Variability of Structurally Constrained and Unconstrained Functional Connectivity in Schizophrenia

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Abstract: Spatial variation in connectivity is an integral aspect of the brain’s architecture. In the absence of this variability, the brain may act as a single homogenous entity without regional specialization. In this study, we investigate the variability in functional links categorized on the basis of the presence of direct

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**INTRODUCTION**

Optimal function of the human brain relies on the cooperation of constituent brain regions. The degree of cooperative activity between two brain regions at rest, measured using functional connectivity [Friston, 1994], varies greatly across the brain. A complete lack of spatial variability in functional connectivity indicates that the entire brain is either acting as a single homogenous unit without regional specialization, or the constituent brain regions are entirely asynchronous, without integration. In contrast, optimum spatial variability in connectional strength reflects simultaneous integration and segregation across distributed brain regions. This diversity forms the core of the overall topology of the complex functional architecture of human brain [Bullmore and Sporns, 2009; Tononi et al., 1994] altered in neuropsychiatric disorders such as schizophrenia [Alexander-Bloch et al., 2010; Fornito et al., 2012; Rubinov and Bullmore, 2013; van den Heuvel et al., 2013].

In healthy controls, presence of structural connectivity (putative axonal linkage assessed using diffusion tensor imaging) strongly predicts the strength of functional connectivity [Damoiseaux and Greicius, 2009; Honey et al., 2009; Skudlarski et al., 2010; van den Heuvel et al., 2009]. Nevertheless, most of the pairwise functional connections exist in the absence of a direct axonal linkage between two regions [Adachi et al., 2011; Damoiseaux and Greicius, 2009; Honey et al., 2009]. In general, functional links that have a structural basis are stronger and involve anatomically more proximal regions [Honey et al., 2009]. In contrast, functional links between regions that do not share direct axonal linkage appear to be weaker, and involve spatially distant regions [Adachi et al., 2011; Honey et al., 2009]. Functional links in the absence of one-to-one axonal connections may emerge from directed polysynaptic connections or shared inputs/outputs involving a third region [Adachi et al., 2011]. Such indirect, weaker functional links may represent a connectional architecture that is physiologically distinct from the links with a more direct structural basis.

Several studies observe a reduction in overall strength of functional connectivity in schizophrenia [Argyelan et al., 2013; Bassett et al., 2012; Lynall et al., 2010], while both increased [Skudlarski et al., 2010; Whitfield-Gabrieli et al., 2009] and decreased [Bluhm et al., 2007; Liang et al., 2006] connectivity involving different regional connections are noted across the brain [Karbasforoushan and Woodward, 2012; Pettersson-Yeo et al., 2011; Rubinov and Bullmore, 2013]. The presence of both hyper- and hypoconnectivity involving different regional connections [Guo et al., 2013; Skudlarski et al., 2010; Venkataraman et al., 2012; Woodward et al., 2012] indicates a large diversity in the distribution of connectivity across the functional links in schizophrenia. If such a diversification lies at the core of the dynamic pathophysiological process defining the presence of schizophrenia in an individual, then for a randomly chosen pair of brain regions, the connectional strength is likely to be less predictable in a patient compared to a healthy control.

The unpredictability of bivariate functional connectivity across the entire brain can be quantified using an index of entropy based on the principles of information theory [Shannon, 1948]. Functional Connectivity Entropy (FCE) is a measure of randomness of the strength of functional connectivity, across all possible pairwise interregional connections in the brain [Yao et al., 2013]. This is different from...
the unpredictability or complexity across time that has been investigated previously in schizophrenia [Bassett et al., 2012; Fernández et al., 2013]. Higher FCE represents the presence of increased variability in connectional strength and has been observed in association with healthy ageing [Yao et al., 2013]. On the basis of a computational model, altered FCE has been attributed to a reduction in the pool of excitatory neurons [Yao et al., 2013], an observation that is highly relevant to the study of schizophrenia [Anticevic et al., 2013].

Recent observations also suggest that weak (rather than strong) [Bassett et al., 2012] and long-distance (rather than short distance) functional links [Guo et al., 2013] are crucial to discriminate patients with schizophrenia from controls. This raises the possibility that the connectional architecture involving functional links that have no direct structural connectivity may be preferentially affected in schizophrenia.

Here, we tested the hypothesis that the randomness of functional connectivity (measured using FCE) is abnormal in schizophrenia, and this abnormality is specific to functional links with no direct structural connectivity. We also investigated the relationship between FCE and symptom burden in patients. Given its onset around adolescence, various dynamic changes coinciding with brain maturation take place in the initial few years in the course of schizophrenia, giving rise to several unstable and inconsistent neurobiological patterns that stabilized during the later stage of the illness [Pantelis et al., 2009]. In particular, the architecture of functional connectivity in patients is significantly affected by duration of illness. Patients with longer duration of illness show reduced segregation and integration [Liu et al., 2008], and reduced connectivity among core brain hubs [Collin et al., 2013]. In light of these observations, we expected a moderating effect of illness duration on FCE.

**METHODS**

**Participants**

This sample has been previously reported in our earlier study [Palaniyappan et al., 2013]. Thirty-four patients satisfying DSM-IV criteria for schizophrenia (n = 28) or schizoaffective disorder (n = 6) and 32 age, gender and parental socioeconomic status [Rose and Pevalin, 2003] matched healthy controls were included. Patients were recruited from the community-based mental health teams (including Early Intervention in Psychosis teams) in Nottinghamshire and Leicestershire, UK. The diagnosis was made in a clinical consensus meeting in accordance with the procedure of Leckman et al. [1982], using all available information including a review of case files and a standardized clinical interview (SSPI) [Liddle et al., 2002]. All patients were in a stable phase of illness (defined as a change of no more than ten points in their Global Assessment of Function score, assessed 6 weeks prior and immediately prior to study participation). The study was given ethical approval by the National Research Ethics Committee, Derbyshire, UK. All volunteers gave written informed consent. Clinical and demographic characteristics of this sample are presented in Table I.

Patients were interviewed on the same day as the scan by a research psychiatrist (LP and VB) and clinical severity of psychosis over the previous week was assessed on the basis of total SSPI scores. Duration of illness was estimated from the time of reported onset of psychotic symptoms (Criterion A of DSM-IV schizophrenia), on the basis of information from case notes and clinical interview. We divided the patient sample into early stage (<5 years duration) and later-stage (>5 years) illness, based on the clinical notion of “critical period” of psychosis during which interventions can modify outcome [Crumlish et al., 2009; McGorry, 2002; McGorry et al., 2008] and clinical stability is achieved [Bleuler, 1978]. Thirty-two out of 34 patients were receiving antipsychotic treatment at the time of scan. The chlorpromazine equivalent doses were calculated using data presented by Woods [2003] and Chong et al. [2000] (for clozapine). In addition, 25 mg risperidone depot injection every 14 days was considered equivalent to 4 mg oral risperidone per day, in accordance with the recommendation of the British National Formulary [Joint Formulary Committee, 2013].

**MRI ACQUISITION**

All scans were conducted at Sir Peter Mansfield Magnetic Resonance Centre, University of Nottingham using 3T Philips Achieva MR Scanner (Philips Medical Systems, The Netherlands). A structural T1 image was acquired along with a series of diffusion-weighted images and 10-
Three types of structural paths defined using diffusion tractography. (a) whole-brain fiber connection network construction by Diffusion Tensor Imaging data; (b) Heschl gyrus (HES) and Insula (INS) are directly connected with each other through white matter tracts. HES ‒ INS link is a primary structural path. (c) Heschl gyrus and Postcentral gyrus (PoCG) are indirectly linked (discontinuous arrow) through white matter tracts (continuous arrows) that connect HES and PoCG with the Insula. Though there are no direct connections between HES and PoCG, INS acts as a common third region. HES ‒ PoCG link is termed as a secondary structural path. (d) Heschl gyrus and Precentral gyrus (PreCG) are indirectly linked (discontinuous arrow) through white matter tracts (continuous arrows) that connect HES with INS, INS with PoCG and PoCG with PreCG. There are no direct connections or a single common third region between HES and PreCG. HES ‒ PreCG link is termed as a tertiary structural path. [Color figure can be viewed in the online issue, which is available at wileyonlinelibrary.com.]

**DTI PROCESSING**

For DTI images, we first used FMRIB Software Library v5.0 (http://fsl.fmrib.ox.ac.uk/fsl) [Jenkinson et al., 2012] to remove the eddy-current and extract the brain mask from the B0 image. Then, we used TrackVis [Wang et al., 2007] to obtain the fiber images by the deterministic tracking method, with anatomical regions defined using the automated anatomical labeling atlas (AAL) convention on the basis of coregistered T1 image from each subject. This enabled us to determine the presence of streamlines connecting every pair of brain regions. All the processes were performed using the PANDA suite [Cui et al., 2013].

If two brain regions A and B have one or more streamlines directly connecting each other, then this link AB is regarded as a direct or primary structural path. In contrast, if two brain regions X and Y have no direct connections, but share a direct connection with a common third region Z, then the link XY can be regarded as indirect secondary structural path. If two brain regions K and L have neither direct connections, nor a common third region, but are connected to each other indirectly by virtue of being connected a common direct link (for example, K connects directly to A, A connects directly to B, B connects directly to L), then the link KL can be regarded as indirect tertiary structural path. To study the effect of the minimum streamline threshold, we varied the minimum from one to three and repeated the primary analysis (group comparisons). The results, shown in Supporting Information Material 6, indicated that the structural paths were robust across these thresholds.

Thus, for every subject, we identified primary, secondary, and tertiary paths on the basis of the DTI fiber streamlines. An illustration of these paths is shown in Figure 1.
fMRI Processing

Weighted summation of the dual-echo images produced a single set of low-artefact functional images [Posse et al., 1999]. Retrospective physiological correction was then performed [Glover et al., 2000]. The first five volumes of fMRI datasets were discarded, to allow for scanner stabilization. Further processing was then conducted by Statistical Parametric Mapping (SPM8) (http://www.fil.ion.ucl.ac.uk/spm) [Penny et al., 2011] and Data Processing Assistant for Resting-State fMRI [Chao-Gan and Yu-Feng, 2010]. After slice-timing correction and realignment to the middle volume, the functional scans were spatially normalized using the unified segmentation approach to a standard template (Montreal Neurological Institute) and resampled to 3 × 3 × 3 mm³. Data was then smoothed using a Gaussian kernel of 8 mm full-width at half-maximum, detrended, and then passed through a band-pass filter (0.01–0.08 Hz) to reduce low-frequency drift and high-frequency physiological noise.

Nuisance covariates including head motion, global mean signals, white matter signals and cerebrospinal signals were regressed out from the timeseries. Regional time series were extracted in each of the 90 AAL [Tzourio-Mazoyer et al., 2002] based brain regions by averaging the signals of all voxels within a region. The names of the 90 brain regions are listed in Supporting Information Table ST1, along with further details of fMRI preprocessing in SM2, SF1 and the characteristics of excluded subjects in SM3.

STATISTICAL INFERENCE

We applied linear regression [Seber and Lee, 2012] to remove age, sex and dose effects for patients and controls. We used Cohen’s d [Cohen, 1988] to quantify the effect size of the mean differences.

RESULTS

Structural Paths

Using the path definitions based on DTI, we noted that in 96% of subjects (64 out of 66), all 4,005 possible structural links could be classified into primary, secondary or tertiary structural paths. In 27 subjects (14 patients and 13 controls), there were no more than seven pairs of regions that required more than two intermediate linked structures to form an indirect path. On average we obtained 27.8% primary, 68.9% secondary, and 3.4% tertiary structural paths in controls. In patients, 27.4% primary, 69.5% secondary and 3.1% tertiary structural paths were noted, with no significant difference between the two groups (Supporting Information Figure SF2). Of the 4005 possible connections, 3774 were in the same category (primary, secondary, or tertiary) when comparing patients and controls. A substantial degree of agreement was noted across the three categories of connections when the two groups of subjects were compared. 95.4% of primary, 91.5% of secondary, and 75.6% links are classified in the same group in both patients and controls.

We also examined the anatomical distribution of the primary, secondary, and tertiary paths using the conventional distribution of Resting State Networks observed in previous studies. This is presented in Supporting Information 2.

Changes in FCE

Patients did not significantly differ in the whole brain FCE from controls when considered as a single group (P = 0.5). But when stratified according to illness duration, whole brain FCE was significantly higher in patients with early stage illness (P = 0.01, d = 0.91) and lower in those with chronic illness (P = 2 × 10⁻⁶, d = 1.69) in comparison with age and sex-matched controls.

Both patients in the early stage of schizophrenia (P = 0.0002, d = 1.46) and chronic illness (P = 1 × 10⁻¹⁷, d = 4.66) showed a showed a significant reduction in the FCE of tertiary paths when compared to age and sex-matched controls. FCE of secondary paths was increased in those with early stage schizophrenia (P = 0.003, d = 1.12), but significantly reduced in patients at a more chronic stage of illness (P = 2 × 10⁻⁶, d = 1.47). FCE of primary paths showed a trend towards an increase in patients with chronic illness (P = 0.084, d = 0.44) but not in those with early stage illness when compared to matched controls (Fig. 2 panel B).

In the early stage of schizophrenia, higher whole brain FCE was associated with higher symptom burden measured using the total SSPI score (r = 0.705, P = 0.034); this was mostly driven by the increased FCE of secondary paths which was significantly related to the symptom burden (r = 0.74, P = 0.023), while the FCE of primary and tertiary paths did not show any significant relationships. In those with chronic illness, lower whole brain FCE was associated with higher symptom burden (r = −0.543, P = 0.024); this was driven by the FCE of both primary (r = −0.64, P = 0.006) and secondary paths (r = −0.51, P = 0.036) being negatively correlated with the symptom burden, while tertiary paths showed no significant relationship (Fig. 3).

DISCUSSION

When compared with healthy controls, the whole brain FCE is higher in the early stages of schizophrenia indicating an increase in unpredictability and spatial randomness of functional connectivity, especially in those with more severe illness. A pronounced reduction in whole-brain FCE is seen in the later stages of schizophrenia, indicating that the underlying pathophysiological process may be associated with an indistinct, nonspecific mode of brain operation at rest. As
hypothesized originally, we note that the most significant changes in the randomness of functional connectivity in schizophrenia occur in indirect paths that do not have a direct axonal linkage. In particular, we note that the variability of connectivity in tertiary paths shows a striking reduction in both early and late stages of schizophrenia, not altered by the severity of symptoms, suggesting that this is an invariant, “trait-like” feature of the illness.

FCE is reduced in tertiary paths in both groups, and across all pathways and in the whole brain in those with chronic illness. One exception to the observation of predominant FCE reduction is the increased FCE in secondary pathways that annuls the effect of FCE reduction seen in tertiary links, contributing to a lack of group difference with controls, when whole brain FCE is considered in the early stage illness. One may speculate that the increase in FCE seen in the secondary pathways in early stages may reflect a compensatory effort against the process driving a predominant reduction in FCE; the observation that those with more severe illness in early stages show a higher FCE in secondary pathways, supports this speculation.

Patients with chronic schizophrenia showed reduced variability in secondary paths, with a trend towards reduced variability in primary paths as well. Both of these changes in the primary (most significant) and secondary tertiary ones comparing with controls. Right panel: In patients ≥5 years, the FCE was close to significantly decreased (0.03, \( P = 0.0843 \), \( d = 0.44 \)) in primary pathways, significantly decreased (0.136, \( P = 2 \times 10^{-5} \), \( d = 1.47 \)) in secondary and most significantly decreased (0.634, \( P = 1 \times 10^{-17} \), \( d = 4.66 \)) in tertiary when compared with controls. All the effects of age, sex, and dose have been removed. [Color figure can be viewed in the online issue, which is available at wileyonlinelibrary.com.]

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**Figure 2.**

FCE in different fiber pathway. (a): In patients <5 years, the FCE was slightly increased in patients (0.0743, \( P = 0.01 \), \( d = 0.91 \)), and significantly decreased (0.135, \( P = 2 \times 10^{-16} \), \( d = 1.69 \)) in patients ≥5 years comparing with controls. (b): Left panel: In patients <5 years, the functional connectivity (FCE) entropy was almost unchanged (−0.00717, \( P = 0.605 \), \( d = 0.10 \)) in primary pathways, significantly increased (−0.103, \( P = 0.003 \), \( d = 1.12 \)) in secondary ones and significantly decreased (0.187, \( P = 0.0002 \), \( d = 1.46 \)) in tertiary ones comparing with controls. Right panel: In patients ≥5 years, the FCE was close to significantly decreased (0.03, \( P = 0.0843 \), \( d = 0.44 \)) in primary pathways, significantly decreased (0.136, \( P = 2 \times 10^{-5} \), \( d = 1.47 \)) in secondary and most significantly decreased (0.634, \( P = 1 \times 10^{-17} \), \( d = 4.66 \)) in tertiary when compared with controls. All the effects of age, sex, and dose have been removed. [Color figure can be viewed in the online issue, which is available at wileyonlinelibrary.com.]
schizophrenia, the issue of spatial diversity of functional connections was investigated in two previous studies. Both Lynall et al., [2010] and Bassett et al., [2012] quantified the average of the variance in the pairwise connections within each brain region (column-wise variance across brain regions in the functional connectivity matrix) and reported an increase in the dispersion of functional connectivity in schizophrenia. Unlike variance, entropy is an index of diversification and uncertainty that does not depend on the absolute values of the underlying metric (in this case, functional connectivity), and uses more information about the probability distribution than variance, especially relevant in the presence of fat-tailed distributions [Schwarz and McGonigle, 2011; Skudlarski et al., 2010]. Higher entropy within a system suggests that the whole provides more information than the sum of its parts [Tononi et al., 1994]. Variance does not capture the unpredictability when the values taken by a variable are similar to each other (i.e., narrow distribution range) [Bach and Dolan, 2012]. FCE is a more intuitive index of uncertainty when a subtle randomization is expected, as in schizophrenia. Our observations suggest that FCE is a relevant measure in this context that is sensitive to both the clinical stage and severity of schizophrenia.

Unlike the indirect paths, the primary structural paths showed no significant alteration in the early stages and only a mild reduction in FCE reaching trend level significance in the chronic illness. If one assumes that the pathophysiology of schizophrenia primarily affects the functional relationship among brain regions, then this observation suggests that the relationship among brain regions that have a direct axonal linkage is less susceptible, at least in the early stages of the illness. In fact, several crucial alterations in functional connectivity relevant to the pathophysiology of schizophrenia have been noted between regions that do not have direct anatomical linkage. To consider an example, consistent functional connectivity between insula and anterior cingulate cortex has been observed using resting state fMRI, constituting a Salience Network [Menon and Uddin, 2010; Seeley et al., 2007; Taylor et al., 2009]. But to date, no consistent direct axonal linkage has been noted between these two regions [Cerliani et al., 2012; Cloutman et al., 2012; van den Heuvel et al., 2009]. Several studies have now shown that the abnormalities within this “indirect” functional network, is a key feature of psychosis [Manoliu et al., 2013; Palaniyappan et al., 2013; Pu et al., 2012; White et al., 2010]. As shown in the Supporting Information 1, the largest proportion of tertiary paths involved a network that included the insula. Focused investigation of the anatomical distribution of indirect functional links is warranted in future studies.

Poor spatial coherence between structural and functional connectivity has been previously reported in schizophrenia [Skudlarski et al., 2010], implying functional dysconnectivity involving anatomically unconnected regions. Indirect
paths indicate the presence of weaker links in the functional connectivity architecture [Goni et al., 2014]. While most previous investigations of the connectome used thresholding procedures to discarded weaker links from analysis, the importance of weak links is being increasingly appreciated in recent times [Gallos et al., 2012; Schwarz and McGonigle, 2011]. The integrity of weaker links relate to global cognitive capacity in healthy individuals [Santaronechi et al., 2014]. Bassett et al. [2012] have also shown that in schizophrenia abnormalities involving the weaker links are likely to be more specific to the illness process and relate to both cognitive abnormalities and symptom severity in schizophrenia. Our results also suggest that while there is a substantial overlap in the anatomical similarity of primary and secondary paths between patients and controls, 24.4% of all identified tertiary pathways were present only in one group of subjects, highlighting that the weaker, indirect links are more prone to spatial redistribution, in addition to reduced variability in the strength of connectivity.

Functional connectivity of secondary but not tertiary paths can be attributed to connections mediated by a third region either by directed polysynaptic linkage [Lu et al., 2011] or shared afferents [Adachi et al., 2011]. As tertiary pathways do not share common afferent/efferent links, the observed connectivity in these paths are likely to be related to the emergent properties of the network-level architecture [Adachi et al., 2011; Goni et al., 2014]. Uncovering the processes beyond physical connections that bring about functional correlation across time to “bind” a distributed set of brain regions is likely to be a crucial step to understand the pathophysiology of psychosis.

The novelty of our study includes the categorization of functional links using DTI-based structural paths, the measurement of an unbiased estimate of unpredictability associated with functional connectivity, and the investigation of the effect of illness stage and severity on the structure-function relationship. Several limitations must be considered when interpreting our observations. To study the effect of illness duration, we used an arbitrary cut-off point of 5 years. From the multitude of neuroimaging studies investigating the effect of illness duration, no single period has emerged as the time of stabilization of brain changes after the onset. Nevertheless, existing evidence suggests that amount of dynamic (supposedly progressive) changes are most pronounced immediately before onset and in the first five years of illness. We studied a medicated sample of patients; antipsychotic treatment alters functional connectivity of brain regions in schizophrenia [Bolding et al., 2012; Stephan et al., 2001]. Given the suspected confound, we adjusted for the current dose of prescribed antipsychotics. We did not have information on cumulative antipsychotic exposure, but unlike volumetric changes that become more pronounced with cumulative exposure, functional dysconnectivity normalizes or returns to baseline levels on using antipsychotics [Abbott et al., 2013]. As a result, functional dysconnectivity in medicated samples are likely to be an underestimate of the true effect. In the absence of investigations focusing on the effect of antipsychotics on indirect functional links, caution is warranted when interpreting these results. Further, when computing the spatial variability of functional connectivity, we have assumed stationarity of the functional architecture within the time scale of fMRI. The unpredictability introduced by nonstationarity (dynamic “making and breaking” of functional dependencies) [Messe et al., 2014] is beyond the scope of this work.

**CONCLUSION**

To conclude, in a healthy brain, the resting connectional architecture (system as a whole) accommodates a higher degree of variability in functional connections (constituent parts), but in established schizophrenia the variability of constituents that make up this system is limited. Our findings address a critical gap in the literature linking structure and function in schizophrenia, and demonstrate for the first time that the abnormal state of functional connectivity preferentially affects structurally unconstrained links in schizophrenia. It also raises the question of a continuum of dysconnectivity ranging from less direct (structurally unconstrained) to more direct (structurally constrained) brain pathways underlying the progressive clinical stages of schizophrenia.

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