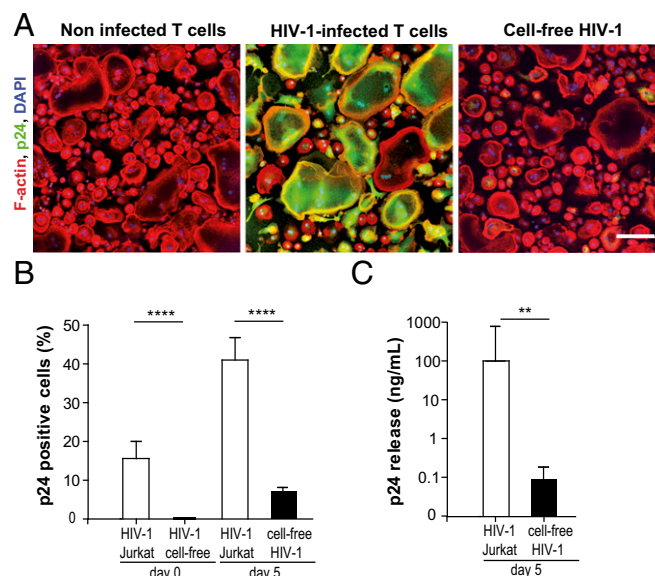


Fig. 3. HIV-1 is transmitted from infected T lymphocytes to OC. (A–C) OC have been in contact for 6 h with noninfected Jurkat T lymphocytes (Left), with HIV-1-infected T lymphocytes (Center), or with cell-free viral particles produced by T cells during 6 h (Right); they were washed, then harvested immediately (day 0) or 5 d later (day 5). (A) Representative mosaic of 3×3 confocal fields of original magnification 20 \times , after staining for p24 (green), F-actin (phalloidin-Texas red, red), and nuclei (DAPI, blue) at day 5. (Scale bar, 50 μ m.) (B and C) Quantifications (mean \pm SD, $n = 8$ donors) of the percentage of p24⁺ cells evaluated by IF (B) immediately (day 0) or 5 d (day 5) postinfection and of p24 release in the supernatants determined by ELISA after 5 d (day 5) (C). ** $P \leq 0.01$; **** $P \leq 0.0001$.

OC by using autologous infected T lymphocytes as virus donor cells (Fig. S3 B and C).

These results show that OC are not only infected by cell-to-cell transfer via infected T cells, but it is also more efficient than infection by cell-free virions.

HIV-1 Infection Enhances OC Precursor Migration and OC Bone Resorption Activity. In HIV-1 transgenic rats, severe bone loss has been correlated to an increase in the number and size of OC (18). This increase could reflect an enhanced recruitment of OC precursors or a stimulated OC differentiation. We tested both hypotheses. It is known that migration of OC precursors from blood to bones requires proteases *in vivo* (31), and that defects in the 3D protease-dependent mesenchymal migration of these cells *in vitro* correlates with lower recruitment of OC to bones (16). For these reasons, we assessed whether HIV-1 infection alters OC precursor mesenchymal migration (32, 33). Human OC precursors were infected at day 1 of differentiation, seeded at day 2 on Matrigel, and migration was measured 48 h later (Fig. 4). Of note, OC precursors inside Matrigel exhibited the characteristic elongated shape of the mesenchymal migration (32). The percentage of migrating cells and the distance covered by the cells were both significantly increased upon HIV-1 infection.

Next, to test whether HIV-1 infection affects OC differentiation, we examined the consequences of infection on the extent of OC fusion and bone resorption activity. HIV-1 infection significantly enhanced cell fusion, as measured by the fusion index and the area covered by infected versus noninfected OC (Fig. 5A), the percentage of TRAP⁺ multinucleated cells, and the number of nuclei per multinucleated cell (Fig. S4 A and B). When cells were treated with Maraviroc before infection, the fusion index was reduced to a similar level to controls (Fig. S4C). Moreover, to explore the effects of OC infection on bone resorption activity, we characterized the morphology of the resorption lacunae. The total bone resorption area increased upon HIV-1 infection (Fig. 5 B–D) and resorption pits displayed profound morphological modifications (Fig. 5 B–F). Infected OC generated resorption pits that appeared deeper ($28 \pm 1.2 \mu\text{m}$ vs. $17 \pm 0.7 \mu\text{m}$ for controls, $***P < 0.0001$) (Fig. 5C) and less elongated (Fig. 5E) than those of noninfected OC, which form resorption trails reminiscent of “inchworm-like migration” (34). These modifications correlated with a significant up-regulation of the bone

volume resorbed per pit in the HIV-1 infection context (Fig. 5F). Furthermore, we also found a significant increase in the concentration of the C-terminal telopeptide of type 1 collagen (CTX) released in the supernatants and used as an additional marker of bone resorption (Fig. 5G).

Osteolytic activity is mediated by acidic dissolution of the minerals and enzymatic digestion of the organic components (35). HIV-1 infection enhanced these two activities, as evidenced by the increase in the capacity of OC to dissolve minerals (Fig. 5H) and release TRAP (Fig. 5I). No variation in protein expression and activity was noticed regarding secreted cathepsin K and MMP9 (Fig. 5 J and K).

Finally, we examined OC attachment/detachment, a critical factor for bone degradation (12), given that OC resorption partly proceeds through a succession of migratory phases alternating with bone resorption stationary phases (12, 36). When infected, OC were more resistant to detachment induced by Accutase treatment than noninfected counterparts (Fig. 5L). This increased adhesion likely slows down OC motility on bone, which should contribute to the modified morphology of resorption lacunae and to the higher bone degradation activity.

All in all, these results indicate that HIV-1 infection enhances the 3D migration of OC precursors, which may favor recruitment of OC to bones, as well as the adhesion and bone resorption activity of mature OC.

HIV-1 Infection Alters the Architecture of the SZ and Activates Src Kinase. Because the SZ has been related to adhesion and bone degradation capacities of OC (11, 37), we characterized the architecture of this structure in infected OC. We observed that in OC seeded on bones, the number of cells harboring SZ was increased (Fig. 6 A and B). This effect was not duplicated by infection with *Mycobacterium tuberculosis*, *Francisella novicida*, or *Salmonella typhimurium* (infection rates $\geq 50\%$ for each bacteria, percentage of cells harboring SZ: 18 ± 9 for *M. tuberculosis*, 22 ± 6 for *F. novicida*, 25 ± 4 for *S. typhimurium* vs. 25 ± 10 in noninfected OC, mean \pm SD, $n = 4$), suggesting that this parameter was not generally influenced by OC infection. In addition, the size of the SZ was increased (Fig. 6 A and C), delineating an area corresponding to $25 \pm 1\%$ of the cell surface of infected OC vs. $18 \pm 2\%$ in control cells (Fig. 6D). We also noticed that the fluorescence intensity of F-actin was higher in the SZ of infected cells (Fig. 6 A and E) and the F-actin core of podosomes, the basal element of the SZ, was larger (Fig. 6 F and G). The tyrosine kinase Src plays a key role in bone homeostasis by controlling the formation and stability of the SZ and the rate of actin turnover within OC podosomes (14, 16, 38). Consequently, OC from *Src*^{−/−} mice do not assemble functional SZ and the mice exhibit a strong osteopetrotic phenotype (15). Interestingly, we showed that in the context of HIV-1-infected OC, the activity of Src kinase family was enhanced as measured by phosphorylation of the regulatory tyrosine (Fig. 6H).

These results show that the increase in bone adhesion and resorption observed in infected OC is associated with larger and more numerous SZ as well as higher Src kinase activity.

The Viral Factor Nef Is Involved in HIV-1-Induced Effects on OC Both *In Vitro* and *In Vivo*. To better understand the viral mechanisms involved in HIV-1-induced effects in OC, we focused on the viral accessory factor Nef because it is known, among other functions, to modulate F-actin organization and to stimulate both the kinase activity of Src (39–44) and cell fusion (45). To this end, we used *wt* HIV-1 and *nef*-deleted HIV-1 ($\delta\text{nefHIV-1}$) ADA strains that present the same level of infectivity in macrophages (33, 45–47) and in OC (Fig. S5A). Importantly, the viral particles produced by OC infected with the *wt* or mutant strains showed the same level of infectivity ($26 \pm 7\%$ of p24⁺ TZM-bl cells for the *wt* strain vs. $22 \pm 5\%$ for $\delta\text{nefHIV-1}$, $n = 4$). As shown in Fig. 7A,

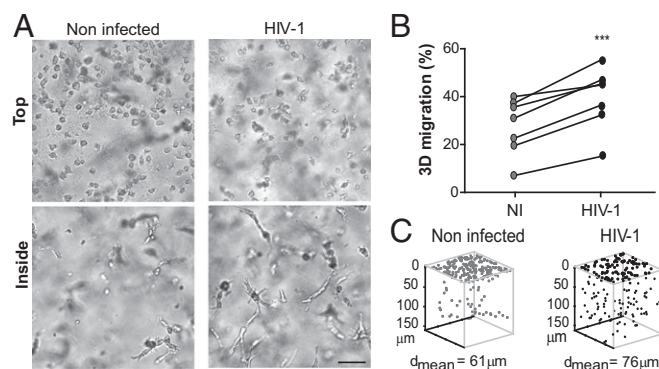


Fig. 4. HIV-1 enhances 3D migration of OC precursors in Matrigel. OC precursors were infected or not at day 1, seeded at day 2 on thick layers of Matrigel polymerized in transwell chambers, and migration was measured 48 h later. (A) Representative brightfield images of live cells either at the surface (top) or within the matrix (inside), taken after 48 h of migration using an inverted video microscope. (Scale bar, 50 μm .) (B) The percentage of migrating cells was measured (mean \pm SEM, $n = 7$ donors) (NI, noninfected). $***P \leq 0.001$. (C) Three-dimensional positions of OC precursors in the matrix and mean distance of migration (d_{mean}) from a representative experiment of seven are shown using the TopCat software.

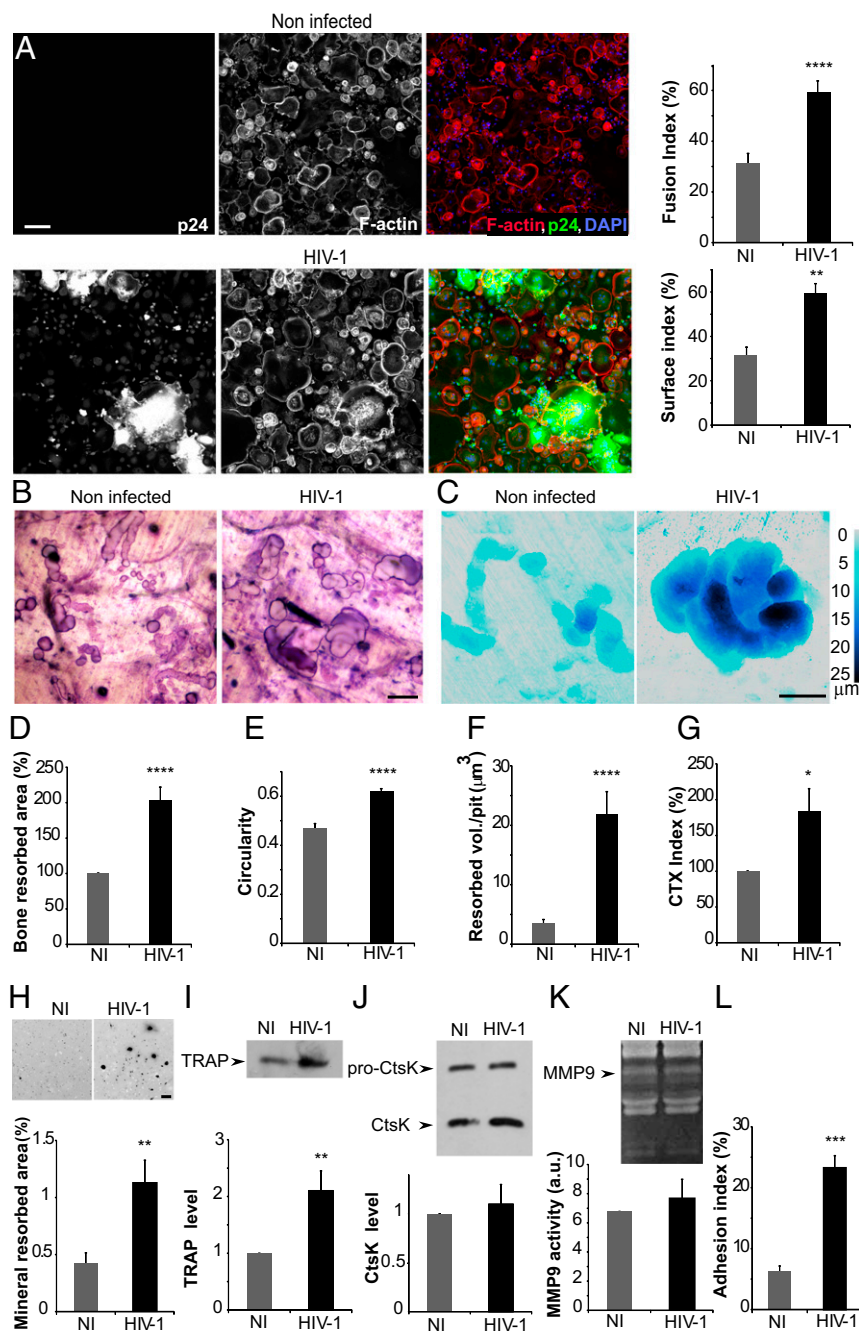
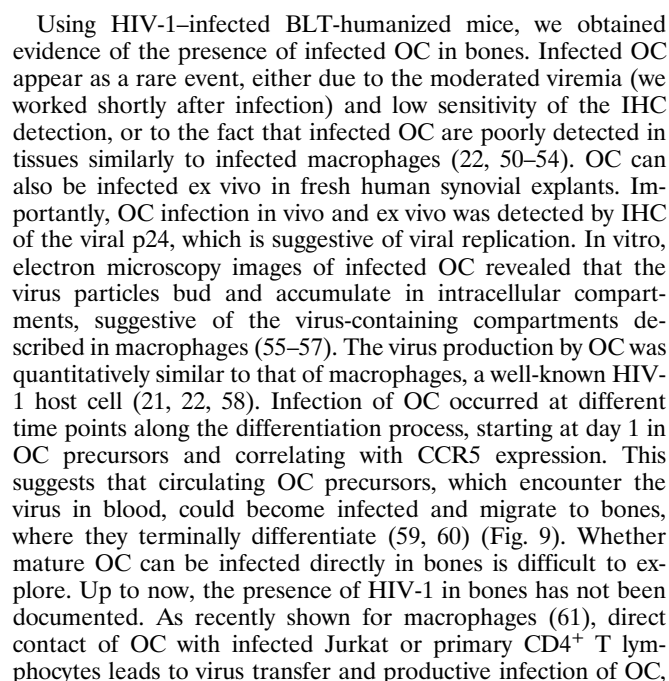


Fig. 5. HIV-1 enhances OC fusion, osteolytic activity, and adhesion of OC. (A) HIV-1 triggers OC fusion. Ten days postinfection, cells were stained for p24 (green), F-actin (red), and nuclei (blue). Images from a mosaic of 4×4 confocal fields of original magnification 20 \times . (Scale bar, 150 μm .) The cell fusion index (nuclei number in multinucleated cells/total nuclei number) and the surface percentage covered by multinucleated cells (surface index) were measured for >3,000 cells in each condition, $n = 6$ donors. Results are expressed as mean \pm SEM. (B–G) HIV-1 infection increases the bone resorption activity of OC. Day 10-noninfected and HIV-1-infected OC were seeded on cortical bone slides for 48 h. Then, OC were removed, the supernatants collected, and the bone slices stained with Toluidine blue to visualize resorption pits in purple. Data were obtained from six donors. (B) Representative brightfield images of bone-resorption pits (purple). (Scale bar, 20 μm .) (C) Representative confocal images of pits. Color codes for the depth of resorption pits (color scale bar). (Scale bar, 20 μm .) Quantification of the percentage of degradation area (D) and circularity (E) from brightfield images and resorbed volume (F) from confocal images are shown. In D, the degradation area of noninfected OC, normalized to 100%, corresponded to 9% of the total area. (G) The concentration of CTX in the supernatant was measured by ELISA and normalized to 100% for non-infected cells (mean CTX concentration = 1,790 pg/mL), $n = 6$ donors. (H–K) Effects of infection on mineral dissolution and extracellular osteolytic enzymes. (H) Day 10-noninfected and HIV-1-infected OC were seeded on crystalline inorganic calcium phosphate-coated multiwells. The cells were removed and the wells stained to reveal demineralized area (black). (Scale bar, 50 μm .) Graph shows the area covered by mineral dissolution pits from 10 fields per condition and per donor, $n = 3$ donors. (I–K) The supernatants of noninfected and HIV-1-infected OC seeded on glass were collected at day 10. TRAP and CtsK expression levels (Western blot, I and J) and MMP9 activity (zymography analysis, K) were quantified. Protein loading has been controlled by Coomassie blue staining, $n = 4$ donors. (L) HIV-1 infection increases OC adhesive properties. Noninfected and HIV-1-infected OC were removed at day 10 with Accutase for 10 min and the percentage of remaining adherent cells quantified by counting nuclei (adhesion index). Graph represents average of five fields per condition from $n = 3$ donors (NI, noninfected). * $P \leq 0.05$; ** $P \leq 0.01$; *** $P \leq 0.001$; **** $P \leq 0.0001$.

the 3D mesenchymal migration of OC precursors infected with $\delta\text{nefHIV-1}$ was similar to noninfected cells. In mature OC, Src kinase phosphorylation, bone resorption activity, percentage of cells with SZ, fusion index, and SZ area were reduced in $\delta\text{nefHIV-1}$ -infected OC in comparison with cells infected with the *wt* virus (Fig. 7 B–F).

Next, we considered performing ectopic expression of Nef-GFP in OC, but we encountered a technical limitation; only 5% of Nef-expressing cells were obtained, precluding a rigorous quantification of SZ size and bone resorption activity. Nonetheless, when the transfected cells were plated on glass, we observed a fraction of Nef-GFP localized at the SZ on bones and that podosomes occupied a larger area than those of control cells (Fig. 7 G and H), thus mimicking the results obtained with OC infected with the *wt* virus (Fig. 6 F and G).

Next, we took advantage of transgenic (Tg) mice expressing Nef under the regulatory sequence of the human CD4C gene to overcome the transfection difficulty of human OC. The CD4C regulatory element drives Nef expression in CD4 $^{+}$ T cells, macrophages, and dendritic cells (48, 49), and also in OC (Fig. S5B). In fact, in the absence of any available Nef antibody for IHC, we used CD4C/HIV-1 $^{\text{GFP}}$ Tg mice. We verified that the GFP was expressed in all OC, indicating that CD4C drives the expression of ectopic genes in OC (Fig. S5B). To characterize the effects of Nef expression in OC, OC precursors were isolated from bone marrow of Nef-Tg and non-Tg mice and differentiated ex vivo. While the fusion of OC from Nef-Tg mice was not modified compared with OC derived from control mice (Fig. 8A), the bone resorption (Fig. 8B) and the width of F-actin staining in the SZ were enhanced (Fig. 8 C and D), indicating that Nef expression is sufficient to increase the osteolytic activity of OC.



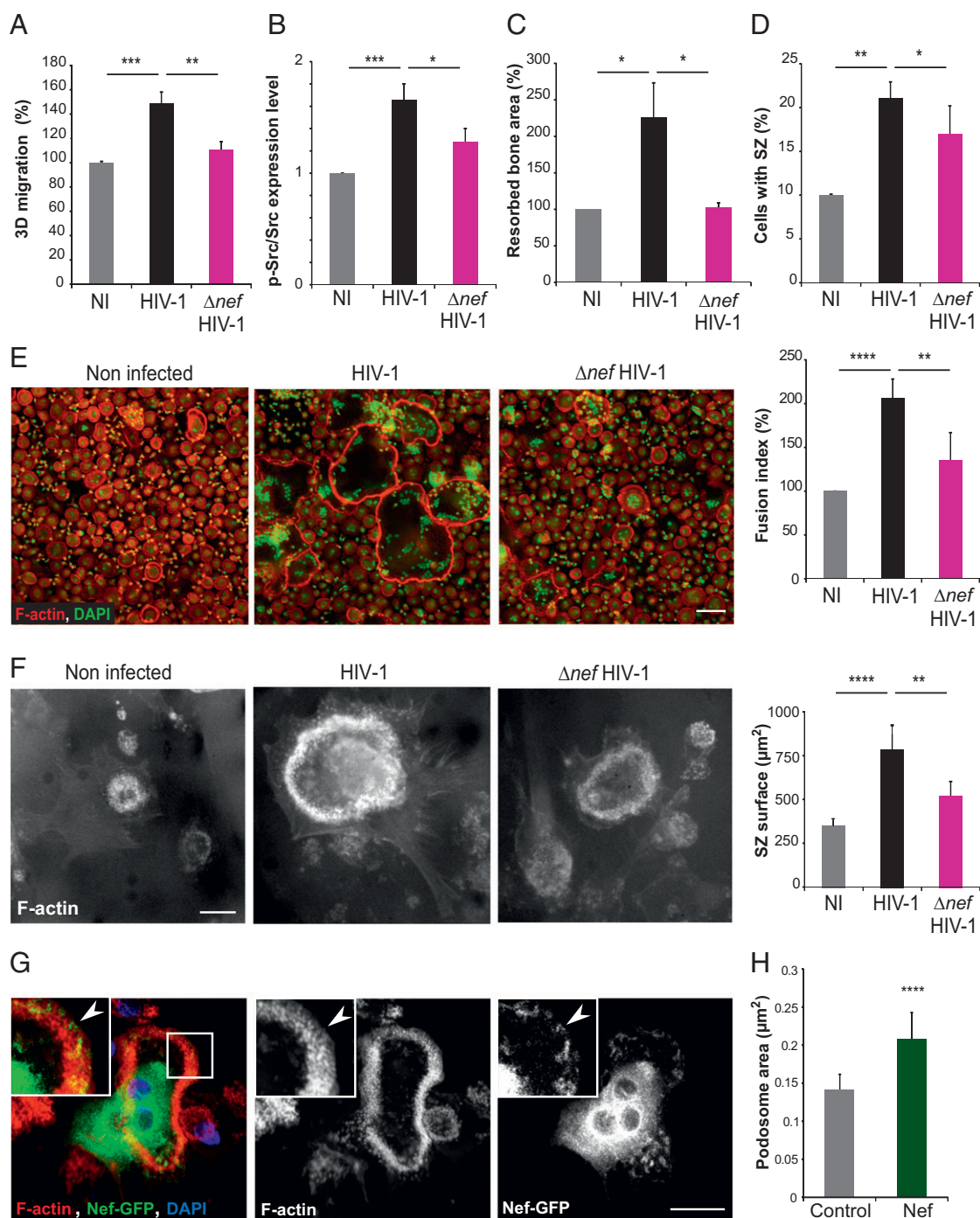


Fig. 7. HIV-1 effects on differentiation and function of OC involve the viral protein Nef. (A–F) Nef is necessary for HIV-1-induced effects in OC. Human OC precursors (A) or mature OC (B–F) were infected with wt HIV-1 or with delta *nef*HIV-1 (NI, noninfected). (A) Percentage of migrating OC precursors after 48 h measured as in Fig. 4, $n = 4$ donors. (B) Quantification of Western blot analyses on whole-cell lysates using antibodies against the phospho-Tyr416 of Src kinases, Src and Actin as in Fig. 6H. Results are expressed as mean \pm SEM, $n = 6$ donors. Quantification of (C) resorbed bone area ($n = 4$), (D) the percentage of cells forming SZ ($n = 4$ donors, 300 cells per donor), (E) the fusion index ($>3,000$ cells per condition, $n = 6$ donors) illustrated by mosaics of 4×4 confocal fields (F-actin in red and nuclei in green), (F) the SZ surface in OC seeded on bones and stained for F-actin (phalloidin), ($n = 3$ donors, >25 SZ). (Scale bars, 150 μm in E, 10 μm in F.) Results are expressed as mean \pm SEM. (G and H) Expression of Nef-GFP in OC. OC were transfected with Nef_{5F2}-GFP or GFP (control) and stained for F-actin (phalloidin, red) and nuclei (blue). (G) A fraction of Nef localizes at the SZ. Confocal images of OC expressing Nef_{5F2}-GFP. Arrowheads show colocalization of Nef-GFP with F-actin at the SZ. (Scale bar, 10 μm , Insets, 2 \times zoom.) (H) Nef expression increases the size of individual podosomes. Automated quantification of the F-actin fluorescence area of individual podosomes. Mean \pm SEM, $n = 4$ donors ($>2,000$ podosomes from over five cells per donor). $*P \leq 0.05$; $**P \leq 0.01$; $***P \leq 0.001$, $****P \leq 0.0001$.

which is clearly more efficient than infection by cell-free viruses. This is likely to be the physio-pathological route to infect OC in situ, which would be consistent with data showing that cell-to-cell

infection is critical for efficient viral spread in vitro and in vivo (25, 29, 61–69). Altogether, these results show that OC are host cells for HIV-1.

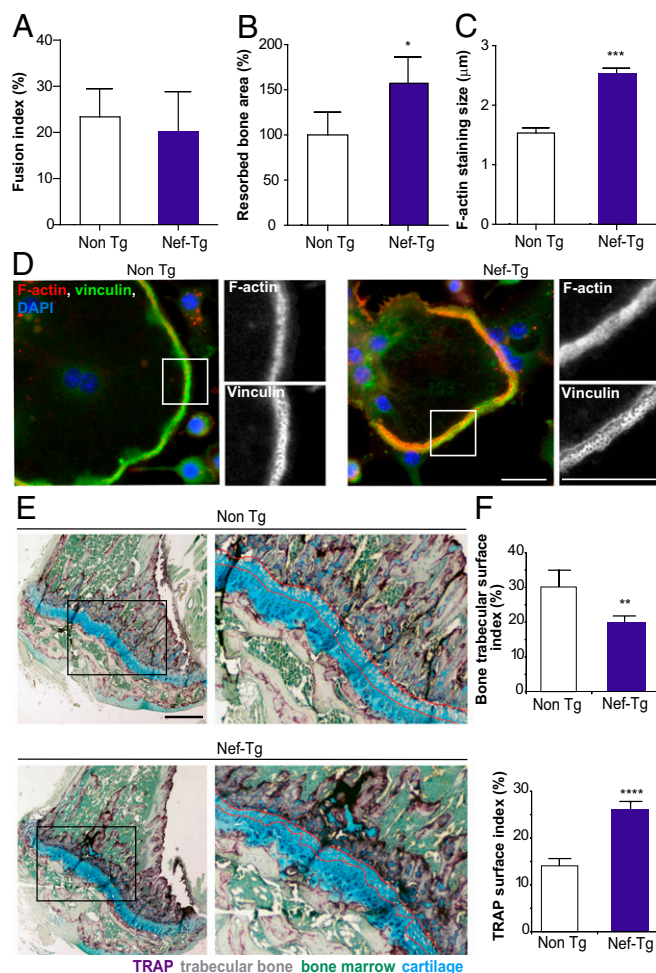


Fig. 8. In Nef-Tg mice, the osteolytic activity of OC and bone defects are enhanced. (A–D) OC differentiated ex vivo from Tg-mice are more osteolytic. OC were differentiated from bone marrow precursors isolated from Nef-Tg and non Tg mice and the fusion index (A), the bone resorption area (B), and the F-actin belt thickness (C) were quantified (50 SZ per condition, $n = 3$ mice per genotype). (D) Representative images of belt structures of OC from non Tg and Nef-Tg mice stained for F-actin (red), vinculin (green) and DAPI (blue). Enlarged frames, 2 \times zoom. (Scale bars, 10 μ m.) (E and F) Nef-Tg mice exhibit bone defects. (E) Representative histological sections of tibia from 7-wk-old mice stained for TRAP to visualize OC (purple), and counterstained with Methyl green and Alcian blue: the bone tissue appears in gray, nuclei in green (corresponding to the nuclei of bone marrow cells), and cartilage in blue. (Scale bar, 200 μ m.) Enlarged frames: $\times 4$ zoom. (F) Quantification of the surface occupied by trabecular bone and surface occupied by TRAP-positive signal in three separate histological sections per mouse ($n = 3$ mice per genotype) are shown. * $P \leq 0.05$; ** $P \leq 0.01$; *** $P \leq 0.001$, **** $P \leq 0.0001$.

Regarding the consequence of HIV-1 infection on OC function, along with this study, there is another report describing that the bone resorption activity of infected OC is exacerbated (20). Herein, we revealed potential mechanisms involved in enhanced bone resorption that concern the structure and function of the SZ, the cell structure instrumental for bone resorption. SZ are larger in HIV-1-infected OC and, consequently, they can degrade larger bone areas. This is in line with the formation of giant OC, which contain twice as more nuclei when they are infected. This exacerbated bone degradation by HIV-1-infected OC also resulted from an increased demineralization process combined with an enhanced secretion of TRAP. The SZ is made of densely connected podosomes, F-actin-rich cell structures involved in cell adhesion, mechanosensing, and cell migration

(70). SZ assembly and patterning are under the control of Src (14, 38). Interestingly, we report that the Src kinase activity was activated in infected OC and that podosomes were enlarged, as visualized by an increase of F-actin staining. In human macrophages, modifications of the F-actin content in individual podosomes have been correlated with fluctuations of protrusion forces exerted onto the extracellular matrix by these cell structures (71). In OC, these modified podosomes may promote more efficient sealing to bone surface. Indeed, we observed stronger adhesion for HIV-1-infected OC compared with noninfected OC. Because the SZ is a barrier that limits the diffusion of acidic and proteolytic molecules released in the resorption lacunae (11–14), increased adhesion would likely enhance the efficiency of containment and favor bone resorption (72). From these results, we propose that modification of several parameters of the SZ (i.e., increased size, adhesion, and degradative activity) contribute to the enhanced osteolytic activity and to the modifications of the topography of resorption pits on infection (12, 36, 73). Pharmacological destabilization of the SZ would reduce the impact of HIV-1 on bone degradation. Several ongoing therapeutic strategies, including the inhibition of the SZ component DOCK5 or cathepsin K activity, are being developed to reduce osteoporotic syndromes while preserving OC viability and differentiation, and thus bone homeostasis (17, 74, 75). We also noticed that OC precursors displayed an enhanced ability to migrate when infected with HIV-1. Interestingly, the efficiency of OC precursor migration has been correlated to OC density in bones (10, 16). Therefore, in addition to enhanced bone resorption activity of infected OC, increased migration of OC precursors should favor OC recruitment to bones, as depicted in Fig. 9, contributing to bone disorders in infected patients.

Nef is a crucial determinant of viral pathogenesis and disease progression. It is known to physically interact with several host proteins to control their activity at the benefit of the virus. Namely, it regulates intracellular protein trafficking (76), actin cytoskeleton (41), cell–cell fusion (45), cell migration (33, 42, 77, 78), and the kinase activity of several members of the Src family (40). In infected OC, all these effects could contribute to the enhanced bone resorption activity that is observed in the present study. In vitro experiments show that Nef, in part located at the SZ, was necessary for all of the HIV-induced effects. The role of Nef was also revealed in vivo in CD4C-Nef-Tg mice that exhibit reduced bone density and an increase of the surface occupied by OC-TRAP staining, suggesting an increase in recruitment and differentiation of OC. Similarly, in HIV-1-Tg rats, a model involving the global transgenic expression of a nonreplicative HIV-1, reduced bone mass was reported, which correlated with a high OC-TRAP staining (18). OC derived from the bone marrow of CD4C-Nef-Tg mice resorbed more and exhibited wider SZ, mimicking the results obtained with human OC infected with HIV-1. Therefore, it is likely that OC participate in the bone remodeling defects evidenced in Nef-Tg mice. Because these mice express Nef in CD4⁺ cells, including T cells, macrophages, OC, and dendritic cells, we propose that the observed bone defects are due, at least in part, to OC expressing Nef in addition to disrupted immune responses, which are known to participate in bone homeostasis (6, 18, 48, 79). Although we do not exclude potential contribution by other viral proteins (80, 81), our results reveal Nef as an essential mediator of the HIV-1 effect on bones (Fig. 9).

It remains to be shown how the virus benefits from manipulating OC. Although OC are giant cells, they do not produce more viral particles than macrophages and these virions exhibit the same infectivity. In contrast with T cells, the cell viability of infected OC is not affected and we can suspect that these infected cells may survive for a long time in bones. Moreover, drug delivery to bones is limited by the unique anatomical features of

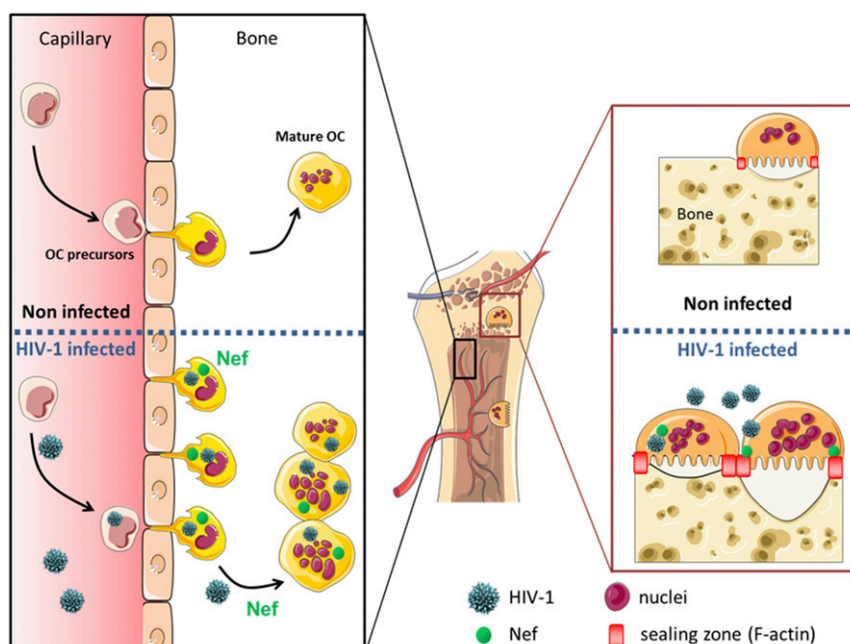


Fig. 9. Graphical abstract proposed to explain HIV-1-induced bone defects in patients. HIV-1 infection affects OC precursor recruitment to bones and the OC differentiation process. These effects, dependent on the viral protein Nef, result in more numerous and more osteolytic OC exhibiting larger and denser SZ.

this tissue (82). Therefore, a putative advantage for the virus may consist in the use of OC as viral reservoirs to hide and survive.

In conclusion, OC are host target cells for HIV-1 that become more osteolytic as a consequence of larger and more degradative SZ. We propose that infected OC participate in bone disorders encountered in HIV-1-infected patients and may constitute a reservoir for the virus. The viral protein Nef appears as a key regulator of the bone resorption activity of OC infected by HIV-1. In summary, this study provides a better understanding of the underlying causes of bone loss following HIV-1 infection.

Materials and Methods

Materials and methods, cell culture and transfection, HIV-1 infection of BLT-humanized mice, viral transfer from infected T cells to primary human OC, cell-free infection of OC and macrophages, and histological analyses of mice bones and of human synovial tissues, are described in *SI Materials and Methods*. Additional materials and methods included flow cytometry analysis, immunoblotting, gel zymography, TRAP staining, IF and transmission electron microscopy analyses, 3D migration, adhesion and resorption assays, chemicals and antibodies, and statistical analysis are also available in *SI Materials and Methods*. Human monocytes were provided by Etablissement Français du Sang, Toulouse, France, under contract 21/PLER/TOU/IPBS01/2013-0042. Experiments with CD4/C HIV Nef mice were approved by the Institutional Animal Ethics committee (Laboratory of Molecular Biology, Clinical Research Institute of Montreal, Montreal, QC, Canada), and experiments with female BLT mice were per-

formed in accordance with guidelines and regulations implemented by the Massachusetts General Hospital Institutional Animal Care and Use Committee.

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