

Brief Research Communication

Association of 5HT_{2A} Receptor Gene Polymorphism and Alcohol Abuse With Behavior Problems

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This study investigated the association between T/C polymorphism, at position 102, of the 5-hydroxytryptamine 2A receptor gene and alcoholism with and without behavior problems. Eighty-five subjects (45 men, 40 women) with alcohol abuse, 75 subjects (51 men, 24 women) with alcohol dependence, and 70 normal control subjects (21 men, 49 women) participated in the study. The results show that the frequency of the homozygous T102 genotype was significantly lower in the group of male alcohol abuse with behavior problems than in the female group ($\chi^2 = 4.072$, $df = 1$, $P < 0.05$) and the allele frequency of T102 was also lower in the male group than in the female group ($\chi^2 = 4.187$, $df = 1$, $P < 0.05$). Of the male alcohol abuse subjects, the group with behavior problems was found to have lower frequencies of the T102 allele than the group without behavior problems ($\chi^2 = 4.328$, $df = 1$, $P < 0.05$). In conclusion, this study demonstrates that alcoholism is heterogeneous and male alcohol abuse with behavioral problems was associated with T/C 102 polymorphism of the 5HT_{2A} receptor gene. *Am. J. Med. Genet. (Neuropsychiatr. Genet.)* 96:797–800, 2000.

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INTRODUCTION

Genetic factors are found to be important in the etiological complex of alcoholism [Cloninger et al., 1981; Sigvardsson et al., 1996]. Type 1 alcoholism, defined by Cloninger et al. [1981], was found to have both environmental and genetic risk factors, and Type 2 alcoholism, the severe form of alcoholism, was found to have a more emphasized genetic etiological factor [Sigvardsson et al., 1996]. Serotonin was found to play important roles in mediating drinking behavior through various mechanisms [Meltzer, 1990; Le Marquand et al., 1994; Virkkunen and Linnoila, 1997; Maurel et al., 1999]. Activation of various subtypes of serotonin receptors, such as 5HT_{2A}, 5-HT_{2C}, and 5-HT_{1B} reduce ethanol consumption [Maurel et al., 1999]. Clinical pharmacological agents increasing serotonergic activities were found to have a modest degree, but not consistent across studies, of action in reducing alcohol intake [Lejoyeux, 1996; Garbutt et al., 1999].

Low serotonergic function was responsible for poor impulse control, such as aggressiveness and self-injurious behaviors [Roy, 1990]. Alcohol-dependent individuals often show poor impulse control behavior [Linnoila et al., 1989]. There is a consistent finding that a proportion of alcoholism patients have association with behavior symptoms assigned as Type 2 alcoholism [Cloninger et al., 1981], Type B alcoholism [Barbor et al., 1992], and deviant-behavior type alcoholism [Hwu et al., 1992]. This behavior-disordered subtype of alcoholism is possibly based on the pathophysiological mechanism of reduced serotonergic functions.

Based on this background, a promising area of study is the search for genes related to serotonin transmission, responsible for vulnerability to alcoholism [Hill et al., 1999]. Serotonin transporter gene (HTT) polymorphism (HTTLPR) was not found to be associated with alcohol dependence [Edenberg et al., 1998]. The LL genotype of the HTT was found to be associated with a low level of alcohol response which was a vulnerability marker of alcoholism [Schuckit et al., 1999]. Nakamura et al. [1999] reported a positive association of alcohol dependence with a (A/G) polymorphism of the promoter region of 5HT_{2A} receptor gene located at chromosome 13q14-21 [Hsieh et al., 1990]. The present study was designed with the hypothesis that there is an association between alcoholism with behavior problems and

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receptor gene T/C polymorphism, at position 102, of the serotonin receptor subtype 2A (HTR2A) [Hsieh et al., 1990; Warren et al., 1993].

METHODS AND SAMPLES

The sample of this study consisted of 85 subjects (45 men, 40 women) with alcohol abuse (AA), 75 subjects (51 men, 24 women) with alcohol dependence (AD), and 70 normal control subjects (21 men, 49 women), after obtaining informed consent. The diagnoses of alcohol abuse and alcohol dependence by DSM-III criteria [APA, 1980], were assessed using the alcoholism section of the Chinese modified version of the Diagnostic Interview Schedule (DIS-CM) [Hwu et al., 1984, 1986; Lin et al., 1995] with satisfactory interrater reliability. Eleven items of childhood-adolescent behavioral symptoms included in this study were repeated truancy, repeated fighting with fists, ever fighting with arms, running away from home, lying, stealing, damaging other's property, being cruel to animals, attacking others without notice, setting fires, and robbery. In this study, those subjects who had three or more items of behavioral symptoms were coded as "with a behavior problem."

RESULTS

Data were obtained by using personal interviews performed by well-trained research interviewers. Peripheral blood samples were drawn and the DNA of white blood cells was extracted using the method of Sambrook et al. [1989] with modifications. Genotyping of the T/C polymorphism at position 102 of HTR2A was done by polymerase chain reaction (PCR) [Warren et al., 1993; Arranz et al., 1995] using the sense primer of 5'-TCT GCT ACA AGT TCT GGC TT-3', and the anti-sense primer of 5'-CTG CAG CTT TTT CTA GGG-3'. The 102T allele (T102) showed a PCR product fragment of 342 bp and the 102C allele (C102) showed two PCR product fragments of 126 bp and 216 bp after digestion with restriction enzyme of MspI.

The allele frequencies T102 and C102 in the AA, AD, and normal control groups were 72.9%, 69.3%, and 69.3%, and 27.1%, 30.7%, and 30.7%, respectively. The homozygous genotypes of T102/T102 and C102/C102 had frequencies of 50.6%, 44.0%, and 44.3%, and 4.7%, 5.3%, and 5.7% in the AA, AD, and normal control groups, respectively. No significant differences in genotype or allele distributions between study groups were found.

Table I shows the frequencies of genotypes and alleles in the three study groups with behavioral problems. The group of male alcohol abuse subjects with behavior problems was found to have lower frequencies of the genotype of homozygous T102 and to have higher frequencies of the genotype of homozygous C102 in contrast with the male groups of both alcohol dependence and normal control. The difference in frequencies of homozygous T102 genotype between the other two genotypes (T/C and C/C) between the male and female alcohol abuse subjects was statistically significant ($\chi^2 = 4.072$, $df = 1$, $P < 0.05$). The frequency of T102 was significantly lower in the male alcohol abuse subjects than in the female subjects with behavior problems ($\chi^2 = 4.187$, $df = 1$, $P < 0.05$).

Table II shows that the frequency of T102 in the male alcohol abuse group without behavioral problems was 79.5% and this was significantly higher ($\chi^2 = 4.328$, $df = 1$, $P < 0.05$) than that of the group with behavioral problems (59.1%). In the female alcohol abuse subjects, the frequencies of allele T102 in the groups with and without behavioral problems were similar (85.0% vs. 73.3%).

DISCUSSION AND CONCLUSION

This study used 11 behavior problem symptoms to evaluate the behavior problems in childhood and adolescence. The presence of a behavior problem was defined as having at least three symptom items. This is an arbitrary criterion for definition of having behavioral problems to indicate the tendency of having poor

TABLE I. Genotypes and Allele Frequencies of T/C 102 Polymorphism of the Receptor Gene of Serotonin Receptor 2A in Study Groups of Alcohol Abuse, Alcohol Dependence, and Normal Controls With Behavioral Problem

Genotypes	Alcoholism with behavioral problem								
	Alcohol abuse			Alcohol dependence			Normal controls		
	Male (%)	Female (%)	Total (%)	Male (%)	Female (%)	Total (%)	Male (%)	Female (%)	Total (%)
T102/T102	7 ¹ (31.9)	7 ¹ (70.0)	14 (53.8)	15 (50.0)	3 (33.3)	18 (46.2)	2 (50.0)	5 (45.5)	7 (46.7)
T102/C102	12 (54.5)	3 (30.0)	15 (46.9)	13 (43.3)	6 (66.7)	19 (48.7)	2 (50.0)	5 (45.5)	7 (46.7)
C102/C102	3 (13.6)	0 (0.0)	3 (9.4)	2 (6.7)	0 (0.0)	2 (5.1)	0 (0.0)	1 (9.0)	1 (6.6)
Total	22	10	32	30	9	39	4	11	15
Alleles									
T102	26 ² (59.1)	17 ² (85.0)	43 (67.2)	43 (71.7)	12 (66.7)	55 (70.5)	6 (75.0)	15 (68.2)	21 (70.0)
C102	18 (40.9)	3 (15.0)	21 (32.8)	17 (28.3)	6 (33.3)	23 (29.5)	2 (25.0)	7 (31.8)	9 (30.0)
Total	44	20	64	60	18	78	8	22	30

¹T102/T102 vs. (T102/C102 and C102/C102) by sex: $\chi^2 = 4.072$, $df = 1$, $P < 0.05$.

²(T102 vs. C102) by sex: $\chi^2 = 4.187$, $df = 1$, $P < 0.05$.

TABLE II. Comparison of Genotype and Allele Frequencies of T/C 102 Polymorphism of the Receptor Gene of Serotonin Receptor 2A Between Study Groups With Behavior Problem (GWB) and Without Behavioral Problems (WOB) in Male and Female Alcohol Abuse Cases

Genotypes	Alcohol abuse			
	Males		Females	
	GWB(%)	GOB(%)	GWB(%)	GOB(%)
T102/T102	7 (31.9)	13 (59.1)	7 (70.0)	15 (50.0)
T102/C102	12 (54.5)	9 (40.9)	3 (30.3)	14 (46.7)
C102/C102	3 (13.6)	0 (0.0)	0 (0.0)	1 (3.3)
Total	22	22	10	30
Alleles				
T102	26 (59.1)	35 (79.5)	17 (85.0)	44 (73.3)
C102	18 (40.9)*	9 (20.5)*	3 (15.0)	16 (26.7)
Total	44	44	20	60

* $\chi^2 = 4.328$, $df = 1$, $P < 0.05$.

impulse control which might be related to serotonin dysregulation [Roy et al., 1990]. This study shows that there is a significant difference in the allele frequencies of T102 and C102 between male alcohol abuse subjects with and without behavioral problems. The polymorphism of T/C 102 does not involve change in the amino acids of the protein. The best speculation for the significance of T/C polymorphism at the molecular biological level in male alcohol abuse may involve the affected translation through the secondary structure and the stability of the mRNA which, in turn, may decrease the functional activity of this receptor in male subjects and cause impulsive behaviors associated with alcohol abuse.

The study results emphasize that alcohol abuse defined by DSM-III is heterogeneous in nature and can be differentiated by gender and by the presence or absence of behavior problems. Moreover, the significance of behavior problems in alcohol abuse and alcohol dependence was also different. The polymorphism of MAOA gene [Sullivan et al., 1990] was found to be associated with Type 2 alcoholism, mainly of alcohol dependence cases, defined by Cloninger et al. [1981]. These data support the idea that the diagnostic categories of alcohol abuse and dependence are heterogeneous nosologically. The location of the HTR_{2A} is near the gene for esterase D [Hsieh et al., 1990], and it was found to be associated with alcoholism [Tanna et al., 1998]. However, this finding was not replicated in another study [Wesner et al., 1991]. This inconsistent finding might be due to different study samples. Moreover, the inconsistent findings of the therapeutic effect of serotonergic medications [Lejoyeux, 1996; Garbutt et al., 1999] in alcoholism could also be explained by the heterogeneity of alcoholism. Future studies of this kind may have to consider the presence of behavior problems and the different diagnostic categories of alcohol abuse and dependence.

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