

IL-4 and IL-12 Polymorphisms are Associated with Response to Suplatast Tosilate, a Th2 Cytokine Inhibitor, in Patients with Atopic Dermatitis

Hideki Nagase¹, Yoshinori Nakachi¹, Keiji Ishida¹, Mamoru Kiniwa^{*1}, Satoshi Takeuchi², Ichiro Katayama³, Yoshinari Matsumoto⁴, Fukumi Furukawa⁵, Shin Morizane⁶, Sakae Kaneko⁷, Yoshiki Tokura⁸, Motoi Takenaka⁹, Yutaka Hatano¹⁰ and Yoshiki Miyachi¹¹

¹Tokushima Research Center, Taiho Pharmaceutical Co., Ltd., Tokushima, Japan

²Department of Dermatology, Graduate School of Medical Sciences, Kyushu University, Fukuoka, Japan

³Department of Dermatology, Osaka University Graduate School of Medicine, Osaka, Japan

⁴Department of Dermatology, Aichi Medical University School of Medicine, Aichi, Japan

⁵Department of Dermatology, Wakayama Medical University, Wakayama, Japan

⁶Department of Dermatology, Okayama University Graduate School of Medicine, Dentistry and Pharmaceutical Sciences, Okayama, Japan

⁷Department of Dermatology, Shimane University Faculty of Medicine, Shimane, Japan

⁸Department of Dermatology, University of Occupational and Environmental Health, Kitakyushu, Japan (Present affiliation: Department of Dermatology, Hamamatsu University School of Medicine, Hamamatsu, Japan)

⁹Department of Dermatology and Allergology, Nagasaki University Hospital, Nagasaki, Japan

¹⁰Department of Dermatology, Faculty of Medicine, Oita University, Oita, Japan

¹¹Department of Dermatology, Kyoto University Graduate School of Medicine, Kyoto, Japan

Abstract: Th2-related immune and inflammatory responses have been implicated in the pathogenesis of atopic dermatitis (AD), but few clinical lines of evidence have been reported regarding how and whether Th2-related responses are associated with other risk factors in the treatment of AD patients. In this study, the associations between the polymorphisms of genes related to the pathophysiology of AD and the efficacy of suplatast tosilate, an oral immunomodulator known to downregulate Th2-related allergic responses, were analyzed in adult patients with chronic AD. Patients were recruited from our previous study, where suplatast tosilate was evaluated for its efficacy when used in combination with topical steroids. The genotypes of 35 single nucleotide polymorphisms (SNPs) of 27 genes related to AD pathogenesis were then determined in 17 responders and 18 non-responders, as defined by the improvement rate in their AD skin scores. While no significant difference in the patient background was observed between responders and non-responders, significant associations were noted between the response to treatment with suplatast tosilate and three SNPs of IL-4 (-590C/T: $P=0.04$, -33C/T: $P=0.04$) and IL-12B (1188A/C: $P=0.03$), but not for the other SNPs. Of note, ethnic differences in the genotype frequencies of IL-4 -590C/T and IL-12B 1188A/C SNPs were found. In conclusion, the present results raise the possibility that AD patients who tend to produce more IL-4 and IL-12 may be susceptible to suplatast tosilate treatment and that ethnic variations should be considered to further understand the role of Th2-related responses.

Keywords: Atopic dermatitis, single nucleotide polymorphism, IL-4, IL-12B, Th1, Th2, suplatast tosilate, ethnic difference.

INTRODUCTION

AD is one of the most common chronic inflammatory diseases of the skin. While its causes and mechanisms have not yet been fully elucidated, its pathogenesis has been

characterized by altered skin barrier function and immune dysregulation. Recent genome-wide association studies or polymorphism analyses of candidate genes have further extended the view that patients with AD have genetically determined risk factors that affect the skin barrier and immune responses [1-6]. It should be noted, however, that few clinical lines of evidence have been reported regarding how these factors interact with each other and, more importantly, whether the down- or up-regulation of any gene or factor may affect the treatment of AD patients.

*Address correspondence to this author at the Tokushima Research Center, Taiho Pharmaceutical Co., Ltd., 224-2, Ebisuno Hiraishi, Kawauchi-cho, Tokushima, 771-0194, Japan; Tel: +81-88-665-6014; Fax: +81-88-665-6554; E-mail: makiniwa@taiho.co.jp

In the present study, we focused on suplatast tosilate, which has been approved in Japan as an oral immunomodulator for AD, allergic rhinitis, and asthma [7-10]. This agent has been used as an adjunct to or in combination with topical anti-inflammatory and immunosuppressive agents and has been shown to reduce the requirements for these agents [11, 12]. Suplatast tosilate has been referred to as a Th2 cytokine inhibitor because 1) it was discovered based on its capacity to suppress murine IgE formation without affecting cellular immune responses [13], 2) it suppressed IL-4 and IL-5 production by Th2 cells [14], 3) it suppressed allergic inflammation *via* indirect actions on Th2 cells as well as direct actions on eosinophils [15-19], and 4) it affected certain biomarkers such as Th1/Th2 ratio during the treatment of allergic asthmatic patients [20]. Although its mechanism of action remains to be identified at a molecular level, it is reasonable to assume that its efficacy in allergic patients may be ascribed to the modulation of the Th2-related immune and inflammatory responses.

We previously performed a clinical study investigating the efficacy of suplatast tosilate in adult chronic AD patients who also received topical steroids but had responded poorly to adjunct therapy such as antihistamines [21, 22]. This study revealed that suplatast tosilate markedly improved the AD skin symptom scores in a certain group of patients, but not in all patients, enabling us to recruit suplatast tosilate responders and non-responders and to analyze the polymorphism of genes related to the pathophysiology of AD retrospectively. The results implicated the efficacy of suplatast tosilate in the modulation of both Th1 and Th2 responses in AD therapy and suggested ethnic variations in the pathogenesis of AD.

MATERIALS AND METHODS

Study Subjects

One hundred and sixty-three patients with atopic dermatitis whose conditions had been poorly controlled by other anti-allergic drugs at 11 hospitals between 2004 and 2006 were enrolled in our previous study [21, 22]. The Th2 cytokine inhibitor suplatast tosilate (300 mg/day) was administered to these patients. After 4 weeks or more of treatment, the skin symptom score was determined to evaluate the usefulness of suplatast tosilate. The effectiveness of suplatast tosilate was evaluated by comparing the improvement rate relative to the baseline skin symptom score [21, 22]. To classify the patients into a highly effective group and a poorly effective group, we divided all the cases into three groups based on the improvement rate of the skin symptom scores. The one-third of cases with the highest improvement rates were defined as “responders”, and the one-third of cases with the lowest improvement rates were defined as “non-responders”. As a result, 54 “responder” patients had an improvement rate of more than 58%, and 53 “non-responder” patients had an improvement rate of less than 25%. From these patients, a total of 35 patients (17 responders and 18 non-responders) provided informed consent to undergo SNP genotyping using DNA from a blood sample. The improvement rate for the 17 high-responders with an improvement rate of more than 58% for their skin symptom scores was $68.8 \pm 12.2\%$. On the other hand, that of the 18 non-responders with an improvement

rate of less than 25% was $1.4 \pm 22.8\%$. This study was approved by the ethics committees of Kyoto University Graduate School of Medicine, Kyushu University, Osaka University Graduate School of Medicine, Aichi Medical University School of Medicine, Wakayama Medical University, Okayama University Graduate School of Medicine, Shimane University, University of Occupational and Environmental Health, Nagasaki University, Oita University and Taiho Pharmaceutical Co., Ltd.

SNP Genotyping Analysis

In each patient, 8.5 mL of blood was collected into a PAXgene Blood DNA tube (Qiagen Inc. CA, USA). DNA from the blood sample was purified using a PAXgene Blood DNA kit (Qiagen Inc.). The genotypes of 35 single nucleotide polymorphisms (SNPs) of 27 allergy-related genes were determined using a TaqMan real-time polymerase chain reaction (PCR) (TaqMan SNP Genotyping Assays; Life Technologies, CA, USA) or direct sequencing methods. The following 27 genes were investigated in this study: Th2 cytokines or cytokines promoting Th2 cells (IL-4, IL-5, IL-13, and TSLP), Th1 cytokine (IFN- γ), cytokines promoting Th1 cells (IL-12B and IL-18), proinflammatory cytokine (IL-17), chemokines (CCL5, CCL11, and CCL17), some of their receptors (IL-4R, IL-5R, IL-12R, IL-22R, IL-23R, and IFN- γ R), other genes related to the symptoms of AD or atopic disease (IL-31R, Chymase, NGF, NGFR, Fc ϵ R1 α , Fc ϵ R1 β , HRH1, HRH4, and LTC4S), and filaggrin (FLG). Thirty-one of the 35 SNPs were genotyped using the Applied Biosystems 7900HT Real-Time PCR System. The assay IDs of the 31 SNPs are shown in Table 1. The remaining four SNPs without a commercial TaqMan SNP genotyping assay were genotyped by direct sequencing at Takara Bio Inc. (Mie, Japan).

Statistical Analysis

The associations between the clinical parameters and the response to treatment with suplatast tosilate were evaluated using a likelihood ratio chi-square test for categorical data or the Student t-test for continuous data. The effect of suplatast tosilate on the clinical parameters was evaluated using a paired *t*-test. Hardy-Weinberg equilibrium of the genotyping results was evaluated using a Pearson's chi-square test. A likelihood ratio chi-square test was used to analyze the associations between the genotype of each SNP and the response to treatment with suplatast tosilate. All the analyses were performed using the statistical software JMP 7.0.1 and the SAS statistical package, version 9.1.3 (SAS Institute Inc., Cary, NC, USA). Differences were considered significant when $P < 0.05$. Because this was the study with a small sample size, a correction for multiple comparisons was not made.

RESULTS

Patient Characteristics and Effect of Suplatast Tosilate on Th2-Related Parameters

No significant differences between the responders and the non-responders were observed with regard to sex, age, and the pre-treatment values for the skin symptom scores, IgE levels or eosinophil counts (Table 2). Note that suplatast tosilate treatment did not affect the serum IgE level or the

Table 1. SNP list and Genotyping Methods

Gene Symbol	SNP name	rs ID	Methods	Assay ID
IL4	-590C/T	rs2243250	RT-PCR*	C__16176216_10
	-33C/T	rs2070874	RT-PCR	C__16176215_10
IL4R	Gln551Arg	rs1801275	RT-PCR	C__2351160_20
	Ile50Val	rs1805010	RT-PCR	C__2769554_10
IL5	-703C/T	rs2069812	RT-PCR	C__16274150_10
IL5RA	-80G/A	rs2290608	RT-PCR	C__15885096_10
IL12B	1188A/C	rs3212227	RT-PCR	C__2084293_10
IL12RB1	-2C/T	rs436857	RT-PCR	C__795468_1_
IL13	2044G/A	rs20541	RT-PCR	C__2259921_20
	-1112C/T	rs1800925	RT-PCR	C__8932056_10
IL17F	7488T/C	rs763780	RT-PCR	C__2234166_10
IL18	113T/G	rs360718	RT-PCR	C__2898461_10
IL22RA1	Arg518Gly	rs3795299	RT-PCR	C__440166_10
IL23R	Gln3His	rs1884444	RT-PCR	C__11728603_10
IL31RA	Ser497Asn	rs161704	RT-PCR	C__2839337_10
IFNG	874T/A	rs2430561	DS†	-
IFNGR2	Arg64Gln	rs9808753	RT-PCR	C__2443413_1_
CCL5	-28C/G	rs2280788	RT-PCR	C__15874396_20
	-403G/A	rs2107538	RT-PCR	C__15874407_10
CCL11	-384A/G	rs17809012	RT-PCR	C__2590323_10
CCL17	-431C/T	rs223828	RT-PCR	C__2392392_10
TSLP	-847C/T	rs3806933	RT-PCR	C__3166722_10
NGFB	Ala35Val	rs6330	RT-PCR	C__2525309_10
NGFR	Ser205Leu	rs2072446	RT-PCR	C__15870920_10
FCER1A	-66T/C	rs2251746	RT-PCR	C__1840470_20
MS4A2	-109C/T	rs1441586	RT-PCR	C__1842226_10
	Glu237Gly	rs569108	RT-PCR	C__900116_10
CMA1	-1897G/A	rs1800875	RT-PCR	C__2796262_10
HRH1	-17C/T	rs901865	RT-PCR	C__25471612_10
HRH4	Ala138Val	rs11665084	RT-PCR	C__3161321_20
LTC4S	-444A/C	rs730012	RT-PCR	C__644967_10
FLG	Tyr2194His	rs2184953	DS	-
	Tyr3105Asp	rs2065958	DS	-
	13347G/A	rs12730241	RT-PCR	C__31910001_10
	13586C/T	rs11204976	DS	-

*RT-PCR: reverse transcription polymerase chain reaction,

†DS: direct sequencing.

eosinophil count in either the responders or the non-responders (Table 3).

Genotype and Allele Frequencies

The genotyping results and the allele frequencies of the 35 SNPs are shown in Table 4. Among 34 SNPs (excluding

the Filaggrin Tyr3105Asp SNP), the genotype frequencies were the values expected under Hardy-Weinberg equilibrium, and no significant difference was observed between the minor allele frequencies reported in this study and those shown in the SNPper (<http://snpper.chip.org/bio/snpper-enter>) or NCBI databases (<http://www.ncbi.nlm.nih.gov/snp/>) (data not shown), indicating that the genotyping

Table 2. Clinical Characteristics of the Patients

Parameters	Responders (n=17)	Non-Responders (n=18)	P-Value*
Sex			
Male/Female	11/6	10/8	0.58
Age	37.1±17.5	32.3±12.6	0.36
(Range)	(18-79)	(13-62)	
Skin symptom score (Pre)	8.6±2.8	10.4±4.0	0.12
(Range)	(4-12)	(3-17)	
IgE level (Pre)	8645±8429	16356±21988	0.20
(IU/mL)			
Eosinophil count (Pre)	615±387	581±329	0.80
(cells/mm ³)			
Eosinophil count (Pre)	10.3±5.1	8.6±3.6	0.34
(%)			

*The P-values for categorical data and for continuous data were obtained using the likelihood ratio chi-square test and Student t-test, respectively.

Table 3. Effect of Suplatast Tosilate on Skin Symptom Score, Serum IgE Level, and Eosinophil Count in Atopic Dermatitis Patients

Parameters		N	Pre-Treatment	Post-Treatment	P-Value*
Skin symptom score	Responders	17	8.6±2.8	2.8±1.5	<0.001
	Non-responders	18	10.4±4.0	10.2±4.2	0.61
IgE level (IU/mL)	Responders	16	8645±8429	7935±8605	0.29
	Non-responders	14	16356±21988	15047±19624	0.09
Eosinophil (cells/mm ³)	Responders	15	615±387	591±306	0.72
	Non-responders	14	581±329	722±527	0.28
Eosinophil (%)	Responders	15	10.3±5.1	9.1±5.4	0.18
	Non-responders	13	8.6±3.6	10.4±6.4	0.29

*P-values were obtained using a paired t-test.

analysis was performed accurately. Regarding the genotyping results for the Filaggrin Tyr3105Asp SNP, the genotype frequency deviated from the values expected from Hardy-Weinberg equilibrium, and the minor allele frequency was significantly higher than that shown in the SNPper database (data not shown). This analysis may not have been performed accurately because the primer design region for the Filaggrin Tyr3105Asp SNP was highly homologous with other genes.

Associations between Genotype and Response to Treatment with Suplatast Tosilate

Significant associations between the response to treatment with suplatast tosilate and three SNPs of IL-4 (-590C/T: $P=0.04$, -33C/T: $P=0.04$) and IL-12B (1188A/C: $P=0.03$) were observed among the 35 SNPs genotyped in this study (Table 4). The genotyping results of two SNPs (-590C/T and -33C/T) within the IL-4 promoter region were perfectly identical among the 35 patients in this study. Therefore, the IL-4 -590C/T SNP, as a representative of the two SNPs of IL-4 that showed a significant association with the response, was used in further analyses. For the IL-4 -

590C/T SNP, patients with a C/C genotype had a significantly lower response rate to suplatast tosilate than those with a T/T or T/C genotype (0.0% vs 53.1%, $P=0.04$) (Table 5). On the other hand, for the IL-12B 1188A/C SNP, patients with an A/A genotype had a significantly lower response rate to suplatast tosilate than those with a C/C or C/A genotype (12.5% vs 59.3%, $P=0.02$, odds ratio=10.2) (Table 5). In a combination analysis of the two SNPs (IL-4 -590C/T and IL-12B 1188A/C), patients with either IL-4 -590C/T C/C or IL-12B 1188A/C A/A had a significantly lower response rate than those with other genotype combinations (10.0% vs 64.0%, $P=0.002$, odds ratio=16.0) (Table 5).

Ethnic Differences in Genotype Frequency of IL-4 -590C/T and IL-12B 1188A/C SNPs

The genotype frequencies of IL-4 -590C/T T/T and IL-12B 1188A/C C/C, which were associated with a strong response to suplatast tosilate, were significantly higher among Asians than among Europeans based on the results of the HapMap project (Table 6, <http://www.ncbi.nlm.nih.gov/snp/>).

Table 4. Genotype Frequency and Association Between Genotype and Response to Treatment with Suplatast Tosilate

SNP Name (SNP ID)	Genotype	Frequency (%)		P- Value*	Frequency (%)				P- Value†
		Total (n=35)			Responders (n=17)		Non-Responders (n=18)		
IL4 -590C/T (rs2243250)	T/T	13	(37%)	0.28	9	(53%)	4	(22%)	0.04
	T/C	19	(54%)		8	(47%)	11	(61%)	
	C/C	3	(9%)		0	(0%)	3	(17%)	
IL4 -33C/T (rs2070874)	T/T	13	(37%)	0.28	9	(53%)	4	(22%)	0.04
	T/C	19	(54%)		8	(47%)	11	(61%)	
	C/C	3	(9%)		0	(0%)	3	(17%)	
IL4R Gln551Arg (rs1801275)	A/A	30	(86%)	0.65	13	(76%)	17	(94%)	0.12
	A/G	5	(14%)		4	(24%)	1	(6%)	
	G/G	0	(0%)		0	(0%)	0	(0%)	
IL4R Ile50Val (rs1805010)	G/G	16	(46%)	0.72	7	(41%)	9	(50%)	0.67
	G/A	16	(46%)		9	(53%)	7	(39%)	
	A/A	3	(9%)		1	(6%)	2	(11%)	
IL5 -703C/T (rs2069812)	T/T	13	(37%)	0.49	8	(47%)	5	(28%)	0.49
	T/C	15	(43%)		6	(35%)	9	(50%)	
	C/C	7	(20%)		3	(18%)	4	(22%)	
IL5RA -80G/A (rs2290608)	G/G	21	(60%)	0.87	12	(71%)	9	(50%)	0.17
	G/A	12	(34%)		5	(29%)	7	(39%)	
	A/A	2	(6%)		0	(0%)	2	(11%)	
IL12B 1188A/C (rs3212227)	C/C	8	(23%)	0.61	6	(35%)	2	(11%)	0.03
	C/A	19	(54%)		10	(59%)	9	(50%)	
	A/A	8	(23%)		1	(6%)	7	(39%)	
IL12RB1 -2C/T (rs436857)	C/C	24	(69%)	0.27	12	(71%)	12	(67%)	0.80
	C/T	11	(31%)		5	(29%)	6	(33%)	
	T/T	0	(0%)		0	(0%)	0	(0%)	
IL13 2044G/A (rs20541)	G/G	15	(43%)	0.93	7	(41%)	8	(44%)	0.51
	G/A	16	(46%)		7	(41%)	9	(50%)	
	A/A	4	(11%)		3	(18%)	1	(6%)	
IL13 -1112C/T (rs1800925)	C/C	20	(57%)	0.95	11	(65%)	9	(50%)	0.65
	C/T	13	(37%)		5	(29%)	8	(44%)	
	T/T	2	(6%)		1	(6%)	1	(6%)	
IL17F 7488T/C (rs763780)	T/T	27	(77%)	0.45	13	(76%)	14	(78%)	0.93
	T/C	8	(23%)		4	(24%)	4	(22%)	
	C/C	0	(0%)		0	(0%)	0	(0%)	
IL18 113T/G (rs360718)	T/T	26	(74%)	0.69	14	(82%)	12	(67%)	0.37
	T/G	8	(23%)		3	(18%)	5	(28%)	
	G/G	1	(3%)		0	(0%)	1	(6%)	
IL22RA1 Arg518Gly (rs3795299)	G/G	19	(54%)	0.08	10	(59%)	9	(50%)	0.60
	G/C	16	(46%)		7	(41%)	9	(50%)	
	C/C	0	(0%)		0	(0%)	0	(0%)	

(Table 4) contd.....

SNP Name (SNP ID)	Genotype	Frequency (%)		P- Value*	Frequency (%)		P- Value†		
		Total (n=35)			Responders (n=17)	Non-Responders (n=18)			
IL23R	T/T	11	(31%)		4	(24%)	7	(39%)	
Gln3His	T/G	20	(57%)		10	(59%)	10	(56%)	
(rs1884444)	G/G	4	(11%)	0.26	3	(18%)	1	(6%)	0.40
IL31RA	G/G	9	(26%)		6	(35%)	3	(17%)	
Ser497Asn	G/A	17	(49%)		6	(35%)	11	(61%)	
(rs161704)	A/A	9	(26%)	0.87	5	(29%)	4	(22%)	0.27
IFNG 874T/A	A/A	24	(69%)		12	(71%)	12	(67%)	
(rs2430561)	A/T	9	(26%)		5	(29%)	4	(22%)	
	T/T	2	(6%)	0.38	0	(0%)	2	(11%)	0.24
IFNGR2	A/A	10	(29%)		3	(18%)	7	(39%)	
Arg64Gln	A/G	18	(51%)		9	(53%)	9	(50%)	
(rs9808753)	G/G	7	(20%)	0.83	5	(29%)	2	(11%)	0.23
CCL5 -28C/G	C/C	26	(74%)		12	(71%)	14	(78%)	
(rs2280788)	C/G	8	(23%)		5	(29%)	3	(17%)	
	G/G	1	(3%)	0.69	0	(0%)	1	(6%)	0.37
CCL5 -403G/A	C/C	16	(46%)		9	(53%)	7	(39%)	
(rs2107538)	C/T	14	(40%)		7	(41%)	7	(39%)	
	T/T	5	(14%)	0.51	1	(6%)	4	(22%)	0.34
CCL11	A/A	17	(49%)		9	(53%)	8	(44%)	
-384A/G	A/G	15	(43%)		6	(35%)	9	(50%)	
(rs17809012)	G/G	3	(9%)	0.90	2	(12%)	1	(6%)	0.61
CCL17	C/C	11	(31%)		8	(47%)	3	(17%)	
-431C/T	C/T	14	(40%)		5	(29%)	9	(50%)	
(rs223828)	T/T	10	(29%)	0.24	4	(24%)	6	(33%)	0.14
TSLP -847C/T	C/C	20	(57%)		11	(65%)	9	(50%)	
(rs3806933)	C/T	11	(31%)		4	(24%)	7	(39%)	
	T/T	4	(11%)	0.22	2	(12%)	2	(11%)	0.61
NGFB	C/C	17	(49%)		7	(41%)	10	(56%)	
Ala35Val	C/T	15	(43%)		7	(41%)	8	(44%)	
(rs6330)	T/T	3	(9%)	0.90	3	(18%)	0	(0%)	0.09
NGFR	C/C	29	(83%)		16	(94%)	13	(72%)	
Ser205Leu	C/T	5	(14%)		1	(6%)	4	(22%)	
(rs2072446)	T/T	1	(3%)	0.22	0	(0%)	1	(6%)	0.17
FCER1A	T/T	30	(86%)		16	(94%)	14	(78%)	
-66T/C	T/C	5	(14%)		1	(6%)	4	(22%)	
(rs2251746)	C/C	0	(0%)	0.65	0	(0%)	0	(0%)	0.15
MS4A2	T/T	19	(54%)		11	(65%)	8	(44%)	
-109C/T	T/C	13	(37%)		5	(29%)	8	(44%)	
(rs1441586)	C/C	3	(9%)	0.72	1	(6%)	2	(11%)	0.48
MS4A2	T/T	25	(71%)		14	(82%)	11	(61%)	
Glu237Gly	T/C	10	(29%)		3	(18%)	7	(39%)	
(rs569108)	C/C	0	(0%)	0.32	0	(0%)	0	(0%)	0.16

(Table 4) contd.....

SNP Name (SNP ID)	Genotype	Frequency (%)		P- Value*	Frequency (%)		P- Value†
		Total (n=35)			Responders (n=17)	Non-Responders (n=18)	
CMA1 -1897G/A (rs1800875)	G/G	17 (49%)		0.48	7 (41%)	10 (56%)	0.69
	G/A	16 (46%)			9 (53%)	7 (39%)	
	A/A	2 (6%)			1 (6%)	1 (6%)	
HRH1 -17C/T (rs901865)	G/G	31 (89%)		0.72	15 (88%)	16 (89%)	0.95
	G/A	4 (11%)			2 (12%)	2 (11%)	
	A/A	0 (0%)			0 (0%)	0 (0%)	
HRH4 Ala138Val (rs11665084)	C/C	31 (89%)		0.72	16 (94%)	15 (83%)	0.31
	C/T	4 (11%)			1 (6%)	3 (17%)	
	T/T	0 (0%)			0 (0%)	0 (0%)	
LTC4S -444A/C (rs730012)	A/A	21 (60%)		0.54	11 (65%)	10 (56%)	0.35
	A/C	13 (37%)			5 (29%)	8 (44%)	
	C/C	1 (3%)			1 (6%)	0 (0%)	
FLG Tyr2194His (rs2184953)	C/C	9 (26%)		0.20	4 (24%)	5 (28%)	0.85
	C/T	21 (60%)			10 (59%)	11 (61%)	
	T/T	5 (14%)			3 (18%)	2 (11%)	
FLG Tyr3105Asp (rs2065958)	G/G	9 (26%)		<0.001	4 (24%)	5 (28%)	0.77
	G/T	26 (74%)			13 (76%)	13 (72%)	
	T/T	0 (0%)			0 (0%)	0 (0%)	
FLG 13347G/A (rs12730241)	A/A	9 (26%)		0.20	4 (24%)	5 (28%)	0.85
	A/G	21 (60%)			10 (59%)	11 (61%)	
	G/G	5 (14%)			3 (18%)	2 (11%)	
FLG 13586C/T (rs11204976)	A/A	9 (26%)		0.20	4 (24%)	5 (28%)	0.85
	A/G	21 (60%)			10 (59%)	11 (61%)	
	G/G	5 (14%)			3 (18%)	2 (11%)	

*P-values were obtained using the Hardy Weinberg equilibrium (HWE) test.

†P-values were obtained using the likelihood ratio chi-square test.

Table 5. Association Between Genotype Combinations and Response to Treatment with Suplatast Tosilate

IL4 -590C/T	IL12B 1188A/C	Response Rate, %	P-Value*	Odds Ratio (95% CI)
T/T, T/C		53.1 (17/32)	0.04	-
C/C		0.0 (0/3)		
	C/C, C/A	59.3 (16/27)		
	A/A	12.5 (1/8)	0.02	(1.1-94.8)
T/T, T/C and	C/C, C/A	64.0 (16/25)	0.002	16.0
C/C or	A/A	10.0 (1/10)		

*P-values were obtained using the likelihood ratio chi-square test.

DISCUSSION

AD is increasingly recognized as a complex disease, since multifunctional cells and factors interact with each other in its pathogenesis. In the present study, based on the down-regulation of the Th2 inflammatory response by suplatast tosilate, we performed a pilot study to implicate Th2 suppression in the pathogenesis of AD by

retrospectively analyzing the polymorphisms of genes reportedly associated with AD or atopic diseases. To this end, patients were recruited from our previous study, in which the efficacy of suplatast tosilate was investigated in adult AD patients whose conditions had been poorly controlled by treatment with topical steroids and adjunct anti-allergic agents, including antihistamines. We then examined the association of suplatast tosilate efficacy and 35

Table 6. Distribution of IL-4 -590C/T and IL-12B 1188A/C Genotypes Among Japanese and European Populations

SNP	Response to Suplatast Tosilate	Japanese		P-Value† (vs Present Study)	European		P-Value† (vs Present Study)
		Present Study (n=35)	HapMap project* (n=85,86)		HapMap Project* (n=113)		
IL-4 -590C/T							
T/T	high	13 (37%)	45 (53%)	0.26	2 (2%)	<0.001	
T/C	medium	19 (54%)	33 (39%)		27 (24%)		
C/C	low	3 (9%)	7 (8%)		84 (74%)		
IL-12 1188A/C							
C/C	high	8 (23%)	28 (33%)	0.56	1 (1%)	<0.001	
C/A	medium	19 (54%)	40 (47%)		41 (36%)		
A/A	low	8 (23%)	18 (21%)		71 (63%)		

*<http://www.ncbi.nlm.nih.gov/snp/>

†P-values were obtained using the likelihood ratio chi-square test.

SNPs of 27 allergy-related genes. Of note, no significant difference was seen in the minor allele frequencies of each SNP, with the exception of one SNP (Filaggrin Tyr3105Asp) with a low reliability, based on a comparison with an SNP control database for the Japanese population (SNPper or NCBI). These findings suggest that no genetic polymorphisms associated with susceptibility to AD were observed among the SNPs investigated in this study.

The results of the present SNP genotyping analysis clearly indicated that suplatast tosilate efficacy was associated with IL-4 and IL-12B polymorphisms, but not with any of the others, indicating that patients with a T/T or T/C genotype for the IL-4 -590C/T SNP as well as a C/C or C/A genotype for the IL-12B 1188A/C SNP responded to suplatast tosilate treatment at a significantly higher rate. Moreover, patients with both the IL-4 -590 T/T or T/C genotype and the IL-12B 1188 C/C or C/A genotype had a significantly higher response rate than those with other genotype combinations. Of note, SNPs in the regulatory sequences of genes are associated with the varying production of relevant cytokines. In fact, the promoter sequence of IL-4 -590T was reported to show a greater binding to nuclear transcription factors than that of -590C [23, 24]. Nakashima *et al.*, reported that PBMC with a -590T/T or -590T/C genotype produced higher levels of IL-4 than those with a -590C/C genotype [25]. In general, IL-4 is a major Th2 cytokine that plays an essential role in the class-switching of B cells to IgE-producing cells, Th2 cell differentiation, and the initial phase of tissue inflammation during the Th2-dominant phase of atopic diseases. Recently, IL-4 has been also reported to play an important role in regulating skin homeostasis and innate barrier function in AD lesions [26]. In fact, Burchard *et al.*, has reported that the sequence variant in the IL-4 promoter region is associated with the asthma FEV1 (Forced expiratory volume in one second) [27]. It is reasonable, therefore, to assume that patients with the IL-4 -590T allele tend to develop a Th2-dominated immune response, leading to the susceptibility of these patients to suplatast tosilate which down-regulates Th2-related responses including skin manifestations.

Suplatast tosilate has been reported to suppress IgE formation and eosinophil counts presumably through inhibition of Th2 cytokine in basic experiments and in several clinical studies in patients with atopic asthma [13-20]. However, no significant decreases in Th2-related parameters, such as the total

IgE level and the eosinophil count, were observed after suplatast tosilate treatment even among the responders in this study. The reason for this contradiction is clearly unknown, but, as a possibility, a Th2-dependency may be decreased during the chronic phase of AD by the effect such as infections. In this sense, of great interest is the observation that patients with the IL-12B -1188 C/C genotype responded to suplatast tosilate treatment at a high rate. The IL-12B -1188 C allele was reported to result in IL-12B mRNA with a lower transcriptional activity and stability than that for the A allele [28]. Although IL-12 p40 encoded by IL-12B is a component of IL-12 (a p40 and p35 heterodimer) [29], the IL-12 p40 homodimer was reported to function as an antagonist of IL-12 action [30, 31]. In fact, PBMC with a -1188IL-12B C/C genotype having a high response rate to suplatast tosilate reportedly produced significantly higher levels of biologically active IL-12 upon stimulation with LPS or PPD than those of other genotypes with the A allele [32]. IL-12 is known to induce IFN- γ production [33], to suppress IgE synthesis [34], and to promote Th1 cell maturation [33]. The results, therefore, suggest that the efficacy of suplatast tosilate is associated with the Th1-immune response that also underlies the AD symptoms. This finding is partly in accordance with an observation by Matsui *et al.*, [35], who reported that PBMC from suplatast tosilate responders with childhood asthma produced higher amounts of IFN- γ than those of suplatast tosilate non-responders. The findings are also more directly in accordance with those by Murakami *et al.*, [36], who found that suplatast tosilate treatment significantly suppressed the elevated expression of IL-4, IL-5, and IFN- γ mRNA in caspase-1 transgenic mice that spontaneously developed AD-like dermatitis. They also observed that suplatast tosilate treatment significantly decreased the expression of IL-18, which induced Th1 responses in synergy with IL-12, possibly explaining the mechanism of suplatast tosilate efficacy for Th1-related responses.

The present results suggest that both Th2- and Th1-related responses may play more important roles in AD patients who responded to suplatast tosilate treatment than other factors, and ethnic differences may exist in the mechanism of chronic AD, since the allele frequencies of the IL-4 and IL-12B polymorphisms associated with the efficacy of suplatast tosilate differ significantly in the Japanese population compared with the European population. It should be noted, however, that these

data were based on a retrospective analysis of a small sample size. Therefore, these results have to be confirmed in a large-scale prospective study. Moreover, further large-scale investigations taking ethnic differences into account are also needed to clarify the exact mechanism of chronic AD.

CONFLICT OF INTEREST

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