The changes of expressive levels of IL-17A, STAT3, and RORγt in different invasive pulmonary aspergillosis mice

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Background: T helper 17 (Th17) lymphocytes play an important role in Aspergillus adaptive immune response against Aspergillus fumigatus, but there is little attention focused on the different types of immunosuppressive models in which invasive pulmonary aspergillosis (IPA) develops. In addition, the expression levels of signal transducer and activator of transcription 3 (STAT3)/retinoic acid-related orphan nuclear receptor gamma (RORγt)/interleukin (IL)-17A signaling pathway, which is involved in the regulation of Th17 cells, as well as whether there are differences between two types of IPA mice models, remain unknown.

Materials and methods: Six to eight weeks old female BALB/c mice were treated with cortisone acetate or cyclophosphamide to establish the immunosuppressive mice models, and then, A. fumigatus inoculum was injected to form the IPA groups and sterile saline was injected to form the control groups. Flow cytometry was performed to measure the proportion of Th17 cells in CD4+ T cells in the peripheral blood, spleen, and lung of the mice. The expression of IL-17A, RORγt, and STAT3 mRNA was detected by real-time polymerase chain reaction. Concentrations of IL-6 in the plasma and bronchoalveolar lavage fluid were measured by enzyme-linked immunosorbent assay.

Results: The proportion of Th17 in the peripheral blood and lung tissue in neutropenic IPA mice showed a more significant increase than in non-neutropenic IPA mice (P<0.01). The IL-6 protein also showed the same trend in plasma and bronchoalveolar lavage fluid (P<0.01). Compared with the control groups, the expression of IL-17A at mRNA level in the lung was significantly increased, while RORγt/STAT3 mRNA was significantly decreased in the IPA groups (P<0.01).

Conclusion: The expression of RORγt and STAT3 mRNA in the lung tissue in both groups was significantly decreased. IL-17 may play a negative role in the defense against Aspergillus through uprating IL-6.

Keywords: non-neutropenic invasive pulmonary aspergillosis, neutropenic invasive pulmonary aspergillosis, Th17, IL-17A, RORγt, STAT3

Introduction

Aspergillus fumigatus is the material cause of invasive pulmonary aspergillosis (IPA), which is related to a high fatality rate even with common antifungal therapy. Patients with prolonged neutropenia, allogeneic hematopoietic stem cell transplant, solid organ transplant, inherited or acquired immunodeficiency, or corticosteroid use are known to have a high risk of IPA.1 However, recently, numerous studies found that patients with severe pulmonary dysfunction or COPD are also susceptible to IPA in the absence of neutropenia.2 Monocytes, macrophages, and neutrophils constitute the innate immunity defense against A. fumigatus. It is well known that T helper 17
RORγt is one of the most important transcription factors for Th17 differentiation. The activation of RORγt is regulated by a series of regulatory factors under normal conditions. It was confirmed that STAT3 is one of the most important regulators of Th17 cells as well as whether there are differences between two types of IP A mice models, are not yet clear. Therefore, in this study, we attempt to establish two patterns of IP A mice models treated with cortisone acetate and cyclophosphamide, respectively, to investigate the changes in STAT3, RORγt, and IL-17A expression in pulmonary, as well as the IL-6 protein concentration in blood and bronchoalveolar lavage fluid (BALF).

Materials and methods
Fungal isolates
A clinical A. fumigatus isolate from a pulmonary aspergillosis patient treated at the First Affiliated Hospital of Wenzhou Medical University (Wenzhou, China), which was provided by the microbiology laboratory was used in our study. A. fumigatus was grown on Sabouraud dextrose agar plates for 5 days at 37°C to prepare the inoculum, and then, conidia were collected by flushing the plates with 10 mL sterile saline within 0.1% Tween 80 and then filtering through four layers of sterile gauze. After the inoculum was adjusted to the required concentration (1×10^7/mL) using a hemocytometer, the conidial suspension was stored at 4°C.

Source of mice
Six to eight weeks old female BALB/c mice weighing 18–22 g (Slaker lab, Shanghai, China) were used in this study. All procedures involving mice were approved by the institutional animal use and care committee of Wenzhou Medical University and infection, playing a positive role against A. fumigatus. However, several papers reported that Th17 cells do not have a role in the defense against Aspergillus or may even have a damaging effect. Despite this, Th17 lymphocytes do play a role in the adaptive immune response against A. fumigatus.

The differentiation and immune function of Th17 cells are regulated by many molecules, of which interleukin (IL)-6 and transforming growth factor-β play a crucial role through the JAK/signal transducer and activator of transcription 3 (STAT3) signaling pathway. Retinoic acid-related orphan nuclear receptor γt (RORγt) is an essential transcription factor for Th17 differentiation. The activation of RORγt is regulated by a series of regulatory factors under normal conditions. It was confirmed that STAT3 is one of the most important RORγt regulatory factors.

Experimental mice models have been developed for research of the role of innate immunity in IPA. However, there is little attention focused on the different immunosuppressive models in which IPA occurs. Moreover, the expression levels of STAT3/RORγt/IL-17 signaling pathway in the IPA mice model, as well as whether there are differences between two types of IPA mice models, are not yet clear. Therefore, in this study, we attempt to establish two patterns of IPA mice models treated with cortisone acetate and cyclophosphamide, respectively, to investigate the changes in STAT3, RORγt, and IL-17A expression in pulmonary, as well as the IL-6 protein concentration in blood and bronchoalveolar lavage fluid (BALF).

**Mouse model of IPA**
After 1 week of adaptive feeding, the mice were randomly divided into five groups of 12 mice as experiment or control groups. Mice in group A were treated with cortisone acetate (Aladdin, Shanghai, China) at 500 mg/kg subcutaneously every other day, starting on Day 4 relative to infection and finishing on Day +3, and served as the non-neutropenic IPA model. Mice in group B were treated with cyclophosphamide (Baxter Oncology GmbH, Halle, Germany) administered intraperitoneally twice at 150 mg/kg on Day 4 and Day –1 and served as the neutropenic IPA model. Mice in groups C and D were only treated with cortisone acetate or A. fumigatus at the same dose as in group A, respectively. Mice in group E were saline-treated immunocompetent mice without immunosuppression and infection. All animals were anesthetized by intraperitoneal injection of pentobarbital (Aladdin) on Day 0 and then intratracheally injected with 30 µL conidial suspension (groups A, B, and D) or sterile saline (groups C and E). Then, 250 mg of tetracycline (Aladdin) was dissolved in 1 L of sterile water to prevent bacterial infections. In the survival studies, six mice from each group were monitored for survival.

After 72 hours of infection, the mice were sacrificed after they were anasthetized and their peripheral blood, BALF, lung, and spleen were harvested. The IPA model was identified through the lung tissue histopathologic analysis and fungal culture. The sections were stained with HE for tissue examination and periodic acid-Schiff and periodic Schiff-methenamine silver stain for fungus detection.

**Peripheral blood and BALF**
Plasma was collected and stored at −80°C after centrifuging the anticoagulant peripheral blood at 3,000 rpm for 5 minutes, and the setting blood cells were used for flow cytometry after hemolysis with Lysing Buffer (BD Biosciences, San Jose, CA, USA). The chest of mice was exposed to ligate the left main bronchus, and then, its right lung was washed out with 0.5 mL cold PBS twice, aspirating three times each. Then, the supernatant was collected and stored at −80°C after centrifuging at 3,000 rpm for 10 minutes at 4°C.

**Flow cytometry**
The lungs and spleens of mice were carefully removed, and a single cell suspension was generated using a 200-mesh sieve. Then the lymphocytes were separated with a mouse lymphocyte separation solution (TBD science, Tianjin, China). Cells from lungs, spleens, and blood were all incubated at
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37°C in RM1640 (Thermo Fisher Scientific, Waltham, MA, USA) containing Phorbol 12-myristate 13-acetate/ionomycin mixture and Brefeldin A/monensin mixture (Multisciences, Hangzhou, China). The stimulated cells were washed and incubated with fluorescein isothiocyanate Rat Anti-Mouse CD4 (BD Biosciences). For intracellular staining, Cytofix/ Cytoperm Soln Kit (BD Biosciences) was used and the cells were stained with PE Rat Anti-Mouse IL-17A (BD Biosciences), after washing with Wash/Perm solution. Isotype controls (BD Biosciences) were used to determine gates for positive antibody responses. Th17 proportion was analyzed using a flow cytometer (BD Biosciences).

Quantitative real-time polymerase chain reaction (qPCR)
The IL-17A, RORγt, and STAT3 mRNA expression in the lung tissue and pulmonary A. fumigatus fungal burden were detected by qPCR. The lung tissues of mice were homogenized in liquid nitrogen, and total RNA was extracted by TRizol Reagent (Thermo Fisher Scientific) following the manufacturer’s instructions. Samples with an OD260/280 ratio of 1.8:2.0 were used to generate cDNA by reverse transcription.

A total of 3 µg of RNA was reverse-transcribed to cDNA with RevertAid First Strand cDNA Synthesis Kit (Thermo Fisher Scientific). qPCR for IL-17A, RORγt, and STAT3 mRNA was performed on Bio-Rad CFX 96 (Bio-Rad Laboratories Inc., Hercules, CA, USA) using the ChamQ Universal SYBR qPCR Master Mix (Vazyme, Nanjing, China) as the detecting probe. The products were also compared with GAPDH as the loading control. The primer sequences used are shown in Table 1.

Cytokine quantification by enzyme-linked immunosorbent assay
Cytokine production in plasma and BALF was measured by enzyme-linked immunosorbent assay according to the manufacturer’s directions (Multisciences).

Statistical analysis
All measurement data are shown as mean and standard errors of the mean in this study, and statistical analysis was performed using an unpaired t-test or one-way analysis of variance (GraphPad Prism Software). A P-value <0.05 was considered significant.

Results
Survival
The survival curves (Figure 1) were established in various models using the same concentration of inoculum (10⁷ conidia/mouse). Infected immunocompetent mice all survived, whereas 100% of non-neutropenic IPA mice died between Days +3 and +5. Infection of neutropenic mice resulted in 100% mortality between Days +2 and +4.

Lung injury and fungal culture
The differences in pulmonary injury were assessed by histological examination performed on lung samples collected 72

Figure 1 Survival of immunocompetent and immunosuppressed mice after intratracheal infection with conidia of Aspergillus fumigatus. Notes: Six animals per group were infected with conidia in four independent experiments. Corticosteroid-treated without infection mice (C) and immunocompetent infected mice (D) all survived, whereas 100% of non-neutropenic IPA mice (A) died between Days +3 and +5. Infection of neutropenic mice (B) resulted in 100% mortality between Days +2 and +4. There was no significant difference between the survival of group A and group B mice (P=0.18).

Abbreviation: IPA, invasive pulmonary aspergillosis.

Table 1 Sequences of primers in real-time PCR

<table>
<thead>
<tr>
<th>Primer</th>
<th>Forward (5′–3′)</th>
<th>Reverse (5′–3′)</th>
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<tr>
<td>GAPDH</td>
<td>GGTGTGTCTTCCTCCGACCTTCA</td>
<td>TGGTCCAGGGTTTCTTACC</td>
</tr>
<tr>
<td>IL-17A</td>
<td>TGCTCAATGCGGAGGGAAAAGC</td>
<td>ACCCTGAAAGTGAGGGGCA</td>
</tr>
<tr>
<td>RORγt</td>
<td>CAGTCCCCCAGGATGACCCATA</td>
<td>CTGCGCTGAAAGGTCCCTCAA</td>
</tr>
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<td>STAT3</td>
<td>GTGGAGAACCTCCGAGGAGA</td>
<td>GCCAGCTCAGTCGAAATGCT</td>
</tr>
<tr>
<td>AF</td>
<td>CGCGAATGATACCTACCTAGGG</td>
<td>TCCGTCCAGGGAGG</td>
</tr>
</tbody>
</table>

Abbreviations: AF, Aspergillus fumigatus; PCR, polymerase chain reaction; RORγt, retinoic acid receptor-related orphan receptor gamma; STAT3, signal transducer and activator of transcription 3; IL, interleukin.
hours after infection of mice in different groups (Figure 2). The lungs of non-neutropenic IPA mice showed hemorrhagic necrosis pneumonia with neutrophil infiltration. Destruction of bronchi and alveoli was also evident (Figure 2Aa-1). A small amount of hyphae of *A. fumigatus* was observed para-tracheal (Figure 2Ab-1, Ac-1). In contrast, for neutropenic mice, we also observed unsystematic pneumonia without inflammatory exudate including polymorphonuclear neutrophil or other cells in the alveoli with hemorrhagic necrosis (Figure 2Ba-2). Alveoli and vessels were invaded by a mass of hyphae of *A. fumigatus* (Figure 2Bb-2, Bc-2).

Mice in both IPA groups showed visible *A. fumigatus* in the lung tissue culture, while in those in immunocompetent infected group, it could be rarely seen.

### Th17 proportion in peripheral blood, spleen, and lung

The Th17 cells (CD4+, IL-17+) proportion in the peripheral blood, spleen, and lung tissue of both patterns of IPA mice was significantly increased compared with that in saline-treated immunocompetent mice. There was no difference between different control groups. The increase in peripheral blood and lungs of neutropenic IPA mice was more significant than that in non-neutropenic ones (*P*<0.01; Figure 3).

#### qPCR analysis of the expression of IL-17, RORγt, STAT3 mRNA, and pulmonary fungal burden in the lung tissue

The expression level of these genes was given as the ratio with respect to saline-treated immunocompetent mice (Figure 4). Levels of GAPDH were analyzed as the control and showed no statistical differences. Overall, there was a generalized reduction in RORγt expression in immunosuppressive mice vs immunocompetent mice, and more significant reduction was seen in IPA groups. Both IPA mice showed a significant reduction in STAT3/RORγt expression and increases in IL-17A expression in response to infection, compared with saline controls (*P*<0.01). There was no significant difference between the two patterns of IPA mice, as well as the three control groups.

![Figure 2](https://example.com/figure2.png)

**Figure 2** Histopathological analysis (×400) in the lung tissue from different groups.

**Notes:** The lungs of non-neutropenic IPA mice (A) showed large foci of pneumonia with the destruction of bronchi and alveoli. Hemorrhagic necrosis with neutrophil infiltration was also observed using the HE stain (a-1). Paratracheal hyphae of *Aspergillus fumigatus* was observed in small numbers with PAS (b-1) and PASM (c-1) stains. In contrast, for neutropenic IPA mice (B), we also observed diffuse pneumonia with edema and congestion within the alveoli, but with no inflammatory exudate involving PMN or other cells (a-2). Of note was that alveoli and parenchyma were invaded by numerous hyphae of *A. fumigatus* (b-2, c-2). There was no significant lesion seen in the lung tissue of corticosteroid-treated without infection mice (C), immunocompetent infected mice (D), and saline immunocompetent mice (E). The lungs of group (C-E) with HE stain (a-3-a-5), PAS stain (b-3-b-5) and PASM stain (c-3-c-5), respectively.

**Abbreviations:** IPA, invasive pulmonary aspergillosis; PAS, periodic acid-Schiff; PASM, periodic Schiff-methenamine; PMN, polymorphonuclear neutrophil.
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The expression of IL-17A, STAT3 and RORγt in IPA mice

We used a real-time PCR that targets a previously described 67 bp segment of a 28S ribosomal RNA coding DNA to detect *Aspergillus* DNA.\(^{13}\) It can be seen that the *A. fumigatus* fungal burden in both IPA mice were significantly higher than in the immunocompetent infected mice (*P* < 0.01). Neutropenic IPA mice showed higher fungal burden compared with the non-neutropenic IPA mice (Figure 5).

**IL-6 concentration in BALF and plasma**

Compared with the saline-treated immunocompetent mice, IL-6 concentrations in the plasma and BALF of neutropenic mice showed a more significant increase than in non-neutropenic mice (*P* < 0.01). Immunocompetent infected mice showed a slight increase of IL-6 in BALF and no obvious change in plasma (Figure 6).

**Discussion**

Th17 cells can specifically secrete IL-17 at high levels, which was reported by Harrington et al for the first time in 2005.\(^{14}\) The differentiation of naïve T cells is not only regulated by cytokines from the environment but also by their inherent procedure. As a novel type of T lymphocytes, Th17 cells have been shown to be regulated positively or negatively by many cytokines. However, the specific regula-
tion mechanism of their lineage differentiation has not been well understood.

In this study, we demonstrated that the STAT3/RORγt expression level was significantly decreased in immunosuppressive infected mice compared with that in saline immunocompetent mice. However, the proportions of Th17 cells and the expression level of IL-17 were significantly increased, which was consistent with the results of Armstrong et al.15

JAK/STAT pathway plays an important role in the differentiation of Th17 cells. Among the STATs, STAT3 is the most active protein which can be activated by numerous cytokines including growth hormone, IL-6 family cytokines, and so on. IL-6 had been shown activating the STAT3/RORγt signaling pathway, which can induce Th17 cells secreting IL-6, IL-17 and other cytokines.8 Previous studies confirmed that RORγt is an indispensable specific transcription factor for IL-17A synthesis in Th17 cells.9,16

The above results of our study indicated that STAT3/RORγt signaling pathway was obviously inhibited. On the contrary, the proportions of Th17 cells and the expression level of IL-17 were significantly increased.

Therefore, two aspects of this problem have to be discussed. One possibility is that the result may suggest that RORγt is not that essential a regulatory factor to Th17. Yang et al found that RORγt deficiency did not absolutely forbid Th17 cytokine expression. They reported that Th17 cytokine was also regulated by another related nuclear receptor RORα,
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but it is still unclear whether RORγt or RORα directly participates in the transcriptional regulation of IL-17A gene.17 Another possibility is that Th17 immunity is, indeed, inhibited in IPA mice. All the CD4+ T cells were inhibited in the immunosuppressed mice, resulting in reduction in the absolute number of Th17 cells, though its proportion increased. Furthermore, the high expression of IL-17A in the lung tissue is not exclusively produced by Th17 cells. Studies have shown that despite the fact that most of the recent literature describes IL-17 as a T cell-secreted cytokine, much of the IL-17 released during an inflammatory response is produced by innate immune cells.18

Moreover, IL-6 was significantly increased in the plasma and BALF of IPA mice in our study, which is in agreement with the study of Heldt et al based on clinical patients with IPA.19 IL-6, a proinflammatory cytokine, is not only a promoter of STAT3/RORγt signaling pathway but also a key downstream target of IL-17.20 Wang et al demonstrated that IL-17 can induce IL-6 production to promote tumor growth.21

Comparing the non-neutropenic IPA mice and the neutropenic IPA mice, the latter had more severe disease with the same inoculum, manifesting as more severe lung injury, higher pulmonary fungal burden, as well as higher IL-6 protein levels and Th17 proportion in the lung and blood. So, we considered that through uprating IL-6, IL-17 may play a negative role in the defense against Aspergillus. This is consistent with the results of a study showing that the cytokine IL-17A can promote the growth of A. fumigatus in vivo and in vitro.6

Unlike autoimmune diseases, researchers have not reached an agreement on the immune function of Th17 cells in defense of A. fumigatus at present, which may perform a two-way effect of promoting infection15 and resisting infection22 in different immune status. Further studies to clarify the role of Th17 cells in IPA hosts are necessary.

There were limitations in our study. We detected the STAT3/RORγt/IL-17A mRNA based on the entire lung tissue which contained various cells, such as epithelial cells, endothelial cells, lymphocytes, and many primary immune cells, such as macrophages. Detection of STAT3/RORγt signaling pathway expression and IL-17A mRNA by extracting total RNA from monocytes in the lung may be the better way of specifically evaluating Th17 functional changes in two types of IPA mice.

Conclusion

Our study demonstrated that the STAT3/RORγt expression level was significantly decreased in immunosuppressive infected mice compared with the saline immunocompetent mice. However, the proportions of Th17 cells and the expression level of IL-17 were significantly increased. We considered that IL-17 may play a negative role in the defense against Aspergillus through uprating IL-6.

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Disclosure
The authors report no conflicts of interest in this work.

References