Association between pediatric autoimmune neuropsychiatric disorders associated with streptococcal infections disease and tumor necrosis factor-α gene −308 g/a, −850 c/t polymorphisms in 4-12-year-old children in Adana/Turkey

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OBJECTIVES: Pediatric Autoimmune Neuropsychiatric Disorders Associated with Streptococcal Infections (PANDAS) is a newly defined disease in neuropsychiatry and occurs with an autoimmune mechanism after Group A Beta Hemolytic Streptococcus (GABHS) infection. Tumor necrosis factor (TNF), encoded by TNF-α gene has an important role in the apoptotic mechanisms of autoimmune diseases. Recently, TNF-α polymorphisms and autoimmune/psychiatric disorders have been reported to be related. In this regard, we focused on to investigate a possible relation between the TNF-α gene promoter region−308 G/A and −850 C/T polymorphisms and PANDAS.

MATERIALS AND METHODS: In this study, ages of PANDAS patient and control groups were ranging from 4 years to 12-year-old. “Patient group” includes childhood onset PANDAS patients (n = 42) and “control group” includes healthy children (n = 58). Diagnoses have been carried out according to Diagnostic and Statistical Manual of Mental Disorder (DSM-IV) criteria with Affective Disorders and Schizophrenia-Present and Lifetime (KSAD-S-PL) and Children Yale-Brown Obsessive Compulsive Scale Moreover, PANDAS criteria established by the American National Psychiatry Institute have been employed for diagnoses. For identifying polymorphisms; Polymerase Chain Reaction, Restriction Fragment Length Polymorphism and Polyacrylamid Gel Electrophoresis were used.

RESULTS AND DISCUSSION: For −308 polymorphism, 37 of 42 PANDAS patients’ results and for −850 C/T polymorphism, 38 of 42 PANDAS patients’ results were obtained. According to our statistical analysis there is a positive relationship between PANDAS patients for −308 G/A polymorphism but not for −850 C/T polymorphism. There is no positive relationship between −308 G/A polymorphism and anti-strep-tolsyn O (ASO) titers and no relationship between −850 C/T polymorphism and ASO titers. We found, however, positive relationship between genders of patients (boys) and the disease. According to our results, we propose that the AA polymorphism of −308 G/A polymorphism can be used as a molecular indicator for PANDAS.

Key words: Autoimmune, pediatric autoimmune neuropsychiatric disorders associated with streptococcal, polymorphism, streptococcus, tumor necrosis factor-α gene

Introduction

Obsessive-compulsive disorder (OCD), tic disorders, a subgroup of childhood-onset OCDs called Pediatric Autoimmune Neuropsychiatric Disorders Associated with Streptococcal (PANDAS) are pre-pubertal onset autoimmune neuropsychiatric disorders associated with the basal ganglia abnormalities in which antibodies produced against Group A Beta Hemolytic Streptococcus (GABHS) infections cross react with neuron epitopes. Therefore, these disorders are referred
as post-streptococcal movement disorders.\cite{2-7} OCD and tic disorders may be triggered by stress, anxiety, and illnesses (for example; GABHS infections).\cite{8} GABHS infections, comorbidities, and tics are characteristic features in the etiology of OCD, PANDAS, and Sydenham’s chorea.\cite{9-10} Due to the damage of basal ganglia in a period of days to weeks following a GABHS infection, tics, and obsessive compulsive symptoms occur in patients.\cite{9} The association between Sydenham’s chorea and streptococcus was first stated by Taranta and Stollerman in 1956 however, association between OCD and Sydenham’s chorea was discovered by some workers.\cite{11,12} In the following decades, clinicians, and researchers continued to perform studies for neuropsychiatric disorders.\cite{12,13} Recent studies suggest that 50-80% of OCD cases have a childhood onset.\cite{9,14-21} Childhood onset OCD patients usually have a positive history of childhood tonsillitis, tic disorders and more antistreptolysin O (ASO) titer than adult onset OCD patients.\cite{22,23} In addition, obsessions and compulsions are the common symptoms for both of them.\cite{24} Prevalence of OCD among adolescents is 0.4%.\cite{6} Gender of patients can be effective or non-effective factors for PANDAS. It is proposed that mostly boys are affected compared to girls (3:2, boys:girls ratio) and age of onset for boys is generally earlier.\cite{14} Immune system of patient is affected as the disease progress. In the patient’s immune system, first B lymphocytes which are susceptible to M group of antigens neutralize streptococcus via macrophages. Following this activation some B lymphocytes pass over blood-brain barrier and associate with the cell membrane of neurons due to epitope similarity.\cite{18,25-28} Outcome of this association is the neuronal destruction of the patient’s apoptosis mechanisms. Therefore, the loss of neurons in the brain part which is supposed to coordinate the movement and active memory cannot function properly thereby neuropsychiatric signs like tics would be unavoidable.

Tumor necrosis factor (TNF-α) gene was selected because it is an important cytokine secreted by monocyte, phagocyte T-and B-lymphocytes, and natural killer cells and it is association with autoimmune-based neuropsychiatric disorders has been proposed previously. Moreover, primary incentive for us to research on this topic is that there has not been any research performed on TNF-α, which is a part of Major Histocompatibility (MHC) complex and autoimmune-based PANDAS disorder. Secondly, TNF-α-308 G/A and – 850 C/T polymorphisms have been shown to be related with autoimmune-based disorders which makes it plausible to think that these polymorphisms could also be related to PANDAS. Despite other researches on this issue, the relationship between neuropsychiatric disorders specifically PANDAS and GABHS infections are not completely understood yet. One reason for this uncertainty is the triggering of other existing factors.\cite{29}

**Materials and Methods**

In this study, 42 children with PANDAS from Cukurova University Medical Faculty Child and Adolescent Psychiatry Department and 58 healthy children from Cukurova University Medical Faculty Social Pediatrics Department were included. Ages of patient and control groups’ ages were ranging from 4-year to 12-year-old. Patients were diagnosed by Cukurova University Medical Faculty Child and Adolescent Psychiatry Department according to Diagnostic and Statistical Manual of Mental Disorder (DSM-IV) criteria with Kiddie Schedule for Affective Disorders and Shizophrenia-Present and Lifetime Version (KSAD-S-PL). Children’s Yale-Brown Obsessive Compulsive Scale was used to evaluate the severity of PANDAS symptoms. Demographic data and previous infection histories of patients and their families including tonsillitis infection history (more than 6 tonsillitis attacks/year), prophylactic antibiotic/penicillin use, tonsillectomy, ASO titers, symptom aggravation associated with infections were obtained. The Ethics Committee of the Cukurova University Medical Faculty approved this study. Informed consent obtained from the patient and control groups’ parents before collecting blood samples. To identify TNF-α gene polymorphisms; first we performed deoxyribonucleic acid (DNA) isolation from the blood; DNA of the patients was isolated by using the Miller’s salting out method.\cite{30} –308 G/A and – 850 C/T regions were amplified with Polymerase Chain Reaction. Polymorphisms were genotyped by using restriction enzymes with Restriction Fragment
Length Polymorphism (RFLP) and Polyacrylamid Gel Electrophoresis (PAGE) methods.[13] Cycle conditions of the reaction mixture were carried out in a thermal cycler using the thermal profile given below for identifying −308 G/A polymorphism; initial denaturation at 94°C for 1 min, denaturation at 94°C for 15 s, annealing at 62°C for 30 s, primer extension at 72°C for 15 s for 40 cycles and a final extension at 72°C for 1 h. For −308 G/A polymorphism, the 142 bp product was amplified with following primers; F-5’-GGGACACACACAG CATCAAGG–3’ and R–5’-AATAGGTTTTTGAAGGCCATG-3’. 126 bp digested product was referred as polymorphic AA genotype. The non-digested 142 bp product was referred as wild-type GG genotype. An allele generates a restriction site for Nco I restriction enzyme. This polymorphism includes only one base variation (G → A) distinguished by Nco I on 8% polyacrylamide gel. Thermal profile for identifying −850 C/T polymorphism; initial denaturation at 94°C for 3 min, denaturation at 94°C for 20 s, annealing at 68°C for 30 s, primer extension at 72°C for 1 min for 40 cycles and final extension at 72°C for 1 h. For −850 C/T polymorphism, the 131 bp product was amplified with following primers; F-5’AAGTCGAGTATGGGGACCCCCCTTA‑3’ and R-5’‑CCCCAGTGTGTGGCCATATCTTCTT‑3’. 106 bp digested product was referred as wild-type CC polymorphism. The non-digested 131 bp product was referred as polymorphic TT polymorphism. This polymorphism includes only one base variation (C → T) distinguished by Hind II enzyme on 8% polyacrylamide gel. Polymorphisms were genotyped using available restriction enzymes with RFLP and PAGE methods.[31‑33]

ASO is considered in Streptolysin O enzyme units (TODD). Values above 200 are accepted as significant and referred as a new indicator of infection.[34] Actually, normal value of ASO is 200 but in our study, cut-off value was taken as 400 because children in the patient group had already consulted to out-patient clinic during active infection period. Serum ASO titers of patients were determined using the Beckman Delta Nephelometor in Cukurova University Medical Faculty Balcali Hospital Central Laboratory Biochemistry Department. This laboratory is accredited by Joint Commission International.

Statistical analysis for comparing the prevalence of polymorphic genotypes between PANDAS patient and control groups were carried out using the Chi-square Test. In PANDAS patients, to evaluate the effect of ASO titer values on the polymorphic genotypes Kruskal-Wallis Test and for multiple comparisons Bonferroni adjusted Mann Whitney U Test was employed. Statistical analysis was performed using SPSS 15.0 software and the level of statistical significance was accepted as \( P < 0.05 \).

### Results

The current study was carried out to investigate the effect of polymorphic genotypes in the progress of PANDAS. For −308 G/A polymorphism, 37 of 42 PANDAS patients' results were obtained [Table 1]. In the patient group, for −308 G/A polymorphism, we detected 5 polymorphic GA (13.5%) and 32 polymorphic AA (86.5%) but no normal GG polymorphism. In the control group, however, five polymorphic GA (8.6%) and 53 normal (91.4%) GG genotypes were detected [Table 1].

For −850 C/T polymorphism, 38 of 42 PANDAS patients' results were obtained [Table 1]. In the PANDAS patients group, 16 normal CC polymorphism (42.1%), 19 polymorphic CT polymorphism (50%), and 3 TT (7.9%) genotypes were detected. In the control group, 19 polymorphic CT genotype (32.8%) and 7 polymorphic TT genotype (12.1%) were detected. However, the majority of genotypes consist of 32 normal CC genotypes (55.2%). Thirty two of the PANDAS patients (76%) and 30 of the control group (51.7%) were boys [Table 1].

According to our statistical results, there is a positive association between the PANDAS patients and − 308 G/A

| Polymorphisms | PANDAS (%) | Control (%) | \( P \) |
|---------------|------------|-------------|------|
| GA 308        |            |             | <0.001 |
| GG            | 0 (0)      | 53 (91.4)   |      |
| GA            | 5 (13.5)   | 5 (8.6)     |      |
| AA            | 32 (86.5)  | 0 (0)       |      |
| CT 850        |            |             | >0.001 |
| CC            | 16 (42.1)  | 32 (55.2)   |      |
| CT            | 19 (50)    | 19 (32.8)   |      |
| TT            | 3 (7.9)    | 7 (12.1)    |      |
| Gender (G/B)  | 10/32 (23.8)/(76.2) | 28/30 (48.3)/(51.7) | <0.001 |

PANDAS: Pediatric autoimmune neuropsychiatric disorders associated with streptococcal infections, GA: Guanine-Adenine, GG: Guanine - Guanine, AA: Adenine-Adenine, CC: Cytosine- Cytosine, CT: Cytosine- Thymine, TT: Thymine-Thymine, G/B: Girl/Boy
polymorphism (Chi-square value = 84.486, \( P < 0.001 \)) but no association for –850 C/T polymorphism was observed (Chi-square value = 2.892, \( P = 0.235 \)). In addition, we evaluated ASO titers of the patients to see if high levels of ASO titers affect the disease prognosis. We also compared the PANDAS patients’ ASO titers; however, we could not compare them with ASO titers of the control group since we had only ASO titer values of the patient group. There was no positive association between –308 G/A polymorphism-ASO titers (Mann Whitney = \( Z = −0.489, P = 0.651 \)) and –850 C/T polymorphism-ASO titers (Kruskal-Wallis Chi-square value = 0.496, \( P = 0.780 \)). We, however, found a positive relationship between disease and gender of the PANDAS patients (boys) (Chi-square value = 13.027, \( P = 0.001 \)). We know that boys are generally affected more than girls with PANDAS8 but we could not find a positive relationship between gender of the patients and ASO titers (Mann-Whitney = \( Z = −0.753, P = 0.457 \)). In addition to these factors, frequent infections are very strong determinants and typical features of PANDAS.

Discussion

In our country, GABHS infections occurring in childhood period are much more common than other countries.\(^{[34–36]}\) Some of the patients delay treatment because of not going to the doctor in time or don’t take enough care for their disease. Because of unconscious use of antibiotics, suppression of immune system, we took into consideration these conditions and decided to study. We know GABHS infections are associated with OCD/ PANDAS symptoms.\(^{[35–38]}\) We studied two polymorphisms of TNF-\( \alpha \) gene and ASO titers of patients. According to our statistical results, we have demonstrated a positive association between TNF-\( \alpha \) –308 G/A and the PANDAS susceptibility but there is no association for – 850 C/T polymorphism with the PANDAS. In different parts of the world, studies conducted for different OCD patients have variable results. Hounie’s \textit{et al.} results are in consistent with our results.\(^{[37]}\) According to their report, –308 G/A polymorphism is strongly associated with OCD.\(^{[20]}\) Rheumatic fever (RF) is a disease that occurs after GABHS infections. RF/rheumatic heart disease is an inflammatory disease that occurs after GABHS infections. It is believed to be caused by similar mechanism of PANDAS. Another study supporting our study is made by Rajendranath \textit{et al.} with RF patients. They found that – 308 G/A polymorphism is associated with RF.\(^{[35]}\) In contrast to, Zai’s \textit{et al.} who stated that TNF-\( \alpha \) gene – 308 G/A polymorphism is not associated with OCD in Canadian patients.\(^{[38]}\) Especially, we can see the remarkable effects of – 308 G/A polymorphism on phenotype-genotype relations because of the increased frequency of polymorphic genotypes in the PANDAS patients. If the PANDAS patients are not treated for longer periods, schizophrenic symptoms may be unavoidable because of the damage in particular parts of the brain. Boin \textit{et al.} found significant association between – 308 G/A polymorphism and schizophrenia.\(^{[39]}\) We didn’t find statistically significant difference among ASO titers, –308 G/A and – 850 C/T polymorphism. In other words, ASO is not a determinant for both polymorphisms. According to some investigators; there hasn’t been any relationship between ASO titers among children with OCD only, OCD + PANDAS, OCD + Chronic Tic Disorder (CTD) patients and the control group.\(^{[9–10]}\) In contrast, the results of another study revealed positive relationship between ASO titer and severity of tic disorder for 150 consecutive children presenting with tics.\(^{[23,40–42]}\) In another study of 105 patients diagnosed with CTD, OCD or attention-deficit hyperactivity disorder (ADHD), ASO titers were definitive. They found that ASO titers of ADHD patients were significantly increased.\(^{[24,25]}\) Furthermore, we found that boys are affected more compared to girls and gender of a patient was a significant factor for the disease. However, if we consider gender frequency for boys and girls who were not equal to each other, it seems like boys has more tendency for PANDAS genotypically. We think that it will be more appropriate to reach this conclusion when the number of boys and girls are equal to each other with broader sampling groups. As a secondary reason, for this difference could be the fact that the 6p21 chromosomal region where MHC locus resides along with TNF-\( \alpha \) is a fragile site which support the findings that this syndrome is autoimmune-based.\(^{[43,44]}\) Moreover, considering the fact that FRAFES-associated mental retardation protein (FMR 2) and FAM II A genes located on fragile X E site (\{(folic acid sensitive site of Xq28, ...
FRAXE) and FRAXF loci on X chromosome, respectively, are fragile sites associated with mental retardation, we could say that chromosomes other than X chromosome are also important on the higher tendency of boys to have PANDAS.[43,44] Through the new comprehension of the diseases, we can say epigenetic mechanism plays role for inactivation of some genes therefore, we have to look in a larger perspective for the disease. In many multifactorial and single gene neurodegenerative disorders, the symptoms are similar, but also these are the common characteristics of polygenic traits.[45] In addition, there are symptoms which differ from patient-to-patient. For that reason, simultaneous expression profiles of immune system genes should be investigated as well. The most compelling aspect of our research is the finding that TNF-α −308 promoter region AA polymorphism was detected in 86.5% of patients but not in controls which makes it a possible indicator to be evaluated in pre-disposition to PANDAS, however, due to the low sampling number our work should be supported with further research employing higher sampling numbers. In conclusion, we propose that TNF-α −308 AA polymorphism can be regarded as a definitive and molecular indicator of PANDAS which should be supported by further research on this issue.

Conclusion

We propose that TNF-α −308 AA polymorphism can be regarded as a definitive and molecular indicator of PANDAS, however, our findings should be supported by further research on this issue.

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