Impaired NK Cell Activation and Chemotaxis toward Dendritic Cells Exposed to Complement-Opsonized HIV-1

Rada Ellegård, Elisa Crisci, Jonas Andersson, Esaki M. Shankar, Sofia Nyström, Jorma Hinkula and Marie Larsson

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Impaired NK Cell Activation and Chemotaxis toward Mucosal Neighbor Cells Exposed to Complement-Opsonized HIV-1

Rada Ellegård,* Elisa Crisci,* Jonas Andersson,* Esaki M. Shankar, † Sofia Nyström,* Jorma Hinkula,* and Marie Larsson*

Mucosa resident dendritic cells (DCs) may represent one of the first immune cells that HIV-1 encounters during sexual transmission. The virions in body fluids can be opsonized with complement factors because of HIV-mediated triggering of the complement cascade, and this appears to influence numerous aspects of the immune defense targeting the virus. One key attribute of host defense is the ability to attract immune cells to the site of infection. In this study, we investigated whether the opsonization of HIV with complement (C-HIV) or a mixture of complement and Abs (CI-HIV) affected the cytokine and chemokine responses generated by DCs, as well as their ability to attract other immune cells. We found that the expression levels of CXCL8, CXCL10, CCL3, and CCL17 were lowered after exposure to either C-HIV or CI-HIV relative to free HIV (F-HIV). DCs exposed to F-HIV induced higher cell migration, consisting mainly of NK cells, compared with opsonized virus, and the chemotaxis of NK cells was dependent on CCL3 and CXCL10. NK cell exposure to supernatants derived from HIV-exposed DCs showed that F-HIV induced phenotypic activation (e.g., increased levels of TIM3, CD69, and CD25) and effector function (e.g., production of IFNγ and killing of target cells) in NK cells, whereas C-HIV and CI-HIV did not. The impaired migration of NK cells is consistent with C-HIV and CI-HIV having a negative impact on NK cell function. Furthermore, the chemokine profile of F-HIV-exposed DCs was different from that of C-HIV and CI-HIV, which suggests that the opsonization of virus with complement factors influences the recruitment of immune cells to the site of infection.

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The online version of this article contains supplemental material.

Abbreviations used in this article: AT2, aldrithiol-2; C-HIV, complement-opsonized HIV; CI-HIV, complement- and Ab-opsonized HIV; DC, dendritic cell; F-HIV, free HIV; MFI, multiple of infection; qPCR, quantitative PCR.

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Our results demonstrated that DC interaction with C-HIV impaired the recruitment of NK cells, as well as the NK cell activation, which may contribute to the failure of innate immune responses to control HIV at the site of initial mucosa infection.

Materials and Methods
Preparation and culturing of DCs
Monocyte-derived DCs were prepared and cultured as described previously (19). In brief, PBMCs were separated from whole blood from healthy volunteers (ethical permit EPN 173-07). DC progenitors were enriched by adhesion of PBMCs to plastic tissue culture plates. The cells were cultured in RPMI 1640 with l-glutamine supplemented with 10 mM HEPES, 20 μg/ml gentamycin (all from Fisher Scientific, Leicestershire, U.K.), 100 IU/ml recombinant human GM-CSF (Immunex, Seattle, WA), 300 U/ml recombinant human IL-4 (R&D Systems, Minneapolis, MN) and 1% human plasma for 5 d.

Virus generation and exposure assay
Infectious HIV-1MAD (lot nos. 4143, 4238, 4235), aldrithiol-2 (AT2)-inactivated HIV-1MAD (lot no. 4243) and infectious HIVMN (lot no. 4091) were produced from SUP-T1/CCR5 cells and purified by sucrose gradient banding, essentially as described previously (20). In addition, a primary isolate (lot no. 4293) was used. This virus was derived from the limiting dilution clone HIV-1 THRO/A66-R5 CL 29, cell line number 506 (A.K.A. [CLN506]). The clone was sequenced and found to be largely identical to the original sequence, and it was confirmed that it retained its R5 tropism.

The virus was incubated with either an equal volume of RPMI 1640 (Sigma-Aldrich, Stockholm, Sweden) to generate (F-HIV) single donor human serum supplemented with 25% veronal buffer, or human serum supplemented with 25% veronal buffer and 2 mg/ml p24, of F-HIV, C-HIV, or CI-HIV, or mock. After 6 to 24 h, the supernatants were harvested, and the cells were prepared for quantitative PCR (qPCR) analysis.

Cytometric bead array
Protein levels of CXCL8, CXCL10, and CCL3 in cell supernatants were assessed using a commercial Cytometric Bead Array (BD Biosciences, Stockholm, Sweden) performed on a BD FACSCanto II flow cytometer (BD Biosciences) and analyzed using FCAP Array version 3 software (BD Biosciences) according to the manufacturer’s protocols.

Immune cell migration assay
Cell-free supernatants collected from DCs 24 h after exposure to HIV or mock were placed in the bottom wells of a Transwell migration plate (Costar Corning, Lowell, MA). PBMCs were added to the upper wells, and the plates were incubated for 90 min to allow the cells to migrate. The migrated cells were collected, counted, and immunostained using the following Abs: VioBlue-CD19 (Miltenyi Biotec, Stockholm, Sweden) and FITC-CD56, PeCy5-CD3, PE-CD8, and APC-CD4 (BD, Stockholm, Sweden). The cells were run on a BD FACScanto II flow cytometer and analyzed using FlowJo Software (TreeStar, Ashland, OR). CD3+CD4+ cells were defined as CD4+ T cells, CD3+CD8+ as CD8+ T cells, CD56+ cells as NK cells, CD19+ cells as B cells, and CD14+ cells as monocytes. To examine the factors responsible for the chemotaxis of immune cells, the assay was performed as above with the exception that neutralizing Abs targeting CCL3, CXCL8, CXCL10 or an isotype control (R&D Systems) were added to the DC supernatants before allowing the PBMCs to migrate toward them.

NK cell purification and flow cytometry
Human NK cells were purified from PBMCs using negative selection (NK Cell Isolation Kit; Miltenyi Biotec, Stockholm, Sweden). The NK cells were incubated with supernatants derived from DCs exposed to mock, F-HIV, C-HIV, or CI-HIV for 24 h and subsequently stained with FITC-CD25, PeCy5-CD49, APC-TIM3, PerCP-HLADR, and Alexa Fluor 647-CCR7 mAbs (BD, Stockholm, Sweden). Samples were acquired on a BD FACScanto II flow cytometer and analyzed using FlowJo software.

Total RNA extraction, reverse transcription, and qPCR
Total RNA was extracted using Mini or Micro RNA purification kits from Bioline (London, U.K.). Reverse transcription was performed using the SuperScript III Reverse Transcriptase First Strand cDNA Synthesis Kit (Life Technologies, Stockholm, Sweden) according to the manufacturer’s protocols. qPCR was performed with SYBR Green master mix (Bioline) using a CFX96 Real-Time System (Bio-Rad, Solna, Sweden) according to manufacturer’s protocols. Primers targeting β-actin and GAPDH were used as housekeeping genes for reference as described by Vandesompele et al. (21). Primers targeting IL-2, IL-10, IL-15, CCL3, CCL13, CCL17, CCL18, CCL22, CXCL8, CXCL10, granzyme B, and perforin were used (Supplemental Table I). All primers were purchased from CyberGene AB (Stockholm, Sweden). Samples were run in triplicate and compared with both housekeeping and target genes on the same plate.

Killing assay
K562 cells (ATCC, Manassas, VA), a cell line negative for MHC class I, were cultured in RPMI 1640 with l-glutamine supplemented with 10 mM HEPES, 20 μg/ml gentamycin, and 10% FBS (all from Fisher Scientific, Leicestershire, U.K.) and used as target cells for NK killing. DCs were exposed to F-HIV, C-HIV, or mock for 24 h and the supernatants harvested as described above. Purified NK cells from the same donor were exposed to these supernatants for 24 h and then cocultured with K562 cells stained with CFSE (Molecular Probes, Life Technologies, Stockholm, Sweden) for 6 h at a 1:8 ratio. NK cells were treated with RPMI or 15 μg/ml PHA and 300 U/ml IL-2 as negative and positive controls, respectively. The frequency of CFSE-labeled K562 cells undergoing cell death was determined using a BD FACScanto II flow cytometer and analyzed using FlowJo software (TreeStar).

Statistical analysis
The ΔΔCt2 method, as described by Rieu and Powers (22), was used to normalize the variations between qPCR plates. Later, the samples were normalized to have a mock treated control of 1. All results were analyzed using GraphPad 5.0 software (GraphPad Software, La Jolla, CA) with repeated measures ANOVA followed by Bonferroni post test; p < 0.05 was considered statistically significant. n denotes the number of independent experiments performed with cells derived from a different healthy individual.

Results
Complement opsonization of HIV results in decreased production of HIV-induced chemokines by DCs
Complement opsonization of HIV has been shown by several studies, including our own, to increase virus uptake and infection of immature DCs in a CR3-dependent manner while suppressing inflammatory responses (9–12, 23). In this study, we extended these studies to examine the effects of complement opsonization on the ability of HIV to activate DC expression of chemotactic factors. The expression of an array of CC- and CXC-chemokine ligands associated with leukocyte migration—namely CCL3, CCL13, CCL17, CCL18, CCL22, CXCL8, and CXCL10—were examined by qPCR in DCs exposed to 8 MOI HIV-1MAD, either as F-HIV, C-HIV, or CI-HIV virions. CXCL10 expression peaked 6 h after exposure, while CCL3, CCL17, CCL22, and CXCL8 peaked 12 h after exposure (Fig. 1A). The levels of all these factors were significantly higher for F-HIV compared with C-HIV and CI-HIV (Fig. 1A). The HIV-induced upregulation of CXCL10, CCL3, and CXCL8 was considerable and visible for all virus groups, whereas the upregulation of CCL17 and CCL22 was less pronounced and visible only for F-HIV (Fig. 1A). Expression of CCL13 or CCL18 did not rise significantly above mock levels for any of the virus-exposed groups at any time point (Fig. 1A).

Cytometric Bead Array analysis showed that F-HIV induced significantly higher levels of CXCL10 compared with C-HIV and CI-HIV (Fig. 1B). In addition, F-HIV induced significantly higher expression of CCL3 compared with mock treated samples, while C-HIV and CI-HIV did not (Fig. 1B). Expression of CXCL8 was significantly upregulated for all virus-exposed groups (Fig. 1B).

When lower viral titers were used, expression of CCL3, CXCL8, and CXCL10 decreased, but the profiles remained the same, with F-HIV giving rise to a higher expression than C-HIV or CI-HIV (Supplemental Fig. 1A).
The expression of CCL3, CXCL8, and CXCL10 were also examined when AT2- HIVBaL, a chemically inactivated virus with the ability to bind and fuse with a target cell but not to replicate (24), the X4-virus HIVMN or a primary R5-tropic isolate were used instead of infectious HIVBaL. AT2-HIVBaL resulted in an almost identical profile as the infectious virus, indicating that infection of the DCs was not necessary to induce the cytokine expression (Fig. 1C). HIVMN also displayed a similar profile, with a lower cytokine expression after exposure to C-HIV compared with F-HIV, although the expression levels were somewhat lower and the difference was less pronounced compared with HIVBaL (Fig. 1C). The primary isolate gave rise to lower CCL3, but higher CXCL10 responses compared with HIVBaL, with F-HIV giving rise to higher expression compared with the complement-opsonized virus (Fig. 1C).

**DCs exposed to F-HIV had the ability to recruit immune cells, especially NK cells, whereas DCs exposed to complement-opsonized HIV did not.**

We next examined whether the differences in chemotactic ligand expression between DCs exposed to F-HIV and DCs exposed to opsonized HIV affected their ability to attract other immune cells. The ability to induce immune cell chemotaxis differed greatly between supernatants derived from immature DCs exposed to F-HIV, C-HIV, and CI-HIV (Fig. 2A). NK cells were the cell type most affected by HIV opsonization status, with a large number of cells migrating toward supernatants derived from DCs exposed to F-HIV, whereas no migration was induced by supernatants from cultures exposed to C-HIV and CI-HIV (Fig. 2A). CD4+ and CD8+ T cell migration was increased for F-HIV compared with groups treated with C-HIV and CI-HIV, although the levels of migrating CD8+ T cells were very low (Fig. 2A). No significant difference was found between free and opsonized virus in regard to the number of migrating monocytes and B cells (Fig. 2A).

To determine which chemokines were responsible for inducing migration, neutralizing Abs targeting CCL3, CXCL8, and CXCL10 or a matching isotype control were added to the supernatants prior to the migration assay. Blocking CCL3 or CXCL10 inhibited the migration of NK cells, whereas blocking CXCL8 did not (Fig. 2B, Supplemental Fig. 2). Migration of monocytes was decreased when CCL3 or CXCL8 was neutralized, whereas migration of T cells and B cells was not significantly affected by any of the neutralizing Abs (Supplemental Fig. 2).

**Supernatants from DCs exposed to F-HIV induced partial NK cell activation, whereas supernatants from DCs exposed to C-HIV did not.**

As the greatest effects on migration were seen for NK cells, we evaluated the effects of supernatants from F-HIV-, C-HIV-, and CI-HIV–exposed DCs on NK cell phenotype and activation status.
The levels of NK cell activation markers CD69, CD25, TIM3, HLA DR, and CCR7 were assessed by staining the NK cells after 24 h of exposure to the DC supernatants followed by flow cytometry analysis. The expression of activation markers CD69, CD25, TIM3, and HLA DR were significantly increased in NK cells exposed to supernatants derived from DCs treated with F-HIV, whereas C-HIV and CI-HIV had minimal effects on their expression levels (Fig. 3).

The gene expression levels of cytokines IFN-γ, IL-2, IL-15, and IL-10 in the NK cells were also examined 6 and 24 after exposure to the supernatants. We found that F-HIV significantly increased IFN-γ gene expression at both time points, whereas C-HIV and CI-HIV had no significant effects compared with supernatants from mock treated DCs (Fig. 4). This profile was also visible when the IFN-γ protein concentration in the NK cell supernatants 24 h after exposure was assessed with ELISA (Fig. 4A). Only relatively low IL-10 and IL-2, and no IL-15, gene upregulation was detected for NK cells exposed to F-HIV supernatants 6 h after exposure (Fig. 4B), and levels of all three factors were extremely low or borderline negative 24 h after exposure (data not shown).

**Complement opsonization of HIV resulted in decreased cytotoxicity in NK cells cultured in supernatants from HIV-exposed DCs**

A few chemokines, such as CCL3, have been shown to enhance the cytolytic activity of NK cells (25, 26); therefore, we assessed the NK cell cytotoxicity after exposure to supernatants from HIV-exposed DCs. The gene expression of granzyme B and perforin and the ability to kill target cells were examined. NK cell gene expression levels of granzyme B and perforin were increased for all HIV groups, but the highest levels were seen for F-HIV (Fig. 5A). When assessing the ability to kill the cell line K562, commonly used as target cells to study NK cell killing, we found that the NK cells exposed to supernatants from DCs exposed to F-HIV increased their killing ability, whereas C-HIV and CI-HIV supernatants induced the same levels of killing that mock treatment did (Fig. 5B).

**Discussion**

Complement opsonization of HIV increases viral internalization and infection of immature DCs (9–12, 23). In addition, we recently found that the inflammatory and antiviral responses induced by HIV in immature DCs are decreased when the virus is complement opsonized, resulting in enhanced infection (13). In this study, we show that the interaction between immature DCs and CI-HIV results in decreased production of chemokines and the ability to attract other immune cells, particularly NK cells. In addition, the activation and cytotoxic ability of the NK cells was reduced when they were cultured in supernatants from DCs exposed to CI-HIV compared with supernatants from DCs exposed to F-HIV. Seeing that HIV can be complement opsonized in vivo (8), this could adversely influence the body’s ability to fight the viral infection. NK cells are important because they kill virus-infected cells and enhance and modulate DC-induced immunity (15, 27), and the reduced ability of immature DCs exposed to CI-HIV to attract and enhance other immune cells, particularly NK cells.
activate these cells could favor viral establishment and persistence.

Inflammatory and chemotactic chemokines are normally produced during infection, and they determine the migration of immune cells to the site and exert direct antiviral properties (28, 29). The exposure of DCs to F-HIV activated the production of multiple chemotactic factors, whereas opsonization decreased these signals, affecting migration of CD4+ and CD8+ T cells, and especially NK cells. This can be explained in part by the decreased production of CCL3, CCL17, CXCL8, and CXCL10 produced by the DCs exposed to complement-opsonized HIV, cytokines that have chemotactic properties for both T cells and NK cells (25, 28, 30–32). A potential lack or suboptimal levels of these factors could be beneficial for HIV, because these chemokines can have other anti-HIV roles in addition to attracting cells with the ability to kill virus-infected cells (29, 33).

Most NK cells have the ability to migrate in response to several chemokines, such as CCL2, CCL3, CCL4, CCL5, CCL8, CXCL9, CXCL10, and CXCL11 (25, 34–36), because of the expression of their cognate receptors (e.g., CXCR1, CXCR2, CXCR3, CXCR4, CX3CR1) on these cells (34, 37). In our system, DCs exposed to F-HIV significantly upregulated gene expression of CCL3 and CXCL10 compared to mock, whereas the CXCL8 levels were reduced (25, 34, 36). This chemokine is chemotactic for NK cells (31), it may help to recruit cells with the ability to kill virus-infected cells (41). CXCL10 attracts activated T cells and NK cells and stimulates monocytes (32). The impaired ability of DCs to respond to HIV exposure by upregulating CXCL10 when the virus was complement-opsonized led to decreased NK cell migration, and this could aid viral establishment in the host.

All virus-exposed DCs secreted comparable amounts of CXCL8, whereas F-HIV induced a significantly higher expression of CCL3 and CXCL10 than complement-opsonized virus did. The NK cells in our system were able to migrate in response to the CCL3 and CXCL10 produced by DCs exposed to F-HIV, whereas the lower cytokine levels induced by complement-opsonized virus failed to induce a higher NK cell migration than the mock treated samples. Previous findings are inconsistent concerning the ability of CXCL8 to induce NK cell migration (25, 34). However, in our system, this

FIGURE 4. Complement opsonization of HIV-impaired IFN-γ expression in NK cells exposed to supernatants derived from HIV-exposed DCs. Supernatants were collected from DCs exposed to F-HIV, C-HIV, CI-HIV, or mock for 24 h, after which the NK cells were evaluated for IFN-γ gene expression using qPCR, and the release of IFN-γ protein was measured using ELISA. ELISA results are shown as one representative experiment and as all experiments normalized to F-HIV (n = 5). (A) NK cells were exposed to supernatants collected from HIV-exposed DCs for 6 h and gene expression of IFN-γ, IL-2, IL-10, and IL-15 were determined using qPCR (n = 3). Results were tested for statistical significance using repeated measures ANOVA followed by Bonferroni post test. *p < 0.05.

FIGURE 5. Complement opsonization of HIV impaired the killing ability of NK cells exposed to supernatants from HIV-exposed DCs. Purified NK cells were cultured in supernatants collected from DCs exposed to F-HIV, C-HIV, CI-HIV, or mock for 24 h. (A) The NK cells were cultured for Granzyme B and Perforin gene expression using qPCR (n = 4). (B) The NK cells were exposed to supernatants from HIV-exposed DCs for 6 h and gene expression of IFN-γ, IL-2, IL-10, and IL-15 were determined using qPCR (n = 3). Results were tested for statistical significance using repeated measures ANOVA followed by Bonferroni post test. *p < 0.05 (n = 4).
citokine did not appear to affect NK cell chemotaxis. The levels of B cell and monocyte migration were less affected by opsonization of the virions compared with NK cells and T cells. Chemokines such as CCL3 and CXCL10 have been reported to be involved in monocyte migration during HIV transmission (42, 43), and we found these factors to be involved in the monocyte migration in our system as well.

In addition to the number of cells migrating to the site of infection, NK-mediated protection is also dependent on the activation status and functionality of the recruited cells. Exposure of NK cells to factors produced by DCs at sites of inflammation and infection can lead to their activation with changes in phenotype and function (44–46) and affect NK cell–mediated destruction of infected cells (47). At the site of genital infection, NK cell migration and activity are an important defense against infections such as HSV2 (14) and NK cell cytotoxicity, and the ability to secrete antiviral cytokines has been correlated with protection in HIV-exposed noninfected individuals (17). NK cells may help to limit the initial HIV infection (41), whereas viremia leads to decreased numbers and impaired functionality of NK cells (48). Importantly, preservation of NK activity has been correlated with better disease outcome (49, 50). The activation status of the NK cells can affect interactions between NK cells and DCs and affect DC maturation with potential to modulate the adaptive immune response (51, 52), and NK cell–DC cross-talk in the periphery is crucial for the formation of optimal Th1 adaptive immunity (53).

We found that supernatants from DCs exposed to F-HIV induced a partial activation of NK cells with increased expression of TIM3, CD25, CD69, HLA DR, and CCR7, whereas supernatants from DCs exposed to complement-opsonized HIV failed to do so. In addition, the ability of the DCs to induce IFN-γ production in the NK cells was suppressed when HIV was complement opsonized. The importance of NK cells is evident in many viral infections in which they help to clear the infection by killing infected cells and when impaired (i.e., low levels or dysfunctional NK cells), the host fails to clear the infection (41). Likewise, in vivo depletion of NK cells in mice infected with *Citrobacter rodentium* gave rise to a higher bacterial load and disseminated systemic infection (54, 55). NK cells control viral infections by killing virus-infected cells, an activity impacted by E:T ratio (56). The lower number of NK cells mobilized by complement-opsonized HIV should limit NK-mediated–killing of infected cells at the site of infection. In addition, exposure to supernatant from DCs exposed to F-HIV enhanced the gene expression of granzyme B and perforin and the killing of target cells, whereas exposure to supernatants from DCs exposed to complement-opsonized virus did not. CCL3, CCL5, and CXCL10 are reported to be effective at inducing the cytolytic activity of NK cells (25, 26); therefore, the NK cell exposure to CXCL10 and CCL3 produced by the DCs exposed to F-HIV should be responsible for their enhanced killing ability.

We have previously shown that C-HIV inhibits induction of inflammatory cytokines in DCs via CR3 interaction (13), and we show that this mechanism also affected the release of chemotactic chemokines and thereby the attraction of immune cells. In a recent SIV study, it was shown that a surprisingly low number of NK cells migrated to the site of infection after vaginal challenge, and the NK cells that did migrate lacked markers associated with activation and cytotoxicity (57). One explanation behind this could be the ability of HIV to exploit complement opsonization as a means to inhibit the secretion of chemokines and other factors by DCs that recruit and activate NK cells. The evasion of NK cell cytotoxicity and secretion of antiviral cytokines, such as IFN-γ, is likely beneficial for HIV and aids in the establishment of infection. This mechanism may also be important to consider when designing mucosal HIV vaccines, as the addition of an adjuvant that overrides the suppression of NK mobilization achieved by complement could prove beneficial for optimal host immune responses.

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**Disclosures**

The authors have no financial conflicts of interest.

**References**


### Supplementary Table 1. Primer Sequences.

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### Supplementary Figure 1. RNA expression of CCL3, CXCL8 and CXCL10 using different concentrations of HIV. DCs were exposed to MOI 0.08-8 HIV for 6h and the mRNA expression of CCL3, CXCL8 and CXCL10 were determined using qPCR.
Supplementary Figure 2. Role of CCL3, CXCL8 and CXCL10 in the migration of PBMCs towards dendritic cells exposed to free and complement opsonized HIV. PBMCs were allowed to migrate for 90min towards supernatants collected from DCs exposed to free HIV (F-HIV), complement opsonized HIV (C-HIV), complement and antibody opsonized HIV (CI-HIV) or mock for 24h. To determine which
cytokines were responsible for the migration of NK cells towards supernatants from DCs exposed to F-HIV for 24h, the supernatants were pre-treated with neutralizing antibodies targeting CCL3, CXCL8, CXCL10 or an isotype control prior to the migration assay. The type of cells migrating, ie (A) NK cells (defined as CD56+), (B) monocytes (CD14+), (C) B cells (CD19+), (D) CD4 T cells (CD3+CD4+) and (E) CD8 T cells (CD3+CD8+) were determined by flow cytometry. Results were tested for statistical significance using repeated measures ANOVA followed by Bonferroni post test. *p<0.05, compared to isotype control. (n=4).