

Serotonin Transporter Promoter Polymorphism Genotype Is Associated With Behavioral Disinhibition and Negative Affect in Children of Alcoholics

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Background: Serotonergic (5-HT) dysfunction has been implicated in the etiology of both behavioral disinhibition (BD) and negative affect (NA). This work extends our previous finding of relationships between whole blood 5-HT and both BD and NA in pubescent, but not prepubescent, children of alcoholics and continues examination of a hypothesized role of 5-HT dysfunction in alcoholism risk. The long and short (L and S) variants of the 5-HT transporter gene-linked polymorphic region (5-HTTLPR) are responsible for differing transcriptional efficiencies in 5-HT uptake. Although associations have been found between the SS 5-HTTLPR genotype and severe alcoholism and neuroticism, recent reports describe relationships between the LL genotype and both low level of response to alcohol and alcoholism diagnosis and a predominance of the LL genotype in early-onset alcoholics.

Methods: This report is from an ongoing prospective study of the development of risk for alcoholism and other problematic outcomes in a sample of families classified by father's alcoholism subtype. This study examines relationships between 5-HTTLPR genotype and both child BD (Child Behavior Checklist Aggressive Behavior) and NA (Child Behavior Checklist Anxious/Depressed) in offspring from 47 families.

Results: Results showed significantly higher levels of BD and NA in the 16 children with the LL genotype than the 46 SS or SL children.

Conclusions: Behaviors of undercontrol, which occur at increased rates in children of alcoholics, may be genetically influenced through the regulation of the 5-HT transporter. Due to the small sample size and the preliminary nature of our findings, replication is necessary.

Key Words: Serotonin Transporter, Behavioral Disinhibition, Negative Affect, Children of Alcoholics, Genetic.

SEROTONERGIC (5-HT) DYSFUNCTION as measured by central and peripheral measures has been consistently implicated in both behavioral disinhibition (BD; impulsive aggression and type II alcoholism) and negative affect (NA; depression and anxiety) in clinical and nonpatient samples of adults and children (Brown and Linnoila, 1990; Coccaro et al., 1997; Fils-Aime et al., 1996; Johnson et al., 2000; Lesch et al., 1996; Linnoila et al., 1983;

Manuck et al., 1998; Nobile et al., 1999; Pfeffer et al., 1998; Sander et al., 1997; Stoff and Vitiello, 1996; Twitchell et al., 2000; Virkkunen et al., 1996). Many psychiatric genetic studies have suggested that a functional polymorphism in the regulatory region of the 5-HT transporter gene-linked polymorphic region (5-HTTLPR) may be a strong candidate gene for behavioral disorders such as alcoholism, anxiety, and depression (Greenberg et al., 2000; Hallikainen et al., 1999; Kaufman et al., 1998; Lesch et al., 1996; Lichtenmann et al., 2000; Nobile et al., 1999; Sander et al., 1997; Schuckit et al., 1999; Turker et al., 1998), although some studies have produced negative results (Edenberg et al., 1998; Gelernter et al., 1998; Jorm et al., 1998; Ohara et al., 1998).

Although children of alcoholics (COAs) are at increased risk for behavioral and affective dysregulation (Sher, 1991, 1997), few studies have examined the potential role 5-HT may play in their development (Twitchell et al., 1998, 2000). Furthermore, to our knowledge, no study has examined associations between 5-HTTLPR genotype and both BD and NA, two characteristics that may place these children at risk for the development of alcoholism. Several studies have documented 5-HT dysfunction in men with positive family histories of alcoholism (LeMarquand et al., 1999;

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Rausch et al., 1991). Additionally, in a study of abstinent adult alcoholics and their adult and young children, increased 5-HT uptake was found in the alcoholics compared with controls, as well as in the alcoholics' young children (mean age, 11 ± 1.1 years), most of whom had not ever consumed alcohol, in comparison with age- and sex-matched control children (Ernouf et al., 1993).

Previously in our own study of COAs, we found lower whole blood 5-HT content in six children who exhibited overt behavior problems as measured by clinical range Child Behavior Checklist (CBCL; Achenbach, 1991) Total Behavior Problem scores in comparison with 38 children with CBCL Total Behavior Problem scores in the normal or borderline range (Twitchell et al., 1998). In a subsequent study of 62 children (original sample with the addition of 18 children) subtyped by paternal alcoholism classification, results indicated that puberty moderated relationships between 5-HT and both BD and NA, with relationships observed in pubescent, but not prepubescent, children (Twitchell et al., 2000).

5-HTTLPR GENOTYPE IS RELATED TO FUNCTIONAL DIFFERENCES IN 5-HT UPTAKE

It has been determined that the long and short (L and S) variants of the 5-HTTLPR are responsible for differing transcriptional efficiencies in 5-HT uptake, one component of 5-HT functioning (Crawchik and Goldman, 2000). The LL variant has been associated with three times greater 5-HT uptake in lymphocytes (Lesch et al., 1996) and blood platelets (Greenberg et al., 1999), in comparison to the SS or heterozygous SL variants. Consequently, the LL variant is expected to be related to increased 5-HT transporter number and function, reduced levels of intrasynaptic 5-HT, and reduced overall 5-HT function (Johnson, 2000). In a recent study with depressed children and adolescents which examined blood platelet measures of 5-HT function, control children with the LL genotype had greater initial 5-HT uptake than control children with SS or SL genotypes (Nobile et al., 1999).

5-HT transporter function in the brain has also been used as a model system for studying potential functional differences among 5-HTTLPR genotypes. In postmortem midbrain samples of ethanol users, 5-HT transporter binding and messenger RNA levels were found to vary by 5-HTTLPR genotype in patterns consistent with the previously reported pattern of the LL genotype being associated with higher function than SS or SL genotypes (Little et al., 1998). In vivo imaging of 5-HT transporter availability in alcoholics and healthy controls further supports functional differences between those with the LL genotype in comparison to those with either the SS or LS genotype, with an interaction observed between diagnosis and 5-HTTLPR genotype on 5-HT transporter availability (Heinz et al., 2000). Among controls, the LL genotype was related to an increase in 5-HT transporter availability in comparison to S

carriers. Among alcoholics, the LL genotype was associated with reduced 5-HT transporter availability. However, in a recent study of major depression and suicide, no relationship was observed between the 5-HTTLPR genotype and level of 5-HT transporter binding in the prefrontal cortex (Mann et al., 2000).

5-HTTLPR GENOTYPE AS A CANDIDATE GENE FOR NEGATIVE AFFECT AND BEHAVIORAL DISINHIBITION

Numerous studies have followed the initial demonstration that the S allele is related to a trait associated with anxiety, hostility, and depression (Lesch et al., 1996), although many studies have failed to replicate this small effect size finding (Gelernter et al., 1998; Jorm et al., 1998; Ohara et al., 1998). The relationship between 5-HT dysfunction and impulsive aggression is also well documented (Linnoila et al., 1983; Virkkunen et al., 1989). That 5-HT dysfunction is implicated in several psychiatric disorders in which behavioral and affective dysregulation is a key component (e.g., obsessive-compulsive disorder, depression, and antisocial alcoholism) suggests that any factor that modulates 5-HT function may be considered an important candidate gene (Stoltenberg and Burmeister, 2000).

Data regarding the relationships between 5-HTTLPR genotype and alcoholism are complicated and often conflicting (Edenberg et al., 1998; Hallikainen et al., 1999; Hammoumi et al., 1999; Johnson et al., 2000; Jorm et al., 1998; Sander et al., 1998). Although a large candidate gene family-based association analysis found no association between 5-HTTLPR genotype and alcoholism diagnosis (Edenberg et al., 1998), recent results from another large candidate gene family-based association analysis provide support for allelic association of the 5-HTTLPR short variant with alcohol dependence (Lichtermann et al., 2000). These discrepancies may reflect the heterogeneity of the disorder, the complexity inherent in psychiatric genetic studies, or both of these. Given that alcoholism is a heterogeneous and polygenic disorder, it may be necessary to study subtypes separately. For example, type II, or antisocial, early-onset alcoholism (Cloninger et al., 1981), which frequently includes impulsive aggressive behavior, has been found to have a stronger genetic basis, perhaps due to the presence of impulsive aggression, a behavior that has been reliably associated with 5-HT dysfunction.

To follow up on our previous findings of relationships between a peripheral index of 5-HT function and behavior, this study was designed to examine 5-HTTLPR as a candidate gene for behavioral and affective dysregulation in COAs. Given the reported associations between 5-HTTLPR genotype and NA, low level of response to alcohol, and alcoholism, we predicted that relationships would be found between 5-HTTLPR genotype and both BD and NA.

Table 1. Sampling Design

Father's alcoholism classification	Child 5-HTTLPR genotype			
	LL (n = 16)	SL (n = 33)	SS (n = 13)	SS/SL (n = 46)
Antisocial alcoholic (n = 17)	Sons = 4	Sons = 8	Sons = 4	Sons = 12
	Daughters = 1	Daughters = 4	Daughters = 0	Daughters = 4
Nonantisocial alcoholic (n = 20)	Sons = 5	Sons = 12	Sons = 2	Sons = 14
	Daughters = 2	Daughters = 2	Daughters = 3	Daughters = 5
Nonalcoholic control (n = 10)	Sons = 2	Sons = 5	Sons = 3	Sons = 8
	Daughters = 2	Daughters = 2	Daughters = 1	Daughters = 3

METHODS

Sample

Subjects were 62 non-Hispanic white children without fetal alcohol syndrome between the ages of 7 and 16 years (45 boys, 17 girls; mean age, 10.9 ± 2.0 years) from a subsample consisting of 47 lower- to lower-middle-class families drawn from the ongoing Michigan State University–University of Michigan Longitudinal Study (Fitzgerald et al., 1995; Zucker, 1987; Zucker et al., 1996, 1997). The larger study from which this sample is drawn is observing a population-based sample of COAs, both biological parents, and demographically similar nonalcoholic control families at 3-year intervals, beginning when the male target children were ages 3 to 5 years and the female target children were between ages 3 and 11 years. Paternal alcoholism diagnosis was assessed with Feighner diagnostic criteria (Feighner et al., 1972) within a structured clinical interview format. Inclusion criteria for the larger study involved having a biological son between 3.0 and 5.0 years of age living in an intact family with both biological parents. Maternal alcoholism was neither inclusionary nor exclusionary, although a subset of the women married to antisocial alcoholic men (41%) met a lifetime DSM-III-R alcohol abuse or dependence diagnosis.

Design for Alcoholism Subtyping. The sampling design for this study is presented in the left column of Table 1. Families were subtyped by paternal alcoholism and antisocial personality disorder diagnoses in earlier analytic work with the longitudinal study (cf Poon et al., 2000; Wong et al., 1999; antisocial alcoholism, nonantisocial alcoholism, nonalcoholic control). For this study, we deliberately selected families to achieve an approximately equal distribution across the three groups. Children were later categorized by 5-HTTLPR genotype. Fifteen of the 47 families had both a son and a daughter in the appropriate age range; in these instances, both children were included.

5-HTTLPR Genotype. Genotyping methods have been described previously. Please see Hanna et al. (1998) for a full description of methods used here. There is some evidence for dominance of the S allele (Greenberg et al., 1999; Lesch et al., 1996). Consequently, SS and SL genotypes have frequently been combined. In our analyses we use this approach, which also has the benefit of increasing statistical power. In addition, we provide data for child behavioral measures for all three genotypes so that the data are more informative to other researchers.

Assessing Child Behavioral Disinhibition and Negative Affect

The 4- to 18-year-old version of the CBCL (Achenbach, 1991) was used to gather child behavioral data. The CBCL is the most commonly used questionnaire for clinical classification of child behavior problems in the United States. Maternal ratings obtained at the most recent wave of data collection were used. Test-retest reliability of item scores on the CBCL ranging from 0.95 at a 1-week interval to 0.84 at a 3-month interval and adequate construct validity have been demonstrated (Achenbach, 1991).

Child BD. The Aggressive Behavior scale from the CBCL was used as a behavioral phenotype indicator of BD (i.e., Argues a lot; Cruelty, bullying, or meanness to others; Gets in many fights; Screams a lot; Temper tantrums or hot temper; Threatens people).

Child NA. The CBCL Anxious/Depressed scale was used as a behavioral phenotype indicator of NA (i.e., Complains of loneliness; Cries a lot;

Nervous, highstrung or tense; Unhappy, sad or depressed; Too fearful or anxious).

Evaluating Child Alcohol Consumption

Children reported their alcohol use on a brief questionnaire that was administered privately and confidentially. Nine children (six boys, three girls) reported having consumed alcohol at some point in their lives (15%). However, all of these children denied having consumed any alcohol within the past 4 weeks. Therefore, this sample is composed of children who have not yet begun regular drinking.

Medication Screening

At the time of blood draw, the primary caretaking parent (95% mothers) completed a child health history that assessed child's height, weight, and the presence of any psychotropic medications taken within the past month.

Demographic Variables

Families were characterized by demographic questionnaire measures of socioeconomic status, parental years of education, family income, and child age. Parental socioeconomic status was coded with the Revised Duncan Socioeconomic Index (Duncan TSE12; Stevens and Featherman, 1981). Parental education was measured by the total years of academic or vocational education achieved (scores ranged from 9 to 20 years of education).

Procedure

Approval from the appropriate institutional review boards was obtained prior to study implementation. Medication screening and blood sampling were performed by GRT during an in-home session. Informed consent was obtained from all study participants before data collection. All subjects donated their blood samples and were paid for filling out the questionnaires.

Statistical Analysis

Group differences were tested with independent sample *t*-tests. All reported probabilities are two tailed.

RESULTS

The sociodemographic characteristics of the sample are shown in Table 2.

5-HTTLPR Genotype is Related to Both Behavioral Disinhibition and Negative Affect in COAs

The *t*-tests showed significantly higher levels of BD and NA in the 16 children with the LL genotype ($t = -2.35, df = 60, p < 0.05; t = -2.52, p < 0.05$, respectively) than the

Table 2. Sociodemographic Characteristics of Sample ($N = 62$)

Sociodemographics	Mean	SD
Child age (yr)	10.88	2.03
Father's years of education	14.03	2.52
Mother's years of education	13.47	2.24
Father's SES ^a	37.61	21.25
Mother's SES ^a	30.56	12.92
Family income (US \$)	46,250.00	25,316.00

^a Duncan TSE12 (Stevens and Featherman, 1981); SES, socioeconomic status.

46 SS or SL children. When analyses were re-run with COAs only, the pattern of results was the same.

The presence of 15 sibling pairs within the 47 families violates the assumption of independence. To address this issue, five different random samples were drawn such that only one child from each family was included. Four of these five samples yielded significant relationships for both BD and NA. For the remaining sample, the results were not significant, although the p values were 0.10 and 0.07 for Aggressive Behavior and Anxious/Depressed, respectively.

Post hoc exploratory analyses evaluated the relationships between 5-HTTLPR genotype and the remaining nine CBCL scales and showed significant differences between LL and SS or SL genotypes for Internalizing, Social Problems, Externalizing, and Total Behavior Problems. The same pattern of findings emerged such that the LL genotype children exhibited higher means, indicating more problems on each CBCL scale (Table 3). The relationship between child 5-HTTLPR genotype and any child alcohol consumption was evaluated with χ^2 . Of the LL genotype children, 31.3% had consumed alcohol at some point in their lives. In comparison, only 8.7% of the SS or SL genotype children had consumed alcohol at any point in their lives. This relationship was significant ($\chi^2 = 4.87$, $df = 1$, $p < 0.05$).

DISCUSSION

The primary finding was that child 5-HTTLPR genotype was related to both BD and NA with the LL variant of the 5-HTTLPR genotype associated with increased levels of BD and NA. This finding lends support to the hypothesis that behavioral undercontrol found at increased rates in COAs may be genetically influenced through the regulation of the 5-HT transporter.

Our results are consistent with findings in the adult literature suggesting that 5-HT function plays a role in a spectrum of behaviors marked by undercontrol. Furthermore, the finding of a relationship between 5-HTTLPR genotype and BD in children at risk for later problems in behavioral and affective regulation is consistent with the accepted theory of 5-HT's role as a general inhibitor of motoric behaviors such as impulsive aggression. Similarly, our finding of a relationship between 5-HTTLPR genotype and NA is consistent with the more recent suggestion that 5-HT may also play a role in affective expression (Lesch et

al., 1996; Lesch and Mossner, 1998; Lucki, 1998; Mazzanti et al., 1998).

Post hoc exploratory analyses provided further support for these conclusions. Independent sample t -tests on all CBCL scales revealed significant relationships between 5-HTTLPR genotype and 6 of the 11 CBCL scales, with relationships consistently observed in the areas of behavioral (Aggressive Behavior, Externalizing, Total Behavior Problems) and affective (Anxious/Depressed, Internalizing) dysregulation. A similar relationship was observed for child genotype and social functioning (Social Problems), providing further support for the direct and moderating effect of 5-HT on social behavior in both human (Knutson et al., 1998; Kruesi et al., 1990) and nonhuman (Higley et al., 1996; Mehlman et al., 1995; Raleigh et al., 1983, 1985) primates.

Furthermore, a χ^2 analysis of child 5-HTTLPR genotype and alcohol consumption indicated that significantly more LL genotype than SS or SL genotype children reported that they had already consumed alcohol (i.e., they had a significantly earlier onset of alcohol use). Taken together, our findings of greater behavioral and affective dysregulation and higher rates of first-time alcohol consumption in children with the LL genotype of 5-HTTLPR provide additional support for the hypothesized role of 5-HT as a potential neurobiological substrate for alcohol-related risk factors and potential development of early-onset, antisocial alcoholism. The genotypic characterization of these children early in their lives documents important biological differences and supports observing these children for the occurrence of problem drinking or alcoholism in early adolescence and adulthood.

A limitation of this study was that it did not use a family-based approach (e.g., quantitative transmission/disequilibrium test; Abecasis et al., 2000). In addition, given the size of the study sample, the observed statistical power to detect differences between genotypes was low (i.e., less than 50%; two-way $\alpha = 0.05$) for the majority of the CBCL scales. Although the sample is composed of children from a relatively ethnically homogeneous group (non-Hispanic, white families), we cannot rule out population stratification (Pritchard et al., 2000). Consequently, these findings should be considered to be preliminary and interpreted with caution. Finally, although we report associations between the LL genotype of 5-HTTLPR and BD and NA, some studies have reported relationships between the SS genotype and neuroticism (Lesch et al., 1996), alcoholism risk (Lichtermann et al., 2000), and antisocial alcoholism (Hallikainen et al., 1999; Sander et al., 1998). It is important to note that our findings are consistent with those of Schuckit et al. (1999), who report a relationship between the LL genotype of 5-HTTLPR and both alcoholism risk as measured by low level of response to alcohol and subsequent alcoholism diagnosis and the recent finding of a predominance of the LL variant among early-onset alcoholics (Ishiguro et al., 1999; Johnson, 2000).

Table 3. Child Behavioral Measures as a Function of 5-HTTLPR Genotype

Variable	5-HTTLPR genotype								t-test, <i>df</i> = 60; LL vs. SS/SL
	LL (<i>n</i> = 16)		SL (<i>n</i> = 33)		SS (<i>n</i> = 13)		SS/SL (<i>n</i> = 46)		
	Mean	SD	Mean	SD	Mean	SD	Mean	SD	
Child dependent variables: CBCL									
Aggressive Behavior	12.69	7.12	9.12	6.52	6.31	4.63	8.33	6.13	-2.35, <i>p</i> < 0.05
Anxious/Depressed	5.19	4.31	2.82	3.05	2.62	2.66	2.76	2.91	-2.52, <i>p</i> < 0.05
Child post hoc variables: CBCL									
Total Behavior Problems	35.19	21.63	24.52	16.16	19.23	11.28	23.02	15.01	-2.50, <i>p</i> < 0.05
Externalizing	14.63	8.81	10.94	7.62	7.38	5.39	9.93	7.19	-2.12, <i>p</i> < 0.05
Delinquent Behavior	1.94	1.95	1.82	1.53	1.08	1.19	1.61	1.47	-0.71, <i>p</i> = 0.48
Internalizing	9.13	6.86	5.64	5.06	5.69	5.04	5.65	5.00	-2.17, <i>p</i> < 0.05
Withdrawn	2.00	1.75	1.70	1.96	2.00	1.96	1.78	1.94	-0.40, <i>p</i> = 0.69
Somatic Complaints	2.00	1.86	1.30	1.63	1.15	2.03	1.26	1.73	-1.44, <i>p</i> = 0.15
Social Problems	3.50	2.90	1.88	1.76	1.54	1.39	1.78	1.66	-2.90, <i>p</i> < 0.05
Thought Problems	0.63	1.50	0.52	0.83	0.23	0.60	0.43	0.78	-0.65, <i>p</i> = 0.52
Attention Problems	3.88	2.60	2.91	2.63	2.69	3.43	2.85	2.84	-1.27, <i>p</i> = 0.21

In future studies, examination of other 5-HT and neurochemical measures would be helpful in discerning the risks for alcoholism and related behavioral and affective dysregulation found in COAs. Other child/adolescent studies are also necessary to replicate these findings. That these findings were observed in a medication-free, community-recruited sample, and in an age range in which regular drinking has not yet begun, suggests that this association may generalize to the overall population.

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