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A structural MRI study in monozygotic twins concordant or discordant for attention/hyperactivity problems: Evidence for genetic and environmental heterogeneity in the developing brain \mathbb{X}

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Several structural brain abnormalities have been reported in patients with Attention Deficit Hyperactivity Disorder (ADHD). However, the etiology of these brain changes is still unclear. To investigate genetic and environmental influences on ADHD related neurobiological changes, we performed Voxel-Based Morphometry on MRI scans from monozygotic (MZ) twins selected from a large longitudinal population database to be highly concordant or highly discordant for ratings on the Child Behavior Checklist Attention Problem scale (CBCL-AP). Children scoring low on the CBCL-AP are at low risk for ADHD, whereas children scoring high on this scale are at high-risk for ADHD. Brain differences between concordant high-risk twin pairs and concordant low-risk twin pairs likely reflect the genetic risk for ADHD; brain differences between the low-risk and high-risk twins from discordant MZ twin pairs reflect the environmental risk for ADHD. A major difference between comparisons of high and low-risk twins from concordant pairs and high/low twins from discordant pairs was found for the prefrontal lobes. The concordant high-risk pairs showed volume loss in orbitofrontal subdivisions. High-risk members from the discordant twin pairs exhibited volume reduction in the right inferior dorsolateral prefontal cortex. In addition, the posterior corpus callosum was compromised in concordant high-risk pairs, only. Our findings indicate that inattention and hyperactivity symptoms are associated with anatomical abnormalities of a distributed actionattentional network. Different brain areas of this network appear to be

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affected in inattention/hyperactivity caused by genetic (i.e., high concordant MZ pairs) vs. environmental (i.e., high-low discordant MZ pairs) risk factors. These results provide clues that further our understanding of brain alterations in ADHD. © 2007 Elsevier Inc. All rights reserved.

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Introduction

In this report we present an MRI study in monozygotic (MZ) twins carefully selected from a large database of longitudinal surveys on problem behavior to investigate genetic and environmental influences on structural brain changes underlying symptoms of inattentiveness and hyperactivity. Twins were selected based on extreme scores on the attention problem (AP) scale of the Child Behavior Checklist (CBCL4/18: [Verhulst et al., 1996\)](#page-16-0). Despite its name, this widely used inventory for the assessment of childhood behavioral disturbances contains not only items on attention problems but also questions related to hyperactivity/ impulsivity. The CBCL-AP scale has been shown to correlate well with the clinical evaluation of Attention Deficit Hyperactivity Disorder (ADHD) according to the Diagnostic and Statistical Manual of Mental Disorders-4th edition (DSM-IV: [American](#page-14-0) [Psychiatric Association, 1994](#page-14-0)). Recently, in a population of 1006 Dutch twins, a correlation between CBCL-AP and DSM-IVADHD scores of 0.6 was found ([Derks et al., 2006b\)](#page-14-0). Twins with high CBCL-AP scores may therefore be considered at risk for ADHD and we expect that our work provides clues for understanding structural brain abnormalities observed in children at risk for

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ADHD as well as in clinical samples of patients diagnosed with ADHD.

ADHD is a common neuropsychiatric disorder characterized by developmentally inappropriate symptoms of inattention, impulsivity and hyperactivity [\(American Psychiatric Association, 1994](#page-14-0)). ADHD is generally referred to as a childhood onset disorder with prevalence estimated at about 3–5% of the child population [\(Goldman et al.,](#page-15-0) [1998; Hudziak et al., 2000\)](#page-15-0). Recent follow-up studies have indicated that the behavioral abnormalities in ADHD may persist into adulthood, although symptoms at later ages may not be persistent enough to meet a DSM-IV diagnosis [\(Faraone et al., 2006](#page-14-0)).

Results from multiple twin studies, in many countries, indicate that the major influence (80%: [Hudziak et al., 2005; Neuman et al.,](#page-15-0) [1999; Rietveld et al., 2004;](#page-15-0) up to 90%: [Thapar et al., 1999](#page-16-0)) on attention problems and ADHD is genetic. There are a variety of theories about the neurobiology of ADHD currently in the literature. One theory that many consider central to ADHD pathophysiology is that ADHD symptoms reflect a deficit in executive function subserved by the frontal lobes and striatal brain regions ([Barkley, 1997; Pennington and Ozonoff, 1996; Sergeant et](#page-14-0) [al., 2002; Zametkin et al., 1990\)](#page-14-0), with possibly the cerebellum playing a role in modulations ([Giedd et al., 2001](#page-15-0)). The dopamine hypothesis of ADHD is based on the prediction that individuals with ADHD have impaired dopamine function in the mesolimbic dopamine system. Dopamine plays an important role in prefrontalsubcortical connectivity and furthermore, medicines shown to be effective in the treatment of ADHD such as methylphenidate enhances activity of frontal and striatal brain regions [\(Lou et al.,](#page-15-0) [1989; Vaidya and Gabrieli, 1999](#page-15-0)). Because of these putative relations, multiple candidate gene studies of the dopamine system (dopamine transporter, dopamine 4 receptor, dopamine 5 receptor) have been investigated. These reveal conflicting results with the strongest evidence of a relation between the DAT gene and ADHD, with a more complicated picture for DRD4 [\(Auerbach et al., 2001;](#page-14-0) [Durston et al., 2005](#page-14-0)). In addition to dopamine, other studies have also implicated genes for regulation of norepinephrine, nicotine, and others, involved in arousal and attentive behavior (for an overview see [Voeller, 2004\)](#page-16-0). As promising as the genetic findings have been, it is true that no clear relation between a single gene and the pathophysiology of ADHD has been discovered. The field has settled on a point, that ADHD is most likely an example of a complex disorder influenced in part by multiple genetic and environmental factors.

The study of environmental factors or of the interaction of genetic and environmental factors, presents an important step forward for our field. However, it is difficult to define what exactly an environmental factor is, partly because they have been poorly studied, but also because many so called environmental factors are themselves influenced by genetic factors [\(Eaves et al., 2005](#page-14-0)). Environmental risks may include those that disrupt normal fetal brain development such as maternal alcohol abuse or smoking during pregnancy ([Huizink and Mulder, 2006; Linnet et al., 2003](#page-15-0)). There is also evidence that maternal alcohol and smoking increase the risk for ADHD through an interaction with the DAT1 genotype [\(Brookes et al., 2006; Kahn et al., 2003\)](#page-14-0). Premature birth combined with low birth weight also constitutes an increased risk for ADHD [\(Bhutta and Anand, 2001\)](#page-14-0).

The dopamine system, particularly in the basal ganglia is highly susceptible to hypoxia and relations of ADHD behavior and hypoxia at birth have been reported ([Lou et al., 2004\)](#page-15-0). In addition, ADHD symptoms are seen after childhood stroke, particularly with lesions in the basal ganglia, especially the putamen ([Max et al.,](#page-15-0) [2002](#page-15-0)), and orbitofrontal and mesial prefrontal cortices ([Max et al.,](#page-15-0) [2005a](#page-15-0)), and traumatic brain injury affecting basal ganglia and orbitofrontal regions ([Gerring et al., 2000; Herskovits et al., 1999;](#page-15-0) [Max et al., 2005b\)](#page-15-0). Pheno-copies can also result from epilepsy [\(Schubert, 2005\)](#page-16-0), and various other neurological insults such as meningitis and autoimmune disorders ([Peterson et al., 2000; Wait](#page-15-0) [et al., 2002\)](#page-15-0).

Information on the contribution of environmental influences to ADHD risk comes from twin studies. To date, although contribution of shared environmental factors have shown to be important in a wide variety of human emotion, behavior, and activities ([Hudziak et al., 2003\)](#page-15-0), there have been no twin studies that provide evidence for the contribution of shared environment on ADHD. Rather, all unique, or non-shared environmental influences, defined as environmental factors that 'lead twins to be different' have been identified in most studies. Thus, much of the focus on the study of environmental mediators of ADHD must necessarily focus on unique insults. For example, despite general knowledge about environmental risk factors contributing to ADHD, except for cases with clear brain lesions, the exact neurobiological injuries or pathways mediating environmentally driven ADHD are still unclear. In addition, it is largely unknown whether environmental adversity affects brain structure and function in a similar way as genetic risk factors do, or whether the genetic and environmental etiologies follow partly different pathways. Also, environmental risk factors may act indirectly through the genome, for example by post-genomic modifications via methylation processes [\(Fraga et al., 2005\)](#page-14-0). To date there is only one study that employed a twin design to examine brain areas linked to ADHD of environmental origin with specific control of genetic factors. In this study by [Castellanos et al. \(2003\)](#page-14-0), anatomic brain magnetic resonance images (MRI) were collected from monozygotic (MZ) twins discordant for ADHD. Since MZ twins are genetically identical, the presence of ADHD in one twin but not the co-twin must originate from differential exposure to environmental risk factors, either directly or through environmental modification of gene expression. Interestingly, after computing volumes of brain areas selected from the MR images, it was found that the affected twins had smaller caudate volumes than their unaffected co-twins. This is in line with the current working hypothesis of a deficit in frontal–striatal processing with ADHD and furthermore suggests that the causal pathways leading to ADHD of environmental origin may, partially, overlap with those in ADHD of genetic origin.

To further our understanding of the genetic and environmental influence on brain defects in ADHD, we present a structural brain imaging study on a sample of twin pairs discordant for the risk of ADHD compared to pairs concordant for high risk of ADHD and pairs concordant for low risk of ADHD. Twin pairs were recruited based on longitudinal data on the CBCL-AP scale available at ages 7, 10 and 12 years. For the concordant groups, twin pairs were selected in which both members scored at either high or low extremes of the CBCL-AP. Inattention and hyperactivity traits are highly heritable and an influence of shared family environment has hardly ever been found [\(Rietveld et al., 2003](#page-15-0)). Therefore, these groups of concordant twins likely represent subjects with either high or low genetic vulnerability for ADHD and differences between these groups are expected to highlight brain structures related to the genetically mediated risk for ADHD. The discordant group consisted of twin pairs in which one member scored high

and the other low on the CBCL-AP. Intrapair differences for these twins are expected to indicate brain structures modified by adverse environmental exposure for the twins at high risk for ADHD but not their low-risk twin counterparts ([De Geus et al., 2006\)](#page-14-0). Instead of a region-of-interest approach we assessed differences in anatomy across the whole gray and white matter of the brain on a voxel by voxel basis using Voxel-Based Morphometry (VBM: [Ashburner and Friston, 2000, 2001\)](#page-14-0).

Methods

Participants

All twins were recruited from the Netherlands Twin Register (NTR: [Boomsma et al., 2002](#page-14-0)). The sample consisted of twins whose mothers had completed the NTR surveys at ages 0, 2, 3, 5, 7, 10, and/or 12 years $(n=6150 \text{ pairs})$. At ages 3, 7, 10 and 12 years the surveys included the CBCL/4-18. For the current purposes, we considered only twins with maternal CBCL-AP ratings from at least two time-points ($n=2887$ pairs) at ages 7, 10 or 12 who were classified as monozygotic with a questionnaire method ($n=1058$ pairs). Each child was classified as either at high risk or low risk for ADHD based on his/her standardized CBCL-AP T-scores (mean=50, $SD=10$) at ages 7, 10 and 12. T-scores were calculated separately in boys and girls. High-risk children had a T-score above 60 at all available time-points and a T-score above 65 at least once. Low-risk children had a T-score below 55 at all available time-points. Prior research by our groups and others has shown this to be a sensitive and specific technique to identify children at risk for clinical ADHD ([Chen et al., 1994;](#page-14-0) [Derks et al., 2006a; Hudziak et al., 2004; Lehn et al., 2007\)](#page-14-0). Using the CBCL-AP data, we identified twin pairs who were concordant high (both twins high CBCL-AP scores), concordant low (both twins low CBCL-AP scores) and discordant (one twin high and co-twin low CBCL-AP score). Two concordant low twin pairs were selected for each discordant pair, matched on sex, zygosity, parental socio-economic status, date of birth, handedness and maternal age. This selection yielded a total of 82 MZ twin pairs; 27 AP concordant high, 37 concordant low and 18 discordant. After the selection, the family of one discordant twin pair discontinued participation. This twin pair and the two associated concordant pairs were excluded. The remaining 79 twin families were invited for an interview with the mother of the twins, to which 72 families agreed (20 concordant high, 35 concordant low, 17 discordant), giving a response rate of 90%. Non-participants included families who could not be contacted $(n=2)$, declined $(n=3)$ or were unable to take part for practical reasons $(n=2)$. Data obtained with the interview and from NTR surveys (collected at ages 0, 2, 3, 5, 7, 10 and 12 years) were examined to determine the twins' suitability for participation in the MRI experiments. Twins were excluded with chronic medical or neurological disorders $(n=5)$; spastic disorder, congenital abnormalities, epilepsy, mitochondrial encephalomyopathy), mental disabilities $(n=3;$ pervasive developmental disorder, mental retardation), conditions that presumably would make participation too demanding $(n=2;$ combinations of various mental and physical problems) and use of psychotropic drugs $(n=2;$ Ritalin, anti-depressants). Concordant pairs who matched non-participating discordant pairs were also excluded $(n=6)$. This left 54 MZ twin pairs; 11 concordant high, 29 concordant low and 14 discordant twin pairs.

The families of 31 twin pairs (3 concordant high, 17 concordant low, 11 discordant) agreed to participate in the MRI experiments. The remaining declined $(n=12)$, could not participate due to orthodontic braces $(n=4)$ or could not be contacted $(n=1)$. Of these 31 pairs, 28 completed the MRI protocol (see below). All twins were asked to provide buccal cell samples (using mouth swabs) for DNA extraction. Zygosity testing included Polymerase Chain Reaction (PCR) of 11 highly polymorphic genetic markers. Three discordant twin pairs appeared dizygotic and were excluded, which leaves 25 pairs. The final study population included 3 concordant high, 17 concordant low and 5 discordant MZ twin pairs. To assess ADHD status, the mothers of the twins also completed a phone interview, administered by trained medical students, based on the Dutch version of the DISC-IV Parent Version (DISC-IV-P; [Lehn et al., 2007](#page-15-0)). None of the twins in our final population met the criteria for an ADHD diagnosis according to this clinical interview. Demographics and AP status of the twin sample are summarized in [Table 1.](#page-3-0) All twins and their parents provided written informed consent and the study was approved by the university ethical review board.

Image acquisition

The MRI session took about 30 min in which 3 separate whole head anatomical scans were consecutively acquired. During the entire scan session, the twins remained inside the scanner and were asked to minimize head movement during and between each scan. Scan order of the twins was randomized. Magnetic resonance imaging of the brain was performed on a 1.5 T Sonata MR system (Siemens, Erlangen, Germany) with a standard circularly polarized head coil. The anatomical scans included 160 sagittal slices at 1 mm intervals acquired with a 3D gradient-echo T1-weighted sequence (flip angle 15 \degree ; Repetition Time, TR = 1900 ms; Echo Time, $TE = 3.93$ ms; Inversion Time, $TI = 1100$ ms). In plane resolution was 256×224 pixels (pixel size 1 mm²), gray level resolution was 16 bit.

Volumetric analysis

Prior to volumetric analyses, the integrity of the acquired MR images was visually checked and for each twin, the second and third anatomical scans were co-registered to the first. The origin of each MRI volume was aligned on the anterior commissure landmark. Differences in brain anatomy were assessed on a voxel by voxel basis using the optimized voxel-based morphometry (VBM) technique proposed by [Good et al. \(2001\),](#page-15-0) an extension of the originally introduced standard VBM method ([Ashburner and](#page-14-0) [Friston, 2000, 2001\)](#page-14-0). The automated procedures of VBM were implemented using the MATLAB (The MathWorks, Inc.) VBM tools by Christian Gaser from the University of Jena, Germany, based on original scripts by John Ashburner and SPM2 analysis software (Wellcome Department of Imaging Neuroscience, London, UK). In a first step standard VBM is used to create study a specific T1 MR template for the whole brain and customized gray matter (GM), white matter (WM) and Cerebrospinal Fluid (CSF) priors. The template and priors were computed across the MRI images from all twin samples. For this purpose, each raw T1 MRI image was spatially normalized to a standard T1 template available in SPM2 ([Ashburner et al., 1997;](#page-14-0) [Ashburner and Friston, 1999](#page-14-0)). Subsequently, the three brain compartments were extracted after image segmentation using

Table 1

T-score: mean AP T-scores across 2 or 3 time points (7, 10 and 12 years), computed separately for boys and girls; Aggression T-score: aggression T-scores from CBCL at age 12, separately for boys and girls. Anxious Depressed *T*-score: anxious depression *T*-scores from CBCL at age 12, separately for boys and girls; Birth weight: birth weight of each twin in grams; Incubation time: number of days spend in incubator; Comorbidities: at the time of MRI scanning, interview data were obtained from the mothers of the twins on items regarding dyslexia, social phobia, separation anxiety disorder, specific phobia, generalized anxiety

Depressed T-score: anxious depression T-scores from CBCL at age 12, separately for boys and girls; Birth weight: birth weight of each twin in grams; Incubation time: number of days spend in incubator; Comorbidities: at the time of MRI scanning, interview data were obtained from the mothers of the twins on items regarding dyslexia, social phobia, separation anxiety disorder, specific phobia, generalized anxiety

disorder, depression, dysthymia, oppositional defiant disorder and conduct disorder. For twin pairs 11–17 of the concordant CBCL-AP low controls interview data were not collected.

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voxel by voxel probability mapping with respect to standard GM, WM and CSF priors. Finally, each volume was spatially low pass filtered using a 8 mm full width at half maximum (FWHM) isotropic Gaussian kernel and a smoothed study-specific T1 template and smoothed GM, WM and CSF compartments in standard stereotactic space were obtained by averaging the individual compartments across all twins.

In a following step, the raw MRI images were segmented based on the optimized procedure proposed by [Good et al. \(2001\).](#page-15-0) Compared to standard VBM, the optimized protocol includes a number of additional processing steps aimed at reducing the probability of voxel classification errors. First, the gray matter partition in native, not normalized, space was extracted from the raw T1 images using the newly obtained customized whole-brain T1 template and GM prior. Subsequently, the spatial normalization parameters for the GM compartment with respect to the GM template were estimated. The obtained deformation matrix/ parameter set was then applied to the raw whole-brain MR images. Subsequently, the whole-brain T1 image was resegmented using parametric mapping with respect to the customized priors and the normalized gray and white tissue compartments were extracted. The resolution of the extracted compartment images was 1 mm³ . During segmentation the images were corrected for intensity non-uniformities introduced by the MR scanner. To preserve volumetric information, a modulation step was added. This consists of multiplying each voxel intensity value by the determinant of the Jacobian matrix defining the spatial transformation from the original MR images to the normalized MRI template. In this way intensity values of voxels belonging to brain structures that are expanded during the warp are reduced, while voxel intensities from brain structures that are contracted are increased. Accordingly, voxel intensity values of the modulated images code for differences in regional brain size in the raw MR images. Before statistical analyses, the resultant modulated images were smoothed using a 12 mm FWHM isotropic Gaussian kernel. After smoothing, the image data correspond better with the Gaussian statistics underlying the statistical parametric mapping technique to detect volumetric differences in brain anatomy ([Worsley et al., 1996\)](#page-16-0). Spatial smoothing also has the effect of rendering the data more normally distributed and reduces the influence of inaccuracies in spatial normalization of individual brains on the following morphometric comparisons.

Statistical tests

Anatomical differences in the twin brains related to attention and hyperactivity problems were assessed by means of voxel by voxel parametric mapping using the ANOVA tests implemented in SPM2. Statistical tests were performed on modulated gray and white matter compartments. Tests comparing the concordant high with the concordant low twins were performed with for each twin the 3 scan replications as separate measurements (concordant high: 3 twin pairs with 3 replications in each member = 18 volumes; concordant low: 17 pairs with 3 replications in each member = 102 volumes). In these statistical comparisons a potential influence of twin pair sex may be present given that our concordant high twins were all girls, while the concordant low sample consisted of 10 female and 7 male twin pairs (see [Table 1\)](#page-3-0). We obtained qualitatively similar results, however, when we performed additional VBM control tests with the male twin pairs excluded from the concordant low sample. In the discordant twin pairs a within-subject ANOVA was performed comparing the 3 replicated compartment scans of each twin and co-twin. Dependence of error variances due to inclusion of measurement replications and correlated repeated measures were accounted for through the non-sphericity correction option in SPM2. Volumetric changes were assumed significant at $p < 0.0005$, uncorrected for individual voxels, with minimal cluster size of 2000 voxels.

Results

Global volumes

Bar graphs of [Fig. 1](#page-5-0) illustrate global volumes of gray matter (top) and white matter (bottom) averaged across the 6 individual tissue extractions of each twin pair (2 twins with 3 replications each). Scatter plots above each bar indicate deviations of compartment volumes with respect to the intrapair means. Statistical tests on global volumes revealed no significant differences for overall gray or white matter between the concordant high-risk twins (left bars) and concordant low-risk twins (middle bars) [Gray matter high vs. low, 728.23 ± 58.57 vs. 735.87 ± 33.85 ml: $F(118,1)=0.41, p=0.53$; White matter high vs. low, 411.25 \pm 27.34 vs. 420.27 ± 28.91 ml: $F(118,1) = 1.07$, $p=0.30$]. As expected, systematic volume differences between the twin and co-twin of each pair (assigned randomly) were also absent.

Deviations of individual volumes within the discordant twin pairs (scatter plots on the right) were of similar size to those observed in the concordant twins. However, there appeared to be a systematic difference; in pairs 1, 3, 4 and 5 for gray and 3, 4 and 5 for white tissue, volumes were smaller in the high-risk twin (open circles) compared to the low-risk co-twin (filled circles). These differences did not reach statistical significance [Gray high vs. low risk, 720.08 ± 71.27 vs. 740.03 ± 55.98 ml: $F(4,1) = 2.55$, $p= 0.19$; White high vs. low risk, 414.59 ± 59.30 vs. 425.20 ± 1.5 47.91 ml: $F(4,1) = 2.34, p = 0.20$].

Reductions for overall gray or white matter volume have been reported previously in patients with ADHD ([Carmona et al., 2005;](#page-14-0) [Durston, 2003\)](#page-14-0).

Voxelwise analyses

Concordant high-risk twin pairs vs. concordant low-risk twin pairs (genetic risk)

[Figs. 2 and 3](#page-7-0) and [Table 2](#page-9-0) show clusters of significant regional volumetric decreases and increases from the VBM group comparison between concordant high- and low-risk pairs.

Gray matter. Anatomical locations of clusters from the test on gray tissue can be appreciated in [Fig. 2.](#page-7-0) The high-risk twins exhibited significant gray matter deficits in medial parietal (cluster labeled A), left and right insular (B, C), left anterior temporal (D) and left orbitofrontal cortical areas (E).

Increased gray matter relative to the low-risk twins was found in the right sensorimotor (F) cortex, left and right prefrontal cortices (G, H) and a left mid temporal region (I).

White matter. VBM analysis of white matter indicated more widespread deficits for the concordant high sample ([Fig. 3:](#page-8-0) top). In line with the gray matter findings, a reduction of white matter compared to the low-risk controls was seen in the left orbitofrontal area (area labeled F). Furthermore, the white matter analysis also

Fig. 1. Global volumes of gray matter (top) and white matter (bottom) segmentations. Bars indicate mean compartment volumes across the 6 individual tissue extractions for each pair. Scatter plots indicate deviations of the individual estimates with respect to the intrapair means. For the CBCL-AP high concordants (left) and low concordants (middle), filled and open circles represent the 3 replication measures for twin and co-twin, with random assignment. For the discordants (right), filled and open circles indicate volumes for the CBCL-AP low and high twin.

indicated involvement of the right orbitofrontal region (G), as well as right and left prefrontal regions (areas B and J). In addition to frontal cortex, substantial areas of reduced white matter were found in the left and right temporal lobes (H, I) and in a region encompassing the posterior corpus callosum splenium (area D). Volume reduction of the splenium was accompanied by decrements in posterior white matter tracts, including a right posterior parietal region (A) and the left occiput (C). Finally, an area of reduced white matter was seen in the left cerebellar hemisphere (area E).

Regions of increased white matter [\(Fig. 3](#page-8-0): bottom) included the left parietal cortex (K) and right sensorimotor area (area L: close to a gray matter enhancement observed near this area).

High-risk vs. low-risk twin from discordant pairs (environmental risk)

[Figs. 4 and 5](#page-10-0) and [Table 3](#page-12-0) indicate clusters of significant volumetric decrements and increments in the high- vs. low-risk twin from discordant pairs.

Gray matter. Decreased gray matter in the high-risk children [\(Fig.](#page-10-0) [4](#page-10-0): top) was found in left and right posterior regions including left and right posterior parietal cortices (A, B), left occipitotemporal (C) and medial occipital (G) cortices, and the left cerebellar hemisphere (F). Additional gray matter deficit was found in the anterior part of the right hemisphere temporal lobe (D) and in the right inferior dorsolateral prefrontal cortex (E).

Relatively increased gray matter for high- compared to low-risk twin ([Fig. 4:](#page-10-0) bottom) was observed in a right frontal area (area H), the left dorsolateral prefrontal cortex (J), right cingulate (I) and left anterior temporal lobe (K).

White matter. White matter deficits in the high-risk twin of the discordant pairs ([Fig. 5:](#page-11-0) top) were observed in left hemisphere posterior parietal, mid temporal and sensorimotor areas (E, A, B), the right anterior corpus callosum (C) and right dorsolateral prefrontal cortex (D). Relatively increased myelination for the high-risk twin [\(Fig. 5:](#page-11-0) bottom) was found in left and right frontal brain regions (F and G).

Post hoc tests

To additionally test for the effect of correlated variances due to the inclusion of repeat scans, we also performed the statistical comparisons on averaged scans, across the 3 separate anatomical

Concordants: Gray matter

A: medial Parietal B: L. Insula C: R. Insula D: L. anterior Temporal E: L. Orbitofrontal

F: R. Sensorimotor G: L. PFC H: R. PFC I: L. Temporal

scans for each individual. This provided results qualitatively similar to the analyses on individual repeats; reflecting a trade-off between the benefit of enhanced image quality on one hand and cost in available data points (a data reduction by a factor of 3) on the other hand. To furthermore examine the effect of having twins of pairs together in groups, we also re-evaluated the contrast between concordant high and low AP samples, restricted both to average anatomical images as well as to one twin of each twin pair (two tests were run, one including only the first-born and one with only the second-born twin of each pair). As expected, due to the large negative impact on the available data (a data reduction by a factor of 6; 1 average scan across 3 repeat scans and 1 instead of 2 twins from each pair), the observed contrast test statistics were negatively affected. Nevertheless, the results still indicated involvement of brain areas similar to those observed in the primary tests. Taken together, these additional inferential tests indicate that the inclusion of measurement repeats did not artificially influence our statistical test results.

We did not find any evidence for volume changes of basal ganglia structures including the caudate and putamen, and limited evidence for volume reduction in the cerebellum (a local decrement of left cerebellar white matter in the high concordants and left cerebellar gray matter in the high discordants) and anterior temporal regions (areas of reduced gray matter in the left anterior temporal lobe for the high-risk concordants and right anterior temporal lobe for the high-risk discordants). In earlier studies on ADHD, gray matter volume reductions have been noted for these structures [\(Castellanos et al., 2002; Durston, 2003; Seidman et al.,](#page-14-0) [2005; Sowell et al., 2003\)](#page-14-0). Due to the nonstationary nature of neuroanatomy, brain tissue maps do not have a uniform smoothness which may have affected our statistical analyses [\(Ashburner and Friston, 2000; Worsley et al., 1999](#page-14-0)). In particular the use of a large cluster threshold (like the minimum of 2000 voxels we applied) may result in real sub-threshold group differences that are missed. To take into account a possible influence of spatial non-stationarities, we re-evaluated the gray matter tests applying an individual voxel-based threshold of p <0.05, adjusted for False Discovery Rate (FDR) combined with a cluster extent limited to 50 voxels. The top panel of [Fig. 6](#page-13-0) shows the ANOVA test results for volume reductions in the high vs. low concordants. Application of voxel-based statistics for this test provided no indication for cerebellar or basal ganglia disruptions. Relaxing the cluster extent threshold did, however, result in additional inclusion of left as well as right ([Fig. 6:](#page-13-0) area labeled C) temporal pole regions. In addition, orbitofrontal brain involvement was extended to include not only left but also mid and right orbitofrontal areas (areas labeled D and E). Evidence for bilateral volume reduction in orbitofrontal and temporal lobe regions is in line with our finding on white matter changes (areas labeled F, G, H, I in [Fig. 3](#page-8-0)). The bottom panel of [Fig. 6](#page-13-0) illustrates the within pair test results for gray matter reduction in the high vs. low discordant twins. In addition to an extension of involved brain regions already found in the original test, adjustment of the statistical criterion for

this contrast indicated a cluster of volume change extending over the right insular cortex and posterior part of the right putamen (area labeled F in [Fig. 6;](#page-13-0) bottom). A similar right hemisphere cluster, and an analogous cluster in the left hemisphere, were found in primary as well as post hoc tests for volume reductions in high vs. low concordant children (Fig. 2: areas B and C; [Fig. 6](#page-13-0) (top): areas A and B). In agreement with the primary analyses, we observed no involvement of left temporal lobe regions.

Discussion

We examined the brain anatomy of genetically identical twins concordant for high scores on the CBCL4/18 attention problem scale (CBCL-AP), concordant for low scores on the CBCL-AP and a sample of genetically identical twins discordant for CBCL-AP scores. A group comparison of structural brain MRI images between concordant high and concordant low scoring pairs was expected to indicate morphological changes related to the genetically mediated risk for ADHD. Intrapair differences between MR images of twin pairs highly discordant for the CBCL-AP score were expected to indicate morphological changes related to the environmentally mediated risk for ADHD.

Tests for anatomical changes using Voxel-Based Morphometry (VBM) revealed significant local brain abnormalities. In the concordant high compared to concordant low-risk twins, a prominent volume reduction was evident in the orbitofrontal regions, including significant loss of gray and white matter in the left and significant loss of white matter in the right orbitofrontal cortex. In the discordant comparison, gray matter loss for the highrisk twins was restricted to the right inferior dorsolateral prefrontal cortex, and accompanied by adjacent white matter deficits, including a region of the right anterior corpus callosum rostrum. Both group contrasts also revealed temporal lobe abnormalities. In the concordant high-risk pairs a reduction of gray matter was observed in a region of the left anterior temporal lobe, combined with white matter decrements in left and right mid temporal areas. In addition, there was evidence for gray matter loss in an anterior temporal region of the left as well as the right hemisphere after applying a more liberal statistical threshold. For the high-risk twins of the discordant pairs, gray matter loss was confined to the right anterior temporal cortex, adjacent to the area of gray matter deficit in the inferior dorsolateral prefrontal cortex. In addition to prefrontal and temporal lobe abnormalities, there were changes in the posterior brain. In the concordant high-risk twins, reduced white matter was found in the posterior corpus callosum splenium, together with decrements of retro-callosal medial parietal gray matter and right parietal and left occipital white matter. Additional areas of gray matter loss were observed in right and left insular cortices. In the high-risk twins from discordant pairs, right and left hemisphere gray matter disturbances in occipital, occipitotemporal and posterior parietal regions were found. The posterior corpus callosum appeared normal. Finally, we found an indication for involvement of the cerebellum. In the left cerebellar hemisphere,

Fig. 2. Parametric t-maps for the ANOVA test on regional gray matter volume reductions (top) and increases (bottom) in the CBCL-AP high relative to the AP low concordant twin pairs. On the left; maximal intensity projections of voxels with uncorrected p <0.0005 in axial and sagittal views. On the right; areas of significant difference projected on average MR sections of the 40 concordant twins, with t-values mapped by the color bar. Horizontal dashed lines in the maximal intensity projections indicate the level of the displayed axial MR slices. Dashed white lines in the axial projections indicate positions of the sagittal and coronal MR sections. PFC = prefrontal cortex.

Fig. 3. Statistical maps for the ANOVA test on regional white matter reductions (top) and increases (bottom) in the CBCL-AP high relative to low concordant twin pairs.

an area of white matter loss was found in the high-risk concordant pairs and an area of gray matter loss in the high-risk twin from discordant pairs.

The brain areas with volume loss related to attention/ hyperactivity problems as measured with the CBCL in the present study are consistent with earlier reports on structural brain alterations in ADHD. Several anatomical imaging studies reported ADHD related volume decrements in orbitofrontal ([Carmona et al.,](#page-14-0) [2005; Giedd et al., 2001](#page-14-0)) and dorsolateral prefrontal lobes ([Carmona et al., 2005; Castellanos et al., 1996; Durston et al.,](#page-14-0) [2004; Filipek et al., 1997; Hill et al., 2003; Kates et al., 2002;](#page-14-0) [Overmeyer et al., 2001; Semrud-Clikeman et al., 2000\)](#page-14-0). Orbitofrontal reductions in ADHD children have generally been found largest in the right hemisphere [\(Carmona et al., 2005; Giedd et al.,](#page-14-0)

Regions of significant brain size differences between concordant CBCL-AP high and low twin pairs. Tissue: tissue type tested (gray or white matter); Contrast: test for significant decreases or increases in CBCL-AP high relative to low twins; Cluster label: alphabetical cluster label as displayed in anatomical overlays of [Figs. 2 and 3;](#page-7-0) Peak-coordinates (mm): location of cluster-voxel with largest effect size (in MNI space); Z score: z-value of voxel with largest effect size. p-value: p-value of cluster, corrected for multiple comparisons at cluster-level; # voxels: number of voxels in cluster; Anatomical location: location of cluster. For cases where regional differences correspond to local maxima within a large interconnected cluster, the p-value and voxel-count of this cluster are displayed in single, merged, cells.

[2001](#page-14-0)). One study also reported orbitofrontal volume loss in adult ADHD subjects, localized in the left hemisphere ([Hesslinger et al.,](#page-15-0) [2002](#page-15-0)). Reduced dorsolateral prefrontal cortex volume has been reported with right [\(Castellanos et al., 1996; Durston et al., 2004;](#page-14-0) [Filipek et al., 1997; Hill et al., 2003; Overmeyer et al., 2001;](#page-14-0) [Semrud-Clikeman et al., 2000\)](#page-14-0) as well as left hemisphere [\(Carmona et al., 2005; Kates et al., 2002](#page-14-0)) dominance. Orbitofrontal and dorsolateral prefrontal brain deficits are in line both with earlier and more recent cognitive models of impaired executive function/inhibition in ADHD ([Barkley, 1997; Castellanos et al.,](#page-14-0) [2006](#page-14-0)). The orbitofrontal subdivisions of the prefrontal lobes are presumed to participate in goal directed processing through inhibition of distracting information and impulses, while the dorsolateral prefrontal areas are hypothesized to contribute by supporting action planning and programming and working memory [\(Fuster, 2002\)](#page-15-0). Orbitofrontal and dorsolateral prefrontal cortex dysfunction due to lesions or epilepsy typically results in attention problems and emotional instability and impulsivity, similar to symptoms of ADHD ([Eslinger et al., 2004; Max et al., 2005a,b;](#page-14-0) [Powell et al., 1997](#page-14-0)).

Our finding of volume reductions in temporal and occipitoparietal brain areas and cerebellum, i.e., regions outside those predicted by the current fronto-striatal model of ADHD, also agrees with earlier work (for a review see [Seidman et al., 2005](#page-16-0)). Volume reduction of the temporal lobes in ADHD has been demonstrated in a cross-sectional anatomical MRI study in ADHD children and adolescents by [Castellanos et al. \(2002\).](#page-14-0) In addition, [Sowell et al. \(2003\)](#page-16-0) applied a regional morphometric analysis technique on brain MR images of ADHD children and observed gray matter reductions in both anterior temporal lobes combined with gray matter defects in inferior dorsolateral prefrontal regions. Many recent anatomical MRI studies also confirm the presence of posterior brain deficits, including the posterior corpus callosum splenium [\(Hill et al., 2003; Lyoo et al., 1996; Semrud-Clikeman](#page-15-0) [et al., 1994\)](#page-15-0) and occipital and parietal lobe regions ([Castellanos](#page-14-0) [et al., 2002; Durston et al., 2004; Filipek et al., 1997](#page-14-0)). Volume reduction of the cerebellum has also consistently been reported in previous neuroimaging studies on ADHD children [\(Carmona et al.,](#page-14-0) [2005; Castellanos et al., 1996; Durston, 2003; Hill et al., 2003;](#page-14-0) [Krain and Castellanos, 2006; Seidman et al., 2005](#page-14-0)).

Dysfunction of the cerebellum and occipitoparietal and temporal brain regions may well relate to attentional impairment and hyperactivity. The cerebellum is assumed to be involved in the modulation of frontal–striatal activity ([Giedd et al., 2001](#page-15-0)). The occipital cortex is relevant for visual processing and the posterior corpus callosum for inter-hemispheric transfer of visual information. The posterior parietal cortex provides access to a multimodal representation of extra personal space and is relevant for spatial attention [\(Posner and Petersen, 1990](#page-15-0)). Finally, the anterior temporal lobes are presumed to be higher association areas that regulate access to and binding of perceptual information from different sensory modalities to establish face-object recognition [\(Ungerleider and Haxby, 1994](#page-16-0)). It has been suggested that posterior and temporal brain regions interact with the prefrontal cortex in a large-scale network that controls attentional focus and prevents distracting impulses [\(Mesulam, 1998](#page-15-0)). Our findings indicate that the attentional deficits in our high-risk twins relate to anatomical abnormalities of this proposed action-attentional

Discordants: Gray matter

Fig. 4. Parametric t-maps for the within-subject ANOVA test on regional gray matter reductions (top) and increases (bottom) in the CBCL-AP high compared to low scoring members of the discordant twin pairs. Areas of significant difference are projected on average MR sections of the 10 discordant twins. PFC = prefrontal cortex.

network and furthermore that disruption of this network appears to be located at different brain areas for attention and hyperactivity problems of genetic and environmental etiology. Most prominently,

there was a difference with regard to the prefrontal lobe regions involved; in genetically mediated risk for ADHD mainly the orbitofrontal subdivisions and in environmentally mediated risk

Discordants: White matter

Fig. 5. Statistical maps for the within-subject ANOVA test on white matter deficits (top) and enhancements (bottom) in the CBCL-AP high vs. low discordants. PFC = prefrontal cortex.

mainly the inferior dorsolateral prefrontal cortex showed morphological changes. An additional difference was noted for the posterior brain; decreased volume of the posterior corpus callosum was found for attention/hyperactivity problems of genetic origin, only.

Our findings on brain deficits associated with environmental stressors differ from a recent report on ADHD discordant MZ twins by [Castellanos et al. \(2003\)](#page-14-0). In a region of interest approach including segmentations of frontal, temporal, parietal and occipital lobes as well as caudate and cerebellum, these researchers found ADHD related volume deficits only for the caudate nuclei. In contrast, we observed volume deficits for several neocortical areas and the cerebellum, but not the striatum. This discrepancy can have several explanations. For example, the absence of caudate deficits may indicate that basal ganglia abnormalities are more severe in patients clinically diagnosed with ADHD compared to individuals at strong risk for the disorder based on CBCL-AP scores. Our failure to find caudate changes may also be due to a vulnerability to false negatives, because the current sample (5 discordant pairs) is smaller compared to the Castellanos et al. study (9 discordant pairs). Alternatively, in another recent report [Castellanos et al.](#page-14-0) [\(2002\)](#page-14-0) found that children with ADHD start out with smaller caudate nuclei than normal controls, but that caudate size decreases with maturation and more so for normal controls than ADHD children. Consequently, any caudate size difference during childhood becomes less apparent at later ages. The fact that the twins in our sample were of relatively higher age (11.8–16.9 years) compared to the [Castellanos et al. \(2003\)](#page-14-0) sample (5.6–15.6 years) may therefore also explain the dissimilar findings. Finally, there are differences in the techniques to examine morphologic changes.

Regions of significant brain size differences between CBCL-AP high and low scoring children of discordant twin pairs. Tissue: tissue type tested (gray or white matter); Contrast: test for significant decreases or increases in the AP high twins relative to the AP low co-twins; Cluster label: alphabetical cluster label as displayed in anatomical overlays of [Figs. 4 and 5;](#page-10-0) Peak-coordinates (mm): location of cluster-voxel with largest effect size (in MNI space); Z score: z-value of voxel with largest effect size. p-value: p-value of cluster, corrected for multiple comparisons at cluster-level; # voxels: number of voxels in cluster; Anatomical location: location of cluster. When regional differences correspond to local maxima within a large interconnected cluster, the p-value and voxel-count of this cluster are displayed in single, merged, cells.

Voxel-based approaches like the present one have shown to correlate well with region of interest methods, but complementary use of both comparison types has been advocated ([Giuliani et al.,](#page-15-0) [2005; Kubicki et al., 2002\)](#page-15-0). It should be noted though that a recent neuroimaging study that applied a voxel-based comparison in ADHD children did find evidence for regional basal ganglia deficits ([Overmeyer et al., 2001\)](#page-15-0).

In addition to regional volume loss, our results also indicated clusters of increased brain tissue, in both anterior and posterior brain regions for the concordant high-risk pairs and restricted to the frontal brain for the high-risk twin from discordant pairs. The finding of gray and white tissue increments appears to contrast with earlier ADHD studies. It should be noted, however, that earlier results predominantly relied on region of interest methods. Voxel by voxel approaches likely afford a more detailed examination of local morphological changes. Recently, [Sowell et](#page-16-0) [al. \(2003\)](#page-16-0) used a computational technique and also found locally increased gray matter in ADHD children, mainly restricted to posterior temporal and inferior parietal cortices. Gray/white matter increments may exist for many reasons. Neural tissue enhancement might be introduced by a regional lack of synaptic pruning during brain development [\(Huttenlocher, 1979](#page-15-0)). Alternatively, local volume increments may reflect a plastic adjustment to compensate for neural loss and consequently reduced processing capability of adjacent brain regions. It has also been proposed that disruption of white matter in a certain brain region may result in increased gray matter in the same area [\(Sowell et al., 2003\)](#page-16-0). Conversely, regions with decreased gray matter may be paired with locally enhanced white matter. Our data, however, provide no conclusive evidence of systematic inverse relations between regional gray and white matter volume changes. Additional research is needed to replicate the presence of locally increased cerebral tissue in attention/hyperactivity problems and further our understanding about the neuropathophysiology underlying these morphological changes.

The present sample size does not afford us to draw conclusions about the environmental risk factors for attention/hyperactivity problems. The sample selected for MRI was, however, drawn from a larger population of concordant high and low, and high-low discordant twin pairs selected for behavioral characterization [\(Lehn](#page-15-0) [et al., 2007\)](#page-15-0). Statistical analysis on survey data (maternal, teacher and self reports) from the 19 CBCL-AP discordant pairs in that study, indicated that the within pair difference in attention/ hyperactivity symptomatology existed since toddlerhood (age 3) and persisted at ages 5, 7, 10 and 12. Compared to their low-risk co-twins, the CBCL-AP high discordant twins also reported other behavioral problems. They showed more rule breaking behavior, more social problems and were more aggressive, withdrawn and anxious/depressed. T-scores for aggression and anxious depression in the present MRI subsample (from CBCL at age 12) are listed in [Table 1](#page-3-0). With regard to the environmental conditions predictive of inattention/hyperactivity, it was found that the discordants "at risk" for ADHD experienced significantly more adversity during infancy (lower birth weight and longer incubation). Our subsample reflected a tendency for lower birth weight, but not longer incubation [\(Table 1](#page-3-0)). The behavioral study also indicated that the CBCL-AP high discordants showed disadvantaged maturation (delayed physical growth and lagged early motor development, specifically with regard to developmental milestones of learning to role over and sit upright).

Fig. 6. Post hoc VBM test results with individual voxel-based threshold of p <0.05, adjusted for False Discovery Rate (FDR), and cluster extend threshold limited to 50 voxels. Top panel: ANOVA results for the test of gray matter volume reduction in the CBCL-AP high concordant vs. AP low concordant twins. Bottom panel: within twin pair ANOVA results for the test of gray matter volume reduction in the AP high vs. low member of the discordant twin pairs. In each panel from top to bottom; maximal intensity projections of suprathreshold voxels, suprathreshold voxel clusters projected on selected average MR slices and voxel clusters projected on a smoothed rendering of a standard brain available in SPM2. The t-value mapping is indicated by color bars. Positions of individual MR slices are indicated by red and white axes crossings in maximal intensity projections and MR images, respectively.

In conclusion, the present neuroimaging study suggests that the neurobiological pathways underlying attention and hyperactivity symptoms of genetic or external environmental origin are partly distinct. Behavioral problems for both etiologies appear to reflect reduced function of an action-attentional network of cerebellar, occipital/parietal and temporal brain regions interacting with the prefrontal cortex. This network seems to be compromised at different brain areas, especially within the prefrontal lobes. In attention deficit/hyperactivity of genetic origin the medial orbitofrontal subdivisions and in attention/hyperactivity problems of environmental origin the right inferior dorsolateral prefrontal cortex were abnormal. A difference was also noted for the posterior corpus callosum which was only disrupted in attention/hyperactivity disturbances of genetic origin. These inferences could be made by examining anatomical MRI images of genetically identical twins at "selective" genetic or environmental risk for ADHD as inferred from the CBCL-AP inventory. In the general population, symptoms of inattention and hyperactivity are likely caused by shared genetic and environmental risk factors. Therefore individuals with inattentiveness/hyperactivity may exhibit anatomical abnormalities in different areas of the action-attentional network. This may importantly contribute to the widespread brain deficits and general heterogeneity of these deficits reported in earlier neuroanatomical studies on ADHD patients.

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