

Involvement of Toll-Like Receptor 4 in Innate Immunity to Respiratory Syncytial Virus

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The mammalian Toll-like receptor 4, TLR4, is an important component in the innate immune response to gram-negative bacterial infection. The role of TLR4 in antiviral immunity has been largely unexplored. In this study, the in vivo immune responses to respiratory syncytial virus (RSV) and influenza virus infection were examined in TLR4-deficient (C57BL/10ScNcr) and TLR4-expressing (C57BL/10Sn) mice. TLR4-deficient mice challenged with RSV, but not influenza virus, exhibited impaired natural killer (NK) cell and CD14⁺ cell pulmonary trafficking, deficient NK cell function, impaired interleukin-12 expression, and impaired virus clearance compared to mice expressing TLR4. These findings suggest that Toll signaling pathways have an important role in innate immunity to RSV.

The mammalian Toll-like receptors (TLR), a family of proteins structurally related to *Drosophila* Toll protein, were identified as critical regulators of innate immunity to a variety of microbes, including gram-positive and -negative bacteria, mycobacteria, and fungi (7, 12, 13, 14, 23, 24). Several studies suggest that Toll-like receptor 2 (TLR2) is a signaling receptor for gram-positive bacteria and fungi (2, 7, 24, 27, 32). TLR4 has recently been shown to be the signal-transducing receptor activated by bacterial lipopolysaccharide (LPS); and mice in which the TLR4 gene is either mutated or missing are hyporesponsive to LPS and do not respond with shock to gram-negative bacterial infection (21, 22). The conserved nature of the TLR and their role in innate immunity suggest that other infectious pathogens, such as viruses, might also activate the innate immune response via the Toll signaling pathway. Several studies from our laboratory indicate that the innate immune response is an important component of respiratory syncytial virus (RSV) immunity (29, 30). Recent in vitro evidence that TLR4 and CD14 are involved in the innate immune responses to the RSV F glycoprotein prompted us to investigate the role of TLR4 in the in vivo immune response to RSV infection.

RSV is the single most important cause of lower respiratory tract disease in infants and young children worldwide and is a high priority for vaccine development. Unfortunately, a broad range of approaches toward RSV vaccine development has not yet produced a safe and effective vaccine. RSV is a member of the family *Paramyxoviridae*, existing as an enveloped virus containing a negative-sense, single-stranded RNA genome that encodes 11 proteins. Two major surface viral glycoproteins, the F (fusion) and G (attachment) glycoproteins, are associated

with the induction of neutralizing antibodies and long-term protective immunity (4, 25). The F glycoprotein has been reported to induce primarily a Th1-type immune response, while the G glycoprotein induces primarily a Th2-type immune response contributing to both protective immunity and disease pathogenesis (6, 9, 20). We have recently reported that the G glycoprotein can alter pulmonary trafficking of natural killer (NK) cells and polymorphonuclear cells (PMNs), inhibit Th1-cytokine expression, and alter macrophage inflammatory protein 1 α (MIP-1 α), MIP-1 β , MIP-2, monocyte chemoattractant protein 1 (MCP-1), and interferon-inducible protein 10 (IP-10) chemokine mRNA expression in bronchoalveolar cells (29, 30), suggesting that the innate immune response is important during RSV infection. It is anticipated that a better understanding of viral and host mechanisms that affect RSV immunity might facilitate vaccine development.

In this study, we examine the in vivo innate immune response in TLR4-deficient (TLR4^{null}) C57BL/10ScNcr mice and wild-type C57BL/10Sn mice (TLR4^{wt}) challenged with RSV or influenza virus to address the role of TLR4 in the innate immune response to a respiratory virus infection. We chose to compare RSV and influenza virus for several reasons. Influenza virus, like RSV, is a major respiratory pathogen, causing significant morbidity and mortality in young children, immunocompromised adults, and the elderly. A member of the *Orthomyxoviridae* family, influenza virus is an enveloped negative-stranded RNA virus, and like RSV, it primarily infects the respiratory epithelium, causing cytopathology and inflammation of the respiratory tract. The results of this study indicate that TLR4 is important for activation of the innate immune response to RSV infection and may be important to the pathogenesis of RSV disease.

MATERIALS AND METHODS

Animals. Specific-pathogen-free, 6-to-8-week-old, female C57BL/10ScNcr (TLR4^{null}) (National Cancer Institute, Bethesda, Md.) and C57BL/10Sn (TLR4^{wt}) (Jackson Laboratory, Bar Harbor, Maine) mice were examined. The C57BL/10ScNcr strain is homozygous for a null mutation of the TLR4 gene (21,

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33, 34). A related mouse strain, C57BL/10ScCr (not used in these studies), has a reported defect in interleukin 12 (IL-12)-induced production of gamma interferon (IFN- γ) (16). However, the C57BL/1-ScNcr mice used in the present study were IL-12 responsive and expressed IFN- γ at levels similar to those of the wild type. The C57BL/10Sn strain was used as the control. All studies were performed in accordance with the guidelines of the Institutional Animal Care and Use Committee.

Virus infection. Mice were anesthetized by intraperitoneal administration of Avertin (2,2,2-tribromoethanol, 0.2 ml/g of body weight; Sigma-Aldrich, St. Louis, Mo.) and intranasally (i.n.) challenged with 10^6 PFU of the A2 strain of RSV or 240 HAU of a mouse-adapted strain of influenza A virus (HKx31) in Dulbecco's phosphate-buffered saline (PBS) (GIBCO Laboratories, Grand Island, N.Y.).

Viruses. The A2 strain of RSV was propagated in Vero cells (ATCC CCL 881) as previously described (29). The mouse-adapted HKx31 strain of influenza A virus was cultured in embryonated eggs and harvested as previously described (28).

Collection of BAL cells and preparation of NK cells. Mice were anesthetized with Avertin and exsanguinated by severing the right caudal artery. Bronchoalveolar lavage (BAL) cells were harvested by lavaging the lungs with PBS containing 1% bovine serum albumin (Sigma). Natural killer cells (DX5⁺) were enriched from BAL cells using the MACS separation system (Miltenyi Biotec, Inc., Auburn, Calif.) according to the manufacturer's instructions. Viability was assessed by trypan blue exclusion. The purity of DX5⁺ cell populations ranged from 80 to 90% as determined by flow cytometry (Becton Dickinson, Mountain View, Calif.).

Fluorescence-activated cell sorter analysis. The procedure used for extracellular staining of BAL cells was modified for microculture staining as described previously (30). Fluorescein isothiocyanate-conjugated or phycoerythrin-conjugated anti-CD3 ϵ (145-2C11), anti-CD45R/B220 (RA3-6B2), anti-NK cell (2B4 and DX5), antineutrophil (RB6-8C5), and anti-CD14 (rmC5-3) monoclonal antibodies and isotype antibody controls were used (BD-PharMingen, San Diego, Calif.). Ten thousand events were collected and analyzed using a FACScan and Cell Quest software (Becton Dickinson, San Diego, Calif.). Intracellular cytokine staining was modified for microculture staining as described previously (30). Briefly, BAL cells were incubated in PBS containing Golgi Stop (PharMingen) for 3 h at 37°C to accumulate intracellular cytokines. The cells were washed in PBS, stained with anti-CD3, anti-CD4, or anti-CD8 antibody, fixed, and permeabilized in Cytofix/Cytoperm (PharMingen). Cells were washed and resuspended in an appropriate dilution of anti-IL-2 (JES6-5H4), anti-IL-4 (BVD4-1D11), anti-IL-5 (TRFK5), anti-IL-12 (C15.6), or anti-IFN- γ (XMG1.2) antibody diluted in PBS containing permeabilization buffer, stained, washed, and analyzed as described previously (30) (all from PharMingen). IL-12 expression presented in Table 1 was determined by subtracting the total IL-12 expression by un gated BAL cell populations from IL-12 expression by CD3⁺ BAL cell populations. The total cytokine-expressing CD4⁺ or CD8⁺ cell populations were determined by multiplying the percent cytokine-expressing CD4⁺ or CD8⁺ cells by the total BAL cell population.

NK Cytotoxicity assays. YAC-1 cells (ATCC TIB 160) were used as target cells. The cells were maintained in minimal essential medium (SMEM) (GIBCO Laboratories, Grand Island, N.Y.) containing 10% fetal bovine serum (FBS) (37°C, 8% CO₂). Two different cytotoxicity assays were used to evaluate NK cytotoxicity. YAC-1 target cells were labeled with either 2 μ l of 3 mM DIOC₁₈ (Molecular Probes, Eugene, Ore.) (30 min at 37°C) or 200 μ Ci of ⁵¹Cr (Amersham Pharmacia Biotech, Quebec, Canada) (18 h at 37°C), washed twice with PBS, and resuspended in SMEM containing 10% FBS. BAL cell populations, unfractionated or purified for NK (DX5⁺) cells, were used as effector cells. Effector BAL cell populations, pooled from 5 to 10 mice, were harvested, and NK cytotoxicity tests were performed using a two-color fluorescence assay (L-7010; Molecular Probes) per the manufacturer's directions or by lysis of ⁵¹Cr-labeled target cells. Briefly, effector cells and 10⁴ DIOC₁₈-labeled targets were plated in a 96-well V-bottomed plate (Costar, Cambridge, Mass.) to yield effector-to-target ratios of 40:1, 20:1, 10:1, and 5:1 and incubated for 4 h at 37°C. The percent lysis was calculated as described previously (3). Spontaneous lysis was determined for DIOC₁₈-labeled targets incubated in the absence of effectors. Spontaneous lysis ranged from 4 to 13%. NK cytotoxicity was also determined using a standard ⁵¹Cr release assay as described previously (5). Briefly, effector cells were incubated (4 h at 37°C) with 10⁴ ⁵¹Cr-labeled target cells, in triplicate wells, using 40:1, 20:1, 10:1, and 5:1 E/T ratios. As appropriate, EGTA-MgCl₂ (5 and 10 mM, respectively; Sigma) was added to the corresponding wells. To address the effect of IL-12 on cytotoxicity, BAL cells were harvested from five mice per group, pooled, and cultured for 24 h with or without 2 ng of rIL-12 (R&D Systems, Minneapolis, Minn.)/ml. The cultured cells were incubated with ⁵¹Cr-labeled YAC-1 target cells at 40:1, 20:1, 10:1, and 5:1 effector-to-target ratios for

TABLE 1. Intracellular cytokine expression by BAL cells after primary infection with RSV or influenza virus

Day	Phenotype/ cytokine	Mean % positive IC cytokine expression of BAL cells \pm SEM ^a			
		RSV (wt)	RSV (null)	FLU (wt)	FLU (null)
5	CD3 ⁺ /IL-2	11 \pm 2	10 \pm 1	15 \pm 1	25 \pm 1^b
	CD3 ⁺ /IL-4	12 \pm 1	11 \pm 1	20 \pm 2	28 \pm 1
	CD3 ⁺ /IL-5	10 \pm 0	8 \pm 1	15 \pm 1	18 \pm 2
	CD3 ⁺ /IFN- γ	13 \pm 1	11 \pm 1	22 \pm 2	31 \pm 1^c
	CD3 ⁻ /IL-12	29 \pm 6	7 \pm 1^d	28 \pm 2	6 \pm 4^e
7	CD3 ⁺ /IL-2	6 \pm 2	8 \pm 2	10 \pm 2	8 \pm 1
	CD3 ⁺ /IL-4	8 \pm 4	10 \pm 0	18 \pm 2	13 \pm 2
	CD3 ⁺ /IL-5	6 \pm 1	4 \pm 1	21 \pm 7	19 \pm 2
	CD3 ⁺ /IFN- γ	11 \pm 3	9 \pm 1	26 \pm 1	18 \pm 2
	CD3 ⁻ /IL-12	30 \pm 1	21 \pm 1^f	30 \pm 3	39 \pm 1 ^g
11	CD3 ⁺ /IL-2	7 \pm 1	7 \pm 1	7 \pm 1	4 \pm 1
	CD3 ⁺ /IL-4	8 \pm 1	8 \pm 0	9 \pm 0	4 \pm 1
	CD3 ⁺ /IL-5	13 \pm 0	13 \pm 2	9 \pm 1	6 \pm 1
	CD3 ⁺ /IFN- γ	10 \pm 2	11 \pm 3	11 \pm 1	5 \pm 1
	CD3 ⁻ /IL-12	32 \pm 6	15 \pm 5 ^h	34 \pm 4	35 \pm 5

^a C57BL/10ScNcr (null) and C57BL/10ScSn (wt) mice were i.n. infected with the A2 strain of RSV or the HKx31 influenza A (FLU) virus. BAL samples from three to five mice were examined at days 5, 7, and 11 p.i. The results of two separate experiments are shown. Data are represented as percent positive intracellular (IC) expression of BAL cells \pm SEM. IL-2, -4, and -5, and IFN- γ expression by CD3⁺ cells and IL-12 expression by CD3⁻ cell populations are presented. Bold type indicates significant differences between TLR4^{wt} and TLR4^{null} cytokine expression.

^b P value comparing FLU(wt) to FLU(null) is 0.014.

^c P value comparing FLU(wt) to FLU(null) is 0.029.

^d P value comparing RSV(wt) to RSV(null) is 0.034.

^e P value comparing FLU(wt) to FLU(null) is 0.050.

^f P value comparing RSV(wt) to RSV(null) is 0.011.

^g P value comparing FLU(wt) to FLU(null) is 0.103.

^h P value comparing RSV(wt) to RSV(null) is 0.093.

4 h at 37°C. Spontaneous and maximum ⁵¹Cr releases were determined by incubating target cells with either medium or with 2% saponin in the absence of effector cells. The percent specific ⁵¹Cr release (percent cytotoxicity) was calculated as [(experimental cpm - spontaneous cpm)/(maximum cpm - spontaneous cpm)] \times 100. Spontaneous lysis counts ranged from 317 to 550 cpm. The use of either assay yielded comparable results.

CTL assay. Spleens from TLR4^{wt} and TLR4^{null} mice were harvested at days 5 and 11 post-RSV infection. Spleen cells were restimulated in vitro for 7 days with RSV-infected spleen cells, which had been infected with RSV for 3 h at 37°C in RPMI containing 10% FBS. Secondary in vitro cytolytic cell activity was measured using a standard ⁵¹Cr release assay. SVB6KHA target cells (*H-2^b*) were incubated with RSV (multiplicity of infection, 100) and ⁵¹Cr (200 μ Ci) for 18 h at 37°C, washed two times in RPMI containing 10% FBS, and distributed in V-bottom 96-well plates (Costar) at a concentration of 10⁴ cells/100 μ l. Effector cells were added at an effector-to-target ratio of 50:1 and serially diluted to 2:1 in triplicate. Plates were centrifuged at 1,200 rpm for 2 min and then incubated at 37°C for 5 h. After incubation, 100 μ l of the supernatant was removed, and radioactivity was measured in a gamma counter (Perkin-Elmer Life Sciences, Boston, Mass.). Spontaneous and maximal release was measured by incubating the target cells in media alone or in 10% Triton X-100 detergent, respectively. Specific release of ⁵¹Cr from target cells was calculated as follows: [(experimental cpm - spontaneous cpm)/(maximum cpm - spontaneous cpm)] \times 100.

Virus titers in lung tissue. Lungs were aseptically removed from three to five mice per group at days 3, 5, 7, and 11 post-RSV or influenza virus infection and stored at -70°C until the assay. Identical weights (-0.1 g of tissue) of individual lung samples were homogenized in 1 ml of Dulbecco's PBS (GIBCO), and 10-fold serial dilutions of the lung homogenates were subsequently added to confluent Vero cell monolayers to detect RSV titers or confluent Madin-Darby Canine Kidney (MDCK) cell monolayers to detect influenza virus titers. Following adsorption for 1 to 2 h at 37°C, monolayers were overlaid with either Dulbecco's modified Eagle medium (GIBCO) containing 10% FBS for Vero cells or Dulbecco's modified Eagle medium containing 0.1% BSA (Sigma), 0.1

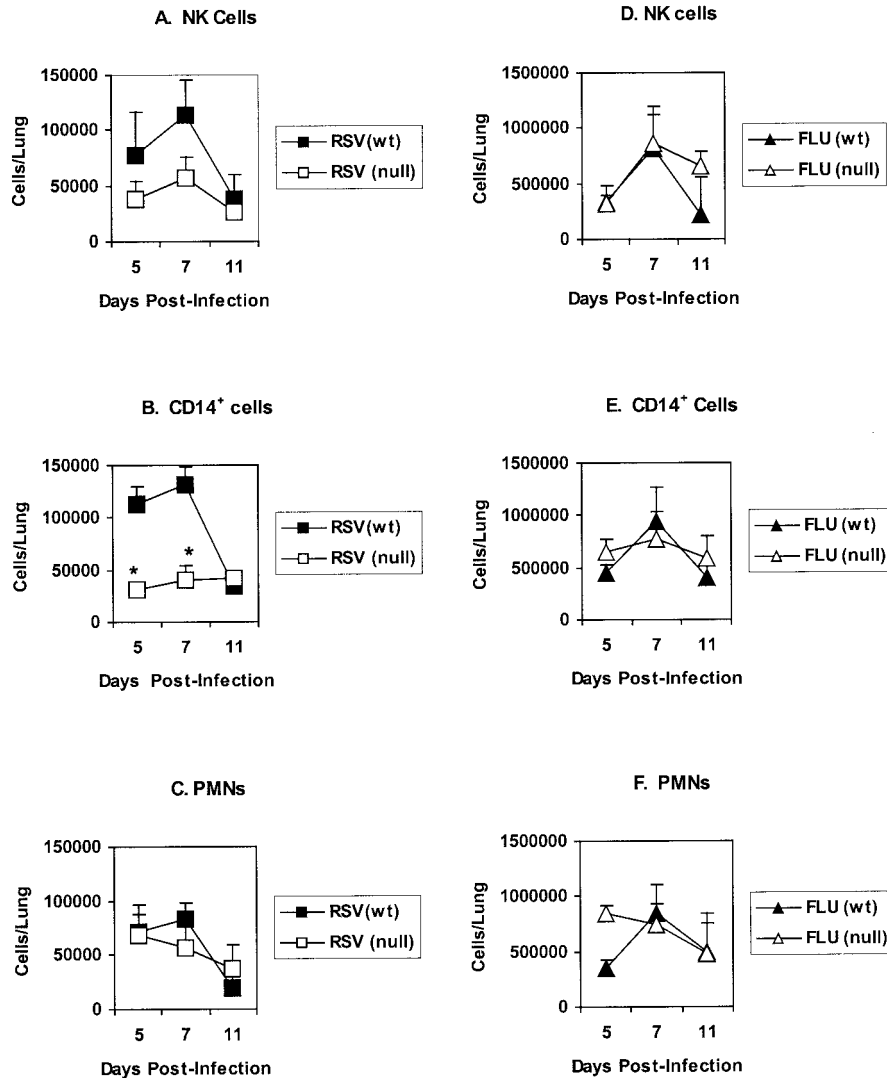


FIG. 1. Decreased pulmonary infiltration of NK cells and CD14⁺ cells in RSV-infected TLR4^{null} mice. Flow-cytometric analysis of BAL cell subsets from TLR4-deficient (TLR4^{null}) and TLR4-expressing (TLR4^{wt}) mice infected i.n. with the A2 strain of RSV virus (RSV) or the HKx31 influenza A virus (FLU). BAL samples were stained with antibodies against NK cells (A and D), CD14⁺ monocytes/macrophages (B and E), and PMNs (C and F). Data are presented as the mean number of cells/lung \pm standard error of the mean at days 5, 7, and 11 p.i. from three independent experiments. Asterisks indicate a significant difference ($P < 0.05$) between TLR4^{null} and TLR4^{wt} mice.

μ g of trypsin (Sigma)/ml, 0.8% agar (BioWhittaker, Rockland, Maine) for MDCK cells. The monolayers were incubated at 37°C for 3 to 4 days, and RSV plaques were enumerated after immunostaining with monoclonal antibodies against the G and F glycoproteins (130-2G and 131-2A, respectively) as described previously (30).

Influenza virus plaques were enumerated after fixing MDCK monolayers with 80% methanol and staining them with 2% crystal violet in 10% ethanol.

Statistical analysis. Statistical significance was determined using a Student *t* test, and a probability of < 0.05 was considered statistically significant.

RESULTS

Decreased NK and CD14⁺ pulmonary cell infiltration in RSV-infected TLR4^{null} mice. To investigate the importance of TLR4 in the immune response to respiratory viral infections, we first examined the phenotype of BAL cells infiltrating the lungs of TLR4-deficient (TLR4^{null}) mice and TLR4-expressing (TLR4^{wt}) control mice following i.n. challenge with RSV or

influenza virus. Flow cytometry was used to identify NK cells (2B4⁺), monocytes and macrophages (CD14⁺), and PMN (RB6-8C5⁺) present in the BAL cell population at days 5, 7, and 11 postinfection (p.i.). Representative data from three separate experiments is presented in Fig. 1. For RSV-infected TLR4^{wt} mice, the pulmonary influx of NK cells (Fig. 1A) and CD14⁺ cells (Fig. 1B) peaked on day 7 p.i. and subsequently declined by day 11 p.i., as was observed following influenza virus infection of TLR4^{wt} mice (Fig. 1D and E). In contrast, RSV-infected TLR4^{null} mice displayed a pattern of NK and CD14⁺ cell infiltration very different from that for TLR4^{wt} mice (Fig. 1A and B). RSV-infected TLR4^{null} mice exhibited decreased infiltration of pulmonary NK cells and CD14⁺ cells ($P < 0.05$) between days 5 and 7 p.i. with RSV (Fig. 1A and 1B); however, no significant differences in the total numbers of pulmonary BAL cells were detected between TLR4^{wt} and

TLR4^{null} mice (1×10^6 to 6×10^5 cells/lung versus 0.85×10^6 to 5×10^5 cells/lung) throughout the period (days 5 to 11 p.i.) examined. In contrast to the case with RSV infection, the numbers of NK and CD14⁺ cells in the BAL of influenza virus-infected TLR4^{wt} and TLR4^{null} mice were comparable (Fig. 1D and E). The altered BAL infiltration by NK and CD14⁺ cells observed in RSV-infected TLR4^{null} mice was not associated with altered numbers of PMN cells or T cells (CD3⁺) present in the BAL after infection with RSV or influenza virus (Fig. 1C and F), and no consistent differences in the percentages of these cell types were detected between TLR4^{wt} and TLR4^{null} mice. Of note, there was a small increase (3 to 5%) in B cells (B220⁺ CD45R⁺) in TLR4^{null} mice compared to wild-type controls; however, this response may be inherent in TLR4^{null} mice, since the small increase in B cells was detected after either RSV or influenza virus infection.

To address another aspect of TLR4 cell activation that may affect cell trafficking, we examined Th1 (IL-2, IFN- γ , and IL-12) and Th2 (IL-4 and IL-5) intracellular cytokine expression by BAL cells in TLR4^{null} and TLR4^{wt} mice infected with RSV or influenza virus (Table 1). Overall, the absence of TLR4 (TLR4^{null}) was associated with reduced IL-12 expression by BAL cells from RSV-infected mice. For RSV-infected TLR4^{null} mice, IL-12 expression by BAL cells was significantly decreased from that observed for TLR4^{wt} mice on day 5 p.i. (7% versus 29%; $P < 0.05$) and day 7 p.i. (21% versus 30%; $P < 0.05$), and by day 11 p.i., IL-12 expression remained decreased but was not statistically different between TLR4^{null} and TLR4^{wt} mice. It is possible that decreased pulmonary CD14⁺ cell infiltration observed after RSV infection, and/or inadequate activation of these cells, may relate to the lowered IL-12 expression observed in TLR4^{null} mice. Although TLR4^{null} mice infected with influenza virus initially exhibited reduced IL-12 expression at day 5 p.i. (6% versus 28%, $P < 0.05$) compared to TLR4^{wt} mice, there was no association of decreased IL-12 expression with decreased CD14⁺ infiltration (Fig. 1) or NK cytotoxicity (data not shown). By day 7 p.i., the level of IL-12 expression for influenza virus-infected TLR4^{null} mice was comparable to that for TLR4^{wt} mice (39% versus 30%) (Table 1) and remained comparable to that for TLR4^{wt} mice throughout the time course (Table 1 and Fig. 2). Interestingly, the percentage of cells expressing Th1 and Th2 cytokines was higher for influenza virus-infected mice than for RSV-infected mice; however, this difference may reflect the increased sensitivity of mice to the mouse-adapted influenza virus infection (31) compared to human RSV infection. Of note, at day 5 p.i., levels of IFN- γ and IL-2 cytokine expression were increased in TLR4^{null} mice relative to TLR4^{wt} mice infected with influenza virus. The absence of TLR4 did not significantly alter Th2 cytokine expression by CD3⁺ cells following RSV or influenza virus infection (Table 1), and neither RSV nor influenza virus infection elicited a predominately Th1- or Th2-type cytokine response (Tables 1 and 2). Of the CD3⁺ BAL cells examined, CD4⁺ cells predominantly expressed both Th1 (IFN- γ and IL-2) and Th2 (IL-4 and IL-5) cytokines (Table 2).

Decreased NK cell cytotoxicity in TLR4^{null} mice after RSV infection. The decreased pulmonary NK cell trafficking in RSV-infected TLR4^{null} mice observed on days 5 and 7 p.i. (Fig. 1A) suggested that differences in NK cytotoxicity may occur

between TLR4^{wt} and TLR4^{null} mice; thus, we examined NK cell cytotoxicity at day 7 post-RSV or influenza virus infection of TLR4^{wt} and TLR4^{null} mice. Examination revealed that NK cells from RSV-infected TLR4^{null} mice are deficient in their ability to lyse YAC-1 target cells compared to cells from RSV-infected TLR4^{wt} mice (Fig. 3). Following RSV infection, the level of NK cytotoxicity in TLR4^{null} mice was significantly diminished (43% at 40:1) from that in TLR4^{wt} mice (60% at 40:1; $P < 0.05$). In contrast, levels of NK cytotoxicity were similar in TLR4^{null} and TLR4^{wt} mice infected with influenza virus (Fig. 3). These results suggest that the innate immune response to influenza virus infection may be less dependent of TLR4 activation, but it is possible that other TLR activation pathways may be involved in this response.

The impaired NK cytotoxicity observed for TLR4^{null} mice infected with RSV might be explained by differences in NK cell numbers (Fig. 1A) and/or by a functional defect in NK cytotoxicity. To examine this possibility, NK cytotoxicity assays were performed using equal numbers of purified NK cells from the BAL of TLR4^{wt} and TLR4^{null} mice (Fig. 4). At day 7 p.i., RSV-infected TLR4^{null} mice had significantly impaired NK cell function compared to TLR4^{wt} mice (Fig. 4). To address one mechanism for this defect, differences in calcium-dependent perforin-mediated cytotoxicity were examined for NK cells from TLR4^{null} mice using the calcium chelator EGTA-Mg²⁺ (Fig. 4). More than 79% of NK cytotoxicity was inhibited in the presence of EGTA-Mg²⁺, suggesting that target cell lysis was mediated primarily through a perforin-dependent mechanism. IL-12 has been shown to enhance NK activity, and RSV-infected TLR4^{null} mice have altered IL-12 expression (Table 1 and Fig. 2). It has also been reported that a substrain of TLR4^{null} mice (C57BL/10ScCr) that was originally derived from the C57BL/10ScNcr strain used in these studies, has a defect in IL-12 responsiveness (16, 33, 34). Therefore, we examined the effect of exogenous addition of 2 ng of IL-12/ml on NK cell cytotoxicity in TLR4^{null} and TLR4^{wt} responses (Fig. 5). NK cytotoxicity in RSV-infected TLR4^{null} mice was restored to wild-type levels by the addition of IL-12 (Fig. 5). Thus, the C57BL/10ScNcr mice used in these studies are IL-12 responsive.

Delayed RSV clearance in TLR4^{null} mice. Since TLR4 appeared to be associated with IL-12 expression, as well as NK cell trafficking and cytotoxicity in RSV (but not influenza virus)-infected animals, we next examined whether the absence of TLR4 altered viral clearance (Fig. 6). Examination revealed that the diminished innate immune response (i.e., decreased IL-12 expression and decreased NK cytotoxicity) in TLR4^{null} mice was associated with a compromised ability to clear acute RSV infection compared to that of RSV-infected TLR4^{wt} mice. The lung titers of RSV in TLR4^{null} mice were higher at all time points examined and were significantly higher at day 11 p.i. ($P < 0.05$) (Fig. 6A). In three of four experiments, mean titers of RSV virus in TLR4^{null} mice had increased 100- to 1,000-fold over titers of virus in TLR4^{wt} mice at day 11 p.i. (Fig. 6A). Notably, the delayed RSV viral clearance in TLR4^{null} mice compared to that in TLR4^{wt} mice correlated with their impaired NK activity (Fig. 3 and 4). In contrast, similar levels of virus clearance were observed for influenza virus-infected TLR4^{null} and TLR4^{wt} mice (Fig. 6B).

CD8⁺ CTL responses for RSV-infected TLR4^{null} mice were comparable to those for TLR4^{wt} mice at day 5 p.i., suggesting

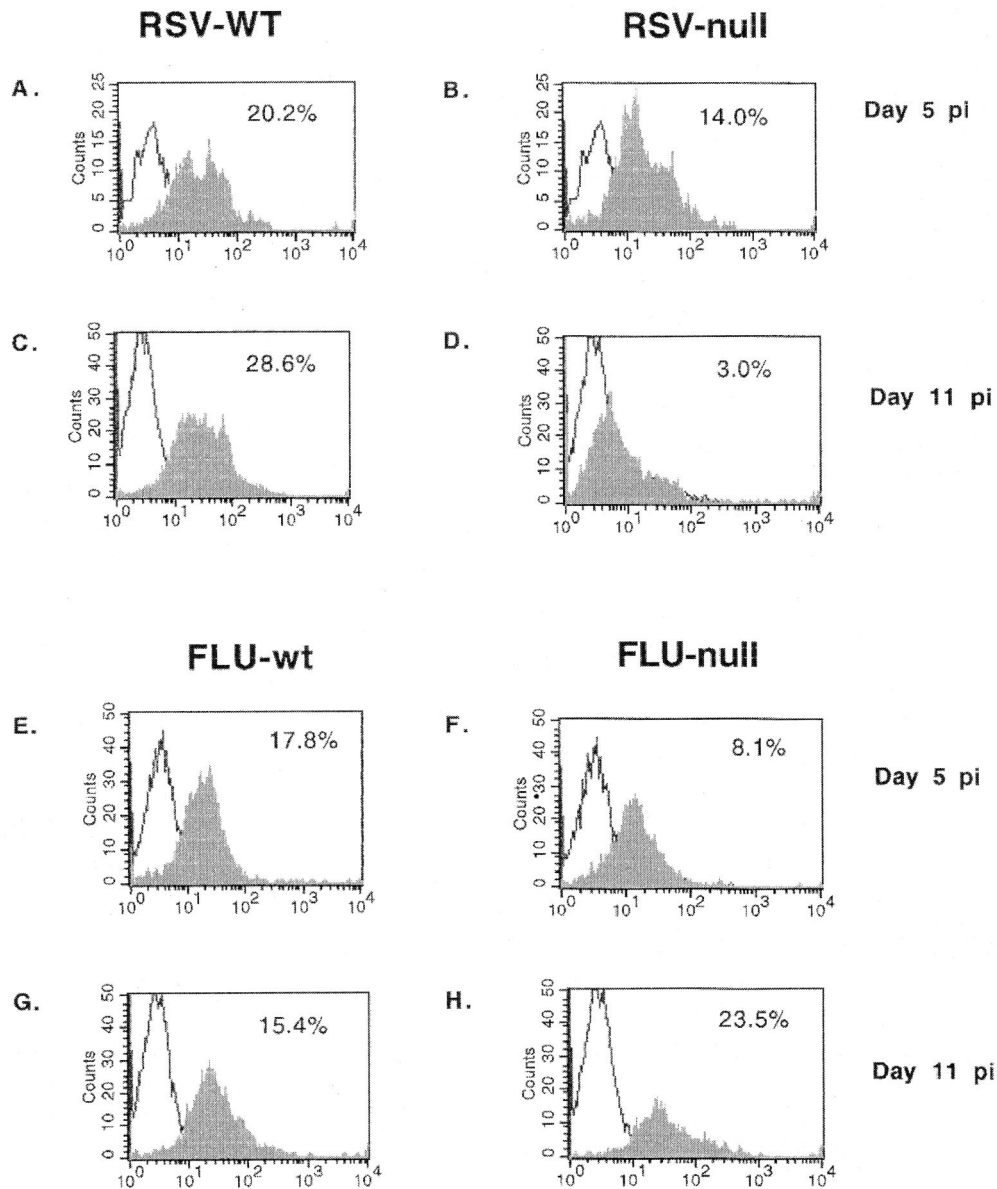


FIG. 2. Decreased expression of IL-12 by BAL cells after RSV infection. BAL samples from RSV- or influenza virus-infected TLR4^{wt} and TLR4^{null} mice were harvested and examined for intracellular IL-12 expression (x axis). IL-12 expression at day 5 post-RSV infection for TLR4^{wt} and TLR4^{null} samples is shown in panels A and B, respectively. IL-12 expression at day 5 post-influenza virus infection for TLR4^{wt} and TLR4^{null} samples is shown in panels E and F, respectively. IL-12 expression at day 11 post-RSV infection for TLR4^{wt} and TLR4^{null} is shown in panels C and D, respectively. IL-12 expression at day 11 post-influenza virus infection for TLR4^{wt} and TLR4^{null} samples is shown in panels G and H, respectively.

that early elevated RSV titers observed in TLR4^{null} mice were not associated with impaired CD8⁺ CTL responses in these mice. Interestingly, the CTL activity in TLR4^{null} mice at day 11 p.i. was modestly increased over the cytolysis observed for TLR4^{wt} mice (Fig. 7). This slight increase in CTL activity may be associated with the clearance of RSV by TLR4^{null} mice, which was observed at day 15 p.i. (data not shown).

DISCUSSION

In this study, we examined the *in vivo* role of TLR4 in the innate immune response to two major respiratory viral patho-

gens, RSV and influenza virus. We demonstrated that TLR4 is important in the innate immune response to RSV infection but does not appear to be as important following influenza virus infection. Acute RSV infection in TLR4^{wt} mice was similar to the pulmonary cell infiltration and pattern and magnitude of cytokine expression that we and others have previously observed in BALB/c mouse studies (8, 10, 17, 31). In contrast, associated with the absence of TLR4 is diminished IL-12 expression and diminished numbers of pulmonary NK cells and CD14⁺ cells in RSV-infected mice. In addition, the limited numbers of NK cells that traffic to the lungs in TLR4^{null} mice in response to RSV infection have significantly impaired cyto-

TABLE 2. Intracellular cytokine expression by CD4⁺ and CD8⁺ BAL cells after primary infection with RSV or influenza virus

Day	Phenotype/cytokine	IC cytokine expression by CD4 ⁺ and CD8 ⁺ BAL cells ± SEM ^a			
		RSV (wt)	RSV (null)	FLU (wt)	FLU (null)
5	CD4 ⁺ /IL-2	18,000 ± 4,700	15,000 ± 5,000	65,000 ± 8,600	25,000 ± 5,000
	CD4 ⁺ /IL-4	13,000 ± 3,000	20,000 ± 6,000	60,000 ± 7,800	50,000 ± 9,000
	CD4 ⁺ /IL-5	15,000 ± 4,000	19,000 ± 6,000	14,000 ± 19,000	99,000 ± 2,000
	CD4 ⁺ /IFN-γ	18,000 ± 5,000	25,000 ± 8,000	39,000 ± 8,000	30,000 ± 6,000
	CD8 ⁺ /IL-2	3,000 ± 900	8,000 ± 2,500	5,900 ± 800	8,000 ± 1,500
	CD8 ⁺ /IL-4	5,000 ± 1,300	6,000 ± 1,800	5,900 ± 800	8,000 ± 1,500
	CD8 ⁺ /IL-5	7,000 ± 1,800	8,000 ± 2,500	24,000 ± 3,000	58,000 ± 11,000
	CD8 ⁺ /IFN-γ	8,000 ± 2,200	15,000 ± 5,000	4,600 ± 1,000	11,000 ± 2,000
7	CD4 ⁺ /IL-2	3,000 ± 580	2,200 ± 840	22,800 ± 7,400	30,000 ± 3,900
	CD4 ⁺ /IL-4	2,700 ± 500	1,800 ± 690	21,300 ± 7,000	28,000 ± 3,580
	CD4 ⁺ /IL-5	1,800 ± 350	2,600 ± 990	30,400 ± 10,000	38,000 ± 4,880
	CD4 ⁺ /IFN-γ	3,300 ± 630	2,200 ± 840	35,000 ± 11,000	46,000 ± 5,850
	CD8 ⁺ /IL-2	900 ± 180	800 ± 300	9,000 ± 3,000	15,000 ± 1,950
	CD8 ⁺ /IL-4	900 ± 180	800 ± 300	7,600 ± 2,000	10,000 ± 1,300
	CD8 ⁺ /IL-5	600 ± 120	1,200 ± 460	11,000 ± 3,000	18,000 ± 2,280
	CD8 ⁺ /IFN-γ	1,500 ± 290	1,000 ± 380	20,000 ± 6,400	28,000 ± 3,580
11	CD4 ⁺ /IL-2	14,000 ± 4,400	7,400 ± 1,600	36,000 ± 12,000	30,500 ± 4,200
	CD4 ⁺ /IL-4	10,000 ± 3,100	6,200 ± 1,300	30,000 ± 10,000	28,000 ± 3,900
	CD4 ⁺ /IL-5	10,000 ± 3,100	5,100 ± 1,100	33,000 ± 11,000	20,000 ± 3,000
	CD4 ⁺ /IFN-γ	11,000 ± 3,500	6,200 ± 1,300	39,000 ± 13,000	33,000 ± 4,600
	CD8 ⁺ /IL-2	3,600 ± 1,000	4,000 ± 800	15,000 ± 5,000	11,100 ± 3,000
	CD8 ⁺ /IL-4	2,600 ± 1,000	2,300 ± 500	12,000 ± 4,000	11,100 ± 3,000
	CD8 ⁺ /IL-5	2,400 ± 800	2,300 ± 500	9,000 ± 3,000	11,100 ± 1,500
	CD8 ⁺ /IFN-γ	5,400 ± 1,700	4,000 ± 800	15,000 ± 5,000	25,000 ± 3,400

^a C57BL/10ScNcr (null) and C57BL/10ScSn (wt) control mice were i.n. infected with the A2 strain of RSV or the HKx31 influenza A (FLU) virus. BAL samples from three mice per group were taken at days 5, 7, and 11 p.i. Data represent total CD4⁺ and CD8⁺ BAL cell IL-2, -4, or -5 or IFN-γ expression ± SEM (as described in Materials and Methods). IC, intracellular.

toxicity, primarily through a defect in perforin-mediated lysis which is reflected by the decreased ability to clear RSV infection in these mice. It is possible that inadequate activation of CD14⁺ cells may contribute to the low IL-12 expression observed at day 5 p.i. for RSV- and influenza virus-infected TLR4^{null} mice and to the delayed virus clearance observed in RSV-infected TLR4^{null} mice (11). Although RSV and influenza virus infection induce similar host cell cytopathologies,

these viruses likely induce different innate immune responses, since influenza virus-infected TLR4^{null} mice did not display altered immune cell trafficking or impaired NK cytotoxicity compared to the case with TLR4^{wt} mice, as was observed following RSV infection of these mice.

The apparent lack of similar deficiencies in the innate immune response observed for influenza virus infection of TLR4^{null} mice, compared to RSV infection of TLR4^{null} mice, suggests that the defects observed are not generalized but are specific to the response to RSV infection, suggesting that RSV stimulates the NK cell and CD14⁺ cell response through the TLR4 pathway. The major surface glycoproteins of RSV (G and F) are likely candidates for inducing TLR4 activation. Recent in vitro studies from our laboratory have demonstrated that the F glycoprotein can stimulate TLR4 activation in a CD14-dependent fashion (11). In contrast, the RSV G glycoprotein appears to suppress rather than promote NK and PMN activation, as indicated by decreased cell trafficking to the lungs of RSV-infected mice and decreased virus clearance (30). Members of our group have also demonstrated that the G glycoprotein alters MIP-1α, MIP-1β, MIP-2, MCP-1, and IP-10 mRNA expression during RSV infection (29). MIP-1α, MIP-1β, MIP-2, MCP-1, and IP-10 are chemokines that contribute to recruitment and trafficking of innate immune cells into the lung (29).

One possible explanation for the diminished NK cytotoxicity

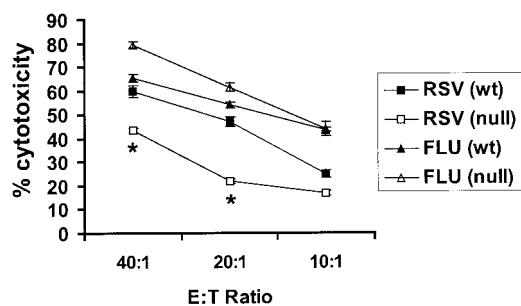


FIG. 3. Impaired NK cell activation in TLR4^{null} mice after RSV infection. BAL samples from TLR4^{null} and TLR4^{wt} mice were harvested 7 days post-RSV or -influenza virus (FLU) infection. NK lytic activity against YAC-1 target cells was determined. Results are expressed as the mean of three independent experiments ± the standard error of the mean. Asterisks indicate a significant difference (*P* < 0.05) between TLR4^{null} and TLR4^{wt} mice. E:T Ratio, effector-to-target ratio.

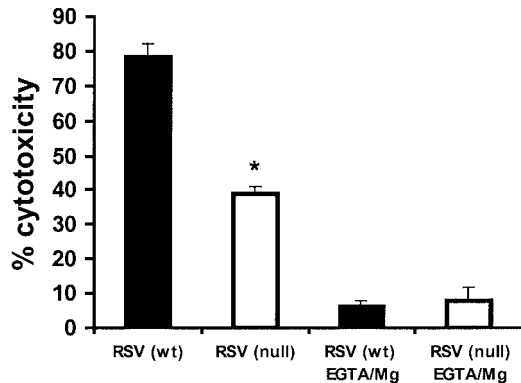


FIG. 4. RSV-induced NK cytotoxicity is perforin dependent. Seven days p.i., BAL samples from TLR4^{null} (open bar) and TLR4^{wt} (closed bar) mice were examined following RSV infection. Samples were purified for NK cells (DX5⁺) (80 to 90% enrichment) by positive selection using the MACS separation system. NK lytic activity against YAC-1 target cells was assessed in the presence or absence of EGTA-MgCl²⁺ at effector-to-target ratios of 40:1 (presented), 20:1, 10:1, and 5:1 (data not shown). The asterisk indicates a significant difference ($P < 0.05$) between TLR4^{null} and TLR4^{wt} mice. The results are representative of two independent experiments.

observed in RSV-infected TLR4^{null} mice may be associated with a failure to activate the MyD88/IRAK/NF- κ B signaling cascade (1, 15). TLR4, the IL-1 receptor, and the IL-18 receptor generate intracellular signaling by a shared molecular protein cascade involving sequential recruitment of MyD88 and IRAK to the receptors, phosphorylation of IRAK, and activation of Traf6. This ultimately leads to translocation of NF- κ B to the nucleus and gene transcription. MyD88-deficient mice have impaired IL-18-mediated NK-cell activation (1). Also, IL-18 is a potent inducer of NK cells and upregulator of perforin-mediated NK activity (18, 19). Similar to MyD88 (-/-) mice, IL-18-deficient mice also display defects in NK cytotoxicity (26). Further, antibody blockade of IL-18 has been shown to result in diminished IFN- γ expression and lymphocyte infiltration of the lungs of mice challenged with another respiratory virus, adenovirus (35). In addition, decreased expression

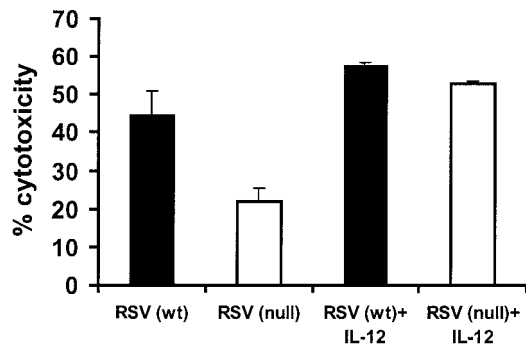
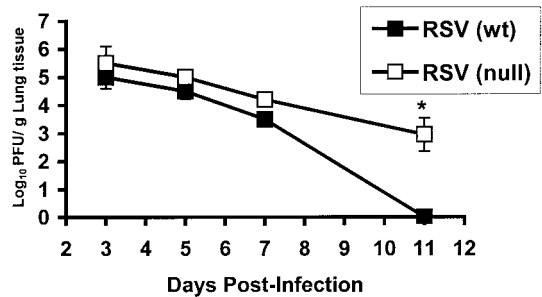


FIG. 5. Addition of IL-12 enhances NK-mediated cytotoxicity in RSV-infected TLR4^{null} mice. BAL samples from TLR4^{null} and TLR4^{wt} mice were examined for cytotoxicity 7 days post-RSV infection. Effector BAL cells were cultured in the presence of 2 ng of IL-12/ml for 24 h. NK lytic activities against YAC-1 target cells at effector-to-target ratios of 40:1 (presented), 20:1, 10:1, and 5:1 (data not shown) were analyzed. A representative experiment is shown.

A



B.

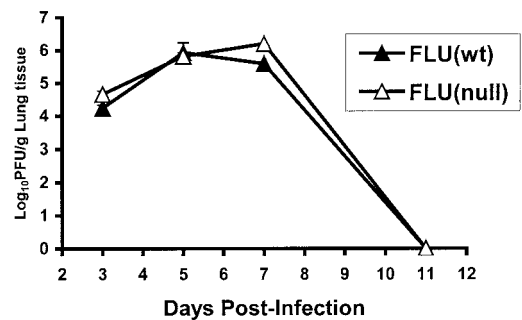


FIG. 6. Delayed RSV clearance in TLR4^{null} mice. The lungs of TLR4^{null} and TLR4^{wt} mice were harvested at days 3, 5, 7, and 11 p.i. with RSV (A) or influenza virus (FLU) (B). The asterisk indicates a significant difference ($P < 0.01$) between TLR4^{null} and TLR4^{wt} mice. The results are expressed as mean log₁₀ PFU/g \pm the standard error of the mean from four separate experiments. In each experiment, three to five mice per group at each time point were analyzed.

of IL-12 in the TLR4^{null} mice infected with RSV, but not influenza virus (Table 1), may also contribute to altered NK cell trafficking and cytotoxicity. IL-12 enhances NK activity and has been shown to act synergistically with IL-18 in the activation of NK cells (26). Our experiments suggest that alteration of NK cell trafficking and NK effector function are more severe in RSV-infected TLR4^{null} mice than in influenza virus-infected

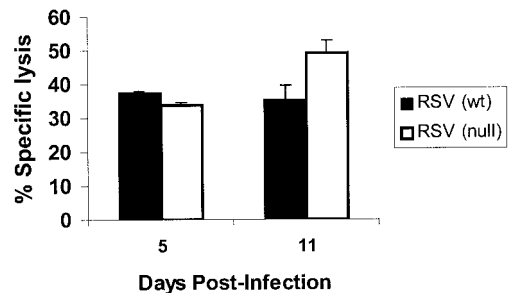


FIG. 7. Kinetics of CTL activity in RSV-infected TLR4^{null} mice. TLR4^{wt} and TLR4^{null} mice were sacrificed on days 5 and 11 post-RSV infection. Data are values for percent specific lysis at an effector-to-target ratio of 50:1 and are representative of two independent experiments.

TLR4^{null} mice. We hypothesize that TLR4^{null} mice may fail to generate an effective NK cell response to RSV because of defects in generating both direct (MyD88/IRAK) and cytokine-mediated (IL-12) signals for NK cell activation. In addition, one study suggests that a related strain of TLR4^{null} mice may also have a defect in IL-12 responsiveness (16). However, our experiments suggest that the C57BL/10ScNcr mice used in the present study are IL-12 responsive. In future studies, we will address the contributions of background genes, such as the IL-12 receptor, on TLR4 function in viral pathogenesis using congenic TLR4 knockout mouse strains.

The study presented here provides strong evidence for the involvement of TLR4 activation in the *in vivo* innate response to nonbacterial microbial pathogens: in particular, RSV. Based upon our *in vitro* studies that show that the RSV F glycoprotein can activate TLR4 (11), we hypothesize that the F glycoprotein is also important for TLR4 activation during the immune response to RSV infection and that TLR4 is an important contributor to the RSV innate immune response. Understanding the mechanisms that contribute to RSV innate immunity may allow new approaches for prevention and/or treatment of RSV-associated disease.

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