

An Associated Research for Genetic Polymorphism of 5-HTTLPR with Post-Traumatic Stress Disorder

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Abstract

The aim of this study was to investigate the influence of a polymorphism in the serotonin transporter gene (5-HTTLPR) in patients diagnosed with post-traumatic stress disorder (PTSD) in a Chinese sample of earthquake survivors. Polymerase chain reaction (PCR) amplification and amplified fragment length polymorphism (AFLP) were performed to type 5-HTTLPR promoter polymorphism in 57 PTSD patients and an equal number of healthy controls. The genotype and allele frequency distribution were analyzed and compared using various statistical methods. The frequency of LL, SL and SS genotypes in patients was found to be 5, 16 and 36 respectively, in comparison to 16, 22 and 19 in healthy controls. Fewer patients tended to be L genotype (22.8%) than controls (47.4%), but the number of patients with the S genotype was higher (77.2%) compared to controls (52.6%). The results show a statistically significant difference in genotype and allele frequency distribution between patients and controls. This research suggests that PTSD symptoms are significantly associated with 5-HTTLPR genetic polymorphism. These results add to the important research of genetics of psychiatric disorders, particularly in a Chinese context that has not been previously studied.

Keywords

Posttraumatic Stress Disorder, Gene Polymorphism, 5-HTTLPR, Genetics

1. Introduction

In recent years, the incidence of Post-Traumatic Stress Disorder (PTSD) has significantly increased due to the frequency of occurrence of natural and man-made

disasters worldwide [1]. According to the diagnostic manual of the American Psychiatric Association, PTSD is a psychiatric disorder that can occur in people who have experienced or witnessed a traumatic event such as a natural disaster, a serious accident, a terrorist act, war/combat, rape, or other violent personal assault [2]. The 4 key symptoms of PTSD include: intrusion symptoms associated with the traumatic event(s); persistent avoidance of stimuli associated with the traumatic event(s) [3]; negative alterations in cognitions and mood associated with the traumatic event(s); and, marked alterations in arousal and reactivity associated with the traumatic event(s) [4]. Because of the higher incidence of PTSD after encountering a traumatic event [5] and the accompanying possibilities of serious detriments to the individual, family and society, this condition has become a popular topic for psychiatric and clinical psychological research in recent years [6]. With the increase of natural disasters occurring among the world and in China, (including earthquakes in Sichuan, Qinghai, Gansu and Yunnan), the incidence of psychological sequelae and more serious condition of PTSD has been increasing in recent years [7]. Therefore, exploring the mechanism of PTSD is important and necessary, to identify effective methods of prevention and treatment [8].

Most biological studies of PTSD have focused on changes in the gray matter nuclei and brain cortex, although recent research interest is being focused on genetics and immunology [9]. Of particular interest is the polymorphism in the serotonin transporter gene (5-HTTLPR) that has been studied in relation to both PTSD and depression [10], as the serotonin transporter (5-HTT) is responsible for terminating serotonin action in the synapsis. Interestingly, the human genetic polymorphism 5-HTTLPR results in a combination of short (S) and long (L) forms. The S genotype restricts the transcriptional activity producing low functional expression of the 5-HTT and thus decreasing serotonin reuptake, while the L genotype does not results in this changes [11]. While current evidence does not fully support a direct effect of the polymorphism 5-HTTLPR on PTSD [12], the present study further aims at exploring the correlation of genotype and allele frequencies of 5-HTTLPR with PTSD in a Chinese population of earthquakes survivors. The study further explores a scientific basis for the clinical diagnosis of PTSD patients based on genetic research of the condition.

2. Material and Methods

2.1. Subjects

All cases in the study group were patients diagnosed with PTSD in the Hainan Province of China during the period from December 2013 to August 2014. This province was selected because the individuals living there had experienced a tragic earthquake several years before. The inclusion criteria were that patients were: 1) aged 35 to 55 years old; 2) right-handed; 3) met the diagnostic criteria for PTSD according to the Diagnostic and Statistical Manual of Mental Disorders, 5th Edition (DSM-5). Patients were excluded if they had a dual diagnosis of

mental retardation, personality disorders, substance dependence or alcohol dependence. Voluntary and informed consent to participate in the research was obtained. In terms of their demographics, the study group consisted of 28 males and 29 females, with a mean age of 46.26 (± 6.40) years, and mean years of schooling of 10.59 (± 3.20) years. The control group consisted of subjects who did not meet any DSM-5 criteria for psychiatric disorder that were recruited. Specifically, the control group consisted of 30 males and 27 females, with a mean age of 42.92 (± 5.98) years, and mean years of schooling of 10.96 (± 3.08) years.

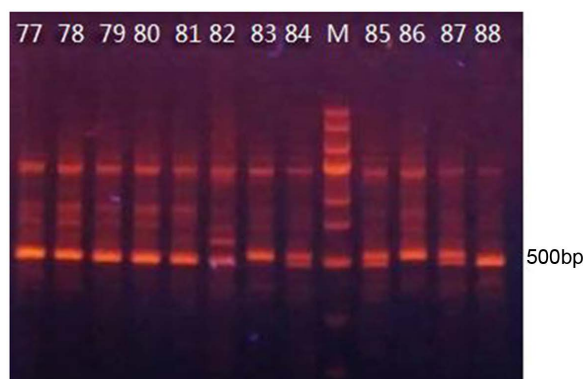
2.2. Methods

Conventional chlorine phenol extraction of genomic human DNA samples was used blood. The PCR primers used for genotyping were: forward 5'-GGCGTTGCCGCTCTGAATGC-3', reverse 5'-GAGGGACTGAGCTGGACAACCAC-3'. PCR was performed using the *Ex Taq* DNA Polymerase reaction (Takara Clontech, BaoBiotechnology) and the PCR product was run in a 6% non-denaturing polyacrylamide gel vertically. Ethidium bromide (EB) detects electrophoresis images of each sample and determines the genotype by using UV gel analysis after staining (**Figure 1(a)**, **Figure 1(b)**).

5-HTTLPR gene polymorphism detection report (**Table 1**, **Table 2**, **Table 3**)



(a)



(b)

Figure 1. Electrophoresis result map.

Table 1. PCR reaction system.

2 mM dNTP	1.5 ul
10 × KOD buffer	1.5 ul
MgSO4	0.6 ul
DMSO	1.5 ul
Primer 5-HTT-F: (10 uM)	1 ul
Primer 5-HTT-R: (10 uM)	1 ul
template	1 ul
Blend Taq-Plus	0.5 ul
ddH2O	6.4 ul
Total	15 ul

Table 2. PCR amplification conditions.

94°C 2 min	} 35 cycle
94°C 20 sec	
60°C 30 sec	
72°C 40 sec	
72°C 10 min	
4°C save	

Table 3. Statistical results.

(a)		
Result type	Sample number	Number of samples (a)
SS type (484/484 bp)	61. 65. 70. 71. 84. 85. 87. 93. 94. 79. 80. 81. 83. 86. 92. 90. 99. 95. 94. 101. 100. 102. 103. 104. 105. 107. 108. 109. 110. 112. 113. 114. 115. 117. 118. 116.	36
LL type (528/528 bp)	73. 76. 77. 78. 91.	5
SL type (484/528 bp)	62. 63. 64. 66. 67. 68. 69. 72. 74. 75. 82. 88. 89. 96. 97. 98.	16
(b)		
Result type	Sample number	Number of samples (a)
SS type (484/484 bp)	1. 2. 3. 4. 5. 6. 7. 8. 9. 10. 11. 12. 13. 14. 15. 16. 17. 18.	19
LL type (528/528 bp)	20. 21. 23. 22. 24. 25. 26. 27. 28. 29. 30. 31. 32. 33. 34. 35.	16
SL type (484/528 bp)	36. 37. 38. 39. 40. 41. 42. 43. 44. 45. 46. 47. 48. 49. 50. 51. 52. 53. 54. 55. 56. 57.	22

First: reagents and instruments

- 1) Reagents: EasyPure[®] Genomic DNA Kit; Blend Taq-Plus BTQ-201
- 2) Instrument: Hema 9600 TE Thermo cycler

Second: the experimental process:

- 1) Blood sample genomic DNA extraction
 - a) Prepare a 56°C water bath in advance

- i) Place 200 μ l of blood sample in a sterile 1.5 ml centrifuge tube.
 - ii) Add 100 μ l of LB2 and 20 μ l of Proteinase K.
 - iii) Incubate at 55°C until complete lysis (3 hours, 2 - 3 times per hour). Add 20 μ l of RNase A to the sample and incubate for 2 minutes at room temperature to remove RNA.
 - iv) Centrifuge at 12,000 rpm for 5 minutes and transfer the supernatant to a sterile centrifuge tube.
 - b) Add 500 μ l of BB2, vortex for 5 seconds, and incubate for 10 minutes at room temperature.
 - c) Add all the solution to the spin column, centrifuge at 12,000 rpm for 30 seconds, and discard the effluent.
 - d) Add 500 μ l of solution CB2, centrifuge at 12,000 rpm for 30 seconds, and discard the effluent.
 - e) Repeat step 4 once.
 - f) Add 500 μ l of solution WB2 (check for the addition of absolute ethanol before use), centrifuge at 12,000 rpm for 30 seconds, and discard the effluent.
 - g) Repeat step 6 once.
 - h) Centrifuge at 12,000 rpm for 2 minutes to completely remove residual WB2.
 - i) Place the spin column in a clean centrifuge tube, add 50 - 200 μ l preheated EB (65°C) in the center of the column, or deionized water (pH > 7.0). Allow to stand at room temperature for 1 minute, centrifuge at 12,000 rpm. In minutes, elute the DNA.
 - j) For more DNA, perform a second elution, add 50 - 200 μ l pre-heated EB (65°C) in the center of the column, or deionized water (pH > 7.0) for 1 minute at room temperature, 12,000 rpm Centrifuge for 1 minute and elute the DNA. The eluted DNA was stored at -20°C.
- 2) PCR and agarose gel electrophoresis identification
 - a) Using the extracted DNA as a template
 - b) Primer: 5-HTT-F: 5'-GGCGTTGCCGCTCTGAATGC-3',
5-HTT-R: 5'-GAGGGACTGAGCTGGACAAAC-3'.
- Three: Experimental result.

2.3. Statistical Analysis

The data comparing the genotype and allele frequencies of two groups was analyzed using the SPSS13.0 statistical software package using the following statistical methods: Hardy-Weinberg (HW) goodness of fit test, frequency analysis, analysis of variance, *t* test, and χ^2 test. $p < 0.05$ was used as the threshold for significance.

3. Results

3.1. The Comparison between the Study Group and the Control Group

The source of the research object is mainly in Hainan Province, a large number

of immigrants from earthquake areas (Sichuan, Qinghai, Gansu and Yunnan) (the resettlement personnel around the Yangtze River Three Gorges Dam in China) settled in Hainan Province, passing our hospital. Patients in the psychological clinic were selected for PTSD; the source of the control group was mainly from immigrants from the earthquake-stricken areas to those who settled in Hainan Province. All participants in the study were willing and informed consent. In terms of demographics, there were no statistical differences in age, gender, and education between the study group and the control group.

As shown in **Table 4**, the demographic characteristics of gender, age and level of education of the study group (with PTSD) and the control group revealed no statistically significant differences ($p > 0.05$).

3.2. Hardy-Weinberg Equilibrium Law Genetic Testing

According to the Hardy-Weinberg equilibrium law, the distribution of the 5-HTTLPR gene is consistent with this genetic equilibrium law with a group representation. **Table 5** shows the frequency of LL, SL and SS genotypes in the study group and control subjects. Specifically, patients were found to be 5, 16 and 36 respectively, in comparison to 16, 22 and 19 in healthy controls. The study and control group test results showed that the mean difference was not significant ($p > 0.05$).

3.3. Comparison of Genotypes and Allele between the Study Group and the Control Group

Table 6 shows the comparison of frequencies of genotypes and allele between the study group and the control group. The table shows that the 5-HTTLPR gene in the PTSD study group had frequencies of 5, 16, 36 respectively in the LL, SL and SS types. In comparison, the control group revealed frequencies of 16, 22, 19 respectively in the LL, SL and SS types. This resulted in a mean difference of 11.96, which it is statistically significant ($p = 0.003$). **Table 6** also shows the allele frequency of the two groups, and reveals that for the PTSD group, the L gene appeared in 22.8% of cases and the S gene appeared in 77.2% of cases. In contrast, for the control group, the L gene appeared in 47.4% of cases, and the S gene appeared in 52.6% of cases. The table shows that a comparison of these frequencies for the study and control group on the allele frequency revealed a mean difference of 15.10 ($p < 0.01$). Thus, both groups showed statistically significant differences between the genotypes and allele frequencies (see **Table 6**).

Table 4. Comparison of demographic characteristics of the study group and the control group.

Groups	n		Age (years)	Years of education (years)
	Male	Female		
The Study Group (PTSD)	28	29	46.26 ± 6.40	10.59 ± 3.20
The control group	30	27	42.92 ± 5.98	10.96 ± 3.08
<i>P</i>	$\chi^2 > 0.05$	$\chi^2 > 0.05$	$t > 0.05$	$t > 0.05$

Table 5. The Hardy-Weinberg equilibrium law genetic testing of 5-HTTLPR gene distribution.

Groups	Genotypes			χ^2	<i>P</i>
	LL	SL	SS		
The Study Group (PTSD)	5	16	36	2.34	>0.05
The control group	16	22	19	2.91	>0.05

Table 6. The comparison of Genotypes and allele between study group and control group comparison [Frequency (%)].

Groups	n	Genotype frequencies (%)			Allele frequency (%)	
		LL	SL	SS	L	S
The Study Group (PTSD)	57	5 (8.8)	16 (28.1)	36 (63.1)	26 (22.8)	88 (77.2)
The control group	57	16 (28.1)	22 (38.6)	19 (33.3)	54 (47.4)	60 (52.6)
			χ^2			
				11.96		15.10
			<i>P</i>			
				0.003		0.000

4. Discussion and Conclusion

The present study demonstrates that PTSD symptoms after exposure to earthquakes in a Chinese sample are associated with the 5-HTTLPR genetic polymorphism. Interestingly, among the study group with PTSD the S genotype was significantly higher when compared to controls, indicating an association between the S genotype and PTSD in this Chinese sample. Notably, our results are consistent with the hypothesis of a gene-by-environment ($G \times E$) interaction that suggests variations in the 5-HTT gene influence psycho-pathological reactions to stressful experiences [13], such as earthquakes.

PTSD is an important post-disaster mental and behavioral disorder with high incidence and prevalence. In addition this disorder has a potentially long duration and poor efficacy characteristics that seriously affect clinical treatment [14]. The increase in natural disasters, war, terrorism, and other precipitating factors of PTSD in recent years has resulted in more research related to the disorder. For example, the occurrence of the Wenchuan earthquake in China's Sichuan Province on May 12, 2008, measuring 8.0 on the Richter scale, created well-deserved attention and needed research about PTSD [15]. The attention has been intensified over the last years due to the increased number of natural disasters occurring in provinces of China.

Currently, despite the considerable amount of research on PTSD and its pathogenesis, the contribution of distinct genetic factors remains unclear. Stress diathesis theories predict that an individual's sensitivity to stressful events depends on their genetic background [16], and such predictions are now increasingly supported by experimental evidence. Much of this research has been focused in the analysis of polymorphisms in single genes or individual sites. One polymorphism that receives increased attention over the years is the serotonin

transporter gene polymorphism (5-HTTLPR). Heils *et al.* first cloned the human 5-HTT coding gene, which is encoded by a single gene SLC6A4 and located on chromosome 17q12 region [17]. Although the 5-HT system has been shown to modulate mood, irritability, and sleep disorders [18], the exact mechanism by which 5-HT modulates PTSD symptoms is still unclear.

The 5-HTTLPR polymorphism occurs in the promoter region of the 5-HTT gene that results in two variations in humans, a short (S) and a long (L) allele. Specifically, the S allele results in a lower expression of the 5-HTT compared with the L allele [19]. In other words, *in vitro* experiments have shown that the L fragment allele has a higher transcriptional activity than the S allele. In addition, cultured brain cells with the LL genotype result in higher 5-HTT mRNA and protein expression than the SS genotype. Thus, changes in 5-HTT protein expression may impair the 5-HT system function and affect individual's emotion [20]. In particular, variations in the functional human polymorphism 5-HTTLPR have been hypothesized to affect risk for PTSD [21]. Interestingly, the 5-HTTLPR human polymorphism has been also associated with neuroticism [22] and increased vulnerability to affective disorders [23], in many but not all studies [24]. For example, the S allele carriers, that have lower expression of 5-HTT, are more vulnerable to depression and suicidality in response to stressful events during adulthood than are individuals carrying the L allele, that have higher expression of 5-HTT [25].

With the aim of investigating the association between PTSD and the 5-HTTLPR polymorphism, we conducted an analysis to shed light on this interaction in a Chinese sample that had suffered from natural disasters in a context that has not been previously studied. By determining the genotype and allele frequency distribution among PTSD patients and healthy control subjects we demonstrate that a higher number of PTSD patients have the S variant when compared to controls. This result suggests that the S allele of 5-HTTLPR is significantly associated with PTSD symptoms and is in line with the mechanistic model of the 5-HTTLPR polymorphism. This model of the human 5-HTTLPR polymorphism suggests that different transcription of the 5-HTT gene creates populations of individuals with lower expression of 5-HTT (S carriers) that exhibited more PTSD symptoms in relation to stressful life events than individuals with higher expression of 5-HTT (L carriers).

Kate *et al.* Prior studies have found that the 5-HTTLPR polymorphism in the promoter region of the serotonin transporter gene (SLC6A4) interacts with stressful life events to increase general risk for PTSD, 5-HTTLPR genotype modified the effect of emotional abuse on PTSD symptom severity. Participants with the low-expression SS genotype who were exposed to emotional abuse had significantly lower reexperiencing and arousal symptom severity scores [26]. Consistent with our results, when examined in a relatively homogenous sample with shared trauma and known prior levels of child and adult trauma, the 5-HTTLPR multimarker genotype may serve as a useful predictor of risk for PTSD-related symptoms in the weeks and months following the trauma [27]. Notably, previous

studies have also shown an association of the 5-HTTLPR with different mental conditions; for example, the SS genotype often appears in patients with severe depression and anxiety disorders [28]. Furthermore, in a study of a large sample of twins it has been reported that individuals containing 5-HTTLPR SS or SL are more prone to depressive reactions of PTSD after stressful events [29]. Other study showed that the SS or SL allele carriers have a higher amygdala activity than the LL allele carriers when the individual encountered fear-related stimuli, and that the anxiety levels were lower in SL carriers when compared to LL carriers. Thus, these results suggest that the S genotype of the 5-HTTLPR may prompt fear and anxiety related symptoms [30]. Furthermore, it has been also shown that individuals with two short alleles (SS or SL) of 5-HTT polymorphisms are more vulnerable to stress than those who have only one or two long alleles (LL) [31], indicating a higher susceptibility risk. Further, mutations in the 5-HTT gene can affect individuals' sensitivity to stress and may play a role in the extent or intensity of the stress reaction. Although few studies has investigated the association between the 5-HTTLPR and susceptibility to PTSD, it has been suggested that the observed amygdala activity improvement after fear exposure therapy is more likely to occur in SS and SL carriers when compared to LL carriers.

From our knowledge, there are not studies that had investigated the relationship between PTSD and 5-HTTLPR in a Chinese population but several lines of evidence suggested that this interaction exist in non-Chinese populations, where it has been demonstrated a correlation between the S allele carriers and PTSD symptoms [31]. The present study addresses this issue and, our results demonstrate that the association between PTSD and 5-HTTLPR also exists in the Chinese population. However, the small sample size of this study presents limitations in the study design and therefore we should be cautious with our own conclusions. For example, it is difficult to rule out the confounding effects of PTSD. Therefore, while the present study contributes importantly to the genetics and psychiatric disorders research in general and specifically in a Chinese context, these results could valuably be replicated, to establish further reliability and validity.

In sum, this epidemiological study provides evidence of a gene-by-environment interaction in a Chinese population, in which their genetic background moderates an individual's response to environmental insults. However, validation and replication of these results will allow us to begin to understand how certain vulnerability factors affects emotional functions to increase the risk for PTSD symptoms later in life. These results may help guide the choice of outcome measures in clinical studies and aid in the identification of molecular mechanisms that may confer vulnerability to PTSD disorders to help us to better categorize and manage this complex disease.

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Conflicts of Interest

The authors declare no conflicts of interest regarding the publication of this paper.

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Highlights

- Our results are consistent with the hypothesis of a gene-by-environment interaction.
- The S genotype was higher in the PTSD group when compared to controls.
- PTSD due to earthquakes in a Chinese sample is associated with the 5-HTTLPR.