Functional connectivity between the amygdala and prefrontal cortex in medication-naive individuals with major depressive disorder

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Introduction

Episodes of major depressive disorder (MDD) are characterized by negative affect and negative biases in processing emotional stimuli, implicating abnormalities in the neural system that processes negative emotional stimuli.1 The prefrontal cortex (PFC) shares extensive connections with the amygdala. These structures are central to a neural system that processes emotional stimuli, especially negatively valenced stimuli.2 Accumulating studies implicate this PFC-amygdala neural system in the disordered emotional processes of individuals with MDD.3

Morphological magnetic resonance imaging (MRI) and conventional functional MRI (fMRI) activation analyses provide evidence of morphological and functional abnormalities within the PFC and the amygdala in adults with MDD.4 We previously found morphological abnormalities in the amygdala in medication-naive individuals with MDD.5 Other
groups have demonstrated excessive responses of the amygdala to negative emotion, especially fearful facial expressions, in medicated individuals with MDD. Studies of the role of the PFC suggest that functional imbalance between the left and right PFC in emotion processing may also be involved in the neuropathophysiology of MDD. 20,21 Consistent with this idea, lesions in the left PFC and left PFC dysfunction, which lead to deficits in the capacity to experience positive affect, a key feature of depression, have been reported in association with depression, particularly during negative emotion processing, possibly indicating greater dysfunction in the left than the right PFC in individuals with MDD.

The conventional fMRI activation studies mentioned previously provide information about the functioning within specific brain regions. To study the ability of brain regions to work together, specialized measures can be used to examine the coordinated activity between brain regions or their “functional connectivity.” Connectivity fMRI (cFMR) assesses activity in different brain regions that are coupled in time. Several cFMR studies have shown connectivity abnormalities in individuals with MDD. For example, studies have shown decreased connectivity between the anterior cingulate cortex (ACC) and amygdala in response to negative stimuli in unmedicated individuals with MDD. Almeida and colleagues reported reduced amygdala–left orbitomedial prefrontal functional connectivity during processing of happy and sad faces in medicated patients with MDD. However, increased connectivity between frontal cortices, including the medial PFC and rostral ACC, and the amygdala when processing negative words has also been reported in depressed patients. The conflicting findings among studies may be related to differences in tasks performed during scanning and/or differences in patient samples, particularly in terms of illness chronicity or medication. Importantly, some studies suggest that antidepressant treatment influences the functioning of the amygdala and frontal cortices and their connectivity in individuals with MDD.

In the present fMRI study, medication-naive individuals with MDD were scanned while processing emotional facial stimuli depicting fearful, happy, and neutral expressions. We assessed the strength of the correlation in time between the blood oxygen level–dependent (BOLD) responses from the amygdala to the PFC. We hypothesized that participants with MDD would demonstrate deficits in functional connectivity between the amygdala and PFC, particularly the left PFC, and especially when processing negative emotional stimuli.

Methods

Participants

We recruited medication-naive patients with MDD aged 19–46 years from the outpatient clinic at the Department of Psychiatry, First Affiliated Hospital of China Medical University and the Mental Health Center of Shenyang. Some of these participants had been included in a previously published resting-state fMRI study. The diagnosis of MDD was confirmed by 2 trained psychiatrists (L.K. and F.W.) using the Structured Clinical Interview for DSM-IV disorders. To be included in our study, individuals with MDD had to fulfill the DSM-IV criteria for MDD; have a current depressive episode; have no comorbid Axis I or II diagnoses; have a score of at least 24 on the 17-item Hamilton Rating Scale for Depression (HAM-D-17); and have no history of psychopharmacotherapy, electroconvulsive therapy or psychotherapy.

We recruited healthy controls from Shenyang, China, via advertisement. Some of these participants had also been included in the previously published resting-state fMRI study. The absence of DSM-IV Axis I disorders in controls was confirmed by 2 independent psychiatrists (L.K. and F.W.) using the Structured Clinical Interview for DSM-IV disorders. Individuals with first-degree family members who had a history of DSM-IV Axis I disorders were excluded.

Additional exclusion criteria for both individuals with MDD and controls were the presence of any MRI contraindications, history of head injury or neurologic disorder, history of drug abuse or dependence, and any concomitant medical disorder. All participants were scanned within 24 hours of initial contact with the research team. The participants provided written informed consent after receiving a detailed description of the study. The Institutional Review Board of the China Medical University approved our study protocol.

MRI data acquisition

The fMRI data were acquired using a GE Signa HDX 3.0 T MRI scanner at the First Affiliated Hospital of China Medical University, Shenyang, China. Head motion was minimized with restraining foam pads. We used a standard head coil for radiofrequency transmission and reception of the nuclear magnetic resonance signal. The fMRI images were acquired using a spin–echo planar imaging sequence, parallel to the anterior commissure–posterior commissure plane with the following scan parameters: repetition time 2000 ms, echo time 40 ms, image matrix 64 × 64, field of view 24 × 24 cm², 35 contiguous slices of 3 mm and without gap.

Emotional face paradigm

During the fMRI runs, each participant completed an event-related facial emotion task, which has been described previously. Participants viewed faces from the Ekman series depicting fearful, happy or neutral expressions, and they were instructed to press a button to make a male–female determination. In brief, 5 male and 5 female faces were each presented for 2 s with interstimulus intervals of 4, 8 or 12 s. Each of the 3 expressions was shown for each individual, for a total of 30 facial stimuli and a run time of 5 min, 6 s. The order of the facial stimuli varied to control for sequential dependencies.

Functional connectivity processing

Statistical Parametric Mapping 8 (SPM8) software (www.fil.ion.ucl.ac.uk/spm) was used for BOLD fMRI preprocessing. We discarded the initial 2 images. The remaining images were corrected for within-scan acquisition time differences
between slices and realigned to the first volume to correct interscan movements. Linear motion (x, y, z planes) for all participants was below 2.5 mm and rotational motion (pitch, roll, yaw) below 2.5 degrees. The fMRI data were then spatially normalized to Montreal Neurological Institute (MNI) space, resampled to $3 \times 3 \times 3$ mm$^3$ and spatially smoothed (8 mm full-width at half-maximum).

The bilateral amygdala seed region of interest (ROI) was defined with the WFU PickAtlas Tool (www.fmri.wfubmc.edu/download.htm). For each participant, we calculated a mean time series for the amygdala seed ROI by averaging the time series for all voxels within the amygdala ROI separately for each facial emotion type (fearful, happy, neutral). We then performed correlational analyses between the amygdala time series and the time series for each brain voxel,$^2$ resulting in 3 correlation maps for each participant, 1 for each facial emotion type. The correlation coefficients in each map were transformed to z values using Fisher r-to-z transformation for further statistical testing.

### Statistical analyses

We used independent-sample $t$ tests and $\chi^2$ tests to compare demographic data, HAMD-17 and Hamilton Anxiety Rating Scale (HARS) scores between the MDD and control groups using SPSS 13.0 software (SPSS Inc.). Group differences in functional connectivity were analyzed using 2-sample (MDD v. control) $t$ tests in SPM, with the functional connectivity correlation coefficients ($r$ values) from the amygdala to all brain voxels as the dependent variables for each variable factor (fearful, happy, neutral). The threshold for contrast maps was set at $p < 0.005$ and a cluster size of at least 729 mm$^3$ (27 voxels) for the hypothesized PFC region; this was equal to a corrected threshold of $p < 0.05$, determined by the Monte Carlo simulation (AlphaSim command line in AFNI [analysis of functional neuroimaging; Cox, 1996]). The PFC was defined with the WFU PickAtlas Tool (www.fmri.wfubmc.edu/download.htm), including Brodmann Areas (BA) 9–12, 24, 25, 32 and 44–47. A linear mixed model was conducted to assess the effects of diagnostic group and the 3 emotional stimulus types on functional connectivity measurement ($z$ values in the PFC regions showing significant differences between the control and MDD groups) with SPSS. In this model, diagnostic group (control v. MDD) represented a between-subjects factor, and emotional type (fear, happy, neutral) was included as a within-subjects factor. The interaction between diagnostic group and emotion was modelled. If the diagnostic group effect and interaction between diagnostic group and emotion demonstrated significance, we performed 2-sample $t$ tests separately for the 3 emotional types in the post hoc analyses, and we considered results to be significant at $p < 0.05$, Bonferroni corrected. We conducted whole brain analyses to explore other possible brain regions not hypothesized a priori. Findings in these regions were considered to be significant at $p < 0.05$, family wise error-corrected for multiple comparisons. We performed post hoc exploratory Pearson correlation analyses in MDD participants to assess the correlation of HAMD-17 and HARS scores with $z$ scores in the PFC regions that were significantly different between the MDD and control groups.

### Results

The MDD group comprised 28 participants with a mean age of 30.6 ± standard deviation (SD) 8.7 years. Half of them were women, and 22 had been included in an earlier study. The mean years of education was 12.9 ± 3.2, the mean duration of illness was 13.0 ± 15.1 months, the mean HAMD-17 score was 28.5 ± 5.2 and the mean HARS score was 20.1 ± 8.3. The control group comprised 30 participants with a mean age of 29.5 ± 8.0 years. Half of them were women, and 23 had been included in a earlier study. All participants were right-handed. There were no significant differences in age ($p = 0.62$), sex ($p > 0.99$) and education ($p = 0.19$) between the MDD and control groups. The MDD group had significantly higher HAMD-17 and HARS scores than the control group ($p < 0.001$, Table 1).

We observed decreased amygdala–left rostral PFC (rPFC; BA 10) functional connectivity in the MDD group compared with the control group for the fearful face condition ( maximal MNI coordinates: $x, y, z = -15, -5, 3$; 37 voxels (999 mm$^3$); $t = 3.65; p < 0.005$, uncorrected; Fig. 1). These findings correspond to a corrected $p < 0.05$ by AlphaSim correction. Analysis of $z$ values in the rPFC showed that the main effect of diagnosis was significant ($F_{1,56} = 6.63, p = 0.010$). In addition, there was a significant group × emotion interaction ($F_{1,56} = 13.31, p = 0.001$). The post hoc 2-sample $t$ tests demonstrated that the contribution to group difference and interaction between group and emotion types was derived mainly from increased $z$ values for the fearful condition ($t_{56} = 3.64, p = 0.003$, Bonferroni corrected) in the MDD compared with the control group. The $z$ values for the other 2 emotion types were not significant (all $p > 0.05$, Bonferroni corrected; Table 2). No significant group decreases in functional connectivity were detected for the happy or neutral conditions. No significant group increases in functional connectivity were detected for any condition. Our whole brain analysis revealed no group differences in altered functional connectivity between the amygdala and other brain regions for any of the conditions. In post hoc correlation analyses, neither HAMD-17 scores nor

### Table 1: Demographic and clinical characteristics of participants with major depressive disorder and healthy controls

<table>
<thead>
<tr>
<th>Characteristic</th>
<th>MDD</th>
<th>Control</th>
<th>$t$ value</th>
<th>$p$ value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Age, yr</td>
<td>30.6 ± 8.7</td>
<td>29.5 ± 8.0</td>
<td>$t_{26} = 0.50$</td>
<td>0.62</td>
</tr>
<tr>
<td>Sex, male:female</td>
<td>14.14</td>
<td>15.15</td>
<td>$\chi^2 = 0.000$</td>
<td>&gt; 0.99</td>
</tr>
<tr>
<td>Education, yr</td>
<td>12.9 ± 3.2</td>
<td>13.9 ± 2.8</td>
<td>$t_{26} = 1.32$</td>
<td>0.19</td>
</tr>
<tr>
<td>HAMD-17 score</td>
<td>28.5 ± 5.2</td>
<td>26.6 ± 1.0</td>
<td>$t = 29.09$</td>
<td>&lt; 0.001</td>
</tr>
<tr>
<td>HARS score</td>
<td>20.1 ± 8.8</td>
<td>11.1 ± 2.5</td>
<td>$t = 12.27$</td>
<td>&lt; 0.001</td>
</tr>
<tr>
<td>Illness duration, mo</td>
<td>13.0 ± 15.1</td>
<td>—</td>
<td>—</td>
<td>—</td>
</tr>
</tbody>
</table>

HAMD-17 = 17-item Hamilton Rating Scale for Depression; HARS = Hamilton Anxiety Rating Scale; MDD = major depressive disorder; SD = standard deviation.

*Unless otherwise indicated.

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HARS scores had significant associations with rPFC functional connectivity in participants with MDD.

Discussion

In this study, we detected a deficit in amygdala–left rPFC functional connectivity in response to fearful face processing in individuals with MDD. To our knowledge, this is the first study to detect such deficits in medication-naive individuals with MDD. Although rPFC is the single cytoarchitectonic subregion of the frontal lobes in the human brain, studies have shown the rPFC plays multiple roles in brain functions. It has been implicated in the integration of information from several emotional and cognitive domains. Recently, Okada and colleagues reported reduced rPFC activation during a verbal fluency task in individuals with remitted MDD. The similar executive dys-function was also reported in an event-related potentials study in an MDD population. The amygdala has been proven to play an important role during emotion processing in animal and human research. Overactivity of the amygdala may correlate with depressive, ruminative thoughts and may be present in the early stages of MDD. Morphological and functional abnormalities within the amygdala in individuals with MDD have also been consistently demonstrated. As it is possible that the decreased functional connectivity of rPFC–amygdala circuitry detected in our MDD group may relate to both executive and emotional dysfunction, future neuroimaging studies using tasks that investigate these functions would be of interest.

The decreased amygdala–rPFC functional connectivity in individuals with MDD detected in our study could reflect a reduction in the PFC’s inhibitory control over the amygdala and could delay the extinction of negative emotion. A study by Petrides and Pandya in the macaque monkey demonstrated that the rPFC has connections with the amygdala, ventral PFC and cingulate cortex, all areas that have been linked with emotion processing. Recently, an fMRI study reported that correct recall of negative faces was associated with increased activity in the amygdala and rPFC, indicating that both areas are intimately involved in the retrieval of emotional stimuli during recognition memory. In the present study, we used the task inducing implicit rather than explicit processing of emotional faces, suggesting that altered functional connectivity between the amygdala and rPFC might play a key role during implicit emotional processing in individuals with MDD and that this dysfunction might contribute to negative automatic thoughts in patients with MDD.

We found abnormality in the functional connectivity to the left, but not to the right, rPFC. The decreased amygdala–left rPFC functional connectivity in response to fearful emotion may be related to the hemispheric asymmetry, which has been demonstrated in individuals with MDD. The balance between the right and left hemispheres is very important to adaptive emotion regulation, and hemispheric asymmetry has been observed in normal affective processing of positive and negative emotions. Previous studies have indicated that the left PFC activates more during the regulation of negative affect and approach-related positive affect. Studies involving depressed individuals have shown decreased activation in the left PFC and increased activation in the right PFC in response to sad mood induction. Our results indicated that the right rPFC might contribute more than the right rPFC to the disturbed emotional processing in individuals with MDD and that abnormalities in hemispheric asymmetry might be related to the pathophysiology of MDD.

Consistent with the previous behavioural, electroencephalogram and functional activation findings of dysfunctional negative emotional processing in individuals with MDD, our finding of decreased amygdala–PFC functional connectivity during fearful face processing and not during happy or neutral face processing supports the involvement of abnormalities in amygdala–PFC functional connectivity in negative emotional processing in individuals with MDD, and it further implicates abnormal negative emotional processing in the neuropathophysiology of MDD. Taken together with the findings of our recent report of

Table 2: Z scores of functional connectivity in rostral prefrontal cortex with 3 conditions

<table>
<thead>
<tr>
<th>Condition</th>
<th>Group; mean ± SD*</th>
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<tbody>
<tr>
<td>MDD, n = 28</td>
<td>Control, n = 30</td>
</tr>
<tr>
<td>Fear</td>
<td>0.36 ± 0.33</td>
</tr>
<tr>
<td>Happiness</td>
<td>0.47 ± 0.27</td>
</tr>
<tr>
<td>Neutral</td>
<td>0.38 ± 0.46</td>
</tr>
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MDD = major depressive disorder; SD = standard deviation.
*Unless otherwise indicated.
decreased amygdala–PFC functional connectivity during a resting-state fMRI in individuals with MDD, our current findings suggest that altered amygdala–PFC functional connectivity may be a key feature of the disorder. These studies also raise interesting questions about the association between amygdala–PFC functional connectivity during the resting state and tasks in individuals with MDD. Activation findings from both resting-state and task-related fMRI suggest that intrinsic resting-state activity may be involved in specific brain circuit engagement to perform a cognitive task and that resting activity can predict subsequent task-evoked brain responses in healthy individuals. To our knowledge, no studies have investigated the association of the functional connectivity within a specific neural circuitry during resting-state and task-related fMRI in individuals with MDD. Unfortunately, we were not able to examine this association within the amygdala–PFC circuitry in this study because of the differences in study design between the present task-related fMRI study and our previously published resting-state fMRI study, such as seed ROI selection (the bilateral amygdala was selected as a single ROI in the present study, whereas the left and right amygdala were selected as 2 separate ROIs in our previous resting-state study). We speculate that altered amygdala–PFC functional connectivity during the resting state might contribute to decreased amygdala–PFC functional connectivity during fearful face processing in individuals with MDD, and that altered functional connectivity during the resting state and negative emotional processing tasks reflect the dysfunctional negative emotional processing observed in individuals with MDD. Future studies with careful design to examine the association of functional connectivity within certain neural circuits during the resting state and emotional face processing tasks in individuals with MDD will be important in understanding the neuropathophysiology of MDD. Furthermore, the association between these functional connectivity disturbances and structural connectivity could be explored using diffusion tensor imaging. The connections between the amygdala and rPFC are bidirectional, but our study did not assess the direction of the functional connectivity. Since individuals were acutely ill at the time of study, we cannot determine whether the abnormality is state- or trait-related or whether successful treatment alters this disturbance.

Limitations

All participants with MDD were already experiencing acute episodes, therefore the findings could not be generalized to the entire MDD population. Future studies of participants in the euthymic state or individuals at risk for MDD are warranted to elucidate the amygdala–left rPF C functional connectivity abnormalities associated with MDD neuropathophysiology or with predisposition to MDD. Furthermore, participants with MDD in the present study had high HARS scores. Although no association between HARS scores and amygdala–rPFC functional connectivity was detected, the abnormalities of functional connectivity within different anxiety levels in individuals with MDD need to be further investigated. In addition, only fear was used as a negative emotion in the study; future studies including more negative emotions, such as sadness, anger and disgust, which have been used in some fMRI studies, are needed to fully understand the neuropathophysiology of MDD.

Conclusion

Our study provides critical evidence supporting abnormalities of amygdala–left rPF C functional connectivity in response to negative emotion in individuals with MDD. The findings in medication-naive individuals with this disorder suggest that they may play an important role in the pathophysiology of MDD.

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Contributors: Y. Tang, G. Fan, H.P. Blumberg, K. Xu and F. Wang designed the study. L. Kong, K. Chen, F. Wu, G. Fan, L. Ren, W. Jiang and Y. Cao acquired the data, which L. Kong, N. Driesen, F. Womer, W. Jiang, and F. Wang analyzed. L. Kong, K. Chen, Y. Tang, N. Driesen, F. Womer, H.P. Blumberg, K. Xu and F. Wang wrote the article. All authors reviewed the article and provided approval for publication.

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