

Relationship between the benefits of suplatast tosilate, a Th2 cytokine inhibitor, and gene polymorphisms in children with bronchial asthma

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Abstract. Although currently available antiasthmatic drugs are effective for many patients with bronchial asthma, some patients do not respond well to medications or exhibit more frequent adverse effects compared to other patients. Antiasthmatic treatment should be tailored individually according to the predispositions and pathophysiological conditions of patients. No reports have been made concerning the relationships between the effects of Th2 cytokine inhibitors and gene polymorphisms. The present study was therefore performed to investigate the relationships between gene polymorphisms known to be involved in allergy and cytokine production in peripheral blood mononuclear cells and the clinical efficacy of suplatast tosilate, a Th2 cytokine inhibitor, to clarify factors determining responses to treatment. A total of 20 children were enrolled in the study. The children were enrolled in a run-in period of 2 weeks and then received suplatast tosilate orally for 8 weeks. The children or their parents were instructed to keep an asthma diary to record changes in signs/symptoms of bronchial asthma before and after treatment. Concentrations of interferon (IFN)- γ and interleukin (IL)-4 in the supernatant were determined using ELISA methods. Using the invader assay method, the genotypes of polymorphisms of the genes were determined. Treatment with suplatast tosilate was more effective in children without the -444 A/C polymorphism of the LTC4 synthase gene and in children without the IL-13 variant R110Q. In children who responded well, production of IFN- γ was significantly increased after treatment. In this study, responses to suplatast tosilate were associated with SNPs of

the LTC4 synthase and IL-13 gene as well as change in the production of IFN- γ before and after drug administration.

Introduction

Allergic diseases such as bronchial asthma and atopic dermatitis develop as a result of the interaction of genetic and environmental factors (1). Several causative candidate genes for allergic diseases have been reported (2-6). Recently, treatment guidelines for bronchial asthma have been published and are utilized in clinical practice. Although antiasthmatic drugs are often effective in many patients with bronchial asthma, some patients may experience adverse effects on non-target organs and may exhibit lack of improvement due to the use of drugs inappropriate for their individual condition.

Although studies have shown that the effects of β_2 -adrenergic agonists (7), leukotriene receptor antagonists (8,9), and inhaled corticosteroids (10) are closely associated with gene polymorphisms, no reports have been made concerning the relationships between the effects of Th2 cytokine inhibitors and gene polymorphisms. In the present study, we investigated the relationships between the effects of suplatast tosilate, a Th2 cytokine inhibitor, and specific genetic polymorphisms and cytokines that are significantly associated with the presence/absence of effects of this drug.

Subjects and methods

A total of 20 children aged 3 to 17 years (14 boys and 6 girls) were enrolled in the study. After obtaining informed consent for participation in the study from the children or their parents, the children were enrolled in a run-in period of 2 weeks, after which blood samples were obtained. The children then took suplatast tosilate orally (6 mg/kg/day) for 8 weeks, and blood samples were taken at the end of the treatment period. Fig. 1 illustrates the study protocol. Before blood collection, the children or their parents received a full explanation of the design, methods, and expected results of the study and consented to blood sampling.

Evaluation of improvement of clinical signs/symptoms of bronchial asthma. The children or their parents were

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Key words: Th2 cytokine inhibitors, gene polymorphism, allergy, cytokine production, interferon- γ , interleukin-4, LTC4 synthase gene, interleukin-13

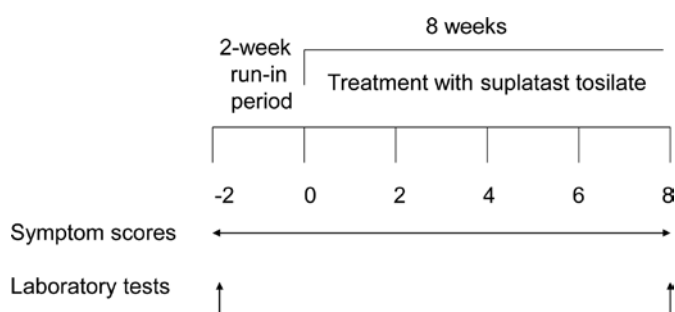


Figure 1. Study protocol. After an initial 2-week run-in period, suplastast tosilate was administered for 8 weeks.

instructed to keep an asthma diary to record changes in signs/symptoms of bronchial asthma before and after treatment. Major, moderate, and minor asthmatic attacks were scored as 9, 6, and 3 points, respectively, and cough as 1 point. Total symptom scores during the 2-week run-in period and the last two weeks of the 8-week treatment period were calculated and compared. Responders and non-responders were defined as children with and without a decrease in symptom score after treatment, respectively. No antiasthmatic drugs other than the study drug were administered during the run-in and observation periods.

Determination of eosinophil count and serum IgE level. Using aliquots of the blood samples obtained before and after treatment, the eosinophil count and serum IgE level were determined.

Determination of production of cytokines by peripheral blood mononuclear cells (PBMCs). PBMCs were isolated from heparinized blood samples using gradient centrifugation in Ficoll-Paque (Pharmacia, Uppsala, Sweden) and suspended at a density of 10^6 cells/ml in RPMI-1640 medium supplemented with 10% heat-inactivated fetal calf serum, 2 mmol L-glutamine, 100 U/ml penicillin and 100 μ g/ml streptomycin. PBMCs were cultured with 10 μ g/ml phytohemagglutinin (PHA) (Gibco BRL, Grand Island, NY, USA) for 24 h at 37°C in the presence of 5% CO₂. After incubation, the concentrations of interferon (IFN)- γ and interleukin (IL)-4 in the supernatant were determined using enzyme-linked immunosorbent assay (ELISA) methods. IFN- γ concentrations were measured with a human IFN- γ ELISA kit (Ohtsuka, Tokyo, Japan); the limit of detection of which was 15.6-1000 pg/ml. IL-4 concentrations were measured with a human IL-4 US ELISA kit (BioSource International, Camarillo, CA, USA); the limit of detection of which was 0.39-25.0 pg/ml.

DNA isolation evaluation of gene polymorphisms. Genomic DNA was extracted from neutrophils with a SepaGene kit (Sankyo Junyaku Co., Ltd., Tokyo, Japan). Using the invader assay method, the genotypes of polymorphisms of genes known to affect the production of IgE and proinflammatory mediators were determined (Table I).

Statistical analysis. Relationships of symptom scores with eosinophil count, serum IgE level, cytokine concentrations,

Table I. Genotype and allele frequencies.

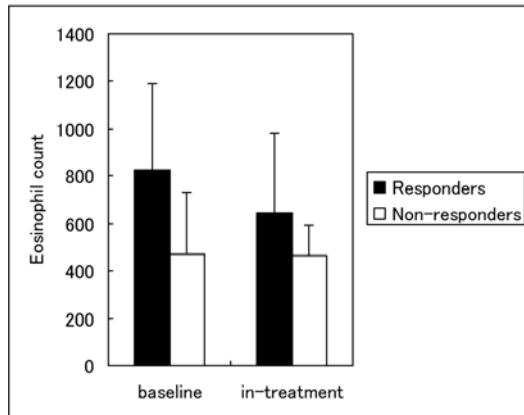
SNP ID	Frequency	%	HWE test
IL-12Rβ1 (c.365T>C)			
Genotype			
T/T	7	35.0	p=0.852
T/C	10	50.0	
C/C	3	15.0	
Allele			
T	24	60.0	
C	16	40.0	
IL-12B (c.3757T>C)			
Genotype			
T/T	6	30.0	p=0.964
T/C	10	50.0	
C/C	4	20.0	
Allele			
T	22	55.0	
C	18	45.0	
IL-12B (c.16078C>T)			
Genotype			
C/C	18	90.0	p=0.004
C/T	1	5.0	
T/T	1	5.0	
Allele			
C	37	92.5	
T	3	7.5	
IL-4Rα (Ile50Val)			
Genotype			
G/G	7	35.0	p=0.154
G/A	12	60.0	
A/A	1	5.0	
Allele			
G	26	65.0	
A	14	35.0	
LTC4S (Glu4Lys)			
Genotype			
G/G	19	95.0	p=0.909
G/A	1	5.0	
A/A	0	0.0	
Allele			
G	39	97.5	
A	1	2.5	
LTC4S (-444A>C)			
Genotype			
A/A	11	55.0	p=0.194
A/C	9	45.0	
C/C	0	0.0	
Allele			
A	31	77.5	
C	9	22.5	
IL-13 (Arg110Gln)			
Genotype			
G/G	9	45.0	p=0.909
G/A	9	45.0	
A/A	2	10.0	
Allele			
G	27	67.5	
A	13	32.5	

Table II. Asthma symptom scores before and after treatment with suplatast tosilate.

	Responders (n=13)	Non-responders (n=7)	p-value
Asthma scores (pre)	13.0±7.5	7.2±12.9	0.1980
Asthma scores (post)	1.8±2.4	9.5±12.9	0.0580
Asthma scores (Δ change)	-11.8±6.7	2.3±3.7	0.0002

P-values were obtained by the t-test.

A



B

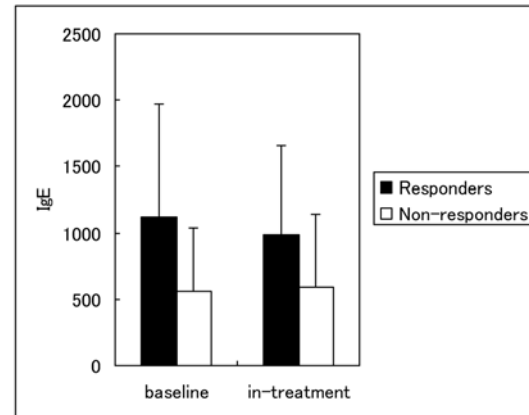


Figure 2. (A) Comparison of the eosinophil count. No significant differences between the responders and non-responders were observed for baseline or in-treatment values or the change in eosinophil count between the baseline and the treatment period. (B) Comparison of the serum IgE level. No significant differences between the responders and non-responders were observed for baseline or in-treatment values or the change in serum IgE level between the baseline and the treatment period.

Table III. Clinical characteristics of the patients.

	Responders (n=13)	Non-responders (n=7)	p-value
Gender (male/female)	8/5	6/1	0.261
Age (years)	7.8±3.8	6.9±3.2	0.563
Range	3-17	3-12	
Duration of asthma (years)	4.1±2.4	4.6±3.7	0.718
Range	1-8	1-10	
Asthma severity			
Mild	8	5	0.658
Moderate	5	2	
Atopic dermatitis			
+	6	2	0.444
-	7	5	
Rhinitis			
+	5	2	0.658
-	8	5	
Asthma type			
Atopic asthma	13	5	0.127
Infectious asthma	0	1	
Mixed asthma	0	1	
Asthma scores (pre)	13.6±7.5	7.2±12.9	0.198

P-values for categorical data were obtained by the Chi-square test; p-values for other data were obtained by the t-test.

and genotypes of SNPs tested were examined using the Chi-square test or t-test.

Results

Evaluation of improvement in clinical symptoms of bronchial asthma. Responses to treatment with suplatast tosilate were assessed based on the descriptions in the asthma diary kept by the participants or their parents during the 2-week run-in period and the 8-week treatment period (Table II). The symptom score during treatment was decreased from baseline in 13 patients (8 boys and 5 girls, mean age of 7.8±3.8 years, referred to as the 'responders') and was unchanged or increased in 7 patients (6 boys and 1 girl, mean age of 6.9±3.2 years, referred to as the 'non-responders'). No significant differences in age, severity of asthma, or prevalence of complications were observed between the responders and non-responders (Table III).

Relationships of the efficacy of treatment with the eosinophil count and serum IgE level. Comparisons of the eosinophil count and serum IgE level were carried out between the responders and non-responders to evaluate the relationships of these parameters with treatment efficacy. As shown in Fig. 2A and B, no significant differences were observed in baseline or in-treatment values or the change in these values between the baseline and the treatment period between the responders and non-responders.

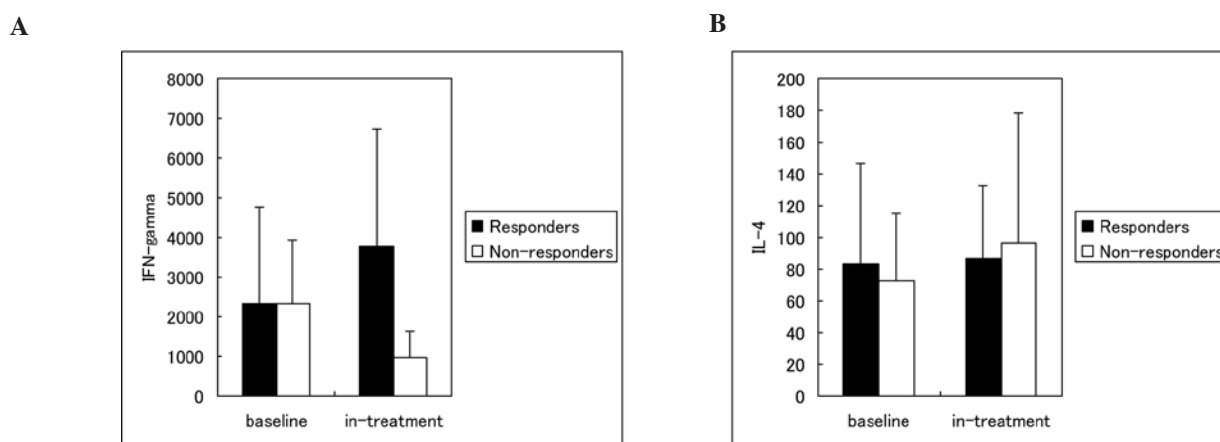


Figure 3. (A) Relationships of treatment efficacy with the amounts of INF- γ produced in peripheral blood mononuclear cells before and after treatment were investigated. The change in INF- γ production between the baseline and treatment periods was significantly greater in the responders than in the non-responders ($p=0.011$). (B) Relationships of treatment efficacy with the amounts of IL-4 produced in peripheral blood mononuclear cells before and after treatment were investigated. Production of IL-4 did not differ significantly between the responders and non-responders.

Table IV. Association between genotype and response to treatment with suplastast tosilate.

	Total no.	Genotype				Allele		
		n (%)	n (%)	n (%)	p-value	n (%)	n (%)	p-value
IL12R β 1(c.365T>C)		T/T	T/C	C/C	0.384	T	C	0.279
Responders	13	4 (30.8)	6 (46.2)	3 (23.0)		14 (53.9)	12 (46.1)	
Non-responders	7	3 (42.9)	4 (57.1)	0 (0.0)		10 (71.4)	4 (28.6)	
IL-12B (c.3757T>C)		T/T	T/C	C/C	0.207	T	C	0.386
Responders	13	4 (30.8)	5 (38.4)	4 (30.8)		13 (50.0)	13 (50.0)	
Non-responders	7	2 (28.6)	5 (71.4)	0 (0.0)		9 (64.3)	5 (35.7)	
IL-12B (c.16078C>T)		C/C	C/T	T/T	0.299	C	T	0.950
Responders	13	12 (92.3)	0 (0.0)	1 (7.7)		24 (92.3)	2 (7.7)	
Non-responders	7	6 (85.7)	1 (14.3)	0 (0.0)		13 (92.9)	1 (7.1)	
IL-4R α (Ile50Val)		G/G	G/A	A/A	0.690	G	A	0.532
Responders	13	4 (30.8)	8 (61.5)	1 (7.7)		16 (61.5)	10 (38.5)	
Non-responders	7	3 (42.9)	4 (57.1)	0 (0.0)		10 (71.4)	4 (28.6)	
LTC4S (Glu4Lys)		G/G	G/A	A/A	0.452	G	A	0.457
Responders	13	12 (92.3)	1 (7.7)	0 (0.0)		25 (96.2)	1 (3.8)	
Non-responders	7	7 (100)	0 (0.0)	0 (0.0)		14 (100)	0 (0.0)	
LTC4S (-444A>C)		A/A	A/C	C/C	0.007	A	C	0.024
Responders	13	10 (76.9)	3 (23.1)	0 (0.0)		23 (88.5)	3 (11.5)	
Non-responders	7	1 (14.3)	6 (85.7)	0 (0.0)		8 (57.1)	6 (42.9)	
IL-13 (Arg110Gln)		G/G	G/A	A/A	0.127	G	A	0.083
Responders	13	8 (61.5)	4 (30.8)	1 (7.7)		6 (23.1)	20 (76.9)	
Non-responders	7	1 (14.3)	5 (71.4)	1 (14.3)		7 (50.0)	7 (50.0)	

Relationships between cytokine production and efficacy of treatment. Relationships of treatment efficacy with the amounts of INF- γ and IL-4 produced by peripheral blood mononuclear cells before and after treatment were investigated (Fig. 3A and

B). The change in INF- γ production between the baseline and treatment periods was significantly larger in the responders than in the non-responders ($p=0.011$). Production of IL-4 did not differ significantly between the responders and non-responders.

Table V. Association between genotype combination and response to treatment with suplatast tosilate.

	Total no.	Genotype combination (1)			Genotype combination (2)		
		n (%)	n (%)	p-value	n (%)	n (%)	p-value
IL12Rβ1 (c.365T>C)		T/T	T/C, C/C	0.589	T/T, T/C	C/C	0.168
Responders	13	4 (30.8)	9 (69.2)		10 (76.9)	3 (23.1)	
Non-responders	7	3 (42.9)	4 (57.1)		7 (100.0)	0 (0.0)	
IL-12B (c.3757T>C)		T/T	T/C, C/C	0.919	T/T, T/C	C/C	0.101
Responders	13	4 (30.8)	9 (69.2)		9 (69.2)	4 (30.8)	
Non-responders	7	2 (28.6)	5 (71.4)		7 (100.0)	0 (0.0)	
IL-12B (c.16078C>T)		C/C	C/T, T/T	0.639	C/C, C/T	T/T	0.452
Responders	13	12 (92.3)	1 (7.7)		12 (92.3)	1 (7.7)	
Non-responders	7	6 (85.7)	1 (14.3)		7 (100.0)	0 (0.0)	
IL-4Rα (Ile50Val)		G/G	G/A, A/A	0.589	G/G, G/A	A/A	0.452
Responders	13	4 (30.8)	9 (69.2)		12 (92.3)	1 (7.7)	
Non-responders	7	3 (42.9)	4 (57.1)		7 (100.0)	0 (0.0)	
LTC4S (Glu4Lys)		G/G	G/A, A/A	0.452	G/G, G/A	A/A	-
Responders	13	12 (92.3)	1 (7.7)		13 (100.0)	0 (0.0)	
Non-responders	7	7 (100)	0 (0.0)		7 (100.0)	0 (0.0)	
LTC4S (-444A>C)		A/A	A/C, C/C	0.007	A/A, A/C	C/C	-
Responders	13	10 (76.9)	3 (23.1)		13 (100.0)	0 (0.0)	
Non-responders	7	1 (14.3)	6 (85.7)		7 (100.0)	0 (0.0)	
IL-13 (Arg110Gln)		G/G	G/A, A/A	0.043	G/G, G/A	A/A	0.639
Responders	13	8 (61.5)	5 (38.5)		12 (92.3)	1 (7.7)	
Non-responders	7	1 (14.3)	6 (85.7)		6 (85.7)	1 (14.3)	

Relationships between SNPs and treatment efficacy. Using the invader assay method (11), the genotypes of SNPs of genes known to affect the production of IgE [interleukin (IL)-12, IL-12 receptor (IL-12R)β1, IFN-γR1, IL-4Rα and IL-13 genes] and mediators (LTC4 synthase gene) were determined in the children enrolled in the study. Notably, the prevalence of the wild-type (A/A) genotype of the LTC4S -444 A/C polymorphism was significantly higher in the responders than in the non-responders (p=0.024 for allele prevalence, Table IV). Comparisons of the prevalence of genotype combinations revealed that the prevalence of the wild-type (G/G) genotype of the IL-13 R110Q polymorphism, compared with the G/A and A/A genotypes, was associated with favorable effects of treatment (p=0.043, Table V).

Discussion

Studies of genetic factors in patients with bronchial asthma have been conducted since the late 1980s (2-6). In the present study on the relationships between genetic polymorphisms and the efficacy of a Th2 cytokine inhibitor in children with bronchial asthma, we investigated the SNP profiles of the IL-12, IL-12Rβ1, IFN-γR1, IL-4Rα and IL-13 genes, which affect IgE production, and LTC4 synthase, which affects the production of proinflammatory mediators, in order to evaluate the

relationships between SNP profiles and response to treatment. In the present study, the LTC4S -444 A/C polymorphism was associated with favorable effects of suplatast tosilate. In 2000, Sampson *et al* reported that asthmatic patients with a variant allele in the LTC4S -444 A/C polymorphism exhibited increased improvement in respiratory function after treatment with leukotriene receptor antagonists (8). In the present study, treatment with the Th2 cytokine inhibitor, suplatast tosilate, (12,13) was more effective in patients with the wild-type -444 A/C polymorphism in the LTC4S gene than in patients with the variant LTC4S gene. It has also been reported that the expression of cysteinyl leukotriene 1 receptors is modified by IFN-γ and IL-13 (14). Evaluation of the relationship between genotype combination and treatment efficacy revealed a relationship between the presence of the wild-type IL-13 gene and efficacy of the Th2 cytokine inhibitor. In addition, the change in IFN-γ production between the baseline and treatment periods was significantly associated with efficacy of treatment. These findings suggest that the relationship between the LTC4S -444 A/C polymorphism and treatment efficacy may be explained by a change in the expression of cysteinyl leukotriene 1 receptors in responders. Further pharmacogenetic studies on gene polymorphism and expression should be performed in order to refine individually tailored treatment for asthmatic patients.

Acknowledgements

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