Association between Dopamine Transporter (DAT1) Genotype, Left-Sided Inattention, and an Enhanced Response to Methylphenidate in Attention-Deficit Hyperactivity Disorder

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A polymorphism of the dopamine transporter gene (DAT1, 10-repeat) is associated with attention-deficit hyperactivity disorder (ADHD) and has been linked to an enhanced response to methylphenidate (MPH). One aspect of the attention deficit in ADHD includes a subtle inattention to left space, resembling that seen after right cerebral hemisphere damage. Since left-sided inattention in ADHD may resolve when treated with MPH, we asked whether left-sided inattention in ADHD was related to DAT1 genotype and the therapeutic efficacy of MPH. A total of 43 ADHD children and their parents were genotyped for the DAT1 3' variable number of tandem repeats polymorphism. The children performed the Landmark Test, a well-validated measure yielding a spatial attentional asymmetry index (leftward to rightward attentional bias). Parents rated their child's response to MPH retrospectively using a three-point scale (no, mediocre or very good response). Additionally, parents used a symptom checklist to rate behavior while on and off medication. A within-family control design determined whether asymmetry indices predicted biased transmission of 10-repeat parental DAT1 alleles and/or response to MPH. It was found that left-sided inattention predicted transmission of the 10-repeat allele from parents to probands and was associated with the severity of ADHD symptomatology. Children rated as achieving a very good response to MPH displayed left-sided inattention, while those rated as achieving a poorer response did not. Our results suggest a subgroup of children with ADHD for whom the 10-repeat DAT1 allele is associated with left-sided inattention. MPH may be most efficacious in this group because it ameliorates a DAT1-mediated hypodopaminergic state.

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INTRODUCTION

Attention-deficit hyperactivity disorder (ADHD), a common childhood disruptive disorder affecting 3–6% of school-aged children worldwide, is characterized by age inappropriate levels of inattention, hyperactivity, and impulsivity. Family, twin, and adoption studies suggest that the disorder is heritable (see Kirley *et al*, 2002 for a review). Pharmacogenetic studies in ADHD suggest that individual differences in stimulant-response may be related to underlying genetic influences (Kirley *et al*, 2003; Hamarman *et al*, 2004). The 10-repeat allele of a variable number of tandem

repeats (VNTR) situated in the 3' untranslated region of the DAT1 gene (mapping to 5p15.3) has been associated with the clinical ADHD phenotype in a number of studies (Cook *et al*, 1995; Gill *et al*, 1997; Daly *et al*, 1999). This variant may confer an enhanced therapeutic response to methylphenidate (MPH) (Kirley *et al*, 2003; but see Winsberg and Comings, 1999; Roman *et al*, 2002). Here, we ask whether an attentional phenotype is related to (a) genetic variation in the DAT1 VNTR and (b) the therapeutic efficacy of MPH in ADHD.

Several lines of converging evidence underscore the relevance of the dopamine transporter gene to ADHD. First, MPH is known to inhibit the dopamine transporter (Volkow *et al*, 1998). The dopamine transporter is highly expressed in the human striatum where it serves as the primary means of dopamine reuptake (Garris and Wightman, 1994). Second, studies of both structural and functional imaging in ADHD have consistently implicated dopamine-rich frontostriatal circuits, particularly on the

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right, in the pathophysiology of ADHD (Casey *et al*, 1997; Vaidya *et al*, 1998; Teicher *et al*, 2000). These abnormalities are ameliorated by treatment with MPH (Vaidya *et al*, 1998; Teicher *et al*, 2000). Third, *in vivo* measurement of DAT with single photon emission computed tomography (SPECT) has demonstrated elevated striatal transporter densities in adults (Dougherty *et al*, 1999; Dresel *et al*, 2000) and children (Cheon *et al*, 2003) with ADHD, compared to controls. In adults, treatment with MPH reduces transporter densities to near-normal levels (Dresel *et al*, 2000; Krause *et al*, 2000). Additionally, there is also evidence that children and adults who are homozygous for the 10-repeat DAT1 may have higher availability of DAT protein in the striatum, relative to those with the 9-repeat/10-repeat genotype (Heinz *et al*, 2000; Cheon *et al*, 2005).

These lines of evidence have led to a biological hypothesis of ADHD under which the 10-repeat DAT1 allele is thought to be associated with a greater abundance of DAT protein, resulting in a relative hypodopaminergia, perhaps particularly within the striatum (Kirley *et al*, 2002). According to this hypothesis, treatment with MPH may be most efficacious in 10-repeat homozygotes because it normalizes DAT density (Heinz *et al*, 2000; Kirley *et al*, 2003). There have, however, been a number of studies that have reported a poorer response to MPH in 10-repeat homozygotes, albeit often in relatively small samples (Rohde *et al*, 2003; Cheon *et al*, 2005).

Recent studies of susceptibility genes for psychiatric disorders have emphasized the utility of quantitative indices for assessing disease risk, termed *endophenotypes*. Endophenotypes are traits that more accurately predict dysfunction in discrete neural systems than conventional clinical phenotypes (Castellanos and Tannock, 2002). A small number of studies have examined whether genetic variants that are thought to confer susceptibility to ADHD are also associated with cognitive impairment (Swanson *et al*, 2000; Manor *et al*, 2002; Loo *et al*, 2003; Langley *et al*, 2004; Bellgrove *et al*, 2005, in press). The approach of linking a genetic risk factor with a cognitive phenotype has considerable heuristic value. Rather than seeking associations between a gene and a rudimentary diagnostic category, this approach measures the association between a gene and a strictly operationalized and objectively measured cognitive process.

In this context, we sought to determine whether the phenomenon of left-sided inattention in ADHD might be related to underlying DAT1 genotype and the therapeutic response to MPH. Left-sided inattention presents as a spatial bias of attention away from the left side. Left-sided inattention arises from dysfunction in any one of a number of right hemisphere cortical (prefrontal, parietal) and subcortical (striatal, thalamic) components of a distributed neural network for spatial attention (Mesulam, 1981). Dysfunction of the right hemisphere network results in more severe and long-lasting spatial inattention than equivalent left hemisphere dysfunction because of the dominance of the right hemisphere for the control of spatial attention: while right hemisphere networks allocate attention to both left and right hemispaces, left hemisphere networks do so only for the right hemisphere (Mesulam, 1981). Consistent with animal studies that have reported neglect consequent upon lesions of the ascending dopami-

nergic pathways (Iversen, 1984), treatment with dopamine agonists reduces the extent of neglect in human subjects (Fleet *et al*, 1987).

A number of studies have reported the presence of attentional asymmetries in ADHD using both clinical and experimental measures of attention (Voeller and Heilman, 1988; Carter *et al*, 1995; Nigg *et al*, 1997; McDonald *et al*, 1999; Sheppard *et al*, 1999). Failure to replicate these effects in some studies (Wood *et al*, 1999; Klimkeit *et al*, 2003), nevertheless, suggests neuropsychological heterogeneity. Left-sided inattention has also been reported in the biological mothers of children with ADHD (Nigg *et al*, 1997), reinforcing its candidacy as an endophenotype. Critically, there is evidence for normalization of left-sided inattention in ADHD with MPH (Nigg *et al*, 1997; Sheppard *et al*, 1999).

This study addressed two distinct but related questions. First, using a family-based genetic association design, we asked whether a continuous measure of spatial attentional asymmetry was associated with DAT1 genotype. Specifically, we first hypothesized that left-sided inattention would be associated with the 10-repeat DAT1 allele in ADHD. Second, we asked whether the same continuous measure of attentional asymmetry was associated with retrospective ratings of the therapeutic response achieved by MPH. We hypothesized that if left-sided inattention may serve as a behavioral assay for a hypodopaminergic state in ADHD, then its existence may relate to the therapeutic response conferred by MPH.

MATERIALS AND METHODS

Participants

A total of 43 right-handed ADHD participants were recruited as part of our ongoing genetic association studies and in accordance with the ethical guidelines of Trinity College Dublin and St James' Hospital, Dublin. DSM-IV diagnoses were confirmed using established diagnostic protocols. From our previously described cohort, ADHD probands (aged 6–16 years) were offered participation in the current study. The current sample is therefore not independent of our previously described cohort (Kirley *et al*, 2002, 2003). We have also recently described the neuropsychological performance, in relation to DAT1 genotype, including a distinct measure of attentional asymmetry, in a smaller subset ($n=22$) of the current cohort (Bellgrove *et al*, in press). Importantly, the current study extends the phenotypic description of our Irish sample using a different measure of attentional asymmetry, a within-family control design and ratings of MPH response. Replication with independent samples will, nevertheless, be required. Exclusion criteria included known neurological conditions including pervasive developmental disorders and epilepsy. Since reading disorder has been associated with attentional asymmetries (Facoetti *et al*, 2001), participants scoring more than $1\frac{1}{2}$ standard deviations below the mean of the reading subtest of the Wide Range Achievement Test (WRAT-3) were also excluded (Table 1). Stimulant medication was withdrawn at least 24 h prior to the neuropsychological testing.

Table 1 Clinical Characteristics of the ADHD Sample

	ADHD DAT1 Genotype Group		Associated significance test
	Low-Risk DAT1 ADHD (n = 21)	High-Risk DAT1 ADHD (n = 22)	
Gender, male, no. (%)	19 (90%)	18 (82%)	$\chi^2(df_1) = 0.67, p = 0.41$
ADHD—combined type	17 (81%)	16 (73%)	$\chi^2(df_1) = 0.41, p = 0.52$
ADHD—inattentive type	3 (14%)	4 (18%)	$\chi^2(df_1) = 0.12, p = 0.73$
ADHD—hyperactive/impulsive type	1 (5%)	2 (9%)	$\chi^2(df_1) = 0.31, p = 0.58$
	Mean (SD)	Mean (SD)	
Age	12.2 (2.7)	12.2 (2.1)	$F(1,41) = 0.01, p = 0.92$
IQ ^a	101.7 (13)	99 (12)	$F(1,39) = 0.48, p = 0.49$
WRAT Reading Standard Score ^b	96 (13)	96 (10)	$F(1,41) = 0.00, p = 0.95$
WRAT Spelling Standard Score	96 (18)	92 (10)	$F(1,41) = 0.76, p = 0.39$
DSM-IV Inattention ^c	72.1 (9.4)	77 (6.4)	$F(1,39) = 4.66, p = 0.04$
DSM-IV Hyperactive/Impulsive	76.3 (12.9)	82.3 (9.8)	$F(1,39) = 2.86, p = 0.09$
DSM-IV Total	75.9 (10.9)	82.2 (5.8)	$F(1,39) = 5.65, p = 0.02$

^aIQ was estimated using a four subtest short form of WISC-III, comprising Picture Completion, Information, Block Design, and Vocabulary. IQs were not available for two participants.

^bReading and Spelling were assessed using the Wide Range Achievement Test (WRAT). Reported scores are the standard scores for each of these subtests.

^cClinical symptomatology was assessed dimensionally using the Conners' Parent Rating Scale-Revised: Long Version. Reported values are age-related normative scores (*t*-scores) for DSM-IV Inattention, Hyperactive/Impulsive, and Total symptomatology. Scores on other dimensions are available from the authors.

Predicting Methylphenidate Response

A total of 36 of these children were currently receiving or had in the past received treatment with MPH. Ratings of MPH response were obtained from parents in two ways: first, parents were asked to rate the therapeutic response achieved with MPH on a three-point scale from 'no response' to 'mediocre' to 'very good'. Second, parents completed the Conners' Parent Rating Scale-Revised: Long Version (CPRS-R:L) (Conners, 1998) twice, retrospectively rating their child's symptoms while 'on' and 'off' MPH. We have previously reported an association between the 10-repeat allele of the DAT1 VNTR and the therapeutic response to MPH using this rating system (Kirley *et al*, 2003).

Testing Attentional Asymmetry: The Landmark Task

In this well-validated and brief (5 min) test, participants judge which end of a pre-bisected line looks shorter to them (Figure 1). Participants performed 20 trials of the Landmark Task. On 10 of these trials, the bisecting line was offset (either to the right or left) allowing accuracy of judgements to be determined. On the remaining 10 trials, the horizontal line was bisected in the middle.

DAT1 Genotyping

DNA was extracted from blood samples or buccal cells using the standard phenol-chloroform procedure from both parents and the ADHD proband in each family. Primer sequence and amplification conditions can be found elsewhere (Kirley *et al*, 2002). Two investigators, who were blind to the identity of the sample, independently scored all genotypes.

Figure 1 The Landmark Task. (a) In the Landmark Task, participants are presented with a pre-bisected line and asked which end of the line is the shorter. Healthy participants tend to show a slight bias of attention to the left owing to the dominance of the right hemisphere for spatial judgements. This scenario leads to a relative inattention to the rightward extent of the line. (b) In patients with right hemisphere lesions and left spatial inattention, there is a bias of attention to the right that arises when the dominant attentional orienting bias of the right hemisphere has been weakened. This scenario leads to a relative inattention to the leftward extent of the line (shaded area) and the subjective experience of the left end of the line as the shorter. Trials on which the left end of the line was nominated as the shorter were designated 'right-biased'. Trials on which the right end of the line was nominated as the shorter were designated 'left-biased'. A continuous measure of spatial bias—the spatial Asymmetry Index—was calculated as $(N_{\text{right-biased trials}} - N_{\text{left-biased trials}})/20$. This yielded values ranging from -1 (leftward spatial bias/right spatial inattention) to $+1$ (rightward spatial bias/left spatial inattention).

Statistical Analyses

Testing the association between DAT1 genotype and attentional asymmetry. In vitro studies indicate that the

10-repeat allele of the DAT1 VNTR may increase DAT expression (see Asherson, 2004 for review). For example, Fuke *et al* (2001) showed that the 10-repeat allele, relative to the 7- or 9-repeat alleles, increased gene expression using a reporter system. Mill *et al* (2002) also reported that mRNA levels in human brain and lymphocyte tissue varied with DAT1 VNTR, being higher in individuals with the 10- vs 9-repeat allele. Based on this evidence of differential gene expression as a function of VNTR alleles and the low frequency of individuals homozygous for the non-10-repeat alleles (less than 5% in the current sample), we compared the Landmark Asymmetry Indices of those ADHD probands without the 10-repeat DAT1 allele and those in possession of one copy of this allele (designated as Low-Risk DAT1) ($n=21$) to those in possession of two copies of this allele (designated as High-Risk DAT1) ($n=22$). Preliminary analyses revealed that performance measures were normally distributed and therefore parametric tests were used in statistical comparisons.

Using a family-based design and a logistic regression adaptation of the transmission disequilibrium test (LR-TDT) (Waldman *et al*, 1999), we also used parental genotype information to examine whether the Landmark Asymmetry Index predicted biased transmission of high-risk (10-repeat), vs low-risk (other alleles), parental DAT1 alleles to ADHD probands. This method is robust against any population stratification effects.

Testing the association between attentional asymmetry and response to MPH. Firstly, Landmark Asymmetry Indices were compared as a function of Medication Response Group (None or Mediocre vs Very Good) using a univariate ANOVA. Secondly, Landmark Asymmetry Indices were compared as a function of the combination of Medication Response Group and DAT1 Genotype (Low-Risk DAT1 vs High-Risk DAT1).

RESULTS

Testing the Association between DAT1 Genotype and Attentional Asymmetry

When symptomatology was measured dimensionally, significant differences emerged between the Low- and High-Risk DAT1 ADHD groups in both DSM-IV Inattentive and Total age-related normative scores (t -scores), as measured by the CPRS-R:L (Table 1). Further, Landmark Asymmetry Indices correlated positively with continuous measures of DSM-IV Inattentiveness ($r(41)=0.35$, $p=0.02$) and DSM-IV Total symptom scores ($r(41)=0.40$, $p=0.01$) but not DSM-IV Hyperactivity/Impulsiveness ($r(41)=0.29$, $p=0.065$). The direction of these correlations suggests that greater symptom severity, particularly with respect to inattentive symptoms, is associated with left-sided inattention.

There was a significant effect of DAT1 Genotype Group on the Asymmetry Index ($F(1,41)=10.9$, $p=0.002$; $r^2=0.19$) that was driven by the right spatial bias/left-sided inattention of the High-Risk DAT1 ADHD group ($M=+0.09$, $SD=0.29$), in comparison to the left spatial bias/right-sided inattention of the Low-Risk DAT1 ADHD group ($M=-0.17$, $SD=0.23$) (Figure 2). Despite the small number of participants not in possession of a 10-repeat

Figure 2 Mean Landmark Asymmetry Indices as a function of ADHD DAT1 Genotype Group. The performance accuracy of the groups on the Landmark task was firstly compared using those trials on which the midpoint of the line was offset. Accuracy was high across both groups (mean accuracies >73%) and not significantly different between the ADHD DAT1 Genotype Groups ($F(1,41)=0.57$, $p=0.45$).

Table 2 Landmark Asymmetry Index (Mean (SD)) as a Function of DAT1 Alleles

	9/9 ($n=2$)	10/other allele ^a ($n=19$)	10/10 ($n=22$)
Landmark Asymmetry Index	-0.35 (0.07)	-0.15 (0.23)	+0.09 (0.29)

^aOne subject possessed an 11-repeat (520 bp) allele and one subject possessed a rare 400 bp allele.

DAT1 allele ($n=2$), regression revealed a parametric effect of increasing number of 10-repeat DAT1 alleles (0, 1, or 2) on Landmark Asymmetry Indices ($F(1,41)=12.23$, $p=0.001$; $r^2=0.21$) (see Table 2). This result suggests an additive effect of possession of the 10-repeat allele.

A logistic regression model was used to examine whether the Landmark Asymmetry Index predicted preferential transmission of high-risk (10-repeat), vs low-risk (other alleles), parental DAT1 alleles. The coding of transmitted alleles from heterozygous parents resulted in 40 informative transmissions. The Landmark Asymmetry Index significantly predicted biased transmission of high-risk, vs low-risk, parental DAT1 alleles ($\chi^2(df=1)=7.43$, $p=0.006$, 37% variance explained). This association was stronger than when using DSM-IV Inattentive ($\chi^2(df=1)=3.6$, $p=0.058$), Hyperactive/Impulsive ($\chi^2(df=1)=2.5$, $p=0.115$), or Total Symptoms ($\chi^2(df=1)=3.6$, $p=0.059$) as continuous predictor variables. Thus, while left spatial inattention is related to dimensional measures of ADHD symptomatology, the former is more strongly associated with DAT1 genotype than the latter. This result satisfies a key assumption of the endophenotype approach.

Testing the Association between Attentional Asymmetry and Response to MPH

Retrospective ratings of MPH response using the three-point scale (No Response vs Mediocre Response vs Very

Good Response) were available for 36 children and adolescents with ADHD. A total of 15 of these participants were rated as achieving No Response or a Mediocre Response, while 21 were rated as achieving a Very Good Response. There was a significant difference in the Landmark Asymmetry Index as a function of Medication Response Group ($F(1,34) = 5.22$, $p = 0.03$, $r^2 = 0.11$), which was driven by the right spatial bias/left-sided inattention of the Very Good Response group ($M = +0.05$, $SD = 0.29$), in comparison to the left spatial bias/right-sided inattention of the No/Mediocre Response group ($M = -0.15$, $SD = 0.19$).

Given the *a priori* prediction that those children achieving a very good response to MPH and possessing two 'high-risk' 10-repeat DAT1 alleles would show left-sided inattention (right bias) on the Landmark Task, we compared the performance of this group (High-Risk DAT1/Very Good Response) ($n = 12$) to the three other Genotype/Medication Response groupings (see Figure 3).

There was a significant effect of DAT1 Genotype/Medication Response Group on the Landmark Asymmetry Index ($F(3,32) = 4.12$, $p = 0.01$, $r^2 = 0.21$) (Figure 3). Figure 3 indicates that Asymmetry Indices became increasingly right biased as a function of DAT1 Genotype/Medication Response Group. As hypothesized, the High-Risk DAT1/Very Good Response group were the only group to display left-sided inattention ($M = +0.13$, $SD = 0.32$), with the Low-Risk DAT1/Mediocre Response Group showing a leftward bias ($M = -0.25$, $SD = 0.11$). Both the Low-Risk DAT1/Very Good Response ($M = -0.07$, $SD = 0.21$) and High-Risk DAT1/Mediocre Response ($M = -0.04$, $SD = 0.21$) groupings had small leftward biases. Pairwise comparisons with Bonferroni corrections revealed that the Asymmetry Indices of the High-Risk DAT1/Very Good Response Group and the Low-Risk DAT1/Mediocre Response Group were significantly different ($p < 0.01$). No other pairwise comparisons were significant.

Changes in symptom severity when on and off MPH, as rated retrospectively by parents using the CPRS-R:L, were available for 32 ADHD participants. The High-Risk DAT1

ADHD group achieved greater symptom reduction than the Low-Risk DAT1 ADHD group in terms of both DSM-IV Hyperactive/Impulsive ($F(1,30) = 5.289$, $p = 0.03$, $r^2 = 0.12$) and DSM-IV Total symptoms ($F(1,30) = 4.40$, $p = 0.04$, $r^2 = 0.13$) but not DSM-IV Inattentive symptoms ($F(1,30) = 1.99$, $p = 0.169$, $r^2 = 0.06$).

DISCUSSION

This study reports that left-sided inattention is related to clinical ADHD symptomatology, the 10-repeat DAT1 allele, and the therapeutic response to MPH. 10-repeat DAT1 homozygotes displayed left-sided inattention, whereas those possessing one or no copies of this allele did not. This result extends preliminary reports of an effect of DAT1 genotype on neuropsychological performance in ADHD (Loo *et al*, 2003; Bellgrove *et al*, in press). Importantly, using a within-family control design, we report that scores on the Landmark Asymmetry Index predict biased transmission of the 10-repeat DAT1 allele from parents to probands. The greater effect of the 10-repeat allele on attentional asymmetry observed in homozygotes, *vs* heterozygotes, suggests additive rather than dominance effects of the 10-repeat allele. As we hypothesized, left-sided inattention was associated with an enhanced therapeutic response to MPH, irrespective of DAT1 status, but was most pronounced in 10-repeat DAT1 homozygotes who achieved a very good response to MPH. Our data support the existence of a subgroup of ADHD that is linked to the 10-repeat allele of the DAT1 VNTR, and is defined (a) in symptom terms, by higher levels of inattentive and total symptomatology (but see Waldman *et al*, 1998); (b) in neuropsychological terms, by left-sided inattention, and perhaps also poor sustained attention (Loo *et al*, 2003); and (c) in pharmacogenetic terms, by an enhanced response to MPH (Kirley *et al*, 2003).

Based on the data reported herein and the previously reviewed literature, we propose the following genetic-neurophysiological mechanism as part of the pathophysiology of ADHD. Increased transporter density (or activity) is associated with the 10-repeat DAT1 allele (or another genetic variant in linkage disequilibrium with the DAT1 VNTR) in ADHD (see Cheon *et al*, 2005). Overactive transporters reduce extracellular dopamine, perhaps particularly within right hemisphere attentional networks. The resultant hypoactivation within right hemisphere systems weakens the attentional bias of the right hemisphere, unmasking the attentional bias of the left hemisphere thus driving spatial attention in a rightward direction (Kinsbourne, 1993). Variation in the DAT1 gene may therefore confer susceptibility to ADHD, in part because of varying effects on the development of brain mechanisms modulating (spatial) attention.

Treatment with MPH may inhibit the transporter, restoring both dopaminergic balance and the reciprocal balance between spatial attentional systems. Accordingly, treatment with MPH may be most effective in those individuals with the greatest transporter densities or activities. Insofar as the Landmark Asymmetry Index is able to act as a behavioral assay for this neural mechanism, then those ADHD individuals presenting with left-sided inattention may be more likely to achieve an enhanced

Figure 3 Mean Landmark Asymmetry Indices as a function of ADHD DAT1 Genotype Group and Medication Response Group. Significant differences existed between the High-Risk DAT1/Very Good Response and Low-Risk DAT1/Mediocre Response groups.

response to MPH. Our data also suggest that knowing the DAT1 genotype of an individual may strengthen this association.

The above model, however, assumes that the DAT1 VNTR modulates transporter densities asymmetrically in ADHD, thus giving rise to the left-sided inattention reported herein. While there is, as yet, little direct evidence for this assumption, a number of studies do suggest a cerebral asymmetry in transporter densities. For example, Laakso *et al* (2000) reported higher striatal dopamine transporter binding in healthy subjects within the right, relative to left, striatum. Cheon *et al* (2003) also reported that DAT binding ratios within the basal ganglia of children with ADHD, relative to controls, were elevated by 51% on the right and 40% on the left. Further, it has recently been demonstrated that the beneficial effect of MPH on attention and impulsivity in ADHD is related to a reduction in the availability of D2/D3 receptors. This reduction in receptor availability is indicative of a pharmacologically evoked increase in extracellular dopamine by blockade of the transporter that is maximal in the right striatum (Rosa-Neto *et al*, 2005). It should also be noted, however, that the results reported herein could also arise from an interaction between DAT1 genotype and its associated alteration in dopaminergic transmission, and structural and/or functional changes within right hemisphere attentional systems (eg Castellanos *et al*, 1994; Casey *et al*, 1997; Bush *et al*, 1999; Sowell *et al*, 2003) that are unrelated to DAT1 genotype.

There are several limitations of this study that require comment. First, the sample size is relatively small for a genetic association study. For example, only two probands possessed the 9/9 genotype. Nevertheless, our results are consistent across analyses focusing on possession and transmission of high-risk alleles. While the latter analysis (LR-TDT) sacrifices power by focusing on transmissions from heterozygous parents only, the results were highly significant and survive correction for multiple comparisons. DAT1 genotype accounted for 19% of the variance in the Landmark Asymmetry Index in the ANOVA-based analysis. This effect on spatial cognition is larger than is typically reported with functional variants, such as the COMT Val/Met polymorphism (Egan *et al*, 2001). Larger collaborative studies using a range of spatial attentional tasks are required to confirm and extend our results. Second, we assessed medication response using a simple nonvalidated three-point scale, and collected retrospective parental ratings. While these results should be viewed as preliminary until replicated within a prospective study, it should be noted that these ratings are unlikely to be biased with respect to DAT1 genotype. Third, we have proposed a biological hypothesis of the relation between left-sided inattention, DAT1 genotype, and MPH response in which dysfunction to attentional systems is centered on the right striatum. While this hypothesis is advanced based upon the known action of stimulants at transporters within the striatum, left-sided inattention can also arise from right parietal or prefrontal lesions (Robertson and Marshall, 1993). We cannot discount the possibility that our behavioral results reflect fronto-parietal dysfunction.

In summary, our results show that left-sided inattention in ADHD is related to both underlying DAT1 genotype and

the therapeutic response conferred by MPH. Our results are internally consistent in demonstrating an attentional endophenotype that relates to ADHD symptomatology, DAT1 genotype, and the therapeutic efficacy of MPH. The further study of spatial (in)attention in ADHD, in relation to DAT1 genotype, may provide a window into the neurobiology of ADHD.

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