RESEARCH ARTICLE

The Role of IL-23/Th17 Pathway in Patients with Primary Immune Thrombocytopenia

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Abstract

Background
Primary immune thrombocytopenia (ITP) is an autoimmune bleeding disorder with an unclear etiology. This study aims to investigate the role of IL-23/Th17 pathway in patients with ITP.

Method
The gene expressions of IL-17, IL-23 and their receptors in ITP patients and healthy controls were analyzed by quantitative real-time PCR. ELISA was used to test the IL-17 and IL-23 levels in plasma. Flow cytometry was used to detect the frequency of Th17 cells. The correlation between plasma IL-23 and IL-17 levels, Th17 cells, platelets were analyzed. The level of Th17-related cytokines was measured by ELISA following stimulation with IL-23. Subsequently, the IL-23 and IL-17 levels were measured in patients post-treatment.

Results
The PBMCs of ITP patients showed increased mRNA expression levels in each of the following: IL-23p19, IL-12p40, IL-23R, IL-12Rβ1, IL-17A, IL-17F, and RORC. In addition, elevated Th17 cells and plasma IL-17, IL-23 levels were also observed in these ITP patients. The correlation between plasma IL-23 and IL-17 levels, Th17 cells, platelets were analyzed. The level of Th17-related cytokines was measured by ELISA following stimulation with IL-23. Subsequently, the IL-23 and IL-17 levels were measured in patients post-treatment.

Conclusion
The IL-23/Th17 pathway may be involved in the pathogenesis of ITP through enhancement of the Th17 response. Moreover, our results suggest that the IL-23/Th17 pathway is a potential therapeutic target in future attempts of ITP treatment.
**Introduction**

Primary immune thrombocytopenia (ITP) is the most common hemorrhagic autoimmune disease, and is characterized by an isolated thrombocytopenia that is not accompanied by any other disorders that may lead to thrombocytopenia[1]. Recent studies have found that cellular immunity deficiency plays an important role in the pathogenesis of ITP, including B-cell activation, T-cell related disorders and antigen-presenting cell functional defect[2–4]. Significantly, it is well-recognized that an abnormal T-cell-mediated response is strongly correlated to the development and progress in ITP[5].

Recent studies have demonstrated that Th17, which is characterized for its production of IL-17, is elevated in ITP patients [6,7]. IL-17 belongs to the IL-17 cytokine family, which contains 6 different IL pro-inflammatory cytokines, from IL-17A to IL-17F. IL-17A is often represented as IL-17. The high degree of similarity between IL-17A and IL-17F is significant, and they also have a common biological function. Increased IL-17 expression has been observed in various autoimmune diseases, such as rheumatoid arthritis (RA)[8] and systemic lupus erythematosus (SLE)[9]. This evidence suggests that IL-17 may be associated with autoimmune diseases.

IL-23, which is mainly secreted by antigen-presenting cells, is a member of the IL-12 family, which includes IL-12, IL-27, and IL-35[10]. IL-23 is a heterodimeric cytokine, comprised a unique p19 subunit and p40 subunit, the latter of which is shared with IL-12. The receptor for IL-23 consists of IL-23R and IL-12Rβ1, the latter of which is also characteristic of IL-12. IL-23 is essential for Th17 differentiation, expansion, and survival by binding to its receptor, thereby activating the signaling pathway [11,12]. Many studies revealed that the IL-23/Th17 pathway is implicated in the pathophysiology of various autoimmune diseases, such as autoimmune arthritis[13], primary biliary cirrhosis[14], and inflammatory bowel disease[15]. But the important role of IL-23 in human ITP, especially in relation to Th17 cells remains unsettled.

In the present study, we used quantitative real-time PCR to investigate the gene expression of the subunits of IL-17, IL-23, and their receptors in ITP patients and healthy controls. Plasma levels of IL-17 and IL-23 were analyzed by ELISA. The frequency of Th17 cells were detected by flow cytometry. Moreover, we explore the function of IL-23 in Th17-associated cytokine production in ITP patients.

**Materials and Methods**

**Patients**

We enrolled 30 patients with confirmed acute ITP who were admitted to the Shanghai Changhai Hospital and Shanghai Tongji Hospital from June 2012 to January 2014. An ITP diagnosis was made according to a report from an international working group [16]. We excluded subjects who had the complications of diabetes, hypertension, cardiovascular diseases, pregnancy, active infection, or autoimmune diseases other than ITP [17]. We followed up on eleven newly diagnosed patients after effective treatment in order to analyze the dynamic change of IL-23. The criteria for assessing response to ITP treatments was according to a previous report [16].

As a control group, we also enrolled 30 healthy subjects (18 females and 12 males; age range: 19–62 years; median age: 36 years). All controls had normal platelet counts and did not receive any steroid therapy. There was no statistical difference between the control group and the ITP group in age and gender (P>0.05).

Our study was approved by the Ethics Committee of Shanghai Changhai Hospital and Shanghai Tongji Hospital (Shanghai, China). Written informed consent was obtained from all subjects.
PBMC isolation and cell culture

Peripheral whole blood was obtained from all subjects. PBMC were isolated using Ficoll-Hypaque density gradient centrifugation. For experiments in vitro, PBMCs from ITP patients were cultured at a concentration of $1 \times 10^6$ cells/mL in RPMI1640 medium (Sigma, R8758) supplemented with 10% fetal bovine serum, 100U/ml penicillin and 100mg/mL streptomycin (Invitrogen, USA) as mentioned previously[18], and stimulated with recombinant IL-23 (R&D Systems, 1290-IL-010) or PRMI1640. Culture supernatant was collected for cytokine analysis. For future flow cytometric analysis, PBMCs were cryopreserved in fetal bovine serum containing 10% dimethyl sulfoxide (DMSO), and stored in liquid nitrogen.

Total RNA extraction and cDNA synthesis

Total RNA was extracted using the RNeasy Mini Kit (Qiagen Inc, 74104) according to the manufacturer’s instruction. RNA yield and purity were determined spectrophotometrically at 260/280 nm and the ratios were in the range of 1.8–2.0. cDNA was synthesized using 1 μg total RNA with random primers and TaqMan Reverse Transcription Kit (ABI, N8080234), and stored at -20°C until use.

Quantitative real-time PCR

All PCR reactions were undertaking on ABI7500. Primers and TaqMan-MGB probes were designed for detecting IL-23p19, IL-12p35, IL-12p40, IL-23R, IL-12Rβ1, IL-12Rβ2, and 18s-RNA mRNA expression as described[14], and 18s-RNA was used as internal control. The mRNA expression for IL-17A, and IL-17F, RORC took place using SYBR Green method, and GAPDH was used as internal control. The primers used were as follows: IL-17A forward 5'-ATGACTCTCGG GAAGACCTCATTG-3'; reverse 5'-TTAGGCCACATGTTGGACAATCGGG-3'; IL-17F forward 5'-GTCACCTTGGGAACCCAACGCG-3'; reverse 5'-CTGACATGGTGGATGACAGGG-3'; RORC forward 5'- GTCCCAGATGCTGTCAAGT-3'; reverse 5'-TGGTTATCGTGGAAGGAC TCA-3'; reverse 5'- CCAG TAGAGGCAGGGATGAT-3'. All samples were tested in triplicate. The CT values of internal control and a target gene were determined. The relative expression for the target gene was given by $2^{-\Delta\Delta CT}$.

Flow Cytometric Analysis

Cryopreserved PBMCs were thawed at 37°C, washed twice with PBS, and stained with trypan blue to determine cell viability. PBMCs were adjusted to a concentration of $1 \times 10^6$/ml in RPMI1640 medium supplemented with 10% fetal bovine serum. The PBMCs were incubated for 4 hours at 37°C, 5% CO₂ in the presence of PMA/Ionomycin mixture (Liankebio, LK-CS1001) and BFA/Monensin mixture (Liankebio, LK-CS1002) according to the manufacturer’s instructions. After incubation, the cells were stained with FITC-conjugated anti-CD3 (BD, USA) and PE-Cy5-conjugated anti-CD8 (Biolegend, 301010) to delimitate CD4+ T cells. Then the cells were stained with PE-conjugated anti-IL-17A (Biolegend, 512306) for Th17 detection after fixation and permeabilization (BD IntraSure Kit, 4043524). Stained cells were tested on a FACS Calibur flow cytometer (BD, USA) and then analyzed using CellQuest software (BD Bioscience).

ELISA

IL-17 and IL-23 levels in plasma were detected using commercial enzyme-linked immunosorbent assay (ELISA) according to the manufacturer’s instructions. Concentrations of IL-6,
IL-17, and IL-21 in culture supernatant were determined using Bio-Plex Pro Human Th17 cytokine assay (BIO-RAD).

Statistical analysis
Statistical analysis was performed using GraphPad Prism 5.0 software (GraphPad Software Inc). Unpaired Student’s t test or Mann-Whitney U test was used, as appropriate, to compare results between groups. Correlations between variables were assessed by Pearson or Spearman correlation coefficients, as appropriate. A paired t-test was used to compare results before and after treatment. A p-value of <0.05 was considered significantly different.

Result
Characteristics of participants
Characteristics of the 30 ITP patients admitted in our study are shown in Table 1. The clinical information of 11 newly diagnosis ITP patients was shown in Table 2. Complete response was observed in 7, response in 4 of the 11 newly diagnosis patients. During the treatment, no bleeding or other obvious complications were observed.

Increased IL-23p19, IL-12p40, IL-23R, IL-12Rβ1, IL-17A, IL-17F, RORC mRNA expression in PBMCs from patients with primary immune thrombocytopenia
As shown in Fig. 1, compared to healthy controls, IL-23 p19 and p40 mRNA expression were both significantly increased in ITP patients, but not IL-12 p35 mRNA (Fig. 1A). Moreover, we analyzed the expression of their relative receptors. We found IL-23R and IL-12Rβ1 mRNA expression increased, but not IL-12Rβ2 (Fig. 1B). The expression of Th17-related RORC, IL-17A and IL-17F also significantly increased (Fig. 1C). There was no statistical difference in these measures between newly diagnosis group and re-activated group.

Increased plasma IL-23, IL-17 level and Th17 cells in patients with primary immune thrombocytopenia
Compared to healthy controls, the IL-23 and IL-17 level in plasma from ITP patients were both significantly increased (Fig. 2). We did not observed any statistical difference in plasma IL-23 and IL-17 levels between newly diagnosis group and re-activated group. In flow cytometry analysis, the percentage of Th17 cells in ITP patient was increased compared to that in healthy controls (Fig. 3).

Correlation between plasma IL-23 and IL-17 level, Th17 cells, platelets
As shown in Fig. 4, IL-23 level in ITP plasma were positively correlated with plasma IL-17 and Th17 cells, while negatively related with platelet counts.

Increased Th17 related cytokines by PBMC from ITP patients after IL-23 stimulation in vitro
In order to further validate the role of IL-23 on Th17 related cytokine production induced by PBMC, we incubated the PBMC from ITP patients with rhIL-23 or medium (as control). We observed that the IL-6, IL-17, and IL-21 levels in culture supernatant increased after rhIL-23 stimulation (Fig. 5). The Th17 related cytokines production in newly diagnosis and re-activated groups were not reached statistical differences.
Decreased plasma IL-23 level after effective treatment

We followed up eleven newly diagnosed ITP patients after admission to this study. We found that the plasma of these patients showed decreased levels of IL-23 and IL-17 after effective treatment (Fig. 6).

Discussion

In the present study, we found that the mRNA expression of IL-23 subunits p19 and p40, IL-23 receptors, IL-17 and RORC is increased in patients with ITP. We also reported that plasma IL-17, IL-23 levels and Th17 cells were increased in ITP patients. A positive correlation exists between plasma IL-23 and IL-17 and Th17 levels. To the best of our knowledge, this is the first

Table 1. ITP patient’s clinical characteristics.

| Patient No | Gender | Age/year | Course/month | Bleeding symptom | PLT/×10^9/on admission | Previous Treatment |
|------------|--------|----------|--------------|------------------|------------------------|--------------------|
| 1          | F      | 18       | 4            | PT, EC           | 19                     | —                  |
| 2          | F      | 24       | 13           | EP, EC           | 23                     | DXM                |
| 3          | M      | 43       | 36           | PT              | 14                     | DXM                |
| 4          | F      | 26       | 20           | PT, EC           | 8                      | Pred               |
| 5          | M      | 44       | 28           | GH              | 11                     | DXM                |
| 6          | M      | 38       | 14           | EC              | 16                     | DXM                |
| 7          | F      | 61       | 7            | PT              | 10                     | —                  |
| 8          | M      | 24       | 1            | PT, EC           | 9                      | —                  |
| 9          | F      | 33       | 2            | GH              | 7                      | —                  |
| 10         | M      | 41       | 7            | PT, EC           | 14                     | —                  |
| 11         | F      | 70       | 31           | EC, EP           | 11                     | Pred               |
| 12         | M      | 51       | 26           | EP              | 6                      | DXM                |
| 13         | M      | 32       | 36           | EP, EC           | 6                      | Pred               |
| 14         | M      | 20       | 7            | EP              | 12                     | —                  |
| 15         | F      | 34       | 6            | EP              | 11                     | —                  |
| 16         | F      | 50       | 29           | EC              | 4                      | DXM                |
| 17         | M      | 27       | 30           | PT              | 10                     | DXM                |
| 18         | F      | 39       | 8            | PT              | 22                     | —                  |
| 19         | F      | 28       | 41           | EC, PT           | 14                     | DXM                |
| 20         | F      | 22       | 22           | PT, GH           | 27                     | Pred               |
| 21         | M      | 25       | 5            | EC, PT           | 21                     | —                  |
| 22         | F      | 19       | 1            | GUH             | 11                     | —                  |
| 23         | F      | 26       | 31           | EP              | 7                      | DXM                |
| 24         | F      | 39       | 26           | PT, EP           | 12                     | DXM                |
| 25         | F      | 47       | 37           | PT              | 10                     | Pred               |
| 26         | F      | 41       | 21           | PT              | 21                     | DXM                |
| 27         | F      | 24       | 9            | GUH             | 16                     | —                  |
| 28         | M      | 36       | 48           | EP, PT           | 25                     | Pred               |
| 29         | M      | 29       | 22           | PT              | 7                      | DXM                |
| 30         | F      | 46       | 14           | PT              | 15                     | DXM                |

F: female; M: male; PT: petechiae; EC: ecchymoses; EP: epistaxis; GUH: genitourinary hemorrhage; GH: gingival hemorrhage; —: no prior treatment; PRED: prednisone; DXM: dexamethasone

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Table 2. Eleven newly diagnosis ITP patients clinical information.

| Patient No | Treatment | Dose(mg/day) | Period of treatment/day | PLT before treatment(×10^9/L) | PLT after treatment(×10^9/L) |
|------------|-----------|--------------|-------------------------|-------------------------------|-----------------------------|
| 1          | DXM       | 40           | 4                       | 19                            | 52                          |
| 7          | DXM       | 40           | 4                       | 10                            | 125                         |
| 8          | DXM       | 40           | 4                       | 9                             | 76                          |
| 9          | DXM       | 40           | 4                       | 7                             | 154                         |
| 10         | DXM       | 40           | 4                       | 14                            | 37                          |
| 14         | DXM       | 40           | 4                       | 12                            | 63                          |
| 15         | DXM       | 40           | 4                       | 11                            | 151                         |
| 18         | DXM       | 40           | 4                       | 22                            | 257                         |
| 21         | DXM       | 40           | 4                       | 21                            | 105                         |
| 22         | DXM       | 40           | 4                       | 11                            | 127                         |
| 27         | DXM       | 40           | 4                       | 16                            | 172                         |

DXM: dexamethasone

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Fig 1. IL-23/Th17 pathway related molecules mRNA relative expression. *P < 0.01.

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time that abnormal IL-23 expression and its correlation with elevated IL-17 were observed in ITP patients.

IL-23, which is known as a pro-inflammatory mediator, maintains the balance between regulatory and effector T-cells, and it is indispensable for promoting autoimmunity through T-cell-mediated inflammation [19–21]. Recently, numerous studies have found that increased expression of IL-23 is involved in the pathogenesis of many autoimmune diseases, such as inflammatory bowel disease, Crohn’s disease (CD)[15], and multiple sclerosis (MS)[22]. IL-23 exerts its biological effects by binding to its receptors. IL-23 receptors consist of IL-23R and IL-12Rβ1, while the receptor for IL-12 consists of the common IL-12Rβ1 and a unique IL-12Rβ2[23]. Our results show that the IL-12 unique subunit p35 and IL-12Rβ2 were not elevated in ITP patients. Therefore, our results showed the increased expression of common IL-12p40 and receptor were caused by IL-23, and not IL-12. Previous study demonstrated that IL-23 can be induced by Toll-like receptor (TLR) agonists and interactions with T cells (CD40/CD40L interaction)[24]. Liu et al found that TLR4 was abnormal expression in ITP patients and demonstrated that TLR4 in combination with FcγRII could cause DCs to express cytokines IL-23 then promote effector Th17 cell responses[25]. These may suggest that the elevated of IL-23 may cause by abnormal TLR, but the exact mechanism need further study.

Th17, a newly defined CD4+ Th subset, has been implicated in the pathogenesis of many diseases, particularly autoimmune disorders[26,27]. Th17 modulates the pro-inflammatory response by producing IL-17, IL-6, IL-21 and other mediators[28,29]. Until now, many studies have revealed that Th17 cell was increased in ITP patients. Ji et al found that the ratio of Treg/Th17 correlated with the disease activity might have prognostic role in ITP[30]. Rocha et al found that increased levels of IL-17A and of Th17-related cytokines contribute to the pathogenesis of ITP[31]. Wang et al found that elevation of IL-17 and IFN-γ may be an important dysregulation of cellular immunity in pediatric patients with chronic ITP[32]. We found that IL-17 was increased both at the mRNA and protein levels in ITP patients, which is consistent with previous studies. RORC, the gene encoding Th17 transcription factor, RORγT, is a specific regulator for Th17 differentiation[33]. In the present study, we found that RORC mRNA expression is increased in ITP patients. These results indicate that Th17 is an abnormal activator in ITP patients. Although it was shown in previous studies[6,34], our work focused on the
Recent studies have shown that Th17 differentiation and activation is regulated by IL-23 [12]. To date, IL-23 is known as the potent cytokine that promotes IL-17 production by Th17 cells. Although IL-6 and TGF-β can also drive the differentiation of Th17 cells from naïve CD4+ T cells [35], IL-23 is the key factor for the maturation and phenotype stabilization of pathogenic Th17 cells [12,36]. Th17 development was suspended in the early activation stage without IL-23 [37]. IL-23 exerts its biological function via its receptor in order to induce Janus family kinase (Jak2) and tyrosine kinase 2 (Tyk2) phosphorylation, which is followed by the activation of STAT1, STAT3, STAT4 and STAT5 [38–40]. Among these, the activation of STAT3 by IL-6 and IL-23 is crucial in Th17 lineage differentiation. Numerous studies clearly suggested that IL-23 promotes the pathogenicity of Th17 cells through several mechanisms: 1.)
maintenance the expression of Th17 signature gene (RORC and IL-17); 2.) induction of effector genes (IL-22, Csf2 and IFN-γ) and down-regulation of repressive genes (IL-2, IL-27 and IL-12); 3.) amplification of its own signal through the up-regulation of IL-23R expression[41]. In the present study, we found a positive correlation between IL-23 and IL-17 levels in plasma from ITP patients, which suggests that IL-17 production is influenced by IL-23.

In order to investigate the role of IL-23 in IL-17 production in ITP, we cultured PBMCs from ITP patients with exogenous rhIL-23 in vitro. Our results showed increased IL-6, IL-17, and IL-21 expression in the culture supernatant following IL-23 stimulation. This suggests that IL-23 is involved in ITP through activation of the Th17 response. We also found a negative relationship between IL-23 levels and platelet counts. To some extent, the platelet counts reflect the disease severity as well as the development of the disease[42]. So IL-23 levels could serve as a potential index to evaluate the disease state.

Moreover, we followed up on 11 patients. Our results showed that both IL-17 and IL-23 decreased after effective treatment. It is widely accepted that glucocorticoids are the first-line treatment approach for ITP in clinic[43]. Several previous studies found that high-dose dexamethasone could correct the T cell subsets and cytokine profiles in ITP [44–46]. Moreover, Li et al [44] found that dexamethasone corrected the T cell subset levels by promoting GATA3 and FOXP3 expression and inhibiting RORC expression. In the present study, we found that IL-17 expression decreased after treatment. This result may be caused by the decreased number of Th17 cells after dexamethasone treatment through depression of RORC.

Based on the above results, we presume that the elevated IL-23 expression in the plasma of ITP patients promoted Th17 responses through RORC up-regulation and the STAT3 signal

Fig 4. Correlation between plasma IL-23 and IL-17, Th17, platelets in ITP patients.

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Fig 5. Th17 related cytokines production in culture supernatant in the presence and absence of IL-23.

Fig 6. Plasma IL-23 and IL-17 level change in ITP patients after effective treatment.
pathway. However, the underlying mechanism requires further investigation, and it is thereby a limitation in our study.

In conclusion, the IL-23/Th17 pathway is engaged in the development of ITP. Monitoring the IL-23 level in plasma may aid in evaluating the ITP disease state. Further studies may focus on the possibility of employing anti-IL-23 drugs to treat ITP patients.

**Author Contributions**

Conceived and designed the experiments: XY AMD BHQ. Performed the experiments: XY LZ. Analyzed the data: XY LZ HW AMD BHQ. Contributed reagents/materials/analysis tools: XY LZ HW YC WWZ AMD BHQ. Wrote the paper: XY LZ AMD BHQ. Sample collection: RRZ CPF.

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