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# Fetal exposure to HIV-1 alters chemokine receptor expression by CD4+T cells and increases susceptibility to HIV-1

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Absolute numbers of lymphocytes are decreased in uninfected infants born to HIV-1-infected women (HIV-1-exposed). Although the exact mechanism is unknown, fetal exposure to maternal HIV-1-infection could prime the immune system and affect T cell trafficking. We compared the expression of chemokine receptors on cord blood CD4<sup>+</sup> T cells from HIV-1-exposed children and healthy controls. At baseline CD4<sup>+</sup> T cells had a largely naïve phenotype. However, stimulation with cytokines resulted in an upregulation of inflammatory response-related chemokine receptors on CD4<sup>+</sup> T cells, with HIV-1-exposed infants having a significantly higher frequency of CD4<sup>+</sup> T cells expressing, in particularly T<sub>h</sub>2 associated chemokine receptors (CCR3 p < 0.01, CCR8 p = 0.03). Numbers of naive CCR7<sup>+</sup> CD4<sup>+</sup> T cells were reduced (p = 0.01) in HIV-1-exposed infants. We further assessed whether the inflammatory phenotype was associated with susceptibility to HIV-1 and detected higher levels of p24 upon in in vitro infection of stimulated CD4<sup>+</sup> T cells of HIV-1-exposed infants. In summary, fetal exposure to HIV-1 primes the immune system in the infant leading to an enhanced immune activation and altered T cell homing, with potential ramifications regarding T cell responses and the acquisition of HIV-1 as an infant.

nnually, more than 1.5 million HIV-1-infected women give birth to 330,000 HIV-1-infected children<sup>1</sup>. Long-term combination antiretroviral therapy (cART) during pregnancy, reduces mother-to-child transmission of HIV-1 from 25% to  $<1\%^2$ . Since 2010, long-term cART during pregnancy is therefore recommended for all HIV-1-infected women, also in resource-poor settings, reducing the number of children infected with HIV-1 at birth<sup>3,4</sup>. However, this also implies that the number of uninfected children born to HIV-1-infected women is rapidly increasing<sup>2,3</sup>.

Although these children are uninfected and considered to be healthy, they are exposed during *in utero* development to maternal HIV-1 and cART. There is accumulating data suggesting that exposure to maternal HIV-1-infection and cART *in utero* has an effect on their immune system and susceptibility to infections<sup>5,6</sup>. Children born to HIV-1-infected mothers with a CD4<sup>+</sup> T cell count less than 200 cells/ $\mu$ L, have reduced lymphocyte counts compared to infants born to mothers with a CD4<sup>+</sup> T cell count above 200 cells/ $\mu$ L<sup>7-9</sup>. Furthermore, exposure to cART *in utero* is associated with reduced numbers of lymphocytes and neutrophils up to at least 8 years of age<sup>7,10,11</sup>. These observations suggest that key components of the immune system of HIV-1-exposed children are substantially altered with clinical consequences; indeed, an increased susceptibility to common infectious diseases<sup>5,12</sup>. Alterations in the infant's immune system could also affect postnatal transmission of HIV-1, as many of these children are breastfed with continuous exposure to HIV-1 after birth up to at least 6 months of age and potentially as an adult.

Toxicity of cART on blood stem cells could affect numbers of circulating  $CD4^+$  T cells; however, this does not explain the correlation between infant and maternal  $CD4^+$  T cell counts<sup>8,9</sup>. Due to *in utero* exposure to HIV-1, altered migratory patterns of  $CD4^+$  T cells from blood into tissues could also influence  $CD4^+$  T cell counts in the circulation of HIV-1-exposed infants. Homing to specific anatomical sites depends on the expression of specific chemokine receptors on  $CD4^+$  T cells<sup>13</sup>. Expression of specific chemokine receptors allows functional characterization of  $CD4^+$  T cells<sup>14-16</sup>. For example, CCR7 is specifically expressed by naive  $CD4^+$  T cells to direct these cells to lymph nodes, where their cognate antigen may be presented<sup>14-16</sup>. Other chemokine receptors are associated





Figure 1 | Chemokine receptor expression on cord blood CD4<sup>+</sup> T cells from HIV-1-infants and healthy controls assessed with flow cytometry. CD4<sup>+</sup> T cells of a healthy control (A) and an HIV-1-exposed infant (B) at birth were analysed with flow cytometry. Expression of CCR7 was high in both groups, whereas inflammatory response-related chemokine receptors CCR5 and CCR8 were nearly absent. (C) High expression of CXCR4 and CCR7 was detected in both groups and did not differ significantly. (D) T<sub>h</sub>1-inflammatory response-related chemokine receptors CCR5 and CCR8 were nearly absent on CD4<sup>+</sup> T cells in both HIV-1-exposed infants and healthy controls. (E) T<sub>h</sub>2- inflammatory response-related chemokine receptors CCR1 and CCR8 were marginally expressed on CD4<sup>+</sup> T cells in both HIV-1-exposed infants and healthy controls, but were significantly higher in HIV-1-exposed infants. CCR6, CCR9 and CCR10 were not depicted as these were expressed at extremely low percentages of CD4<sup>+</sup> T cells in both groups (<0.4%). N = 12 (healthy control) and 10 (HIV-1-exposed infants). Median is depicted with interquartile range (IQR). Statistical analyses with Man-Whitney U-test.

with specific immune responses upon infection, such as CXCR6, in the case of  $T_h1$  responses, and direct  $CD4^+$  T cells towards the infected tissues<sup>14-16</sup>. Furthermore, increased expression of certain chemokine receptors, especially those upregulated during inflammatory responses, also play a role in the pathogenesis of inflammatory diseases<sup>17,18</sup>. The evaluation of chemokine receptor expression on  $CD4^+$  T cells can, therefore, provide insight into the homing pattern as well as the immune response of  $CD4^+$  T cell in an individual and help elucidate the observed differences in  $CD4^+$  T cell count in HIV-1-exposed infants.

In the study reported here, we investigated the expression of chemokine receptors on CD4<sup>+</sup> T cells from cord blood from infants born to HIV-1-infected women and healthy women. Although small, we detected a higher expression of chemokine receptors, normally upregulated under inflammatory conditions already at birth. However, after *in vitro* stimulation with a broad variety of cytokines and stimuli, including IL-1 $\beta$ , chemokine receptors were greatly upregulated and the differences between HIV-1-exposed infants and controls became strikingly evident. Furthermore, the enhanced state of activation of CD4<sup>+</sup> T cells of infants born to HIV-1-infected women was associated with an increased susceptibility to HIV-1 infection compared to healthy controls. Combined, these data not only show an upregulation of chemokine receptors on CD4<sup>+</sup> T cells upon *in utero* 

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HIV-1 exposure, which allows egress from blood in tissues, but also an overall activated state, which affects postnatal susceptibility to HIV-1 and potentially other infections.

#### Results

CD4<sup>+</sup> T cells have a largely naïve phenotype in HIV-1-exposed infants and healthy controls at birth. CD4+ T cells derived from cord blood were analyzed by flow cytometry to determine baseline chemokine receptor expression profiles at birth. CD4<sup>+</sup> T cells from both healthy controls and uninfected HIV-1-exposed infants had a largely naive, undifferentiated phenotype (CD45RO<sup>-</sup>CD27<sup>+</sup>) (data not shown) with high expression of CCR7 and CXCR4 (Figures 1A-C). In accordance, only few CD4<sup>+</sup> T cells expressed chemokine receptors associated with inflammatory responses, such as CCR2, CCR5 and CXCR6 that are predominantly expressed by T<sub>h</sub>1 cells (Figure 1D), and CCR1, CCR3 and CCR8, which are expressed by  $T_h2$  cells as shown in (Figure 1E)<sup>14–16,19</sup>. Although percentages were very small, significantly more CD4<sup>+</sup> T cells from HIV-1-exposed infants expressed chemokine receptors associated with inflammatory responses than from the healthy controls (Figures 1D, E). These data suggest that CD4<sup>+</sup> T cells from both healthy and HIV-1exposed infants have a largely naïve phenotype, however small percentages of CD4 $^+$  T cells of HIV-1-exposed infants maybe activated in utero.

Chemokine receptors upregulated upon infection or inflammation are upregulated upon stimulation of cord blood CD4<sup>+</sup> T cells in vitro. Adult naïve CD4<sup>+</sup> T cells cultured with cytokines and dendritic cells increase expression of inflammatory response-related chemokine receptors<sup>20</sup>. We explored whether upon stimulation with cytokines chemokine receptors could be upregulated on naïve cord blood CD4<sup>+</sup> T cells from healthy infants and HIV-1-exposed infants. The cord blood mononuclear cell fractions of healthy controls and HIV-1-exposed infants were labelled with CFSE to assess proliferation and cultured for 6 days with one of the following stimuli: IL-2, IL-4, IL-7, IL-15, IL1-β, GM-CSF, or a combination of monoclonal antibodies against CD3 and CD28 to simulate TCR stimulated activatin. Chemokine receptor expression on and CFSE dilution in CD4<sup>+</sup> T cells were measured by flow cytometry. In vitro proliferation of cord blood CD4<sup>+</sup> T cells varied among the different cultures; in culture with IL-15 10% (IQR 9-35%) of the CD4<sup>+</sup> T cells had proliferated, whereas 73% (IQR 59-82%) when cultured with IL-4 and almost all CD4<sup>+</sup> T cells (>99%) had undergone multiple cycles of proliferation, when cultured in the presence of monoclonal antibodies against CD3 and CD28 (Figure 2).

Chemokine receptors related to inflammatory responses such as CCR3, CCR5, CCR8 and CXCR6 were upregulated on CD4<sup>+</sup> T cells in cultures with in particular IL1- $\beta$  and GM-CSF (Figure 2). CCR7, expressed by almost all CD4<sup>+</sup> T cells before culture (>98% (median)), was downregulated on proliferating cells upon culture. To distinguish patterns in the regulation of chemokine receptors by the stimuli, we performed a hierarchical clustering between the expression of the chemokine receptors on the CD4<sup>+</sup> T cells and the different stimuli (Figure 3). Two groups were identified: *1*) chemokine receptors mainly expressed by naïve cells such as CCR7 and CXCR4, which were downregulated in cultures with the stimuli; and *2*) chemokine receptors associated with inflammatory responses, i.e. CCR5, CXCR6 and CCR8 which were upregulated upon culture.

Specific stimuli clustered together based on the pattern of expression of specific chemokine receptors on CD4<sup>+</sup> T cells in these cultures. For example, IL-2 and the other  $\gamma c$  chain receptor (CD132)restricted cytokines, IL-4, IL-7 and IL-15 induced similar expression profiles of chemokine receptors. Cultures with IL1- $\beta$  and GM-CSF also clustered together, inducing the highest expression of inflammatory response-related chemokine receptors on CD4<sup>+</sup> T cells.

Thus, stimulation of cord blood CD4<sup>+</sup> T cells with specific stimuli is associated with specific chemokine expression patterns. Moreover, inflammatory response-related chemokine receptors can be upregulated on cord blood CD4<sup>+</sup> T cells.

Increased frequencies of CD4<sup>+</sup> T cells expressing inflammatory response-related chemokine receptors in HIV-1-exposed children compared to healthy controls upon stimulation. In the above analyses we showed that chemokine receptors can be upregulated on cord blood derived CD4<sup>+</sup> T cells and specific patterns are induced by specific stimuli. Next, we compared chemokine receptor regulation on CD4<sup>+</sup> T cells from uninfected HIV-1-exposed infants to those from healthy controls. An ANOVA analysis per chemokine receptor, including all the cultures, showed that upon stimulation in vitro a significantly larger percentage of CD4<sup>+</sup> T cells of HIV-1exposed infants expressed inflammatory response-related chemokine receptors than  $CD4^+$  T cells of control infants; CCR1 (p = 0.04), CCR3 (p < 0.01), CCR4 (p < 0.01), CCR8 (p = 0.03) and CXCR6 (p < 0.01) (Fig. 4A, C). At the same time CCR7 expression was significantly lower on CD4<sup>+</sup> T cells from HIV-1-exposed infants compared to healthy controls after culture (p = 0.01) (Figures 4A– B). The expression of CCR5 did not significantly differ between the groups (p = 0.3). The proliferation capacity as measured by CFSE dilution in CD4<sup>+</sup> T cells from HIV-1-exposed infants was similar to

the controls. These data strongly indicate that upon activation with cytokines CD4<sup>+</sup> T cells of HIV-1-exposed infants behave differently compared to controls; the CD4<sup>+</sup> T cells have an enhanced state of activation as reflected by the increased upregulation of inflammatory response-related chemokine receptors and decreased expression of chemokine receptors expressed by naive CD4<sup>+</sup> T cells<sup>14-16,19</sup>.

Levels of IL-1ß and IL-8 are increased in plasma from HIV-1exposed infants. T cell responses are shaped by signals from innate cells, including cytokines released by macrophages or dendritic cells. We investigated whether levels of soluble factors, such as cytokines in plasma of HIV-1-exposed infants were also indicative of an enhanced immune activation, using a Luminex assay assessing a panel of soluble factors in plasma at birth. Out of the 17 cytokines and chemokines we assessed, most were undetectable in plasma, including IFN- $\alpha$  and IFN- $\gamma$ . However, the cytokines associated with inflammation, IL-1 $\beta$  and IL-8, were detectable and significantly higher in HIV-1-exposed infants than in healthy controls (Figure 5). IP10 (CXCL10) and IL-10 were also detected, however these did not differ between the two groups. These data demonstrate that inflammatory cytokines IL-1ß and IL-8 are increased in plasma of HIV-1-exposed children at birth, supporting the model of an enhanced immune activation in these infants.

Increased susceptibility to HIV-1-infection of CD4<sup>+</sup> T cells from HIV-1-exposed infants compared to controls. Immune activation is critical for HIV-1 infection as highly activated CD4<sup>+</sup> T cells are more susceptible to HIV-1 and supportive of HIV-1 replication, than resting CD4<sup>+</sup> T cells. We therefore explored whether CD4<sup>+</sup> T cells in HIV-1-exposed infants, due to their enhanced activation, would be more permissive to HIV-1 infection compared to CD4<sup>+</sup> T cells from controls. Cord blood mononuclear cells were inoculated directly after isolation with YU2 (R5-strain) for 12 hours or after stimulation for 4 days with IL-2 100 IU/ml or GM-CSF. As described earlier non-stimulated CD4<sup>+</sup> T cells from HIV-1-as well as healthy controls could not be infected with HIV-1 (data not shown)<sup>21</sup>. However, after immune activation by IL-2 or GM-CSF, infection was detected (Figure 6). Strikingly, levels of p24 were significantly higher in cell cultures from HIV-1-exposed infants compared to healthy controls. Together, these data demonstrate that the enhanced immune activation in HIV-1-exposed infants leads to increased HIV-1 susceptibility of CD4<sup>+</sup> T cells.

#### Discussion

Although uninfected HIV-1-exposed children are considered to be healthy, there is accumulating data suggesting that exposure to maternal HIV-1 infection and cART *in utero* has an effect on their immune system<sup>7,10,11</sup>. In this study, we demonstrated that chemokine receptors associated with inflammatory responses were expressed on a significantly larger percentage of CD4<sup>+</sup> T cells from HIV-1exposed infants compared to controls at birth, in particular after culture with stimuli released upon inflammation *in vivo*. At the same time chemokine receptors indicating mainly naïve cells such as CCR7 were expressed by fewer CD4<sup>+</sup> T cells in HIV-1-exposed infants.

These data strongly indicate that, already at baseline but particularly upon stimulation, a larger percentage of  $CD4^+$  T cells of HIV-1exposed infants express chemokine receptors associated with an inflammatory response. *In vivo* a similar situation is expected during common viral or bacterial infections. Consequently, a larger number of  $CD4^+$  T cells egress from blood into immune active sites, such as the intestinal tract, resulting in lower lymphocyte counts in blood and providing a mechanistic explanation for the epidemiological observations in HIV-1-exposed children.





Figure 2 | Chemokine receptor expression on CD4<sup>+</sup> T cells after *in vitro* culture analysed with flow cytometry. Mononuclear cell fractions derived from cord blood of either healthy controls or HIV-1-exposed infants were labelled with CFSE and cultured for 6 days with various stimuli: IL-2, IL-4, IL-7, IL-15, IL-1 $\beta$  GM-CSF and T cell receptor (TCR)-mediated activation with monoclonal antibodies against CD3 and CD28. Expression of chemokine receptors and CFSE was measured by flow cytometry. CD4<sup>+</sup> T cells are shown of a HIV-1-exposed infant after stimulation. Stimulation with monoclonal antibodies against CD3 and CD28 induced proliferation in almost all CD4<sup>+</sup> T cells (>99%), measured by dilution of CFSE. Culture with IL-1 $\beta$  or GM-CSF induced the highest upregulation of CCR5 and CCR8. Proliferating cells in all cultures except in cultures with antibodies against CD3 and CD28 downregulated CCR7 expression.





Figure 3 | Hierarchical clustering of the expression of the chemokine receptors on CD4<sup>+</sup> T cells at baseline and after stimulation. The heatmap shows the hierarchical clustering of the expression levels of the chemokine receptors on CD4<sup>+</sup> T cells at baseline and after culture with various stimuli, measured by flow cytometry. The *-e*- after chemoreceptor indicates HIV-1-exposed infants and the *-c*- healthy controls. CCR7, CXCR4 grouped together, as well as chemokine receptor-related inflammatory responses, i.e. CCR5, CCR8 and CXCR6. The expression of inflammatory response-related chemokine receptors such as CCR5, CCR8 and CXCR6 increased in *in vitro* cultures in particular with IL-1 $\beta$  and GM-CSF, clustering these two cytokines together. N = 7 in both groups.

The increased expression of inflammatory response-related chemokine receptors in HIV-1-infants and inflammatory cytokine levels suggest that the immune activation is enhanced in these infants. Previous data also indicated towards this, with more effector memory CD4<sup>+</sup>CD27<sup>-</sup> T cells and CD4<sup>+</sup> CD38<sup>bright</sup> HLA-DR<sup>bright</sup> cells at birth in HIV-1-exposed infants compared to healthy controls, even though the percentages were small in both groups  $(<1\%)^{9,22,23}$ . The mechanism behind the enhanced immune activation in HIV-1-exposed children is unknown. In HIV-1infected patients, enhanced immune activation is an important characteristic of infection<sup>24,25</sup>. In utero recognition of viral components, which have crossed the placenta or maternal immune inflammatory stimuli could induce infant innate cells, e.g. dendritic cells or macrophages, to produce inflammatory cytokines including IL-1 $\beta$ , which we detected in a higher concentration in plasma of HIV-1-exposed infants compared to healthy infants<sup>26</sup>. These acute phase cytokines could shape the innate and adaptive immune responses towards an inflammatory profile in HIV-1-infants and priming T cells. Furthermore, these data also explain the correlation between  $CD4^+$  T cell counts in the mother and the child, as mothers with decreased counts will have a higher immune activation and viral load, which activates the infant's immune system and affects chemokine receptor expression. As these observations are preserved past the neonatal period, viral exposure *in utero* may lead to epigenetic changes regulating inflammatory responses in infants over longer times and sustaining altered inflammatory responses beyond the duration of viral and immune activation exposure *in utero*<sup>27-29</sup>.

In particular, T<sub>h</sub>2-related chemokine receptors, CCR1, CCR3 and CCR8 were increased on CD4<sup>+</sup> T cells in HIV-1-exposed infants, and not T<sub>h</sub>1-related chemokine receptors, CCR5 and CXCR6. This pattern closely resembles the altered T<sub>h</sub>1/T<sub>h</sub>2 ratio in HIV-1-infected patients, which is also observed in HIV-1-infected pregnant women<sup>30–32</sup>. A preferential T<sub>h</sub>2 response concurs with the higher IgG antibody titers observed in uninfected HIV-1-infants compared to controls<sup>33,34</sup>. However, this may have implications when children are infected with intracellular pathogens, such as viruses when T<sub>h</sub>1 responses are required.





Figure 4 | ANOVA analyses of chemokine receptor expression of CD4<sup>+</sup> T cells from HIV-1-exposed infants and healthy controls. (A) CCR7 was significantly lower on cultured CD4<sup>+</sup> T cells from HIV-1-exposed infants compared to healthy controls in an ANOVA analysis (p = 0.01). (B) CCR5 expression on CD4<sup>+</sup> T cells of HIV-1-exposed infants and healthy controls after a 6 day culture with various stimuli did not differ significantly across all cultures. (C) CCR8 expression on CD4<sup>+</sup> T cells of HIV-1-exposed infants and healthy controls after a 6 day culture with various stimuli was significantly higher across all cultures (p = 0.03). N = 7 in both groups. Medians are depicted with interquartile range.

Furthermore, the enhanced immune activation may have direct clinical implications for these children in the context of mother-tochild transmission of HIV-1 as we also detected an increased susceptibility to HIV-1 infection in uninfected HIV-1-exposed infants. The majority of the children born to HIV-1-infected mothers are breastfed for at least 6 months and consequently are continuously exposed and at risk for viral transmission. Recently, we have shown that already at birth activated memory CD4<sup>+</sup> CCR5<sup>+</sup> T cells, exemplary target cells for HIV-1, are almost exclusively present in the gut. Due to the enhanced immune activation, children exposed to HIV-1 in utero are expected to have greater percentages of these target cells in the gut facilitating HIV-1 transmission during delivery or breastfeeding<sup>35</sup>. Interestingly, although culture with IL1- $\beta$  leads to an increase of CCR5 this did not lead to increase of p24 production, in contrast to IL-2. It is known that IL-2 at several stages of viral replication positively impacts virus production, 1. by increasing nucleotide pools for reverse transcription and 2. increasing NF-KB induced transcription<sup>36-39</sup>. Taken together CCR5 expression is important for infection, but additional factors are needed for successful viral replication.

Our studies were performed on a relatively small number of infants; the mothers had only very mild HIV-1 disease and low viral load, which would limit inflammatory responses in the infant. Nonetheless, our results already show an evident difference in the response of the CD4<sup>+</sup> T cell compartment of these HIV-1-exposed infants upon stimulation. Although we have not investigated this, we conjecture that in regions with a higher microbial load and more advanced HIV-1 disease these differences would probably be more explicit and have clinical implications. Studies investigating uninfected children born to HIV-1 infected women detected more severe sequelae of infections with intracellular pathogens such as CMV in infants<sup>6,12,40</sup>. This suggests that similar to common viral infections, fetal HIV-1 exposure may have clinical consequences<sup>41</sup>.

In conclusion, fetal exposure to maternal HIV-1 infection results in an enhanced immune activation with increased expression of inflammatory response-related chemokine receptors in HIV-1exposed infants. This affects the homing of CD4<sup>+</sup> T cells in tissues and provides a mechanistic explanation for the low blood lymphocyte counts at birth and thereafter in these children. The consequences of this finding reach beyond our research question and are



Figure 5 | Levels of soluble factors in plasma of HIV-1-exposed infants and healthy controls. Cytokines, chemokines and growth factors (IL1- $\beta$  IL-4, IL-5, IL-6, IL-8, IL-10, IL-13, IL-17A, IL-18, IFN- $\alpha$ , IFN- $\gamma$ , IP-10, TNF- $\alpha$ , MIP-1 $\alpha$ , MIP-1 $\beta$ , RANTES and GM-CSF) were measured in plasma at birth from 20 healthy controls and 20 HIV-1-exposed infants by LUMINEX analyses. Cytokines associated with acute phase inflammatory responses, IL-1 $\beta$  and IL-8 were significantly higher in HIV-1-exposed infants. Cytokines related to antiviral or regulatory responses were low and not significantly different in the plasma of both healthy controls and HIV-1-exposed children. Medians are depicted with interquartile range. Statistical analyses performed with Man-Whitney U-test.

expected to have clinical implications, regarding the immune responses generated in HIV-1-exposed infants upon common infections and susceptibility to HIV-1 later in life as well other pathogens.

#### **Methods**

**Subjects.** This study was carried out in the Emma Children's Hospital at the Academic Medical Centre (AMC), Amsterdam, the Netherlands. Prospectively, cord blood samples from uninfected children born to HIV-1-infected women (N = 10) and from children born to healthy mothers (N = 12) were obtained, for the chemokine receptor expression at birth. The study was approved by our local ethics committee and parental consent provided.

cART was initiated in the HIV-1-infected mothers, from 20 weeks gestation and consisted of zidovudine, lamivudine and nelfinavir or lopinavir boosted with ritonavir. All mothers acquired an undetectable viral load before delivery and had a CD4<sup>+</sup> T cell count > 200 cells/µL. The newborn infants were treated for 4 weeks with zidovudine and lamivudine orally, according to national guidelines. HIV-1 infection was diagnosed with a positive HIV-1 RNA PCR in the first 18 months after birth and confirmed at 18 months with a negative HIV-1 p24 antibody test. The uninfected infants born to HIV-1-infected women were included in the study.

All infants were born after 37 weeks of gestational age and had a birth weight above 2500 grams. None of the children suffered from an infection at birth. The children in the comparison group consisted of healthy children born to healthy mothers and were matched for ethnicity.



Figure 6 | *In vitro* HIV-1 infection of CD4<sup>+</sup> T cells from HIV-1-exposed infants and healthy controls. Isolated cord blood mononuclear cells from HIV-1-exposed infants and healthy children were directly infected with YU2 or after a 4 day culture with, IL-2 or GM-CSF. HIV-1 infection was determined using an ELISA for p24 at day 6 after infection. After preculture with IL-2 or GM-CSF significantly higher p24 levels were detected in cultures from HIV-1-exposed infants than in the healthy controls. Median levels of p24 in cultures with IL1- $\beta$  were under the detection limit of the ELISA (25 ng/ml) and as such can not be statistically compared. Median is depicted with inter quartile range. Statistical analyses with Man-Whitney U-test.

**Cell isolation and cultures.** Cord blood mononuclear cells were isolated using Ficoll gradients and were directly stained with monoclonal antibodies for flow cytometry analyses. From 7 HIV-1-exposed infants and 7 controls the remaining cells were used in the cytokine stimulation assay and HIV-1 infection assay. The cells were labeled with CFSE and cultured for 6 days in IMDM, supplemented with 10% heat-inactivated fetal bovine plasma, penicillin (50 U/mL), streptomycin (50 mg/mL) and various stimuli; either IL-2 (100 IU/ml, Chiron Benelux), IL-1 $\beta$  (10 ng/ml), IL-4 (10 ng/ml), and IL-15 (20 ng/nl), GM-CSF (10 ng/ml) (all from R&D systems) or anti-CD2 and anti-CD28 (both from Sanquin).

Flow cytometry. The expression of surface markers on single cells was determined with a FACS CANTO (BD Biosciences) using fluorochrome-conjugated monoclonal antibodies. The antibodies included CD3 (Sanquin), CD4 (Pharmingen), CCR1, CCR2, CCR3, CCR4, CCR5, CCR6, CCR7, CCR8, CCR9, CCR10, CXCR4 and CXCR6 (all from R&D Systems).

HIV-1 infection assay. After isolation from cord blood, mononuclear cells from 7 HIV-1-exposed infants and 7 healthy controls were inoculated directly with YU2 (R5-strain) at a multiplicity of infection of 1 for 12 hours at  $37^{\circ}$ C or after a stimulation for 4 days with IL-2 (100 IU/ml, Chiron Benelux), GM-CSF and IL-1 $\beta$  (10 ng/ml, RnD systems). After removal of virus, the cells were cultured for 6 days in IMDM, supplemented with 10% heat-inactivated fetal bovine plasma, penicillin (50 U/mL), streptomycin (50 mg/mL) and with either IL-2 (100 IU/ml) or GM-CSF (10 ng/ml). Supernatant was analyzed for p24 with a standardized ELISA to determine HIV-1-infection in the mononuclear cell culture.

**Luminex.** A multiparameter Luminex bead assay was used to assess levels of soluble factors indicative a various immune responses; IL1- $\beta$ , IL-4, IL-5, IL-6, IL-8, IL-10, IL-13, IL-17A, IL-18, IFN- $\alpha$ , IFN- $\gamma$ , IP-10, TNF- $\alpha$ , MIP-1 $\alpha$ , MIP-1 $\beta$ , RANTES and GM-CSF, as indicated by the manufacturer (Invitrogen).

Statistical analyses. Flow cytometry data were analyzed with FlowJo (Tree Star Inc) and statistically investigated using Graphpad Prism Software (GraphPad Software Inc.). Medians are depicted with interquartile ranges (IQR) To compare continuous data, a Mann-Whitney U-test was performed between the two groups. In a two-way ANOVA analysis, we compared the expression of specific chemokine receptors between controls and HIV-1-exposed children across all cultures. Tree analyses were performed using Los Alomos tools (http://www.hiv.lanl.gov/content/sequence/HEATMAP/heatmap\_mainpage.html).

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#### **Author contributions**

M.J.B. and T.W.K. are the principal investigators who conceived and designed the study. M.J.B. and K.B. acquired the samples. N.A.K. helped designing the HIV-1 infection experiment. M.J.B., J.L.H. and M.J. performed the experiments. M.J.B. devised and performed the analyses and wrote the first draft of the manuscript with input from all authors supervised by T.W.K. All authors approved the final manuscript revisions.

#### **Additional information**

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