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## Increased Rate of Twins among Affected Sibling Pairs with Autism

To the Editor:

There is consistent evidence from twin and family studies implicating genetic factors in the etiology of autism (MIM 209850), but no specific genes associated with autism have yet been identified. In a recent article in the Journal, Greenberg et al. (2001) reported a striking excess of twin pairs, both MZ and DZ, in the cohort of families with at least two siblings with autism or autismrelated conditions recruited by the Autism Resource Exchange (AGRE) of the Cure Autism Now (CAN) Foundation. The proportion of twins among autistic sib pairs (18%, 30/166) was significantly higher than the expected twinning rate per sib pair (2.4%). Greenberg et al. (2001) demonstrated that to ascribe this excess of twins with autism to a sampling bias would require very large ascertainment factors, which seem unlikely. These findings suggest that being a twin represents a risk factor for autism, and they have important implications for the etiology of autism. However, as Greenberg et al. (2001) pointed out, these results need to be replicated in other data sets. We report here a similar excess of twins in a sample of affected sib pairs recruited by the Paris Autism Research International Sibpair (PARIS) study.

Families with two or more children with autism or autism-related disorders (Asperger syndrome or pervasive developmental disorder not otherwise specified [PDD NOS]) were recruited by the PARIS study at specialized clinical centers in eight countries (Austria, Belgium, France, Israel, Italy, Norway, Sweden, and the United States). Patients were included after undergoing a complete clinical and neuropsychological assessment described elsewhere (Philippe et al. 1999); subjects demonstrated to suffer from organic conditions associated with autism, such as tuberous sclerosis, fragile X syndrome, or other established chromosomal disorders, were excluded from the study. We divided the affected sib pairs into two diagnostic categories: "narrow," when both affected sibs had autism, and "broad," when one or both of the affected sibs had either Asperger syndrome or PDD NOS. Patients in the narrow diagnostic category

fulfilled the DSM-IV and ICD-10 criteria for autistic disorder/childhood autism and met the Autism Diagnostic Interview–Revised algorithm (Lord et al. 1994). All the families were white except one of mixed ethnicity (white/Asian).

To make our results directly comparable to those obtained by Greenberg et al. (2001), in the current analysis we included only families having exactly two affected offspring; families with triplets or with an affected twin pair and one or more affected nontwin siblings were excluded. We also excluded families with mixed twin pairs (one affected twin, one unaffected co-twin, and a nontwin affected sib) and families with half siblings.

Table 1 shows the distribution of affected sib pairs divided according to diagnostic category, twin status, and sex. Table 2 shows the observed proportion of twins in our data set compared both with the population rates reported by Greenberg et al. (2001) and with the rate observed in the AGRE families. In agreement with the results of Greenberg et al. (2001), we observed a remarkably high proportion of MZ twin pairs among affected sib pairs. Of 79 affected sib pairs (narrow + broad diagnoses), 11 were twin pairs (2 DZ and 9 MZ). This represents a 14-fold increase for MZ twins, compared with the population frequency, and is statistically significant ( $P < 10^{-5}$ ). In contrast, we did not observe a significant increase in the proportion of DZ twins.

We did not include the following families in the calculations: (1) one family with affected triplets (two MZ and one DZ); (2) one family with two affected MZ twins plus an affected nontwin; (3) one family with two affected twins (zygosity unknown) plus two affected nontwins; (4) one family with one affected MZ twin, one co-twin deceased during the first year of life, and a non-

Table 1

Distribution of Affected Sib Pairs

	No. in Diagnostic Category <sup>a</sup>					
GROUP	Narrow	Broad	Total			
Singletons	59 (34, 23, 2)	9 (5, 4, 0)	68 (39, 27, 2)			
DZ twins	2 (2, 0, 0)	0	2 (2, 0, 0)			
MZ twins	9 (7, —, 2)	0	9 (7, —, 2)			
Total	70 (43, 23, 4)	9 (5, 4, 0)	79 (48, 27, 4)			

<sup>&</sup>lt;sup>q</sup> Numbers in parentheses indicate breakdown into male-male, mixed sex, and female-female pairs, respectively.

1382 Letter to the Editor

Observed Proportion of Twins Compared with Population Rates and with Results of the Study by Greenberg et al. (2001)									
	Greenberg et al. (2001) <sup>a</sup>		Narrow Diagnosis		Narrow + Broad Diagnoses				
TWIN POPULATION GROUP RATE <sup>b</sup>	Rate Observed	P°	Rate Observed	P°.	Rate Observed	$P^{c}$			
.016	.072 (12/166)	<.00005	.029 (2/70)	N.S.	.025 (2/79)	N.S. <.000001			
	POPULATION RATE <sup>b</sup>	GREENBERG ET A	$\begin{array}{c c} & & & & & & & & \\ & & & & & & & \\ POPULATION & & & & & \\ RATE^b & & Observed & & P^c \\ & .016 & .072 \ (12/166) & < .00005 \end{array}$	GREENBERG ET AL. (2001)a   NARROW DI	$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$				

.157 (11/70)

Table 2

<.000001

.024

All

.181 (30/166)

twin affected sib; and (5) three sets of discordant DZ twins (one affected and one unaffected). Together with the other twin pairs, these families further reinforce the hypothesis that twinning per se is a significant risk factor for autism.

The high proportion of twins among affected sib pairs with autism observed by Greenberg et al. (2001) and replicated in our data set strongly suggests the involvement of biological factors rather than an ascertainment bias. Moreover, different ascertainment methods were used in our study and that of Greenberg et al. (2001): the AGRE families were recruited exclusively via mailings and presentations to autism support groups, whereas the majority of our families were collected by clinicians members of the PARIS study among their clinic cases.

The PARIS study collects affected sib pairs for linkage studies but also parent-offspring trios for association studies. We do not preferentially collect MZ twins, but neither do we turn them away, because these families can be analyzed in association studies. In our data set, only MZ twins were overrepresented among affected sib pairs. Although Greenberg et al. (2001) observed an excess of both DZ and MZ twins among their families, the deviation from population rates was more important among MZ twins (12-fold) than among DZ twins (4fold). As mentioned by Greenberg et al. (2001), the excess of MZ twins compared with DZ twins in autism suggests that the estimates of heritability based on concordance for autism in MZ pairs versus DZ pairs may be overestimated. These intriguing findings also emphasize the need to explore the participation of nongenetic as well as genetic factors in the etiology of autism.

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## **Electronic-Database Information**

<.000001

The accession number and URL for data in this letter are as follows:

Online Mendelian Inheritance in Man (OMIM), http://www .ncbi.nlm.nih.gov/Omim/ (for autism [MIM 209850])

## References

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<sup>&</sup>lt;sup>a</sup> Includes narrow + broad diagnoses.

<sup>&</sup>lt;sup>b</sup> From Greenberg et al. (2001).

<sup>&</sup>lt;sup>c</sup> Two-sided, exact binomial calculations. N.S. = not significant.

Letter to the Editor 1383