Negative mood-induction modulates default mode network resting-state functional connectivity in chronic depression

Citation for published version (APA):

Document status and date:
Published: 15/01/2017

DOI:
10.1016/j.jad.2016.10.022

Document Version:
Publisher's PDF, also known as Version of record

Document license:
Taverne

Please check the document version of this publication:

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Research paper

Negative mood-induction modulates default mode network resting-state functional connectivity in chronic depression

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\textbf{ARTICLE INFO}

\textbf{Keywords:}
Mood-induction
Chronic depression
Resting state fMRI
Default mode network
Posterior cingulate cortex

\textbf{ABSTRACT}

\textbf{Background:} The aim of this study was to investigate the effects of sad mood on default mode network (DMN) resting-state connectivity in persons with chronic major depressive disorder (cMDD).

\textbf{Methods:} Participants with a diagnosis of cMDD (n=18) and age, gender and education level matched participants without a diagnosis of depression (n=18) underwent a resting-state fMRI scan, before and after a sad mood induction. The posterior cingulate cortex (PCC) was used as a seed for DMN functional connectivity across the two resting-state measurements.

\textbf{Results:} Mood ratings decreased in both groups following the sad mood induction procedure. PCC connectivity with the parahippocampal gyrus, the superior temporal gyrus and the anterior inferior temporal cortex increased in cMDD patients following the sad mood induction, whereas it decreased in non-patient controls. PCC connectivity with the anterior prefrontal cortex and the precuneus decreased in cMDD patients following the sad mood induction, whereas it increased in non-patient controls.

\textbf{Limitations:} Limitations of this study include the relatively small sample size and lack of a clinical control group.

\textbf{Conclusions:} These findings are in line with neurobiological models of depression suggesting that the observed changes in DMN connectivity following the sad mood induction might reflect a failure to exert cognitive control over memory retrieval in patients with cMDD.

1. Introduction

Major depressive disorder (MDD) is a relatively common, psychiatric disorder that often takes a chronic course [1]. Recent efforts to understand the neuropathology of depression have focused on connectivity within or between brain networks that can be identified during rest for reviews see e.g.: [2,3]. The default mode network (DMN), a brain network which has well-documented associations with self-referential processing [4], rumination [3,5] and autobiographic memory [6], plays a key role in the neuropathology of MDD [7]. Numerous studies have now demonstrated that patients with MDD show increased DMN resting-state functional connectivity, compared to non-patient controls [2,8].

Although the finding of increased connectivity in the DMN of MDD patients is fairly consistent [2], connectivity within this network might not necessarily be stable. For example, a number of studies have shown that DMN connectivity in depression changes when symptoms improve following treatment with antidepressants [9]. Moreover, changes in mood states might also drive network dynamics. For example, studies in non-depressed [10] and remitted depressed [11] participants have shown that DMN connectivity can be modulated by a sad mood induction. Harrison et al. (2008) used a combination of autobiographical recall and classical music to induce a neutral (scan 1) or sad (scan 2) mood state in healthy participants. The authors found that the

\textbf{http://dx.doi.org/10.1016/j.jad.2016.10.022}

Received 18 December 2015; Received in revised form 6 September 2016; Accepted 21 October 2016
Available online 23 October 2016

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strength of DMN connectivity decreased in the sad vs. neutral mood induction condition [10]. Zamoscik et al. (2014) used a combination of autobiographical recall and classical music to induce sad mood in individuals with remitted depression and healthy controls. The authors found that individuals with remitted depression showed greater connectivity between the PCC and parahippocampal gyri following the sad mood induction compared to healthy controls [11].

Whereas these previous studies reveal important mechanisms as to how DMN connectivity can be modulated by a sad mood induction in healthy or previously depressed individuals, it is unclear how patients with a current MDD modulate connectivity in the DMN following a sad mood induction. Given that connectivity within the DMN is positively correlated with depressive chronicity [12], it is intriguing to study DMN connectivity following a sad mood induction in patients with chronic major depressive disorder (cMDD). Therefore, the aim of the present study was to investigate resting state connectivity of the DMN in patients with a current cMDD and matched non-patient controls before and after a sad mood induction.

Harrison et al. (2008) instructed participants to actively visualize and ruminate about the event recalled. The authors argued that the sad mood-induction might require more cognitive effort from their healthy participants whereas neutral recall might require less cognitive effort. Given previous findings of task induced deactivation of the DMN they expected that the sad vs neutral mood induction would lead to reduced DMN connectivity [10]. However, DMN connectivity in individuals with depression, compared to non-depressed individuals, is generally increased and immersing in a sad mood state might be less effortful for individuals with depression compared to non-depressed individuals. Moreover, the sad-mood induction that was used in the current study was conducted prior to scanning and did not require participants to engage in a task during scanning. Finally, as has been argued by Zamoscik et al. (2014) a negative mood induction can trigger increased self-referential processing in (remitted) depressed patients, increasing DMN connectivity. Based on these previous findings it is hypothesized that compared to non-patient controls, cMDD patients would show increased connectivity in the DMN following a sad mood induction.

2. Method

2.1. Participants

Eighteen patients with chronic major depressive disorder (cMDD) and 18 age, education level and gender matched non-patient controls participated in this study. Non-patient controls were also matched on left/right handedness except for one left handed patient who could not be matched with a left handed control participant. Patients were recruited from a specialized secondary care outpatient unit in the Netherlands (Riagg Maastricht) as part of a treatment study [13]. All patients were scanned before the start of the intervention phase of the treatment study. Non-patient controls were recruited via poster advertisements and a volunteer database. Participants were excluded if they had MRI contraindications.

Inclusion criteria for patients in the treatment study were a current diagnosis of cMDD defined as meeting Diagnostic and Statistical Manual of Mental Disorders fourth edition (DSM-IV) criteria of major depressive disorder for ≥2 years, assessed with the Structured Clinical Interview for axis-I Disorders (SCID-I) by trained interviewers and a Beck Depression Inventory second edition (BDI-II) score of 20 or higher at study entry. Exclusion criteria for patients in the treatment study were the presence of bipolar disorder, psychotic disorders, alcohol or substance dependence, autism spectrum disorders, cluster-A or cluster-B personality disorder and acute suicide risk. One patient with co-morbid borderline personality disorder was not included in the treatment study but is included here. Use of antidepressant medication at study entry was an exclusion criterion, unless patients were stable on medications for three month or longer prior to study entry (n=10; 55.6%). Control participants meeting any current DSM-IV axis-I disorder or lifetime mood disorder, as assessed with the SCID-I interview were excluded.

The study protocol was approved by the medical ethical committee of the academic community hospital Maastricht, the Netherlands (Protocol ID/number: NL 3187.1.06.81.10). After complete description of the study to the participants, written informed consent was obtained. The authors assert that all procedures contributing to this work comply with the ethical standards of the relevant national and institutional committees on human experimentation and with the Helsinki Declaration of 1975, as revised in 2008. Demographic and clinical characteristics of the sample are summarized in Table 1.

2.2. Procedure

This study used a mixed design with group (cMDD, non-patient controls) as between-subject factor and two fMRI runs (pre mood induction, post mood induction) as within subject factor. First, participants underwent a 6 min resting-state scan. Participants then rated their current mood along four visual analogue scales (VAS) ranging from 0 to 100 with higher scores representing more positive mood using an MR compatible joystick. The four VAS scales measured mood on a bipolar continuum between negative and positive, dull and glad, secure and anxious, and sad and happy. After the initial mood ratings, participants received the sad mood induction, consisting of mood suggestive music (“Adagio for Strings” by Samuel Barber played via headphones), played during the anatomical scan, in combination with autobiographical recall of a personal sad event (8 min). This standard protocol reliably induces short lasting sad mood-states [14] and the specific piece of music has successfully been used in our previous studies [15,16]. Directly after the sad mood induction, current mood-state was again assessed along the four VAS scales. Following this, participants underwent the second resting-state fMRI scan (6 min). After the second resting-state scan participants again rated their current mood-state on the four VAS scales.

2.3. Measures

2.3.1. Beck depression inventory

Depressive symptom severity was assessed using the Beck Depression Inventory second edition (BDI-II), a 21-item self-report instrument [17]. In the present study the Dutch version of the BDI-II was used which has been shown to have high internal consistency in a Dutch sample of psychiatric patients (Cronbach’s α=0.92) and ade-

<table>
<thead>
<tr>
<th>Table 1</th>
<th>Demographic and clinical characteristics of the sample.</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>cMDD (N =18)</td>
</tr>
<tr>
<td>Age, M (SD)</td>
<td>41.17 (17.53)</td>
</tr>
<tr>
<td>Gender, n (%)</td>
<td></td>
</tr>
<tr>
<td>female</td>
<td>13 (72.2)</td>
</tr>
<tr>
<td>male</td>
<td>5 (27.8)</td>
</tr>
<tr>
<td>Education level, n (%)</td>
<td></td>
</tr>
<tr>
<td>low</td>
<td>1 (5.6)</td>
</tr>
<tr>
<td>medium</td>
<td>6 (33.3)</td>
</tr>
<tr>
<td>high</td>
<td>11 (61.1)</td>
</tr>
<tr>
<td>BDI-II score, M (SD)</td>
<td>30.50 (9.39)</td>
</tr>
<tr>
<td># Previous episodes, M (SD)</td>
<td>3.38 (2.39)</td>
</tr>
<tr>
<td>Antidepressant use at baseline, n (%)</td>
<td>10 (55.6%)</td>
</tr>
</tbody>
</table>

Note. BDI-II = Beck Depression Inventory second edition; cMDD = chronic Major Depressive Disorder. Information on number of previous episodes was available from 16 patients.
quate construct validity with other depression scales rs between 0.79 and 0.85 [18].

2.4. Image acquisition

Anatomical images were acquired using a Magnetom Allegra 3 T scanner (Siemens Healthcare, Netherlands) located at the Faculty of Psychology and Neuroscience, Maastricht University with a T1-weighted gradient echo (196 slices, TR =2250 ms, TE =26 ms, flip angle =90°, field of view =256 mm and voxel dimensions 1×1×1 mm³). A T2*-weighted functional measurement was acquired using a standard echo-planar imaging (EPI) sequence (TR 2000 ms, TE 30 ms, flip angle =90°, 32 slices, 180 volumes, 3×3×3 mm³). A slice tilt correction of −30 was used to minimize inhomogeneity artefacts [19].

2.5. fMRI data preprocessing

Data pre-processing and analyses were conducted with Brainvoyager QX v2.4 (Brain Innovations, Maastricht, the Netherlands). The first two volumes of the functional images were omitted due to magnetization artifacts. Preprocessing of the remaining functional data included slice time correction using sinc interpolation, 3D motion correction using sinc interpolation, spatial smoothing (Gaussian kernel with full-width-at-half-maximum of 4 mm) and linear trend removal. Individual functional datasets were then co-registered with structural images of the same participant and subsequently normalized using the Talairach and Tournoux transformation procedure [20]. Talairach coordinates were subsequently transformed to Montreal Neurological Institute (MNIs) coordinates based on [21]. Individual anatomical datasets were then averaged to create a group-based anatomical brain mask to exclude voxels belonging to the outside of the brain.

2.6. Resting-state functional connectivity analysis

To determine change in DMN connectivity following the sad mood induction, we conducted a seed based connectivity analysis. The PCC was selected as seed because this region has been shown to reveal connectivity in the DMN most consistently [22]. The anatomical location of the PCC seed was based on the results of a separate self-referential processing task, performed in the same experimental session by the same participants, following the resting-state scans [23]. In this task a whole-brain Random Effects (RFX) ANOVA analyses was used to contrast personality trait words with non-words. This analysis resulted in the functional localization of the cortical midline structures involved in self-referential processing [24], including the PCC. The peak voxel of the left PCC (5 mm sphere around MNI: −9, −56, 25) was selected as seed, which is comparable in location to regions that previous studies have used to reveal the DMN [11,25]. A 5 mm sphere has been previously used to define the PCC as seed area e.g. [26].

In order to perform the group seed based analysis in Brainvoyager QX, single study design matrix (SDM) files were created for each run of each participant using the NeuroElf MATLAB (Mathworks, Inc.) toolbox (www.neuroelf.net) and a resting state analysis toolbox developed at Maastricht University [27,28]. These SDM files included the BOLD time series of the PCC seed region and 122 z-normalized nuisance confound regressors: six 3D head motion parameters and their first derivatives, mean signal from the ventricles and white matter, global signal, and signal oscillations at a frequency above 0.1 Hz (sine-cosine pairs). The resulting SDM files were then used to run an RFX GLM to find voxels that showed correlation with the PCC seed for each run of each participant. The Pearson correlation coefficients (r) maps were then transformed using Fisher’s r-to-z transformation (z=0.5 Ln [(1+r)/(1−r)]), to yield variates that are approximately normally distributed. The resulting F-map showed all voxels that had a significant correlation with the PCC seed at a conservative false discovery rate (FDR) of 0.001 for all participants over all runs. This map was transformed into a DMN mask (Fig. 1), which was used to limit the final main effects and groups x mood induction analyses to the hypothesized DMN.

In the final analysis, first, the main effect of group on the first resting-state scan (pre mood induction) and the main effect of run (resting-state pre/post mood induction) was computed to replicate earlier findings of increased DMN connectivity in depressed patients and to determine the effect of the sad mood induction on DMN connectivity in general, respectively. Second, a repeated measures ANOVA analysis with group (non-patient control vs. cMDD) as subject factor was conducted to test for the hypothesized difference in the effects of the sad mood induction between the two groups. The resulting three F-maps were thresholded at a p-value of 0.01 and a cluster size threshold of 135 mm³ (5 voxels) as determined by the cluster-level statistical threshold estimator plugin (BrainVoyager QX; 1000 Monte Carlo simulations).

![Fig. 1. PCC based functional connectivity mask resembling the default mode network.](image-url)
3. Results

3.1. Behavioral data

3.1.1. Manipulation check

Within group effects of the mood induction were assessed using paired t-tests and between group differences were assessed using repeated measures analysis of variance. To reduce statistical testing, an average mood scale of all four VAS scales was constructed per mood assessment moment (Cronbach’s alpha = 0.91, 0.88, and 0.85). Fig. 2 shows average mood ratings before and after the sad mood induction and after the second resting state scan. Within group comparisons revealed a significant decrease in mood for non-patient controls and cMDD patients, \( t(17) = 6.04, p < 0.001 \) and \( t(17) = 3.18, p = 0.005 \), respectively, indicating the success of the sad mood induction procedure. The decrease in mood following the sad mood induction was slightly stronger in non-patient controls, but the interaction effect was not statistical significant \( F(1, 34) = 3.87, p = 0.057 \). At the end of the experiment, mood state remained at a lower level compared to the pre mood induction assessment for both patients \( t(17) = 2.76, p = 0.01 \) and non-patient controls \( t(17) = 3.46, p < 0.01 \). At each time point there was a significant main effect of group, showing that patients reported a lower mood-state at pre mood induction, \( t(34) = -7.58, p < 0.001 \), post

![Fig. 2. Effects of the mood induction manipulation on self-reported mood ratings Mood ratings prior to the mood induction (T1), after the mood induction (T2) and at the end of the experiment (T3); higher scores represent more positive mood. Error bars represent standard error of the mean.](image1)

![Fig. 3. Group (chronic Major Depressive Disorder/non-patient controls) × mood induction (pre/post) interactions. The y–axis of the graphs depicts the mean z values of functional connectivity of the Posterior Cingulate Cortex with the Default Mode Network. Error bars represent standard error of the mean.](image2)
mood induction, t(34) = −5.11, p < 0.001, and after the second resting-state scan, t(32) = −5.29, p < 0.001.

3.2. fMRI results resting-state functional connectivity

3.2.1. Main effect group

Prior to the sad mood induction cMDD patients, compared to non-patient controls, showed greater connectivity between the PCC and the medial temporal gyrus (52,−68,28), the dorsolateral medial prefrontal cortex (10,61,30), the precuneus (−6,−65,29), the anterior prefrontal cortex (−6,57,30), the PCC (−6,56,10), the superior occipital gyrus (−45,−78,34) and the medial temporal gyrus (−58,−59,15).

3.2.2. Main effect pre to post mood induction

There was an overall significant decrease in connectivity from pre mood induction to post mood induction in the precuneus (−6,−58,35) and in two separate clusters of the parahippocampal gyrus (−13,−33,−1 and −36,−25,−18).

3.2.3. Group on mood induction interaction

Five ROIs with significant group (cMDD/non-patient controls) × mood induction (pre/post) interactions were identified (Fig. 3; Table 2), including the anterior inferior temporal cortex, the superior temporal gyrus, the uncus/anterior parahippocampal gyrus, the anterior prefrontal cortex, and the precuneus. In line with our hypothesis the anterior inferior temporal cortex, the superior temporal gyrus and the uncus/anterior parahippocampal gyrus showed increased connectivity after the sad mood induction in patients with cMDD compared to non-patient controls. However, the precuneus and anterior prefrontal cortex showed the opposite pattern, with increased connectivity after the sad mood induction in non-patient controls compared to the patient group.

4. Discussion

The present study provides the first evidence, to our knowledge, that DMN connectivity in currently depressed participants is modulated by sad mood states. Using a PCC seed, main effect analyses of group prior to the sad mood induction replicated previous findings showing increased DMN connectivity in patients with depression, for a review see for example [29]. The DMN has been linked to self-referential processing, rumination and autobiographic memory and therefore plays an important role in the neuropathology of depression. In this study we focused on connectivity within the DMN using the PCC as seed. Previous studies have also demonstrated increased cerebellar-DMN connectivity in individuals with depression showing that sub-regions of the Cerebellum are linked to the DMN [30]. The main focus of the present study was on the hypothesized increase in DMN connectivity following the sad mood induction in cMDD patients. In line with this hypothesis, PCC connectivity with the anterior inferior temporal cortex, the superior temporal gyrus and the parahippocampal gyrus increased following the sad mood induction in cMDD patients. However, the precuneus, an important DMN node, showed a decrease in connectivity with the PCC in cMDD patients. Note that the simple effect of change in PCC-precuneus connectivity within the cMDD group was marginally significant (p =0.08). Non-patient controls showed an increased PCC connectivity with the anterior prefrontal cortex. These findings are in line with previous studies showing increased DMN connectivity in remitted depressed patients, compared to healthy controls, following a sad mood induction [11] and decreased DMN connectivity in healthy controls following a sad mood induction [10]. Together, empirical findings show that DMN connectivity in depressed patients can be modulated by subjective mood-states.

In general, the present finding of increased DMN connectivity, following the sad mood induction in cMDD patients supports neurobiological models of depression [31]. Importantly, in the current study the sad mood induction procedure was conducted before functional imaging of the second resting-state scan and no effortful cognitive processing was required by participants during the resting state scan. In contrast, non-clinical participants in Harrison et al. (2008) engaged in effortful processing during the resting state scan. Thus their finding of decreased connectivity following the sad vs neutral mood induction might be due to task-induced (sad mood induction) deactivation as the authors discuss [10]. A negative mood induction in (remitted) depressed patients might trigger self-referential processing and rumination, associated with increased DMN connectivity [11]. The increased connectivity following the sad mood induction in patients compared to non-patient controls in the current study therefore likely reflects a relatively natural reaction of patients, after being challenged with sad mood, rather than compliance of the task instruction to recall sad autobiographical events.

We found increased PCC – parahippocampal gyrus connectivity in patients with cMDD following the sad mood induction. Another study focusing on dynamic changes of functional connectivity within the DMN in individuals with depression used a MPFC seed and reported decreased dynamic connectivity with the parahippocampal gyrus potentially reflecting increased self-focus thinking [32]. In line with findings of the current study, another study [11] reported increased PCC - parahippocampal gyrus connectivity following a sad mood induction in previously depressed participants. The increased PCC - parahippocampal gyrus connectivity in patients following the sad mood induction might reflect activation of negative thoughts and rumination in patients as has been suggested previously [11]. The results of the current study therefore provide further evidence for the idea that the parahippocampal gyrus plays a potentially important role in chronic depression [11].

The finding of decreased PCC – precuneus connectivity in patients versus controls following the mood induction was unexpected in the light of previous studies showing increased precuneus – DMN connectivity in patients with (chronic) depression [12]. The precuneus plays a key role in visual mental imagery, episodic memory recall and self-referential processing [33]. Thus, increased PCC – precuneus connectivity in non-patient controls in the present study might reflect episodic autobiographical memory retrieval triggered by the negative mood induction that has not been observed in previous studies not using mood-induction procedures. One previous study in medication

<table>
<thead>
<tr>
<th>Table 2</th>
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<tbody>
<tr>
<td>Region of interest details identified for interaction between group (chronic Major Depressive Disorder vs non-patient controls) and pre and post-mood induction.</td>
</tr>
<tr>
<td>Brain region</td>
</tr>
<tr>
<td>---</td>
</tr>
<tr>
<td>Anterior inferior temporal cortex</td>
</tr>
<tr>
<td>Uncus / parahippocampal gyrus</td>
</tr>
<tr>
<td>Anterior prefrontal cortex</td>
</tr>
<tr>
<td>Superior temporal gyrus</td>
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<tr>
<td>Precuneus</td>
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Note. L/R = Left / Right; Cohen’s d = Mean difference/pooled SD.
The present study suggests that PCC connectivity with a number of DMN regions increases in cMDD patients but not in controls following a sad mood induction. This finding has potential clinical implications. Patients with cMDD might fail to exert cognitive control over memory retrieval processes once sad mood is triggered, as indicated by the decreased PCC – anterior prefrontal cortex connectivity in patients with cMDD. Non-patient controls seem to increase cognitive control processes possibly in an attempt to cope with the sad mood-state, as indicated by an increased PCC connectivity with the anterior prefrontal cortex. Increasing cognitive control in patients with depression either through neurostimulation [45] or through behavioral interventions might be a fruitful avenue in the treatment of cMDD. In conclusion, this study suggests that DMN connectivity in patients with cMDD is modulated by negative mood states.

Role of Funding Source

This work was support by Maastricht University and by the Academic Community Mental Health Centre (RIAGG) Maastricht.

Acknowledgements

We are grateful to our colleagues at Maastricht University who supported this study: Annie Hendriks and Annie Raven helped with patient recruitment; Danielle Jongen assisted with recruitment of matched controls; Thanks are also due to our colleagues at the Riaag Maastricht who helped to recruit patients and conducted clinical interviews for this study.

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