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## Emotion-dependent functional connectivity of the default mode network in adolescent depression

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### Abstract

**Background**—Functional magnetic resonance imaging (fMRI) research suggests that both adult and adolescent major depressive disorder (MDD) is marked by aberrant connectivity of the default mode network (DMN) during resting-state. However, emotional dysregulation is also a key feature of MDD. No studies to date have examined emotion-related DMN pathology in adolescent depression. Comprehensively understanding the dynamics of DMN connectivity across brain states in depressed individuals with short disease histories could provide insight into the etiology of MDD.

**Methods**—We collected fMRI data during an emotion identification task and also during resting-state from 26 medication-free adolescents (13-17 years) with MDD and 37 wellmatched healthy

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controls (HCL). We examined between-group differences in blood oxygenation level-dependent task responses, emotion-dependent, and resting-state functional connectivity of the two primary nodes of the DMN: medial prefrontal cortex (mPFC) and posterior cingulate cortex (PCC). Additionally, we examined between-group differences in DMN functional connectivity and its relationship to depression severity.

**Results**—Relative to HCL, unmedicated MDD adolescents demonstrated reduced mPFC and PCC emotion-related deactivation and greater mPFC and PCC emotion-dependent functional connectivity with precuneus, cingulate gyrus, and striatum/subcallosal cingulate gyrus. Importantly, PCC-subcallosal cingulate connectivity remained inflexibly elevated in MDD versus HCL during resting-state. Lastly, stronger PCC emotion-dependent functional connectivity was associated with greater depression severity and an earlier age of depression onset.

**Conclusions**—Adolescent depression is associated with inflexibly elevated DMN connections. Given recent evidence of DMN maturation throughout adolescence, our findings suggest that early-onset depression adversely impacts normal development of functional brain networks.

### Keywords

functional magnetic resonance imaging; major depressive disorder; default mode network; functional connectivity; psychophysiological interaction; adolescence

## Introduction

The incidence of depression rises alarmingly during adolescence (1, 2), affecting 4-17% of youth worldwide (3-5). Adolescent-onset depression is over four times more likely to lead to a recurrent episode than adult-onset depression (6), and is associated with greater symptom severity, likelihood of relapse, and suicidality across the lifespan (7-9). Despite its clinical significance, relatively few studies have examined the neurobiological underpinnings of depression during this sensitive period of development. The adolescent brain, and especially the circuits supporting emotional processing, continues to undergo neural pruning and maturation (10-12). Thus, investigating the neurobiological mechanisms of emotional processing early in the disease course is critical for elucidating the etiology of depression.

Functional connectivity methods in neuroimaging have led to examinations of intrinsic brain networks in depression and other disorders, in particular the default mode network (DMN) (13-18). The DMN is a network of functionally connected brain regions whose activity becomes synchronously elevated during undirected or internally-directed mental states (e.g., passively resting, recalling the past, imagining the future) but dampens during goal-directed cognition (19). Most studies suggest that the DMN is further comprised of interconnected subsystems that converge on key nodes, with the two most notable being the medial prefrontal cortex (mPFC) and the posterior cingulate cortex (PCC) (19-22). Activity in the mPFC is believed to support self-referential mentation (21, 23-25), while the PCC is a highly connective area that is heavily implicated in memory-related processes, consciousness and awareness (26, 27).

Both the anterior and posterior hubs of the DMN have been linked to social and affective cognition and are anatomically connected to regions involved in emotion generation,

salience monitoring, and self-inspection (16, 21, 23, 28). Aberrant DMN function could therefore lead to dysfunctional self-referential and affective processing in the form of excessively negative self-focus (16, 17, 24). Indeed, altered resting-state connectivity has been demonstrated in depressed adults (29-44), adolescents (40, 45-51), and even children (52, 53). These studies tend to report altered, albeit directionally variable, functional coupling between nodes in the salience network and regions in the DMN in depressed individuals, with one study even reporting no significant differences in resting-state DMN functional connectivity between depressed and healthy adolescents (50). These inconsistencies, particularly in the adolescent literature, are compelling and highlight the need for more thorough examinations of the DMN in this population (51). Moreover, emotional dysregulation is also a core feature of MDD (54, 55). Thus, functional connectivity analyses of the DMN during a task that appropriately probes emotional processing are needed to target brain areas relevant to this feature of depression.

Presently, there are few studies examining the DMN during emotional processing in either depressed adults or adolescents. Ineffective suppression of the DMN during affective processing has been reported in depressed adults (31, 56), with greater mPFC and PCC activation correlating positively with feelings of depression and hopelessness (56). However, these studies did not examine functional connectivity. Prior studies examining emotion-dependent functional connectivity in depression have typically focused their analyses on nodes in the salience network, reporting diminished functional connectivity of the amygdala, anterior cingulate cortex, and mPFC in depressed versus healthy individuals (57-63). While this research suggests that MDD may be characterized by altered connectivity between the anterior hub of the DMN and structures involved in affective and salience processing (13, 16, 17), no work to date has examined both mPFC and PCC connectivity during emotional processing in depressed adolescents. To our knowledge, there are only two studies examining functional connectivity of emotional processing in depressed adolescents and they are limited in their sample sizes and focus only on regions involved in affective processing (57, 58). Furthermore, recent evidence shows that the DMN undergoes developmental changes (64-67). It is therefore imperative that DMN dynamics are investigated in early-onset depression free from chronic illness and pharmacological treatment so that the pathophysiology of this disease at the level of brain networks may be elucidated.

To this end, we recruited a relatively large sample of medication-free adolescents with major depressive disorder (MDD) and a healthy control (HCL) group that was well-matched on several demographic variables. Subjects underwent fMRI scanning during an emotion identification task and also during resting-state. Clusters in the mPFC and PCC exhibited significantly less emotion-related deactivation in MDD compared to HCL. Subsequently, context-dependent and resting-state functional connectivity analyses were conducted using these mPFC and PCC regions as seeds. Based on previous studies implicating aberrant functional connectivity in MDD, we predicted that depressed adolescents would exhibit altered emotion-dependent connectivity of the DMN and that the strength of these connections would correlate positively with depression severity (57-63). Given the role of the DMN in self-referential processing and its association with depression in adults, we hypothesized that emotion-dependent functional connectivity of the DMN would correlate

with clinical measures of rumination (29, 34, 36, 39). Finally, in light of growing evidence showing developmental changes of the DMN (64, 65), we also hypothesized that emotion dependent functional connectivity of the DMN may be associated with age of depression onset.

## Methods and Materials

### Subject recruitment

A total of 83 right-handed adolescents (13-17 years) were recruited for this study. Six subjects (3 MDD) were excluded due to a failure to record behavioral responses in the scanner. We applied rigorous motion outlier thresholds to the remaining subjects (see *Inclusion/exclusion criteria* in the supplement for more details), resulting in whole brain and emotion-dependent functional connectivity results for 63 subjects (26 MDD) and resting-state results for 57 subjects (23 MDD). Depressed adolescents were recruited from psychiatric and primary care clinics in San Diego; HCL were recruited from the same geographic area via e-mail, internet, or flyers. Adolescents from both genders and all ethnicities were allowed to participate and all subjects received financial compensation for their participation. All participating adolescents provided written informed assent and their parent(s)/legal guardian(s) provided written informed consent in accordance with the Declaration of Helsinki and the Institutional Review Boards at the University of California, San Diego (UCSD), UC San Francisco, Rady Children's Hospital in San Diego, and the County of San Diego.

### Diagnostic assessment of subjects

The Schedule for Affective Disorders and Schizophrenia for School-Age Children-Present and Lifetime Version (K-SADS-PL) (68) was administered to all potentially depressed subjects. The computerized Diagnostic Interview Schedule for Children 4.0 (69) and the Diagnostic Predictive Scale (69) were used to screen for the presence of any Axis I diagnoses in HCL.

Depression symptoms were measured with the Children's Depression Rating Scale-Revised (70) and the Beck Depression Inventory-II (BDI-II) (71), anxiety with the Multidimensional Anxiety Scale for Children (MASC) (72), and rumination with the Ruminative Responses Styles (RRS) inventory (73).

The groups were well-matched on age, gender distribution, ethnicity, pubertal status, socioeconomic status, and general intelligence. Details regarding diagnostic assessment, demographic measures, and inclusion/exclusion criteria are presented in the supplement. See Table 1 for a summary of the characteristics of the depressed subjects and Table S1 for a summary of their psychosocial treatment history.

### Experimental stimuli and paradigm

All subjects completed one run each of the emotion identification task and resting-state in the same scan session. Emotional face stimuli have been shown to deactivate regions of the DMN in healthy individuals (74). However, the only other functional connectivity study

using emotional faces to examine adolescent depression not only did not examine the DMN, but also tested only static fearful faces (58). Because dynamic face stimuli are considered more ecologically valid and more robustly activate affective processing regions compared to static face stimuli (75, 76), our emotion identification task employed dynamically morphing faces expressing fear, happiness, and sadness. We used ten standardized faces (5 female) expressing fear, happiness, and sadness (77) morphed with computer graphical manipulation (78, 79). On FACE trials, a screen displaying text of the possible emotions to discern (FEAR, HAPPY, SAD) was presented for 1500ms. Next, a neutral face morphed smoothly to an emotion of prototypical intensity over the span of 3000ms and remained onscreen for an additional 800 ms before the screen turned blank for 700ms, signifying the trial's end (Figure 1A). At stimulus onset, two possible emotion choices were displayed in the bottom left and right corners; subjects were instructed to press one of two buttons corresponding to their choice as soon as they recognized the facial emotion. OVAL trials (6 s per trial), where subjects had to determine if a morphing oval was tilting left or right (maximal tilt angle=10°), were used as a sensorimotor control (Figure 1B). At the end of the scan, a blank screen was presented for 10s. On run contained 80 trials (60 facial trials and 20 oval trials; total run=490s). Response time and accuracy on every trial during scanning was recorded. See also Figure 1 and the supplement.

### Image acquisition

All scanning was conducted on a General Electric 3T MR750 System (General Electric Healthcare, Milwaukee, WI) with Twin Speed gradients and a GE 8-channel head coil at the Center of Functional MRI at UCSD. See the supplement for more details.

### Image analysis

All image processing and analyses were conducted with AFNI (80) and FSL (81). The T1-weighted images were skull-stripped and transformed to MNI152 (Montreal Neurological Institute, Montreal, Quebec, Canada) with an affine transform (82, 83) followed by nonlinear refinement (84). EPI data were slice-time and motion corrected and then aligned to the T1-weighted images using a Localized Pearson Correlation function (85). Next, the EPI data were convolved with a 4.2 mm full-width half-maximum (FWHM) isotropic Gaussian filter and grand mean scaled before being transformed to MNI152 space at 3x3x3 mm resolution. A generalized least squares regression model that estimates the serial correlation of noise with an autoregressive moving average method was used to fit each voxel's time-series. Each stimulus type was included as a regressor-of-interest (FEAR, HAPPY, SAD, OVAL). Each time-series of interest spanned stimulus onset until the first valid (150 ms) response, thus providing a naturally jittered reference function, before being convolved with a gamma-variate function (86). Demeaned motion parameters (3 rotational and 3 translational) and a second order Legendre polynomial were included as nuisance regressors (i.e., baseline). Volumes where the Euclidean norm of the motion derivatives exceeded 0.2 or where more than 10% of voxels exceeded the median absolute deviation of the detrended time-series were censored. Subjects where more than 20% of their volumes were censored were removed from the final analysis (see *Inclusion/exclusion criteria* in the supplement for more details). A general linear test for FACE-OVAL was

computed for each participant. Brain activation was operationally defined as percentage signal change relative to baseline.

### Functional connectivity analyses

Context-dependent functional connectivity analyses were conducted according to prior studies utilizing facial stimuli from our laboratory (58) and others (87-89). Specifically, we employed a seed-based approach using the psychophysiological interaction (PPI) (90) method developed for AFNI (<http://afni.nimh.nih.gov/sscc/gangc/CD-CorrAna.html>), with clusters in the left mPFC and left PCC exhibiting significant between-group differences on the emotion identification task defined as distinct seeds (Figure 2). Details on our functional connectivity analyses can be found in the supplement under *Emotion-dependent functional connectivity (PPI) image analysis* and *Resting-state image analysis*. Additional exploratory PPI analyses were also performed to examine DMN functional connectivity to each emotion (see *Exploratory emotion-dependent functional connectivity analysis* and Tables S4-S6 in supplement).

### Between-group whole brain task analysis

Group differences on the emotion identification task were assessed using a linear mixed effects (LME) model on the estimates from the regression model, with group (MDD, HCL) modeled as a fixed factor and participant modeled as a random factor.

### Between-group emotion-dependent functional connectivity analysis

Group differences in the PPI analysis were assessed using a LME on the Fisher's z-scores of the interaction effect, with group modeled as a fixed factor and participant modeled as a random factor. A LME was run for each seed separately.

### Relating emotion-related activation of mPFC and PCC with elevated emotion-dependent functional connectivity

We correlated mean percentage signal change of emotion-related activation in the mPFC and PCC (i.e., the seeds in the PPI analysis) with the mean beta-values from each of the regions exhibiting significant between-group differences in the PPI analysis with that respective seed using two-tailed tests of the non-parametric Spearman's correlation coefficient ( $r_s$ ). For details, see the supplement.

### Controlling for multiple comparisons

We computed the minimum number of contiguous voxels passing the voxel-wise threshold for a significant group difference that would result in a cluster-wise  $p < 0.05$  corrected threshold at the whole brain level using Monte Carlo simulations: 51 voxels (1377  $\mu\text{L}$ ). For details on our simulation parameters, see the supplement.

### Comparison between emotion-dependent and resting-state functional connectivity

Resting-state functional connectivity values (i.e., Fisher's z-scores) in the regions showing significant between-group differences with either the mPFC or PCC seeds on the PPI analyses were extracted and subjected to a LME with group and brain-state (task-activated,

resting-state) modeled as fixed factors, participant modeled as a random factor, and with the Satterthwaite approximation for degrees of freedom.

### **Correlations between clinical variables and emotion-dependent functional connectivity**

Statistical analyses of all demographic and clinical scales were computed with R (91) and Matlab (Natick, MA). Within the MDD group only, the relationships between CDRS-R, BDI-II, RRS, MASC, and the mean Fisher's z-score within each of the significant regions identified by the between-group PPI analysis were examined with two-tailed tests of Pearson's correlation coefficient ( $r$ ).

## **Results**

### **Sociodemographic, clinical, and behavioral data**

The groups did not differ on age, gender distribution, ethnicity, pubertal status, socioeconomic status, general intelligence, or behavioral performance on the emotion identification task (Table 1, all  $p$ 's>0.30), but differed significantly as expected on depression, anxiety, and rumination measures (Table 1, all  $p$ 's<0.001).

### **Between-group whole brain task results**

The MDD group showed less deactivation in the left mPFC and left PCC (Figure 2) relative to HCL during emotional processing (see also supplement, Figure S1, and Table S2). Emotion-related activation in the mPFC and PCC showed no significant differences between depressed subjects who were undergoing psychosocial therapy at the time of scanning versus those who were not (all  $p$ 's>0.59; see the supplement and Table S7 for more details).

### **Between-group emotion-dependent functional connectivity results**

We seeded the left mPFC and PCC clusters that exhibited significant between-group differences on the emotion identification task in a seed-based PPI analysis (Figure 2). These same seeds were also used in the resting-state functional connectivity analyses. MDD exhibited greater mPFC emotion-dependent functional connectivity with bilateral precuneus (BA 7), cingulate gyrus (BA 23/24), and left inferior parietal lobule/supramarginal gyrus compared to HCL (Figure 3, Table 2). MDD also exhibited greater PCC emotion-dependent functional connectivity with bilateral precuneus (BA 7), right cingulate gyrus (BA 23/24), and left lentiform nucleus/subcallosal cingulate gyrus compared to HCL (Figure 4, Table 3). These areas showed no differences in emotion-dependent functional connectivity between depressed subjects who were undergoing psychosocial therapy at the time of scanning versus those who were not (all  $p$ 's>0.27; see the supplement and Table S6 for more details).

### **Relating emotion-related activation of mPFC and PCC with elevated emotion-dependent functional connectivity**

Greater emotion-related activation in the PCC correlated significantly with stronger PCC emotion-dependent functional connectivity with dorsal striatum/subcallosal cingulate gyrus ( $r_s=0.271$ ,  $p=0.032$ ). Detailed methods and results can be found in the supplement.

### Comparison between emotion-dependent and resting-state functional connectivity

We compared mPFC and PCC-based functional connectivity in areas showing between-group differences from our PPI analysis across brain-states (task-activated, resting-state). We found that PCC functional connectivity with the dorsal striatum/subcallosal cingulate gyrus exhibited a significant interaction between group and brain-state ( $F_{1,54}=5.410$ ,  $p=0.023$ ; Figure 3; Table 3). Post-hoc  $t$ -tests (Fisher's LSD) further revealed that this difference was driven by reduced functional connectivity during the emotion identification task in HCL. While all other areas displayed a significant effect of brain-state (resting-state > task activated, all  $p$ 's<0.01), no other areas showed a significant main effect of group nor a significant interaction between group and brain-state in resting-state functional connectivity (all  $p$ 's>0.05).

### Correlations between clinical variables and emotion-dependent functional connectivity

All correlations were performed between the mean Fisher's  $z$ -scores of every significant cluster arising from the between-group PPI analysis for each DMN seed with levels of depression, rumination, anxiety, and age of depression onset. CDRS-R scores correlated positively with mPFC-cingulate functional connectivity ( $r=0.483$ ,  $p=0.014$ ; Figure S2A) and PCC-cingulate functional connectivity ( $r=0.436$ ,  $p=0.029$ ; Figure S2B). Functional connectivity between the PCC seed and dorsal striatum/subcallosal cingulate gyrus correlated negatively with age of onset ( $r=-0.527$ ,  $p=0.014$ ; Figure S2C). No functional connections showed significant associations with RRS or MASC scores (all  $p$ 's>0.05), suggesting our results are selective to depression.

## Discussion

Examining the dynamics of intrinsic functional networks associated with emotional dysregulation in early-onset depression is critical for elucidating the neurobiological mechanisms of MDD. Here, we examined functional connectivity of one of the major intrinsic brain networks, the default mode network (DMN), during active emotional processing and also during resting-state in a relatively large sample of unmedicated adolescents with MDD and well-matched healthy controls. We hypothesized that depressed adolescents would exhibit altered emotion-dependent connectivity of the DMN compared to healthy controls and that the strength of these aberrant connections would correlate with clinical measures of depression. Our investigation yielded three primary results. First, depressed adolescents exhibited relatively greater mPFC and PCC functional connectivity with areas involved in default mode (precuneus, posterior cingulate gyrus), cognitive executive (middle cingulate gyrus, inferior parietal lobule), and salience and affective processing (dorsal striatum/subcallosal cingulate gyrus). Second, functional connectivity between PCC and dorsal striatum/subcallosal cingulate gyrus remained inflexibly elevated in the depressed group during resting-state. Lastly, stronger PCC emotion-dependent functional connectivity was associated with greater depression severity and an earlier age of depression onset. Taken together, these findings advance a dynamic network model of depression, whereby depressed adolescents exhibit altered intrinsic functional connections that remain inflexibly elevated.



The finding of reduced deactivation in the DMN during emotional processing in depressed adults (24, 56, 92) was replicated in our adolescent sample (Figure 2), providing additional validation of DMN dysfunction in depressive illness and extending the relevance of DMN pathology to adolescent depression. Importantly, the same mPFC and PCC areas exhibiting reduced emotion-related deactivation in MDD coactivated with several regions during affective challenge (Figures 3-4, Tables 3-4). Specifically, depressed adolescents showed comparatively elevated mPFC and PCC functional coupling with default mode regions involved in self-referential processing (precuneus, posterior cingulate gyrus) (19, 25, 93), areas in the central executive network governing attentional selection of external stimuli (middle cingulate gyrus, inferior parietal lobule) (94-103), and affective processing structures best associated with the salience network (subcallosal cingulate gyrus/dorsal striatum) (31, 35, 40, 104-118). In healthy individuals, the DMN is anticorrelated with the central executive and salience networks (119, 120). However, our results demonstrate that these functional connections coactivate strongly together during emotional processing in adolescent depression.

Our emotion-dependent functional connectivity results may appear upon initial examination to differ from other recent studies of depression also employing seed-based psychophysiological interaction (PPI) analyses. Specifically, studies in medication-naïve adults (63) and adolescents (57) with depression reported decreased connectivity between mPFC and amygdala during emotional processing in depressed versus healthy individuals. Similarly, in a PPI study of unmedicated adolescents (58), depressed individuals showed reduced functional connectivity between the subcallosal cingulate area and PCC when evaluating fearful faces (58). However, all of these studies used static images and faces; dynamic face stimuli have been shown to be more ecologically valid than static faces and more robustly activate affective processing regions (75). Our present study uses dynamically morphing faces, thereby representing a more optimal functional assay to probe the neurocircuitry of emotional dysregulation. Additionally, none of the aforementioned studies explicitly examined both the mPFC and PCC. Critically, none of these studies defined their seeds functionally (i.e., they were derived from atlases or based on previously published coordinates). While hypothesis-driven, such a method may be suboptimal when trying to specifically target areas underlying behavioral and cognitive symptoms related to a disease state—namely, emotional dysregulation in depression.

Prior resting-state (37, 40, 46, 49, 50, 121, 122) and PPI studies (58, 59, 62, 63) have suggested that MDD is characterized by alterations within and between intrinsic functional brain networks. Our previously described functional connectivity results support this idea, but our finding that depressed adolescents failed to show brain-state (task-activated versus resting-state) dependent changes in PCC connectivity with the subcallosal cingulate gyrus (Figure 4) also suggests that depression may be characterized by inflexible functional connections. These findings corroborate a recent study in adolescents and young adults (15-24 years) that compared functional connectivity of the subcallosal cingulate area during a cognitive control task and resting-state, and reported inflexible functional connectivity between the subcallosal cingulate area and ventral striatum in the depressed group (107). By targeting emotionally-selective areas of the DMN and comparing DMN patterns across brain-states, our results build from the extant literature to advance a network model of

depression characterized by both altered and inflexible connections of intrinsic brain networks.

Consistent with prior reports of significant associations between clinical measures of depression and resting-state DMN connectivity (40, 48, 49) we found that greater DMN functional connectivity during emotional processing correlated positively with depression severity (Figure S2A and S2B). Contrary to our hypothesis and findings in resting-state adult studies (29, 33, 36, 39), we did not find significant associations with rumination. Rumination as measured by the RRS may not capture negative self-focus as it pertains to adolescent depression (123), or it may reflect more general negative affect (124, 125). Given that emotion-dependent functional connectivity also did not correlate significantly with anxiety suggests that our results are specific to adolescent depression. Supporting this notion is our finding that stronger PCC connectivity during emotional processing was significantly associated with an earlier age of depression onset (Figure S2C). Connectivity between the PCC and parahippocampal cortex during recollection of negative life events has also been shown to increase with the number of depressive episodes, as well as the frequency and subsequent prediction of sad mood in daily life (128). Individuals with an increased genetic risk for developing depression also exhibit altered PCC connectivity with the amygdala during emotional regulation—but not during resting-state (129). These findings in concert with our results suggest that stronger emotion-dependent functional connections between the PCC and limbic regions may be a contributing mechanism to the development of depression. Another interpretation is that depression may impact the normal development of intrinsic brain networks through the manifestation of inflexible connections. The DMN follows a developmental trajectory in humans wherein the coherence between the mPFC and PCC subsystems is initially weak but solidifies by adolescence (64-66). Recent work has even suggested developmental changes selective to anticorrelated networks based in the mPFC during both resting-state (66) and a social emotion task (130). These results indicate that DMN maturation is vulnerable to experiences such as early life stressors and other risk factors associated with the development of depression (131). Under this framework, our findings suggest that early-onset depression adversely impacts normal maturation of intrinsic brain networks.

Although this study is the first to compare emotion-dependent and resting-state functional connectivity in adolescent depression, our results must be interpreted in light of their limitations. Future studies with larger samples are needed to replicate our results and different probes of affective processing are needed to assess the generalizability of our findings. The impact of depression on brain maturation remains a critical research question; future work examining dynamic brain-state patterns in participants spanning a wider age range may yield clearer answers. Finally, longitudinal research is needed to address the limitations of our cross-sectional design and to determine whether the aberrant dynamic DMN patterns we report are an enduring trait of MDD or whether they resolve with successful treatment.

In summary, the findings reported here are notable in several respects. First, no other study has investigated functional connectivity dynamics of the DMN in adolescent depression. Second, our finding of inflexibly elevated DMN functional connectivity in our depressed

sample may reflect core features of depressive illness (e.g., emotional dysregulation) and thus, represent a candidate biomarker for evaluating treatment outcomes. Lastly, our unique study population of medication-free depressed adolescents offers potential insight regarding the influence of depression on normal brain development. Nevertheless, the neurobiological mechanisms by which these alterations occur and the optimal interventions needed requires additional study.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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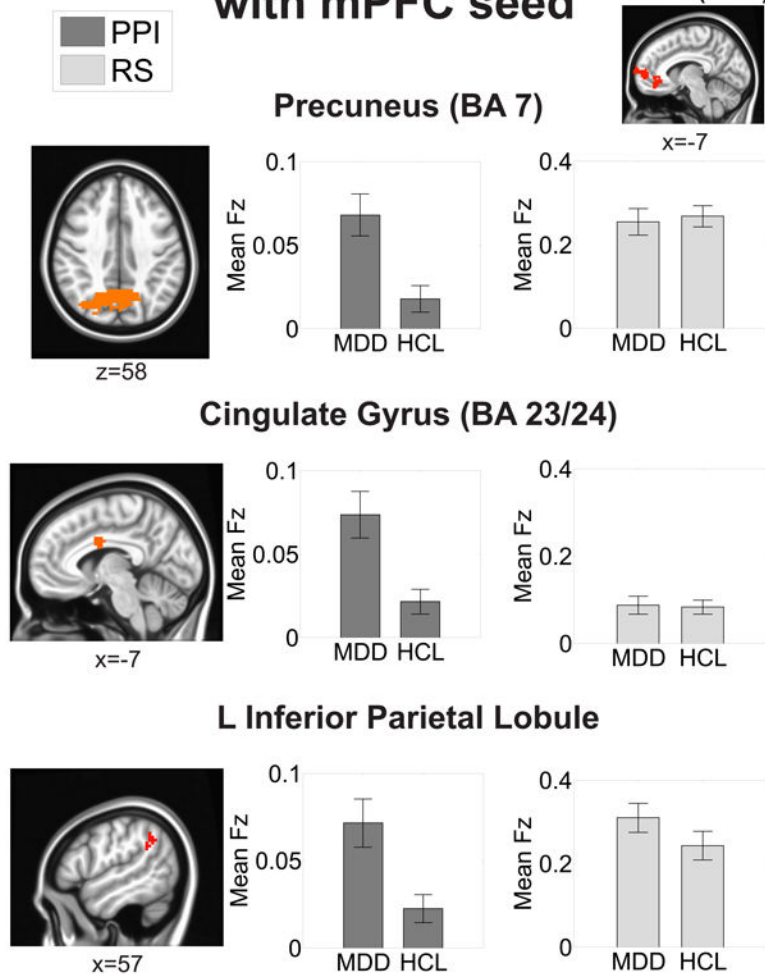
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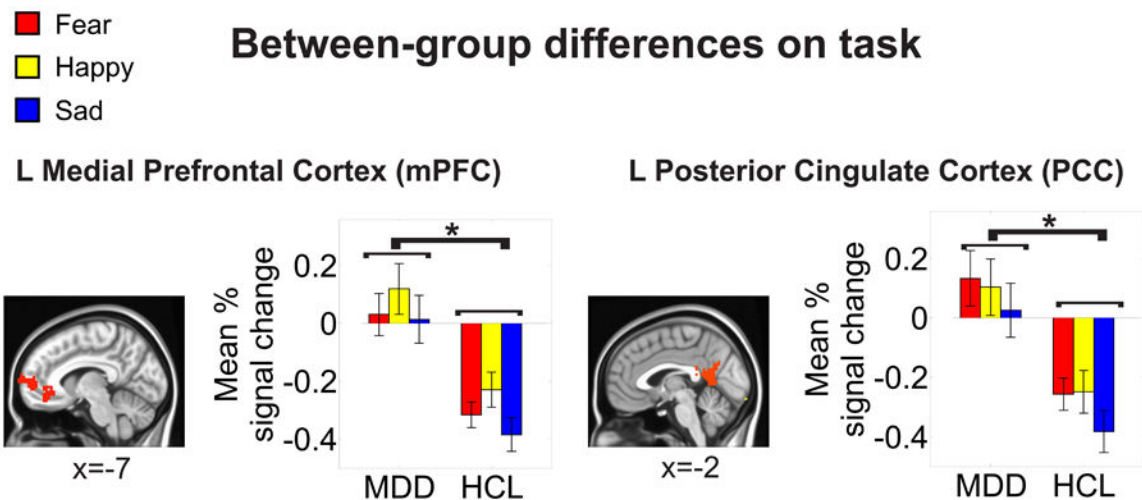
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## Between-group differences in PPI with mPFC seed

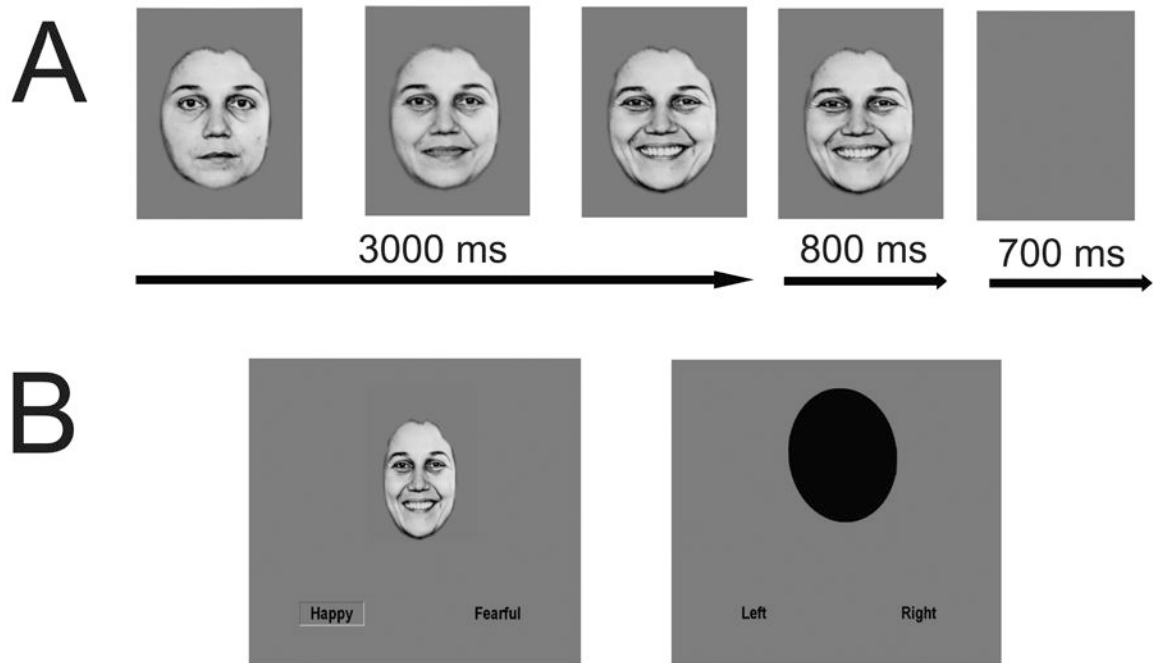


### Figure 1. Emotion identification task with dynamically morphing face stimuli

On FACE trials, a screen displaying text of the three possible emotions to discern (FEAR, HAPPY, SAD) was presented for 1500 ms. Next, a neutral face morphed smoothly and dynamically to an emotion of prototypical intensity over the span of 3000 ms. At maximal emotion intensity, the face then remained on the screen for an additional 800 ms of the trial before the screen turned blank for 700 ms (**Figure 1A** depicts an example face and is enlarged for purposes of clarity). At stimulus onset, two possible emotion choices were displayed in text on the bottom left and right corners; subjects were instructed to press one of two buttons corresponding to the displayed emotion as soon as they recognized the emotion of the face. As a sensorimotor control, we also had OVAL trials (6 s per trial), where subjects had to determine if the top of an oval was tilting to the left or right and make a button response accordingly as soon as they recognized the tilt direction (**Figure 1B** depicts a sample FACE and OVAL trial). See the supplement for more details.



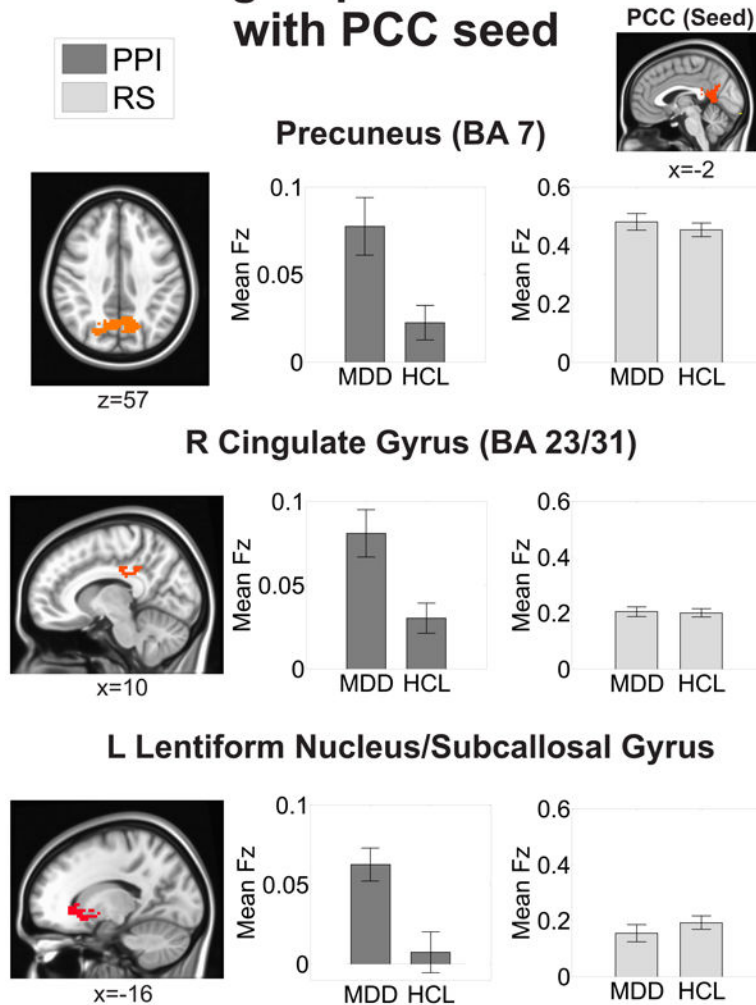
**Figure 2.** Mean activation to each emotion type in the mPFC and PCC regions (i.e., seeds for the PPI analysis) showing significant between-group differences on the task (FACE-OVAL). Mean percentage signal change to each condition (FEAR-OVAL, HAPPY-OVAL, SAD-OVAL) was extracted and subjected to post-hoc *t*-tests (Fisher's LSD). These post-hoc *t*-tests indicated a main effect of group at  $p < 0.05$  (denoted by \*). These areas survived correction for multiple comparisons at a cluster-wise threshold of  $p < 0.05$ . Locations are reported in MNI coordinates (radiological convention). See Figure S1 and Table S2 for a full summary of the between-group differences on the task.



**Figure 3. Regions showing significant between-group differences in emotion-dependent functional connectivity of the mPFC seed**

All areas survived correction for multiple comparisons at a cluster-wise threshold of  $p < 0.05$ . Mean functional connectivity values are reported as Fisher's z-scores (Fz). The mean resting-state (RS) Fz were extracted from the regions exhibiting significant between-group differences on the psychophysiological interaction (PPI) analysis (see *Methods* for more details). Locations are reported in MNI coordinates (radiological convention). See also Table 2.

## Between-group differences in PPI with PCC seed



**Figure 4. Regions showing significant between-group differences in emotion-dependent functional connectivity with the PCC seed**

All areas survived correction for multiple comparisons at a cluster-wise threshold of  $p < 0.05$ . Mean functional connectivity values are reported as Fisher's z-scores (Fz). The mean resting-state (RS) Fz were extracted from the regions exhibiting significant between-group differences on the psychophysiological interaction (PPI) analysis (see *Methods* for more details). Locations are reported in MNI coordinates (radiological convention). See also Table 3.

**Table 1**  
**Summary of subject groups' demographic, clinical characteristics, and behavioral performance of task**

Entries are of the form: mean  $\pm$  SEM. Statistical analyses of between-group differences were conducted with chi-squared tests ( $\chi^2$ ), Student t-tests ( $t$ ), and Wilcoxon Rank Sum test ( $U$ ). All p-values indicate two-tailed statistical significance levels. Details on the scales used can be found in the supplement. A summary of the psychosocial treatment history of our depressed subjects can be found in Table S1. All depressed subjects were naïve to antidepressants at the time of scanning except for two: one had last been exposed 4 months prior to their scan and one had been last exposed 4 years prior to their scan.

Characteristic	Depressed	Control	Statistic	P-value
Number of participants in final analysis	26	37		
Gender (M / F)	7 / 19	14 / 23	$\chi^2=0.819$	0.366
Age at time of scan (years)	16.1 $\pm$ 0.3	16.0 $\pm$ 0.2	$t_{61}=0.29$	0.77
Ethnicity (Asian / Black / Caucasian / Hispanic / Mixed)	3 / 3 / 8 / 10 / 2	4 / 2 / 13 / 14 / 4	$U=9$	0.458
Hollingshead Socioeconomic Score	40 $\pm$ 25.2	29 $\pm$ 16.3	$U=432$	0.36
Tanner Score	4.2 $\pm$ 0.4	4.0 $\pm$ 0.7	$U=509$	0.90
Wechsler Abbreviated Scale of Intelligence (Standardized)	104.2 $\pm$ 4.6	107.3 $\pm$ 3.3	$t_{61}=0.56$	0.58
Beck's Depression Inventory II	28.4 $\pm$ 2.0	3.4 $\pm$ 0.7	$t_{60}=13.10$	<0.0001
Children's Depression Rating Scale-Revised (Standardized)	73.1 $\pm$ 1.8	34.3 $\pm$ 1.2	$t_{61}=18.64$	<0.0001
Multidimensional Anxiety Scale for Children (Standardized)	59.8 $\pm$ 1.8	42.1 $\pm$ 1.4	$t_{60}=7.88$	<0.0001
Ruminative Responses Scale	67.6 $\pm$ 3.3	33.7 $\pm$ 1.7	$t_{51}=9.81$	<0.0001
Age of depression onset (years)	12.28 $\pm$ 0.56			
<b>Comorbidities</b>	<b><i>n</i> (%)</b>			
No comorbidities	9 (34.62%)			
Generalized Anxiety Disorder (GAD)	7 (26.92%)			
Post Traumatic Stress Disorder (PTSD)	3 (11.5%)			
Specific Phobia	2 (7.7%)			
GAD + PTSD	1 (3.8%)			
Anxiety Disorder Not Otherwise Specified (NOS)	1 (3.8%)			
Attention Deficit Hyperactivity Disorder NOS	1 (3.8%)			
Social Phobia	2 (7.7%)			
<b>Behavioral performance on emotion identification task</b>				
Accuracy (%)	80.73 $\pm$ 6.44	80.06 $\pm$ 9.73	$t_{61}=0.053$	0.958
Response Time (s)	2.25 $\pm$ 0.52	2.16 $\pm$ 0.59	$t_{61}=0.110$	0.913

**Table 2**  
**Summary of significant clusters from the between-group emotion-dependent functional connectivity analysis of mPFC**

Regions are the result of a psychophysiological interaction (PPI) method of functional connectivity and corrected for multiple comparisons at a cluster-wise threshold of  $p < 0.05$ . Locations are reported according to the center of mass of cluster in MNI coordinates (radiological convention). Functional connectivity values are reported as mean  $\pm$  SEM. Resting-state (RS) functional connectivity values were extracted from the regions exhibiting significant between-group differences on the PPI analysis (see *Methods* for more details). See also Figure 3.

Region (Brodmann's Area)	Location (x,y,z)	# of voxels	Statistic( $F_{1,61}$ )	MDD (PPI)	HCL (PPI)	MDD(RS)	HCL (RS)
<b>Directionality: MDD &gt; HCL</b>							
Precuneus (BA 7)	8, -67, 38	321	6.39	0.068 $\pm$ 0.01	0.018 $\pm$ 0.008	0.255 $\pm$ 0.03	0.268 $\pm$ 0.03
Cingulate gyrus (BA23/24)	-6, -1, 30	176	6.30	0.073 $\pm$ 0.01	0.021 $\pm$ 0.007	0.087 $\pm$ 0.02	0.083 $\pm$ 0.02
L inferior parietal lobule, supramarginal gyrus	-50, -52, 34	72	5.43	0.071 $\pm$ 0.01	0.022 $\pm$ 0.008	0.310 $\pm$ 0.03	0.242 $\pm$ 0.03



**Table 3**  
**Summary of significant clusters from the between-group emotion-dependent functional connectivity analysis of PCC**

Regions are the result of a psychophysiological (PPI) method of functional connectivity and corrected for multiple comparisons at a cluster-wise threshold of  $p < 0.05$ . Locations are reported according to the center of mass of cluster in MNI coordinates (radiological convention). Functional connectivity values are reported as mean  $\pm$  SEM. Resting-state (RS) functional connectivity values were extracted from the regions exhibiting significant between-group differences on the PPI analysis (see *Methods* for more details). See also Figure 4.

Region(Brodmann's Area)	Location (x,y,z)	# of voxels	Statistic( $F_{1,61}$ )	MDD (PPI)	HCL (PPI)	MDD (RS)	HCL (RS)
<b>Directionality: MDD &gt; HCL</b>							
Precuneus (BA 7)	5, -65, 37	195	6.25	0.078 $\pm$ 0.02	0.023 $\pm$ 0.009	0.481 $\pm$ 0.03	0.454 $\pm$ 0.02
R cingulate gyrus (BA 31/23)	14, -26, 32	179	5.42	0.081 $\pm$ 0.01	0.030 $\pm$ 0.008	0.205 $\pm$ 0.02	0.200 $\pm$ 0.01
L dorsal striatum, subcallosal cingulate gyrus	-17, 17, -4	63	5.49	0.063 $\pm$ 0.01	0.008 $\pm$ 0.01	0.155 $\pm$ 0.03	0.193 $\pm$ 0.02