

5-HT2a Receptor Polymorphism Gene in Bipolar Disorder and Harm Avoidance Personality Trait

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The purpose of this study was to investigate the relationship between bipolar disorder and the harm avoidance personality trait (HA), and the genetic contribution of the polymorphic DNA variation T102C in exon 1 of 5-HTR2a (chromosome 13q14–21) in bipolar disorder and HA personality trait. Forty bipolar patients and 89 normal subjects completed the TPQ questionnaire and were genotyped for 5-HT2a. Bipolar patients scored higher than normal subjects on the HA dimension. However, no contribution of the 5-HTR2a polymorphism on the bipolar disorder or on the HA personality trait emerged. Despite the limited sample size, these results exclude a major effect of the 5-HTR2a polymorphism on bipolar disorder and HA personality trait but not a minor effect. Am. J. Med. Genet. (Neuropsychiatr. Genet.) 96:360–364, 2000. © 2000 Wiley-Liss, Inc.

KEY WORDS: personality; 5-HT2A; association study; harm avoidance; bipolar disorder

INTRODUCTION

Cloninger and collaborators [Cloninger, 1986, 1987a, 1988, 1997; Cloninger et al., 1993] have brought to-

gether neurobiology, genetics, psychophysiology, cognitive processes, and clinical syndromes in an original model of personality. According to this model, personality is the result of specific combinations of heritable and stable dimensions, associated with basic emotional dispositions and learning processes, and related to neuroanatomic and neurochemical processes in the brain. Three distinct personality dimensions have been defined: novelty seeking (NS), harm avoidance (HA), and reward dependence (RD). These dimensions are proposed to reflect activity in the three major brain systems underlying activation, inhibition, and maintenance behaviors. Dopaminergic, serotonergic, and noradrenergic circuits are the neurochemical substrates of these brain systems, respectively. Cloninger hypothesized positive correlations between serotonergic activity and HA, dopaminergic activity and NS, and noradrenergic activity and RD.

The involvement of the serotonergic system in HA has already been suggested [Hansenne et al., 1997; Lesch et al., 1996; Ricketts et al., 1998; Sander et al., 1998] whereas negative results also have been reported in healthy subjects [Ebstein et al., 1997a,b; Gelertner et al., 1998] and depressed patients [Staner et al., 1998]. To our knowledge, the specific contribution of the 5-HT2a receptor polymorphism in HA personality trait has never been investigated. However, negative associations between 5-HT2a receptor polymorphism and bipolar depression were reported [Gutierrez et al., 1995; Mahieu et al., 1997].

Because previous research showed that bipolar patients score higher on the HA dimension compared to normal controls [Osheret et al., 1996; Strakowski et al., 1993; Young et al., 1995], the aims of the present study were to replicate this finding, to investigate the contribution of the 5-HT2a receptor polymorphism in bipolar depression, and to compare the contribution of the

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5-HT2a receptor polymorphism in HA personality traits in bipolar patients and normal subjects.

obtained before drawing venous blood for 5-HTR2a genotyping.

MATERIALS AND METHODS

Tridimensional Personality Questionnaire

The Tridimensional Personality Questionnaire (TPQ) by Cloninger [1987b], a 100-item, true-false test, assesses three higher order dimensions of personality, NS, HA, and RD and 12 lower order traits (see Table I).

Description of the Sample

Fourteen male and 26 female bipolar patients, aged 43.15 years ($SD = 13.87$), were recruited at the Centre Hospitalier de Luxembourg (Grand Duchy of Luxembourg). Caucasian origin was confirmed before completing the TPQ self-questionnaire. Bipolar disorder was diagnosed according to the Research Diagnosis Criteria (RDC) [Spitzer et al., 1978], using the Schedule for Affective Disorders and Schizophrenia-Lifetime Version [Spitzer and Endicott, 1979]. Patients were considered as being in at least a 2-month euthymic period by the clinician in charge. In addition, euthymia was confirmed with SADS part-1 items assessing the mood of the current state and the mood for the 2 previous months (instead of previous week) by an independent rater. This procedure provided material to rule out an RDC affective disorder diagnosis such as hypomania and mania, and major, minor, or intermittent depression in the 2 preceding months. Further, 54 female and 35 male Caucasian control subjects without personal or family history of psychiatric disorder were recruited among medical personnel at Erasme Hospital (Brussels, Belgium). Mean age was 40.82 years ($SD = 12.74$). No significant age difference emerged between patients and controls, $F(1, 127) = .873, P > .05$. The procedure and aims of the study were fully explained to the participants, and written, informed consent was

DNA Analysis

The 5-HT2a receptor gene has been located on chromosome 13 (13q14–21), spans over 20 kb, and consists of three exons interrupted by two introns [Chen et al., 1992]. A 5-HT2a receptor DNA polymorphism (T/C, substitution of C for T at position 102) has been identified within the coding exonic 1 region [Warren et al., 1993]. The T/C substitution does not alter the amino-acid sequence (serine, at position 34) and is unlikely to produce functional effects (silent mutation).

Genomic DNA was isolated from peripheral blood leukocytes using standard salting out procedures. The polymorphism in 5-HT2a was identified by the polymerase chain reaction (PCR) followed by restriction enzyme digestion. Standard PCR was carried out in a 25 μ l volume containing 100 ng genomic DNA, 200 μ M of each dNTP, 1.25 mM $MgCl_2$, 50 pmol of each primer and 0.2 units Goldstar DNA polymerase (Eurogentec, Seraing, Belgium). Published primer sequences were used [Warren et al., 1993]. After an initial denaturation step at 94°C for 2 min, 30 cycles were performed: denaturation at 94°C for 1 min, annealing at 60°C for 1.5 min, and extension at 72°C for 2 min. An additional final extension step was performed at 72°C for 5 min. Twenty μ l of the PCR product was digested overnight at 37°C with 0.1 units/ μ l of *MspI* in a total volume of 25 μ l. Digestion products were visualized by ethidium bromide staining after electrophoresis in a 3% agarose gel.

RESULTS

Personality Differences Between Bipolar Patients and Controls

A MANOVA with scores on the TPQ scales (HA, NS, and RD) as a within subject factor, and group (patients vs. controls) and sex as between subjects factors was conducted. The results revealed significant main effects of TPQ scales, $F(2, 126) = 8.50, P < .001$, and group, $F(2, 127) = 7.38, P = .008$. These main effects were qualified by a significant TPQ Scales \times Group interaction, $F(2, 126) = 11.40, P < .001$. No significant main effect or interaction involving sex emerged. Post hoc analyses showed that bipolar patients scored higher than normal subjects on the HA scales, $F(1, 127) = 23.73, P = .001$. No other significant differences emerged for TPQ scores between bipolar patients and normal subjects. Means for bipolar patients and normal subjects are shown in Table II. Thus, the results support the notion that personality differences, in terms of HA, are observed between euthymic bipolar patients and normal subjects.

Finally, ANOVAs were conducted on the scores of the four subscales (HA1, HA2, HA3, and HA4) that compose the HA dimension (see Table I). Means for this analysis are shown in Table III. The results revealed a significant main effect of group for the HA1, HA2, and HA4 subscales, $F(1, 127) = 10.45, P = .002; F(1, 127)$

TABLE I. TPQ Scales and Subscales

Novelty seeking (NS)
NS1: exploratory excitability versus stoic rigidity (9 items)
NS2: impulsiveness versus reflection (8 items)
NS3: extravagance versus reserve (7 items)
NS4: disorderliness versus regimentation (10 items)
NS = NS1 + NS2 + NS3 + NS4 (24 items)
Harm avoidance (HA)
HA1: anticipatory worry versus uninhibited optimism (10 items)
HA2: fear of uncertainty versus confidence (7 items)
HA3: shyness with strangers versus gregariousness (7 items)
HA4: fatigability and asthenia versus vigor (10 items)
HA = HA1 + HA2 + HA3 + HA4 (34 items)
Reward dependence (RD)
RD1: sentimentality versus insensitiveness (5 items)
RD2: persistence versus irresoluteness (9 items)
RD3: attachment versus detachment (11 items)
RD4: dependence versus independence (5 items)
RD = RD1 + RD2 + RD3 + RD4 (30 items)

TABLE II. Means and Standard Deviations of Harm Avoidance (HA), Novelty Seeking (NS), and Reward Dependence (RD) Scores in Function of Group

TPQ	Group	Means	SD
HA ^a	Bipolar	17.27	7.43
	Controls	11.44	5.72
	Total	13.29	6.82
NS ^b	Bipolar	14.78	4.88
	Controls	16.09	4.44
	Total	13.29	6.82
RD ^b	Bipolar	17.05	4.36
	Controls	17.03	4.00
	Total	17.04	4.10

^aSignificant difference between bipolar patients and control subjects.

^bNo significant difference between bipolar patients and control subjects.

= 15.57, $P < .001$; and $F(1, 127) = 32.65$, $P < .001$, respectively. Inspection of the means showed that bipolar patients scored higher than normal subjects. Furthermore, inspection of the effect size (η^2) revealed that HA4 subscales explained more variance than the HA1 and HA2 ($\eta^2 = .205$, $\eta^2 = .076$, and $\eta^2 = .109$, respectively). Thus, the difference between bipolar patients and normal subjects on HA score is essentially explained by the fatigability–asthenia versus vigor subscales.

TABLE III. Means and Standard Deviations of the Four Subscale Scores of Harm Avoidance (HA) in Function of Group

Group	Subscales	Means	SD
Bipolar	HA1 ^a	4.68	2.74
	HA2 ^a	4.95	1.54
	HA3 ^a	2.80	2.29
	HA4 ^b	4.85	3.04
Control	HA1 ^a	3.20	2.22
	HA2 ^a	3.63	1.85
	HA3 ^a	2.42	1.76
	HA4 ^b	2.19	2.13

^aSignificant difference between bipolar patients and control subjects.

^bNo significant difference between bipolar patients and control subjects.

Contribution of 5-HTR2a Receptor Polymorphism in Bipolar Depression and Harm Avoidance Personality Trait¹

To investigate the contribution of the 5-HTR2a receptor polymorphism in bipolar depression, allele frequencies, genotype count, and homozygotes–heterozygotes distributions of bipolar patients were compared to those of normal subjects by chi-square. The results yielded no significant differences for allele frequencies, genotype count, or homozygotes–heterozygotes distribution, $\chi^2(1) = .824$, $P > .05$; $\chi^2(2) = 1.43$, $P > .05$, and $\chi^2(1) = .342$, $P > .05$, respectively. Frequencies for these analyses are shown in Table IV, and power to detect associations given the sample size is shown in

¹The sample has been tested for Hardy and Weinberg equilibrium. The analyses did not reveal significant differences between the expected values and the observed values. Thus, the sample (patients and controls) can be considered as homogenous.

TABLE IV. Allele Frequencies, Genotype Counts, and Homozygote–Heterozygote Distribution for 5-HTR2a in Bipolar Patients and Normal Subjects

5-HTR2a	Bipolar	Normal
Alleles		
1	32 (0.40)	82 (0.46)
2	48 (0.60)	96 (0.54)
Genotypes		
1–1	5 (0.12)	19 (0.21)
1–2	22 (0.55)	44 (0.49)
2–2	13 (0.33)	26 (0.30)
Homozygotes–heterozygotes		
Homozygotes	18 (0.45)	45 (0.51)
Heterozygotes	22 (0.55)	44 (0.49)

Table V. These analyses suggested no association between bipolar depression and 5-HTR2a receptor polymorphism.

To investigate the contribution of the 5-HTR2a receptor polymorphism on HA in bipolar patients and normal subjects, a MANOVA with group (patient vs. controls) and allele 1 (carriers vs. noncarriers) as between subject factors and a MANOVA with group (patient vs. controls) and allele 2 (carriers vs. noncarriers) as between subject factors were conducted on the general score as well as on subscale scores of the HA dimension. The analyses did not reveal any significant main effect or interaction involving allele 1 or allele 2. Further, to investigate the contribution of the 5-HTR2a genotype, a MANOVA with group (patient vs. controls) and 5-HTR2a genotype (1–1, 1–2 vs. 2–2) as between subject factors was conducted on the general score of the HA dimension as well as on the subscales scores of HA. The results did not reveal any significant main effect or interaction involving genotype. Power to detect a main effect of genotype and a Genotype \times Diagnosis interaction on the HA dimension given the sample size is shown in Table VI.

DISCUSSION

Higher scores on the HA dimension were found for bipolar patients compared to normal subjects. This finding is consistent with those of other investigators [Osheret et al., 1996; Strakowski et al., 1993; Young, 1995]. Moreover, the difference on HA score is essentially due to a higher fatigability and asthenia for bipolar patients. Although our patients were in remis-

TABLE V. Power to Detect Associations Between Allele Frequencies, Genotype Count, Homozygote–Heterozygote Distribution, and the Bipolar Disorder Given the Sample Size*

	Small effect	Medium effect	Large effect
Alleles ^a	0.20	0.92	0.99
Genotypes ^b	0.15	0.87	0.99
Homozygotes			
Heterozygotes ^a	0.20	0.92	0.99

*n = 129. Effect size conventions according to Faul and Erdfelder [1992]: small effect size = 0.10, medium effect size = 0.30, and large effect size = 0.5.

^aChi-square, df = 1.

^bChi-square, df = 2.

TABLE VI. Power to Detect a Main Effect of Genotype, and a Genotype–Diagnosis Interaction on the HA Dimension Given the Sample Size*

	Small effect	Medium effect	Large effect
Main effect of genotype ^a	0.35	0.99	1.00
Genotype by diagnosis interaction ^b	0.23	0.96	1.00

*n = 129. Effect size conventions according to Faul and Erdfelder [1992]: small effect size = 0.02, medium effect size = 0.15, and large effect size = 0.35.

^aMANOVA, df = 1, 127.

^bMANOVA, df = 3, 125.

sion, one cannot exclude the possibility that higher HA scores could also be due to residual symptoms and/or concomitant treatment. No further personality differences were found using the TPQ.

From these results, no genetic contribution of 5-HT2a receptor polymorphism to bipolar disorder or to HA personality trait in normal and bipolar subjects emerged. Our results have to be interpreted with caution, however. First, it must be mentioned that the sample had an adequate statistical power to detect a medium and major effect gene whereas for a minor effect gene the power is small (see Tables V and VI). Thus, although one can exclude medium and major effects of the 5-HTR2a receptor polymorphism in bipolar disorder and HA personality, our sample was too small to detect a minor gene effect [Owen, 1997]. Consequently, one cannot exclude the possibility of a small effect of the 5-HTR2a receptor polymorphism; such an effect could be investigated with a larger sample. Second, complex behavior is most likely the result of the interaction between various neurotransmitter systems. The present study has investigated only the 5-HTR2a receptor polymorphism, and it could be argued that this polymorphism might contribute to personality differences only in the context of interactions with other mediators. In order to interpret personality differences, future research should investigate polygenic models. Third, because the 102-T/C variant does not cause an amino acid substitution in the receptor protein, it is unlikely to affect receptor function or expression, which may be directly related to differences in the HA dimension. On the other hand, it has been shown that this 102-C/T polymorphism is in almost complete linkage disequilibrium with a –1438-G/A promoter polymorphism [Arranz et al., 1998], which might have an effect on receptor expression level. Due to the high linkage disequilibrium between these two markers, it does not seem likely that the –1438-G/A polymorphism largely influences the variance in HA. Furthermore, other known functional polymorphisms seem to be rare in the general population [Gutierrez et al., 1997], making them unlikely candidates for HA variation in the general population.

In conclusion, a relationship between the HA dimension and bipolar disorders was observed in our study. We did not find evidence for any relationship between the bipolar phenotype or HA and a nonfunctional variation in the 5-HTR2A gene. Additional data are needed with other candidate genes to further clarify the rela-

tionships between the serotonergic system and various personality dimensions in mood disorders.

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