

Tryptophan Metabolism and White Matter Integrity in Schizophrenia

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Schizophrenia is associated with abnormalities in the structure and functioning of white matter, but the underlying neuropathology is unclear. We hypothesized that increased tryptophan degradation in the kynurenine pathway could be associated with white matter microstructure and biochemistry, potentially contributing to white matter abnormalities in schizophrenia. To test this, fasting plasma samples were obtained from 37 schizophrenia patients and 38 healthy controls and levels of total tryptophan and its metabolite kynurenine were assessed. The ratio of kynurenine to tryptophan was used as an index of tryptophan catabolic activity in this pathway. White matter structure and function were assessed by diffusion tensor imaging (DTI) and ¹H magnetic resonance spectroscopy (MRS). Tryptophan levels were significantly lower ($p < 0.001$), and kynurenine/tryptophan ratios were correspondingly higher ($p = 0.018$) in patients compared with controls. In patients, lower plasma tryptophan levels corresponded to lower structural integrity (DTI fractional anisotropy) ($r = 0.347$, $p = 0.038$). In both patients and controls, the kynurenine/tryptophan ratio was inversely correlated with frontal white matter glutamate level ($r = -0.391$ and -0.350 respectively, $p = 0.024$ and 0.036). These results provide initial evidence implicating abnormal tryptophan/kynurenine pathway activity in changes to white matter integrity and white matter glutamate in schizophrenia.

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INTRODUCTION

Decreased plasma concentration of tryptophan was initially identified in antipsychotic-free schizophrenia patients over 40 years ago (Manowitz *et al*, 1973). Several subsequent studies have also found reduced plasma tryptophan in schizophrenia patients, although findings of no significant decrease have also been reported (summarized in Table 1). Because over 90% of tryptophan is metabolized through the kynurenine pathway (Schwarcz *et al*, 2012) controlled by rate-limiting enzymes indoleamine 2,3-dioxygenase (IDO) and tryptophan 2,3-dioxygenase (TDO), the reduction of tryptophan in schizophrenia has been attributed to an increased conversion from tryptophan to kynurenine metabolites, as supported by higher kynurenine/tryptophan (KYN/TRP) ratio (Barry *et al*, 2009; Schwieler *et al*, 2015). However, an unanswered question is how abnormal tryptophan metabolism may contribute to brain abnormalities identified in patients with schizophrenia.

IDO regulation of kynurenine pathway metabolism is thought to modulate the activation of myelin-specific T cells (Platten *et al*, 2005). These activated T cells have been shown to generate proinflammatory cytokines that can directly contribute to demyelination and indirectly potentiate antibodies against myelin proteins and also increase perivascular infiltrates (Vass *et al*, 1992; Kroenke and Segal, 2011). Animal studies have established that abnormal elevation of kynurenine in early development may result in decreased dendritic spine density and glutamatergic abnormalities in adults, as well as cognitive deficits resembling those observed in schizophrenia (Pocivavsek *et al*, 2012; Pershing *et al*, 2015). These preclinical data suggest a possible impact of abnormal kynurenine pathway metabolism on cerebral white matter structure and/or biochemistry. A previous study in psychiatrically healthy elderly individuals found that low plasma tryptophan levels were associated with more deep white matter lesions (Yao *et al*, 1999), further supporting a possible link between abnormal tryptophan metabolism that could be due to intake, availability, and/or degradation, and changes in the white matter.

Indeed, accumulating evidence indicates that white matter abnormality is one of the more consistent characteristics of schizophrenia pathophysiology (Kochunov and Hong, 2014;

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Table 1 Summary of Previous Findings Regarding Peripheral Levels of Tryptophan in Schizophrenia

	Control		SZ patient		Effect size	Key findings
	N	TRP ($\mu\text{mol/l}$) (mean \pm SD)	N	TRP ($\mu\text{mol/l}$) (mean \pm SD)		
Barry <i>et al</i> (2009)	36	53.0 \pm 12.6	34	48.8 \pm 12.7	−0.33	↔ Nonfasting plasma TRP in medicated SZ
Carl <i>et al</i> (1992)	13	76.5 \pm 26.6	13	91.2 \pm 12.3	0.71	↔ Plasma TRP in medicated SZ
Fukushima <i>et al</i> (2014)	27	81.3 \pm 17.5	25	101 \pm 22.4	0.99	↑ Nonfasting serum TRP in medicated SZ
Kim <i>et al</i> (2009)	174	69.7 \pm 13.1	71	63.5 \pm 17.2	−0.43	↓ Plasma TRP in unmedicated SZ, TRP tended to normalize with treatment
Koike <i>et al</i> (2014)	38	50.4 \pm 9.1	30	49.6 \pm 10.1	−0.08	↔ Plasma TRP in medicated SZ
Lee <i>et al</i> (2011)	55	40.7 \pm 9.8	159	36.0 \pm 9.9	−0.48	↓ Plasma TRP in treatment-resistant SZ compared with both nontreatment-resistant SZ and HC
Manowitz <i>et al</i> (1973)	15	58.8 \pm 9.5	53	42.1 \pm 10.7	−1.60	↓ Plasma TRP in unmedicated SZ, TRP tended to normalize with medication
Potkin <i>et al</i> , (1983)	17	68.1 \pm 19.6	22	60.2 \pm 13.7	−0.48	↔ Plasma TRP in medicated SZ
Rao <i>et al</i> (1990)	90	85 \pm 26	110	68.2 \pm 22.1	−0.70	↓ Serum TRP in unmedicated SZ
Tortorella <i>et al</i> (2001)	11	291 \pm 102	11	126 \pm 74.3	−1.84	↓ Serum TRP in unmedicated SZ, TRP tended to normalize with clozapine
van der Heijden <i>et al</i> (2005)	73	46.0 \pm 6.1	66	45.5 \pm 8.4	−0.07	↔ Plasma TRP in unmedicated SZ, no change with treatment
Xuan <i>et al</i> (2011)	18	N/A	18	N/A	N/A	↓ Serum TRP in unmedicated SZ; levels tended to normalize in responders to treatment
Yao <i>et al</i> (2010)	30	37.3 ^a	25	31.5 ^a	N/A	↓ Plasma TRP in unmedicated SZ (trend-level), no change with treatment
Total/average ^b	579	66.9	619	53.8	−0.42 ^c	

Abbreviations: HC, healthy control; SZ, schizophrenia; TRP, tryptophan.

^aData were not reported for SD.

^bTotals and averages exclude study by Xuan *et al* (2011).

^cWeighted effect size based on sample size.

Davis *et al*, 2003) and likely contributes to symptoms of schizophrenia through impaired synchronization of brain networks (Bartzokis, 2002). Much of the recent evidence for white matter abnormalities comes from use of diffusion tensor imaging (DTI) that measures the directional diffusion of water. Fractional anisotropy (FA) is a commonly used metric obtained with DTI that is reduced in schizophrenia (Ellison-Wright and Bullmore, 2009) and sensitive to aging, inflammation, and demyelination (Werring *et al*, 1999; Song *et al*, 2005; Kochunov *et al*, 2008). One of our aims is to test the hypothesis that peripheral levels of tryptophan and kynurenine are related to white matter neurobiology in schizophrenia as indexed by cerebral white matter FA.

We also acquired proton magnetic resonance spectroscopy (¹H-MRS) from a frontal white matter region to investigate a potential link between tryptophan metabolism and white matter glutamate levels. Previous MRS studies of glutamate in the white matter in schizophrenia found higher glutamate levels in acute psychosis (Ota *et al*, 2012) and elderly schizophrenia patients (Chang *et al*, 2007). In rodents, systemic administration of kynurenine significantly decreased extracellular glutamate levels in the prefrontal cortex (Wu *et al*, 2010; Konradsson-Geuken *et al*, 2010). However, to our knowledge no studies have examined potential effects of tryptophan/kynurenine on white matter glutamate in schizophrenia. Based on the preclinical data in the prefrontal cortex, we tested the hypothesis that peripheral

levels of tryptophan and kynurenine are related to white matter glutamate levels in schizophrenia patients.

MATERIALS AND METHODS

Participants

Patients ($n=37$, age range 20–58 years; 30 with diagnosis of schizophrenia and 7 with diagnosis of schizoaffective disorder) were recruited from the outpatient clinics at the Maryland Psychiatric Research Center and the neighboring mental health clinics. Healthy controls ($n=38$, age range 20–61 years) were recruited through media advertisements. Demographics of the sample are reported in Table 2. Diagnoses were confirmed with the Structured Clinical Interview (SCID) for DSM-IV in all participants. Major medical and neurological illnesses, history of head injury with cognitive sequelae, and mental retardation were exclusionary. Patients and controls with substance dependence within the past 6 months or current substance abuse (except nicotine) were excluded. Except for four medication-free participants, all schizophrenia patients were on antipsychotic medications, including 8 taking clozapine, 4 taking typical antipsychotics, 17 taking atypicals, and 4 taking a combination of typical and atypical antipsychotics. In addition, 15 of the patients were on an antidepressant medication at the time of study. Controls had no current DSM-IV Axis I diagnoses and no family history of psychosis in the prior two generations.

Table 2 Demographic, Clinical Characteristics, and Group Differences in Plasma Metabolites

	Healthy control (n = 38)	Schizophrenia (n = 37)	Test statistic	P-value
Age (years)	39.7	39.2	t(df = 73) = 0.16	0.87
BMI	26.2	29.5	t(df = 73) = 2.64	0.01
Smoker/nonsmoker	8/30	20/17	$\chi^2 = 8.73$	0.003
Male/female	25/13	28/9	$\chi^2 = 0.88$	0.35
Tryptophan ($\mu\text{mol/l}$)	70.7 \pm 15.2	57.1 \pm 15.2	F(1, 73) = 14.99	<0.001
Kynurenine ($\mu\text{mol/l}$)	1.73 \pm 0.42	1.64 \pm 0.57	F(1, 73) = 0.68	0.41
KYN/TRP ($\mu\text{mol/mmol}$)	24.9 \pm 5.3	29.8 \pm 11.2	F(1, 73) = 5.85	0.018

Abbreviations: BMI, body mass index; KYN/TRP, kynurenine/tryptophan ratio.

Participants gave written informed consent. This study was approved by the University of Maryland IRB.

Clinical Assessments

Overall psychiatric symptoms were assessed with the mean of the 20 item Brief Psychiatric Rating Scale (BPRS); subscales for positive psychosis symptoms and anxiety/depression symptoms were also generated in order to test specific symptom domains (Overall and Gorham, 1962). To explore how cognitive deficits might be related to altered levels of tryptophan and kynurenine metabolites, participants were tested with the Digit Symbol Coding task of the WAIS-3 (Wechsler, 1997) and the Digit Sequencing task from the Brief Assessment of Cognition in Schizophrenia (Keefe et al, 2004) to assess processing speed and working memory, respectively.

Biochemistry

Participants were instructed to fast overnight. Whole blood was collected between 0900 and 1030 h in EDTA-containing tubes (Vacutainer) that were immediately centrifuged at 2500 r.p.m. for 10 min. Plasma was then aliquoted into separate tubes and stored at -80°C until assay. Total tryptophan (free plus protein bound) and kynurenine were measured by reverse-phase high-performance liquid chromatography, as previously described (Widner et al, 1997; Laich et al, 2002), using 3-nitro-L-tyrosine as an internal standard. Tryptophan was detected using an excitation wavelength of 266 nm and an emission wavelength of 366 nm. Kynurenine was detected by UV absorption at 360 nm. Retention times were 4.3 min for tryptophan and 2.3 min for kynurenine. Coefficients of variance for the assay were 1.7% for tryptophan and 4.2% for kynurenine.

Diffusion Tensor Imaging

All imaging was performed at the University of Maryland Center for Brain Imaging Research using a Siemens 3T TRIO MRI (Erlangen, Germany) system equipped with a 32-channel phase array head coil. The DTI data were collected using a single-shot, echo-planar, single refocusing spin-echo, T2-weighted sequence with a spatial resolution of $1.7 \times 1.7 \times 3.0$ mm. The sequence parameters were: TE/TR = 87/8000 ms, FOV = 200 mm, axial slice orientation with 50 slices and no gaps, five b = 0 images, and 64 isotropically

distributed diffusion-weighted directions with b = 700 s/mm². These parameters maximized the contrast to noise ratio for FA measurements (Kochunov et al, 2012). A tract-based spatial statistics (TBSS) method was used for tract-based analysis of diffusion anisotropy (Smith et al, 2006). FA images were created by fitting the diffusion tensor to the motion and eddy current diffusion data. RMSDIFF (Smith et al, 2006) was used to estimate the root mean square movement distance between diffusion sensitized and b = 0 images. All data passed QA control of <3 mm accumulated motion during the scan. In the next step, all FA images were globally spatially normalized to the Johns Hopkins University atlas (Wakana et al, 2004) and then nonlinearly aligned to a groupwise, minimal-deformation target (MDT) brain using the FLIRT method (Kochunov et al, 2001; Smith et al, 2006). Next, individual FA images were averaged to produce a group-average anisotropy image. This image was used to create a groupwise skeleton of white matter tracts. Finally, FA images were thresholded at FA = 0.20 level to eliminate nonwhite matter voxels, and FA values were projected onto the groupwise skeleton of white matter structures. This step accounts for residual misalignment among individual white matter tracts. FA values were assigned to each point along a skeleton using the peak value found within a designated range perpendicular to the skeleton. Whole-brain tract-averaged FA was used as the primary measure for statistical analysis, based on the rationale that if there was a relationship between peripheral measures of tryptophan metabolism and white matter structure, the effect would be global rather than specific to any particular tract. However, as the MRS voxel was in frontal white matter (see below), we also conducted exploratory analyses using FA for corona radiata and genu of the corpus callosum. One patient did not complete DTI.

White Matter MRS

A spectroscopic voxel was placed in the white matter of the forceps minor area of the left hemisphere, corresponding to the left anterior corona radiata (ACR) under the left prefrontal cortex, avoiding CSF and gray matter. The prefrontal white matter MRS data were previously reported in the context of examining the relationship of myo-inositol and FA in the white matter in schizophrenia (Chiappelli et al, 2015); MRS data from all participants involved in the previous study are included here, with no new MRS data added to this analysis, although biochemical data from one

additional schizophrenia patient who did not complete MRS are included here. Single-voxel PRESS localization was utilized with the following parameters: TR/TE = 2000/30 ms, VOI $\sim 3.4 \text{ cm}^3$, NEX = 256, 2048 complex points, 1.2 kHz spectral width, and total scan time $\sim 12 \text{ min}$. A water reference (NEX = 8) was collected and utilized for phasing and eddy current correction. A basis set was simulated using the GAVA software package (Soher *et al*, 2007) that was modified to yield a Lorentzian lineshape instead of its default Gaussian lineshape as previously reported (Rowland *et al*, 2015). This basis set was imported into LCModel, an automated curve fitting software package (Provencher, 1993) for metabolite quantification. Metabolites were corrected for the proportion of the gray matter, white matter, and cerebrospinal fluid (CSF) within the spectroscopic voxel using in-house Matlab code (Gasparovic *et al*, 2006). All metabolite concentrations were relative to the water reference and are reported in institutional units (IUs). The exclusion criteria for these data were: (1) SNR reported by LCModel was ≤ 10 ; (2) FWHM reported by LCModel was ≥ 0.09 ; and (3) metabolite fits with %SD (estimated standard deviations also called Cramer–Rao lower bounds) above > 20 . All spectra were of good quality with a mean FWHM of 0.045 for controls and 0.048 for patients ($t = 1.01$, $p = 0.318$) and a mean SNR of 27.7 for controls and 26.0 for patients ($t = 1.31$, $p = 0.196$). Five metabolites were consistently identified with %SD < 20 and were used in statistical analyses: glutamate (mean %SD = 7.3 (control), 7.4 (patient)), NAA plus NAAG (tNAA; mean %SD = 2.7 (control), 3.1 (patient)), creatine plus phosphocreatine (tCr; mean %SD = 2.9 (control), 3.1 (patient)), glycerophosphocholine plus phosphocholine (tCho; mean %SD = 4.2 (control), 4.3 (patient)), and *myo*-inositol (mean %SD = 5.3 (control), 5.1 (patient)).

Statistical Analyses

Group differences were examined with one-way ANOVA, t -test, and χ^2 , as indicated. Further ANOVA tests were performed to test the influence of smoking and body mass index (BMI) on group differences, as the groups were significantly different for these variables (see below). Correlations between variables were examined with Pearson's correlation coefficients, except for variables that deviated from a normal distribution as determined by Kolmogorov–Smirnov tests, in which case Spearman's rank correlation coefficient was used. Correlational analyses were performed on the entire sample first, and then repeated for patient and control groups separately, except for when both variables in the analyses were significantly different between groups. The α was set at 0.05 and exact p -values are reported. Bonferroni corrections were only applied for whole-sample analyses with white matter metabolites because of the large number of tests. All tests were two tailed.

RESULTS

Demographics

Patient and control groups were balanced for age and gender (Table 2). However, patients had significantly greater BMI

than controls ($p = 0.01$) and were more likely to smoke ($p = 0.003$).

Group Differences

One-way ANOVAs comparing plasma levels of tryptophan, kynurenine, and KYN/TRP ratio between patients and controls found that patients had significantly lower levels of tryptophan ($F(1, 73) = 14.99$, $p < 0.001$; Figure 1) but not kynurenine ($F(1, 73) = 0.68$, $p = 0.41$); KYN/TRP ratio was higher in patients compared with controls ($F(1, 73) = 5.85$, $p = 0.018$; Table 2). Adding smoking status and BMI as covariates, tryptophan remained significantly lower in patients compared with controls ($F(1, 70) = 9.98$, $p = 0.002$). Tryptophan levels did not differ between smokers and nonsmokers ($F(1, 70) = 0.84$, $p = 0.35$), nor was there a smoking \times diagnosis interaction ($F(1, 70) = 1.34$, $p = 0.27$). Females had lower levels of tryptophan than males ($F(1, 71) = 5.57$, $p = 0.021$) but there was no gender \times diagnosis interaction ($F(1, 71) = 0.05$, $p = 0.83$). There was a significant inverse correlation between tryptophan and age in the combined sample ($r = -0.254$, $p = 0.028$) but not separately in either group (both $p > 0.05$). Tryptophan was not related to BMI in either controls ($r = -0.270$, $p = 0.10$) or patients ($r = 0.143$, $p = 0.40$). Based on these results, subsequent analyses focused on testing the potential association of the significantly low tryptophan level and higher KYN/TRP ratios with brain imaging markers.

White Matter Fractional Anisotropy

Whole-brain tract-averaged FA in patients was significantly reduced in patients compared with controls ($t(72) = 2.05$, $p = 0.044$). Plasma fasting tryptophan level was significantly correlated with whole-brain tract-averaged FA in patients ($r = 0.347$, $p = 0.038$, $n = 36$; see Figure 2) but not in controls ($r = 0.182$, $p = 0.27$, $n = 38$). Tryptophan level remained positively related to FA in patients after adding age, smoking status, and gender as covariates in linear regression ($\beta = 0.320$, $p = 0.009$). Tryptophan levels were also positively correlated with FA of corona radiata ($r = 0.330$, $p = 0.050$) and genu of corpus callosum ($r = 0.346$, $p = 0.039$) in

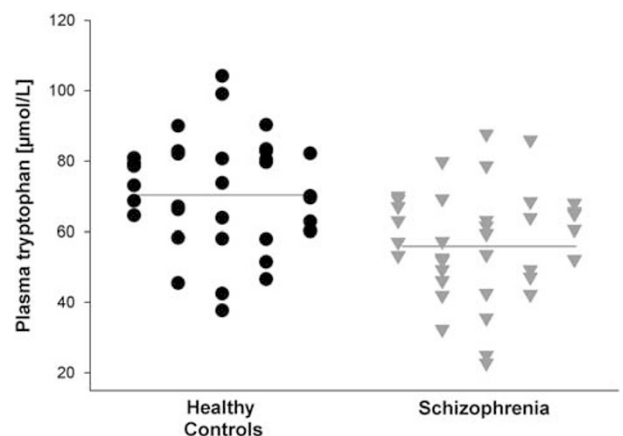


Figure 1 Scatterplot showing difference in plasma tryptophan levels.

patients. KYN/TRP ratio was not significantly correlated with FA in either controls or patients (all $p > 0.05$).

White Matter Neurochemistry

As previously reported (Chiappelli et al, 2015), there were no significant differences between patients and controls in levels of glutamate ($p = 0.81$) or the other metabolites including tCho, tNAA, tCr, or *myo*-inositol in this frontal white matter location (all $p > 0.14$). In the combined sample of patients and controls, with Bonferroni correction for 15 (3 peripheral metabolites tryptophan, kynurenine, and KYN/TRP \times 5 MRS metabolites) analyses (corrected α is $p < 0.0033$), the only significant correlation was a negative correlation between KYN/TRP and frontal white matter glutamate ($r = -0.359$, $p = 0.002$). Examination of this relationship with linear regression found that the KYN/TRP still predicted white matter glutamate after controlling for age, gender, and smoking status ($\beta = -0.248$, $p = 0.025$). Further analysis showed that the negative relationship between KYN/TRP and white matter glutamate was found in both healthy controls ($r = -0.350$, $p = 0.036$) and schizophrenia patients ($r = -0.391$, $p = 0.024$; Figure 3), suggesting that higher

tryptophan to kynurenine conversion was associated with lower glutamate level at this prefrontal white matter region, consistent with animal data found in the prefrontal cortex (Wu et al, 2010; Konradsson-Geuken et al, 2010).

Association with Clinical Variables

Among patients, tryptophan levels were not significantly related to chlorpromazine dose equivalent of antipsychotic medication ($\rho = 0.206$, $p = 0.25$). KYN/TRP ratios were also not related to chlorpromazine dose equivalent ($\rho = -0.022$, $p = 0.90$). Tryptophan levels were not significantly different between patients not on medication at the time of study ($M = 57.9$ (SD = 11.3) $\mu\text{mol/L}$, $n = 4$) and patients on antipsychotic medication ($M = 57.0$ (SD = 15.7), $n = 33$; $t = 0.11$, $p = 0.91$); nor did unmedicated patients differ in KYN/TRP ratio ($t = 0.34$, $p = 0.74$). Tryptophan levels did not differ between patients taking an antidepressant and patients not taking an antidepressant medication ($t(35) = 0.26$, $p = 0.79$). KYN/TRP ratio also was not significantly different based on antidepressant use ($t(35) = 0.72$, $p = 0.48$). There were no significant differences between patients taking clozapine and those not treated with clozapine for tryptophan ($t(35) = 0.20$, $p = 0.84$) or KYN/TRP levels ($t(35) = 0.03$, $p = 0.98$). Tryptophan was not significantly associated with mean BPRS scores ($r = -0.229$, $p = 0.17$) or mean BPRS psychosis subscale scores in the patients ($r = -0.071$, $p = 0.69$). KYN/TRP was also not significantly associated with BPRS scores ($r = 0.20$, $p = 0.23$) or BPRS psychosis subscale scores ($r = -0.004$, $p = 0.98$). There was a nonsignificant trend for lower tryptophan to be associated with higher levels of BPRS anxiety/depression subscale scores in patients ($r = -0.329$, $p = 0.06$). Tryptophan or KYN/TRP were not significantly associated with working memory or processing speed in patients (all $p > 0.31$).

DISCUSSION

This study employed neuroimaging assessments to explore the relationship of tryptophan metabolism to markers of white matter health in schizophrenia. Collectively, our results

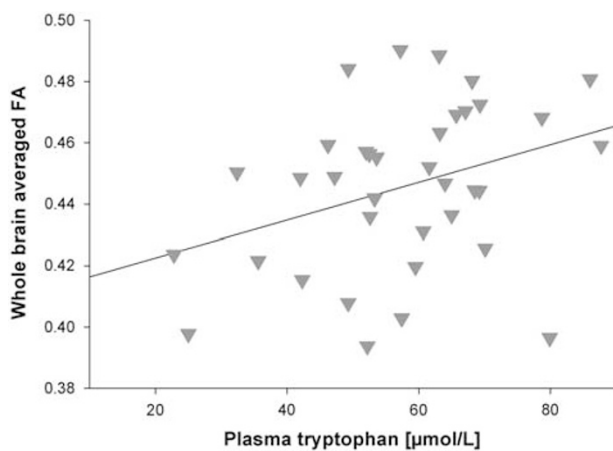


Figure 2 Scatterplot displaying relationship of tryptophan to whole brain-averaged FA in patients.

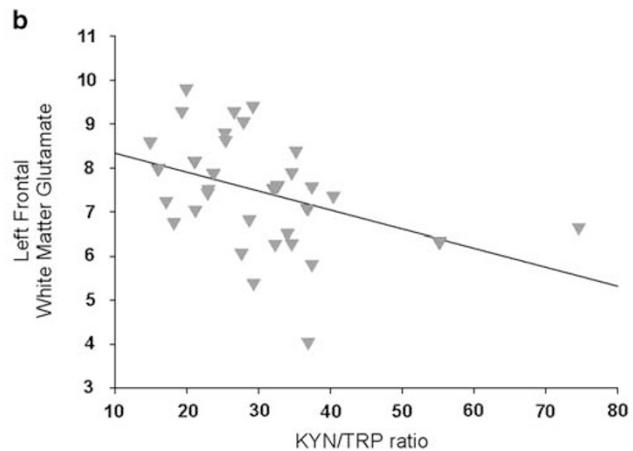
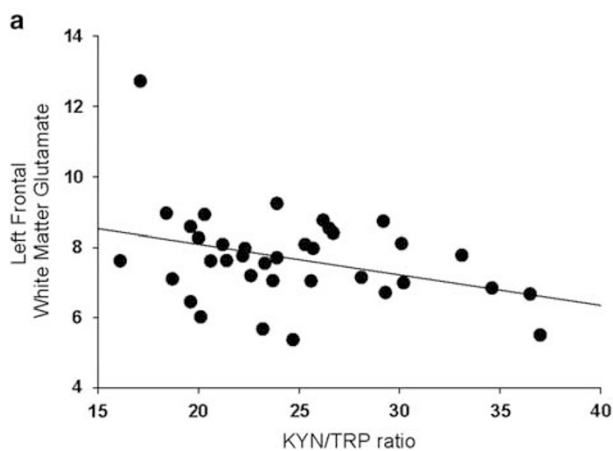


Figure 3 Scatterplots showing relationship between plasma KYN/TRP ratio ($\mu\text{mol/mmol}$) and glutamate levels in left frontal anterior corona radiata (levels reported in IUs) in (a) healthy controls and (b) schizophrenia patients.

suggest a reduced tryptophan level in schizophrenia, and that this may indicate an increased degradation as suggested by increased KYN/TRP ratio. Altered tryptophan metabolism appears to be associated with several white matter imaging markers: lower tryptophan was associated with lower fractional anisotropy of white matter, a nonspecific marker of white matter microstructure, and a higher KYN/TRP ratio was associated with lower glutamate levels in the frontal white matter.

Although the proper interpretation of FA regarding microanatomy and physiology remains disputed (Jones *et al*, 2013), this parameter is robust for clinical significance and reliability. Aging is associated with both reduced FA (Kochunov *et al*, 2008) and decreased plasma concentration of tryptophan (Capuron *et al*, 2011), but in this study tryptophan levels predicted FA in schizophrenia patients even after controlling for the age effect. The causes of white matter abnormalities in schizophrenia remain unclear. One of the possibilities explaining the link between tryptophan level and FA may in part be related to evidence of chronic inflammation in schizophrenia (Miller *et al*, 2011; Upthegrove *et al*, 2014). Proinflammatory cytokines, interferon- γ in particular, upregulate IDO, one of the rate-limiting enzymes involved in conversion of tryptophan to kynurenine (Schwarcz *et al*, 2012). Therefore, inflammatory mediators can upregulate the metabolism of tryptophan to kynurenine, or kynurenine to kynurenic acid and quinolinic acid by glial cells within the brain (Schwarcz *et al*, 2012). Depending on their relative concentration, these metabolites can protect against or exacerbate excitotoxicity, pathological processes that can directly damage myelin sheath and axons. Additional support to this hypothesis is that IDO and kynurenine pathway metabolites regulate the activation of myelin-specific T cells (Platten *et al*, 2005) that can directly contribute to demyelination and indirectly potentiate antibodies against myelin proteins (Kroenke and Segal, 2011; Vass *et al*, 1992). However, in the brain TDO mRNA and protein, but not IDO, is upregulated in schizophrenia (Miller *et al*, 2004; Miller *et al*, 2006). Furthermore, given that fasting tryptophan levels, but not KYN/TRP ratios, were correlated with whole-brain tract-averaged FA in patients, it is also worth considering that this relationship is related to reduced availability of tryptophan. Tryptophan levels are not only influenced by dietary intake, but also by the composition of the gut microbiota, as specific bacteria within the gut can either synthesize tryptophan *de novo*, or metabolize tryptophan, thus controlling its availability to the host (O'Mahony *et al*, 2015). In addition, evidence from preclinical studies indicates that tryptophan and tryptophan metabolites (both bacterially derived and host derived) have immunomodulatory effects (Zelante *et al*, 2013). Thus, although speculative, the findings reported here may be due to a chronic mild inflammatory state in schizophrenia that reduces tryptophan availability and/or promotes tryptophan catabolism to kynurenine metabolites toxic to oligodendrocytes and/or white matter (Sundaram *et al*, 2014).

The relationship between tryptophan levels and mood symptoms may also be mediated through stress reactivity, as stress hormones regulate TDO, another rate-limiting enzyme for tryptophan/kynurenine conversion. In previous work we have found that schizophrenia patients with poor stress tolerance exhibit a robust increase in salivary kynurenic acid

in response to a mild psychological stressor (Chiappelli *et al*, 2014). Furthermore, these patients with poor stress tolerance exhibit a more prolonged cortisol response to the stressor that correlates with lower white matter FA (Nugent *et al*, 2015). Thus, enhanced stress reactivity represents another possible mechanism underlying the observed relationship between low tryptophan levels and impaired white matter structure in schizophrenia patients.

Evidence of altered activity of the kynurenine pathway in schizophrenia includes findings of increased concentration of kynurenine and kynurenic acid in brain tissue (Schwarcz *et al*, 2001; Miller *et al*, 2006; Sathyaikumar *et al*, 2011) and in cerebrospinal fluid of schizophrenia patients (Nilsson *et al*, 2005; Linderholm *et al*, 2012). Kynurenine and kynurenic acid administration to the prefrontal cortex significantly decrease extracellular glutamate levels in rodents (Wu *et al*, 2010; Konradsson-Geuken *et al*, 2010). Here, we found that in both patients and controls, KYN/TRP ratios were inversely correlated with frontal white matter glutamate. We interpret this finding as indicating that higher conversion from tryptophan to kynurenine is associated with lower glutamate in the white matter, parallel to the rodent observation in prefrontal gray matter. Recent work has also uncovered evidence of axon–glial communication within white matter that appears to be based on glutamatergic signaling and may represent a mechanism for activity-dependent modulation of myelination (Kukley *et al*, 2007; Ziskin *et al*, 2007). Although the levels of white matter glutamate measured with MRS cannot be assumed to directly reflect axon–glial signaling, and cannot distinguish between extracellular *vs* intracellular levels, the set of findings reported here raise the possibility that kynurenine pathway activity may have a downstream influence on glutamate-dependent white matter activity.

As this was a cross-sectional study, we cannot draw causal conclusions regarding the pathophysiological mechanism that underlies the relationship between tryptophan metabolism and white matter abnormalities. We did not have data to approximate premorbid IQ of the patients in this study that could provide clues regarding the neurodevelopmental context of our findings. Most patients included in this study were on antipsychotic medications, and the potential confounding effect of these medications cannot be ruled out. However, we found that the chlorpromazine equivalent dose was not related to plasma tryptophan in the patients. This is consistent with an *in vitro* study that found no effect of antipsychotics on tryptophan metabolism in cultured peripheral blood mononuclear cells from schizophrenia patients (Krause *et al*, 2013). Although we excluded participants identified by diagnostic interview as having current substance abuse disorders, we did not routinely perform drug screens before the study, leaving open the possibility that recent illicit drug use could have altered laboratory or imaging results for some participants. Another limitation of this study was that only total tryptophan was measured, and that kynurenine was the only tryptophan metabolite measured; the KYN/TRP ratio may be only a crude index of IDO/TDO activity (Badawy, 2015). Furthermore, ~5 to 10% of tryptophan is metabolized to serotonin (Schwarcz *et al*, 2012), a pathway that is not examined in the current study. Indeed, low levels of tryptophan have been associated with mood disturbances and our data revealed a trend toward low tryptophan being associated with greater anxiety and

depression in schizophrenia patients. We must also consider the limitations posed by only having peripheral and not central measures of tryptophan and kynurenine levels. Tryptophan and kynurenine both cross the blood–brain barrier; the majority of kynurenine in the brain is derived from blood, especially in conditions of systemic inflammation (Kita *et al*, 2002; Schwarcz *et al*, 2012). However, peripheral levels of tryptophan and kynurenine are not necessarily informative about the activity of IDO/TDO activity in the brain, nor do they provide information on the relative balance of kynurenine *versus* kynurenic acid produced by glial cells. Further studies on the relationship of tryptophan metabolism and brain structure will require parallel animal studies to provide mechanistic explanations.

Although the results of previous studies examining levels of tryptophan in schizophrenia patients have been variable, the weighted average of effect size across these studies (-0.42 ; see Table 1) suggests that reduced tryptophan in the peripheral blood in schizophrenia is a replicable finding with modest effect size. Levels of tryptophan can be affected by food intake, exercise, stress, gut microbiota, and some medications (Badawy, 2015). We attempted to limit the impact of some of these factors by collecting blood samples in individuals who had been fasting overnight and who were instructed to avoid strenuous exercise the day before blood collection, although we did not precisely regulate intake, stress, or activity levels. Finally, the exploration of correlations among numerous imaging and peripheral markers presents a high risk for type I error and the results must therefore be considered as only initial evidence of a link between increased tryptophan metabolism and white matter impairment in schizophrenia patients.

In conclusion, this study found preliminary evidence that altered tryptophan metabolism is related to white matter integrity in schizophrenia. Although the data in this study are not sufficient to pinpoint the underlying mechanism of this relationship, it is possible that low tryptophan reflects both decreased availability and increased catabolism due to inflammation; and inflammation may leave patients vulnerable to white matter damage through neuroinflammatory, neurovascular, and neuroendocrine pathways. Further investigation of the mechanisms underlying this relationship may reveal clinically useful biomarkers and potential treatment targets for schizophrenia.

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