

Multicenter Mapping of Structural Network Alterations in Autism

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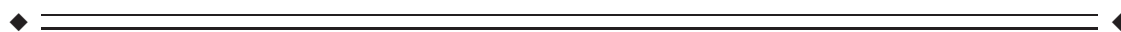
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Abstract: Autism spectrum disorders (ASD) are a group of neurodevelopmental conditions primarily characterized by abnormalities in social cognition. Abundant previous functional MRI studies have shown atypical activity in networks encompassing medial prefrontal cortex (mPFC) and medial parietal regions corresponding to posterior cingulate cortex and precuneus (PCC/PCU). Conversely, studies assessing structural brain anomalies in ASD have been rather inconsistent. The current work evaluated whether structural changes in ASD can be reliably detected in a large multicenter dataset. Our comprehensive structural MRI framework encompassed cortical thickness mapping and structural covariance analysis based on three independent samples comprising individuals with ASD and controls ($n = 220$), selected from the Autism Brain Imaging Data Exchange open-access database. Surface-based analysis revealed increased cortical thickness in ASD relative to controls in mPFC and lateral prefrontal cortex. Clusters encompassing mPFC were embedded in altered inter-regional covariance networks, showing decreased covariance in ASD relative to controls primarily to PCC/PCU and inferior parietal

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regions. Cortical thickness increases and covariance reductions in ASD were consistent, yet of variable effect size, across the different sites evaluated and measurable both in children and adults. Our multi-site study shows regional and network-level structural alterations in mPFC in ASD that, possibly, relate to atypical socio-cognitive functions in this condition. *Hum Brain Mapp* 36:2364–2373, 2015. © 2015 Wiley Periodicals, Inc.

Key words: ASD; ABIDE; neocortex; connectivity; medial PFC; connectome

INTRODUCTION

Autism spectrum disorders (ASD) are a group of life-long neurodevelopmental disorders currently recognized in more than 1% of children (Baird, et al., 2006; CDC, 2014). Core to ASD are impairments in socio-cognitive functioning thought to stem from atypical Theory of Mind (ToM), which is also known as mentalizing (Castelli, et al., 2002; Frith, 2003) as well as sensory abnormalities and restricted, repetitive patterns of behavior, interests, or activities (APA, 2013). Functional MRI studies have shown altered activations in this condition, with somehow diverse location and direction of functional findings. Despite this variability, regions subserving ToM function, such as medial prefrontal cortex (mPFC) and the posterior cingulate cortex and precuneus (PCC/PCU) in the medial parietal lobe, have been relatively consistently implicated in ASD (Di Martino, et al., 2009; Frith, 2003; Kana, et al., 2009; Kennedy and Courchesne, 2008; Lombardo, et al., 2011). In contrast, structural abnormalities of these same areas have been reported less consistently.

Adult studies have provided little consensus regarding the location and direction of *regional* alterations in ASD. Some have reported primarily grey matter reductions in frontal, parietal, and temporal cortices (Hadjikhani, et al., 2007; Scheel, et al., 2011); others have suggested increased grey matter in frontal and temporal regions (Doyle-Thomas, et al., 2013; Duerden, et al., 2012; Ecker, et al., 2012). Likewise, while some studies in children with ASD have shown medial and lateral prefrontal cortical thickness increases (Mak-Fan, et al., 2012; Raznahan, et al., 2010), others reported decreases in lateral orbitofrontal grey matter (Girgis, et al., 2007; Hardan, et al., 2006a). Divergences across studies may relate to the small sample sizes studied in single labs, due to the high costs and challenges associated to recruitment and MRI acquisition in ASD. Another possible reason for inconsistency is that rather than focal abnormalities, dysconnections among large-scale network characterize autism (Minshew and Williams, 2007; Schaefer, et al., 2013).

Approaches allowing examinations of both local and large-scale brain abnormalities are likely better suited for autism research. Cortical thickness analyses meet such criterion. They can provide a detailed picture of the topography of structural alterations in ASD as well as information on *inter-regional* network formation. Specifi-

cally, in recent years, the framework of MRI covariance analysis has been proposed to probe inter-regional networks that likely reflect the coordination of structural growth during development and maturation (Alexander-Bloch, et al., 2013a; Lerch, et al., 2006). While developmental conditions, such as ASD, likely relate to disruptions of morphological coupling across brain regions, only relatively few previous covariance analyses have been conducted in this disorder (Bernhardt, et al., 2014c; Dziobek, et al., 2010; McAlonan, et al., 2005). Importantly, only small samples have been assessed (ASD/Controls in Bernhardt et al.: 18/18; Dziobek et al.: 27/29; McAlonan et al.: 17/17), limiting generalizability of these results.

In the current work, we took advantage of the open-access Autism Brain Imaging Data Exchange database (ABIDE) to overcome limitations related to sample size and facilitate the evaluation of across-site reproducibility (Di Martino, et al., 2014). We carried out a multisite MRI assessment of patterns of regional and inter-regional structural differences in ASD. Analyses were based on three large and independent ABIDE subsamples, selected as they included both children and adults with autism and typical controls. We mapped differences between ASD and controls in regional cortical thickness as well as *inter-regional* structural covariance networks. Given their role in mediating socio-cognitive functions and previous task-based as well as task-free functional findings showing alterations of these regions in ASD (Di Martino, et al., 2009; Di Martino, et al., 2014), we expected to identify alterations in ASD primarily in mPFC and PCC/PCU and their inter-regional relationships. Nevertheless, given that the extent of the abnormalities beyond circuits involved in social cognition is still unclear, our cortical thickness analyses were carried out in an unconstrained fashion across the entire cortical mantle.

MATERIALS AND METHODS

Subjects

We selected a subsample of the ABIDE database, including children and adults (age range = 6.5–50.2 years). Specifically, we selected only sites that included both children and adults, with at least 10 individual datasets per diagnostic group after quality control (please, see below). Moreover, we selected only males given the low

TABLE I. Sample breakdown by site, age, ADOS total score, and full-scale IQ are presented in mean \pm SD (range)

site	Age	ADOS	IQ
USM			
ASD ($n = 52$)	23.6 \pm 7.6 (15–50)	13.6 \pm 3.3 (6–21)	100.6 \pm 16.5 (65–132)
Controls ($n = 40$)	21.5 \pm 7.8 (9–39)	–	115.3 \pm 13.9 (89–148)
PITT			
ASD ($n = 20$)	20.8 \pm 7.3 (12–35)	12.7 \pm 3.0 (8–19)	113 \pm 14 (81–131)
Controls ($n = 22$)	19.7 \pm 6.9 (9–33)	–	110 \pm 9 (95–127)
NYU			
ASD ($n = 35$)	16.8 \pm 7.5 (7–39)	11.3 \pm 4.2 (5–22)	105.3 \pm 13.8 (76–137)
Controls ($n = 51$)	17.5 \pm 6.7 (7–31)	–	115.0 \pm 12.1 (81–139)

prevalence of data from females that were not present in all sites. This left us with an initial sample of 297 individuals from three sites: (1) NYU Langone Medical Center (NYU, $n = 147$, 68/79 ASD/controls); (2) University of Utah, School of Medicine (USM, $n = 101$, 58/43 ASD/controls); (3) University of Pittsburgh, School of Medicine (PITT, $n = 49$, 26/23 ASD/controls).

As detailed at http://fcon_1000.projects.nitric.org/indi/abide, individuals with ASD had DSM-IV-TR diagnosis of Autistic Disorder, Asperger’s Disorder, or Pervasive Developmental Disorder Not-Otherwise-Specified, established by expert clinical opinion aided by ‘gold standard’ diagnostic instruments: the Autism Diagnostic Observation Schedule, ADOS (Lord, et al., 2000), and/or the Autism Diagnostic Interview-Revised, ADI-R (Lord, et al., 1994). In the NYU, PITT, and USM datasets, individuals diagnosed with associated disorders such as Fragile-X or tuberous sclerosis were excluded. Intelligence (full scale IQ, performance IQ and verbal IQ) was measured by WASI, WAIS III, and/or WISC III (Wechsler, 1999). Controls were free of history of mental disorders and matched for age at each site.

MRI Acquisition

High-resolution T1-weighted images were available from all three sites. NYU data were acquired on a 3T Siemens Magnetom Allegra using a 3D-TurboFLASH sequence (TR = 2530 ms; TE = 3.25 ms; TI = 1100 ms; flip angle = 7°; 128 sagittal slices; matrix size = 256 \times 256; FOV = 256 mm; slice thickness = 1.33 mm, yielding a voxel size of 1.3 \times 1.0 \times 1.3 mm³). PITT data were acquired on a 3T Siemens Magnetom Allegra using a 3D-MPRAGE sequence (TR = 2100 ms; TE = 3.93 ms; TI = 1000 ms; flip angle = 7°; 176 sagittal slices; matrix size = 269 \times 269; FOV = 269 mm; slice thickness = 1.05 mm, yielding a voxel size of 1.1 \times 1.1 \times 1.1 mm³). USM data were acquired on a 3T Siemens Magnetom TrioTim using a 3D-MPRAGE sequence (TR = 2300 ms; TE = 2.91 ms; TI = 900 ms; flip angle = 9°; 160 sagittal slices; matrix size = 240 \times 256; FOV = 256 mm; slice thickness = 1.2 mm, yielding a voxel size of 1.0 \times 1.0 \times 1.2 mm³).

MRI-Based Cortical Thickness Measurements

FreeSurfer (Version 5.1.0; <http://surfer.nmr.mgh.harvard.edu>) was used to generate models of the cortical surface and to measure cortical thickness from the T1-weighted images. Previous work has validated FreeSurfer by comparing it with histological analysis (Rosas, et al., 2002) and manual measurements (Kuperberg, et al., 2003). The processing steps have been described in detail elsewhere (Dale, et al., 1999; Fischl, et al., 1999; Han, et al., 2006). Individual surfaces were aligned to an average spherical representation, fsaverage5, improving correspondence of measurement points with regards to (mostly primary and secondary) sulcation patterns. For whole-brain analysis, thickness data were smoothed on the tessellated surfaces using a 20 mm full-width-at-half-maximum Gaussian kernel prior to statistical analysis. Selecting a surface-based kernel reduces measurement noise but preserves the capacity for anatomical localization, as it respects cortical topological features (Lerch and Evans, 2005).

Quality Control and Final Sample Selection

The ABIDE open-access dataset includes structural and functional data of a wide range of image quality. Cortex extractions in each subject were visually inspected and segmentation inaccuracies manually corrected by two raters blind to ASD diagnosis (SLV and BCB). ASD versus control status was assigned to the raters in a randomized fashion. Moreover, subjects with faulty segmentations, movement, or other artifacts were excluded from the study ($n = 77$). This left us with a final sample of 220 individuals (107 ASD, 113 controls).

We automatically measured the signal to noise ratio, based on the QA-tools associated with FreeSurfer (<https://surfer.nmr.mgh.harvard.edu/fswiki/QATools>), providing a rater-independent quantitative index of image quality.

In the final sample, ADOS scores were available for all individuals with ASD (ADOS total: mean \pm SD = 12.3 \pm 3.7, range = 5–22; ADOS social: 8.4 \pm 2.7, 2–14; ADOS communication: 4.3 \pm 1.5, 0–8). ASD and control groups had a comparable age (ASD: 20.9 \pm 8.0 years, 7.2–50.2 years; Controls: 19.3 \pm 7.3 years, 6.5–39.4 years; Difference: $t = -1.48$;

$P = 0.14$). On the other hand, compared to controls individuals with ASD had lower full-scale IQ (ASD: 104 ± 16 , 65–137; Controls: 114 ± 12 , 81–148; Difference: $t = 5.11$; $P < 0.001$), performance IQ (ASD: 106 ± 15 , 72–133; Controls: 112 ± 13 , 67–155; Difference $t = 3.09$; $P < 0.002$), and verbal IQ (ASD: 102 ± 17 , 55–136; Controls: 113 ± 12 , 80–140; Difference $t = 5.67$; $P < 0.001$). For a breakdown by site after quality control, please see Table I.

Statistical Analyses

As in previous structural MRI analyses (Bernhardt, et al., 2014a; Bernhardt, et al., 2014b; Bernhardt, et al., 2014c), we used SurfStat for Matlab [R2010a, The Mathworks, Natick, MA] (Worsley, et al., 2009). Analyses were carried out in an unconstrained fashion, at each neocortical surface point (henceforth, *vertex*).

- a. *Mapping of cortical thickness differences.* We fitted a linear model at each vertex i that assessed between-group differences in thickness T between ASD and controls:

$$T_i = \beta_0 + \beta_1 * Site + \beta_2 * Age + \beta_3 * IQ + \beta_4 * Group$$

Where T_i is the thickness at vertex i , *Site* is a term controlling for site (*i.e.*, NYU, PITT, USM), *Age* a term controlling for age, *IQ* is a term controlling for full-scale IQ, *Group* is the group factor (*i.e.*, ASD and controls).

Post hoc analyses were carried out to evaluate the consistency of findings (see *Results*) within each of the three sites, and within children and adults separately.

In a series of separate post hoc analyses, we repeated the above group comparisons after correcting for global mean thickness, after correcting for signal to noise ratio, and without correction for IQ.

- b. *Covariance network analysis.* We mapped structural covariance networks by seeding from each cluster of significant ASD-specific findings (*from a*) to all other cortical vertices.

$$T_i = \beta_0 + \beta_1 * Site + \beta_2 * Age + \beta_3 * IQ + \beta_4 * Glob + \beta_5 * Seed$$

Seed indicates the mean thickness of a cluster of ASD-specific findings. This analysis was carried out separately in ASD and in controls. The above covariance analyses were restricted to networks ipsilateral to the respective seed regions. As in previous covariance analyses (Bernhardt, et al., 2011; Chen, et al., 2008; He, et al., 2008), we additionally corrected for global mean thickness, *Glob*, in the above model.

Using linear interaction models, we assessed whether ASD diagnosis resulted in a modulation of seed covariance relative to controls.

$$T_i = \beta_0 + \beta_1 * Site + \beta_2 * Age + \beta_3 * IQ + \beta_4 * Glob + \beta_5 * Seed + \beta_6 * Group + \beta_7 * (Seed * Group)$$

Separate post hoc analyses evaluated whether findings were consistent when no correction for IQ and global mean thickness were applied.

- c. *Assessment of age effects.* We assessed the interplay between aging and ASD-specific structural alterations by measuring interactions between *age* and diagnostic *group* on cortical thickness.

$$T_i = \beta_0 + \beta_1 * Site + \beta_2 * Age + \beta_3 * IQ + \beta_4 * Group + \beta_5 * (Group * Age)$$

Analyses were carried out at each vertex; in a separate analysis, we also assessed age effects on the mean thickness of clusters of ASD-specific cortical thickness changes (*from a*).

We assessed interactions between age and ASD diagnosis on structural covariance networks. To this end, we assessed the triple interaction

$$Seed * Age * Group$$

- This contrast was evaluated based on a model that also contained each of these terms as simple effect and their simple interactions, together with the *Site*, *IQ*, and *Glob* control terms. As in *b*), seeds were based on clusters of significant ASD-specific findings.
- d. *Correction for multiple comparisons.* Surface-based findings were corrected using random field theory for non-isotropic images (Worsley, et al., 1999). Specifically, we applied the SurfStat function *SurfStatP.m*, which controlled the chance of reporting a family-wise error (FWE) in a given statistical analysis to $P_{FWE} < 0.05$.

RESULTS

Cortical Thickness Increases in ASD Relative to Controls

Surface-based comparisons of cortical thickness between ASD and controls revealed increased thickness in the former group in bilateral clusters extending from mPFC (ventromedial and dorsomedial PFC) to lateral PFC ($P_{FWE} < 0.05$; Cohens d for thickness increase left PFC = 0.44; right PFC = 0.53). Increases in mPFC thickness were relatively consistent across both hemispheres; conversely, lateral PFC thickness increases were more extensive in the right hemisphere, where thickening was seen in orbitofrontal, ventrolateral and dorsolateral regions; lateral PFC thickening in the left hemisphere was mostly restricted to dorsolateral PFC. There were no findings of decreased thickness in ASD relative to typically developing controls (Fig. 1).

We ran a post hoc analysis on mean thickness in clusters of ASD-related thickening to assess the consistency of our effect across sites. This analysis indicated that bilateral medial and lateral prefrontal cortical thickness increases in

REGIONAL CORTICAL THICKNESS ANALYSIS

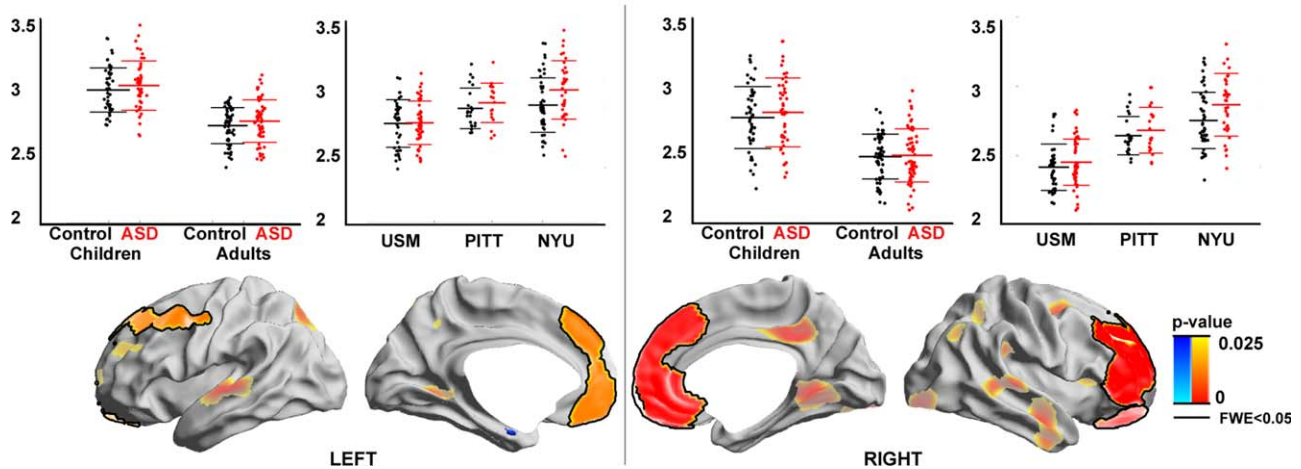


Figure 1.

Regional cortical thickness analysis. ASD-specific cortical thickness increases relative to controls across three sites (in a model that controlled for age, IQ, and site) are shown in red/decreases in blue. Significant regions after multiple comparisons correction at a cluster-level of $FWE < 0.05$ (thresholded using random field theory for nonisotropic images) are surrounded by solid black

outlines. To illustrate trends, findings at $P < 0.025$, uncorrected (no black outlines, semitransparent) are also shown. Inset scatter plots show mean effects for each significant cluster, either within each site (i.e., USM/PITT/NYU) or within children and adults separately.

ASD were consistent across the three sites, albeit with variable effect size (Cohen's d in NYU/ PITT/ USM; left PFC = 0.60/0.44/0.25; right PFC = 0.63/0.40/0.43). Restricting the sample to either children or adults revealed consistent prefrontal thickening in ASD in both age groups (Cohen's d in Children/Adults; left PFC = 0.5/0.42; right PFC = 0.69/0.44).

Running between-group comparisons while additionally controlling for global mean thickness yielded virtually identical patterns of results (post hoc t -value left/right PFC = 3.11/4.13). Likewise, running a model that did not correct for IQ also revealed bilateral PFC thickening (post hoc t -value left/right PFC = 3.60/4.07). Global mean thickness did not differ between groups (ASD/Controls = $2.66 \pm 0.15/2.65 \pm 0.14$; $P > 0.1$ in model correcting for site, age, and IQ).

ANOVA on signal to noise ratio did not indicate any interaction between site and group ($F = 0.17$, $P < 0.84$) or a main effect of group ($F = 2.48$, $P > 0.1$); however, there was a significant effect of site ($F < 19.3$, $P < 0.001$). Running between group-comparisons after controlling for signal-to-noise confirmed the robustness of ASD-specific thickness increases in PFC (post hoc t -value left/right PFC = 3.36/3.93).

Abnormal Inter-Regional Structural Covariance in ASD

We complemented the regional cortical thickness mapping with an assessment of inter-regional structural

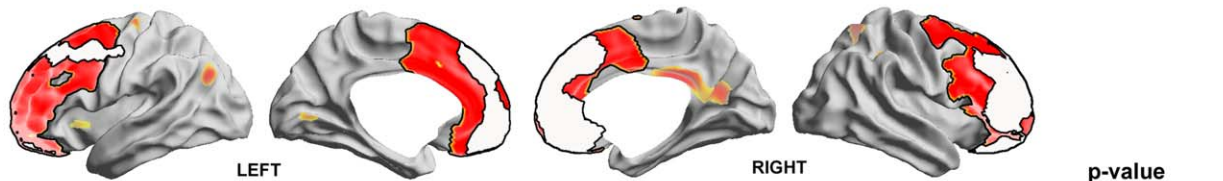
covariance networks (Fig. 2). Networks were centered on the left and right PFC clusters that showed ASD-specific cortical thickness increases in our prior analyses (see Fig. 1).

In controls, the left PFC seed showed high covariance to extended cluster encompassing medial prefrontal, lateral prefrontal, and anterior midcingulate regions as well as trends to a cluster in TPJ. Similarly, the right PFC seed showed covariance increases to medial and lateral prefrontal extending to superior regions as well as trends to PCC/PCU. Group comparisons for covariance did not reveal any differences in the left PFC. However; covariance of the right PFC cluster was markedly reduced in ASD relative to controls, with target regions in right parietal regions encompassing PCC/PCU ($P_{FWE} < 0.05$).

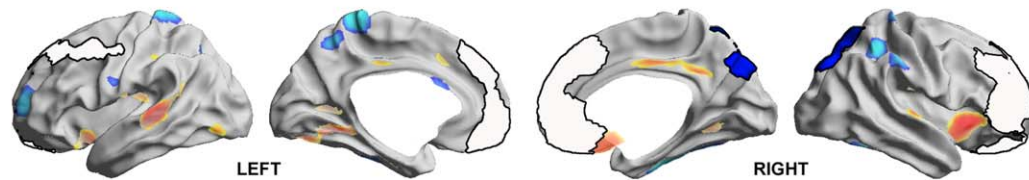
Post hoc analysis, which assessed correlations between mean thickness of the seed in right PFC and the target region in right PCC/PCU, indicated that covariance reductions had a similar direction across all three sites, but again with variable effect (t -value of $Group \times Seed$ interactions in NYU/PITT/USM = 1.7/0.8/1.6). Moreover, similar to our findings in the regional analysis, running covariance without the control for global mean thickness yielded similar results (post hoc t -value = 2.4), as did a covariance analysis between the seed and target when no IQ was corrected (post hoc t -value = 3.8), and when rerunning the covariance analysis while additionally controlling for signal-to-noise confounds (post hoc t -value = 3.7).

ASD-SPECIFIC DISRUPTIONS OF INTER-REGIONAL STRUCTURAL COVARIANCE

A Covariance networks in controls



B Network alterations in ASD



C Interaction

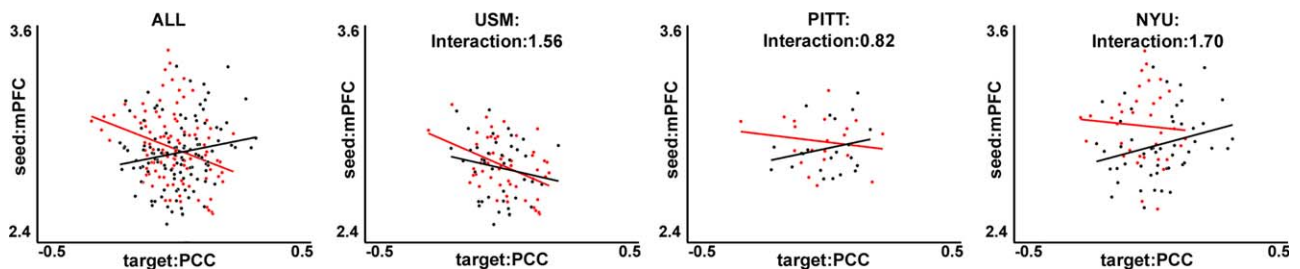


Figure 2.

Inter-regional structural covariance network analysis. Seed regions were determined as clusters of cortical thickness increases in ASD (see Fig. 1). **(A)** Covariance networks in controls. Significant correlations between cortical thickness in the seed and a cortical target region across the sample were interpreted as structural networks. **(B)** Alterations in structural covariance networks between ASD and controls. Blue/Red areas

indicate reductions/increases in seed covariance in ASD relative to controls. **(C)** Scatter plots illustrating the interaction showing in **(B)**, between mean thickness of right PFC seed and mean thickness of right PCC/PCU target region, once for the overall sample and once within each site (*i.e.*, USM/PITT/NYU). Please, see Figure 1 for details on the statistical thresholding.

Network Modulations by Age

Both controls and ASD groups showed marked age-related thinning across multiple cortical regions, surface-based interaction analysis failed to show any significant between-group differences in age-dependent structural change following correction for multiple comparisons. At uncorrected thresholds ($P < 0.025$), trends for more marked age-related thinning in ASD were observed in scattered clusters in left temporal, right prefrontal and midline parietal regions, whereas more protracted age-related thinning

in ASD was seen in insular regions. Similarly, we did not find any significant between-group differences in the modulation of inter-regional covariance by age (Supplementary Figure).

DISCUSSION

We performed a comprehensive examination of cortical thickness and its inter-regional covariance to study structural network alterations based on the three independent

ABIDE subsamples that included both children and adults. In our study, cortical thickness analyses were carried out in a surface-based and unconstrained fashion. Yet, in the light of several previous behavioral and functional studies suggesting an association between ASD and impairments in socio-cognitive functioning, particularly atypical ToM (Castelli, et al., 2002; Hill and Frith, 2003), we expected to reveal structural network abnormalities primarily in regions and inter-regional networks associated with such functional processes, such as mPFC and PCC/PCU (Di Martino, et al., 2009; Di Martino, et al., 2014). We indeed observed cortical thickness increases in individuals with ASD in large portions of bilateral mPFC, extending to lateral PFC in both hemispheres. While of variable effect size, the direction of changes in PFC regions was consistent across all three sites and age groups alike. Complementary covariance analysis revealed that mPFC regions of thickness increases also showed abnormal structural network integration with parietal regions encompassing PCC/PCU. Our findings, therefore, provide multisite evidence for structural regional and connectivity disruptions particularly in areas known to subservise socio-cognitive functioning, such as mPFC and medial parietal cortices.

Previous neuroimaging studies have reported mixed findings in children and adults with autism. In children, brain overgrowth has been reported in frontal and temporal cortices (Courchesne, et al., 2001; Girgis, et al., 2007; Hardan, et al., 2006b), while other studies did not observe any apparent change (Amaral, et al., 2008; Bloss and Courchesne, 2007; Redcay and Courchesne, 2005). MRI studies in adults have also reported rather divergent findings (Doyle-Thomas, et al., 2013; Ecker, et al., 2012; Hadjikhani, et al., 2007; Scheel, et al., 2011). While recent findings across several mouse models of autism have also pointed towards a high variability in structural phenotypes (Ellegood, et al., 2014), diverse findings in human studies could be driven variable inclusion criteria. In the current study, only individuals with ASD supported by ADOS and/or ADI-R were included. Opting for homogeneous subject inclusion criteria may be an important factor, given the somewhat inconsistent nature of previous functional as well as structural findings in this condition. Our series of post hoc thickness comparisons between groups revealed a consistent direction of change between ASD and controls; yet, thickness increases were of variable effect size across the different centers studied. Such variability, which could be due to other variations in cohorts, scanners, and scanning protocols, together with possible limitations in sample size may have, in part, contributed to the diverse pattern seen across previous reports. While the present work also drew attention to significant difference in the signal-to-noise ratio across sites, no clear relationship was found between the ability to detect diagnostic group differences and variations in signal-to-noise ratio. It is important to note that it is difficult to pinpoint the exact effect of signal-to-noise on results, due to

various causes of differences in signal-to-noise such as movement, scanner parameters, and physiological effects.

A recent study (Haar, et al., 2014) on the ABIDE dataset reported rather subtle differences in cortical structure between ASD and controls. At low thresholds, the study of Haar and colleagues also revealed tendencies for rather increases in cortical thickness in ASD—a finding in accordance to the work presented here. The higher effect sizes reported in the current work could be attributed to differences in center- and case-inclusion. Indeed, the current study included those three sites that provided data from adults as well as children, while the study of Haar and colleagues included data from more centers, and preselected cases based on explicit matching criteria. Furthermore divergences may be attributed to whether manual FreeSurfer correction procedures were applied, as in the current work, or not (Haar, et al., 2014).

Direct histopathological validation studies of MRI-derived morphological changes in autism are virtually absent. Notably, a previous *post mortem* study of Casanova and colleagues suggested a smaller width but increased number of *minicolumns*, neuronal assemblies centered on radially oriented pyramidal neurons, in this condition (Casanova, et al., 2002). Further pathological studies of individuals with autism have reported glial abnormalities in the frontal lobe (Morgan, et al., 2010; Vargas, et al., 2005), with one study showing increased microglial density in white and grey matter (Morgan, et al., 2010). Last, histopathological analyses have reported laminar rearrangement and a poorly defined grey and white matter interface (Avino and Hutsler, 2010; Mukaetova-Ladinska, et al., 2004), a common sign of atypical migration and cortical organization during neurodevelopment, starting possibly already at prenatal developmental stages (Stoner, et al., 2014).

Univariate cortical thickness mapping was complemented by MRI covariance analysis to probe inter-regional structural MRI networks (Alexander-Bloch, et al., 2013b; He, et al., 2008; Lerch, et al., 2006). According to its underlying assumptions, cortical regions belonging to the same network would show highly correlated growth during development, due to their high propensity to exchange trophic factors and participate in common molecular signaling pathways (Alexander-Bloch, et al., 2013b). In our sample, we observed decreased covariance of mPFC in ASD relative to controls, particularly to parietal regions, including PCU/PCC. Similar to the regional thickness findings, the direction of effects remained consistent across sites, but of variable effect size. Functional MRI connectivity analysis have consistently reported ASD-related abnormalities in the inter-regional coupling of different divisions in the frontal neocortex (Di Martino, et al., 2014; Just, et al., 2012; Lynch, et al., 2013; Monk, et al., 2009; Redcay, et al., 2013). A previous meta-analysis of 14 diffusion MRI studies has furthermore suggested long-range white matter diffusion abnormalities in individuals with autism, especially in superior longitudinal and uncinate

fasciculi as well as the corpus callosum, suggesting alterations in long-distance frontal structural connectivity in this condition (Aoki, et al., 2013). *Post mortem* studies have supported these findings, showing possible histological underpinnings of atypical connectivity in ASD (Hutsler and Zhang, 2010; Raymond, et al., 1996; Zikopoulos and Barbas, 2010; Zikopoulos and Barbas, 2013). In a recent study, Zikopoulos and Barbas could show fewer large myelinated axons in the deep white matter, which mediate long-range connectivity, in ASD than in controls below regions corresponding to the anterior cingulate cortex and mPFC. On the other hand, the authors reported a concomitant increase in thin myelinated axons in the superficial white matter below the same regions, indicative of excess axonal branching and local hyper-connectivity (Zikopoulos and Barbas, 2010).

Previous research has reported abnormal trajectories of cortical structural development in ASD (Raznahan, et al., 2010; Scheel, et al., 2011; Schumann, et al., 2010; Wallace, et al., 2010). In this work, on the other hand, only trends for an age-dependent modulation of cortical thickness changes were observed. In addition to differences in sample characteristics between our work and previous studies (with differences in the mean age and/or age range), a reduced sensitivity in the current work may have resulted from the absence of an age-stratified sampling scheme when pooling data across centers.

By studying the publically accessible ABIDE dataset, we had the opportunity to investigate structural brain abnormalities in a large, multisite sample of individuals with ASD. Although no direct behavioral markers of social cognition were available in this open access repository, the co-occurrence of thickness increases and covariance disruptions in mPFC is remarkable, given the key role of this region in socio-cognitive functions related to ToM (Frith and Frith, 2006; Kana, et al., 2009). Ultimately, our structural MRI findings appear to support theories that hypothesize structural and connectional anomalies in prefrontal and/or parietal midline regions in ASD that may, likely, relate to atypical social cognition in this condition.

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