Human monocytotropic ehrlichiosis is caused by *Ehrlichia chaffeensis*, a Gram-negative bacterium lacking lipopolysaccharide. We have shown that fatal murine ehrlichiosis is associated with CD8\(^+\) T cell-mediated tissue damage, tumor necrosis factor-\(\alpha\), and interleukin (IL)-10 overproduction, and CD4\(^+\) Th1 hyporesponsiveness. In this study, we examined the relative contributions of natural killer (NK) and NKT cells in *Ehrlichia*-induced toxic shock. Lethal ehrlichial infection in wild-type mice induced a decline in NKT cell numbers, and late expansion and migration of activated NK cells to the liver, a main infection site that coincided with development of hepatic injury. The spatial and temporal changes in NK and NKT cells in lethally infected mice correlated with higher NK cell cytotoxic activity, higher expression of cytotoxic molecules such as granzyme B, higher production of interferon-\(\gamma\) and tumor necrosis factor-\(\alpha\), increased hepatic infiltration with CD8\(^+\)CD11c\(^+\) dendritic cells and CD8\(^+\) T cells, decreased splenic CD4\(^+\) T cells, increased serum concentrations of IL-12p40, IL-18, RANTES, and monocyte chemotactic protein-1, and elevated production of IL-18 by liver mononuclear cells compared with nonlethally infected mice. Depletion of NK cells prevented development of severe liver injury, decreased serum levels of interferon-\(\gamma\), tumor necrosis factor-\(\alpha\), and IL-10, and enhanced bacterial elimination. These data indicate that NK cells promote immunopathology and defective anti-ehrlichial immunity, possibly via decreasing the protective immune response mediated by interferon-\(\gamma\) producing CD4\(^+\)Th1 and NKT cells. (Am J Pathol 2010, 177:766–776; DOI: 10.2353/ajpath.2010.091110)

Human monocytotropic ehrlichiosis (HME) is an emerging tick-borne and a life threatening illness caused by *Ehrlichia chaffeensis*, an obligatory intracellular bacterium.\(^1,2\) HME can manifest as either mild disease with nonspecific flu-like symptoms or severe and fatal disease that presents as toxic shock-like syndrome, multi-organ failure, or adult respiratory distress syndrome.\(^2,6\) Severe HME in immunocompetent patients is thought to be due to immune-mediated pathology, which is attributed to severe inflammation in the absence of large quantities of ehrlichiae in the tissues.\(^3\) Doxycycline treatment is most effective if administered early in the course of illness.\(^1,7\) The lack of complete understanding of the pathogenic mechanisms of *Ehrlichia*-induced toxic shock-like syndrome is a major limitation in successful management of these patients. *Ehrlichia muris* and *Ixodes ovatus Ehrlichia* (IOE), which are closely related to each other and to *E. chaffeensis*, stimulate protective or pathogenic immune responses, respectively, in C57BL/6 mice.\(^5,9\) Intradermal (i.d.) and intraperitoneal (i.p.) infection with a high dose of IOE induces mild and fatal disease, respectively.\(^9\) Using these models, we and others have shown that antigen-specific interferon (IFN)-\(\gamma\) producing CD4\(^+\)Th1 cells and IgG2a antibodies play important roles in protective immunity against *Ehrlichia*.\(^4,10,11\) In contrast, fatal murine ehrlichiosis, which recapitulates the pathological and laboratory manifestations of fatal HME is associated with a suppressed or weak CD4\(^+\)Th1 immune response, marked lymphopenia, high levels of serum tumor necrosis factor (TNF)-\(\alpha\) and interleukin (IL)-10, and severe liver

**Immunopathology and Infectious Diseases**

**Natural Killer Cells Promote Tissue Injury and Systemic Inflammatory Responses During Fatal *Ehrlichia*-Induced Toxic Shock-Like Syndrome**

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injury, characteristics consistent with toxic shock syndrome.\textsuperscript{4,5,12} $\beta_{m^{-/}}$ mice, which are deficient in CD8$^+$ T cells and natural killer T (NK) cells, are resistant to fatal IOE-induced toxic shock when compared with wild-type and TAP$^{-/-}$ mice that are deficient in only CD8$^+$ T cells, suggesting that CD8$^+$ T cells mediate immunopathology and fatal ehrlichiosis.\textsuperscript{12} Interestingly, absence of NK T cells in CD1d$^{-/-}$ mice did not protect mice from fatal ehrlichiosis even though their absence prevented the development of liver injury and systemic inflammation.\textsuperscript{13} LPS-lacking \textit{Ehrlicha} can directly stimulate IFN-\gamma production by NK T cells in a CD1 days-dependent, but toll-like receptor 4-independent, manner.\textsuperscript{14} Further studies showed that NKT cells are critical for clearance of the bacterial burden in vivo, which may account for death of CD1d$^{-/-}$ mice following lethal ehrlichial infection.\textsuperscript{13}

NK cells are abundant in the liver, able to produce high levels of pro- and anti-inflammatory cytokines, and play an important role in innate immunity to microbial pathogens.\textsuperscript{15} NK cells contribute to the capacity of CD8$^+$ T cells to produce IFN-\gamma and to lyse \textit{Listeria monocyto-\textit{genes-} and \textit{Mycobacterium tuberculosis-infected monocytes.} \textsuperscript{16,17} Cross talk between NK cells and dendritic cells (DC) is essential for maturation of DC and activation of NK cells. IL-12, IL-15, and IL-18 produced by activated macrophages or mature DC activate cytotoxic NK cells and stimulate their IFN-\gamma production, which in turn promotes the expansion of CD4$^+$ Th1 cells.\textsuperscript{17–24} While NK cells play a protective role in some viral infections, NK cells cause hepatocyte apoptosis in models of viral hepatitis.\textsuperscript{25,26}

The current study was undertaken to specifically delineate the contributions of NK cells and NKT cells to the immunopathogenesis of \textit{Ehrlicha}-induced toxic shock-like syndrome using animal models of nonlethal and lethal ehrlichiosis caused by i.d. and i.p. inoculation of IOE, respectively. Our data provide a new finding suggesting that NK cells contribute to severe tissue injury and suppressed protective immunity during infection with these LPS-lacking, obligately intracellular bacteria.

\section*{Materials and Methods}

\subsection*{Mice, Bacteria, and Doxycycline Treatment}

Age-matched (8 to 12 weeks old) female wild-type C57BL/6 mice were used in all experiments (Jackson Laboratories, Bar Harbor, Maine). A highly virulent \textit{Ehrlicha} spp. (IOE), originally isolated from \textit{Ixodes ovatus} ticks in Japan, was used in all experiments.\textsuperscript{8,27,28} Mice were inoculated i.d. or i.p. as previously described\textsuperscript{9} with a high dose (~$5 \times 10^4$ ehrlichial copies) of IOE and were then sacrificed at different time points postinfection (p.i.). Antibiotic treatment of IOE-infected mice consisted of doxycycline injections (10 mg/kg body weight, two injections per day) given i.p. for three consecutive days (early treatment at 1, 2, and 3 days p.i. and late treatment at 5, 6, and 7 days p.i.)(Supplemental Figure S1, A–C at http://ajp.amipathol.org). Mice were housed in the animal research center at the University of Texas Medical Branch and Meharry Medical College in accordance with the institutional guidelines for animal welfare and monitored daily for signs of illness as previously described.\textsuperscript{9}

\subsection*{Flow Cytometry and Intracellular Cytokine Staining}

Analysis of the frequency of different cell subsets in liver and spleen including NK, NKT, CD11b$^+$, and CD11c$^+$ cells, as well as the measurement of expression of Fas and Granzyme B on splenic and liver mononuclear cells (LMNCs) was performed directly ex vivo. For intracellular cytokine staining, the cells were cultured at a concentration of $5 \times 10^6$ cells per well along with irradiated naive syngeneic spleen cells ($5 \times 10^6$ per well) as antigen presenting cells. The cells were stimulated \textit{in vitro} with IOE antigen (5 \mu g/ml) for 18 hours followed by 4 hours incubation with brefeldin A (BD Biosciences, San Diego, CA) and then analyzed by flow cytometry. Positive and negative control wells included cells stimulated with 5 \mu g/ml ConA or medium, respectively. Optimal concentrations of the following fluorophore-labeled anti-mouse monoclonal antibodies (BD Pharmingen, San Diego, CA) were used: CD3 (clone 145-2C11), CD11c (clone HL3), CD4 (clone GK.1.5), CD8a (clone 53-6.7), granzyme B (clone 16G6; eBioscience), CD11b (clone M1/70), CD4 (clone RM4-5), CD8 (clone 53-6.7), DX5 (clone 145-2C11), NK1.1 (clone PK136), CD19 (clone 1D3), B220 (clone RA3-6B2), IFN-\gamma (clone XMG1.2), TNF-\alpha (clone MP6-XT22), Fas (clone FAS), and corresponding isotype controls. Lymphocyte or monocyte populations were gated based on forward and side-scatter parameters, cells were analyzed using a BD FACSCalibur (BD Immunocytometry Systems, San Jose, CA) flow cytometer, and data were analyzed using FlowJo (Tree Star Inc., Ashland, OR).

\subsection*{Determination of Cytokine Levels in Serum}

The concentrations of IFN-\gamma, TNF-\alpha, and IL-10 in serum samples were determined by Quantikine enzyme-linked immunosorbent assay (R&D Systems, Minneapolis, MN) according to the manufacturer’s recommendations. Concentrations of several C-C chemokines and cytokines in serum were determined by microsphere multiplexed cytokine immunoassays (Bio-Plex Cytokine Assay, Bio-Rad Laboratories, Hercules, CA) following the manufacturer’s instructions.

\subsection*{Isolation of LMNCs and Hepatic IL-18 Measurement}

LMNCs were isolated and purified using a modified enzymatic dispersal protocol as described previously.\textsuperscript{29} Hepatocytes were removed by differential centrifugation (36 x g) for 1 minute at 4°C, and the final pellet containing LMNCs was resuspended in RPMI 1640 medium containing 10% fetal calf serum and 1% HEPES. LMNCs were purified using Lympholyte M (Cedarlane Laboratories, Burlington, NC). LMNCs were then stimulated with IOE antigen
NK Cell Depletion and NK Cytotoxicity Assays

A nonactivating polyclonal antibody (Ab) against asialo-GM1 (Wako Chemicals USA, Inc.) was used for NK cell depletion. C57BL/6 mice were injected i.p. with 40 μl of 1:100 dilution of anti-asialo GM1 Ab, or normal rabbit IgG isotype Ab on days −1, 2, and 5 of infection. Depletion of NK cells was approximately 95% efficient as determined by flow cytometric analysis of the number of CD3−NK1.1+ or CD3−DX5+ NK cells in the spleen and liver of depleted mice. Injection of anti-asialo GM1 Ab did not affect the number of CD11b+ monocytes and resulted in ≤4% depletion of CD3−NK1.1+ NK T cells as determined by flow cytometry (data not shown). The effect of depletion of NK cells on splenocyte cytotoxic activity was examined by a functional assay using YAC-1 target cells (ATCC, Manassas, VA). For NK cell cytotoxicity assays, splenocytes were isolated from lethally or nonlethally infected mice, uninfected controls, and lethally infected mice treated with anti-asialo GM1 or control Ab. Effector spleen cells were cultured with YAC-1 cells at different effector to target cell ratios. Cells were then collected and stained with propidium iodide and 3,3′-dioctadecyloxacarbocyanine (DiOC) using the LIVE/DEAD Cell-Mediated Cytotoxicity Kit (Molecular Probes, Inc., Eugene, OR) and flow analysis following the manufacturer’s recommendations. The percent specific lysis was determined as follows: 100 × (experimental release - spontaneous release)/(maximum release - spontaneous release). The percent specific lysis was normalized to the number of NK cells in the spleen.

Ehrlichial Load Determination in Collected Organs by Quantitative Real-Time PCR

Bacterial burden was measured in different organs using real-time PCR as previously described. Ehrlichial burdens were determined using an iCycler IQ multicolor real-time detection system (Bio-Rad, Hercules, CA). Primers and probes targeting both E. muris and koxes ovatus Ehrlichia dsb (which encodes a thio-disulfide oxidoreductase gene) and host glyceraldehyde-3-phosphate dehydrogenase genes were used as described before. The comparative cycle threshold method was used to determine the ehrlichial burdens as previously described.

Histopathology and Terminal Deoxynucleotidyl Transferase-Mediated dUTP Nick-End Labeling (TUNEL) Assays

Formalin-fixed, paraffin-embedded sections of liver were stained with H&E. Liver lesions were assessed semiquantitatively by four parameters: hepatocyte damage, number of lesions, size of inflammatory foci, and extent of perivascular inflammation as described before. TUNEL staining was performed on sections that show apoptotic cell death without focal necrosis as described previously.

Statistical Analysis

Data were analyzed using SigmaPlot software (SPSS, Chicago, IL), and P values were determined using the Student’s two-tailed t-test. P values less than 0.01 were considered highly significant, and those less than 0.05 were considered significant. Mouse groups contained three mice unless otherwise indicated, and error bars represent the SD for each group. SE of the mean was used for analysis of combined data from more than one experiment.

Results

Severe Ehrlichiosis Resulted in a Significant Increase in Hepatic CD8+ T Cells, But Not CD4+ T Cells, when Compared with Nonlethal Disease

Our previous studies indicated that fatal Ehrlichia-induced toxic shock-like syndrome is associated with a decreased number of CD4+Th1 cells and expansion of CD8+ T cells. We examined whether migration of CD4+ T cells to liver could account for the decreased total number of CD4+ T cells, using animal models of mild and severe monocytic ehrlichiosis caused by i.d./nonlethal and i.p./lethal inoculation with the same dose of highly virulent IOE, respectively. We chose to examine hepatic immune responses since the liver is the main site of ehrlichial infection and where NK and NKT cells are known to accumulate during other infections. The liver is also the main site of Ehrlichia-mediated tissue injury, which correlates with the marked elevation of liver transaminases in patients with HME. It is also considered the site of deposition of apoptotic bodies since the liver is the main site of ehrlichial infection and where NK and NKT cells are known to accumulate during other infections. The liver is also the main site of Ehrlichia-mediated tissue injury, which correlates with the marked elevation of liver transaminases in patients with HME. It is also considered the site of deposition of apoptotic cells, unlike that for splenic CD8+ T cells, is not due to their migration to the site of infection.
Lethal Ehrlichial Infection Enhanced Expression of Apoptotic and Cytotoxic Molecules in CD4+ and CD8+ T Cells, Respectively, Compared with Nonlethal Infection

We next examined the mechanisms that account for the decreased number of splenic CD4+ T cells observed in this study, as well as the apoptosis of CD4+ T cells and other hepatic cells during lethal infection as observed in our previous studies.5,9 Our data show that CD8+ T cells isolated from i.p./lethally infected mice are more cytotoxic and have greater expression of the intracellular cytotoxic molecule granzyme B, than those isolated from i.d./non-lethally infected mice (7.74% vs. 0.79%, Figure 2A). Surprisingly, splenic CD4+ T cells isolated from i.p./lethally infected mice also expressed significantly higher levels of intracellular granzyme B when compared with those from non-lethally infected mice (2.82% vs. 0.35%, Figure 2B). In addition, i.p./lethally infected mice had significantly greater expression of the apoptotic death receptor Fas on splenic CD4+ T cells compared with i.d./non-lethally infected mice (10.69% vs. 0.43%, Figure 2C).

Lethal Ehrlichial Infection Is Associated with Temporal and Spatial Changes in NKT Cells when Compared with Nonlethal Ehrlichial Infection

We next examined the contributions of NKT cells to mild and severe ehrlichiosis by assessing NKT cell expansion and migration to sites of infection. On day 5 p.i., the percentage (data not shown) and absolute numbers (Figure 3A) of CD3+/NK1.1+ NKT cells (measured by surface staining of CD3 and NK1.1 markers and mononuclear cell yield determined by trypan blue exclusion) in the spleen of i.p./lethally infected mice had begun to decline while those in the spleen and liver of i.d./nonlethally infected mice had begun to expand. On day 7 p.i., the total numbers (Figure 3B) and percentage (Figure 3, C and D) of CD3+NK1.1+ NKT cells in the i.p./lethally infected mice were significantly lower in both spleen and liver when compared with nonlethal infection. Interestingly, early doxycycline treatment on day 1, 2, and 3 p.i. abrogated the infection and protected 78% (7 out of 9) of mice from fatal disease (see Supplemental Figure S1A at
In contrast, late treatment with doxycycline on days 5, 6, 7 p.i. did not protect mice from fatal disease (see Supplemental Figure S1A at http://ajp.amjpathol.org), even though it restored the number of NKT cells in the spleen (see Supplemental Figure S1B at http://ajp.amjpathol.org) and decreased serum levels of cytokines associated with fatal disease such as IFN-γ, TNF-α, and IL-10 (see Supplemental Figure S1C at http://ajp.amjpathol.org). These data suggest that the decreased number of NKT cells observed later in lethal infection could be triggered by the high bacterial burden and further indicate that fatal ehrlichiosis is an immune-mediated disease.

**Lethal Ehrlichial Infection Induced Infiltration of CD11b+ and CD11c+ Cells into the Liver Parenchyma**

We next examined the possibility that increased hepatic infiltration with antigen presenting cells may contribute to inflammatory liver injury and fatal disease. Within the CD3+ NK1.1+ population of LMNCs, we detected significantly greater numbers of CD11b+ (Figure 4A) cells and CD11c+ cells (data not shown) in lethally infected mice when compared with nonlethally or uninfected mice (31.02% compared with 6.19% and 3.67%, respectively). Within this CD3+ NK1.1+ population, the CD11c+ LMNC in lethally infected mice lacked the lineage markers for B cells (CD19) (data not shown), but had a higher percentage of CD11c+ cells expressing CD8α (7.32%) when compared with nonlethally (1.76%) infected and uninfected mice (0.47%) (Figure 4B). These results indicate an expansion of hepatic CD8α+ CD11c+ DC in lethal ehrlichial infection.
Lethal Ehrlichial Infection Stimulated Expansion and Migration of NK Cells to the Liver

We next examined the effect of lethal and nonlethal ehrlichial infection on NK cell activation, expansion, and function. Lethal infection was associated with a significant decrease in the percentages and absolute numbers of NK cells in the spleen compared with nonlethal infection on days 5 (Figure 5, A and E) and 7 (Figures 5, C and G) p.i. In contrast, the percentage and total number of NK cells increased significantly in the liver of lethally infected mice on days 5 (Figure 5, B and F) and 7 (Figures 3D, and 5, D and H) p.i. compared with nonlethally infected mice. Hepatic CD3<sup>−</sup>NK1.1<sup>+</sup> NK cells from lethally infected mice also expressed CD11b, which is an activation marker of NK cells (Figure 4A). These data indicate that lethal ehrlichial infection induces spatial and temporal changes in activation and expansion or migration of NK cells to the liver, the main site of tissue injury.

NK and NKT Cells are a Major Source of Intracellular Granzyme B, IFN-γ, and TNF-α in Lethal Ehrlichiosis

Our previous studies demonstrated a significant association between overproduction of TNF-α, apoptosis of CD4<sup>+</sup>T cells, and tissue injury following lethal ehrlichial infection<sup>4</sup>,<sup>6</sup>,<sup>12</sup> To this end, we examined the cellular source of pro-inflammatory cytokines and cytotoxic molecules in infected mice. We detected significantly higher percentages of splenic NKT cells and NK cells, but not CD4<sup>+</sup> and CD8<sup>+</sup>T cells, producing TNF-α, IFN-γ, and granzyme B in lethally infected mice when compared with nonlethally or uninfected mice on day 7 p.i (Figure 6, A–C). However, since the number of NKT cells decreased substantially in lethally infected mice on day 7 p.i. (Figure 3B), these data indicate that NK cells are the major cellular source of TNF-α, IFN-γ, and granzyme B in lethally infected mice.
Lethal Infection Was Associated with Higher Serum Levels of IL-12p40, IL-18, MCP-1, and RANTES, When Compared with Nonlethal Infection

We measured the serum levels of several cytokines and chemokines in i.d./nonlethally and i.p./lethally infected mice that could contribute to substantial hepatic expansion and/or migration of NK cells, CD8\(^+\) T cells, DC, and monocytes to the liver. Compared with nonlethal infection, lethal ehrlichial infection was associated with significantly higher serum levels of IL-18, MCP-1, and RANTES (Figure 7A) on day 7 p.i. Interestingly, although the serum levels of IL-12p70 were not significantly different between mild and severe disease (data not shown), lethal disease was associated with a higher level of serum IL-12p40 than nonlethal disease (Figure 7A).

IL-18, together with IL-12p40, is a potent activator and inducer of cytotoxic NK cells and NKT cells in the liver and has been implicated in the pathogenesis of septic shock caused by other bacterial and viral infections.\(^{26,33,34}\) We examined here whether this cytokine is produced during severe ehrlichiosis. Strikingly, LMNCs (Figure 7B) and splenocytes (data not shown) from lethally infected mice produced significantly greater amounts of IL-18 than those from i.d./nonlethally infected mice. Data are representative of three mice per group, and similar results were obtained in three independent experiments. *P < 0.05, **P < 0.01.

**NK Cell Cytotoxic Function and Cytokine Production Are Enhanced in Lethal Ehrlichial Infection**

Next, we investigated whether lethal ehrlichial infection induced greater cytotoxic function of NK cells. To explore this possibility, the killing of YAC-1 cells by NK cells and the production of different cytokines were examined. Compared with splenocytes from uninfected or nonlethally infected mice, NK cells from the spleen of lethally infected mice produced significantly greater amounts of IL-18 than those from i.d./nonlethally infected mice. Data are representative of three mice per group, and similar results were obtained in three independent experiments. *P < 0.05, **P < 0.01.
NK cells in lethally infected mice significantly decreased the serum levels of IFN-γ, TNF-α, and IL-10 compared with lethally infected, control antibody-treated mice (Figure 9A). Collectively, these data suggest that NK cell cytotoxic functions are enhanced in i.p./lethally infected mice and that NK cells are major producers of pro-and anti-inflammatory cytokines observed during Ehrlichia-induced toxic shock-like syndrome.4,5

Enhanced Protective Immunity Against Ehrlichia and Attenuation of Tissue Injury in NK Cell-Depleted Mice

We measured the effect of NK cell depletion on the bacterial burden in lethally infected mice. Surprisingly, NK cell depletion resulted in a dramatic decrease in bacterial burden in the liver, lung, and spleen of lethally infected mice on day 7 p.i., when compared with similarly infected, control antibody-treated mice (Figure 9B). On day 9 p.i., a decreased bacterial burden was also evident in the liver, but not in the lung or spleen, which corresponds to a repopulation of NK cells in these organs (data not shown). These data suggest that NK cells suppress protective anti-ehrlichial immunity.

Next, we investigated the role of NK cells in the development of hepatic immunopathology following lethal ehrlichial infection. NK cell depletion abrogated liver injury, inflammation, and apoptosis as detected by histopathology (Figure 10, A and B) and TUNEL assays (Figure 10, C–F) when compared with lethally infected sham control mice on day 7 p.i. However, on day 9 p.i., both NK cell-depleted and control antibody-treated mice developed secondary necrosis, followed by death on day 11 p.i. Interestingly, mortality of the NK cell-depleted mice was accompanied by repopulation of CD3+ NK1.1+ NK cells in the spleen and liver as measured by flow cytometry (data not shown). These data demonstrate that the late cytokine storm and the extent of liver injury in response to lethal IOE infection are dependent on NK cells.

Discussion

The present study demonstrates that NK cell depletion in lethally infected mice improved protective immunity against virulent Ehrlichia and prevented the development of liver injury, providing evidence for a major pathogenic role of NK cells in fatal ehrlichiosis. Our murine models of mild and fatal ehrlichiosis clearly indicate that pathogenic host immune responses mediate tissue injury and fatal disease. Our studies show a differential localization or expansion of NK cells in the spleen and the liver of nonlethally and lethally infected mice. Higher numbers of NK cells in the liver, but not in the spleen, of lethally infected mice could be due to proliferation of hepatic NK cells or their trafficking to the Ehrlichia-infected microenvironment. Elevated production of IL-18, one of the main cytokines that promotes activation and expansion of NK cells,35 by LMNCs suggests that the hepatic microenvironment enhances NK expansion and proliferation. In addition, increased production of several chemokines including RANTES and MCP-1 that are potent NK cell and...
T cell chemoattractants\textsuperscript{36,37} suggest that enhanced trafficking of NK cells to the liver may contribute to severe tissue injury and host cell death in lethal disease. This conclusion is consistent with previous studies showing higher levels of RANTES in patients with fatal Mediterranean spotted fever rickettsiosis compared with patients with mild disease,\textsuperscript{38} as well as higher serum levels of CCL2 (MCP-1) in fatal murine primary and secondary ehrlichiosis compared with mild ehrlichiosis where blocking CCL2 ameliorates fatal disease.\textsuperscript{39} In an influenza model where severe disease is due to T cell-mediated injury, it has been suggested that TNF-\(\alpha\) regulates the production of MCP-1 and possibly other chemokines by host cells.\textsuperscript{40,41} Thus, it is possible that during fatal primary ehrlichiosis the production of TNF-\(\alpha\) by NK cells (Figure 6A), as well as by CD8\(^+\) T cells as we have shown before,\textsuperscript{5,12} promotes increased chemokine production and chemokine-dependent inflammation resulting in liver pathology. Migration of NK cells from the spleen and bone marrow into the liver has also been described in concanavalin A-induced hepatitis.\textsuperscript{34} NK cells have also been shown to migrate from the blood and spleen into the inflamed peritoneal cavity during polymicrobial sepsis induced by cecal ligation and puncture.\textsuperscript{42} In addition to the chemotactic effects of those chemokines, it is also possible that elevated levels of MCP-1 and RANTES, as observed in the current study, may contribute to the observed lymphopenia in fatal murine ehrlichiosis, mainly by reducing CD4\(^+\) T cell proliferation as suggested by other studies. For example, increased production of MCP-1 and RANTES during infection with Anaplasma marginale, the causative agent of human granulocytic ehrlichiosis (currently known as human anaplasmosis), was suggested as a potential mechanism that might play a role in bone marrow suppression and resulting cytopenias in human anaplasmosis.\textsuperscript{43,44} Other studies indicated that high concentrations of RANTES could enhance T cell apoptosis.\textsuperscript{45} Our further studies will examine the roles of these chemokines in the pathogenesis of fatal HME.

Our data clearly show that NK cells play a key role in tissue injury in severe and fatal disease. The mechanisms by which NK cells potentiate tissue injury after lethal ehrlichial infection are not completely understood. However, it is possible that NK cells mediate fatal ehrlichiosis by enhancing local and systemic inflammation and direct cytotoxic killing of host cells, as evidenced by the abrogation of systemic pro-inflammatory cytokine production (Figure 6, A–C and 9A) and minimal tissue damage (Figure 10) following NK cell depletion. In addition, NK cells may enhance the generation of pathogenic cytotoxic CD8\(^+\) T cells, which in turn promote tissue damage and apoptosis of protective CD4\(^+\) T cells as we have shown.

Figure 10. NK cells played a role in severe liver injury/apoptosis in lethal disease. On day seven p.i., livers were harvested from i.p. lethally infected, NK cell-depleted mice (A, C, and E), infected, sham-depleted mice (B, D, and F), or uninfected mice injected with PBS only (data not shown). H&E staining showed minimal hepatic pathology in the infected NK cell-depleted mice (A) compared with infected sham-depleted mice, which had multiple foci of host cell death (arrows). B: TUNEL assays showed substantially lower numbers of apoptotic cells in the infected NK cell-depleted mice (C and E) with fewer than one to two apoptotic cells observed per \(\times 20\) field compared with to 30 to 50 apoptotic cells per \(\times 20\) field in the infected sham-depleted control mice (D and F). Uninfected control mice had only one apoptotic cell per ten fields. Data are representative of several liver sections from three mice per group. Similar results were obtained in two independent experiments.
previously.\textsuperscript{12} In support of this conclusion, we observed increased cytotoxicity in splenic CD8\textsuperscript{+} T cells and greater infiltration of CD8\textsuperscript{+} T cells in the liver of lethally infected mice (Figures 1 and 2).

Depletion of NK cells in i.p./lethally infected mice resulted in more effective clearance of \textit{Ehrlichia}. These results are similar to our previous study showing decreased ehrlichial burden in β\textsubscript{2}m\textsuperscript{−/−} mice;\textsuperscript{12} however, differs from that observed in lethally infected CD1d\textsuperscript{−/−} mice, where absence of NK cells results in an increased bacterial burden.\textsuperscript{13} CD4\textsuperscript{+} Th1 and NKT cells mediate protection against \textit{Ehrlichia} through IFN-γ production. Our current and previous data\textsuperscript{1,12} show that fatal disease is associated with suppressed CD4\textsuperscript{+} T cell proliferation and apoptosis of CD4\textsuperscript{+} T cells. These observations were also associated with a significant increase in the number of intrahepatic NK cells (Figure 3D, 4A, and 5H). Therefore, it is possible that NK cells inhibit protective immunity against \textit{Ehrlichia} in lethal ehrlichiosis through either a direct or an indirect mechanism that causes apoptosis and/or suppression of IFN-γ producing CD4\textsuperscript{+}Th1 cells and NKT cells. Decreased IL-10 levels in NK cell depleted mice (Figure 9A) suggest that NK cell-mediated production of IL-10 may play a role in the generation of the CD4\textsuperscript{+}Th1 hyporesponsiveness in fatal ehrlichiosis.

Unexpectedly, NK cell-deficient mice succumbed to lethal disease similar to infected sham-depleted controls. Mortality of NK cell-depleted mice and sham-depleted controls was associated with the development of hepatocellular necrosis on day 9 p.i. and repopulation of NK cells in the spleen (data not shown). Repopulation of NK cells could be due to incomplete depletion of NK cells and/or restoration of their numbers, as NK cells are known to arise locally from hepatic hematopoietic stem cells in response to constitutive and inducible expression of IL-15 during other infections.\textsuperscript{46} Our data show that the peak increase in numbers of NK cells during severe ehrlichiosis occurs on day 7 p.i., when the experimentally administered anti-asialo GM1 antibody concentration in vivo might not have been sufficient to maintain depletion of the large numbers of expanded NK cells.

Cross talk between NK cells and antigen presenting cells, mainly DC, is critical for NK cell activation as well as DC maturation and induction of the acquired immune response against infections with several intracellular pathogens.\textsuperscript{47–52} Our results demonstrated substantial infiltration or expansion of CD8α\textsuperscript{+} DC in the liver, but not in the spleen, of lethally infected mice when compared with i.d./nonlethal infection. CD8α\textsuperscript{+} DCs produce large amounts of IL-12 and IL-18 and are responsible for activation and mobilization of NK cells\textsuperscript{22,24} and cross-priming of CD8\textsuperscript{+} T cells \textit{in vivo}.\textsuperscript{19} Our data demonstrated that lethal \textit{Ehrlichia} infection induced increased production of IL-18 by LMNCs as well as elevated serum levels of IL-12p40 and IL-18 (Figure 7A). Although, the cellular sources of these cytokines are not yet known, it is possible that hepatic CD8α\textsuperscript{+} DC produce IL-18 and IL-12p40 during lethal ehrlichial infection, which in turn enhances activation and cytotoxic functions of NK cells and CD8\textsuperscript{+} T cells leading to tissue injury. This conclusion is supported by studies showing that IL-18 plays a pathogenic role in several infectious diseases including LPS-induced multi-organ failure and sepsis where high levels of IL-18 correlate with the severity of hepatic damage, while absence of IL-18 enhances resistance to LPS-induced sepsis.\textsuperscript{15} In conclusion, these findings provide new information regarding the contribution of NK cells to the pathogenesis of fatal ehrlichiosis and the potential mechanisms by which NK cells facilitate the local inflammatory response leading to severe organ pathology during \textit{Ehrlichia}-induced toxic shock-like syndrome.

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