양극성 장애 환자에서 CTLA-4 유전자 다형성

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Polymorphism of CTLA-4 Gene in Patients with Bipolar Disorder

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ABSTRACT

bjective: Bipolar disorder is known to have strong genetic background and cellular immune activation. Based on the hypothesis that abnormalities of normal inhibitory control of T cell immunity can contribute to the pathophysiology of bipolar disorder, we investigated the relationship between the first exon at position +49(A/G) polymorphism of cytotoxic T lymphocyte antigen 4(CTLA4) gene and bipolar disorder.

Method: Among the Korean patients diagnosed as bipolar disorder according to DSM - , 90 patients without serious medical illness, neurologic illness, hormonal disorder, or concomitant mental illness were selected. The normal control group consisted of 149 age - and sex - matched subjects without current or past history of autoimmune diseases or mental disorder. DNA was extracted from whole blood and the exon 1 region of CTLA - 4 gene was amplified by polymerase chain reaction. Gene typing was performed using single strand conformation polymorphism.

Results: There were no significant differences in genotype frequencies of G/G, G/A, and A/A between the patients with bipolar disorder and the control group(48.9% vs 46.3%, 44.4% vs 39.6%, and 6.7% vs 14.1%, respectively). There were no significant differences in allelic frequencies of G and A between the patients with bipolar disorder and the control group(71.1% vs 66.1%, and 28.9% vs 33.9%, respectively).

Conclusion: This study did not show the association of exon 1 polymorphism of CTLA - 4 gene with bipolar disorder.

KEY WORDS : CTLA - 4 \cdot Polymorphism \cdot Bipolar disorder.



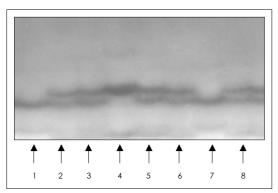
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가 가		alanine		
가	.5)	・ CTLA - 4 가		
가 ,		+49 CTLA - 4 exon 1		
・ , フト ⁷⁾⁸⁾	가 ⁶⁾	연구 대상 및 방법		
, (soluble interleukin - 2 receptor : sll 6R フト phytohemaggl		1. 연구 대상 가 DSM - ²⁴⁾		
9)10)		, , , 1 90 . 149		
Cytotoxic T lymphocyte antigen - T surface molecule) ,	· 4(CTLA - 4) (cell (antigen - prese-	가 ,		
nting cell) B7 . ¹¹⁾ CTLA - 4 B7 가 T		2. CTLA-4 유전자 형별분석		
(negative signal) ,	. ¹²⁾ T	1) DNA 추출 (Becton Dickin-		
CTLA - 4 T (negative (homeostasis) . 13)14), CTLA -	regulation) T 4가 T	son Vacutainer System, USA) 10ml , Proteinase K가 DNA		
, CTLA -	4 가 . CTLA - 4	2) Polymerase chain reaction(PCR) CTLA - 4 exon 1 + 49 A/G 2ul DNA 10X buffer (500mM KCI, 100mM Tris - HC1 pH 8.3, 15mM		
2q33 7,195 , 7h		MgCl ₂) 2ul, 20uM oligonucleotide primers(5 '-GCT CTA CTT CCT GAA GAC CT - 3 ', 5 '- AGT CTC ACT CAC CTT TGC AG - 3 ') 2ul, 2mM dNTP 2ul, Taq DNA polymerase(5U/ul : Boehringer Man-		
, exon 1 +49(A 가 lupus erythematosus) exon 1 +49 A/G	(systemic -22)23) threonine	nnheim, Germany) 0.2ul D/W 7\\ 20ul . Thermal cycler(GeneAmp PCR System 9600, Perkin Elmer, Foster City, USA) 94 4 , 94 45		
		(denaturation), 62 45		

(annealing), 72 45 (elongation) 35 3) Single strand conformation polymorphism (SSCP) PCR (single strand) DNA 5 95 stop buffer (95% formamide: 10mM NaOH: 0.25% bromophenol blue and 0.25% xylene cyanol) 14 MDETM gel(BioWhittaker Molecular Applications, Rockland, NY, USA) silver stain DNA 1). 3. 통계 분석 SAS/PC version 1) 6.0 2 0.05



2)

Fig. 1. The SSCP finding of the CTLA-4(A/G)dimorphism: Lanes 1 and 7: CTLA-4*G/G: Lanes 2, 3, 5, 6 and 8 : CTLA-4*G/A : Lane 4 : CTLA-4*A/A.

Table 1. Frequencies of the genotypes and alleles of the CTLA-4 gene

	Genotypes ^a			Allelesb	
Group	G/G	G/A	A/A	G	Α
	n(%)	n(%)	n(%)	n(%)	n(%)
Double on to					

Patients 44(48.9) 40(44.4) 6(6.7) 128(71.1) 52(28.9) (n=90)Controls 69 (46.3) 59 (39.6) 21 (14.1) 197 (66.1) 101 (33.9)

a: ²=3.14, d.f.=2, p=0.208 b: ²=1.29, d.f.=1, p=0.256

(power) 0.05 13% (w=0.13)26) (power=0.80)

걜 과

 2 =0.59, p= 0.74) 2 =2.03, p= 0.36) Hardy - Weinberg 가 CTLA-4 A/A 가 G/G, G/A (48.9% 46.3%, 44.4% 39.6%, 6.7% 14.1%)(1). CTLA-4 G 가 (, 66.1%, 28.9% 71.1% 33.9%)(1).

卫 찰

CTLA-4 exon 3 3 - untranslated re-.¹⁷⁾ exon gion 642 (AT)n 1 + 49(A/G)16) (promoter) - 318(C/T) 가 .²²⁾²³⁾ Grave 's CTLA-4 20)21) disease, 19)

CTLA-4 CTLA - 4 exon 1(+49)

가 CTLA-4 (association) 가 CTLA - 4 27)28) CTLA-4

27)

중심 단어: CTLA - 4 ·

가 가

,²⁹⁾ T

CTLA-4

가

가

가

interferon -

가

(mitogenic)

. 32) 기 (negative immune regulation)

CTLA - 4

가

13%

(w=0.13) (0.80) , (odds ratio) 1.83

(odds ratio)

,

, CTLA - 4 exon 1

, 가

가

CTLA - 4 exon 1

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