No Association between the Response to Methylphenidate and DRD4 Gene Polymorphism in Korean Attention Deficit Hyperactivity Disorder: A Case Control Study

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Objective: Recently the relationship between alleles frequency distribution, drug response and the attention deficit hyperactivity disorder (ADHD), has been actively researched. We investigated the association between the genetic type, alleles and drug response for the dopamine receptor D4 (DRD4) gene in ADHD patients in Korea.

Methods: One hundred fourteen patients diagnosed with ADHD according to the the Diagnostic and Statistical Manual of Mental Disorders version IV (DSM-IV) diagnostic criteria were selected for the study. The clinical features of patients were confirmed by Korean version of Conners' parent rating scale, Attention deficit Diagnostic System, Korean version of Spielberger state—trait anxiety scale. Blood samples were taken from the 198 subjects, DNA was extracted from blood lymphocytes, PCR was performed for DRD4 Polymorphism. Alleles, genotype frequencies, the Clinical Global Impression (CGI) improvement score were compared using the chi-square test. Korean ADHD Rating Scale (K-ARS) and CGI severity scores were compared using the t-test. Results: In comparing the ADHD with 4/4 repeats group and without the ADHD with 4/4 repeats group, no significant difference

was seen between the DRD4 genetic type, alleles distribution, and CGI drug response.

Conclusion: As a result, it is viewed that there is no relationship between ADHD and DRD4, but final decision is indefinite. Follow up studies with larger patient or pure subgroups are expected.

KEY WORDS: Attention deficit hyperactivity disorder; DRD4 polymorphism; Case control studies; Drug response; Methy-lphenidate,

INTRODUCTION

Attention deficit hyperactivity disorder (ADHD), which is a neurodevelopmental disorder characterized by inattention, hyperactivity and impulsivity behavior, is a common disorder that has about 3-5% prevalence rate among prepubertal elementary school children (American Psychiatric Association, 1994). Recent family study shows that ADHD has an 80-90% hereditability, 11 accordingly molecular genetic studies are being actively carried out. Most of the ADHD studies are being done on the enzyme or receptor related genes in the dopaminergic neurotransmission pathways, because symptoms of ADHD show a 70% improvement upon treatment with the psy-

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chostimulants such as methylphenidate (MPH).²⁾ The dopamine receptor D4 (DRD4) gene has also been regarded as a leading candidate gene for ADHD. DRD4 maps to chromosome 11p15.5 and displays a high degree of variability at the 48-bp variable number tandem repeat (VNTR) on exon 3, which can be 2-11 times.³⁾

Association between the 7 repeat allele of a VNTR within the coding region of DRD4 and ADHD has been studied by several groups. LaHoste *et al.*⁴⁾ were the first report an increased frequency of DRD4 7 repeats (22% vs. 9%) in ADHD children. These findings were subsequently replicated in Coucasian children mainly. But there were few studies conducted in Asian ADHD children, no 7 repeats allele has been founded.^{5,6)} Leung *et al.*⁵⁾ reported firstly in eastern area a significantly increased prevalence of the 2-repeat allele at DRD4 in Chinese ADHD children, whereas none of them had the DRD4 7-repeat allele. Seeger *et al.*⁷⁾ reported that the presence of the 7-repeat allele of the DRD4 gene seems to be associated with a poor response to MPH treatment in ADHD. In addition,

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Hamarman et al. 8) reported that ADHD children possessing the DRD4 7-repeat allele require higher doses of MPH for symptom improvement and symptom normalization, and suggested the possibility that the 7-repeat allele of the DRD4 gene VNTR polymorphism correlates with treatment outcome. Also a few previous studies have reported that the most frequent allele and genotype of DRD4 in the Korean population were the 4-repeat allele and 4/4, respectively. 9,10) One preliminary report suggested that ADHD children having the 7-repeat allele but not having a 4-repeat allele of the DRD4 exon-III VNTR polymorphism showed a worse response to treatment with MPH than ADHD children having the 4-repeat allele, owing to the decreased function of the D4 receptor. 11) It was particularly suggested that the 4-repeat allele at DRD4 could be a predictor of the response to treatment. The aim of the present study was to further investigate the association between the DRD4 VNTR polymorphism and treatment response in groups of ADHD and healthy controls children drawn from the genetically homogenous Korean population.

METHODS

Subjects

The patient group individuals were recruited from the ADHD outpatient clinic at the Child and Adolescent Psychiatric Division of the Psychiatric Department of Dankook University Hospital, Cheonan, Korea. We diagnosed ADHD according to the Diagnostic and Statistical Manual of Mental Disorders version IV (DSM-IV)¹²⁾ and Schedule for Affective Disorders and Schizophrenia for School-Age Children-Present and Lifetime Version (K-SADS-PL). 13) All the subjects were Korean. Subjects were excluded from the study if there was any evidence of conduct disorder, mood disorder, anxiety disorder, Tourette's syndrome, pervasive development disorder, mental retardation (IQ \leq 70) and neurological disorders, including epilepsy. All the interviews were done by a well-trained child psychiatrist. The ADHD patients were administered the Attention-deficit Diagnosis System (ADS), the Korean version of the modified computerized test of variables of attention (TOVA) for obtaining the accurate diagnosis. 14) The Korean version of the Conner's rating scales were used in the present study to obtain parental reports of their children's behavior. 15) Subjects were included from our sample if they had a score over two standard deviations from the norm on the tests for ADS (T-score > 70). ADHD had a lot of cormorbid disorders,

as depressive disorder and anxiety disorder. So we excluded children with the high score of depressive symptoms and anxiety symptoms. Subjects with high anxiety scores (Spielberger trait/state anxiety scale score >47/49) on the Korean version of Spielberger trait-state anxiety scale for children were excluded, and subjects with high depression scores (Kovacs depression inventory score >29) on Kovacs depression inventory for children were also excluded. In this study we used as control group with no psychiatric disorder. These include 84 elementary students from Pyeongtaek in Korea, about 200,000 populations, screened to exclude depression, anxiety disorder, ADHD. Finally, 198 members that consisted of the affected children (114) and volunteers for normal control (84) were included in our study.

The efficacy measure was defined as a change in scores on the Korean ADHD Rating Scale (K-ARS)¹⁸⁾ and Clinical Global Impression (CGI) score¹⁹⁾ over time. The mean changes in the K-ARS, CGI severity score from baseline (week 0) to the end of treatment (week 8) and CGI improvement score (week 8) were also evaluated.

Procedures

The study protocol was approved by the institutional review board, and written informed consent was obtained. After structured interview, physical examination, and laboratory testing, eligible participants were assigned to drug. OROS MPH was dosed using a forced titration strategy starting at 18 mg/day and increased weekly in 9 mg/day increments, the maximum dose being 36 mg/day based on tolerability. No other psychotropic medications were permitted during the study. Efficacy and safety evaluations were also performed at baseline and 8 weeks.

DNA Extraction and Genotyping

Standard high salt extraction methods were used to isolate DNA from the whole blood lymphocytes. The primer sequence used to amplify the 48 base pair sequence of the VNTR polymorphic loci were the same as previously reported. Thirty-five cycles were conducted that consisted of denaturation at 94°C for 60 sec, annealing at 60°C for 60 sec and extension at 72°C for 60 sec. An initial denaturing step at 94°C for 10 min and a last extension step at 72°C for 10 min were also added. The reaction volumes were 50 ml, and they consisted of 0.1 μ g of genomic DNA, 10 pmol of each primer, 10 mM dNTPs, 0.5 Unit Taq polymerase, 10 Taq buffer and distilled water. 8 μ l of PCR products were run on 8% TAE (0.04 M Tris-acetate, 0.01 M EDTA) agarose gel. The amplified PCR products

were electrophoresised at 50 volt for 1 hour 20 min; they were stained with Ethidium bromide solution and visualized under UV light.

Statistical Analysis

The allele, genotype frequencies, and CGI improvement scores were compared using the chi-square test for the case control comparison analyses and CGI severity scores and K-ARS were compared using the t-test, using the SPSS program ver. 15.0 (SPSS Inc., Chicago, IL, USA). Statistical significance was set at p values less than 0.05.

RESULTS

Demographic Characteristics of the Subjects

Of the total 114 cases with ADHD group, 90 (78.9%) were boys and 24 (21.1%) girls. Their ages ranged from 6 years to 14 years with a mean age of 8.9±2.2 (standard deviation, SD) years. Abbreviated Conners' scale mean score was 18.9±3.3 (SD). Of the total 84 cases with control group, 51 (60.7%) were boys and 33 (39.3%) girls. Their ages ranged from 6 years to 14 years with a mean age of 10.2±1.3 (SD) years, abbreviated Connors' scale mean score was 5.9±4.5 (SD).

Comparison of the Frequency of the Genotypes and Alleles with Genetic Polymorphism between the Control Group and the ADHD Group

Genotype frequencies in the ADHD sample were 0.9% 2/2 repeats, 0.9% 2/3 repeats, 21.9% 2/4 repeats, 1.8% 2/6 repeats, 0.9% 3/4 repeats, 69.3% 4/4 repeats, 4.4% 4/5 repeats. Genotype frequencies of the control sample were 2.4% 2/2 repeats, 34.5% 2/4 repeats, 1.2% 2/6 repeats, 57.1% 4/4 repeats, 4.8% 4/5 repeats (χ^2 =6.24, degree of freedom [df]=6, p=0.40). In otherwise the 4/4 genotype frequencies in the ADHD sample were 69.3%, non 4/4 genotype frequencies were 30.7%. The 4/4 genotype frequencies in the control sample were 57.1%, non 4/4 genotype frequencies were 42.9%. The chi-square test result suggested that the 4/4 repeats genotype was borderline significantly higher trend in ADHD children than the non 4/4 repeats genotype (χ^2 =3.11, df=1, p=0.08) (Table 1). Genotype frequencies of DRD4 were consistent with those expected under Hardy-Weinberg equilibrium (HWE).

Allele frequencies in the ADHD sample were 13.2% 2 repeats, 0.9% 3 repeats, 82.9% 4 repeats, 2.2% 5 repeats, 0.9% 6 repeats. Allele frequencies of the control sample were 20.2% 2 repeats, 76.8% 4 repeats, 2.4% 5 repeats,

Table 1. Comparison between the 4/4 genotype at DRD4 genotype

Dating soals	Genotyp	oe, n (%)	2	p value*
Rating scale	Control	ADHD	χ	
The 4/4 genotype The non 4/4 genotype	48 (57.1) 36 (42.9)	79 (69.3) 35 (30.7)	3.11	0.08

 $[\]chi^2$ -test by Fisher's exact test, *If p<0.05 is a significant, and if p>0.05, p<0.1 is a borderline significant.

0.6% 6 repeats. The chi-square test result suggested that the 4 repeats allele was not significant in ADHD children than the non 4 repeats allele ($\chi^2=2.28$, df=5, p=0.13).

Comparison of the CGI Improvement with Genetic Polymorphism between the Control Group and the ADHD Group

According to the CGI improvement of children as assessed by their parents, we found that 74.7% of subjects with the 4/4 genotype at DRD4 showed a good response (above moderate improvement) to OROS MPH treatment, but 74.3% of subjects with the non 4/4 genotype at DRD4 showed a good response to MPH treatment ($\chi^2 = 1.47$, df=1, and p=0.68). The CGI severity score of the ADHD subjects with 4/4 genotype decreased from 5.1 ± 0.8 (base) to 3.4 ± 1.3 (8 weeks), the ADHD subjects with non 4/4genotype decreased from 5.0±0.9 (base) to 3.2±1.3 (8 weeks) respectively at baseline (t-test value=0.91, p=0.34) and after 8 weeks (t-test value=0.33, p=0.57) of MPH treatment (Table 2). Also the K-ARS overall score of the ADHD subjects with 4/4 genotype decreased from 30.1±7.3 (base) to 18.3±6.0 (8 weeks), the ADHD subjects with 4/4 genotype decreased from 29.5±8.6 (base) to 18.5±6.8 (8 weeks) respectively at baseline (t-test value=0.10, p=0.75) and after 8 weeks (t-test value=0.03, p=0.87) of MPH treatment (Table 2).

DISCUSSION

There have been few studies that defined the association of ADHD and the 4-repeat allele of DRD4. Barr et al.²¹⁾ conducted a transmission disequilibrium test (TDT) analysis to determine the association between ADHD and DRD4 and found significant evidence for the transmission of the 7-repeat allele and nontransmission of the 4-repeat allele. Leung et al. 5) have compared the transmitted/not transmitted groups of 32 Chinese ADHD children and their family by using the TDT; they detected a significant difference in the 4-repeat allele of DRD4. Kim et al. 6 reported a nonsignificant association between the DRD4 exon III

Table 2. Comparison between the 4/4 genotype at DRD4 and the response of the ADHD children to the drug treatment according to the CGI, K-ARS

Rating scale -	CGI improvement*		CGI severity		K-ARS total	
	Poor	Good	Base [†]	8 weeks [†]		8 weeks ¹
ADHD with 4/4 genotype	20 (25.4)	59 (74.7)	5.1±0.8	3.4±1.3	30.1±7.3	18.3±6.0
ADHD with non 4/4 genotype	9 (25.7)	26 (74.3)	5.0±0.9	3.2 ± 1.3	29.5±8.6	18.5±6.8

Values are presented as number (%) or mean±standard deviation. *The χ^2 -test (by Fisher's exact test) value=1.47; degree of freedom=1, p value=0.69; † Student's t-test value=0.91, p=0.34; † Student's t-test value=0.33, p=0.57; $^{\$}$ Student's t-test value=0.10, p=0.75; † Student's t-test value=0.03, p=0.87. p<0.05. DRD4, dopamine receptor D4; ADHD, attention deficit hyperactivity disorder; CGI, Clinical Global Impression; K-ARS, Korean ADHD Rating Scale.

48-bp VNTR polymorphism and 126 Korean ADHD children and their family by using the TDT. Also Cheon et al.²²⁾ detected a significant association between the DRD4 exon III 48-bp VNTR polymorphism and the response to MPH treatment in ADHD children. Those subjects homozygous for the 4-repeat allele at DRD4 showed a better response to MPH treatment than those who were not homozygous for this allele. In our study, the relation of the distribution of the DRD4, drug response and ADHD was examined, and an association was not detected. These findings are relatively not consistent with the preliminary results of Sallee et al. 11) study and Cheon et al. 22) study in Korea. Though there was a nonsignificant difference of DRD4 genotype frequencies between patient group and control group, there was not a difference in DRD4 allele frequencies. Such a result can suggest that ADHD is not associated with the polymorphism of the DRD4 gene in Koreans. However, since our result was obtained from a very small number, it is not reasonable to consider that this result represents all the ADHD children and their families in Korea. It has also been suggested that ADHD is a polygenic disorder and that genes involved in the metabolism of dopamine, norepinephrine, and other neurotransmitters may only contribute small phenotypic variance. Hence, it is difficult to make a conclusion. In addition, the result may have come about due to the heterogeneity of the various genes involved in ADHD. In other words, the point is that due to the expression of the various alleles of DRD4, by the multiple genetic interactions, a specific effect of a single gene may not be detected. Case control studies could also present population stratification. Otherwise family-based association methods could overcome this effect, particularly given the wide variation in the DRD4 allele among human populations.²³⁾ In the studies performed by Swanson et al., 23) it has been reported that the polymorphism of the DRD4 candidate gene and ADHD disorder were not associated, and as the major causality of such a result, there is a difference for the distribution of the alleles among geographical regions and races. Due to

these regional and racial differences, there could be wide differences for the frequency of alleles, and in our study, the frequency of the alleles in ADHD children was somewhat different from the frequency reported in other countries. The chi-square test result suggested that frequencies of the 4 repeats allele and CGI improvement scores were not significant in ADHD children than the non 4 repeats allele. However, larger sample size is necessary for controlling the confounding factors. Lastly, the relatively small sample size limits the generalization of our findings.

The limitations of our study are that first, the number of the study subjects was small, due to the power of our sample. The study was performed on 114 children who were diagnosed at outpatient clinics and the 84 normal control group. However, it was a small number, and so the range of statistical errors may be large. Second, in our study, the comparison with family group was not performed; thus, it could not be considered that the difference of the distribution from other races with checking the distribution of a normal group. Although our study population is composed of single ethnicity-Koreans, as was our previous genetic study in Korean children, our study's case control design may have a more diverse population. In our study, the comparison was performed with applying the case control method that do not considered the genetic influence within a family, and so the false negative results were increased.

Future gene association studies could be anticipated in the pure ADHD group with excluding comorbid disorder, in male children only, in the subtype of the hyperactive behavior group, in the group that responded well to MPH.²⁴ In the future, a study comparing the clinical differences in a larger number of ADHD patients by classifying them into small groups is required.

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