

Polymorphism screening of the neurotrophin receptor p75 gene and association analysis with schizophrenia

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Abstract

Based on both neurodevelopmental and neurodegenerative hypotheses in the etiology of schizophrenia, neurotrophic factors and their receptors may be involved in the pathogenesis of schizophrenia. In line with this, we previously reported a possible association between polymorphisms of the BDNF and NTF3 genes and the disorder. Here we searched for polymorphisms in the neurotrophin receptor p75 gene for genomic DNA from patients with schizophrenia. We detected one silent and one missense, nucleotide substitutions in the coding region. We performed an association analysis with respect to the missense polymorphism in a Japanese sample of 116 schizophrenics and 152 controls, matched for sex and age distributions. However, there was no significant difference in the genotype or allele frequency between the two groups, suggesting that the examined missense polymorphism is unlikely to play a major role in giving susceptibility to schizophrenia.

Introduction

Based on both neurodevelopmental and neurodegenerative hypotheses in the etiology of schizophrenia, neurotrophic factors and their receptors may be involved in the pathogenesis of schizophrenia. We previously reported a possible association between polymorphisms of the BDNF 1)

and NTF3 2,3) genes and the disorder. To further explore the possible role of the neurotrophins, we searched for polymorphisms in the gene encoding p75, a low affinity receptor for neurotrophin family. A detected missense polymorphism was examined for an association with schizizophrenia.

Methods

Polymorphism screening

Genomic DNA from 20 patients with schizophrenia was used. Six sets of primers were designed encompassing amino-acid coding region of the p75 gene. Polymorphisms were screened by direct sequencing (forward and reverse) of the PCR products.

Association analysis

Subjects for association analysis were 145 schizophrenics and 152 controls matched for age and sex distributions. Genotyping for a Ser/Leu missense polymorphism was done by PCR with primers 5'-gctaaaagggaggagtgggggaag-3' (p75E4F) and 5'-ttcaggtcaaggtcacagcaaagtct-3' (p75E4R) followed by digestion with a restriction enzyme BanII. To visualize polyacrylamide gel electrophoresis and etidium bromide staining was employed.

Results and Discussion

Two polymorphisms were found in the putative exon 4: one missense (Ser/Leu)

and the other silent, nucleotide substitutions. In an association analysis for the missense (Ser/Leu) polymorphism (Table), there was no significant difference in the genotype or allele distribution between the patients and controls.

We performed polymorphism screening for the aminoacid coding region of the p75 gene. However, we obtained no evidence for the involvement of the p75 gene in the pathogenesis of schizophrenia.

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390-2, 1994.

Table Genotype and allele frequencies for a missense polymorphism of the p75 gene among patients with schizophrenia and controls

	n	Genotype distribution			Allele frequency	
		S/S	S/L	L/L	S	L
Schizophr.	145	0.81	0.17	0.02	0.90	0.10
Controls	152	0.80	0.16	0.04	0.88	0.12
			p=0.64			p=0.56

Ⅲ. 研究成果の刊行に関する一覧表 (主要なものを選択)

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