



Published in final edited form as:

*Prog Neuropsychopharmacol Biol Psychiatry*. 2013 January 10; 40: 187–192. doi:10.1016/j.pnpbp.2012.08.003.

## Alterations in the Cerebral White Matter of Genetic High Risk Offspring of Patients with Schizophrenia Spectrum Disorder

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### Abstract

Alterations in White Matter (WM) may be seen in young relatives at risk and may underlie vulnerability to Schizophrenia. We were interested in exploring which of the WM regions were altered in adolescent offspring at familial risk for Schizophrenia. We examined structural alterations in the offspring of subjects with Schizophrenia or Schizoaffective disorder (HR; n=65; 36 males) and healthy controls (HC; n=80: 37 males) matched for age and education. MRI images were collected using a GE 1.5T scanner at the University of Pittsburgh Medical Center. Image processing was done using FreeSurfer (MGH) by an experienced rater blind to clinical data. We used multivariate analysis of covariance, with intracranial volume ( $p > 0.05$ ) and age as covariates. High Risk offspring had significant reductions in total WM, hemispheric WM and WM within left parietal and left cingulate cortices. Male offspring had more pronounced right hemisphere WM reductions than females.

### Keywords

Schizophrenia; Genetic High-Risk; Offspring; FreeSurfer; MRI

## 1. Introduction

Schizophrenia represents a group of probably etiologically heterogeneous, severe mental disorders the neurobiological underpinnings of which are not yet fully understood (Keshavan et al, 2008a; MacDonald and Schulz, 2009). Schizophrenia has been thought to result from “disconnectivity” of white matter systems (Friston and Frith, 1995). It has been proposed that alterations in the white matter connectivity may give rise to the some of the symptoms found in schizophrenia. For example, Crow (1998) suggested that auditory hallucinations could arise from aberrant communication between language centers in the left

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#### Disclosure/Conflict of Interest Statement

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and right temporal cortices. Hubl and colleagues reported alterations in lateral parts of the arcuate fasciculus in patients with hallucinations (Hubl et al. 2004). Aberrations in white matter connectivity between distributed systems could also lead to disorders of self-monitoring which in turn could lead to auditory hallucinations (Silbersweig and Stern, 1996, 1998).

Morphometric studies using ROI methodology have shown that in addition to the gray matter abnormalities, regional and global white matter volumes also have been compromised in schizophrenia (reviewed in Shenton et al., 2001). This is supported by several studies using different methodologies such as voxel based Morphometry [Giuliani et al, 2005; Meda et al., 2008], deformation based Morphometry (Davatzikos, 2005) and ADC (Apparent diffusion coefficient) based Morphometry (Ardekani et al., 2005). These various methods have shown, albeit with differing results, that white matter volumes in the brain are altered in schizophrenia. A recent review and meta-analysis (Olabi et al., 2011) of 928 patients and 867 controls examining 32 brain regions showed that patients' annual WM volume reduction in several brain regions was  $-0.32\%$  in the frontal,  $-0.32\%$  in the parietal, and  $-0.39\%$  temporal lobes.

While there is growing evidence that schizophrenia subjects show changes in white matter (Kubicki et al., 2007), (Bloemen et al, 2010), there is very little known about whether these alterations are related to underlying liability to the illness. Family history remains one of the strongest etiologic factors in schizophrenia, with an estimated heritability of almost 80% (Mcgrath et al., 2008). The study of young relatives of patients with schizophrenia therefore offers a unique window into the premorbid liability to the illness (Keshavan et al, 2008b). The Preclinical (premorbid and prodromal) phase of the illness has been investigated in studies (Lawrie et al., 2001; Lymer et al., 2006; Lawrie et al., 2008) that have shown white matter alterations.

Although the evidence for white matter involvement is accumulating, it is unclear whether this involvement is limited to certain regions or whether they are widespread, and whether the familial susceptibility may have a role in the neurodevelopmental aspects of WM. To assess the potential role of WM in disrupted neurodevelopmental processes, we examined volumetric alterations in the white matter of familial HR subjects and Healthy controls. Few studies have looked at total WM volume, hemispheric WM volumes and regional WM volumes across both hemispheres in high risk subjects. These studies have not yielded consistent results (reviewed in Agnew-Blais & Seidman, in press; reviewed in Agnew-Blais & Seidman, 2012; Boos et al., 2007; meta-analysis).

We hypothesized that given their familial susceptibility for the illness, subjects at high risk for Schizophrenia would show volumetric WM alterations compared to age and sex matched healthy controls.

## 2. METHODS

### 2.1. Participants

The study was conducted at the Western Psychiatric Institute and Clinic, Pittsburgh. Sixty five racially diverse adolescents or young adult offspring (OS) of of schizophrenia probands and eighty healthy controls (HC) were recruited. The overall sample characteristics have been described in previous reports (Francis et al, 2011). Twenty eight OS subjects had one parent with Schizoaffective disorder (SZA) and thirty seven OS subjects has one parent with SZ. Of these, twenty three fathers had a diagnosis of SZ/SZA, while forty two mothers had a diagnosis of SZ/SZA. Offspring of parents with schizophrenia or schizoaffective disorder were recruited by approaching patients in the clinic and through advertisements. Healthy

Controls were recruited through advertisements in the same community as OS subjects. Diagnostic assessments of HC and OS and parental diagnoses of schizophrenia or schizoaffective disorder used the structured clinical interviews for DSM-IV diagnoses (SCID)(First et al., 1995) and were confirmed using consensus meetings led by senior diagnosticians (M.S.K and D.M). Participants with an IQ < 80, lifetime evidence of a psychotic disorder, exposure to antipsychotic medications or anti-depressant medications, current or recent (within the previous month) substance use disorder, significant neurological or unstable medical conditions were excluded. All participants signed informed consent after the study was fully explained to them. For participants < 18 years of age, consent was provided by the parent or guardian, and the subjects provided informed assent. The study was approved by the University of Pittsburgh Institutional Review Board.

## 2.2. Image Acquisition

MRI scans were obtained on subjects using a GE 1.5T whole body scanner (GE Medical Systems, Milwaukee, Wisconsin). The detailed scanning protocol has been described in an earlier publication (Gilbert et al, 2001). The scans were T1 weighted images: three-dimensional spoiled gradient recalled (SPGR), acquired in a steady-state pulse sequence (124 coronal slices, 1.5 mm thickness, TE=5 msec, TR=25 msec, acquisition matrix=256×192, FOV=24 cm, flip angle 40°). Approximately 10% of the MRI scans with radio frequency inhomogeneity defects and motion artifacts were not included in the analysis.

## 2.3 Image Analysis

**2.3.1 Semi-automated morphometric analysis using FreeSurfer**—We used FreeSurfer (FS) 4.0.5 (64 bit version) (Massachusetts General Hospital, (Fischl et al, 2004; Desikan et al, 2006; Dale et al, 1999; Fischl et al, 1999; van der Kouwe et al, 2008;); running on Linux for morphometric analysis. FS, a semiautomated brain image morphometric software, has been used to study the brain morphology of several illnesses including schizophrenia (Fischl and Wald, 2007; Fischl et al, 2002;). FS has 3 automated stages (Fischl et al, 2004), each followed by manual image editing by an Image analyst (AF). Image processing included motion correction of volumetric T1-weighted images, removal of non-brain tissue using a hybrid watershed/surface deformation procedure (Segonne et al., 2004), automated Talairach transformation, segmentation of the subcortical white matter and deep gray matter volumetric structures (Fischl et al., 2002, 2004b), intensity normalization, tessellation of the gray matter white matter boundary, automated topology correction (Fischl et al., 2001; Segonne et al., 2007), and surface deformation following intensity gradients to optimally place the gray/white and gray/CSF borders at the location where the greatest shift in intensity defines the transition to the other tissue class (Dale et al., 1999). Detailed pictures of WM segmentation may be seen in Salat et al, (2009).

We chose FreeSurfer's semi-automated morphometric analyses for specific reasons. Inter-subject variability and random error inherent in manual techniques is minimized by automated methods. A second reason is the economy of time for completion. By contrast, the possibility of systematic error in automated approaches can be detected and corrected by rigorous manual editing, such as that followed in this study. Each scan was carefully checked for errors of template registration, skull strip, segmentation and parcellation at each stage of the image processing. We performed Inter-Rater reliability between AF and NT (see acknowledgements) on the 33 WM regions of ten brains and obtained an average inter class correlation value = .99 (WM) and .96 (GM).

Our objective was to examine early WM volume alterations in the regions of the left and right hemispheres of non-psychotic High risk subjects and controls. Since two decades of

research studies have consistently implicated regions within the frontal, temporal and parietal, and cingulate WM in the etiopathogenesis of SZ (reviewed in Shenton et al, 2001), we examined the WM volumes in these lobes. Although we did not include the other regions within the MANCOVA, we present ANCOVA analysis of the WM of all the other regions in Tables 1 and 2. The corpus callosum was not included since it was published elsewhere (Francis et al, 2011).

## 2.4. Statistical Analysis

White matter volumes were normally distributed [Shapiro–Wilk’s test (W statistic,  $p > 0.1$ )]. Multivariate analyses of covariance (MANCOVA) were used with Group and Sex as categorical predictors and Age and ICV as covariates to test differences in white matter volumes between offspring and healthy controls. We first compared total, left and right hemispheric WM across groups. We next performed several MANCOVA analyses on each of the lobes (right and left Frontal, Temporal, Parietal lobes, Motor-Somatosensory cortex and Cingulate cortices) as dependent variables. Tables 2 and 3 show the regions included in the MANCOVAs. Significant MANCOVAs were followed by univariate ANCOVAs to identify specific regional volumetric deficits. Univariate ANCOVAs were carried out on other regions in secondary analysis. Bonferroni corrections were used on the MANCOVAs and ANCOVA tests to control for multiple comparisons. We derived a Bonferroni corrected p value by dividing 0.05 by the number of MANCOVAs and the univariate tests within each MANCOVA. This alpha value was then used to determine whether each p value survived the correction or not.

## 3. RESULTS

### 3.1. Total and Hemispheric White Matter

OS subjects had significantly reduced total white matter  $F(1, 145) = 15.54$   $p < 0.0001$ .

OS subjects had significantly less WM in the left hemisphere [HR - 204734.22 mm<sup>3</sup> (SD) 27586] [HC - 218976.6 mm<sup>3</sup> (SD) 27728.85] ( $F(1, 145) = 14.83$   $p < 0.0001$ ) and right hemisphere [HR-206716.82 mm<sup>3</sup> (SD) 28635.97] [HC - 221522.54 mm<sup>3</sup> (SD) 27932.82] ( $F(1, 145) = 15.46$   $p < 0.0001$ ).

### 3.2. Lobar effects

The MANCOVAs showed that left parietal [ $F(6,142) = 3.42$   $p < 0.004$ ], left motor-somatosensory [ $F(3,142) = 3.3$   $p < 0.02$ ] left cingulate [ $F(4,141) = 3.44$   $p < 0.01$ ] and right temporal [ $F(8,141) = 2.5$   $p < 0.01$ ], right parietal [ $F(6,142) = 2.5$   $p < 0.02$ ] had significantly reduced volume in High risk offspring compared to healthy controls. Only the left parietal cortex WM and left cingulate WM survived the correction for multiple MANCOVAs. Tables 2 and 3 shows the ANCOVA analysis for all the white matter regions in high risk subjects when compared to healthy control subjects.

### 3.3. Gender, Age and White Matter

There was a significant group X sex interaction in the right hemisphere WM volume  $F(1, 145) = 4.6408$ ,  $p < 0.032$ . Although HR subjects had smaller RH WM volumes than their healthy counterparts, this appeared to be more pronounced in the males than females. There were no significant sex by group interaction in the left hemisphere WM. White matter correlated positively with age for all assessed regions but these correlations did not significantly differ between the groups.

## 4. Discussion

White matter constitutes the anatomical infrastructure for inter and intra hemispheric connectivity. In a sample of non-psychotic genetic high risk offspring and age matched healthy controls, our study showed that total WM volume in the right hemisphere was reduced by 6.68 % and the left hemisphere was reduced by 6.5 % in high risk subjects indicative of a developing vulnerability to the illness. In addition, male HR brains showed a greater reduction in the right hemisphere WM when compared to females. Our study showed that WM in the posterior regions of the brain namely left parietal cortex, and left cingulate cortex were significantly reduced in HR subjects. A study done on 22q11 deletion syndrome, which is a risk model for Schizophrenia, showed similar WM reductions in the parietal, temporal and occipital areas of the brain (da Silva Alves et al, 2011). It is possible that such a reduction of WM volume early in the preclinical phase may present as a decrease in the white matter fiber density in schizophrenia as shown in several DTI studies (Karlsgodt et al, 2009, Bloemen et al, 2010). In a separate study on the same sample, we showed that the splenium volume of the Corpus Callosum (Francis et al, 2011) was significantly reduced lending evidence that the inter-hemispheric connectivity between temporo-parietal regions could also be altered in this sample.

ROI based studies have shown that frontal WM is compromised in SZ (Brier et al., 1992; Buchanan et al, 2004; Sanfilipo et al, 2000; Wible et al., 2001; Hulshoff-Pol 2002; Mathalon et al, 2003 and Buchsbaum et al, 2006). In a meta-analysis of 17 VBM studies of WM, Di et al, (2009) showed WM alterations in the right and medial frontal WM regions which was consistent with the ROI studies. In addition, they showed that the internal capsule WM is also altered. This is consistent with several DTI studies (Szeszko et al., 2005; Kubicki et al, 2005, 2007; Buchsbaum et al., 2006). In summary, WM alterations in the left hemisphere in HR subjects observed in this study lend evidence to the concept of a loss of connectivity between regions as a possible marker for susceptibility in schizophrenia. It is possible that such abnormalities may underlie a later misconnection syndrome as hypothesized by several investigators of white matter involvement in schizophrenia (Karlsgodt et al, 2009). More studies are warranted since genetic influences on WM volume and their relationship with cognitive abilities in genetic High risk subjects and healthy individuals are only being presently unraveled. While morphometric studies of WM such as the present study provide important clues to altered short and long range connectivity of WM fasciculi in the brain, Diffusion Tensor Imaging studies and fMRI connectivity studies are more likely to definitively inform these issues.

## Acknowledgments

### Role of Funding Source

MH064023 (MSK)

MH045203 (MSK)

NARSAD Established Investigator award (MSK)

We are grateful to Diana Mermon MS for her assistance in clinical assessments and recruitment. Dr. Vaibhav Diwadkar helped with diagnostic ascertainments. Neeraj Tandon helped with the reliability analysis of FreeSurfer measurements.

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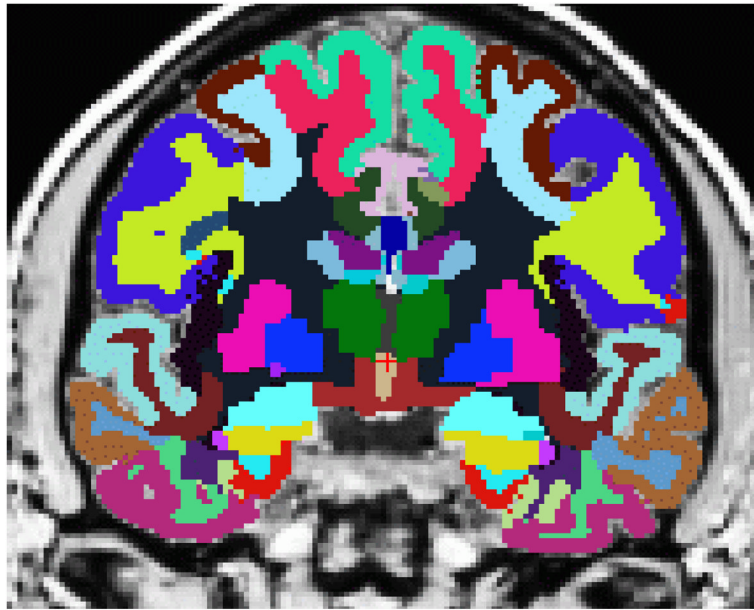
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### Highlights

- Our study compares WM volumes in Genetic High Risk against Control subjects.
- WM volume in the R hemisphere was reduced by 6.68 % in high risk subjects.
- Left hemisphere WM was also reduced by 6.5 %.
- Male HR brains showed a greater reduction in the R hemisphere WM.
- WM volumes of L parietal lobe, and L cingulate cortex were reduced in HR subjects.



**Fig 1.**  
FreeSurfer White Matter Segmentation:

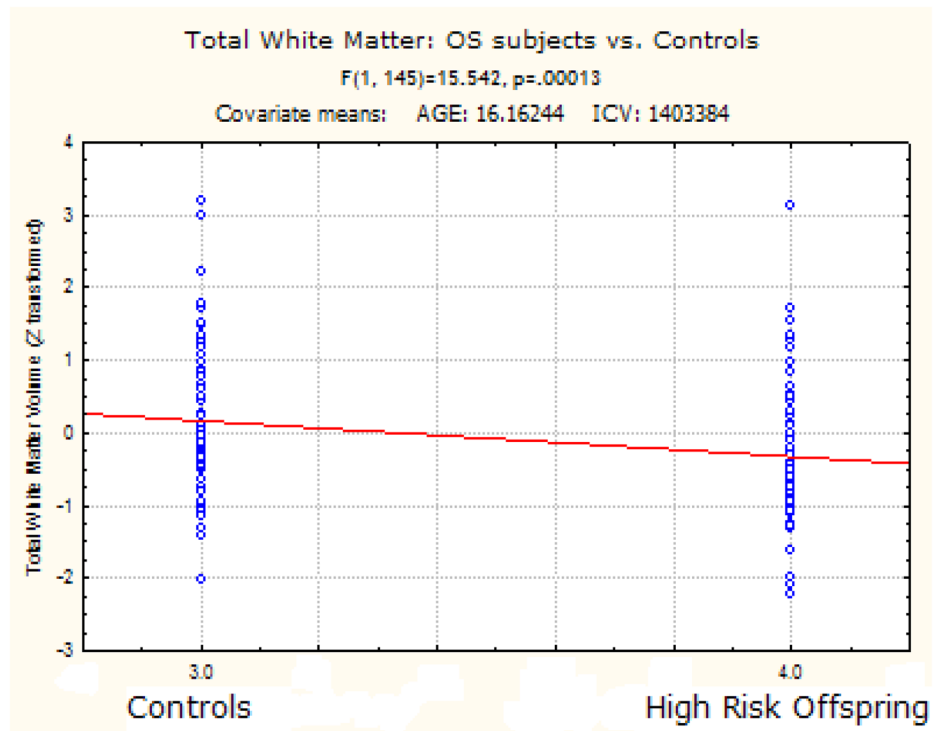


Fig 2.

**Table 1**

## Demographic Information of Subject Populations

Variable	Offspring (N = 65)	Control (N = 80)	<i>P</i> <sup>b</sup>
	<i>M</i> ( <i>SD</i> )/ <i>N</i> (%)	<i>M</i> ( <i>SD</i> )/ <i>N</i> (%)	
Age	16.30 (3.40)	16.62 (3.65)	0.60
Male	38 (54%)	24 (38%)	0.08
Caucasian	34 (49%)	41 (65%)	0.08
Education (yrs)	9.72 (3.33)	10.18 (3.29)	0.43

<sup>b</sup>Fisher's exact test or independent *t*-test, two-tailed, for significant differences between OS and HC participants.

Table 2

LEFT HEMISPHERE – ALL Regional White Matter

LOBES	GYRI	Mean Volume in mm <sup>3</sup>				DF	F	P	Partial Eta Squared
		Controls		Offspring					
		Mean	SD	Mean	SD				
Frontal	Lateral Orbitofrontal	7162.1	1142.2	6652.9	937.6	145	7.48	<b>0.007</b>	0.05
	Medial Orbitofrontal	3706.6	758.7	3388.2	676.7	145	7.16	<b>0.008</b>	0.05
	Caudal Middle Frontal	6486.1	1180.1	6123.6	1214	145	3.86	<b>0.05</b>	0.02
	Rostral Middle Frontal	12192.2	2297.2	11521.7	2044.2	145	3.18	<b>0.07</b>	0.02
	Pars Orbitalis	862.5	261.7	809.7	222.9	145	1.21	<b>.274</b>	0.01
	Pars Opercularis	3569.3	572.8	3405.9	666.2	145	1.64	<b>.203</b>	0.01
	Pars Triangularis	3491.7	683.9	3314.5	666.7	145	1.38	<b>.281</b>	0.009
	Superior Frontal	17649.3	2588.7	16731.9	2879	145	4.19	<b>0.043</b>	0.02
	Frontal Pole	306.4	168.6	302.5	154.1	145	0.197	<b>0.658</b>	0.001
Temporal	Middle temporal	5358.65	975.2	4931.02	1004.5	145	5.25	<b>0.02</b>	0.03
	Inferior temporal	6138.25	1135.5	5565.35	1263.9	145	8.43	<b>0.004</b>	0.056
	Planum Temporale	1413.43	192.8	1336.97	191.9	145	3.94	<b>0.04</b>	0.026
	Superior Temporal	8026.61	1274	7599.6	1286	145	3.03	<b>0.081</b>	0.021
	Entorhinal	463.05	192.2	401.9	154.4	145	3.19	<b>0.07</b>	0.02
	Fusiform	5796.6	927	5365	931.6	145	6.55	<b>0.012</b>	0.04
	Parahippocampal	1278	353.3	1177.9	244.3	145	3.034	<b>0.084</b>	0.02
	Temporal Pole	568.2	247.4	541.9	173.4	145	0.498	<b>0.491</b>	0.003
	Supramarginal	8637.91	1357.6	7839.29	1338.2	145	13.99	<b>0.001*</b>	0.091
Parietal*	Superior parietal	12723.86	1936.6	11603.38	2027.4	145	13.68	<b>0.001*</b>	0.089
	Angular	9511.95	1662.2	9050.17	1805.9	145	2.57	<b>.111</b>	0.018
	Cuneus	2527.4	530.9	2414.7	651.7	145	.671	<b>.414</b>	0.004
Motor Somatosensory	Pericalcarine	3230.6	728.1	3006.3	767.2	145	2.146	<b>0.145</b>	0.015
	Precuneus	8873.9	1557.8	8409.8	1493.8	145	3.40	<b>0.067</b>	0.02
	Post Central	7901.84	1371.4	7252.89	1160.2	145	9.36	<b>0.003</b>	0.063

LOBES	GYRI	Mean Volume in mm <sup>3</sup>				DF	F	P	Partial Eta Squared
		Controls		Offspring					
		Mean	SD	Mean	SD				
	Pre Central	13247.04	1892.5	12603.02	1874.8	145	2.83	<b>0.09</b>	0.094
	Paracentral	3813	780	3697.1	722	145	0.28	<b>0.6</b>	0.001
Cingulate *	Isthmus cingulate	2681.67	468.8	2519.37	422.4	145	4.27	<b>0.03</b> *	0.032
	Posterior cingulate	3980.04	485.1	3689.91	505.5	145	11.37	<b>0.001</b> *	0.075
	Caudal Anterior Cingulate	2176.74	372.3	2080.2	352.6	145	1.84	<b>.177</b>	0.013
	Rostral Anterior Cingulate	1787.92	361.1	1762.22	335.6	145	.129	<b>.720</b>	0.0009
Occipital	Lateral Occipital	10752.3	1943.1	9726.9	2233.3	145	7.51	<b>0.007</b>	0.05
	Lingual	5208.7	1096	4948.7	1001.5	145	.867	<b>.353</b>	0.006

\* -survived correction for multiple comparisons

Table 3

## RIGHT HEMISPHERE – ALL Regional White Matter

LOBES	GYRI	Mean Volume in mm <sup>3</sup>				DF	F	P	Partial Eta Squared
		Controls		Offspring					
		Mean	SD	Mean	SD				
Frontal	Lateral Orbitofrontal	6907.8	1055	6610.4	864.3	145	2.82	<b>0.09</b>	0.019
	Medial Orbitofrontal	3953.2	677.9	3853.4	683.1	145	.67	<b>0.41</b>	0.004
	Caudal Middle Frontal	5741.1	1019.5	5569.6	1060	145	0.74	<b>.38</b>	0.005
	Rostral Middle Frontal	13049	2571.6	11893.8	2199.1	145	8.56	<b>0.003</b>	0.05
	Pars Orbitalis	1026.8	281	965.9	229.6	145	1.42	<b>0.234</b>	0.01
	Pars Opercularis	3493.7	644.2	3346.9	630.3	145	1.62	<b>0.204</b>	0.01
	Pars Triangularis	3097.3	663.3	2823.2	587.2	145	7.41	<b>0.007</b>	0.05
	Superior Frontal	18001.4	2894	17109.8	2787.8	145	3.6	<b>0.05</b>	0.025
	Frontal Pole	421.4	176	397.1	181.5	145	.743	<b>.390</b>	0.005
	Middle temporal	5972.9	1057.6	5367.5	964.4	145	13.75	<b>0.0003</b>	0.09
	Inferior temporal	5662.8	1188	5123.5	1080.8	145	7.06	<b>0.008</b>	0.04
	Planum Temporale	972.2	143.7	935.8	139.6	145	1.44	<b>0.23</b>	0.003
Temporal	Superior Temporal	7228.4	1083	7030.9	1098.7	145	0.58	<b>0.44</b>	0.004
	Entorhinal	551.4	219.6	532.2	174.1	145	.449	<b>.504</b>	0.003
	Fusiform	5601.5	791.2	5215.5	1077.2	145	3.68	<b>0.057</b>	0.02
	Parahippocampal	1347	232.6	1271.5	253.4	145	3.11	<b>0.08</b>	0.02
	Temporal Pole	647.4	223.2	611	166.8	145	1.30	<b>.256</b>	0.009
	Supramarginal	8401.2	1321.4	7778.6	1353.6	145	13.97	<b>0.0002</b>	0.06
	Superior parietal	12076.1	1812.9	11444.5	2108.8	145	3.67	<b>0.05</b>	0.02
	Angular	11891.7	2160.6	10638.9	2207.8	145	9.85	<b>0.002</b>	0.09
	Cuneus	2553.8	630.5	2369.9	606.6	145	2.04	<b>.155</b>	0.014
	Pericalcarine	3218	868.3	3071.8	865.2	145	.412	<b>.522</b>	0.002
	Precuneus	8979.6	1561	8514.6	1603.6	145	2.93	<b>0.089</b>	0.02
	Post Central	7435.4	1106.9	7007.7	1228.2	145	4.87	<b>0.02</b>	0.03
Pre Central	13649.1	2194.3	13119.9	1999.6	145	0.84	<b>0.361</b>	0.006	
Motor Somatosensory									

LOBES	GYRI	Mean Volume in mm <sup>3</sup>				DF	F	P	Partial Eta Squared
		Controls		Offspring					
		Mean	SD	Mean	SD				
Cingulate	Paracentral	4854.2	914.8	4731	794.5	145	.35	.554	0.002
	Isthmus cingulate	2584.3	405.9	2415.1	413	145	6.27	0.013	0.04
	Posterior cingulate	3884.3	576.6	3762.4	475.8	145	1.28	0.259	0.009
	Caudal Anterior Cingulate	2310.7	336	2258	425.9	145	.311	.578	0.002
	Rostral Anterior Cingulate	1388.5	301.5	1372.2	344.4	145	.100	0.753	0.0007
Occipital	Lateral Occipital	10355.3	1964.9	9692.2	1953.7	145	2.70	.102	0.01
	Lingual	4938.7	1052.4	4508.4	956.1	145	3.93	0.049	0.02