Respiratory syncytial virus infection modifies and accelerates pulmonary disease via DC activation and migration

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ABSTRACT

In the present studies, we have established that RSV can elicit a more pathogenic environment dependent on improper DC-associated sensitization. Our initial studies demonstrated that RSV, but not influenza, infection during an allergen exposure into the airway induced a more severe allergen response. The RSV-induced exacerbation included an increased Th2 cytokine response and pathophysiology as monitored by AHR and mucus overproduction. DCs played a central role in the allergen-induced responses, as instilling RSV-infected BMDC into the airway could recapitulate a live virus challenge. With the use of CCR6-/- mice that have a primary defect in the recruitment of mDC subsets, reduced exacerbation of disease was observed when RSV was administered along with allergen. Furthermore, sensitization of mice with RSV-infected BMDC into the airway produced a more severe immune response to a live virus challenge. Subsequently, using RSV-infected BMDC from CCR7-/- mice (that do not migrate efficiently to LNs) to sensitize the exacerbated response demonstrated that the response was dependent on DC migration to the LN. Finally, the ability of RSV-infected DCs to elicit an exacerbated, allergen-induced pathogenic response could be maintained for as long as 3 weeks, suggesting that RSV-infected DCs themselves created an altered immune environment that impacts off-target mucosal responses that could have prolonged effects. J. Leukoc. Biol. 94: 5-15; 2013.

Introduction

The induction of immune responses in the host often determines the nature and the severity of the ensuing illness during infectious diseases and may even dictate future local immune responses. Studies have shown that patients with severe respiratory viral infections have an increased risk for the development of chronic pulmonary diseases [1–5]. A number of respiratory viruses have been implicated in the induction of pulmonary diseases, including infections with RSV, rhinovirus, influenza, parainfluenza, and adenovirus. How the interactions of specific viruses, underlying immune conditions, genetic predisposition, and local mucosal environments contribute to disease severity is not entirely clear. Recent studies in children [6] and adults [7] support this contention related to severe asthmatic exacerbations. Understanding the mechanisms that promote inappropriate immune responses to viral infections may lead to better therapy. RSV infections are especially detrimental and have been associated with early and persistent pulmonary disease, especially in premature infants.

The activation of the innate immune response, including antigen-presenting DCs, and recruitment of T cells that amplify and skew the immune response toward more intense pathology likely contribute to more severe disease and clinical crisis in asthmatic patients. Whereas various immune responses to viral infections may have common mechanisms of activation, RSV appears to promote responses that on their own can cause severe pulmonary problems [8–10]. Recent evidence suggests that RSV also has a significant role in elderly patient populations, as well as in patients with chronic obstructive pulmonary disease [11]. In addition, the specific mechanism(s) of immune regulation identified in RSV studies may be relevant to other viral infections that also must be recognized and properly cleared to prevent a more pathogenic disease progression.

A number of studies have shown that RSV infection of DCs significantly alters the ability of mDCs to express class II and costimulatory molecules, causes the preferential expression of IL-10, and impairs the induction of a Th1 response [12–16]. Correspondingly, mucosal mDCs have been shown to preferentially skew immune responses in the lung toward a more pathogenic Th2 phenotype and lead to severe asthma-like responses, including increase airway hyper-responsiveness and

Abbreviations: AHR=airway hyper-reactivity, BMDC=bone marrow-derived DC, CRA=cockroach allergen, mDC=myeloid DC, NS=influenza nonstructural, PAS=periodic acid Schiff, pDC=plasmacytoid DC, qPCR=quantitative PCR, RIG-I=retinoic acid-inducible gene 1, RSV=respiratory syncytial virus

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mucus overexpression [17–19]. The migration of DC subsets to the draining LNs indicates that lung-derived and LN resident DCs are involved in the presentation of antigen to CD4 and CD8 T cells [20]. The present work examines whether RSV alters the immune response to a simultaneous allergen sensitization and secondly, whether RSV-infected DCs played a role in this process. Our results show that RSV-infected DCs contribute to this altered immune environment and also that the DCs need to migrate back to the draining LNs for these perturbations to occur. We hypothesize that viral infections provide an inappropriate maturation stimulus that induces the DC presentation of environmental allergens, thus increasing the sensitization, and accelerates the chronic allergic responses.

MATERIALS AND METHODS

Animals and virus

Female BALB/c/J and C57BL/6 mice, 6–8 weeks of age, were obtained from The Jackson Laboratory (Bar Harbor, ME, USA). Animals with specific genetic deletions of CCR6 and CCR7 were graciously supplied by Professor Sergio Lira at Mount Sinai Medical School (New York, NY, USA). All mice were housed under specific pathogen-free conditions within the animal care facility at the University of Michigan (Ann Arbor, MI, USA). The RSV A strain was derived from a clinical isolate at the University of Michigan and was propagated and titered as Line 19 [21, 22]. The mouseadapted influenza virus (Strain A/PR/8/34) was purchased from American Type Culture Collection (Manassas, VA, USA) and was used at a nonlethal dose following an initial experiment to determine an infectious dose that was tolerable by all mice (i.e., 100% survival).

Allergen sensitization

In our first set of experiments, the DC-instilled animals were simultaneously given a CRA (HollisterSteir, Spokane, WA, USA) sensitization into the trachea. Fourteen days later, a second administration of allergen (1 μ g) was given into the nasal passages, followed by two intratrachea allergen challenges (4 μ g/mouse) on Days 19 and 21 of the protocol. As our allergen is a skin-test allergen for clinical use, it has very low or no endotoxin and therefore, produces a strong Th2 response without induction of IFN or IL-17.

In a second protocol designed to determine whether RSV-infected DCs have a long-term alteration of the pulmonary immune environment, an intranasal administration of CRA was given 7 days (or 21 days) after the RSV-infected DC instillation. Subsequently, the animals were given intratrachea allergen challenges on Days 13 (or 27 days) and 15 (or 29 days) post-DC administration. Twenty-four hours after the final allergen challenge, AHR was assessed, and the lungs and LNs of mice were harvested for analysis.

Isolation and RSV infection of mouse DCs

BM cells were flushed from the femurs of mice. After RBC lysis, BM cells were cultured at 1×10^6 cells/ml in RMPI medium (RPMI 1640, 10% FCS, 1 μ M Na pyruvate, 2 mM L-glutamine, 100 μ g/ml streptomycin, 100 U/ml penicillin) containing 10 ng/ml GM-CSF (R&D Systems, Minneapolis, MN, USA), and fresh medium was added after 3, 6, and 8 days. On Day 10, DCs were harvested. These cells represent a mDC population with an intermediate maturation status, as most cells were CD11c⁺ (90%) and also expressed CD11b (95%). DCs were washed and plated at 2×10^6 /ml and then infected with RSV (~0.8×10⁶ PFU/ml) or medium alone overnight at 37°C.

CFSE staining of DCs

After RSV infection, 4 \times 10⁶ DCs/ml were stained with the green fluorescent dye CFSE (Molecular Probes, Eugene, OR, USA) at a final concentra-

tion of 2.5 μ M for 5 min at room temperature. After extensive washing, RSV-infected or control CFSE⁺ DCs (0.5×10^6 cells/50 μ l) were injected intratracheally on Day 0 after ketamin/xylazin anesthesia. Hereafter, 1, 2, or 3 days later, lung draining mediastinal LNs were isolated and dispersed using 0.2% collagenase (Type IV, Sigma-Aldrich, St. Louis, MO, USA) in RPMI 1640 with 10% FCS at 37°C for 45 min. Single-cell dispersions were used for flow cytometric analysis.

Flow cytometric analysis of LNs

After blocking nonspecific binding by FcR, cells were stained with antimouse CD11C, CD11b, MHC II, CD80, CD86 or CD40, 7-amino actinomycin D, and Annexin V (all from PharMingen, San Diego, CA, USA; and eBioscience, San Diego, CA, USA). Cells were analyzed on a Cytomics FC500 flow cytometer (Beckman-Coulter, Brea, CA, USA).

RSV infection and allergen sensitization (see Fig. 1)

Mice were infected with RSV (~1×10⁵ PFU/mice) on Day -1 and the next day, were given a CRA (HollisterStier; 3 μ g/mouse) sensitization into the trachea. Fourteen days later, a second administration of allergen (1.5 μ g/mouse) was given intranasally, followed by two intratrachea allergen challenges (4 μ g/mouse) on Days 19 and 21. Twenty-four hours after the final allergen challenge, AHR was assessed, and the lungs and LNs of mice were harvested for analysis.

DC transfer and RSV or CRA exposure

After extensive washing, RSV-infected or control DCs $(2 \times 10^5 \text{ cells}/50 \ \mu\text{l})$ were injected intratracheally on Day 0 after ketamin/xylazin anesthesia. Whereas we cannot rule out completely that the virus was not transferred, when we examined the lungs at 2–4 days post-DC transfer (a time that is normally peak virus propagation), we do not find any evidence by PCR of viral mRNA expression. Thus, the magnitude of any low-level infection must be very limited.

RSV infection

On Day 7, mice were an esthetized lightly and given RSV (1×10^5 PFU/mouse) by intratracheal infection, and after 8 additional days, the animals were analyzed for AHR, cytokine responses, and pathologic changes.

Measurement of airway responses

Airway reactivity in anesthetized mice was measured as described previously [23]. Briefly, mice were anesthetized with sodium pentobarbital, and the trachea was cannulated and ventilated using a pump ventilator in a mouse plethysmograph. After baseline measurements, mice were injected i.v. with 2.5 μ g methacholine (Sigma-Aldrich), and the peak airway resistance was recorded.

LN restimulation assays

Single-cell suspensions of lung draining LNs were made by isolating draining lung LNs, pushing cells through a nylon mesh using a syringe, and then lysing RBCs. Cells were cultured at a concentration of 6×10^6 cells/ml in the presence of CRA (2 µg/ml) or medium and incubated at 37°C. After 2 days, supernatants were isolated and stored at -80° C until further analysis. A standardized sandwich ELISA (R&D Systems) or Bio-Plex (Bio-Rad Laboratories, Hercules, CA, USA) was performed to measure cytokine and chemokine levels in culture supernatants.

Histology

Lungs were maintained in formalin for a further 24 h before being processed into paraffin using standard histological techniques. Lung tissue sections were stained with H&E for analysis of inflammatory cell accumulation and alcian blue/PAS stain for assessment of mucus production.

Statistical analysis

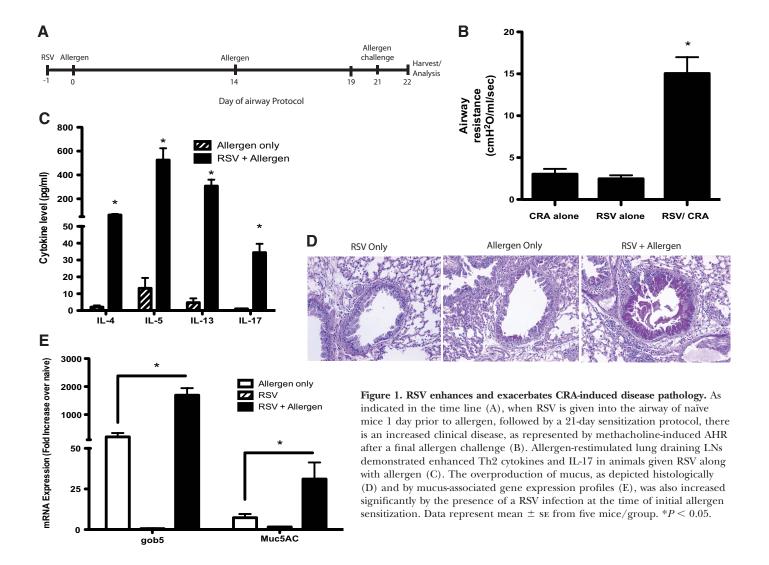
Significance was determined using one-way ANOVA with 95% confidence intervals, followed by a Student Neuman Keuhl's post-test. With a P value of <0.05, differences were considered significant.

RESULTS

RSV infection enhances allergen sensitization and severe pathology associated with mDC accumulation and responses

A strong correlation in patient populations has been found between early pulmonary viral infection and subsequent development of allergic asthma [24–27]. Although the mechanism is unknown, it may be that the coincidental exposure of the viral infection along with an environmental allergen lead to the most severe disease. To address this latter concept, we examined whether instilling allergen along with viral infection would alter the development/sensitization of the allergen response. In our studies, the infection of mice with RSV (1×10^5 PFU/mouse) along with allergen sensitization into the airway clearly elicited a more severe allergic disease after an additional challenge with the allergen 21 days later (**Fig. 1A**). These data include enhanced AHR responses to methacholine (Fig. 1B), increased potentially pathogenic Th cytokines, IL-4, IL-5, IL-13, and IL-17 (Fig. 1C), more intense goblet cell development in the upper airways (Fig. 1D), and increased expression of mucus-associated genes gob5 and muc5ac (Fig. 1E). In contrast, when animals were infected with H1N1 influenza virus and simultaneously sensitized to allergen using a similar protocol as RSV, we observed a decrease in allergen-specific cytokine production and no change in the mucus gene expression or the histologic appearance (Supplemental Fig. 1). These data suggest that the RSV, but not influenza, infection leads to increased sensitization to allergen and caused an intensified local skewing response [28]. Thus, not all viral infections can promote an enhanced allergic airway response.

To determine the contribution of the mDC to the intensified response, BMDCs were infected with RSV in vitro (MOI, 1.0). This procedure addressed the effects of the RSV infection of airways, as DCs do not propagate RSV but did sensitize animals to the responses as described above. To address virus propagation from DCs that were transferred, we examined a



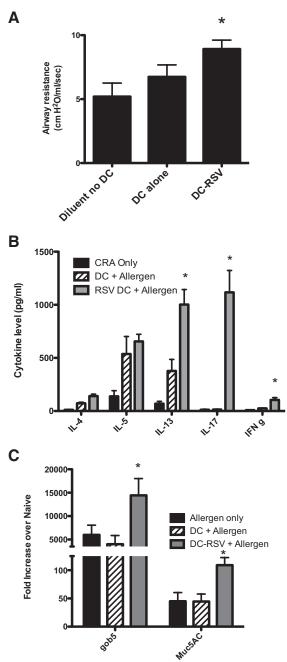


Figure 2. RSV-infected DC instilled into the airways of mice can replace a live RSV infection for exacerbation of allergen-induced responses. After an overnight infection, the washed DCs were transferred into the lungs of naïve mice by intratracheal administration, along with CRA. With the use of the same timeline as in Fig. 1, mice were challenged with CRA by intranasal (Day 14) and intratrachea delivery (Days 19 and 21), and 24 h after final allergen exposure, theresponses were assessed. The RSV-infected DCs induced a significant increase in AHR (A) compared with results with no DC + allergen or (continued) uninfected DC + allergen. When lung draining LN cells were restimulated in vitro with allergen at the end of the in vivo responses, changes in secreted cytokines were observed in the RSV-DCtreated mice compared with the other groups, as assessed in 48-h culture supernatants by Bio-Plex analysis (B). There was also a significant increase in mucus-associated genes in whole lung qPCR (C). Data represent mean \pm sE from five mice/group. *P < 0.05.

time-course analysis (24-72 h postinfection) of RSV-infected DCs in vitro and found no virus propagation from DCs by plaque analysis at any time-point based on continuous decreasing levels of plaqued virus (0.04% of input at 24 h, 0.004% by 48 h, and none by 72 h; Supplemental Fig. 2A). However, when we examined RSV protein mRNA expression in lungs of mice at 2 days post-DC transfers, we found some evidence of RSV infection based on qPCR results, although at a barely detectable level compared with a live virus infection (Supplemental Fig. 2B). The data in Fig. 2 demonstrate that RSV-infected DC leads to an exacerbation of the allergic response with a significantly increased AHR response compared with controls (Fig. 2A). Additionally, IL-13, IL-17, and IFN were increased significantly, and mucus gene expression was also increased compared with controls (Fig. 2B and C). Interestingly, it was only in the presence of RSV-infected DCs that the allergenrechallenged LN cells produced IL-17 (Fig. 2B). Moreover, simply delivering additional DCs into the airway at the time of allergen sensitization enhanced the Th2 cytokine sensitization; however, the activation of those DCs with RSV prior to instillation induced an even more intense and broader profile response (Fig. 2B). Thus, the DCs enhance the pathologic response, not only by promoting allergen sensitization but also by inducing potentially pathogenic IL-17. Characterization of the in vitro-infected DC demonstrated that RSV up-regulated CD80, CD86, and CD40 (Supplemental Fig. 2C) but did not induce apoptotic or necrotic cell death (Supplemental Fig. 2D).

To further address whether the recruitment and accumulation of mDC subsets are necessary for the development of the RSV-induced enhanced allergic responses described above, CCR6-/- mice were used. Studies from several laboratories, including our own, have demonstrated that CCR6-/- animals have a defect in mDC accumulation, but not T cells, within the lungs of RSV-infected or allergen-sensitized animals [29-35]. The infection of WT and CCR6-/- mice with RSV, along with allergen sensitization, was repeated as described above in Fig. 1A. As published previously, we also observed decreases in CD11b/CD11c+ DC accumulation but no difference in T cell populations that accumulated in the sensitized and challenged mice (data not shown). Mice rechallenged at Day 21 demonstrated a striking reduction of RSV-induced exacerbation of AHR in the CCR6-/- mice compared with WT control mice (Fig. 3A). Also, the CCR6-/- animals histologically displayed a significant reduction in overall inflammation and reduction in mucus production, and this was supported by quantification of mucus gene expression (Fig. 3B). Finally, when cytokine responses were examined, there was a decrease in the expression of Th2 cytokines in the RSV-exacerbated responses (Fig. 3C). However, whereas IL-17 was up-regulated by the RSV exposure, it was not reduced in the CCR6-/- mice (data not shown).

RSV infection predisposes animals to more severe pulmonary responses via DC activation and LN localization

Previous data have indicated that the severity of RSV-induced pathology can be associated with the presence of defined DC subsets recruited to the lung [14, 36–39]. Specifically, local

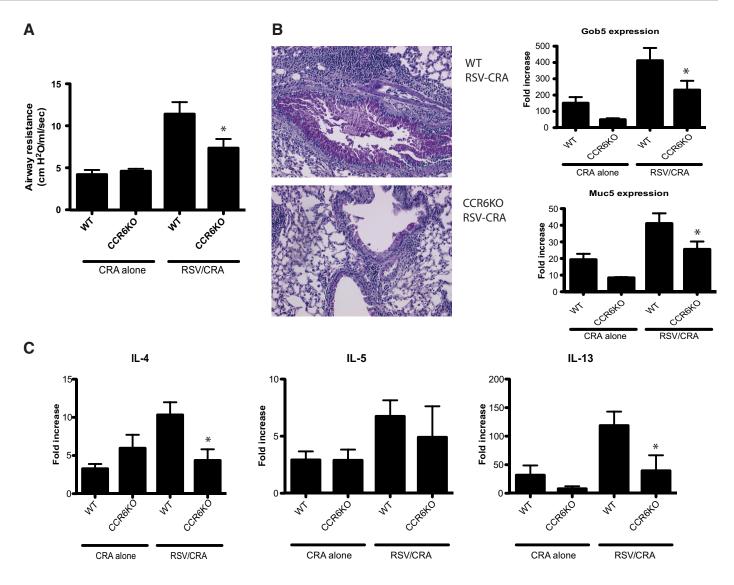


Figure 3. Attenuation of RSV exacerbated allergen sensitization in CCR6-/- mice. With the use of the sensitization protocol established in Fig. 1, WT or CCR6-/- [knockout (KO)] was sensitized to CRA, with or without RSV infection. Mice were challenged with CRA intranasally (Day 14) and intratrachelly (Days 19 and 21), and the responses were assessed. Twenty-four hours after final allergen exposure, mice were examined for methacholine-induced AHR (A). Differences in mucus hypersecretion in RSV-enhanced allergic disease were depicted by examining histology and mucus-associated genes muc5ac and gob5 by qPCR of lung mRNA (B). There was also a significant increase in Th2 genes, IL-4 and IL-13, in lung qPCR (C). Data represent mean \pm sE from five mice/group. *P < 0.05.

mucosal (lung) mDCs were shown to be responsible for skewing the immune responses toward a modified Th cell response that becomes more pathogenic [15, 17–19, 40]. Thus, we examined whether RSV-infected DCs also skewed a RSV infection challenge, as observed above with allergen. To examine this aspect specifically, BMDCs (CD11c/CD11b+) were cultured overnight, with or without RSV (MOI of 1), in vitro to allow infection, subsequently washed free of excess virus, and instilled into the trachea of naïve mice (2×10^5 DC/mouse). After 7 days, the mice were infected with RSV, and the immune response was assayed to determine if the infected BMDCs altered the immune environment. The data in **Fig. 4A** demonstrate that those animals that received RSV-infected BMDCs had an enhanced IL-4, IL-5, and IL-13 Th2-induced immune response, whereas IL-17 and IFN- γ production were down-regulated. This was accompanied by increases in mucus overexpression and airway pathology (Fig. 4B and C). Our studies also observed a decrease in RSV-specific mRNA, as might be expected with a local sensitization (data not shown) [41]. Thus, this response suggests that exogenously added DCs alter immune responses to the virus and resulted in increased pathology via Th2 cytokine overproduction in agreement with other studies [42–45].

We next determined whether RSV infection enhances migration of DCs to the LN. The data in **Fig. 5A** depict flow cytometry of the draining LNs from the instilled mice at different time-points, indicating that RSV-infected DCs traffic better to the LNs than do uninfected DCs. Thus, RSV does activate DCs

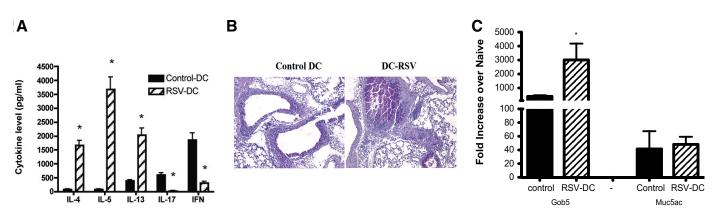


Figure 4. Airway sensitization of mice with RSV-infected DCs elicited a Th2-skewed immune response upon a RSV infection challenge. RSV-infected BMDC (2×10^5 /mouse) were instilled into the airways of naïve mice and infected with live RSV (1×10^5 PFU/mouse), 7 days after BMDC instillation. Supernatants from RSV-restimulated lung draining LN cells were analyzed for cytokines by Bio-Plex (A). Lungs were taken 8 days after the additional RSV infection (15 days later, the DC-RSV instillation) and were stained with PAS (B) and assessed by qPCR for mucus genes (C). Data represent mean \pm sE from five mice/group. *P < 0.05.

and enhances their migration into the draining LNs of mice, potentially facilitating sensitization of the immune response.

Our next series of studies was set up to determine whether the trafficking of the RSV-infected DCs to the LN was required for the altered responses, as previous studies have demonstrated that lung as well as LN resident DCs are involved in RSV-specific CD4 and CD8 T cell activation [20]. To address this question, studies used a similar system as described above with BMDCs from WT or CCR7-/- mice. To verify that the CCR7-/- DCs were impaired in their ability to traffic to draining LNs, as reported [46], the nodes were harvested at 24 h and 48 h postinstillation of CFSE-labeled, RSV-infected DCs. The data in Fig. 5B confirmed that DCs from CCR7-/animals do not traffic efficiently to the LNs after RSV infection. Seven days after the instillation of RSV-infected DC into the lungs of naïve mice, the animals were infected with RSV. Following 8 days of infection, the draining LNs were harvested, and the isolated cells rechallenged in vitro with a RSV infection. The resulting response indicated that when CCR7-/- DCs were used for airway sensitization, there was a significant reduction of Th2 cytokines with no change in IL-17 expression (Fig. 5C). Thus, the exacerbated immune response appears to be, at least partially, dependent on LN accumulation of the instilled, RSV-infected DCs and also enhances the pathogenic response to subsequent challenges. These responses should help identify the locational requirements for altering the immune responses during respiratory viral infections that may enhance allergen responses.

RSV-infected DC enhancement of allergic responses does not need to be simultaneous

We next examined whether the RSV-infected DCs have a longterm effect on allergen sensitization. Isolated DCs were infected with RSV and instilled into the airway as above. After 7 days of sensitization, the animals were given an allergen sensitization into the airway via intranasal delivery. Following an additional 6 days, the animals were again challenged twice with intratracheal allergen, 48 h apart. This suboptimal aller-

gen-sensitization process led to increases in AHR responses (Fig. 6A) and increases in Th2 cytokines IL-4, IL-5, and IL-13 (Fig. 6B), with no allergen-specific increase in IL-17 detected (data not shown). Histologic studies of PAS-stained lungs showed that although the mice receiving DCs alone displayed significant inflammatory cell infiltration, a significantly more mucogenic response was observed in animals given RSV-infected DCs prior to the allergen (Fig. 6C). Furthermore, when we waited for 21 days after RSV-infected DC instillation, prior to the allergen sensitization, the Th2 cytokine response continued to be enhanced by the RSV-DC transfer, with increased pathology characterized by mucus overproduction (Supplemental Fig. 3). Thus, these data indicate that RSV-infected DCs accelerated the allergen-sensitization process and caused Th2 cytokine skewing, leading to more severe disease, even when given up to 21 days prior to allergen exposure.

DISCUSSION

In these studies, we have begun to address several important mechanistic questions, including: are there detrimental effects of an antiviral immune response within the pulmonary environment that lead to a more pathogenic response to noninfectious (allergen) stimuli? Can innate immune cells (DCs) dictate the nature of a subsequent immune response? Although the present studies have only begun to address these issues, the data do correlate with clinical observations, suggesting that early viral infections can lead to long-term effects that alter subsequent immune responses [26, 27, 47, 48]. The role of RSV for development of childhood pulmonary disease has been epidemiologically controversial [9, 25, 47, 49-54], and several reports have begun to address the role of RSV in model allergen systems [55-61]. In fact, a recent publication has indicated that RSV can break an induced, antigen-specific tolerance response by altering regulatory T cell function [62]. Although RSV itself appears to promote a dysregulated innate immune response via specific NS proteins (NS1 and NS2) [63-66], it may be that the virus infection is a stimulus for en-

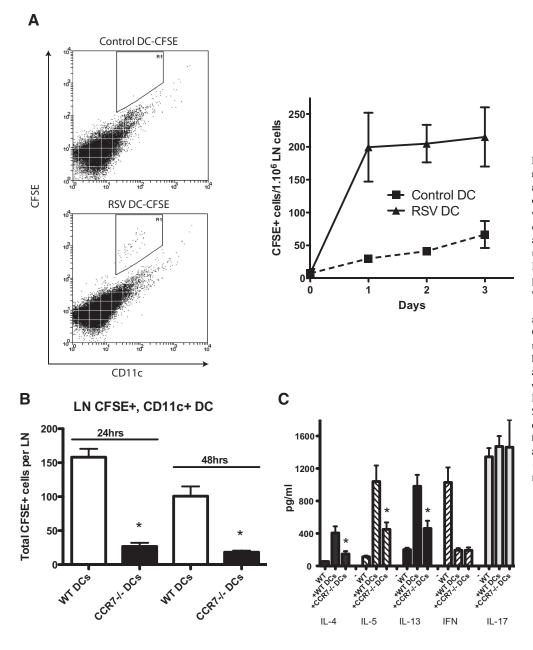
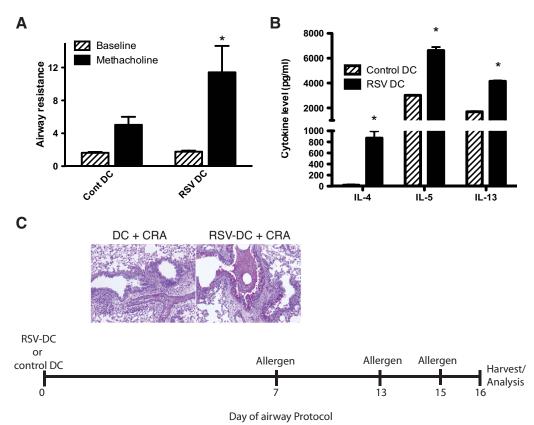


Figure 5. BMDC from CCR6-/mice do not migrate to draining LNs and fail to sensitize mice for an exacerbated RSV infection response. BMDCs were cultured overnight, with or without RSV, and stained with CFSE. After an injection of DCs into mice by intratrachea injection, lung draining LNs were isolated at different timepoints, and single-cell dispersions were used for flow cytometry analysis (A). Left panel is flow cytometry plot at 24 h time-point. R1 represents CD11c⁺ CFSE-positive cells. Similar results were observed at 48-h and 72-h later time-points (right panel). WT and CCR7-/- BMDCs were infected with RSV and were stained with CFSE. Lung draining LNs were isolated after 24 h and 48 h and performed flow cytometric analysis (B). Supernatants from RSV-restimulated LN cells were analyzed for cytokines by Bio-Plex (C). Data represent mean \pm sE from five mice/group. *P < 0.05.

hancing sensitization and leads to the altered immune responses. This was manifested in our studies using the DC sensitization model, where the active infection was replaced with RSV-infected DCs that propagate undetectable levels of active virions. In contrast, influenza infection did not induce a similar state of exacerbated disease and in fact, inhibited the skewing of allergen-specific responses. This latter change may be associated with differences in key mediators that are induced by the two viruses, such as type I IFN, which is highly induce by influenza [67] but inhibited by RSV NS1 and NS2 proteins [63, 65, 68]. With the use of DCs from CCR7-/- mice, studies established that migration into the LN was one necessary step for the altered immune response to a subsequent RSV infection. In patient populations, especially immune "inexperienced" infants, early viral infection may aid in the improper sensitization of individuals to environmental allergens. This effect, along with a genetic predisposition to an altered pulmonary environment, could lead to an enhanced pathologic response within the lung. The results, showing that the infected DC-induced immune deviation lasted 1 week and as long as 21 days, suggest that these cells are primarily involved in the process, either directly via APC function or indirectly by mediator release, such as IL-10 [69, 70]. Although the use of BMDCs is not entirely physiologic and the relevance to lung DC unclear, the studies help to identify a potential cellular mechanism of how RSV can alter an immune environment. Future studies will be needed to resolve these issues and to test exactly how long this effect persists.

One of the primary implications of our data is the relative lack of specificity of the altered pulmonary immune response

Figure 6. RSV-infected BMDCinduced allergen exacerbation is long-lived and does not need to be simultaneous. The BMDCs were infected with RSV (MOI=1.0) and instilled into naïve Balb/c mice. After 1 week, the mice were exposed to CRA (1 μ g/mouse) by intranasal instillation with repeated instillations, 6 and 8 days later (Days 13 and 15 of the protocol). Twenty-four hours after final allergen exposure (Day 16), the animals were assessed for AHR (A) by methacholine challenge. Cont, Control. Draining LNs were prepared to single-cell suspension and restimulated with allergen (2 μ g/ ml) and 48-h culture supernatant assessed for cytokines by Bio-Plex analysis (B). Formalin-preserved lung tissue (left lobe) was assessed for mucus and goblet cell hypertrophy by PAS staining of histology sections (C). Data represent mean \pm sE from five mice/group. *P < 0.05.



that extends its influence on a noninfectious allergen response. This raises the possibility that specific viral infections preferentially alter the immune environment as a result of a modified immune response. As mentioned above, this may be a result of specific viral proteins, such as the NS proteins of RSV that can inhibit key innate cytokines, and/or the entry of RSV via a nonendosomic route [71]. This latter concept is intriguing, as a primary mechanism for type I IFN production during RSV infection in epithelial cells is via the cytoplasmic sensor RIG-I that recognizes dsRNA [72, 73]. Recent studies have shown that the NS2 protein of RSV antagonizes IFN-B production induced by RIG-I [74]. Other studies have clearly demonstrated a skewed pathogenic environment in the absence of STAT1, an essential downstream signaling molecule for type I IFN expression [75-78]. Thus, DC activation and mediator release are affected directly by RSV infection. Subsequently, TLRs found in the endosome are also activated and contribute to the innate immune response during RSV infection [79-81]. Other studies have suggested that RSV G protein can inhibit TLR-induced activation [82]. Interestingly, when the RIG-I pathway and MyD88-dependent TLR pathways were blocked in mitochondrial antiviral-signaling protein/ MyD88 double-knockout mice, no alteration in cytotoxic T cell development and viral clearance was observed [83]. Likewise, the use of any one of several knockout mice deficient in TLR or MyD88 signaling did not alter viral clearance; however, significantly altered immune responses and increased pathogenesis during RSV infection were observed [79, 80, 84-86]. Thus, there may be multiple mechanisms for immune response alteration during RSV infection that can influence the direction and/or quality of the immune responses and therefore, the pathologic outcome.

The cytokine profile induced with a primary RSV-infected DC sensitization into the airway was a skewed Th2 response with decreases in IFN and IL-17 upon a RSV infection. A different profile was recognized when mice were infected with RSV coincident with allergen sensitization with an allergenspecific increase in Th2 and Th17 cytokines. Although one could explain these responses by the ability of RSV to cause enhanced DC migration to the LNs (as demonstrated with the CCR7 - / - DC studies), the ability to achieve enhanced allergen-induced Th2 responses, 1-3 weeks after RSV-DC instillation into the airway, is more difficult to explain. Previous studies have clearly suggested that pulmonary mDC populations have a preferential ability to skew the local immune responses toward a Th2-type immune environment [17-19, 87]. However, as described in a recent study by Guerrero-Plata et al. [88], pDCs also appear to be altered upon RSV infection. This latter effect may be important, as pDCs have a pivotal role in establishing an appropriate anti-RSV response and are necessary for maintaining a toleragenic immune environment against locally administered antigens/allergens [36, 38, 39]. Recent evidence has suggested that the NS proteins from RSV also suppress the maturation of DCs, including costimulatory proteins CD80, CD83, and CD86, and are associated with previously described reductions in type I IFN production [16]. Thus, the altered immune environment that was characterized by elevated Th cytokine production may have been caused by

altered DC maturation. Although our studies have focused on using a BMDC that best resembles a monocyte-derived DC subset (CD11b+/CD11c+), there are other important subsets in the lung, especially the CD103+ subset that appears to be relevant in RSV infection [20]. The CD103+ subset is migratory and has been suggested to better promote an antiviral Th1type cytotoxic T cell response as a result of its ability to crosspresent antigens [20, 89-93]. In contrast, a more recent study indicated that pulmonary CD103+ DCs were better able than CD11b+ DCs at promoting Th2 cytokine responses when an inert antigen (OVA) was used [94]. Thus, offsetting the balance of the subset ratios by adding a CD11b+/CD11c+ DCs may represent a highly inflamed environment that promotes the recruitment of more "inflammatory"-type DCs that migrate into the lung and initiate pathologic responses. In our studies, when we examined the CFSE trafficking of infected DCs into the LN, we had no evidence that the CD103+ cells were involved in the process, but we cannot rule out their contribution to the process. However, the use of CCR6-/- mice, which have altered inflammatory CD11b/CD11c DC accumulation after RSV and allergen exposures [29, 34, 35], further supports the concept that altering DC subset ratios can affect the outcome of the pulmonary response. Interestingly, our studies observed no alteration in IL-17 production in CCR6-/- or CCR7-/- studies in the presence of RSV, but we have not pursued this further at this time.

The data presented in these studies suggest that alteration of the pulmonary immune environment by RSV infection may have an underlying effect on the progression of disease pathology. Although several inter-related mechanisms may be induced, these studies suggest that RSV infections may predispose the host to an altered and modified disease phenotype through activation of DC subsets that may allow immune responses to allergens to be altered. Future studies will explore the nature of the RSV-mediated DC alteration.

AUTHORSHIP

S.J. contributed to design, writing, and execution. J.S. contributed to the design and execution. L.E.K. contributed to the execution of some of the experiments. N.W.L. contributed to the writing, execution, and design of the studies.

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