



Associations Between *PPP1R1B* Gene Polymorphisms and Anxiety Levels in the Chinese Population

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Dear Editor:

Early in 1983, DARPP-32, a phosphoprotein regulated by dopamine (DA) and cAMP, encoded by the *PPP1R1B* gene, was identified as a mediator of postsynaptic DA receptor signaling in striatal neurons [1]. Later, studies on postmortem brains using the quantitative polymerase chain reaction, immunohistochemical staining, and immunoblotting demonstrated significantly changed expression of DARPP-32 in the dorsolateral prefrontal cortex (DLPFC) of patients with mental disorders such as schizophrenia and bipolar disorder [2, 3]. At the same time, several gene association studies demonstrated that *PPP1R1B* polymorphisms or their haplotypes are associated with schizophrenia and substance dependence [4, 5]. However, no studies have tested for an association between *PPP1R1B* polymorphisms and anxiety levels or anxiety disorders. It has been demonstrated that the *PPP1R1B* gene is expressed in brain regions closely associated with anxiety, such as the amygdala and hippocampus [6, 7]. Moreover, DARPP-32 is closely associated with the DA and glutamine neurotransmitter systems that mediate anxious emotion and behavior. The *PPP1R1B* gene and DARPP-32 may thus play a role in anxiety through these neural pathways.

Here, we investigated in a healthy Han Chinese population whether genetic variants in the *PPP1R1B* gene are associated with state anxiety (a current emotional state) and

trait anxiety (a consistent emotional state) using the 40-item State-Trait Anxiety Inventory (STAI). A previous study by our team has demonstrated significant associations between the *PPP1R1B* intron single nucleotide polymorphisms (SNPs) rs12601930C/T, rs879606A/G, and rs3764352A/G, and personality traits measured by the Tridimensional Personality Questionnaire in healthy Han Chinese [8]. Because anxiety, especially trait anxiety, is considered to be an important and enduring dimension of personality, we examined its association with the above three SNPs based on their previously-established association with personality traits. A total of 571 unrelated healthy Han Chinese volunteers (272 males, 299 females) were recruited. The age range was 19–58 years (30.20 ± 7.82 years). All protocols were approved by the Ethics Committee of China Medical University. Written consent was given by all participants and venous blood samples were taken. The polymorphism genotyping experiments were performed according to the methods described in our previous studies [4, 8, 9].

The genotype counts and frequencies for the rs12601930C/T, rs879606A/G, and rs3764352A/G polymorphisms are shown in Table 1. The genotype distributions did not deviate from Hardy-Weinberg equilibrium. We did not find any statistical difference in the state-anxiety dimension for the rs12601930C/T, rs879606A/G, and rs3764352A/G genotypes, but statistical differences in the trait-anxiety dimension for the rs879606A/G ($F = 5.207$, $P = 0.006$) and rs3764352A/G ($F = 3.960$, $P = 0.020$) genotypes (Table 1). We also compared the anxiety scores among the three genotypes of each SNP between genders (Table 1). In the female participants, there were statistical associations of the rs879606A/G ($F = 4.977$, $P = 0.007$) and rs3764352A/G ($F = 3.404$, $P = 0.035$) genotypes with trait-anxiety levels, while no such associations were

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Table 1 Effects of *PPPIR1B* gene polymorphisms on STAI scores.

SNP	Genotype	Total <i>n</i> (%)	State anxiety	Trait anxiety	Male <i>n</i> (%)	State anxiety Male	Trait anxiety Male	Female <i>n</i> (%)	State anxiety Female	Trait anxiety Female
rs12601930	C/C	287 (50.26%)	37.51 ± 7.96	38.53 ± 6.97	117 (43.01%)	37.59 ± 7.12	39.15 ± 6.59	170 (56.86%)	37.45 ± 8.50	38.10 ± 7.20
	C/T	234 (40.98%)	37.93 ± 7.69	38.86 ± 6.43	127 (46.69%)	37.24 ± 7.82	38.43 ± 6.36	107 (35.79%)	38.75 ± 7.49	39.37 ± 6.51
	T/T	50 (8.76%)	37.22 ± 9.36	39.54 ± 7.69	28 (10.29%)	36.71 ± 8.31	39.82 ± 6.48	22 (7.36%)	37.86 ± 10.72	39.18 ± 9.16
	<i>F</i>		0.259	0.515		0.171	0.715		0.793	1.120
rs879606	<i>P</i> value		0.772	0.598		0.843	0.490		0.454	0.328
	A/A	130 (22.77%)	37.97 ± 8.48	38.85 ± 7.07	74 (27.21%)	38.46 ± 8.60	39.23 ± 7.32	56 (18.73%)	37.32 ± 8.37	38.34 ± 6.74
	A/G	291 (50.96%)	38.12 ± 7.84	39.47 ± 6.84	140 (51.47%)	37.27 ± 7.14	39.11 ± 6.16	151 (50.50%)	38.90 ± 8.39	39.81 ± 7.41
	G/G	150 (26.27%)	36.49 ± 7.70	37.28 ± 6.34	58 (21.32%)	36.05 ± 7.01	37.90 ± 6.01	92 (30.77%)	36.76 ± 8.13	36.89 ± 6.55
rs3764352	<i>F</i>		2.210	5.207		1.669	0.865		2.094	4.977
	<i>P</i> value		0.111	0.006		0.190	0.422		0.125	0.007
	A/A	138 (24.17%)	36.58 ± 7.86	37.36 ± 6.37	47 (17.28%)	36.26 ± 7.09	37.89 ± 6.07	91 (30.43%)	36.75 ± 8.26	37.09 ± 6.53
	A/G	286 (50.09%)	37.78 ± 7.49	39.08 ± 6.34	143 (52.57%)	37.10 ± 7.02	39.08 ± 6.10	143 (47.83%)	38.45 ± 7.89	39.07 ± 6.60
rs3764352	G/G	147 (25.74%)	38.43 ± 8.89	39.43 ± 7.89	82 (30.15%)	38.35 ± 8.62	39.10 ± 7.28	65 (21.74%)	38.52 ± 9.28	39.85 ± 8.64
	<i>F</i>		1.988	3.960		1.294	0.663		1.360	3.404
	<i>P</i> value		0.138	0.020		0.276	0.516		0.258	0.035

Data of STAI scores are mean ± SD.

The values highlighted in bold indicate statistical significance $P < 0.05$.

found between the three genotypes of each SNP and state-anxiety levels. There were no associations between these genotypes and either state- or trait-anxiety levels in the male participants.

No previous studies have demonstrated a direct link between *PPP1R1B*/DARPP-32 and anxiety despite some indirect evidence. There is a large amount of anatomical, physiological, pharmacological, and behavioral evidence for the importance of DARPP-32 regulation in the DA system [1]. Clinical studies on the influence of gene variations in the DA system on anxiety have demonstrated that polymorphisms of the DA transporter catechol-O-methylaminotransferase and monoamine oxidase type A genes are associated with anxiety spectrum disorders [10–12]. Therefore, the *PPP1R1B* gene and DARPP-32 may play a role in normal and/or pathological anxiety through DA neuronal pathways. Moreover, animal experiments have demonstrated that physiological or psychosocial stress induces anxious behaviors through altered *PPP1R1B* gene expression [13, 14]. It is known that psychological factors including cognition, reinforcement learning, emotion processing, personality traits, and defense mechanisms act as mediators or moderators of the effects of stressors on individual anxiety responses. Of interest, the rs879606 and rs3764352 genotypes, demonstrated to be associated with anxiety in the present study, are related to cognition [5], personality traits [8], and defense mechanisms [9], while the rs879606G/rs3764352T/rs907094A haplotype is involved in associative emotional learning [15]. Moreover, the rs907094 genotype of the *PPP1R1B* gene has been associated with reward learning [16] and the processing of anger [17]. The *PPP1R1B* gene and DARPP-32 may thus indirectly regulate stress-induced anxiety levels through the above mediation processes.

In the present study, we investigated the possible association between three *PPP1R1B* polymorphisms and trait- and state-anxiety dimensions in healthy Han Chinese participants, and demonstrated significant differences in the trait-anxiety dimension among the rs879606A/G and rs3764352A/G genotypes, especially in the female participants. Our results are consistent with previous findings and further complement them. Recently, Kunii *et al.* [2] found that the expression of a truncated DARPP-32 transcript (t-DARPP-32) is increased in the DLPFC of patients with schizophrenia or bipolar disorder and is strongly associated with the rs879606, rs90974, and rs3764352 SNPs, as well as the previously identified 7-SNP haplotype [5], indicating that variations in the *PPP1R1B* gene may affect the abundance of t-DARPP-32 mRNA and reflect the potential molecular mechanisms underlying some mental disorders. Therefore, further studies examining the expression of *PPP1R1B* mRNA as affected by variations in the *PPP1R1B* gene are necessary to gain a more complete

understanding of the effects of *PPP1R1B* allelic variants on anxiety.

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