

# Relationship Between Interleukin-6 –174G/C Genetic Variant and Efficacy of Methotrexate Treatment in Psoriatic Arthritis Patients

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**Introduction:** The purpose of the study was to investigate whether single-nucleotide polymorphisms (SNPs) IL-6 –174 G/C and IL-6R Asp358Ala are associated with susceptibility to psoriatic arthritis (PsA) or affect response to treatment with methotrexate (MTX).

**Patients and Methods:** Seventy-four patients diagnosed with PsA and qualified for MTX treatment were enrolled to the study. The control group consisted of 120 healthy individuals. Polymorphisms IL-6 –174 G/C and IL-6R Asp358Ala were genotyped using a polymerase chain reaction (PCR) amplification employing LightSNiP assays.

**Results:** A significant association between the IL-6 –174 CC genotype and an improved clinical outcome of MTX therapy was observed. A good response was more frequently observed among PsA patients bearing the IL-6 –174 CC genotype than patients with the GC or GG genotypes ( $P = 0.007$ ). On the other hand, patients carrying the IL-6 –174 GC genotype less frequently responded to MTX treatment as compared to patients with other genotypes ( $P = 0.006$ ). With respect to the IL-6R Asp358Ala SNP, there were no significant differences in genotype and allelic frequencies in relation to clinical outcome of MTX treatment. No association was found between the IL-6 –174 G/C or IL-6R Asp358Ala SNPs and PsA susceptibility.

**Conclusion:** Results from this study provide evidence that the IL-6 –174 G/C polymorphism might influence efficacy of MTX treatment.

**Keywords:** psoriatic arthritis, methotrexate treatment, IL-6 –174G/C polymorphism, IL-6R Asp358Ala polymorphism

## Plain Language Summary

Psoriatic arthritis (PsA) is a disease caused by the immune system attacking its own tissues. PsA is a form of inflammatory arthritis and it affects joints as well as skin.

A first-line drug for a treatment of PsA is methotrexate (MTX). This drug can be very effective on some, however a substantial group of patients responds to the therapy partially or not at all.

The exact cause of PsA is not known, but it is well established that a genetic background plays an important role in the disease development.

Also, it is supposed that genetic factors may influence response to the PsA treatment.

The aim of this research was to investigate whether changes in genes encoding the interleukin-6 (IL-6) and interleukin-6 receptor (IL-6R) might have an effect on the response to MTX in PsA patients. Two variants within these genes were selected: IL 6 –174 G/C in the IL-6 gene and IL-6R Asp358Ala in the IL-6R gene.

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Interleukin-6 belongs to a group of small proteins called cytokines. Cytokines are produced by numerous immune system cells and are important in the regulation of immunity and inflammation.

The research was performed on 74 PsA patients treated with MTX.

The results showed that patients with the C variant of the IL-6 -174 G/C gene responded significantly better to the therapy with MTX.

This study showed that the IL-6 -174 G/C genetic variant could affect the response to MTX in PsA patients.

## Introduction

Psoriatic arthritis (PsA) is a multifactorial chronic inflammatory disease manifested by joint inflammation and cutaneous psoriasis.

The precise etiology of PsA remains unknown, however both genetic and environmental factors contribute to development of this immune-mediated disease. Twin and family studies demonstrated strong heritability of PsA and pointed to a strong genetic component underlying the etiology of this disease.<sup>1</sup> Conventional synthetic disease modifying anti-rheumatic drugs (csDMARDs), including methotrexate (MTX), constitute the first line therapy for PsA. This immunomodulatory agent is one of the most commonly prescribed medications for this disorder, however patients display variable responses to treatment.<sup>2,3</sup> Although the exact mechanism of action of MTX in the context of PsA is not completely understood, it has been reported that therapy with this drug results in decreased production of interleukin-6 (IL-6).<sup>4,5</sup>

Interleukin-6 is a multifunctional cytokine characterized by a broad range of biologic activities, comprising immune regulation, inflammation and oncogenesis.<sup>6</sup> Synthesis of acute-phase protein<sup>7</sup> and transition between acute and chronic inflammation is stimulated by IL-6.<sup>8</sup> Also, this cytokine exerts an effect on bone metabolism and mediates bone resorption through stimulation of osteoclast cell formation.<sup>9</sup> Moreover, IL-6 is associated with increased vascular endothelial growth factor (VEGF) expression, leading to pannus development. In addition, IL-6 is involved in inflammatory process by inducing differentiation of activated B cells into antibody-producing cells, differentiation of CD8<sup>+</sup> T cells into cytotoxic T cells and by playing a crucial role in Th17 development. The IL-6 interacts with its specific interleukin-6 receptor (IL-6R) and the IL-6-IL-6R complex associates with glycoprotein 130 subunits (gp130), leading to activation of signaling cascade.<sup>10</sup> The receptors for IL-6 exist as soluble as well

as membrane-bound forms. Signaling through soluble IL-6R, called the trans-signaling pathway, is responsible for pro-inflammatory activities of IL-6. On the other hand, the pathway via the membrane-bound IL-6R, known as classic-signaling, mediates anti-inflammatory and regenerative properties of this cytokine.<sup>11</sup>

The production of IL-6 protein might be partially determined on the genetic level by polymorphisms located within the promoter region of its gene. The IL-6 -174 polymorphism has been implicated in the regulation of the IL-6 levels. This SNP consists of a G to C substitution at position -174 in the promoter region of the IL-6 gene. The G allele has been associated with increased transcriptional activity and higher levels of IL-6 in in vitro and in vivo studies.<sup>12</sup> The IL-6R Asp358Ala has also been demonstrated to affect levels of circulating soluble IL-6 receptor (sIL-6R).<sup>13</sup> It has been established that the IL-6R C allele is associated with increased sIL-6R levels. The presence of the IL-6R CC genotype correlated with higher sIL-6R concentrations in patients diagnosed with asthma, diabetes as well as coronary heart disease.<sup>14–17</sup> Moreover, healthy individuals carrying the IL-6R C allele displayed substantially increased sIL-6R levels.<sup>14,15,18</sup> The relevance of the IL-6 -174 polymorphism in pathogenesis of autoimmune disorders was examined in numerous studies. However, this polymorphism has not been widely studied in the context of its potential impact on PsA pathogenesis and only two studies regarding the role of this SNP in context of the disease exist. To date, there are also no studies evaluating a possible role of functional interleukin-6 receptor (IL-6R) Asp358Ala (rs2228145) polymorphism in PsA pathology. A brief summary of studies performed to date regarding associations of these polymorphisms and arthritis as well as dermal diseases can be found in [Table 1](#).

The objective of this study was to investigate whether single-nucleotide polymorphisms (SNPs) IL-6 -174 G/C (rs1800795) and IL-6R Asp358Ala (rs2228145) are associated with susceptibility to PsA or affect response to treatment with MTX.

## Methods

### Patients

In total, 74 patients with PsA from the Department of Rheumatology and Internal Medicine of Wrocław Medical University were included in the study. All the patients were diagnosed with PsA according to the

**Table 1** Summary of Studies Performed Up to Date in Regard to Associations of the IL-6 -174 G/C and IL-6R Asp358Ala Polymorphisms and Arthritis as Well as Dermal Diseases

Disease	Effect	References
<b>IL-6 -174</b>		
RA susceptibility	Lack of association with disease risk	[19]
RA susceptibility	GG genotype associated with disease risk	[20]
RA susceptibility	C allele associated with increased disease risk	[21]
Susceptibility to juvenile idiopathic arthritis	G allele associated with disease risk	[12,22]
Disease severity in RA	G allele correlated with higher disease activity	[23]
Disease severity in RA	CC genotype correlated with higher disease activity	[24]
Severity and progression of bone erosive damage	G allele correlated with increased progression of bone	[25]
Radiographic damage in RA	GG genotype correlated with increased radiographic damage	[26]
Hypertension in RA	Lack of association	[27]
Endothelial dysfunction in RA	GG genotype associated with more severe endothelial dysfunction	[28]
Risk of cardiovascular disease in RA	Lack of association	[29]
Risk of cardiovascular disease in RA	C allele correlated with cardiovascular disease risk	[30]
Response to anti-TNF therapy in RA	G allele associated with better response	[19,31,32]
Response to MTX therapy in RA patients	GG genotype correlated with worse response	[33,34]
Response to rituximab in RA	CC genotype associated with worse response	[35]
Disease activity and clinical pattern in PsA	GG genotype associated with peripheral form of PsA	[36]
PsA susceptibility and severity	Lack of association	[37]
Psoriasis susceptibility and response to phototherapy	GG genotype associated with disease risk	[38]
Response to anti-TNF treatment in psoriasis	C allele correlated with better response	[39]
Psoriasis susceptibility	Lack of association	[40]
Psoriasis susceptibility	GG genotype associated with decreased risk of disease	[41]
<b>IL-6R Asp358Ala</b>		
Response to biological therapy in RA	Lack of association	[35,42–44]
Risk of cardiovascular disease in RA	Lack of association	[29,45]
RA susceptibility	C allele associated with decreased risk	[46]
Disease activity in RA	C allele in patients correlated with lower disease activity	[47]
Susceptibility to systemic sclerosis	Lack of association	[48]
Susceptibility to atopic dermatitis	Allele C correlated with increased risk of disease	[49]

Classification Criteria for Psoriatic Arthritis (CASPAR) and were naïve to MTX or other csDMARDs. The control group consisted of 120 age and gender matched healthy blood donors without any family history of autoimmune disorders. All the participants were of Caucasian ancestry.

Demographic data and complete medical history were collected from each patient. The clinical and laboratory evaluation included a count of swollen and tender joints (SJC and TJC, respectively), erythrocyte sedimentation rate (ESR), C-reactive protein (CRP), rheumatoid factor (RF), visual analogue scale (VAS; range, 0 to 100 mm) of pain, health assessment questionnaire (HAQ), global health assessment by the patient and the physician. Disease activity was measured using disease activity score (DAS28). Psoriasis severity was assessed using the psoriasis area severity index (PASI).

Criteria excluding patients from this study comprised of: age below 18; coexistence of other systemic diseases of connective tissue besides PsA; clinically significant impairment of hepatic and renal function; infections with hepatotropic viruses; infections resistant to therapy; ongoing history of cancer or uncontrolled diabetes; alcohol abuse; pregnancy or breastfeeding; prior treatment with csDMARDs or biologic agents; insufficient clinical records; unwillingness or inability to cooperate.

The main characteristics of patients are summarized in Table 2. Thirty-one subjects were females and the mean  $\pm$  SD age of the patients was  $46.0 \pm 10.9$  years. PsA was classified as oligoarticular in 12 patients and polyarticular in 62 patients. All the patients were negative for rheumatoid factor. The mean age at the onset of psoriasis was  $29.4 \pm 14.2$  years. Mean age at PsA onset was  $37.5 \pm 12.1$

**Table 2** Demographic and Clinical Characteristics of the Study Subjects

PsA Patients	N = 74
Sex [females/males (% of females)]	31/43 (41.9)
Age (years) [mean (±SD)]	46.0 (±10.9)
BMI [mean (±SD) % overweight/% obese]	27.6 (±5.2) 70.3/28.4
Current smokers [number of patients (%)]	10 (13.5)
Alcohol intake [mean units per week (±SD)]	4.7 (±3.6)
Psoriasis onset (years) [mean (±SD)]	29.4 (±14.2)
PsA duration (years) [mean (±SD)]	5.3 (±4.2)
Arthritis onset (years) [mean (±SD)]	37.5 (±12.1)
Arthritis pattern [number of patients (%)]	
Oligoarthritis	12 (16.2)
Polyarthritis	62 (83.8)
Tender joint count [mean (±SD)]	13.2 (±4.9)
Swollen joint count [mean (±SD)]	12.7 (±4.9)
Dactylitis presence [number of patients (%)]	25 (33.8)
CRP mg/L [mean (±SD)]	9.0 (±12.3)
ESR mm/h [mean (±SD)]	20.1 (±19.7)
DAS28 [mean (±SD)]	5.6 (±1.0)
PASI score [mean (±SD)]	14.6 (±3.7)
VAS of pain [mean (±SD)]	64.3 (±15.5)
Type of psoriasis [number of patients (%)]	
Type 1 (<40 years)	54 (73.0)
Type 2 (≥40 years)	20 (27.0)

**Abbreviations:** DAS28, disease activity score 28; CRP, C-reactive protein; ESR, erythrocyte sedimentation rate; PASI, psoriasis area and severity index; VAS, visual analogue scale; SD, standard deviation.

years. Fifty-four patients had type 1 psoriasis, the rest had type 2 psoriasis.

The patients were administered with MTX dose of 25 mg/week orally. All the patients supplemented folic acid dose of 10 mg/week during treatment. Efficacy of MTX treatment was evaluated using the American College of Rheumatology definition of 20% improvement (ACR20).<sup>50</sup> Patients were considered as ACR20 responders when they achieved an improvement of at least 20% in both tender and swollen joint counts (TJC and SJC), as well as at least 20% improvement in 3 of the 5 following measures: the physician's global assessment of disease activity, the patient's global assessment of the disease activity, the patient's functional ability assessment and acute-phase reactant value (CRP). As a secondary outcome measure, psoriatic arthritis response criteria (PsARC) were employed. These criteria comprise of: TJC, SJC (assessing 68 and 66 joints, respectively, including distal interphalangeal joints of hands and feet), PtGA, and PhGA. Improvement by at least 30% in at least 2 of 4

measurements, one of which had to be a joint count, without decrease in any measurement, was defined as a positive outcome of therapy.<sup>51,52</sup> In addition, Dactylitis Severity Score (DSS) was also used as a secondary outcome measure. Each digit with dactylitis was classified in accordance to a four level scale from 0 to 3 (0 = no dactylitis, 1 = mild dactylitis, 2 = moderate dactylitis, 3 = severe dactylitis), where scores greater than 0 indicate a presence of dactylitis and a total score was calculated as a sum of scores for all 20 digits.

The study was approved by Wrocław Medical University Ethics Committee and written informed consents were obtained from all participants. The study was conducted in accordance with the Declaration of Helsinki.

## IL-6 and IL-6R Genotyping

For the present study the selection of genetic variants within the IL-6 as well as IL-6R genes was based on available literature analysis as well as search results from HapMap and NCBI dbSNP databases. Information regarding predicted functional consequences of SNPs was obtained using SNPinfo Web Server.<sup>53</sup> Studied SNPs were characterized with minor allele frequencies above 10% (1000 Genomes Project).<sup>54</sup>

Genomic DNA was isolated from peripheral blood taken on EDTA using Maxwell 16 Blood DNA Purification Kit (Promega Corp., Madison, WI, USA) following the instruction of the manufacturer.

The IL-6 -174 G/C (rs1800795) and IL-6R Asp358Ala (rs2228145) genetic variants were determined by real-time polymerase chain reaction (PCR) amplifications and melting-curve analysis using LightSNiP typing assay (TIB-MolBiol, Berlin, Germany). The real-time PCRs were carried out on a LightCycler 480 Real-Time PCR system (Roche Applied Science, Mannheim, Germany) in accordance to the manufacturer's recommendations. The reagents used for each reaction included 1.6 µL MgCl<sub>2</sub>, 14.4 µL H<sub>2</sub>O, 1 µL of LightSNiP reagent mix (containing premixed primers and probes specific for each variant; designed and manufactured by TIB Molbiol) and 2 µL of FastStart DNA Master HybProbe (Roche Diagnostics), which were set up in a final volume of 20 µL containing 1 µL of DNA solution. The cycling conditions were as follows: initial denaturation at 95 °C for 10 minutes, followed by 45 cycles of denaturation at 95 °C for 10 seconds, annealing at 60 °C for 10 seconds, and extension at 72 °C for 15 seconds. Fluorescence was assessed at the end of each annealing phase. The melting curve analysis of the amplification products was performed at 95 °C for 30

seconds, 40 °C for 2 minutes, and 75 °C with a ramp of 1.5 °C/s and continuous acquisition of data.

## Statistical Analysis

Genotype frequencies in cases and controls were tested for Hardy–Weinberg equilibrium using Population Genetics package for R. Means and standard deviations were calculated for continuous data and proportions for categorical ones. Comparisons between nonparametric variables were performed using the Mann–Whitney *U*-test. Fishers exact test was used to compare categorical variables. A *p* value < 0.05 was considered statistically significant. All statistical calculations were performed with R software environment (version 3.2.3; x86\_64-pc-linux-gnu).<sup>55</sup>

## Results

### Genotype and Allele Frequencies of the IL-6 –174 G/C and IL-6R Asp358Ala Polymorphisms in PsA Patients and Healthy Individuals

The genotype and allele frequencies of the IL-6 –174 G/C and IL-6R Asp358Ala polymorphisms observed in patients and controls are presented in Table 3.

Distribution of the genotypes and alleles of the IL-6 –174 G/C and IL-6R Asp358Ala polymorphisms among patients and controls were in agreement with the Hardy–Weinberg equilibrium.

The frequencies of the IL-6 –174 CC, IL-6 –174 GC, and IL-6 –174 GG genotypes within the patient group were 20.8%, 58.3%, and 20.8%, respectively, and 19.6%, 56.3%,

24.1%, respectively, in the control group. The genotype and allele frequencies of the IL-6 –174 G/C polymorphism did not differ between PsA patients and healthy controls.

With respect to the IL-6R Asp358Ala SNP, the frequencies of the AA, AC, and CC genotypes were 29.7%, 62.2%, and 8.1%, respectively, within the patient group, and 37.5%, 52.5%, 10.0%, respectively, in the control group. There were no significant differences between distribution of the IL-6R Asp358Ala genotypes and alleles in PsA patients and controls.

### Analysis of Clinical Features of PsA Patients in Relation to the IL-6 –174 G/C and IL-6R Asp358Ala Genetic Variants

The relationships of the IL-6 –174 G/C and IL-6R Asp358Ala genetic variants with baseline clinical parameters, including DAS28, CRP, ESR, TJC, SJC, and pain intensity, as well as presence of dactylitis, were examined and their results are shown in Table 4.

The genotype and allele frequencies of the IL-6 –174 G/C polymorphism did not differ between PsA patients with respect to CRP or ESR level. No significant association between the IL-6 –174 G/C polymorphism and DAS28 was also observed. No significant differences were also observed in TJC, SJC, PASI score or pain intensity with respect to genotypes and alleles of the IL-6 –174 G/C. Similarly, no relations between studied polymorphisms and presence of dactylitis were found.

Considering the IL-6R Asp358Ala polymorphism, there were no differences in genotype and allele distributions among PsA patients concerning CRP or ESR level. No correlation between the IL-6R Asp358Ala polymorphism and TJC, SJC or pain intensity has been observed. The analysis also revealed no statistically significant differences between DAS28 score or presence of dactylitis and the IL-6R Asp358Ala genetic variant.

### Relationships Between the IL-6 –174 G/C and IL-6R Asp358Ala Gene Polymorphisms and Clinical Outcome of Treatment with MTX

The clinical outcome of MTX treatment with respect to the IL-6 –174 G/C and IL-6R Asp358Ala genetic variants is presented in Tables 5 and 6.

There was a significant association between the IL-6 –174 G/C polymorphism and efficacy of MTX therapy. A response to MTX in accordance to the ACR20 criteria was more frequently observed among patients bearing the

**Table 3** The IL-6 –174 G/C and IL-6R Asp358Ala Genotypes and Alleles Frequencies in PsA Patients and Healthy Subjects

	PsA Patients Number (%)	Controls Number (%)
<b>IL-6 –174</b>		
C	72 (50.3)	107 (47.8)
G	71 (49.7)	117 (52.2)
CC	15 (20.8)	22 (19.6)
GC	42 (58.3)	63 (56.3)
GG	15 (20.8)	27 (24.1)
<b>IL-6R Asp358Ala</b>		
A	89 (60.5)	153 (63.8)
C	58 (39.5)	87 (36.3)
AA	22 (29.7)	45 (37.5)
AC	46 (62.2)	63 (52.5)
CC	6 (8.1)	12 (10.0)



**Table 4** Clinical Features at Baseline According to the IL-6 –174 G/C and IL-6R Asp358Ala Genotypes and Alleles Distribution of PsA Patients

	CRP Mean (±SD)	DAS28 Mean (±SD)	ESR Mean (±SD)	TJC Mean (±SD)	SJC Mean (±SD)	VAS Mean (±SD)	DSS Number (%)
<b>IL-6 –174</b>							
CC	5.2 (±6.7)	5.1 (±0.8)	11.8 (±7.8)	12.4 (±7.8)	11.6 (±7.7)	62.0 (±16.0)	2 (2.7)
GC	12.1 (±14.8)	5.8 (±1.0)	21.3 (±16.5)	13.6 (±5.0)	13.1 (±5.1)	63.8 (±15.6)	19 (25.7)
GG	7.8 (±14.5)	5.3 (±0.7)	16.8 (±11.9)	12.2 (±2.6)	11.2 (±3.0)	52.0 (±11.7)	4 (5.4)
<b>IL-6R Asp358Ala</b>							
AA	11.9 (±14.3)	5.8 (±1.2)	27.4 (±26.9)	13.9 (±4.4)	11.6 (±6.0)	61.8 (±14.8)	8 (10.8)
AC	10.0 (±13.2)	5.5 (±0.9)	18.0 (±16.6)	12.9 (±4.9)	12.1 (±4.6)	64.7 (±15.6)	16 (21.6)
CC	2.7 (±2.4)	5.8 (±0.9)	16.8 (±13.3)	13.8 (±4.8)	13.4 (±5.1)	63.3 (±14.9)	1 (1.4)

**Abbreviations:** DAS28, disease activity score 28; CRP, C-reactive protein; ESR, erythrocyte sedimentation rate; TJC, tender joint count; SJC, swollen joint count; VAS, visual analogue scale; DSS, dactylitis severity score; SD, standard deviation.

**Table 5** Genotypes and Alleles Frequencies of the IL-6 –174 G/C and IL-6R Asp358Ala Polymorphisms in Psoriatic Arthritis Patients in Relation to Response in Accordance to the ACR20 and PsARC Criteria

	ACR20		PsARC	
	Responders Number (%)	Non-Responders Number (%)	Responders Number (%)	Non-Responders Number (%)
<b>IL-6 –174</b>				
C	37 (58.7)	26 (48.1)	30 (57.7)	40 (47.6)
G	26 (41.3)	28 (51.9)	22 (42.3)	44 (52.4)
CC	12 (37.5) <sup>a</sup>	3 (8.3) <sup>a</sup>	10 (38.5) <sup>c</sup>	5 (11.9) <sup>c</sup>
GC	13 (40.6) <sup>b</sup>	27 (75.0) <sup>b</sup>	10 (38.5) <sup>d</sup>	30 (71.4) <sup>d</sup>
GG	7 (21.9)	6 (16.7)	6 (23.1)	7 (16.7)
<b>IL-6R Asp358Ala</b>				
A	40 (65.6)	40 (55.6)	36 (64.3)	45 (57.7)
C	21 (34.4)	32 (44.4)	20 (35.7)	33 (42.3)
AA	12 (38.7)	8 (22.2)	10 (35.7)	10 (25.6)
AC	17 (54.8)	24 (66.7)	16 (57.1)	25 (64.1)
CC	2 (6.5)	4 (11.1)	2 (7.1)	4 (10.3)

**Notes:** <sup>a</sup>CC vs GC+GG,  $P = 0.007$ , OR = 6.42, 95% CI (1.49, 39.73); <sup>b</sup>GC vs CC+GG,  $P = 0.006$ , OR = 0.23, 95% CI (0.07, 0.72); <sup>c</sup>CC vs GC+GG,  $P = 0.016$ , OR = 4.51, 95% CI (1.18, 19.72); <sup>d</sup>GC vs CC+GG,  $P = 0.011$ , OR = 0.26, 95% CI (0.08, 0.79).

**Abbreviations:** OR, odds ratio; 95% CI, 95% confidence interval.

IL-6 –174 CC genotype than in patients with the IL-6 –174 GC or GG genotypes ( $P = 0.007$ ; OR = 6.42; 95% CI = (1.49, 39.73)). On the other hand, patients carrying the IL-6 –174 GC genotype responded less frequently to MTX treatment, as judged by the ACR20 criteria and compared to patients with the IL-6 –174 CC or GG genotypes ( $P = 0.006$ ; OR = 0.23; 95% CI (0.07, 0.72)). Similar, though less significant, trends were observed when a response to MTX was classified by the PsARC criteria. Patients with the IL-6 –174 CC genotype were more frequent among responders' group in comparison to

other genotypes ( $P = 0.016$ ; OR = 4.51; 95% CI (1.18, 19.72)). Also, patients carrying the IL-6 –174 GC genotype responded to the treatment in accordance to the PsARC classification less frequently than other patients ( $P = 0.011$ ; OR = 0.26; 95% CI (0.08, 0.79)). No associations have been observed between the studied polymorphisms and response to MTX measured as the dactylitis severity score.

With respect to the IL-6R Asp358Ala polymorphism, there were no significant differences in genotype and allelic frequencies in relation to response to MTX treatment.

**Table 6** Genotypes Frequencies of the IL-6 –174 G/C and IL-6R Asp358Ala Polymorphisms in Psoriatic Arthritis Patients in Relation to Response Assessed by the Dactylitis Severity Score

	Baseline DSS Mean ( $\pm$ SD)	Post Treatment DSS Mean ( $\pm$ SD)
<b>IL-6 –174</b>		
CC	7.0 ( $\pm$ 5.0)	3.0 ( $\pm$ 3.0)
GC	6.3 ( $\pm$ 2.1)	1.0 ( $\pm$ 1.4)
GG	8.0 ( $\pm$ 5.8)	3.0 ( $\pm$ 2.2)
<b>IL-6R Asp358Ala</b>		
AA	5.0 ( $\pm$ 1.7)	1.3 ( $\pm$ 1.3)
AC	7.5 ( $\pm$ 3.7)	1.7 ( $\pm$ 2.2)
CC	6.0 ( $\pm$ 0.0)	0.0 ( $\pm$ 0.0)

**Abbreviations:** DSS, dactylitis severity score; SD, standard deviation.

## Discussion

The IL-33 cytokine plays a crucial role in development of persistent inflammation, acting as a potent mediator of an inflammatory response. The IL-6 is involved in the pathogenesis of rheumatoid arthritis (RA) and psoriasis, disorders that share common pathobiology with PsA. The pathological role of IL-6 in RA development has been highlighted by studies conducted on murine arthritis models. Blockade of IL-6 signaling by gene knockout or administration of anti-IL-6 or anti-IL-6R antibodies leads to suppression of the disease development.<sup>56–58</sup> This beneficial effect has been confirmed on human organisms, resulting in the introduction of a humanized anti-IL-6 receptor monoclonal antibody Tocilizumab as a novel drug in RA therapy. Levels of circulating IL-6 are elevated both in RA and psoriasis as well as in numerous other autoimmune and chronic inflammatory diseases. It suggests that deregulation of signaling pathways mediated by IL-6 may contribute to autoimmune pathology.

The obtained results revealed an association of the IL-6 –174 G/C polymorphism with clinical outcome in PsA patients treated with MTX. Individuals carrying the IL-6 –174 G allele responded less effectively to the therapy than those with the IL-6 –174 CC genotype. These observations are in agreement with studies reporting an association of the IL-6 –174 G/C polymorphism with RA. The IL-6 –174 GG genotype was more common in RA patients displaying increased radiographic damage.<sup>26</sup> Also, the IL-6 –174 G allele was associated with susceptibility to juvenile RA and presence of the IL-6 –174 GG genotype correlated with higher pain scores in JIA patients.<sup>12</sup>

In accordance to our knowledge, only two studies to date investigated the possible impact of the IL-6 –174 G/C polymorphism in context of PsA.<sup>36,37</sup> In the study performed by Cubino et al the IL-6 –174 GG genotype was associated with a peripheral form of PsA.<sup>36</sup> However, in a study conducted by Balding et al no associations were observed, neither with disease susceptibility, nor clinical characteristics.<sup>37</sup> With respect to the possible influence of the IL-6 –174 G/C genetic variant on MTX efficacy, no study involving PsA patients has been performed to date. However, two groups examined the association of the IL-6 –174 G/C polymorphism with efficacy of the MTX treatment in patients with RA.<sup>33,34</sup> Pawlik et al reported significantly worse response to MTX therapy among patients possessing the IL-6 –174 GG genotype.<sup>33</sup> In addition, in a study by Ruiz et al the IL-6 –174 GG genotype was associated with a higher risk of failure in therapeutic response to MTX or leflunomide (lefl).<sup>34</sup> However, after stratifying patients' groups in accordance to MTX or lefl, a significant association was observed only in patients treated with lefl.

The results of the present study are concordant with previous reports concerning the Polish population in context of the role of the IL-6 –174 G/C polymorphism in the pathogenesis of immunological disorders.<sup>23,38,40</sup> In a study by Pawlik et al patients homozygous for the IL-6 –174 G allele were characterized with higher disease severity assessed by DAS28 score.<sup>23</sup> Also, the IL-6 –174 GG genotype was overrepresented in patients diagnosed with psoriasis in comparison to a control group and this genotype was associated with almost a twofold increased risk of psoriasis.<sup>38</sup> However, another study did not report any significant differences between Polish patients with psoriasis and healthy controls.<sup>42</sup> On the other hand, the association of the IL-6 –174 GG genotype with susceptibility to psoriasis and response to anti-TNF therapy was reported by di Renzo et al.<sup>39</sup> In line with these findings, in a study conducted by Boca et al the IL-6 –174 CC genotype was less frequent in the patients' group and conferred protection against psoriasis.<sup>41</sup>

This polymorphism was also implicated in systemic lupus erythematosus (SLE) pathogenesis and recent meta-analysis indicated that the IL-6 –174 G allele predisposes, particularly in the European population, to SLE development.<sup>59</sup> Furthermore, a significantly increased frequency of the IL-6 –174 GG genotype was observed in patients with diabetes mellitus.<sup>60,61</sup> Also, the same genotype correlated with disease activity in systemic sclerosis

patients.<sup>62</sup> On the other hand, no relationships were detected between the IL-6 -174 G/C polymorphism and development of RA or type I diabetes.<sup>63</sup> Data from the present study also did not expose any significant associations between this SNP and susceptibility to PsA.

Regarding the IL-6R Asp358Ala SNP, the results derived from the present study suggest a lack of association between this polymorphism and predisposition to PsA or response to MTX treatment. In agreement with these results, in previous studies on RA patients there were no significant relationships between the IL-6R Asp358Ala genetic variants and clinical outcome of therapy, neither with rituximab, tocilizumab, nor anti-TNF agents.<sup>35,42,43</sup> Also, no significant correlations were detected between this SNP and susceptibility to systemic sclerosis.<sup>48</sup> However, some studies reported an association of the IL-6R Asp358Ala allele C with increased risk of atopic dermatitis and decreased risk of RA.<sup>46,49</sup>

It should be noted that the relatively limited sample size constitutes a major limitation of the present study that could result in a statistical power insufficient to detect small or moderate effects. Therefore, further studies on larger patient cohorts are necessary to confirm findings of the present study.

In conclusion, this is the first report regarding the possible impact of IL-6 -174 G/C polymorphism on the efficacy of methotrexate treatment in PsA. The results from the present study suggest an involvement of the IL-6 -174 G/C polymorphism in pathomechanisms determining the PsA. The presence of the IL-6 -174 CC genotype among PsA patients predisposed to a favorable response to MTX therapy. With respect to the IL-6R rs2228145 polymorphism the obtained data did not reveal any significant associations with PsA susceptibility or clinical outcome of MTX treatment, indicating that this SNP may not be involved in the genetic background of PsA. However, the results demonstrated in this study require validation in larger cohorts as well as in populations of different ethnic origin.

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## Disclosure

The authors report no conflicts of interest in this work.

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