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Human peritoneal macrophages from ascitic fluid can be infected by a broad range of human immunodeficiency virus-type 1 isolates

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Abstract

Macrophages are major HIV target cells. They support both productive and latent HIV-1 infection. Susceptibility of primary macrophages to HIV depends on the anatomical location and activation state of the cells. We demonstrate that peritoneal macrophages (PMs) are abundant in ascitic fluid of patients with liver cirrhosis and are susceptible to HIV-1 infection. PMs expressed CD68, a differentiation marker, exhibited phagocytic activity, and survived in culture for 2 months without additional growth factors. Freshly-isolated PMs were susceptible to HIV-1 R5 strains but not to X4-T cell line adapted (TCLA) strains. Interestingly, after 7 days in culture, PMs acquired susceptibility to X4-TCLA strains. HIV entry inhibitors, TAK779 and AMD3100, blocked HIV infection of PMs, indicating that infection by R5 and X4 strains was mediated by CCR5 and CXCR4, respectively. Although PMs did not express detectable cell surface levels of CXCR4 and CCR5, they did express mRNAs of these HIV co-receptors and responded to stimulation by their natural ligands, SDF-1 α and RANTES. PMs were susceptible to HIV-1 X4, R5, and X4R5 primary isolates. PMs after 7 days in culture produced greater amounts of X4 and X4R5 HIV than freshly-isolated PMs. The day-7 PMs were more susceptible to R5 infection in a single-cycle infection assay, but there was no increase in viral production in a multiple-round infection assay. The level of CXCR4 mRNA and production of CC-chemokines (MIP-1 α , MIP-1 β and RANTES) increased significantly during 7 days in culture. Our results indicate that PMs are susceptible to receptor-mediated infection by a broad range of HIV strains. These primary macrophages could provide a valuable system for investigating the role of primary macrophages in HIV pathogenesis.

Keywords

Human immunodeficiency virus (HIV-1); peritoneal macrophages; CCR5; CXCR4

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Introduction

Macrophages are one of the major cell types in which HIV can both productively replicate and persist in a latent state¹. Macrophages play an important role in HIV-1 pathogenesis because of their ability to generate progeny virions, resistance to HIV-induced killing, and long lifespan¹⁻⁴. In some settings, tissue macrophages contribute significantly to viral load. High levels of HIV-1 RNA are present in macrophages of rhesus macaques infected by a pathogenic simian immunodeficiency/HIV type 1 chimeric virus (SHIV-1) that eliminates CD4⁺ T cells⁵. Macrophages in liver and brain are thought to be the source of the high levels of HIV-1 DNA that are present in certain AIDS patients with very low CD4⁺ T cell counts^{6, 7}. Opportunistic infections, which occur more frequently during the advanced stages of AIDS, increase the amount of HIV produced by macrophage, suggesting that activated macrophages may be a major source of the X4-utilizing variants that often emerge in patients with advanced disease^{8, 9}. Further studies regarding HIV-infected macrophages are needed, as this information may lead to more effective strategies for HIV eradication¹⁻⁴.

The susceptibility of macrophages to HIV-1 infection varies depending on their location in the body and on the strain of the virus. HIV-infected Kupffer cells (hepatic macrophages) are present in patients with AIDS^{6, 7, 10}, and these cells are susceptible to HIV infection *in vitro*¹¹. Macrophages in human tonsils can be infected *ex vivo* by primary HIV-1 X4 and X4R5 dual tropic viruses but not X4-T cell-line adapted (TCLA) strains¹². Alveolar macrophages are susceptible to R5 and X4R5 primary isolates¹³, but intestinal macrophages are not susceptible to HIV-1 infection *in vitro*¹⁴.

Experimental investigation of the interactions between HIV-1 and macrophages has been impeded by the difficulty of isolating human primary macrophages. Monocyte-derived macrophages (MDMs) are often used to study HIV-1 infection; however, the functional properties of MDMs vary depending on the methods used for isolation and cultivation¹⁵⁻¹⁷, and it is unclear to what extent MDMs model the characteristics of tissue macrophages. MDMs typically express abundant CCR5 and minimal CXCR4^{18, 19-21}. They can be infected by both primary X4 and X4R5 dual tropic viruses, but not by X4-TCLA strains.

Primary cultures of peritoneal macrophages (PMs) are a potential system for investigating interactions between HIV-1 and primary macrophages. These cells, which reside in the peritoneal cavity, have distinctive properties, including the ability to suppress T-lymphocyte activation²². PMs from peritoneal fluid of women undergoing diagnostic laparoscopy are known to be susceptible to R5 HIV-1_{BaL}²³; however, the susceptibility of PMs to other HIV strains is unknown and merits investigation. In this study, we developed methods for preparing, culturing, and cryopreserving large numbers of PMs from ascitic fluid of patients with liver cirrhosis. We demonstrate that PMs supported receptor-mediated HIV-1 infection, including that of X4-TCLA strains. HIV production in PMs with X4 and X4R5 primary isolates was enhanced in PMs during 7 days in culture. The level of CXCR4 mRNA was significantly increased in these cultures compared to that in day-1 PMs. Our study indicates that HIV-1 infection of PMs from ascitic fluid provides a novel and useful model for identifying the factors that modulate macrophage susceptibility to HIV-1 infection.

Materials and Methods

Sources of peritoneal macrophages (PMs) and peripheral blood mononuclear cells (PBMCs)

Ascitic fluid (AF) was collected under sterile conditions from patients with liver cirrhosis and refractory ascites who were undergoing therapeutic large volume paracentesis. With approval from the Mount Sinai School of Medicine (MSSM) IRB, subjects gave written consent for

medical record review and AF collection. The etiology of cirrhosis was alcoholic liver disease or hepatitis C virus (HCV) infection; subjects did not have bacterial peritonitis. Ascitic mononuclear cells (AMCs) were prepared by centrifugation at 250 $\times g$ for 15 min followed by Ficoll-Hypaque gradient centrifugation. AMCs at 10×10^6 /ml were cryopreserved by slow freezing cells in media containing 10% (v/v) DMSO and 90% (v/v) FBS at -80°C overnight, followed by storage in liquid nitrogen. AMCs were recovered from frozen stocks by quick thawing at 37°C and transferring in warm RPMI media with 10% FBS followed by centrifugation to remove DMSO. Cells were plated immediately, as described below.

PMs were prepared by plating AMCs in a 48-well plate at 1×10^5 cells per well, culturing for 16 h to allow adherence, and vigorously washing four times with PBS to remove non-adherent cells. Cultures were maintained in RPMI with 10% FBS. To deplete CD3 cells from AMCs prior to plating, freshly-isolated AMCs were treated with 10% human serum for 10 min at room temperature, incubated with CD3 magnetic beads (Miltenyi Biotec, Auburn, CA), and passed over a column. The efficiency of CD3 depletion was confirmed by FACS analysis.

PBMCs from normal healthy blood donors were isolated by Ficoll-Hypaque gradient centrifugation. Monocytes were isolated from PBMCs using CD14 isolation kits (Miltenyi Biotec), placed in dishes coated with human serum. Cells were cultured in RPMI with 20% FBS for 10 days and allowed to differentiate into monocyte-derived macrophages (MDMs).

FACS analysis

Adherent cells were incubated at 4°C in ice-cold PBS for 15 min and then detached using cell scrapers. For surface molecule staining, cells were stained with mAbs conjugated with phycoerythrin (PE), allophycocyanin (APC), or fluorescein isothiocyanate (FITC) (BD Pharmingen). For the intracellular staining of CD68, macrophages were fixed with 2% paraformaldehyde and permeabilized with 0.2% saponin before staining. Appropriate isotype controls were included in all assays. Stained samples were analyzed on FACSCalibur flow cytometer using CellQuest (BD). Results were analyzed with FlowJo software (BD).

Immunofluorescence microscopy

AMCs were plated in dishes containing cover slips or in 8-well chamber slides and cultured in RPMI 1640 with 10% FBS overnight followed by extensive washing. Cells were fixed with 2% paraformaldehyde for 20 min. After washing with PBS, cells were permeabilized with 0.5% Triton X-100 for 10 min, blocked with 3% BSA/0.3% Triton X-100 in PBS for 30 min, incubated with appropriate primary antibodies for 1 h, and washed four times with washing solution (0.3% Triton X-100 in PBS) for 10 min per wash. Cells were then incubated with appropriate secondary antibodies for 1 h, washed four times with washing buffer and then mounted with VECTASHIELD® HardSet™ mounting medium with DAPI. For CD68 detection, anti-CD68 antibody (KP1 at 1:300, DAKO) was used followed by a secondary goat anti-mouse antibody conjugated with Texas-Red. For double staining with HIV p24 and CD68, mouse anti-HIV p24 mAb (clone #24-2) from AIDS Research and Reference Reagent Program (ARRRP), Division of AIDS, NIAID, NIH at a dilution of 1:200 and rabbit anti-human CD68 antibody (Santa Cruz Biotech, Santa Cruz, CA) at a dilution of 1:100 were used followed by donkey anti-mouse Alexa-594 and goat anti-rabbit Alexa-488 at a dilution of 1:400. Images were captured using Zeiss Axiophot2 and analyzed with Image J (<http://rsb.info.nih.gov/ij>).

Phagocytosis

Cells were cultured with FluoSphere carboxylate-modified microspheres (1.0 μm in diameter; Invitrogen, Carlsbad, CA) in RPMI with 10% FBS at 37°C for 2 h, washed with cold PBS, fixed, and permeabilized.

HIV-1 infection

For use in single-cycle infection assays, replication-defective HIV-1_{HXB2} and HIV-1_{JR-FL} Env-pseudotyped reporter viruses expressing luciferase were produced in HEK293T cells, as described previously^{24–26}. Virus particles at approximately 8 ng of HIV p24 per sample in a 48-well plate were used. Macrophages were exposed to HIV-1_{JR-FL} or HIV-1_{HXB2} pseudotyped luciferase reporter viruses for 2 h at 37°C. Unbound virus was then removed by washing. Cells were incubated for 48 h, and lysed with Passive Lysis Buffer (Promega Inc, Madison, WI). Luciferase activity was measured using Promega's luciferase assay systems. In multiple-round infection assays, HIV-1 primary isolates (provided by ARRRP and The UNAIDS Network for HIV-1 Isolation and Characterization, and the DAIDS, NIAID) prepared from PHA-activated PBMCs were incubated with cells for 2 h. Cells were then washed with media and maintained in RPMI containing 10% FBS. HIV-1 p24 levels in the media were measured using the HIV-1 p24 ELISA kit (SAIC Frederick).

Immunoblotting analysis

PMs were pre-treated with or without AMD3100 or TAK 779, specific inhibitors for CXCR4 or CCR5, respectively for 1 h. Cells were then exposed to a CXCR4 ligand, SDF-1 α (Peprotech, Rocky Hill, NJ) at 300 ng/ml, or a CCR5 ligand, RANTES (R&D Systems, Minneapolis, MN) at 200 ng/ml in the presence or absence of inhibitors for 15 min. Whole-cell extracts (WCE) were prepared by lysis of cells in 20 mM HEPES buffer (pH 7.9) containing 0.2% NP-40, 10% glycerol, 200 mM NaCl, 0.1 mM EDTA, 1 mM dithiothreitol, 1 mM sodium orthovanadate, 0.5 mM phenylmethylsulfonyl fluoride, and a protease inhibitor cocktail (Roche Applied Science, Indianapolis, IN). The protein concentration was determined using the Bio-Rad Protein Assay. MAPK (p44/p42) activity was analyzed by immunoblotting on PDVF membranes, as described previously²⁷. Blots were probed with rabbit polyclonal antibodies against phospho-p44/p42 MAPK (Cell Signaling Technology), washed, and reacted with horseradish peroxidase-linked goat anti-rabbit antibody (KPL, Inc. Gaithersburg, MD). Bands were visualized using the Amersham ECL kit. Blots were incubated with RestoreTM Western blot stripping buffer (Thermo Fisher Scientific, Rockford, IL), and re-probed with antibodies against p44/p42 MAPK proteins.

Real-time PCR analysis of CCR5, CXCR4, and CD4 mRNAs

Total RNA was isolated from cells using Qiagen RNeasy[®]Total RNA Mini Kit (Qiagen, Valencia CA) and treated with RNase-free DNase I. To synthesize first-strand cDNA, 500 ng of total RNA, oligo d(T)₁₆ (Invitrogen) at 25 μ g/ml and dNTP at 0.5 mM were incubated at 65°C for 5 min and quick-chilled on ice. Reverse transcription (RT) was performed at 42°C for 50 min using SuperScriptTM II (Invitrogen). The PCR reaction contained cDNA equivalent to 25 ng of RNA input, 200 nM of primer sets and SYBR Green Master Mix (QIAGEN) in an ABI PRISM 7900HT Sequence Detection System (Applied Biosystems, Foster City, CA). Each reaction was performed in triplicate. The primer sequences were: CD4 forward (5'-AAGGGGATACAGTGGAAGTAC-3'), CD4 reverse (5'-GGACCTTTAGTTAAGAAGGAGCC-3'); CXCR4 forward (5'-TACACCGAGGAA ATGGGCTCA-3'), CXCR4 reverse (5'-TTCTTCACGGAAACAGGGTTC-3'); CCR5 forward (5'-AGGGCTGTGAGGCTTATC TTC-3'), CCR5 reverse (5'-CACCTGCATAGCTTGGTCCA-3'); β -actin forward (5'-GTGGACTTGGGAGAGGACTG-3'); β -actin reverse (5'-ACTGGAACGGTGAAGGTGAC-3'). PCR conditions included denaturation at 95°C for 10 minutes followed by 40 cycles of 95°C for 30 seconds, 55°C for 30 seconds, and 72°C for 30 seconds. Quantification of PCR products was normalized using the housekeeping β -actin gene. Relative quantification of gene expression was calculated by using a Δ Ct (Ct, threshold cycle of real-time PCR) method according to the following formula: $2^{-\Delta\text{CT}} = [2^{-(\text{sample Ct} - \beta\text{-actin Ct})}]$.

Luminex fluorescent-bead assay

CC-chemokines measurement was performed by using a Human Chemokine Five-Plex Kit (Invitrogen, Carlsbad, CA) according to the manufacturer's instructions.

Statistical analysis

Statistical analysis was performed using the two-tailed Student's *t* test, with significance set at a *p*-value of <0.05.

Results

Macrophages are the most abundant cell type in AMCs

AMCs were prepared from the ascitic fluid of patients with liver cirrhosis. The yield was approximately $3-9 \times 10^9$ cells for each 3-9 liter specimen. As determined by FACS analysis, over 80% of the AMCs expressed CD68, a marker of differentiated macrophages (Table 1), indicating that ascitic fluid is an abundant source of mature macrophages. In contrast, less than 30% of the PBMCs were CD68⁺ macrophages. The macrophages remained viable and functional following a freeze-thaw cycle of the AMCs (not shown). Comparable results were obtained whenever frozen or freshly-isolated AMCs were used for preparation of peritoneal macrophages (PMs). Experiments, such as those using CD3-depleted cell cultures, were only performed on freshly-isolated cells.

PMs were isolated from AMCs by selecting for adherent cells after 16 h attachment. The majority of the adherent cells had classical macrophage morphology and expressed CD68 (Fig. 1A). FACS analysis revealed that more than 95% of the adherent cells were CD68⁺ (Fig. 1B). The macrophages remained viable for more than two months (data not shown) and efficiently phagocytosed fluorescent beads (Fig. 1C). They produced high levels of TNF α in response to LPS treatment, as did monocyte-derived macrophages (MDMs) (data not shown).

Expression of cell surface CD4, CXCR4 and CCR5 on peritoneal macrophages

PMs were obtained by plating AMCs for 16 h and removing non-adherent cells. FACS analysis revealed that these cells expressed cell surface CD4, albeit at levels far below those of MDMs or freshly-isolated PBMCs (Fig. 2A). Similar to intestinal macrophages and alveolar macrophages^{13, 14}, PMs expressed little, if any, cell surface CXCR4 or CCR5. In contrast, MDMs expressed both cell surface CXCR4 and CCR5 and PBMCs expressed a high level of CXCR4, but undetectable CCR5 (Fig. 2A).

To determine whether PMs expressed functional CXCR4 and CCR5, PMs were exposed to SDF-1 α or RANTES, high affinity ligands for CXCR4 and CCR5, respectively, followed by assessment of MAPK phosphorylation (Fig. 2B). Cells were also treated with small molecule inhibitors of CXCR4 and CCR5, AMD3100 and TAK779, respectively. SDF-1 α and RANTES induced phosphorylation of p44/p42 MAPK in PMs. MAPK activation was blocked by AMD3100 or TAK779 (Fig. 2B). This result demonstrates that PMs express functional CXCR4 and CCR5, even though these receptors could not be detected by FACS analysis.

Freshly-isolated PMs are susceptible to the R5 strain, HIV-1_{BaL}, but not to the X4 T cell-line adapted strain (TCLA), HIV-1_{IIIB}

Freshly-isolated PMs were exposed to the R5 strain, HIV-1_{BaL}, and to the X4 strain, HIV-1_{IIIB}, at multiplicities of infection (MOIs) of 0.01 and 0.1 for 2 h. Unbound virus was removed by washing, and the cells were maintained in RPMI containing 10% FBS. HIV-1 virus particles released into media were measured by HIV p24 ELISA. Freshly-isolated PMs

supported replication of an R5 strain, HIV-1_{BaL}, but were resistant to an X4-TCLA strain, HIV-1_{IIIIB} (Fig. 3A).

To confirm that PMs support receptor-mediated HIV infection, AMCs were depleted of CD3⁺ T cells prior to plating and the HIV susceptibility of the purified PM cultures was examined in a single-cycle infection assay. After overnight plating and selection of adherent cells, PMs were exposed to HIV luciferase reporter viruses pseudotyped with envelopes derived from either the R5 strain HIV-1_{JR-FL} or the X4-TCLA strain HIV-1_{HXB2}. Unbound viruses were removed by washing. Cells were treated with entry inhibitors before and during HIV infection. After cells were cultured for 48 h at 37°C, luciferase activity was measured. PM cultures lacking CD3⁺ T cells were susceptible to the R5 HIV-1 and were resistant to the X4-TCLA virus (Fig. 3B). Infection by the R5 virus was blocked by TAK779, indicating that viral entry was mediated by CCR5. The PMs produced a much lower signal from the pseudotyped R5 virus than MDMs (Fig. 3C). FACS analysis confirmed that the CD3⁺-depleted PM population did not contain CD3⁺ T cells (Fig. 3D).

Peritoneal macrophages acquire susceptibility to the HIV-1 laboratory X4-TCLA strain, HIV-1_{IIIIB}

Because PMs survive for several weeks *in vitro*, it is possible to evaluate changes in HIV susceptibility over time. To seek evidence of altered susceptibility, we cultured PMs in RPMI containing 10% FBS for 7 days and then exposed them to the R5 strain, HIV-1_{BaL}, or the X4-TCLA strain, HIV-1_{IIIIB}, at an MOI 0.1, as described above. Day-7 PMs were susceptible to HIV-1_{BaL} (Fig. 4A). Interestingly, these PMs had acquired susceptibility to HIV-1_{IIIIB}. Comparable amounts of the R5 strain, HIV-1_{BaL}, and the X4-TCLA strain, HIV-1_{IIIIB} HIV were produced from PM cultures, regardless of whether they were derived from AMCs with or without CD3 depletion (Figs 4A and 4B). These results establish that PMs developed susceptibility to the X4-TCLA strain during 7 days in culture. As expected, MDMs were susceptible to HIV-1_{BaL} and released large quantities of virus particles; however, they were not susceptible to HIV-1_{IIIIB} (Fig. 4C).

To further confirm these results, single-cycle infection assays were performed on day-7 PMs prepared from CD3-depleted AMCs (Fig. 4D). Similar to the results in the multiple-round infection assays, these PMs were susceptible to infection with both R5 and X4-TCLA strains (Fig. 4D). The signal from the X4 virus was much lower than that from the R5 virus even though similar amounts of HIV p24 were used during viral exposure. The susceptibility of day-7 PM cultures to the X4-TCLA strain was also shown by immunostaining. In these colocalization experiments, PMs infected by HIV-1_{IIIIB} reacted with antibodies against HIV p24 and CD68 (Fig. 4E) and PMs infected by a pseudotyped HIV-1_{HXB2} GFP-expressing virus expressed GFP and reacted with anti-CD68 antibodies (data not shown).

HIV infection of day-7 PMs by HIV-1 X4 or R5 virus was blocked by AMD3100 or TAK779, respectively, indicating that HIV infection was mediated through HIV co-receptors. Interestingly, in the single-cycle infection assay, the R5 signal from day-7 PMs was significantly higher than that of day-1 cultures (Figs. 3B and 4D); however, in the multiple-cycle infection assay, there was no significant difference in R5 HIV-1_{BaL} production between the two (Figs. 3A, 4A and 4B).

Peritoneal macrophages are susceptible to primary isolates of X4, R5 and X4R5 HIV-1

Day-1 and day-7 PMs were exposed to HIV-1 X4 (92HT599), R5 (92IN905), or dual tropic X4R5 (92RW009) primary isolates for 2 h, washed and cultured in complete media. HIV production was determined by HIV p24 ELISA. Productive HIV infection could be detected

in both day-1 and day-7 PMs (Fig 5). Significant enhancement of viral production occurred in day-7 PMs infected with X4 and dual-tropic X4R5 viruses, but not with R5 viruses (Fig. 5).

CCR5, CXCR4, and CD4 gene expression is up-regulated in PMs during 7 days in culture

It has been shown that over-expression of CXCR4 but not CD4 renders MDMs susceptible to X4-TCLA strains, indicating that high levels of CXCR4 can overcome the block to infection²⁰. Additionally, over-expression of CD4 enhances susceptibility to primary isolates of X4 viruses. These results suggest that higher expression of HIV-1 receptors, and/or enhancement of signal transduction pathways linked to these receptors may underlie the increased susceptibility to X4-TCLA strains that occurred during 7 days in culture. Because PMs become firmly attached to their support surface over time, it was not possible to detach the day-7 PMs for FACS analysis of HIV-1 receptors without causing cell damage. As an alternative, we analyzed mRNA levels of CD4, CCR5 and CXCR4 by real-time PCR in day-1 and day-7 PMs. Although expression of cell surface CCR5 and CXCR4 on the day-1 PMs was undetectable by FACS analysis (Fig. 2), these cells contained detectable levels of CD4, CCR5 and CXCR4 mRNAs in day-1 PMs (Fig. 6). CXCR4 mRNA appeared to be less abundant than CCR5 and CD4 mRNAs. Importantly, the level of CXCR4 mRNA was significantly increased, by approximately 13-fold, in the day-7 cells compared to the day-1 cells. This induction might account for the acquired susceptibility to X4-TCLA HIV-1 observed in the day-7 cells.

PMs produce CC-chemokines during 7 days in culture

While day-7 PMs were more susceptible to HIV R5 infection in a single-cycle infection assay than day-1 PMs, there was no increase in HIV production in the day-7 PMs in a multiple-round infection assay. It is possible that CC-chemokines, natural HIV R5 entry inhibitors, might be induced in cultured PMs. CC-chemokines could block HIV infection in a multiple round infection but enhance viral infection by activation of macrophages in a single-cycle infection assay. To investigate CC-chemokine accumulation in the cell culture media, Luminex assays were used to measure MIP-1 α , MIP-1 β , and RANTES in the supernatants of PMs cultured for 1 or 7 days without a change of the media. Levels of MIP-1 α and MIP-1 β were significantly increased, by 5- to 30-fold, in the day-7 supernatants (Fig. 7A). RANTES was also increased, but to a lesser extent.

HIV infection enhances production of CC-chemokines (reviewed in^{28, 29}). To determine whether HIV infection further increased CC-chemokine production, cells were exposed to HIV-1_{BaL} at MOIs 0.01 and 0.1, washed, and then cultured in RPMI with 10% FBS. One-half volume of the supernatants was collected at various times and the levels of HIV p24 and CC-chemokines were measured. Fresh complete media were added back at each collection. Similar to the results obtained in the study above, which measured the cumulative amounts of these factors, the levels of MIP-1 α , MIP-1 β , and RANTES were increased in the supernatants of day-4 cultures. While HIV at an MOI 0.01 did not affect CC-chemokines production, HIV at an MOI of 0.1 promoted CC-chemokine production (Fig. 7B). An increase in the production of CC-chemokines in PMs after several days in culture may have contributed to the lack of increased HIV production in the day-7 PMs with exposure to replication competent R5 virus.

Discussion

We demonstrated that ascitic fluid from cirrhotic patients contains a large number of primary, differentiated macrophages. These macrophages were phagocytic, produced TNF α in response to LPS, and could be maintained in culture for over two months with only FBS and no additional growth factors or cytokines. Importantly, they are susceptible to infection by a broad range of HIV-1 isolates *in vitro*. It is likely that they differ physiologically from the PMs present in healthy individuals, but they have the advantage of being available in large quantities in a waste

material (ascitic fluid) that is collected without surgery. While patient-to-patient variability is an inherent property of human material, we obtained consistent results when using PMs from different patients. Our data indicate that cultured PMs are a valuable system for investigating interactions between HIV-1 and primary macrophages. These cells also provide a relevant model for investigating HIV-1 infection of macrophages in patients with end stage liver disease due to HCV infection. In the United States, about 30% of HIV-infected patients are also infected with HCV. End stage liver disease is an increasingly important cause of death in the post-HAART era, but few experimental systems are available for investigating HIV/HCV co-infection. Because macrophages are reported to support HCV replication³⁰, PMs could serve as a potential *in vitro* model for HIV/HCV co-infection of primary macrophages.

The ease of maintaining PMs in cell culture allowed us to examine HIV-1 receptor and co-receptor expression over time. Freshly-isolated (day-1) PMs contained mRNAs for CD4 and HIV co-receptors. Stimulation of these PMs with ligands for CXCR4 and CCR5 induced MAPK activation. These cells were susceptible to X4, R5 and X4R5 primary isolates although they did not express detectable cell surface CCR5 or CXCR4. Intestinal macrophages are resistant to HIV infection and alveolar macrophages are susceptible to R5 and X4R5 primary isolates, although neither of these tissue macrophages expresses detectable levels of cell surface CCR5 or CXCR4^{14, 31, 32}. These results show that susceptibility to HIV infection does not correlate with expression of cell surface CCR5 and CXCR4, as determined by FACS analysis³². HIV infection of PMs was mediated by CCR5 and CXCR4 as TAK779 and AMD3100, specific inhibitors for CCR5 and CXCR4, respectively, blocked viral infection. Interestingly, day-7 PMs were susceptible to X4-TCLA strains. Five-day old cultures of Kupffer cells are the only primary tissue macrophages previously reported to be susceptible to X4-TCLA strains¹¹.

The restriction of the X4-TCLA strain has been attributed to the low level of CXCR4 combined with the unfavorable status of cell signaling pathways (reviewed in¹⁸). An increase in the level of cell surface CXCR4 on MDMs by transduction of CXCR4 or treatment with TGF- β increases susceptibility to X4-TCLA or X4-using dual tropic virus, respectively^{20, 33}, suggesting that the levels of CXCR4 and CXCR4-mediated cell signaling modulate the restriction of X4-TCLA strains and X4-using dual tropic viruses. Post-entry restriction of X4-TCLA strains in MDMs or alveolar macrophages occurs at the step of nuclear import^{32, 34}. Here we demonstrated that day-1 PMs were resistant to the X4-TCLA virus but day-7 PM cultures were sensitive. Analysis of the mRNA level of CCR5, CXCR4, and CD4 revealed that expression of CXCR4 was significantly increased in PMs after culturing for 7 days, which could contribute to the acquired susceptibility to the X4-TCLA strain.

R5 viruses are preferentially transmitted during infection of a new host and they dominate during the initial stages of infection^{35, 36–38}. Over time, X4 and dual-tropic viruses arise in ~50% individuals infected with subtype B^{39, 40}. Interestingly, our results demonstrated that day-7 PM cultures produced greater amounts of primary isolates of X4 and X4R5 viruses than day-1 cultures, while production of the R5 viruses was unaltered. Contact with the plastic substrate, oxidative stress, and exposure to soluble factors produced by PMs may have induced changes in permissiveness to the X4 and dual tropic strains. Human PMs produce fibronectin, cytokines, CC-chemokines (MIP-1 α , MIP-1 β and RANTES), and growth factors without exogenous stimulation^{41, 42}. In MDMs, CD40 ligands increase susceptibility to X4-TCLA strains and decrease R5 HIV infection⁴³. Induction of CC-chemokines, MIP-1 α , MIP-1 β and RANTES, contributes to the decrease in R5 HIV infection of MDMs in response to CD40 ligands or LPS^{44, 43, 45–47}. CC-chemokines can activate cell signaling pathways and enhance HIV-1 infection⁴⁸; however, they can also block entry of R5 HIV-1 and decrease susceptibility⁴⁹. Indeed, we found that CC-chemokines were induced in PMs during culture, which could

explain why HIV infection was increased in 7 day-old PM cultures in a single-cycle infection assay but the HIV p24 level was not elevated in a multiple-round infection assay.

The intensity of macrophage infection is strongly influenced by clinical status^{8, 50}. Inflammation, neoplasia, and opportunistic infections increase the viral load of HIV-1-infected tissue macrophages. Macrophage infection is often associated with advanced disease. PMs provide an opportunity to rigorously investigate whether susceptibility to HIV-1 infection is heightened by exposure to inflammatory cytokines and/or by microbial infection. Our unpublished data indicated that HIV DNA was detectable in PMs of patients with undetectable HIV-1 viral load. Analysis of the HIV-1 envelope sequences in these cells could shed light on the potential of macrophages to provide an HIV-1 reservoir.

In summary, large quantities of PMs can be obtained from ascitic fluid. These cells are susceptible to receptor-mediated HIV-1 infection. PMs provide a new *in vitro* model for molecular studies of interactions between HIV-1 and primary macrophages. Understanding the cellular determinants of HIV-1 susceptibility may provide insights into the increased permissiveness of human tissue macrophages to X4-using strains that has been reported during late-stage HIV-1 disease, establishing the foundation for developing new interventions.

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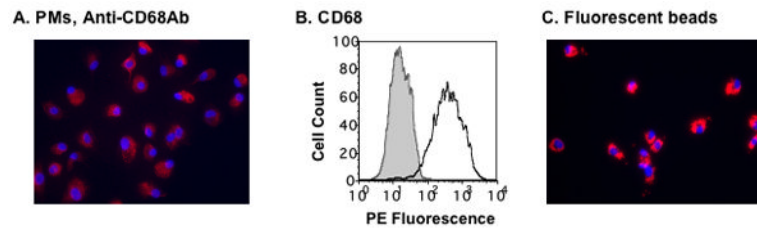


Figure 1. Peritoneal macrophages (PMs) are highly abundant in ascitic fluid

(A) PMs were prepared by plating AMCs ($1-2 \times 10^6$ per well) in 6-well plates overnight, followed by washing the cultures four times with PBS. Adherent cells, PMs, were cultured in RPMI with 10% FBS and stained with a mouse monoclonal antibody against CD68, a marker of differentiated macrophages. (B) The expression of CD68 in day-1 PMs was determined by FACS analysis. The gray area represents the signal from cells stained with isotype control antibody, whereas the solid line indicates the signal from cells stained with CD68 antibody. (C) Day-1 PMs phagocytosed Fluoresbrite particles (6.0 μ m in diameter). Panels A, and C are 20x magnification. The results represent experiments carried out on cells from three or more patients.

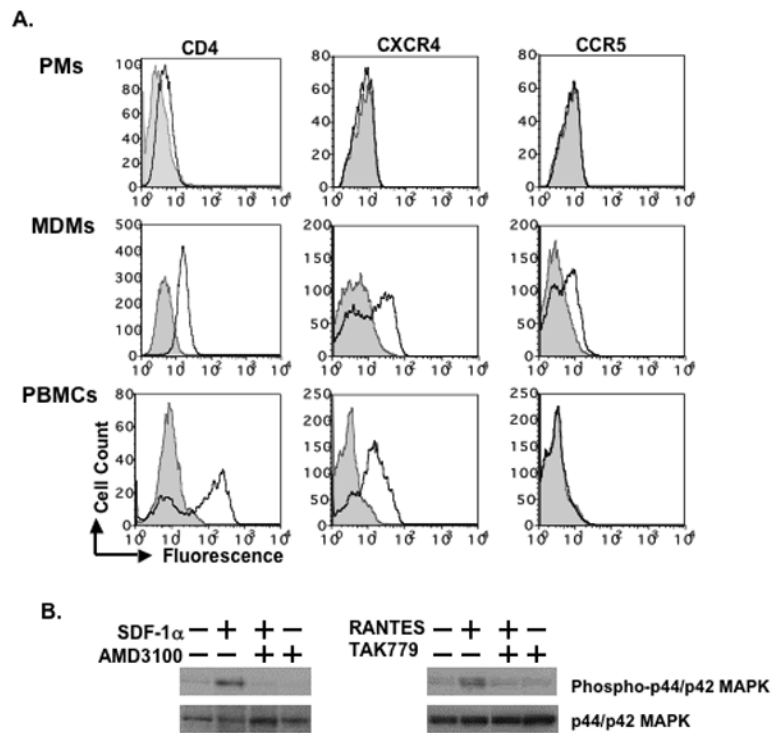
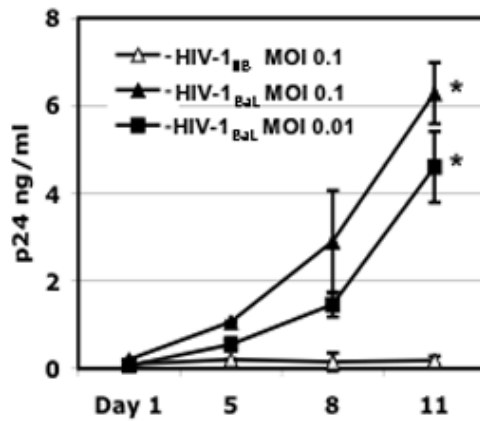


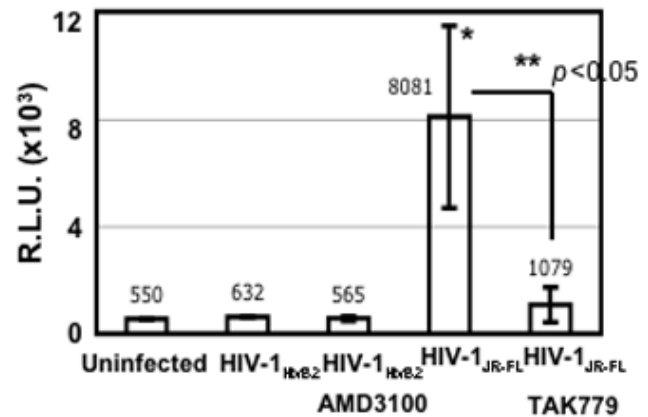
Figure 2. Cell surface expression of CD4, CXCR4 and CCR5 on day-1 PMs

(A) Cell surface expression of CD4, CXCR4 and CCR5 was determined by FACS analysis. MDMs and PBMCs were included as a comparison. The gray area represents the signal from cells stained with isotype control antibody, whereas the solid line indicates the signal from cells stained with CD4, CXCR4 or CCR5 antibody. (B) To examine whether PMs expressed functional CXCR4 or CCR5, cells were exposed to SDF-1 α at 300 ng/ml or RANTES at 200 ng/ml for 15 min, respectively. PMs were also pre-treated with or without specific inhibitors of CXCR4 and CCR5 (AMD3100 or TAK779 at 10 μ M, respectively) for 1 h followed by stimulation with SDF-1 α or RANTES in the presence of inhibitors. Unstimulated PMs were included as controls. Whole cell extracts were prepared and phosphorylation of MAPK (p44/p42) was determined by Western blotting. Blots were then stripped and re-probed with antibodies against p44/p42 MAPK. The results represent three independent experiments using PMs from different patients. Similar data were obtained when day-1 PMs from frozen or freshly-isolated AMCs were used.

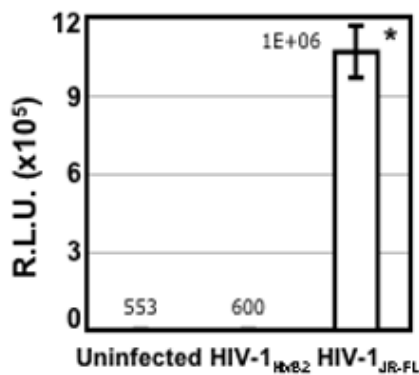
A. Freshly-isolated PMs



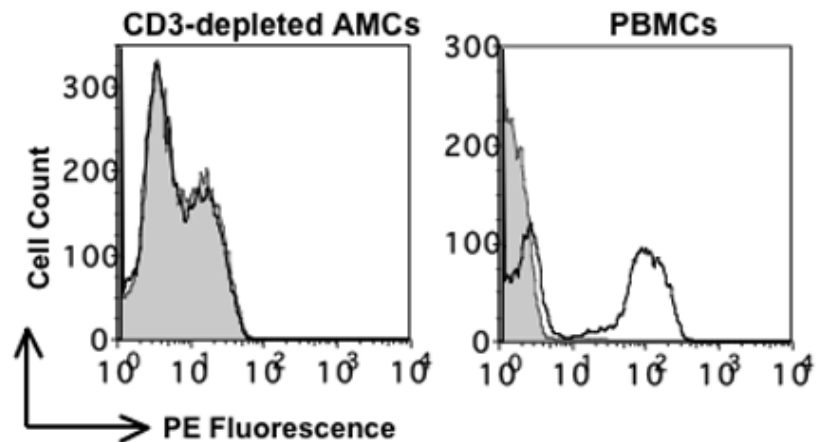
B. Freshly-isolated PMs from CD3-depleted AMCs



C. MDM

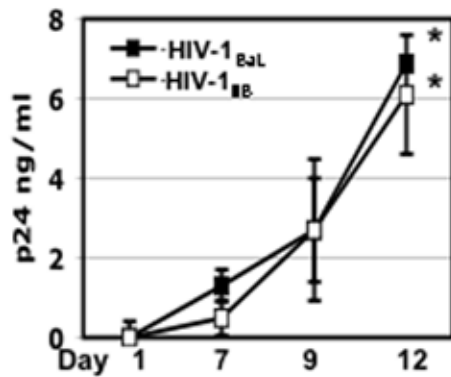


D. Anti-CD3 Ab staining

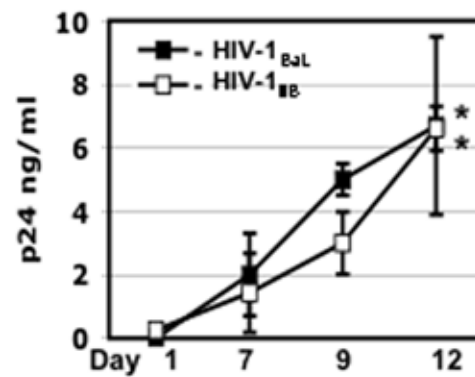
Figure 3. Day-1 PMs are susceptible to R5 HIV-1_{BaL} but not to X4 HIV-1_{IIIIB}

(A) Freshly-isolated (day-1) PMs were exposed to HIV-1_{BaL} or HIV-1_{IIIIB} for 2 h. Unbound virus was removed by washing with PBS, and cells were cultured in RPMI with 10% FBS without additional growth factors. HIV-1 infection was monitored by measuring virus particles in media using an HIV-1 p24 ELISA. The difference in the HIV p24 level from HIV-1_{BaL}-infected cells at day 1 vs day 11 after viral infection was significant ($*p < 0.05$). (B) Freshly-isolated PMs from CD3-depleted AMCs were exposed to pseudotyped X4 HIV-1_{HxB2} or R5 HIV-1_{JR-FL} luciferase reporter viruses for 2 h. Cells were washed and cultured for 48 h before measurement of luciferase activity. To determine sensitivity to inhibitors, PMs were treated with entry inhibitors (AMD3100 or TAK-799 at 10 μ M) for 1 h before viral exposure. Inhibitors were added back to the culture during and after viral exposure. The mean value of luciferase activity is shown. $*p < 0.05$, uninfected control vs HIV-1_{JR-FL}-exposed cells; $**p < 0.05$ for HIV-1_{JR-FL}-exposed cells in the presence or absence of TAK-799. (C) MDMs were exposed to pseudotyped X4 HIV-1_{HxB2} or R5 HIV-1_{JR-FL} luciferase reporter virus and HIV infection was determined at 48h after infection. $*p < 0.05$, uninfected control vs HIV-1_{JR-FL}-exposed cells. (D) The CD3⁺ population in AMCs after CD3 depletion was determined by FACS analysis. Freshly-isolated PBMCs were included as a comparison. For experiments in panels A to C, data are mean \pm SD of triplicate samples and represent three independent experiments from different donors.

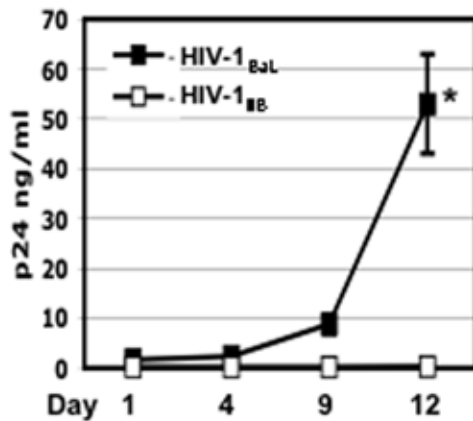
A. PMs, 7 days



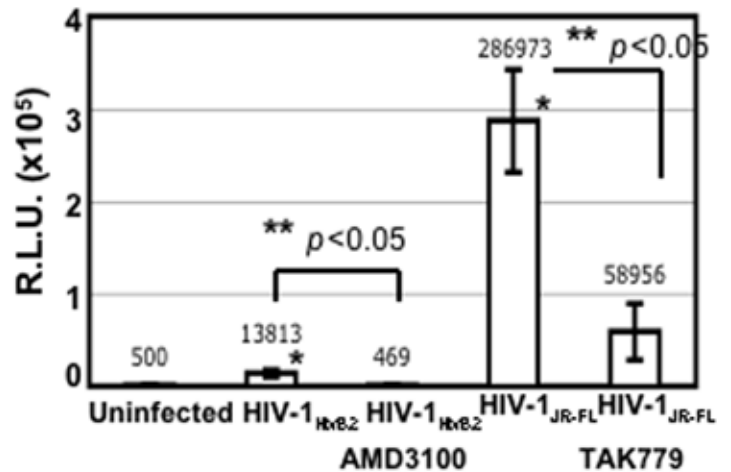
B. PMs from CD3-depleted AMCs, 7 days



C. MDMs



D. PMs from CD3-depleted AMCs, 7 days



E.

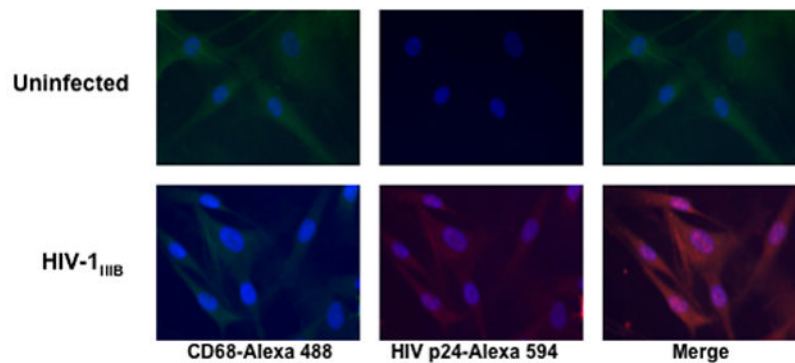
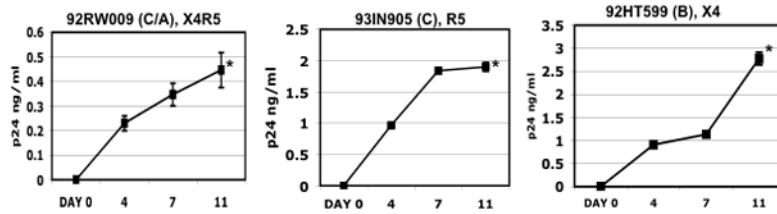


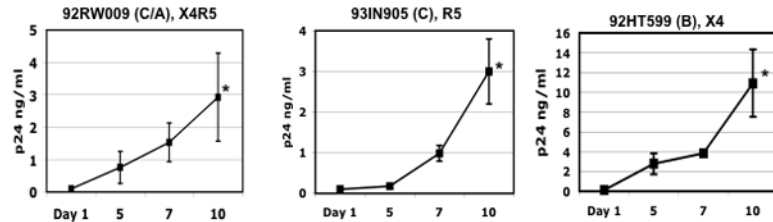
Figure 4. PMs acquire susceptibility to the HIV-1 laboratory X4-TCLA strain
 (A and B) PMs from AMCs with or without CD3 depletion were cultured for 7 days before exposure to HIV-1_{BaL} or HIV-1_{IIB} for 2 h. Unbound virus was removed by washing with PBS,

and cells were cultured in RPMI with 10% FBS. HIV-1 infection was determined by HIV-1 p24 ELISA. The difference in the HIV p24 level at day 1 vs day 12 after infection was significant for both strains ($*p < 0.05$). (C) As a comparison, MDMs were exposed to HIV-1_{BaL} or HIV-1_{IIIIB} and HIV production was determined by ELISA. $*p < 0.05$, HIV-1_{BaL} at day 1 vs day 12 after infection. (D) PMs from CD3-depleted AMCs were cultured for 7 days before exposure to pseudotyped X4 HIV-1_{HXB2} or R5 HIV-1_{JR-FL} luciferase reporter viruses for 2 h. Cells were washed and cultured for 48 h before measurement of luciferase activity. To determine sensitivity to inhibitors, PMs were treated with entry inhibitors (AMD3100 at 10 μ M and TAK-799 at 10 μ M) for 1 h before viral exposure. Inhibitors were added back to the culture during and after viral exposure. The difference in the luciferase activity between uninfected cells and HIV-1_{HXB2} PMs ($*p < 0.05$) as well as HIV-infected PMs with or without inhibitors ($**p < 0.05$) was significant as determined by two-tailed student *t* test. (E) PMs were cultured for 7 days before exposure to HIV-1_{IIIIB} at an MOI 0.1. Unbound virus was removed by washing; cells were cultured for 10 days and stained with antibodies against HIV-1 p24 and CD68 and appropriate secondary antibodies (20x). For experiments in panels A to D, data are the mean \pm SD of triplicate samples and represent at least three independent experiments using PMs from different patients.

A. Freshly-isolated PMs



B. PMs cultured for 7 days

**Figure 5. PMs are susceptible to HIV-1 X4, R5, X4R5 primary isolates**

Freshly-isolated (day-1 PMs) (Panel A) or day-7 cultures of PMs (Panel B) were exposed to HIV-1 X4, R5 and X4R5 primary isolates for 2 h at 37°C and then unbound virus was removed by washing. The names and tropisms of virus isolates are indicated and viral genotypes are shown in parentheses. HIV-1 production was determined by HIV p24 ELISA. Data are the mean±SD of triplicate samples and represent two independent experiments using peritoneal macrophages from different patients. The difference in the HIV p24 level at day 1 vs day 10 or 11 after infection was significant (* $p < 0.05$).

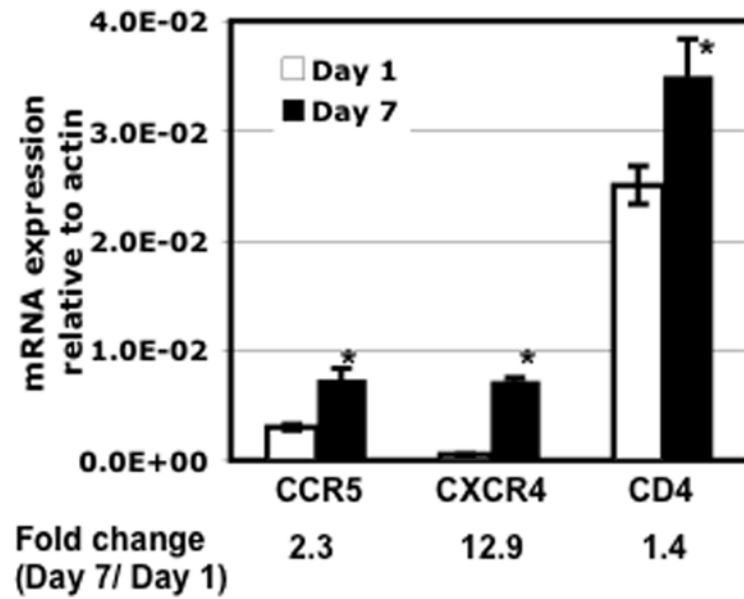


Figure 6. CD4, CCR5 and CXCR4 gene expression is up-regulated in PMs after 7 days in culture
Total RNA was prepared from PMs after culturing for 1 or 7 days. The levels of CD4, CCR5, and CXCR4 mRNA were determined by real-time PCR analysis. The results represent three independent experiments using PMs from different patients. The difference in the level of gene expression from cells after culturing for 1 day vs 7 days was significant ($*p < 0.05$).

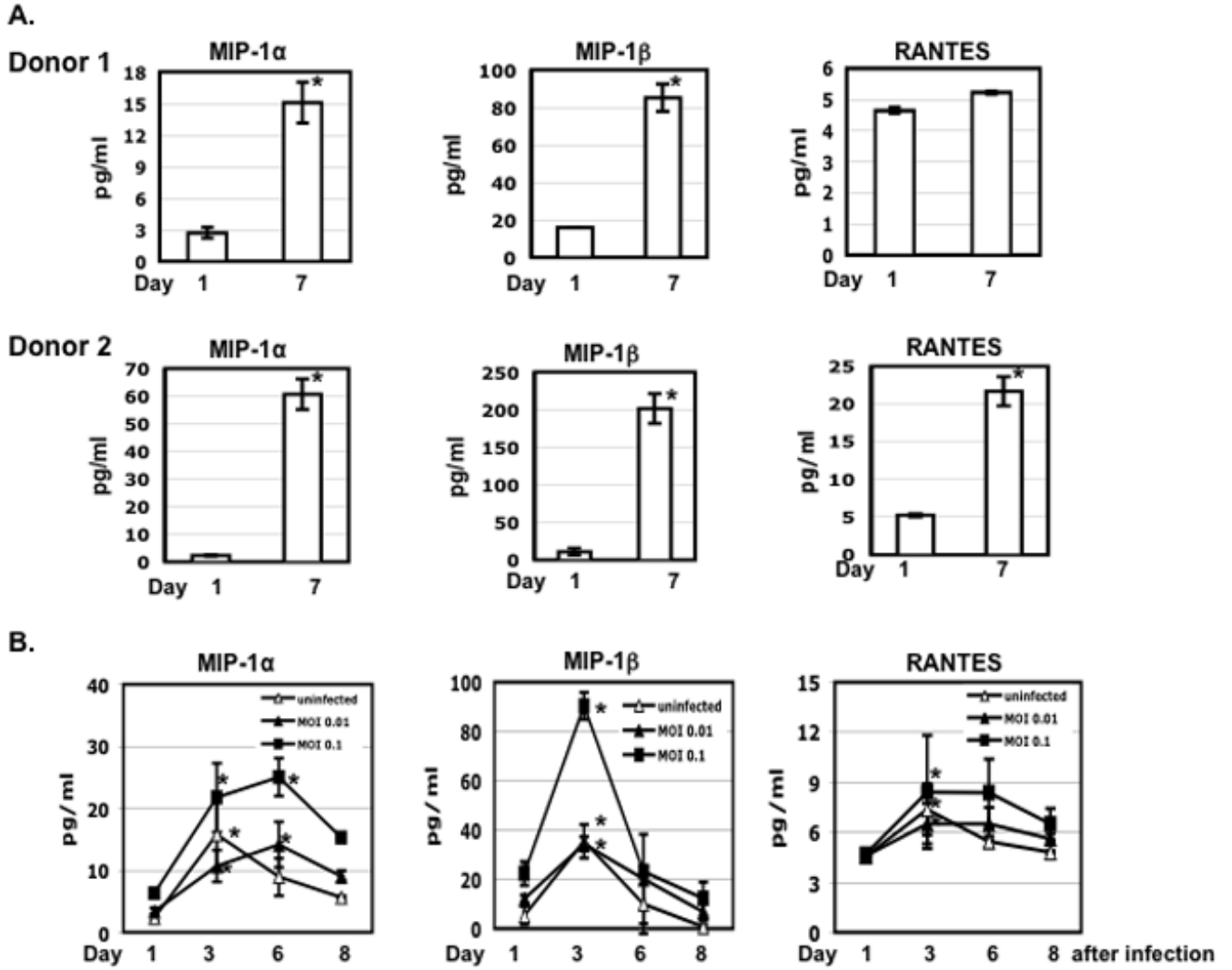


Figure 7. CC-chemokines are induced in PMs after culture

(A) AMC_s (1×10^6 per well) were plated in a 6-well plate, cultured for 16 h to allow adherence, and washed four times with PBS to remove non-adherent cells. Media from PMs after 1 or 7 days in culture were collected and the level of chemokines were determined by Luminex assays. (B) PMs were prepared by plating AMC_s (1×10^5 cells per well) in a 48-well plate for overnight following by washing. Cells were exposed to R5 HIV-1_{BaL} at an MOI 0.01 or 0.1 for 2 h, washed and cultured in RPMI with 10% FBS. Uninfected cells were included as a comparison. One-half volume of media was collected at the indicated time points to determine the levels of chemokines and HIV p24. Fresh media were added back at each time point. Significant differences between the level of chemokines at day 1 vs day 4 or 7 after culture (panel A) or infection (panel B) are noted ($*p < 0.05$). The results represented four independent experiments from different patients.

Table 1**Cell populations in AMCs**

Freshly isolated AMCs were subjected for cell surface staining for CD3, CD14, CD19. For CD68 staining, cells were fixed and permeabilized. Results of FACS analysis of AMCs from two donors with liver diseases (LD1 and LD2) and PBMCs from two healthy donors (HD1 and HD2) were shown.

	AMCs LD1	AMCs LD2	PBMCs HD1	PBMCs HD2
CD3	12%	10%	36.5%	36%
CD14	70%	74%	15%	4.23%
CD19	0.03%	0.07%	2.38%	4.52%
CD68	82%	85%	26%	12%