

ORIGINAL ARTICLE

Topologically convergent and divergent functional connectivity patterns in unmedicated unipolar depression and bipolar disorder

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Bipolar disorder (BD), particularly BD II, is frequently misdiagnosed as unipolar depression (UD), leading to inappropriate treatment and poor clinical outcomes. Although depressive symptoms may be expressed similarly in UD and BD, the similarities and differences in the architecture of brain functional networks between the two disorders are still unknown. In this study, we hypothesized that UD and BD II patients would show convergent and divergent patterns of disrupted topological organization of the functional connectome, especially in the default mode network (DMN) and the limbic network. Brain resting-state functional magnetic resonance imaging (fMRI) data were acquired from 32 UD-unmedicated patients, 31 unmedicated BD II patients (current episode depressed) and 43 healthy subjects. Using graph theory, we systematically studied the topological organization of their whole-brain functional networks at the following three levels: whole brain, modularity and node. First, both the UD and BD II patients showed increased characteristic path length and decreased global efficiency compared with the controls. Second, both the UD and BD II patients showed disrupted intramodular connectivity within the DMN and limbic system network. Third, decreased nodal characteristics (nodal strength and nodal efficiency) were found predominantly in brain regions in the DMN, limbic network and cerebellum of both the UD and BD II patients, whereas differences between the UD and BD II patients in the nodal characteristics were also observed in the precuneus and temporal pole. Convergent deficits in the topological organization of the whole brain, DMN and limbic networks may reflect overlapping pathophysiological processes in unipolar and bipolar depression. Our discovery of divergent regional connectivity that supports emotion processing could help to identify biomarkers that will aid in differentiating these disorders.

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INTRODUCTION

Affective disorders, such as unipolar depression (UD, or major depressive disorder) and bipolar disorder (BD), are highly prevalent and debilitating conditions associated with high suicide rates and a heavy social burden.¹ Although BD consists of recurring episodes of mania/hypomania and depression, the depressive episodes are the most common mood manifestation of the illness.² Unfortunately, when BD manifests as the depressed state, it is often misdiagnosed as UD, leading to inappropriate treatment, poor clinical outcomes and greater health-care costs.³ It is, therefore, crucial to identify objective and reliable neurobiological markers to help differentiate these disorders, especially in individuals presenting during depressive episodes.

Accumulating evidence suggests widespread brain structural and functional alterations in both UD and BD (mostly BD I), such as the prefrontal cortex, limbic system, ventral striatum, insula and thalamus.^{4–8} However, direct comparisons of neuroimaging measures between UD and BD depression are sparse and the results are inconclusive. Structural neuroimaging studies suggest

more widespread abnormalities in white matter connectivity and white matter hyperintensities in BD than in UD.^{3,6,9–11} Task-based functional magnetic resonance imaging (fMRI) studies reported patterns of functional abnormalities in the emotion processing, reward and emotion regulation neural circuits in UD that were distinct from those in BD.^{3,4,6,12,13} These abnormalities may help distinguish BD from UD depression. Recently, resting-state fMRI (R-fMRI) has become widely used, as well. By using resting-state functional connectivity (FC) analysis, several studies found FC alterations in the default mode network (DMN) and affective network in different types of affective disorders.^{5,8,14,15} Moreover, Marchand *et al.*¹⁶ found that the right posterior cingulate cortical FC not only differentiates unipolar from bipolar depression but may also mediate differences in neural processes associated with symptom expression (for example, depression severity and current suicidal ideation) between unipolar and bipolar individuals. However, Anand *et al.*¹⁷ and Wang *et al.*¹⁸ failed to observe a significant difference in the resting-state FC between these two disorders, a finding that might imply an overlap in the

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neuropathology of depression. However, some of the extant studies included BD- and UD-depressed groups that were not well matched clinically (such as in the duration of illness or the number of episodes, as well as in the levels of depressive symptomatology), some included individuals taking a diverse array of medications and some included both BD I and BD II individuals.⁶ Notably, there are two established subtypes of BD: type I and type II. The subtype BD II is distinguished from BD I mainly by the absence of full-blown manic episodes. Most neuroimaging studies have involved bipolar I patients, but comparatively few studies have focused on bipolar II disorder.⁸ However, a growing body of evidence suggests that there may be neurobiological differences between BD I and BD II.^{19–21} Therefore, paying attention to the pathophysiology and the neurobiological mechanisms underlying BD II is important.

Graph theory provides a powerful mathematical framework for describing the topological organization of brain networks consisting of nodes (that is, brain regions) and edges (that is, physical or functional connections between brain regions).²² Using graph theoretical methods, researchers revealed that the intrinsic activity of a healthy human brain tends to demonstrate efficient, 'small-world' networks that have an optimal balance between local specialization and global integration for information processing.²³ Optimal small-world networks have a high level of local clustering (that is, nodes are often connected to their neighbors), combined with short path lengths (that is, it takes few steps from any node to any other node in the network) and a low network cost (that is, the mean physical distance between connected nodes is considerably less than is the case in a random network).²³ Several studies have found disrupted small-world networks, modular architecture and abnormal regional characteristics in various brain diseases, such as Alzheimer's disease,²⁴ attention-deficit/hyperactivity disorder²⁵ and schizophrenia.²⁶ Until now, only a few studies have examined the topology of the whole-brain functional network in UD using R-fMRI and graph theory, and those studies that did were inconsistent in the network topological properties that they reported.^{27–31} With respect to BD, only one study showed a disrupted functional topological architecture of the resting-state brain network in BD with hypomanic/manic and depressed episodes (both BD I and II included).³²

To our best knowledge, no study has directly compared the topological organization of large-scale whole-brain functional networks between UD and BD II patients. In this study, we acquired R-fMRI data from currently unmedicated UD- and BD II-depressed patients as well as from healthy controls. Then, we analyzed the topological organization of the whole-brain intrinsic functional networks and determined between-group differences in topological parameters. We hypothesized that UD and BD II patients would show convergent and divergent patterns of disrupted topological organization of the functional connectome, especially in the DMN and the limbic network, both of which support emotion and cognitive functioning. Furthermore, we used a receiver operator characteristic (ROC) analysis to investigate whether these topological organizations showed abnormalities which can discriminate between unipolar and bipolar depression.

MATERIALS AND METHODS

Participants

A total of 32 currently depressed patients with bipolar II disorder and 31 currently depressed patients with UD were recruited from the Psychiatry Department of the First Affiliated Hospital of Jinan University, Guangzhou, China. The patients were 18–54 years old. The diagnoses of UD and BD type II were made according to the Structured Clinical Interview for DSM-IV by two experienced psychiatrists. The clinical state for each patient was assessed using the 24-item Hamilton Depression Rating Scale (HAM-D) and the Young Mania Rating Scale during the 7-day period prior to the R-fMRI

scan. The inclusion criterion for the patients with UD was a total HAM-D-24 score >21, whereas for the depressed patients with BD II was a total Young Mania Rating Scale score <7 and total HAM-D-24 score >21. The exclusion criteria were patients with other Axis-I psychiatric disorders (except for UD, BD and anxiety disorders), a history of organic brain disorders, neurological disorders, mental retardation, cardiovascular diseases, alcohol/substance abuse or dependence, pregnancy or any physical illness. None of the patients had ever received electroconvulsive therapy prior to participating in the study. In addition, all the patients with UD had no family history of BD. At the time of scanning, all the patients were either medication-naïve or had been unmedicated for at least 5 months.

We also recruited 43 healthy subjects as the controls via local advertisements. They were carefully screened using a diagnostic interview, the Structured Clinical Interview for DSM-IV-Nonpatient Edition, to rule out the presence of current or past psychiatric illness. Further exclusion criteria for the healthy controls were any history of psychiatric illness in first-degree relatives and current or past significant medical or neurological illness.

All the subjects were right-handed according to their self-report. In addition, all the included subjects were determined to have no brain abnormalities on conventional MRI by two experienced radiologists. The study was approved by the Ethics Committee of the First Affiliated Hospital of Jinan University, China. Each subject signed a written informed consent form after a full written and verbal explanation of the study.

Data acquisition

All MRI data were obtained on a GE Discovery MR750 3.0 T System with an eight-channel, phased-array head coil. The R-fMRI data were acquired using a single-shot gradient-echo echo planar imaging sequence with the following parameters: repetition time = 2000 ms, echo time = 25 ms, flip angle = 90°, field of view = 240 mm × 240 mm, data matrix = 64 × 64, thickness/gap = 3.0/1.0 mm, 35 axial slices covering the whole brain and 210 volumes acquired in 7 min. During the scanning, each subject was asked to keep their eyes closed, to relax the mind, but not to fall asleep and not to think of anything in particular. In addition, two routine scans using axial T1-weighted fluid attenuation inversion recovery and fast spin-echo T2-weighted MR sequences were also applied to obtain brain images to confirm the absence of any brain structural abnormalities.

Data preprocessing

The R-fMRI data were preprocessed using SPM8 (<http://www.fil.ion.ucl.ac.uk/spm/>) and DPARSF³³ (<http://restfmri.net/forum/DPARSF>). Preprocessing comprised the removal of the first 10 volumes, slice timing correction for the remaining 200 volumes, realignment to the first volume for head motion correction, spatial normalization to the Montreal Neurological Institute space with 3 × 3 × 3 mm³ voxels, signal linear detrending and temporal band-pass filtering (0.01–0.08 Hz). All the subjects in this study satisfied our criteria for head motion, displacement < 1.5 mm in any plane and rotation < 1.5° in any direction.

Constructing brain functional networks

Using the GREYNA toolbox³⁴ (<http://www.nitrc.org/projects/greyna/>), we constructed the brain functional network for each subject according to the automated anatomical labeling (AAL) template.³⁵ First, we randomly parcellated the regions of the AAL template into 1024 ROIs with equal volume size (referred to as the AAL-1024 template) by using the algorithm developed by Zalesky *et al.*,³⁶ each ROI being defined as a node of the network. For a given ROI in each subject, we obtained its time series by averaging the time courses of all the voxels within the ROI and performed a linear regression to remove the effects of the following nuisance covariates: the Friston-24 parameters of head motion (six head motion parameters, six head motion parameters one time point before and the twelve corresponding squared items)³⁷ and the signals of the whole brain, white matter and cerebrospinal fluid. Second, for each node, we first used the residual of its time series to calculate Pearson's correlation coefficients with the other 1023 ROIs and then repeated this process for all the ROIs. Thus, we obtained a 1024 by 1024 symmetric correlation matrix for each subject and the corresponding *P*-value for each matrix element for each subject. Each *P*-value was the probability of getting a correlation as large as the observed value by random chance when the true correlation is zero. Third, we employed a thresholding procedure on the correlation matrix for each subject using two previous studies.^{24,38} Specifically, internodal

correlations surviving a threshold of $P < 0.05$ (Bonferroni correction) were retained or, otherwise, were set to 0. Of note, during the calculations, negative correlations were excluded, given their ambiguous interpretation and detrimental effects on test–retest reliability.^{39–42} Finally, by taking the remaining Pearson’s coefficients as the edge weights, we obtained a weighted 1024 by 1024 FC matrix for each subject, which was used in the subsequent analyses. We also displayed the mean FC and one subject’s FC pattern from each group as examples to help to clarify the FC matrix (Supplementary Figure S2).

Network analysis

Global parameters. Graph theory was used to analyze the topological properties of the human brain functional networks. We used four parameters, the clustering coefficient (C_w), characteristic path length (L_w), global efficiency (E_{glob}) and local efficiency (E_{loc}),⁴³ to characterize the global topological properties of the brain networks. The C_w and E_{loc} indicate the functional segregation in the brain in that C_w shows the extent of the local interconnectivity or cliquishness in a network and E_{loc} measures the ability to transfer local information. L_w and E_{glob} indicate the functional integration in the brain in that L_w measures the harmonic length between pairs and quantifies the ability of information to propagate in parallel and E_{glob} measures the ability to transfer global information. Their definitions and descriptions are provided in Supplementary Table S1. In addition, we utilized three parameters: the normalized clustering coefficient (γ), normalized weighted characteristic path length (λ) and small worldness (δ) to measure the small-world property. Typically, a small-world network satisfies the following criteria: $\gamma > > 1$ and $\lambda \approx 1$, or $\sigma = \gamma/\lambda > 1$.⁴⁴

Modular parameters. We calculated the modularity of the brain network using an optimized algorithm,⁴⁵ based on the average FC matrix of the healthy controls. Modularity is defined as the extent to which a network can be decomposed into subnetworks that are more connected within modules than between modules. The definition of the modularity index Q is also given in Supplementary Table S1.

On the basis of the detected modules, we calculated the values of the intra- and intermodular FC for each subject. The strength of the intramodular FC was defined as the mean of all the internodal FCs within the selected module, whereas the strength of intermodular FC was defined as the mean of the FCs between any two nodes in the two selected modules. For each module, we also used the intramodular FC to estimate the values of the four global parameters (C_w , L_w , E_{loc} and E_{glob}) to characterize the model’s topological properties.

Nodal parameters. We also used two parameters, nodal strength (S_i^w) and nodal efficiency (E_i^w) to characterize the nodal properties of the brain networks. S_i^w quantifies the extent to which a node is relevant to the graph, and E_i^w is the ability to transfer information from one node to others. Their definitions and descriptions are also provided in Supplementary Table S1.

Statistical analysis

Group effect analysis. A non-parametric permutation test, which inherently accounts for multiple comparisons,^{46,47} was used to detect differences between the UD, BD II and control groups in each of the demographic information, clinical variables, global parameters (C_w , L_w , E_{loc} , E_{glob} , γ , λ and δ), intra- and intermodular FCs and nodal parameters (S_i^w and E_i^w). In the calculations, we took age, gender and age–gender interaction as nuisance covariates and regressed them out. The adjective ‘permutation’ applied to parametric tests (t -tests, analysis of variance) indicates that we permuted the group to determine the P -value of the corresponding t or F statistic rather than using the traditional, Gaussian-based P -values. A detailed description of the permutation test is presented in the Supplementary Materials. The false discovery rate (FDR) method⁴⁸ was used to correct for multiple comparisons when needed.

Once a significant group effect was observed for a parameter, we estimated its effect size (η^2) according to Cohen’s definition,⁴⁹ according to which the levels of small, medium and large effect size correspond to 0.01, 0.06 and 0.14, respectively. In addition, we performed *post hoc* tests to determine the between-group differences for each of the network parameters (10 000 permutations).

On the basis of the individual brain functional networks, we also performed a network-based statistic analysis³⁶ to identify any subnetworks, or clusters of connections, in which each edge showed significant

differences between the UD patients and the control group or between the BD patients and the control group. This approach can control the family-wise error rate when mass-univariate testing is performed at every connection in the subnetwork.³⁶ The description of the network-based statistic approach is presented in the Supplementary Materials.

Correlations between the network parameters and clinical variables. For the network parameters that showed a significant group effect, we performed a partial correlation analysis to describe their relationships with the clinical variables in each patient group ($P < 0.05$, FDR-corrected). These clinical variables included the HAMD scores, number of episodes, onset age of illness and duration of illness.

Robustness

Robustness is a key issue in network analysis.^{50,51} To determine whether the observed group differences in the topological parameters reflect true differences rather than artifacts, we repeated the network analyses using each of the following approaches. First, we used both the positive and negative correlations to construct the brain network. Second, we constructed the binary network. Third, we constructed the network based on a sparsity range. Finally, we considered the effect of the global signal. Specifically, we constructed the network based on the positive correlations and then constructed it based on both the positive and negative correlations without regressing out the global signal.

Network classification

For those network parameters showing a significant group effect, we plotted the ROC curve to determine which of these parameters could clearly distinguish the UD patients from the controls, the BD II patients from the controls and the BD II from the UD patients. The ROC curve is a fundamental plot in signal detection theory^{52,53} and is widely used in medical science and neuroimaging studies. A ROC curve, a scatter plot showing the relationship between the false-positive rates and the true-positive rates, describes the relationship between the underlying distribution of the places where signals were absent and the places where signals were present. This analysis was performed using a public MATLAB code (<http://www.mathworks.com/matlabcentral/fileexchange/199500-roc-curve>; by Giuseppe Cardillo).

RESULTS

Demographic information

Table 1 lists the demographic and clinical characteristics of the three groups. No significant differences were detected between the three groups in age, gender, education levels or the mean framewise displacement parameter. In addition, there was no significant difference in depression severity, illness duration or onset age of illness between the two depressed groups.

Global parameters

Figure 1 shows the global parameters (C_w , L_w , E_{loc} , E_{glob} , γ , λ and δ) for the UD, BD and control groups. Statistical analyses revealed significant differences in L_w and E_{glob} between the three groups ($P < 0.05$). *Post hoc* comparisons revealed that both the UD and BD II groups had a significantly increased L_w and a significantly decreased E_{glob} , compared with the controls ($P < 0.05$). However, no differences were found between the UD and BD II groups in these parameters (see Supplementary Table S2 for more information).

Modularity

In the controls, we identified seven modules ($Q = 0.6803$): the sensorimotor network, cerebellum, visual network, limbic network, frontoparietal network, DMN and salience network, which are shown in Supplementary Figure S3. The limbic network primarily includes the bilateral hippocampus, parahippocampus, thalamus, anterior cingulate gyrus, temporal pole, caudate and amygdala. The DMN primarily involves the bilateral medial frontal gyrus, superior frontal gyrus, precuneus, posterior cingulate gyrus,

Table 1. Demographics and clinical characteristics of the patients with UD, patients with BD and HCs

Parameters	UD (n = 31)	BD II (n = 32)	HC (n = 43)	P-value
Age (years)	30.41 ± 9.85	28.22 ± 10.13	30.19 ± 11.11	0.23 ^a
Gender	10 M/21 F	13 M/19 F	17 M / 26 F	0.39 ^b
Number of episodes	1.93 ± 2.19	2.16 ± 1.19	N/A	0.35 ^c
Age of illness onset (years)	26.41 ± 10.72	24.58 ± 10.42	N/A	0.26 ^c
Duration of illness (months)	40.68 ± 50.41	39.14 ± 56.62	N/A	0.46 ^c
HAMD-24	26.76 ± 5.58	26.23 ± 6.24	N/A	0.37 ^c
YRMS	2.31 ± 3.69	1.58 ± 1.50	N/A	0.16 ^c
Mean FD (mm)	0.09 ± 0.06	0.09 ± 0.04	0.08 ± 0.03	0.42 ^a

Abbreviations: ANOVA, analysis of variance; BD, bipolar disorder; FD, framewise displacement; HAMD, Hamilton Depression Scale; HC, healthy control; N/A, not applicable; UD, unipolar depression; YRMS, Young Mania Rating Scale. ^aThe P-value was obtained from a permutation ANOVA analysis. ^bThe P-value was obtained from Pearson's χ^2 -test. ^cThe P-value was calculated from a permutation two-sample t-test.

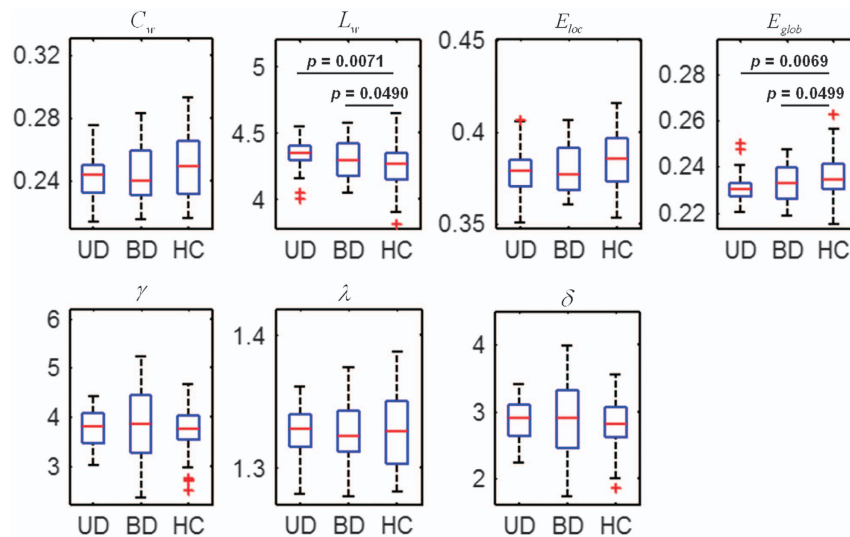


Figure 1. Box plots showing statistical comparisons in the global parameters between the patients with UD, patients with BDs and HCs. The symbol of '+' in red color indicates outliers. Significant group effects were observed in L_w ($P=0.0335$) and E_{glob} ($P=0.028$). δ , small worldness; γ , normalized clustering coefficient; λ , normalized shortest path length; BD, bipolar disorder; C_w , weighted clustering coefficient; E_{glob} , weighted global efficiency; E_{loc} , weighted local efficiency; HC, healthy control; L_w , weighted characteristic path length; UD, unipolar depression.

angular gyrus, middle temporal gyrus and inferior temporal gyrus. The sensorimotor network includes the bilateral precentral gyrus and postcentral gyrus. The visual network includes the bilateral occipital gyrus. The frontoparietal network involves the bilateral inferior parietal gyrus, middle frontal gyrus, inferior frontal gyrus, superior frontal gyrus and middle temporal gyrus. The salience network involves the bilateral insular and cingulate gyrus. These were consistent with previous studies.^{54,55}

We calculated four global parameters (C_w , L_w , E_{loc} and E_{glob}) for each module as well as the intra- and intermodular FCs for each subject. The statistical analysis for these seven modules, except for the DMN and the limbic network, showed no significant group effect for any of these four parameters. Figure 2a shows that for the DMN, the four global parameters were all significantly different between the three groups ($P < 0.05$, FDR-corrected; Supplementary Table S2). *Post hoc* comparisons revealed that both the UD and BD II groups had a significantly decreased C_w , E_{loc} and E_{glob} , but significantly increased L_w , compared with the controls. Figure 2b shows that, for the limbic network, the four global parameters were significantly different between the three groups ($P < 0.05$, FDR-corrected; Supplementary Table S2). *Post hoc* comparisons found that both the UD and BD II groups had a

significantly decreased C_w , E_{loc} and E_{glob} , but a significantly increased L_w , compared with the controls. However, no differences in these parameters were found between the UD and BD II groups.

Figure 2c shows that, of the seven modules, the intramodular FCs in the DMN and in the limbic network were significantly different between the three groups ($P < 0.05$, FDR-corrected; Supplementary Table S3). *Post hoc* comparisons revealed that the UD and BD II groups had significantly decreased intramodular FCs in both the DMN and the limbic network compared with the controls. However, no differences in the intramodular FC in the DMN and the limbic network were found between the UD and BD II groups. For the other modules, we found no significant group effect in the intramodular FC.

The seven modules gave 28 intermodular FCs ($C_7^2 = 21$ intermodular FCs). However, no significant group effect survived after FDR correction ($P < 0.05$, FDR-corrected).

Nodal parameters

Figure 3 shows the brain regions with a significant group effect in the nodal parameters (S_i^w and E_i^w) between the three groups ($P < 0.05$, FDR-corrected). We found a significant group effect in

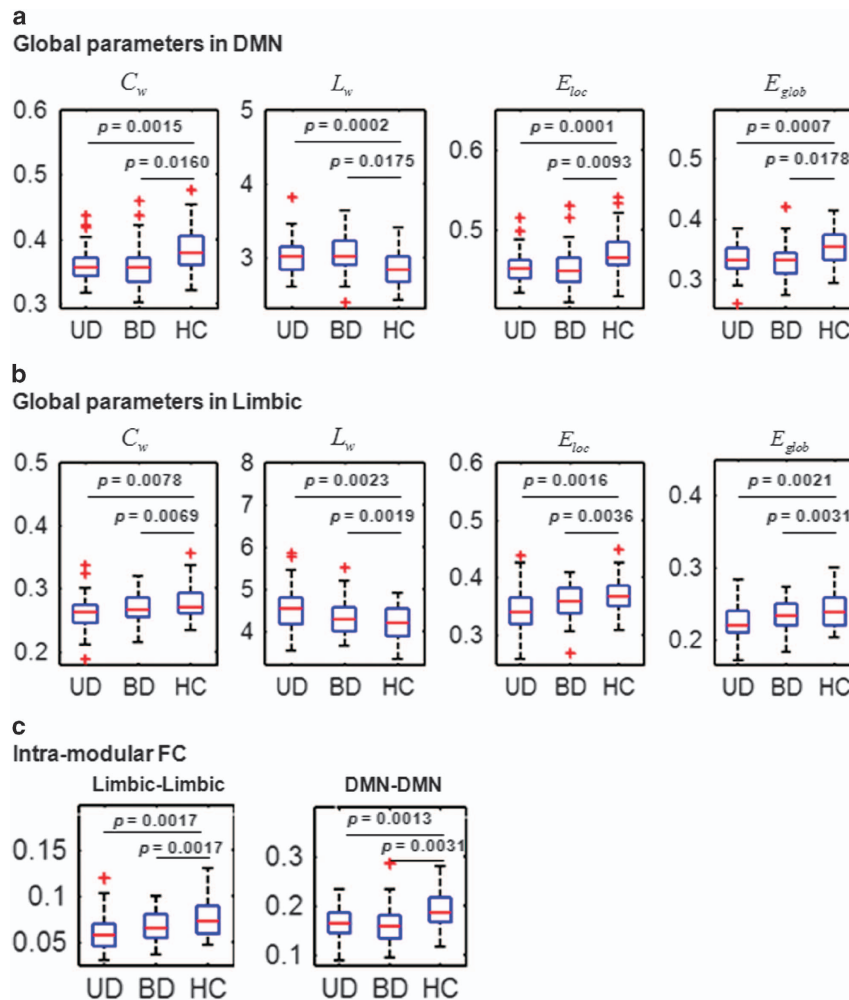


Figure 2. Box plots showing statistical comparisons in the global parameters and intramodular FC between the patients with UD, patients with BDs and HCs (permutation ANOVA, $P < 0.05$, corrected) in the modules of the DMN and the limbic network. The symbol of '+' in red color indicates outliers. **(a)** For the DMN, we observed significant group effects in C_w ($P = 0.0147$), L_w ($P = 0.0037$), E_{loc} ($P = 0.0045$) and E_{glob} ($P = 0.0026$). **(b)** For the limbic network, we observed significant group effects in C_w ($P = 0.0068$), L_w ($P = 0.0008$), E_{loc} ($P = 0.0011$) and E_{glob} ($P = 0.0019$). **(c)** For the intramodular FC, we observed significant group effects in the DMN ($P = 0.0020$) and the limbic network ($P = 0.0064$). δ , small worldness; γ , normalized clustering coefficient; λ , normalized shortest path length; ANOVA, analysis of variance; BD, bipolar disorder; C_w , weighted clustering coefficient; DMN, default mode network; E_{glob} , weighted global efficiency; E_{loc} , weighted local efficiency; FC, functional connectivity; HC, healthy control; L_w , weighted characteristic path length; UD, unipolar depression.

S_i^w in two regions (Figure 3a), which are located in the right superior temporal pole and left posterior lobe of the cerebellum. *Post hoc* comparisons revealed that, in the right superior temporal pole, the UD group had a significantly decreased S_i^w compared with the BD II group and with the control group. In the left cerebellum posterior lobe, both the UD and BD II groups had a significantly decreased S_i^w compared with the controls. In addition, we detected a significant group effect in E_i^w in six regions (Figure 3b), which are located in the bilateral medial superior frontal gyri, bilateral precuneus, right middle temporal pole and left middle cingulum. *Post hoc* comparisons revealed that in the right precuneus both the UD and BD II groups had a significantly decreased E_i^w compared with the controls and the UD group had a significantly increased E_i^w compared with the BD II group. In the bilateral medial superior frontal gyri and left middle cingulum, both the UD and BD II groups had a significantly decreased E_i^w compared with the control group. In addition, in the right middle temporal pole and left precuneus, the UD group showed a significantly increased E_i^w compared with the BD II and control groups. Furthermore, we noticed that most of these regions (1/2 for the S_i^w and 6/6 for the E_i^w) are included in the DMN and

the limbic network (see Supplementary Table S4 for more information).

Relationship between network parameters and clinical variables

For the correlations between network parameters and clinical variables, no significant correlation results survived after FDR correction ($P < 0.05$, FDR-corrected). We have also displayed the correlation results without correction in Supplementary Figure S5.

Robustness

Table 2 lists the statistical results between the UD, BD II and control groups using four other data-processing strategies. These calculations indicated that most of our findings reported above were relatively reliable across the various strategies. Specifically, significant differences were always found in the the L_w and E_{glob} between the UD, BD II and control groups.

Classification

For the network parameters showing significant group effects, we plotted their ROC curves to examine their ability to distinguish the

Table 2. Robustness of the statistical comparisons of the global parameters for the brain functional networks obtained with different strategies between the UD, BDs and control groups ($P < 0.05$)

Different processing strategies	C_w	L_w	E_{loc}	E_{glob}	δ
<i>Global regressing</i>					
Main processing strategy	-	S	-	S	>1
Sparsity	-	S	-	S	>1
No global regressing	-	S	-	-	>1
Both positive and negative	-	S	-	S	>1
Binary	-	S	-	S	>1
<i>No global regressing</i>					
Positive	-	S	-	-	>1
Both positive and negative	S	S	-	S	>1

Abbreviations: δ , small worldness; BD, bipolar disorder; C_w , weighted clustering coefficient; E_{glob} , weighted global efficiency; E_{loc} , weighted local efficiency; FC, functional connectivity; L_w , weighted characteristic path length; R-fMRI, resting-state functional magnetic resonance imaging; UD, unipolar depression. Bold values are significant at $P < 0.05$. Notes: The 'S' indicates a significant group effect. '-' indicates no significant group effect. 'Sparsity' stands for estimating global parameters in the threshold range of 0.10–0.35 at an interval of 0.01 by using a measure of sparsity (the ratio between the total number of edges and the maximum possible number of edges in a network). We applied these thresholds to each of the FC matrices. 'No global regressing' refers to a network analysis based on a connectivity matrix that was constructed from the R-fMRI data without regressing out the global signal. 'Both positive and negative' refers to a network analysis based on a connectivity matrix that included both positive and negative internodal correlations. 'Positive' refers to a network analysis based on a connectivity matrix that only included positive internodal correlations. In addition, 'Binary' refers to a connectivity matrix that was converted into a binary matrix at a selected threshold when constructing the network. Bold values are significant at $P < 0.05$.

DISCUSSION

By studying these unmedicated UD patients and unmedicated BD II-depressed patients, we detected abnormal topological parameters of the brain functional networks without the potentially confounding influence of psychotropic medications such as lithium or antidepressants. The main findings are as follows: (1) at the global level, both the UD and the BD II patients showed a significantly increased L_w and a decreased E_{glob} ; (2) at the module level, both the UD and BD II patients had disrupted intramodular connectivity within the DMN and limbic network; (3) at the nodal level, both the UD and BD II patients showed impaired nodal parameters predominantly in regions of the DMN, limbic network and cerebellum; (4) the UD and BD II groups showed differences in the nodal parameters in the precuneus and temporal pole. To our knowledge, this study is the first to report topologically convergent and divergent patterns of the whole-brain functional networks between UD and BD II patients. These findings may contribute to our understanding of the neuropathological mechanisms in these two disorders.

Similar global-parameter changes in UD and BD

In this study, the whole-brain functional networks in the UD, BD II and control groups showed small-world properties that were consistent with previous findings in healthy adults.^{22,56} We also found several global topological properties of the whole-brain networks, including increased characteristic path length L_w and decreased global efficiency E_{glob} (Figure 1), that showed the same change in direction in the UD and BD II groups compared with the controls. These findings, which were consistent with previous neuroimaging studies of brain functional and structural networks in UD patients^{28,57,58} and BD patients,^{32,59,60} may reflect the

disrupted global integration of the whole-brain functional networks in the two disorders. We also noticed inconsistency between this study and several previous studies in the changes in the global properties in the UD patients. For example, Zhang *et al.*³¹ found decreased characteristic path length L_w and increased global efficiency E_{glob} in first-episode drug-naive UD patients by measuring the partial correlation coefficients of the R-fMRI signals between 90 cortical and subcortical regions; this is the opposite of what we found. In addition, several R-fMRI studies found no differences in global topological properties between UD patients and healthy controls.^{61–63} These inconsistencies may have been caused by any of several factors, including sample heterogeneity (age, depressive episode, medication and disease duration), modality used (for example, EEG/MEG; structural, diffusion and functional MRI), selection of the network matrix (binary or weighted) or the definitions of node and edge weight.

Similar modular-parameter changes in UD and BD

In this study, both the UD and BD II groups had decreased clustering coefficients C_w , local efficiencies E_{loc} , global efficiencies E_{glob} and increased characteristic path lengths L_w within two modules: the DMN and the limbic network (Figures 2a and b). This may indicate disrupted effective segregation and integration of information processing within these networks. In addition, the UD and BD II groups had decreased intramodular FCs in the DMN and limbic network (Figure 2c). These findings suggest that UD and BD patients share similar disruptions of their intrinsic modular structures. In fact, previous studies have suggested that the DMN is involved in a diverse array of functions, such as episodic memory, self-relevant mental processing, monitoring the external environment, remembering the past and planning the future.^{64,65} Several studies have also found DMN abnormality in UD^{5,31,64,66} and BD^{6,67} patients. Aberrant DMN function could lead to dysfunctional self-referential and affective processing in the form of an excessively negative self-focus.^{68,69} The limbic system is the central part of the 'emotional brain' circuitry and is dedicated to processing and regulating emotion or participating in emotional expression.^{70,71} Previous studies have reported brain structural and functional abnormalities in the limbic system supporting emotion dysregulation in UD and BD patients,^{4,72} and several studies found dysfunctional limbic circuits in depression with clinical remission, regardless of the specific treatment modality.^{72,73} Taken together, our findings of disruptions in the DMN and the limbic network in UD and BD II may contribute to core deficits in cognitive and affective functioning that are believed to underlie clinical symptoms.

Similar and distinct nodal parameter changes in UD and BD

In addition to the global and modular levels, we found that both the UD and BD II groups showed decreased nodal strength and nodal efficiency predominantly in the DMN and the limbic network, including the bilateral precuneus, left middle cingulum, bilateral superior frontal gyri and right temporal pole (Figure 3 and Supplementary Table S4), which suggests impaired regional connectivity in these regions. These results are consistent with several previous studies related to UD and BD, which reported that these regions exhibited abnormalities in FC and local activation,^{5,6,8,74} glucose metabolism⁷⁵ and gray matter morphology.⁶ In addition, we found decreased nodal strength in the left posterior lobe of the cerebellum in both the UD and BD II groups compared with the controls (Figure 3a). In fact, other studies have shown cerebellum abnormalities, including reduced gray matter volume,^{76–78} decreased spontaneous activity^{18,78,79} and decreased FC,¹⁸ in both UD and BD patients. A recent R-fMRI study also found increased cerebellar–DMN connectivity in depressed patients compared with healthy controls.⁸⁰ Therefore, our finding of impaired connectivity in the posterior lobe of the

cerebellum provides additional evidence for the involvement of cerebellar dysfunction in the pathophysiology of UD and BD.

We also found significant differences in nodal parameters (nodal strength and nodal efficiency) between the UD and BD II patients in the bilateral precuneus and right temporal pole (Figure 3), suggesting pathophysiological differences between UD and BD. Specifically, BD patients may display more significant abnormalities than UD patients in these regions. A further ROC analysis showed that abnormal nodal strength in the right superior temporal pole was specific to the UD group (Supplementary Table S5), whereas abnormal nodal efficiency in the right precuneus was specific to the BD II group (Supplementary Table S6). Nevertheless, a larger sample size is warranted in future studies to replicate our results and to extend the generalizability of our results. We also found that nodal efficiency in the precuneus had high classification power in discriminating the BD group from the UD group (Supplementary Table S7). These results are partially consistent with previous studies. A R-fMRI study found that the increase in the intrinsic connectivity of the amygdala with the temporal poles was inversely correlated with depression severity in UD.⁸¹ Task-based fMRI studies found increased amygdala activity in response to angry⁸² and sad^{12,83} faces, but decreased amygdala activity in response to positive^{12,83} faces in UD compared with participants with BD. These changes in the amygdala may represent a potential neural basis for the increased negative and reduced positive emotional attentional bias in UD. In addition, a structural MRI study found a gray matter volume difference between UD and BD in the amygdala.⁷⁷ Thus, these findings suggest that abnormalities in the temporal pole and amygdala supporting emotion processing may be specific to UD, although the underlying mechanisms remain to be clarified.^{77,84} In addition, Marchand *et al.*¹⁶ suggested that the FC of the right posterior cingulate cortex can be used to differentiate bipolar type II and UD. Recently, a voxel-based meta-analysis of diffusion tensor imaging studies revealed a greater decrease in fractional anisotropy in the left posterior cingulum in BD compared with UD.⁸⁵ A number of studies support the importance of the posterior cingulum and precuneus/posterior cingulate cortex in neurocognitive functions such as memory, attention and planning.⁸⁶ Therefore, these studies, combined with our findings of reduced nodal efficiency in the precuneus in BD, suggest that cognitive deficits may be more representative of BD,^{85,87} although a degree of cognitive decline has been described in both UD and BD.⁸⁸ Finally, our results of differences in nodal parameters in the precuneus and temporal pole may aid in distinguishing the two types of depression.

Limitations

This study has several limitations. First, we cannot predict whether some patients with depression diagnosed as UD will later switch to BD, although the patients with UD in this study had no family history of BD. In addition, we tracked the UD patients after we acquired the data and found that no UD patient had switched to BD by the time of the submission of this manuscript. Second, this study did not consider the effect of brain structural changes on the network topologies in the UD or BD patients, which may bias the findings of the present study. Several previous studies suggested that brain structural changes may have an impact on the results of FC and network analysis.²⁸ Third, we only examined patients with BD II, thereby avoiding the potential confounds of the bipolarity subtype differences in brain functional networks but also limiting the generalizability of our findings to BD I. Fourth, although the participants were unmedicated for at least 5 months prior to the scan, each group may have had variable levels of past exposure. Thus, it is difficult to estimate the potential contribution of lifetime exposure to medication in relation to the results. Fifth, because clinical symptomatology could change over a period of

7 days, collecting the measurements of clinical variables such as the HAMD and Young Mania Rating Scale scores over a 7-day period is another limitation. Sixth, because of selection bias,⁸⁹ the ROC results should be interpreted as descriptive and exploratory, requiring confirmation in an independent data set. Further studies utilizing optimized analytical strategies and multiple imaging modalities are needed to detect more sensitive and robust biomarkers for UD and BD. Finally, like all R-fMRI studies, our results are based on the FC analysis, which is only an exploratory technique and requires confirmation with task-based or experimental work.

In conclusion, our study revealed convergent deficits in the topological organization in the whole brain and in the DMN and the limbic network in UD and BD II. These deficits may reflect overlapping pathophysiological processes in unipolar and bipolar depression. In addition, divergent regional connectivity supporting emotion processing could potentially provide biomarkers that would aid in differentiating between these disorders.

CONFLICT OF INTEREST

The authors declare no conflict of interest.

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