

A controlled family study of children with DSM-IV bipolar-I disorder and psychiatric co-morbidity

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Background. To estimate the spectrum of familial risk for psychopathology in first-degree relatives of children with unabridged DSM-IV bipolar-I disorder (BP-I).

Method. We conducted a blinded, controlled family study using structured diagnostic interviews of 157 children with BP-I probands ($n=487$ first-degree relatives), 162 attention deficit hyperactivity disorder (ADHD) (without BP-I) probands ($n=511$ first-degree relatives), and 136 healthy control (without ADHD or BP-I) probands ($n=411$ first-degree relatives).

Results. The morbid risk (MR) of BP-I disorder in relatives of BP-I probands (MR=0.18) was increased 4-fold [95% confidence interval (CI) 2.3–6.9, $p<0.001$] over the risk to relatives of control probands (MR=0.05) and 3.5-fold (95% CI 2.1–5.8, $p<0.001$) over the risk to relatives of ADHD probands (MR=0.06). In addition, relatives of children with BP-I disorder had high rates of psychosis, major depression, multiple anxiety disorders, substance use disorders, ADHD and antisocial disorders compared with relatives of control probands. Only the effect for antisocial disorders lost significance after accounted for by the corresponding diagnosis in the proband. Familial rates of ADHD did not differ between ADHD and BP-I probands.

Conclusions. Our results document an increased familial risk for BP-I disorder in relatives of pediatric probands with DSM-IV BP-I. Relatives of probands with BP-I were also at increased risk for other psychiatric disorders frequently associated with pediatric BP-I. These results support the validity of the diagnosis of BP-I in children as defined by DSM-IV. More work is needed to better understand the nature of the association between these disorders in probands and relatives.

Received 29 January 2009; Revised 3 September 2009; Accepted 6 September 2009; First published online 6 November 2009

Key words: Bipolar, children, family study.

Introduction

A converging body of evidence indicates that a sizeable minority of children and adolescents in clinic and research settings satisfy DSM-IV diagnostic criteria for bipolar disorder (Mick *et al.* 2003; Perlis *et al.* 2004). This literature also documents that pediatric bipolar disorder is extremely morbid and commonly associated with significant functional impairment in multiple domains including increased risks for psychiatric hospitalization, antisocial behaviors, addictions and suicidal ideation (Wozniak *et al.* 1995a; Geller *et al.* 2000a; Biederman *et al.* 2004; Birmaher *et al.* 2006). In parallel to pediatric studies, an emerging literature in adults documents that as many as 65% of adults with

bipolar disorder have an onset of their disorder in childhood and adolescence, indicating that onset in childhood and adolescence is a common feature of this disorder (Perlis *et al.* 2004). Despite these compelling findings, questions remain as to the validity of pediatric bipolar disorder.

A cornerstone of establishing the validity of a psychiatric disorder is demonstrating that relatives of diagnosed individuals (i.e. the proband) are at an increased risk for the same disorder (Robins & Guze, 1970). That bipolar disorder in adults is highly familial has been known since the middle of the twentieth century (Tsuang & Faraone, 1990; Faraone *et al.* 2003). In a recent review, Craddock & Forty (2006) estimated the risk for bipolar disorder in the siblings of adult probands to be 5–10% and that the heritability of the disorder ranges from 0.80 to 0.90. In contrast to a rich literature on family studies of adult bipolar disorder, a much more limited literature exists on the familiarity

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of pediatric bipolar disorder. In an uncontrolled study, Dwyer & Delong (1985) showed an excess of bipolar disorder in the relatives of 20 out-patient children with DSM-III diagnosed bipolar disorder. The excess risk for bipolar disorder in first-degree relatives was replicated in subsequent family studies of child probands with DSM-III (Strober *et al.* 1988; Kutcher & Marton, 1991; Neuman *et al.* 1997), DSM-III-R (Wozniak *et al.* 1995*b*) and DSM-IV bipolar disorder (Findling *et al.* 2001; Geller *et al.* 2006; Brotman *et al.* 2007; Wilens *et al.* 2007), which reported ranges of bipolar disorder in relatives of pediatric bipolar disorder probands ranging from 12 to 35% with the risk of unipolar depression ranging from 15 to 42%.

However, despite their clear contributions, the existing family studies of pediatric bipolar disorder suffer from several methodological limitations. Three studies relied on family history methods rather than directly interviewing relatives and half of the available studies examined only parents or adult relatives (Strober *et al.* 1988; Kutcher & Marton, 1991; Neuman *et al.* 1997; Findling *et al.* 2001; Brotman *et al.* 2007; Wilens *et al.* 2007). Of the four studies utilizing DSM-IV criteria, two (Geller *et al.* 2006; Brotman *et al.* 2007) restricted recruitment of probands to children meeting a 'narrow' phenotype (Leibenluft *et al.* 2003) excluding other children who may have otherwise fully met DSM-IV bipolar-I (BP-I) disorder.

Furthermore, despite the fact that high rates of psychiatric co-morbidity have been consistently reported in youth with bipolar disorder, the extant literature on family studies of pediatric bipolar disorder probands has seldom systematically assessed other psychiatric disorders beyond mood disorders and those studies that did assess other psychiatric disorders did not account for a potential impact of psychiatric co-morbidity in probands on the risk of psychiatric morbidity in relatives.

The main aim of the present study was to re-evaluate the familiarity of pediatric BP-I disorder attending to the limitations of the extant literature using a large family study sample. To this end, we conducted a familial risk analysis comparing structured diagnostic interview derived data from all first-degree relatives of pediatric probands with DSM-IV BP-I disorder attending to psychiatric co-morbidity in probands and relatives. Comparisons were made with findings in first-degree relatives of probands with attention deficit hyperactivity disorder (ADHD) and control probands without BP-I or ADHD. We hypothesized that first-degree relatives of probands with pediatric BP-I would be at increased risk for BP-I compared with relatives of ADHD probands and non-bipolar, non-ADHD control probands. Additionally, based upon patterns of psychiatric co-morbidity in

children with BP-I (Wozniak *et al.* 1995*b*; Geller *et al.* 2000*b*; Findling *et al.* 2001; Mick *et al.* 2003; Brotman *et al.* 2007), we also hypothesized that relatives of pediatric BP-I probands would be at increased risk for disruptive behavior disorders, anxiety disorders, addictive disorders and psychosis. To the best of our knowledge this study represents one of the largest and most comprehensive family studies of pediatric bipolar disorder.

Method

Subjects

Families were recruited and assessed at the Clinical and Research Program in Pediatric Psychopharmacology and Adult ADHD at Massachusetts General Hospital. Probands were recruited for studies of bipolar probands 6–17 years of age of both genders (Wozniak, 2005) and ADHD or non-ADHD control probands 6–17 years of age of both genders (Biederman *et al.* 1992, 1999, 2006*a, b*). All studies were sampled from the same source population and used the same assessment methodology regardless of the disorder used to classify subjects as cases. All study procedures were reviewed and approved by the subcommittee for human subjects of our institution. All subjects' parents or guardians signed written informed consent forms and children older than 7 years of age signed written assent forms.

We recruited 157 BP-I probands and their 487 first-degree relatives for the family study of pediatric bipolar disorder. From 522 families participating in our case-control ADHD family studies we randomly selected 162 non-bipolar ADHD (511 first-degree relatives) and 136 non-bipolar non-ADHD control probands (411 first-degree relatives) so that the age and gender distribution was similar to that of the BP-I probands. ADHD probands with co-morbid bipolar disorder were not included in the present analyses.

Ascertainment method

Potential BP-I probands were ascertained from our clinical service, referrals from local clinicians or self-referral in response to advertisements. To avoid biasing our sample toward familial cases of bipolar disorder, all probands were ascertained blind to the diagnostic status of their relatives. Subjects were administered a phone screen, reviewing symptoms of DSM-IV BP-I and, if criteria were met, were scheduled for a face-to-face structured diagnostic interview (described below). In addition to the structured diagnostic interview it is the routine for the PI (J.W.) to perform a clinical interview that includes both the

proband and his or her parents in order to confirm the diagnosis of bipolar disorder using the Schedule for Affective Disorders and Schizophrenia for School-Age Children (KSADS) mania module and we make every effort to ensure that this interview occurs with every proband. We have published data on the convergence of these clinical interviews with our structured interview diagnosis on the first 69 cases. We report 97% agreement between the structured interview and clinical diagnosis in this analysis of 69 children (Wozniak *et al.* 2003).

As previously reported (Biederman *et al.* 1992, 1999; Wozniak *et al.* 2005) ADHD cases were identified from either a major academic medical center, where we selected ADHD subjects from referrals to a pediatric psychopharmacology program, or from a major Health Maintenance Organization, in which ADHD subjects were selected from pediatric clinic out-patients. Controls were ascertained from out-patients referred for routine physical examinations to pediatric medical clinics at each setting, identified from their computerized records as not having ADHD. Screening procedures were similar to those described for the recruitment of the bipolar probands with the exception that we queried about ADHD (and not bipolar disorder) in the initial telephone screening and each proband was not assessed clinically.

Diagnostic procedures

Psychiatric assessments of subjects younger than 18 years were made with the KSADS-E (epidemiologic version; Orvaschel, 1994) and assessments of adult family members were made with the Structured Clinical Interview for DSM-IV (SCID; First *et al.* 1997) supplemented with modules from the KSADS-E to cover childhood disorders. Diagnoses were based on independent interviews with mothers and direct interviews with children older than 12 years of age. Data were combined such that endorsement of a diagnosis by either reporter resulted in a positive diagnosis.

Interviews with both the KSADS and the SCID were conducted by extensively trained and supervised psychometricians with undergraduate degrees in psychology. This training involved several weeks of classroom instruction of interview mechanics, diagnostic criteria and coding algorithms. They also observed interviews by experienced raters and clinicians and were observed while conducting interviews during the final training period. In addition, all diagnoses were reviewed by a sign-off committee of experienced board-certified child and adolescent psychiatrists or clinical psychologists. The committee members were blind to the subjects' ascertainment group, ascertainment site and data collected from other family

members. We computed κ coefficients of agreement by having experienced clinicians diagnose subjects from audiotaped interviews made by the assessment staff. Based on 500 interviews, the median κ coefficient between raters and clinicians was 0.99 and for individual diagnoses was ADHD (0.88), conduct disorder (CD) (1.0), major depression (1.0), mania (0.95), separation anxiety (1.0), agoraphobia (1.0), panic (0.95), substance use disorder (1.0) and tics/Tourette's (0.89). The median agreement between individual clinicians and the clinical review committee chaired by the PI was 0.87 and for individual diagnoses was ADHD (1.0), CD (1.0), major depression (1.0), bipolar (0.78), separation anxiety (0.89), agoraphobia (0.80), panic (0.77), substance use disorder (1.0) and tics/Tourette's (0.68).

Children were diagnosed with BP-I disorder according to DSM-IV criteria. The DSM-IV requires subjects to meet criterion A for a distinct period of extreme and persistently elevated, expansive or irritable mood lasting at least 1 week, plus criterion B, manifested by three (four if the mood is irritable only) of seven symptoms during the period of mood disturbance. To ensure that the B criterion symptoms were concurrent with A criterion mood disturbance, subjects were directed to focus on the worst or most impairing episode of mood disturbance while being assessed for the presence of the confirmatory B criterion symptoms. That is, the subject was asked to consider the time during which the screen was at its worst for the purposes of determining whether the remaining symptoms were also evident at the same time as the screening item. Also recorded was the onset of first episode, the number of episodes, offset of last episode and total duration of illness. Any subject meeting criteria for BP-II or BP-NOS was not included in this study. To gauge a distinct episode our interviewers asked for 'a distinct period (of at least 1 week) of extreme and persistently elevated, expansive or irritable mood' and further required that the irritability endorsed in this module is 'super' and 'extreme'.

Statistical analysis

We analyzed censored time-to-failure data (i.e. onset of disorder if the disorder is present or age at interview if the disorder was not present) with survival analysis methods to weight the contribution of each family member by their age at assessment. This is necessary because we assessed both child and adult relatives and the simple prevalence of disorder in relatives may underestimate the true risk since children have not yet progressed through the entire window of risk. Thus, we report estimates of morbid risk (MR) calculated from Kaplan–Meier cumulative failure function.

Table 1. Clinical and demographic characteristics

	BP-I families	ADHD families	Control families	Statistic
Total	<i>n</i> = 644	<i>n</i> = 673	<i>n</i> = 547	
SES	1.8 ± 0.9 ^a	1.8 ± 0.9 ^a	1.6 ± 0.8	<i>F</i> (2, 451) = 5.3, <i>p</i> = 0.005
Race/Ethnicity				
Caucasian	594 (92) ^{a,b}	667 (99)	536 (98)	$\chi^2(6) = 56.7$, <i>p</i> < 0.001
African-American	32 (5)	6 (1)	7 (1)	
More than 1	12 (2)	0 (0)	0 (0)	
Unknown	6 (1)	0 (0)	4 (1)	
Probands	<i>n</i> = 157	<i>n</i> = 162	<i>n</i> = 136	
Age (years)	10.5 ± 3.2	10.6 ± 3.0	10.7 ± 3.0	<i>F</i> (2, 452) = 0.2, <i>p</i> = 0.8
Gender				$\chi^2(2) = 2.1$, <i>p</i> = 0.3
Female	125	121	99	
Male	80	75	73	
Past GAF	40.6 ± 5.9 ^{a,b}	50.7 ± 7.3 ^a	70.5 ± 8.5	<i>F</i> (2, 452) = 630.7, <i>p</i> < 0.001
Current GAF	46.2 ± 6.3 ^{a,b}	57.4 ± 8.2 ^a	73.3 ± 7.3	<i>F</i> (2, 452) = 505.0, <i>p</i> < 0.001
Parents	<i>n</i> = 301	<i>n</i> = 323	<i>n</i> = 269	
Age (years)	42.3 ± 6.6	41.3 ± 6.4	41.6 ± 5.8	<i>F</i> (2, 884) = 2.0, <i>p</i> = 0.1
Gender				$\chi^2(2) = 0.3$, <i>p</i> = 0.9
Female	144	161	133	
Male	48	50	49	
Past GAF	52.2 ± 9.7 ^{a,b}	56.9 ± 12.6 ^a	63.5 ± 12.4	<i>F</i> (2, 875) = 65.9, <i>p</i> < 0.001
Current GAF	63.4 ± 7.8 ^{a,b}	68.5 ± 9.5 ^a	72.9 ± 7.9	<i>F</i> (2, 834) = 85.5, <i>p</i> < 0.001
Siblings	<i>n</i> = 186	<i>n</i> = 188	<i>n</i> = 142	
Age (years)	11.6 ± 5.5 ^b	13.7 ± 5.8	12.9 ± 5.1	<i>F</i> (2, 511) = 7.2, <i>p</i> = 0.001
Gender				$\chi^2(2) = 0.8$, <i>p</i> = 0.7
Female	98	103	74	
Male	51	55	52	
Past GAF	57.7 ± 9.4 ^{a,b}	61.8 ± 12.0 ^a	65.9 ± 10.7	<i>F</i> (2, 504) = 23.3, <i>p</i> < 0.001
Current GAF	62.6 ± 7.7 ^{a,b}	67.9 ± 10.8 ^a	71.1 ± 8.4	<i>F</i> (2, 503) = 35.9, <i>p</i> < 0.001

BP-I, Bipolar-I, ADHD, attention deficit hyperactivity disorder; SES, socio-economic status; GAF, global assessment of function.

^a *p* < 0.05 *v.* control; ^b *p* < 0.05 *v.* ADHD.

Hazards ratios (HRs) and their 95% confidence intervals were estimated from Cox proportional hazard models to test for differences between groups of relatives. To estimate the independent risk of additional psychiatric morbidity in relatives, congruent proband co-morbidity was included in family risk models. For example, in estimating the relative increase in familial risk of anxiety, we modeled the risk of anxiety in relatives as a function of group status (BP-I, ADHD and control), co-morbid anxiety in proband and any other confounders of interest.

To account for non-independence within families, we adjusted variance estimates of these Cox models with Huber's (1967) formula as implemented in Stata (Rogers, 1993) to produce *p* values that are robust to distributional assumptions. Other demographic data (e.g. age, gender, etc) were analyzed with one-way

analysis of variance or Pearson's χ^2 test. All statistical tests were two-tailed and any *p* values < 0.05 were considered statistically significant.

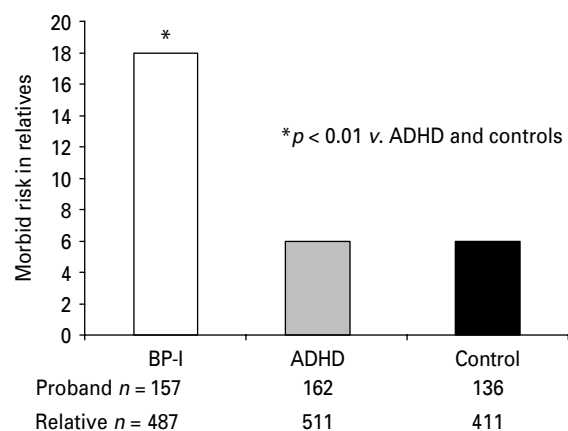
Results

Demographic and clinical characteristics of the sample are presented in Table 1. In total, 80% of the BP-I probands were male. There were small but statistically significant differences in the ethnic and socio-economic backgrounds of the families. The control had higher socio-economic status (SES) and the BP-I families had more ethnic diversity. Accordingly, all subsequent tests were adjusted for SES and race. There were no differences in the age or gender of the BP-I, ADHD and control probands (by design, see Ascertainment method). The BP-I probands were more

Table 2. Psychiatric co-morbidity in proband children

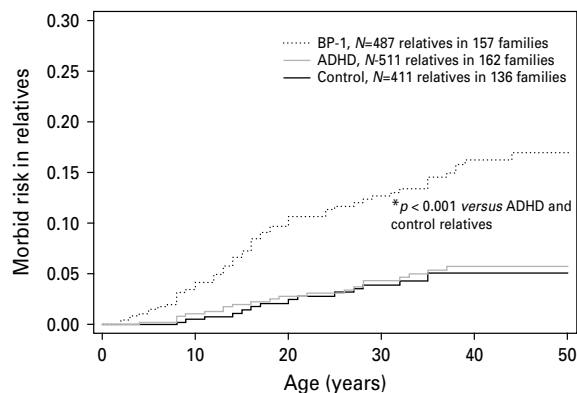
	BP-I (<i>n</i> = 157) <i>n</i> (%)	ADHD (<i>n</i> = 162) <i>n</i> (%)	Control (<i>n</i> = 136) <i>n</i> (%)	Statistic
Psychosis	51 (33)	– ^c	– ^c	
Major depression	131 (83) ^{a,b}	61 (37) ^a	10 (7)	$\chi^2(2) = 175.5, p < 0.001$
Multiple (2) anxiety disorders	100 (64) ^{a,b}	43 (27) ^a	6 (4)	$\chi^2(2) = 120.7, p < 0.001$
ADHD	133 (85)	– ^c	– ^c	
Oppositional defiant disorder	141 (90) ^{a,b}	87 (54) ^a	8 (6)	$\chi^2(2) = 205.9, p < 0.001$
Conduct disorder	80 (51) ^{a,b}	24 (15) ^a	2 (2)	$\chi^2(2) = 109.9, p < 0.001$
Substance (alcohol or drug) use disorder (abuse or dependence)	18 (12) ^{a,b}	5 (3)	1 (1)	$\chi^2(2) = 19.2, p < 0.001$

BP-I, Bipolar-I.

^a $p < 0.05$ *v.* controls; ^b $p < 0.05$ *v.* attention deficit hyperactivity disorder (ADHD).^c Potential probands with psychosis were excluded during ascertainment of ADHD and control families.**Fig. 1.** Familial risk of bipolar-I (BP-I) disorder in first-degree relatives. ADHD, attention deficit hyperactivity disorder.

impaired than both ADHD and control children according to past and current global assessment of functioning (GAF) score (Table 1). There were no meaningful demographic characteristics differences between the parents or siblings of the BP-I, ADHD and control probands, but relatives of BP-I probands were more impaired according to both lifetime and current GAF scores than the relatives of both the ADHD and control probands.

The clinical presentation of BP-I disorder in probands was characterized by early onset (5.8 ± 3.4 years), rapid cycling (22.4 ± 61.6 episodes) and a chronic course (3.6 ± 3.3 years in duration). As shown in Table 2, BP-I in probands was predominantly mixed with co-occurring depression ($n = 131$, 83%) and probands with BP-I were at increased risk of multiple (≥ 2) anxiety disorders, disruptive behavior disorders and substance use disorder relative to both the ADHD and control probands. Although statistical comparisons could not be made between these groups for

**Fig. 2.** Bipolar-I (BP-I) disorder in first-degree relatives.

ADHD or psychosis due to the inclusion/exclusion criteria of the ADHD family studies, both of these disorders were over-represented in BP-I probands.

The age-dependent cumulative MR of BP-I disorder in relatives is illustrated in Figs 1 and 2. The risk of BP-I disorder in relatives of BP-I probands was statistically significantly higher compared with the relatives of both the ADHD (HR = 3.1 (1.8–5.5); $p < 0.0001$) and the healthy control probands (HR = 3.3; (1.9–5.5); $p < 0.0001$). In contrast, the relatives of ADHD probands were not at increased risk of BP-I compared with relatives of control probands (HR = 1.0; (0.5–1.9); $p = 0.9$). In this context, the HR indexes the relative risk for a disorder in the relative given the proband diagnoses. For example, the HR for BP-I in relatives of BP-I *versus* relatives of ADHD probands was 3.6, which means that there was an age-corrected 3.6-fold increase of BP-I among the relatives of BP-I probands compared with relatives of ADHD probands. Controlling for psychiatric co-morbidity in probands

Table 3. Psychiatric morbidity in first-degree relatives of pediatric bipolar disorder, attention deficit hyperactivity disorder (ADHD) and control probands

	BP-I (<i>n</i> = 487) MR (95% CI)	ADHD (<i>n</i> = 511) MR (95% CI)	Control (<i>n</i> = 411) MR (95% CI)
Psychosis	0.07 (0.05–0.11) ^{b,c}	0.01 (0.005–0.03)	–
Major depression	0.49 (0.44–0.55) ^{a,b,c}	0.39 (0.34–0.45) ^{a,c}	0.22 (0.18–0.27)
Multiple (2) anxiety disorders	0.38 (0.32–0.44) ^{a,b,c,d}	0.19 (0.15–0.23) ^a	0.13 (0.10–0.18)
Substance (alcohol or drug) use disorder (abuse or dependence)	0.57 (0.52–0.63) ^{a,b,c,d}	0.40 (0.35–0.46)	0.28 (0.23–0.33)
ADHD	0.23 (0.19–0.27) ^{a,c}	0.20 (0.17–0.24) ^{a,c}	0.07 (0.04–0.09)
Oppositional defiant disorder	0.21 (0.17–0.25) ^{a,b}	0.12 (0.09–0.16) ^a	0.07 (0.05–0.10)
Conduct disorder/Antisocial personality disorder	0.17 (0.13–0.21) ^a	0.16 (0.12–0.20) ^a	0.06 (0.04–0.10)

MR, Morbid risk; CI, confidence interval; HR, hazard ratio.

All results corrected for family socio-economic status and race.

^a $p < 0.05$ *v.* controls relatives;

^b $p < 0.05$ *v.* ADHD relatives;

^c $p < 0.05$ *v.* controls relatives after correcting for concordant psychiatric co-morbidity in the proband (e.g. the analysis of ADHD in the relatives was corrected for the presence of ADHD in the proband);

^d $p < 0.05$ *v.* ADHD relatives after correcting for concordant psychiatric co-morbidity in the proband.

[oppositional defiant disorder (ODD), CD, major depression, multiple anxiety disorders and substance use disorder] did not impact the statistical significance or magnitude of the HRs comparing the relatives of BP-I probands with relatives of ADHD (corrected HR = 3.8 (1.9–7.6, $p < 0.0001$) or control (corrected HR = 4.0 (1.6–10.1, $p < 0.0001$) probands.

The MR of additional psychiatric disorders in the first-degree relatives of BP-I, ADHD and control probands are presented in Table 3. Relatives of BP-I probands were at increased risk of psychosis, major depression, multiple anxiety disorders, substance use disorders, ADHD, ODD and antisocial CD or antisocial personality disorder (ASPD) compared with relatives of control probands. In addition, in comparison with relatives of ADHD probands, relatives of BP-I probands were also at increased risk of major depression, multiple anxiety disorders, substance use disorders and ODD (Table 3). However, in models adjusting for the same psychiatric co-morbidity in probands, the relatives of BP-I probands were no longer at increased risk for ODD nor for CD/ASPD compared with relatives of control probands, nor for ODD and CD/ASPD compared with relatives of ADHD probands (all p 's > 0.05). Thus, BP-I in probands was independently associated with major depression, multiple anxiety disorders substance use disorders and ADHD in comparison with controls. Relatives of BP-I probands were at statistically significant increased risk of psychosis, multiple anxiety disorders and substance use disorder compared with

relatives of ADHD probands independently of the psychiatric co-morbidity with these disorders in probands.

To determine if our findings of familial transmission were moderated by age, we augmented our statistical models by adding the interaction of age (≤ 12 years *v.* > 12 years) by proband group. We found no statistically significant interaction, which indicates that the magnitude of familial transmission was not moderated by age group.

Discussion

Particular strengths of this study include its large sample size, the comprehensive scope of psychopathology examined in both probands and relatives and the use of both psychopathological (ADHD) and healthy control comparison groups. By assessing a wide range of psychiatric conditions in these data, we could adjust for psychiatric co-morbidity in probands when estimating the familiarity of BP-I in their first-degree relatives and estimate the familial risk of additional psychiatric disorders in relatives of BP-I child probands, while also adjusting for psychiatric co-morbidity in the probands. The diagnosis of pediatric bipolar disorder continues to confound clinicians and researchers, with questions remaining as to its validity. Following the logic of Robins & Guze (1970), family studies provide data external to the clinical picture, which can support the validity of a diagnosis. The present study is unique in several ways: (1) it is

the largest family study of this disorder; (2) it includes a psychopathological control group; (3) it ascertains subjects based on unmodified DSM criteria; (4) it is the first to focus on the familiarity of additional psychopathological conditions in relatives.

In our sample, 80% of the BP-I probands are male. This is consistent with previous reports of male preponderance by Geller *et al.* (2008) 67%, Findling *et al.* (2001) 71.1% and Luckenbaugh *et al.* (2009) 70%. Only Birmaher *et al.* (2009) found a nearly equal gender representation for BP-I subjects of 53.5%.

The significantly elevated MR of BP-I in relatives was not appreciably changed after controlling for psychiatric co-morbidity in probands. This was so despite the high rates of co-morbid ADHD, ODD, major depression and anxiety disorders in children with BP-I disorder, as has been previously documented by several research groups (Wozniak & Biederman, 1995; Wozniak *et al.* 1995a; Geller *et al.* 2000b; Findling *et al.* 2001; Mick *et al.* 2003; Brotman *et al.* 2007). Moreover, the similar magnitude and statistical significance of the corrected and uncorrected HRs suggests that co-morbidity in probands had little impact on the familiarity of BP-I disorder. This finding cannot be explained by any theory that posits BP-I disorder in children to be an epiphenomenon of another disorder.

Although relatives of BP-I probands were at increased risk for major depression compared with relatives of controls, the risk for major depression was not distinguishable between relatives of BP-I and relatives of ADHD probands. These results are consistent with previous studies (Kutcher & Marton, 1991; Wozniak *et al.* 1995a, b; Geller *et al.* 2006; Brotman *et al.* 2007). Based on our previous work examining the nature of the association between ADHD and major depression, similarities in the risk for major depression between ADHD and BP-I families could be explained by research suggesting that ADHD and major depression may share familial risk factors (Faraone & Biederman, 1997; Biederman *et al.* 1998, 2008).

Our findings that the elevated rates of antisocial disorders in relatives of BP-I probands was accounted for by these disorders in the proband are consistent with findings reported by Wozniak *et al.* (2001). These investigators also found that the relatives of BP-I in probands were not at increased risk for antisocial disorders after accounting for this co-morbidity in probands.

The association between BP-I and ADHD in families has been the subject of prior investigation. Rende *et al.* (2007) reported that 33% of children with BP-I disorder had a family history of ADHD and, although the MR of ADHD was not specifically estimated, Geller *et al.* (2006) found that relatives with ADHD, of child bipolar probands, were at increased risk for bipolar

disorder. The familial relationship between ADHD and BP-I observed in these studies of BP-I probands (Geller *et al.* 2006; Rende *et al.* 2007) is consistent with our previous family studies of ADHD and BP-I (Wozniak *et al.* 1995b; Faraone *et al.* 1997, 2001). That these disorders may also share familial risk factors could explain the association of the dopamine transporter gene with both bipolar disorder (Greenwood *et al.* 2006; Mick *et al.* 2007) and ADHD (Brookes *et al.* 2006a, b; Asherson *et al.* 2007). More work is needed with larger samples of non-ADHD BP-I probands and their relatives to fully parse the familiarity of ADHD and pediatric BP-I.

Our finding of increased familial risks for substance use disorders and anxiety disorders in relatives of pediatric BP-I probands is consistent with the literature (Dwyer & Delong, 1985; Strober *et al.* 1988; Brotman *et al.* 2007). Wozniak *et al.* (2002) also found an excess of anxiety disorders in relatives of pediatric bipolar probands but only among those of probands who also suffered from anxiety disorders. Similarly, prior work has shown an excess of substance use disorders in child BP-I probands and their relatives (Biederman *et al.* 2000a, b). The consistency of the familial risk for psychopathology in the current study with that of the extant literature is particularly noteworthy considering the variability in study methodology (i.e. ascertainment criteria, sample size, study design) and suggests that the familiarity of pediatric BP-I disorder may be as highly reproducible as it is for adult BP-I disorder (Tsuang & Faraone, 1990).

Our findings should be considered in the context of methodological limitations. Despite the large sample size, full stratification by psychiatric co-morbidity would have resulted in cells with small sample size (e.g. 24 non-ADHD BP-I probands). For our structured interviews, both the KSADS and the SCID, we used extensively trained interviewers with undergraduate degrees in psychology, rather than clinician raters. Although we did not administer structured diagnostic interviews directly to children younger than 12 years of age, a clinical diagnosis of BP-I in probands was corroborated by clinical assessment by an expert clinician prior to study inclusion (Wozniak *et al.* 2003). Also, we did not concurrently enroll comparison families but relied instead upon existing samples of ADHD and non-ADHD families. However, because all subjects were recruited from the same catchment area using the same ascertainment schema and research assessments, it is unlikely that the sample definition accounts for the findings presented here. Finally, because this sample was clinically referred and primarily Caucasian, these results may not generalize to non-referred children or to families of other ethnicities.

With these considerations in mind, we report that relatives of pediatric probands with DSM-IV defined BP-I disorder were at significantly increased risk for BP-I disorder compared with relatives of both ADHD and control probands. In addition, we found that pediatric BP-I disorder was associated with an increased familial risk of syndrome-congruent psychiatric co-morbidity such as of major depression, substance use disorder, anxiety disorders and psychosis. High rates of antisocial disorders were noted among the relatives of BP-I probands who also suffered from these co-morbidities. These results are consistent with the literature documenting the familiarity of pediatric bipolar disorder and suggest that DSM-IV BP-I disorder diagnostic criteria applied to children is a valid clinical entity worthy of further clinical and scientific attention. While the familial component of the Robins & Guze (1970) criteria for a valid psychiatric condition may have been met, prospective follow-up studies, genetic association studies and neuroimaging studies are needed to further characterize the prognostic course and neurobiological underpinnings and causes of DSM-IV defined pediatric bipolar disorder.

Acknowledgements

This work was supported by National Institutes of Health (NIH) grants K08MH001503 and R01MH066237 to Dr Wozniak, R01MH050657 and R01HD036317 to Dr Biederman and K01MH065523 to Dr Mick. This work was also supported by a grant from the Susan G. Berk Endowed Fund for Juvenile Bipolar Disorder. The Heinz C. Prechter Bipolar Research Fund and the support of members of the MGH Pediatric Psychopharmacology Council. All sponsors of this research supported only the collection of the data and played no other role in the design, analysis or interpretation of these findings. Dr Eric Mick had full access to all of the data in the study and takes responsibility for the integrity of the data and the accuracy of the data analysis.

Declaration of Interest

Dr Eric Mick receives research support from the following sources and is on an advisory board for the following sources: McNeil Pediatrics, Ortho-McNeil Janssen Scientific Affairs, Pfizer, Shire Pharmaceuticals and the National Institute of Mental Health (NIMH) and has had an advisory or consulting relationship with Pfizer and Shire Pharmaceutical. Dr Joseph Biederman is currently receiving research support from the following sources: Alza, AstraZeneca, Bristol Myers Squibb, Eli Lilly and

Co., Janssen Pharmaceuticals Inc., McNeil, Merck, Organon, Otsuka, Shire, NIMH and NICHHD. Dr Biederman is currently a consultant/advisory board member for the following pharmaceutical companies: Janssen, McNeil, Novartis and Shire. He is currently a speaker for the following speaker's bureaux: Janssen, McNeil, Novartis, Shire and UCB Pharma, Inc. In previous years, Dr. Biederman has received research support, consultation fees or speaker's fees for/ from the following additional sources: Abbott, AstraZeneca, Celltech, Cephalon, Eli Lilly and Co., Esai, Forest, Glaxo, Gliatech, NARSAD, NIDA, New River, Novartis, Noven, Neurosearch, Pfizer, Pharmacia, The Prechter Foundation, The Stanley Foundation and Wyeth. Dr Michael Monuteaux was a speaker in a symposia sponsored by Shire, Inc. Dr Faraone receives research support from and consults to Shire Pharmaceutical Development. He receives grant support from Eli Lilly and the NIH. Dr Wozniak has been a speaker for McNeil, Primedia/ MGH Psychiatry Academy, on the Advisory Board for Pfizer and Shire and received research support from NIMH, McNeil, Shire and Lilly. Her spouse, John Winkelman MD, PhD, has been on the Speakers Bureau for Boehringer-Ingelheim, Cephalon, Glaxo-SmithKline, King, Sanofi-Aventis, Sepracor, Takeda, on the Advisory Board for Axon Labs, Boehringer-Ingelheim, GlaxoSmithKline, Jazz Pharmaceuticals, Novartis, Neurogen, Novadel Pharma, Pfizer, UCB (Schwarz) Pharma, Sepracor, Takeda and received research support from Boehringer-Ingelheim, Glaxo-SmithKline, UCB (Schwarz) Pharma, Sepracor.

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