Tissue and Serum Inflammatory Cytokine Levels in Korean Psoriasis Patients: A Comparison between Plaque and Guttate Psoriasis

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Background: The phenotypic heterogeneity of psoriasis could be explained by the alternate activation of either T-helper (Th)-1- or Th-17-related cytokines. However, evidence directly supporting this hypothesis is scarce.

Objective: To characterize the expression of Th-1- and Th-17-related cytokines according to the morphological psoriasis phenotype: guttate vs. plaque.

Methods: In this study, we enrolled 68 patients exhibiting either guttate or plaque psoriasis, and 10 healthy controls. To avoid age-related bias, age matching was performed for each group. Circulating levels of interferon (IFN)-γ, interleukin (IL)-1RA, IL-2, IL-12p40, IL-17A, IL-22, and IL-23 were measured with an enzyme-linked immunosorbent assay (ELISA). Psoriasis-affected tissue was obtained through biopsy sampling from the eight patients who exhibited the most typical morphology. Levels of IL-1RA, IL-12p40, IL-17, IL-22, and IL-23 in the psoriasis tissue samples were measured with western blot analysis.

Results: ELISAs of the serum samples showed higher levels of inflammatory cytokines such as IL-1RA, IL-2, IL-23, and IFN-γ in patients with psoriasis than in healthy controls. However, the inflammatory cytokine levels did not differ significantly between guttate and plaque psoriasis patients. Western blot analysis of psoriatic tissue revealed higher protein levels of Th-1- and Th-17-related cytokines in patients than in healthy controls. The levels of IL-12p40 and IL-23 were unexpectedly higher in plaque tissue than in guttate tissue.

Conclusion: The morphological phenotype of psoriasis does not appear to be determined by a specific activation of either the Th-1 or Th-17 pathway. Rather, the cytokine profile influences disease activity and is altered according to the status of the lesion (early or chronic).

Keywords: Cytokines, Phenotype, Psoriasis

INTRODUCTION

Psoriasis is a chronic inflammatory disorder of the skin that is known to be mediated by immune alterations. Measurements of T-cell activity after antigen challenge, therapeutic inhibition of activated T-cells, and cell transfer and explantation have been performed in a large number of studies, with the results demonstrating that psoriasis is a T-cell-mediated disorder. It was previously assumed that T-helper (Th)-1 cells play a dominant role in the initiation and maintenance of psoriasis along with interferon (IFN)-γ, interleukin (IL)-2, and IL-12. It has recently been demonstrated that a new subset of T cells, Th-17 cells, are also important in the pathogenesis of psoriasis. The secretion of IFN-α from plasmacytoid dendritic cells (DCs) is one of the earliest events driving the inflammatory eruption in psoriasis. IFN-α stimulates DCs, which produce tumor necrosis factor (TNF)-α and inducible nitric oxide synthase, to
secrete IL-23\(^7\). IL-23 stimulates Th-17 cells to produce Th-17-related cytokines (henceforth referred to as Th-17 cytokines), thus increasing Th-17 cell survival and proliferation. The Th-17-derived cytokines include IL-17A, IL-17F, IL-6, TNF-\(\alpha\), and IL-22\(^{10,11}\). IL-22 has been shown to induce hyperplasia and abnormal differentiation of keratinocytes, ultimately resulting in plaque formation\(^{12}\).

Psoriasis manifests with a heterogeneous phenotype and exhibits a wide range of clinical activities; it has also been suggested that one phenotype can transform into another. Christophers\(^13\) recently proposed that rerouting cutaneous inflammation from the IL-12 pathway toward the IL-23 pathway, and vice versa, has a bearing on both the diversity of phenotypes and their instability. However, very few studies have explored the heterogeneity of the psoriasis phenotype.

In this study, we measured the protein levels of the Th-1- and Th-17 cytokines in human psoriasis tissue and serum by using western blot and enzyme-linked immunosorbent assay (ELISA), respectively. In addition, differences in the levels of cytokines between the guttate and plaque types were investigated.

**MATERIALS AND METHODS**

**Patients and controls**

Sixty-eight patients with psoriasis (38 with plaque psoriasis and 30 with guttate psoriasis) were enrolled for the analysis of their serum with an ELISA. To avoid age bias, we tried to enroll age-matched patients with guttate and plaque psoriasis. Eight patients exhibiting the typical guttate or plaque morphology were enrolled to participate in the western blot analysis aimed at exploring differences according to morphological phenotype. The diagnosis of psoriasis was confirmed by using clinical and histopathological criteria. The following major inclusion criteria were implemented: no local or systemic treatment for at least 4 weeks before study entry; no significant infection or immune suppression; and no history of specific medical diagnoses with renal, hepatic, cardiovascular, pulmonary, rheumatic, or endocrine involvement. The patients were divided into two groups according to their plaque morphology: guttate or plaque. The guttate group comprised patients whose lesions were papules with diameters of <1 cm, whereas the plaque group comprised those with at least one lesion with a long axis of >5 cm. The Psoriasis Area and Severity Index (PASI) was used to evaluate the clinical severity of the disease. The control group comprised healthy volunteers with no history of specific medical diagnoses with renal, hepatic, cardiovascular, pulmonary, rheumatic, or endocrine involvement. This study was conducted in accordance with the guidelines of the Helsinki Conference and the Korean Good Clinical Practice, with the participants’ rights and safety taking precedence. Approval for the study was obtained from the institutional review board of Konkuk University. All patients gave their informed consent to participate before screening.

**Western blot analysis of tissue samples**

Skin biopsy samples (4 mm) were collected from the lesions of patients with psoriasis (four guttate and four plaque lesional biopsies), and from the healthy residual skin of patients after lipoma excision, which was neither inflamed nor pathologic. The samples were immediately stored at \(-70^\circ\text{C}\) until use.

Tissue proteins were prepared in radioimmunoprecipitation assay buffer (1× phosphate-buffered saline, 1% NP-40, 0.5% sodium deoxycholate, 0.1% sodium dodecyl sulfate [SDS; Sigma-Aldrich, St. Louis, MO, USA], and 10 \(\mu\)g/ml phenylmethylsulfonyl fluoride). The proteins (50 \(\mu\)g/lane) were separated on NuPAGE 4% to 12% Bis-Tris polyacrylamide gels (Invitrogen, Carlsbad, CA, USA) and then transferred electrophoretically to immunoblot polyvinylidene fluoride membranes. The membranes were incubated with the following specific antibodies for 2 hours at room temperature: anti-human IL-1RA (Santa Cruz Biotechnology, Santa Cruz, CA, USA), anti-IL-22 (Abcam Technologies, Cambridge, MA, USA), anti-IL-23 (Abcam Technologies), anti-IL-12B/IL-12p40 (MyBiosource, San Diego, CA, USA), and anti-IL17 (R&D Systems, Minneapolis, MN, USA). The anti-IL-12B/IL-12p40 antibody used in this study detects only IL-12p40 (i.e., not IL-23p40). The membranes were then washed and incubated with horseradish-peroxidase-conjugated secondary antibodies (Santa Cruz Biotechnology). The signal was visualized with an enhanced chemiluminescence detection kit (Promea, Madison, WI, USA) by using x-ray films. The blots were stripped by using a stripping buffer (100 mM \(\beta\)-mercaptoethanol, 2% SDS, and 62.5 mM Tris-HCl [pH 6.7]) and reprobed with anti-\(\beta\)-actin antibody (Sigma-Aldrich) as loading controls. Multi-Gauge ver. 3.1 software from Fuji Film (Tokyo, Japan) was used to quantify the resulting bands, and the data were normalized with \(\beta\)-actin as a reference.

**Assays for serum samples**

Venous blood samples (5～10 ml) were collected from patients with psoriasis (30 with guttate psoriasis and 38 with plaque psoriasis) into vacuum tubes in sterile conditions. The serum was extracted from the samples through centrifugal spinning, and immediately frozen at \(-70^\circ\text{C}\) until analysis. Multiple cytokine analyses were conducted by
using xMAP technology (Luminex, Austin, TX, USA) to measure the serum levels of IFN-γ, IL-1RA, IL-2, IL-12p40, IL-17A, IL-22, and IL-23. The Milliplex MAP multiplex assay was performed by using a 96-well microplate format according to the manufacturer’s instructions (Millipore, Billerica, MA, USA). Briefly, each of the bead solutions was transferred into a mixing vial and adjusted to a volume of 3 ml with bead diluents. Internal controls, in the range 0 to 10,000 pg/ml for each cytokine, were included with every assay. After the addition of sera and beads, the mixture was incubated overnight at 4°C. Detection antibodies and streptavidin-phycoerythrin were successively added at room temperature for 30 minutes, and the plate was analyzed with the Luminex 200 system.

Statistical analysis

Data analyses were performed by using SAS software (ver. 9.3; SAS Institute, Cary, NC, USA). The three groups were compared by using the Kruskal-Wallis test. In addition, the Wilcoxon rank-sum test was performed to compare the data for the two groups. The cutoff for statistical significance was set at \( p < 0.05 \).

RESULTS

Patient demographics

The mean age of the 68 patients was 36.6±14.2 years (mean±standard deviation [SD]), and that of controls was 32.58±10.0 years \( (p>0.05) \); among the guttate group, plaque group, and healthy controls, age showed no significant difference after age matching (37.6±14.8 years vs. 35.7±13.9 years vs. 30.2±7.4 years, \( p>0.05 \)). The mean age of eight patients from whom psoriatic tissue was obtained for western blot analysis was 37.0±13.0 years, and that of controls was 51.0±3.9 years \( (p<0.05) \); there was a significant difference between the guttate group, plaque group, and healthy controls (41.0±17.8 years vs. 33.0±6.0 years vs. 51.0±3.9 years, \( p<0.05 \)).

The mean PASI for the 68 patients was 8.2±4.0 points (mean±SD), and was significantly higher in the plaque group (9.1±3.4 points) than in the guttate group (7.2±3.5 points, \( p<0.05 \)). The mean PASI of the eight participants from whom psoriatic tissue was obtained for western blot analysis was 11.6±3.6 points; the plaque group had a

### Table 1. Demographics of the healthy controls and psoriasis patients

| Variable       | Healthy control | Guttate psoriasis | Plaque psoriasis | \( p \)-value |
|----------------|-----------------|-------------------|------------------|--------------|
| Western blot analysis |
| No. of case | 4               | 4                 | 4                |              |
| Age (yr)  | 51.0±3.9        | 41.0±17.8         | 33.0±6.0         | 0.002*       |
| PASI      | 10.1±3.8        | 13.2±3.0          | 3.12             |              |
| ELISA     |
| No. of case | 10              | 30                | 38               |              |
| Age (yr)  | 30.2±7.4        | 37.6±14.8         | 35.7±13.9        | 0.397        |
| PASI      | 7.2±3.5         | 9.1±4.3           | 0.045*           |              |

Values are presented as number only or mean±standard deviation. Kruskal-Wallis test comparing age among healthy controls and patients with guttate and plaque psoriasis. Wilcoxon rank-sum test comparing duration and PASI between the patients with guttate and plaque psoriasis. ELISA: enzyme-linked immunosorbent assay, PASI: Psoriasis Area and Severity Index. * \( p<0.05 \).

### Table 2. Comparison of cytokine levels between healthy controls and psoriasis patients

| Variable       | Healthy controls (n=4) | Psoriasis patients (n=8) | \( p \)-value |
|----------------|------------------------|--------------------------|--------------|
| IL-1RA/\( \beta \)-actin | 0.48±0.07              | 0.77±0.08                | 0.0008*      |
| IL-23/\( \beta \)-actin  | 0.00±0.00              | 0.26±0.10                | 0.0003*      |
| IL-22/\( \beta \)-actin  | 0.19±0.05              | 0.52±0.15                | 0.0011*      |
| IL-12p40/\( \beta \)-actin | 0.40±0.04             | 1.01±0.18                | 0.0008*      |
| IL-17/\( \beta \)-actin  | 0.21±0.03              | 0.73±0.06                | 0.0003*      |

Values are presented as mean±standard deviation. The experimental Western blot data are normalized relative to the \( \beta \)-actin reference. Wilcoxon rank-sum test comparing the protein level of tissue inflammatory cytokines between the healthy controls and psoriasis patients. IL: interleukin, IL-1RA: IL-1 receptor antagonist. * \( p<0.05 \).
Table 3. Comparison of cytokine levels between guttate and plaque psoriasis patients

| Variable         | Guttate psoriasis (n=4) | Plaque psoriasis (n=4) | p-value |
|------------------|-------------------------|------------------------|---------|
| IL-1RA/β-actin   | 0.82±0.07               | 0.72±0.06              | 0.0833  |
| IL-23/β-actin    | 0.19±0.07               | 0.32±0.07              | 0.0209* |
| IL-22/β-actin    | 0.48±0.20               | 0.56±0.10              | 0.5637  |
| IL-12p40/β-actin | 0.88±0.16               | 1.14±0.05              | 0.0209* |
| IL-17/β-actin    | 0.72±0.08               | 0.74±0.05              | 0.5637  |

Values are presented as mean±standard deviation. The experimental Western blot data are normalized relative to the β-actin reference. Wilcoxon rank-sum test comparing the protein level of tissue inflammatory cytokines between the patients with guttate and plaque psoriasis. *p<0.05.

Table 4. Comparison of serum cytokine levels between healthy controls and psoriasis patients

| Variable          | Healthy controls (n=10) | Psoriasis patients (n=68) | p-value |
|-------------------|-------------------------|---------------------------|---------|
| IL-1RA (pg/ml)    | 1.76±3.18               | 4.72±5.85                 | 0.0441* |
| IL-2 (pg/ml)      | 1.00±1.69               | 1.07±0.66                 | 0.0453* |
| IL-12p40 (pg/ml)  | 1.60±2.84               | 3.38±3.90                 | 0.1386  |
| IL-17A (pg/ml)    | 1.42±2.36               | 1.73±3.23                 | 0.6809  |
| IL-22 (pg/ml)     | 0.11±0.02               | 0.15±0.11                 | 0.6688  |
| IL-23 (pg/ml)     | 0.99±0.23               | 2.04±1.12                 | <0.0001*|
| IFN-γ (pg/ml)     | 1.32±1.92               | 4.71±3.49                 | 0.0003* |

Values are presented as mean±standard deviation. Wilcoxon rank-sum test comparing the serum level of inflammatory cytokines between healthy controls and psoriasis patients. IL: interleukin, IL-1RA: IL-1 receptor antagonist. *p<0.05.

Table 5. Comparison of serum inflammatory cytokine levels between patients with guttate and plaque psoriasis

| Variable          | Guttate psoriasis (n=30) | Plaque psoriasis (n=38) | p-value |
|-------------------|--------------------------|-------------------------|---------|
| IL-1RA (pg/ml)    | 5.01±5.47                | 4.49±6.19               | 0.4589  |
| IL-2 (pg/ml)      | 1.12±0.72                | 1.04±0.61               | 0.7897  |
| IL-12p40 (pg/ml)  | 2.84±3.73                | 3.80±4.03               | 0.2881  |
| IL-17A (pg/ml)    | 1.40±2.71                | 1.99±3.60               | 0.4752  |
| IL-22 (pg/ml)     | 0.15±0.11                | 0.16±0.10               | 0.4311  |
| IL-23 (pg/ml)     | 2.07±1.26                | 2.02±1.02               | 0.7716  |
| IFN-γ (pg/ml)     | 4.77±3.94                | 4.66±3.14               | 0.9457  |

Values are presented as mean±standard deviation. IL: interleukin, IL-1RA: IL-1 receptor antagonist, IFN: interferon. Wilcoxon rank-sum test comparing the serum levels of inflammatory cytokines between patients with guttate and plaque psoriasis.

A Comparison between Plaque and Guttate Psoriasis

Psoriasis manifests with various morphological phenotypes, and in each form the morphology may alter with the disease activity. The question of what explains the difference between these morphological phenotypes is a long-standing one. It was assumed that the morphological diversity of psoriasis is related to differences in inflammatory cytokines, particularly Th-1- and Th-17-related cytokines. The accumulated evidence concerning inflammatory cytokines and the histopathology among different psoriatic phenotypes suggests that a combination of inflammatory pathways routing from IL-12/IFN-γ to IL-23/IL-17, and vice versa, could determine the psoriasis phenotype. Christophers proposed that Th-1-predominant activation represents stable plaque morphology, and that Th-17 activation explains the inflammatory plaque types such as guttate or pustular psoriasis. However, studies of cytokine profiles relative to morphological phenotypes have yielded conflicting results. To examine the inflammatory psoriasis types such as guttate or pustular psoriasis, we used reverse transcription polymerase chain reaction (RT-PCR) and ELISA. Our western blot analysis showed that both Th-1- and Th-17-related cytokine levels were significantly elevated in patient tissue compared with those in healthy control

DISCUSSION

Psoriasis manifests with various morphological phenotypes, and in each form the morphology may alter with the disease activity. The question of what explains the difference between these morphological phenotypes is a long-standing one. It was assumed that the morphological diversity of psoriasis is related to differences in inflammatory cytokines, particularly Th-1- and Th-17-related cytokines. The accumulated evidence concerning inflammatory cytokines and the histopathology among different psoriatic phenotypes suggests that a combination of inflammatory pathways routing from IL-12/IFN-γ to IL-23/IL-17, and vice versa, could determine the psoriasis phenotype. Christophers proposed that Th-1-predominant activation represents stable plaque morphology, and that Th-17 activation explains the inflammatory plaque types such as guttate or pustular psoriasis. However, studies of cytokine profiles relative to morphological phenotypes have yielded conflicting results. To examine the inflammatory psoriasis types such as guttate or pustular psoriasis, we used reverse transcription polymerase chain reaction (RT-PCR) and ELISA. Our western blot analysis showed that both Th-1- and Th-17-related cytokine levels were significantly elevated in patient tissue compared with those in healthy control.
tissue, confirming that both the Th-1 and Th-17 pathways are associated with the pathogenesis of psoriasis regardless of morphology.

In contrast to previous studies, our serum inflammatory cytokine data showed no difference in the levels of inflammatory Th-1 and Th-17 cytokines between the guttate and plaque groups. There has been discrepancy in the findings of analyses of inflammatory cytokines relative to morphological phenotype. Yan et al. and Choe et al. compared serum inflammatory cytokine levels between guttate and plaque psoriasis or early inflammatory and stable psoriasis by using ELISA, and found that IL-17 levels were higher in the former. By contrast, Yilmaz et al. performed cytokine analysis on inflammatory psoriatic tissue by using RT-PCR, and reported that the expression of tissue IL-17 mRNA was not significantly elevated in guttate psoriasis compared with that in plaque psoriasis.

In the present study, we also performed western blot analysis with human psoriasis tissue. Our results showed a discrepancy between the results of the tissue and serum samples when comparing the guttate and plaque groups; in the serum, there were no significant differences in the levels of inflammatory Th-1 and Th-17 cytokines between the two groups, whereas, unexpectedly, the expression of Th-17 cytokines such as IL-23 was higher in plaque tissue than in guttate tissue. In a recent review by Christophers et al., it was postulated that during the progression of stable psoriasis, two types of inflammation may coexist in the same patient: an IL-17/Th17-dominated autoinflammation in highly inflammatory early lesions, and a Th1-dominated late-phase inflammation that prevails in stable psoriasis plaques. In other words, it means that each lesion even in the same patient shows a different disease activity, which leads to a different immune pathway predominance in each lesion. Thus, the results of the present study conclusively support the hypothesis that the cytokine profile is not associated with morphological phenotype, but rather with whether the lesions are early or chronic.

Overall, the IL-23/IL-17 pathway did not predominate in guttate psoriasis relative to plaque psoriasis. However, a limitation of this study is the relatively small number of patients involved, and thus this finding needs to be confirmed in future studies involving larger samples. In conclusion, the findings of this study suggest that both the Th-1 and Th-17 pathways are associated with the pathogenesis of psoriasis regardless of the morphological phenotype. Our results support the theory that the predominant pathway, either IL-23/IL-17 or IL-12/IFN-γ, is determined by the status of the lesion as either early or chronic.

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