ORIGINAL RESEARCH



Ruminative reflection is associated with anticorrelations between the orbitofrontal cortex and the default mode network in depression: implications for repetitive transcranial magnetic stimulation

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Accepted: 1 November 2021 / Published online: 3 December 2021
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Abstract

Patients with depression who ruminate repeatedly focus on depressive thoughts; however, there are two cognitive subtypes of rumination, reflection and brooding, each associated with different prognoses. Reflection involves problem-solving and is associated with positive outcomes, whereas brooding involves passive, negative, comparison with other people and is associated with poor outcomes. Rumination has also been related to atypical functional hyperconnectivity between the default mode network and subgenual prefrontal cortex. Repetitive pulse transcranial magnetic stimulation of the prefrontal cortex has been shown to alter functional connectivity, suggesting that the abnormal connectivity associated with rumination could potentially be altered. This study examined potential repetitive pulse transcranial magnetic stimulation prefrontal cortical targets that could modulate one or both of these rumination subtypes. Forty-three patients who took part in a trial of repetitive pulse transcranial magnetic stimulation completed the Rumination Response Scale questionnaire and resting-state functional magnetic resonance imaging. Seed to voxel functional connectivity analyses identified an anticorrelation between the left lateral orbitofrontal cortex (-44, 26, -8; k = 172) with the default mode network-subgenual region in relation to higher levels of reflection. Parallel analyses were not significant for brooding or the RRS total score. These findings extend previous studies of rumination and identify a potential mechanistic model for symptom-based neuromodulation of rumination.

 $\textbf{Keywords} \ \ Rumination \cdot Reflection \cdot Treatment-resistant \ depression \cdot Default \ mode \ network \cdot Repetitive \ transcranial \ magnetic \ stimulation$

Introduction

Rumination, a disabling repetitive focus on symptoms of depression, has been related to abnormally high functional connectivity (FC) between the subgenual prefrontal cortex and the default mode network (DMN; Hamilton et al., 2015). There is an evolving literature indicating that repetitive pulse transcranial magnetic stimulation (rTMS) can alter FC in major depressive disorder (MDD; for a review see Beynel et al., 2020; Fox et al., 2014; Philip et al., 2018). Many early models of rTMS depression therapy hypothesize that the

dorsolateral prefrontal cortex is dysfunctional in exerting cognitive control over other regions that mediate emotions (e.g., Lantrip et al., 2017). Whereas rTMS of the frontal lobe is an FDA approved approach to treat MDD, the most effective frontal targets within that region for different patients are not established. A number of studies demonstrate that frontal regions with the strongest negative correlation (anticorrelation) with the subgenual cingulate are relatively more effective targets in reducing depression (Cash et al., 2017; Fox et al., 2012; Mir-Moghtadaei et al., 2015; Weigand et al., 2018; Williams et al., 2018). These finding are consistent with brain activity in depressed patients, including abnormally high activity in the subgenual region (Berlim et al., 2014; Drobisz & Damborska, 2019; Mayberg et al., 1999; Morishita et al., 2014) and low activity in the

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frontal lobe (Baxter et al., 1989; George & Wassermann, 1994; Martinot et al., 2010). Thus, targeting regions with high anticorrelation between these regions may normalize this relationship. Identifying which frontal regions are functionally connected with the DMN-subgenual complex of regions (DMN-subgenual), and which are also associated with severity of rumination, could identify cortical regions that could potentially modulate this elevated connectivity and alter rumination. One challenge however, is that rumination is comprised of two distinct cognitive subtypes—reflection and brooding—that are both associated with worse current depression severity but different longitudinal outcomes (Nolen-Hoeksema & Morrow, 1991; Treynor et al., 2003). Reflection is an active problem-solving thought process that predicts later decreases in depression severity. Whereas increasing a form of rumination to treat depression may seem counterintuitive, reflection is believed to be compensatory to support recovery. In contrast, brooding is a passive comparison with an unachievable standard that predicts later increases in depression severity (Treynor et al., 2003). Knowing which of these subtypes of rumination are associated abnormal DMN-subgenual FC is thus important for directing clinical neuromodulation. In sum, if elevated DMN-subgenual FC plays a mechanistic role in rumination then identifying which rumination subprocess is related and which cortical target is connected will be important for modulating this system.

We aimed to perform a resting state fMRI analysis of data from a cohort of patients with treatment-resistant depression undergoing rTMS therapy (Yesavage et al., 2018) and identify a prefrontal cortical region in which the functional connection to the brain areas related to rumination (i.e., DMN-subgenual) is related to the self-reported level of rumination (i.e., reflection and/or brooding). Since reducing depression involves targeting a frontal region anticorrelated with a subgenual region, we hypothesized that a similar pattern of FC would be related to a process like reflection that is associated with reduced depression (Treynor et al., 2003). Specifically, the strength of self-reported reflection should be correlated with the strength of the frontal anticorrelation with the DMN-subgenual region. Conversely, since brooding is associated with increased depression (Treynor et al., 2003), the strength of self-reported brooding should be correlated with the strength of the frontal positive correlation with the DMN-subgenual region. To test these hypotheses, we defined a region of interest (ROI) based on a meta-analysis of rumination studies of MDD (Hamilton et al., 2015). This region of interest was comprised of two core regions of the DMN (posterior cingulate cortex and medial prefrontal cortex; Fox et al., 2005) and a subgenual region (Hamilton et al., 2015). We then used a seed to voxel analysis (Whitfield-Gabrieli & Nieto-Castanon, 2012) testing whether there was a cortical region, likely in the prefrontal cortex, whose

correlation with this ROI was related to level of self-reported reflection and/or brooding. This cortical region could serve as a potential candidate region to target for modulation (Fox et al., 2014) and further supported the model of rumination as involving an interaction between a frontal lobe subregion and the DMN-subgenual region.

Methods

Participants

Forty-three Veterans with treatment-resistant depression (33 male; mean age = 54.98 [SD = 12.27] years, mean education = 13.9 [SD = 1.96] years; see Table 1 for demographic data), were enrolled in this sub-study as part of a larger prospective, randomized, sham-controlled rTMS for depression clinical trial (ClinicalTrials.gov ID: NCT01191333). A detailed description of the full inclusion and exclusion criteria for the trial was previously published (Mi et al., 2017; Yesavage et al., 2018). As part of the clinical trial, the clinician administered Hamilton Depression Rating Scale (Hamilton, 1960) and Clinician-Administered PTSD Scale for the DSM-IV (Blake et al., 1995) were completed to confirm the presence of major depressive disorder and to identify posttraumatic stress disorder (28% of the participants in this study met criteria for PTSD, see Table 1). These

Table 1 Demographic and questionnaire descriptive statistics

Characteristics	Mean (SD)		
Age (years)	54.98 (12.27)		
Education (years)	13.9 (1.96)		
RRS-Reflection	11.28 (3.35)		
RRS-Brooding	12.4 (3.79)		
BDI-II	23.3 (10.26)		
PCL-M	42.67 (17.23)		
fMRI Post-Active Treatment (months)	21.07 (9.34)		
	N (percentage)		
Sex			
Male	33 (76.7%)		
Female	10 (23.3%)		
Handedness			
Right	38 (88.4%)		
Left	5 (11.6%)		
fMRI Post-Active Treatment	7 (16%)		
PTSD Diagnosis	12 (28%)		

RRS Rumination Response Scale, reflection and brooding subscales, BDI-II Beck Depression Inventory-II, PCL-M Posttraumatic Stress Disorder Checklist, Military Version, fMRI Post-Active Treatment subjects with fMRI scan acquired after active repetitive pulse transcranial magnetic stimulation treatment; PTSD Diagnosis is determined from the Clinician-Administered PTSD Scale



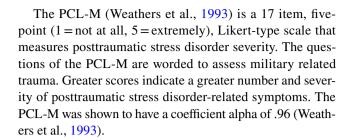
measures were sometimes completed at a remote time from MRI data acquisition, so that to enable evaluation of changes in symptom severity, brief self-report measures of depression (BDI-II) and posttraumatic stress disorder (PCL-M) were completed close in time to the MRI data acquisition, see questionnaires for additional detail. Participants were selected from patients who were willing and able to undergo fMRI data acquisition. All participants were screened for known neurological, comorbid psychiatric, and vascular risk factors or any medication which might affect vascular reactivity or cognitive performance.

Questionnaires

Participants completed the Rumination Response Scale (RRS; Nolen-Hoeksema & Morrow, 1991), Beck Depression Inventory-II (BDI-II; Beck et al., 1996), and the Posttraumatic Stress Disorder Checklist, Military Version (PCL-M; Weathers et al., 1993) within one week of fMRI data acquisition.

The RRS (Treynor et al., 2003) contains three subscales, two of which are unique to the cognitive subtypes of rumination- reflection and brooding. The third subscale contains depression related items and was not evaluated in this study since it was redundant with other more focused measures of depression (e.g., BDI-II). The RRS asks participants to indicate how often they think about or do mental and behavioral activities when they are depressed on a scale from low (1 = almost never) to high (4 = almost always) frequency. The reflection and brooding subscales include 5-items per subscale. The reflection subscale is thought to measure an attempt to decrease depression severity through problemsolving, with questions such as "Go away by yourself and think about why you feel this way." The brooding subscale is thought to measure a passive comparison with others and unrealistic standards (Treynor et al., 2003). The brooding subscale includes items such as "Think about a recent situation and wishing it could have gone better." The reflection subscale was shown to have a coefficient alpha of .72 and the brooding subscale was shown to have a coefficient alpha of .77 (Treynor et al., 2003). The RRS total score combines all three subscales into a composite score reflecting the construct of rumination. The RRS total score was shown to have a coefficient alpha of .90 (Treynor et al., 2003).

The BDI-II (Beck et al., 1996) is a 21-item, four-point, Likert-type scale that measures depression symptom severity over the past two weeks. Higher total scores are indicative of an increased number and severity of current depression symptoms. Scores from 0 to 13 indicate minimal depression, 14–19 indicates mild depression, 20–28 indicates moderate depression, and 29–63 indicates severe depression. The BDI-II has was shown to have a coefficient alpha of .92 (Beck et al., 1996).



Data collection

Scanning was performed exclusively on 3 T MRIs across five imaging centers as follows: Siemens Verio (Pittsburgh, PA), Siemens Trio (Charleston, SC and Salt Lake City, UT), Siemens Skyra (San Francisco, CA), and a GE 3 T Discovery 750 scanner (Palo Alto, CA). All Siemens sites used a 12-channel head coil with the exception of San Francisco which used a 32-channel head coil, and the GE site used an 8-channel head coil. High-resolution structural MRI (T1) of approximately 1mm³ in-plane resolution was collected using the protocols from the Alzheimer's Disease Neuroimaging Initiative (http://adni.loni.ucla.edu). The echoplanar fMRI pulse sequences were adapted from the functional biomedical informatics research network (fBIRN; Brown et al., 2011; Greve et al., 2011). Functional MRI scans were collected in an axial plane using an echoplanar sequence (TR 2 s, TE 30 ms, 77-degree flip angle, image resolution $3.44 \times 3.44 \times 3.5$ mm, 30 4-mm slices) of eight minutes duration (240 repetitions). Seven participants completed six-minute runs (180 repetitions) prior to the decision to lengthen the sequence to improve data robustness. Before commencing data collection an fBIRN (Keator et al., 2016) phantom and a traveling subject were scanned and cross site image quality was evaluated by our lead MR physicist.

fMRI preprocessing and FC analyses

Analyses were performed using the CONN Toolbox, version 18a (Chai et al., 2012; Whitfield-Gabrieli & Nieto-Castanon, 2012), which used Statistical Parametric Mapping, version 12 (http://www.fil.ion.ucl.ac.uk/spm/). Preprocessing of resting-state fMRI used the standard CONN pipeline, which included realignment and unwarping, centering to 0, 0, 0 coordinates, outlier detection, slice timing correction, segmentation and normalization into MNI space, and smoothing to 6-mm FWHM. Denoising to avoid spurious correlations caused by head motion and other signal changes unrelated to brain activity was completed. Problematic time points during the scan were removed using the Artifact Detection Tools (https://www.nitrc.org/projects/artifact_detect). Outlier images were removed if the head displacement in x, y, or z direction was greater than 0.9 mm from the previous frame, or if the global mean intensity in the image was



greater than 5 standard deviations from the mean image intensity for the entire scan. Global signal regression, a widely used preprocessing method, was not used because it can produce negative correlations that can influence the presence of anticorrelations (Murphy et al., 2009) and contribute to spurious positive correlations (Saad et al., 2012). Additionally, with the use of the Artifact Detection Tools software the benefit that global signal regression offers for motion correction (Ciric et al., 2017; Parkes et al., 2018) was less germane. Instead, the anatomical CompCor approach of noise reduction was used (Behzadi et al., 2007). Anatomical volumes were segmented into grey matter, white matter, and cerebrospinal fluid areas, and the resulting white matter and cerebrospinal fluid masks were eroded (one voxel erosion) to minimize partial volume effects. Nuisance variables were modeled as temporal covariates and removed from the blood oxygen level-dependent (BOLD) time series functional data using linear regression. These covariates included estimated subject motion (3 rotation and 3 translation parameters, plus another 6 parameters representing their first-order temporal derivatives) and the BOLD time series data outside the subject-specific grey matter mask (i.e., 3 temporal PCA components from the subject-specific white matter mask and 3 temporal PCA parameters from the cerebrospinal fluid mask). The resulting BOLD time series was band-pass filtered (0.008 Hz < f < 0.08 Hz).

DMN seed definition

The seed ROI was defined to reflect the DMN and the abnormally high functionally connected subgenual prefrontal cortex reported in individuals with MDD (Hamilton et al., 2015). Two commonly used DMN nodes, the posterior cingulate cortex (-5, -49, 40) and medial prefrontal cortex (-1, 47, -4; Fox et al., 2005), were combined with the subgenual prefrontal cortex (0, 26, -10) as a seed region. The WFU PickAtlas (Maldjian et al., 2004; Maldjian et al., 2003) was used to create 6-mm radius ROIs in MNI space for each of these three regions.

Subject-level seed to voxel analyses

Subject-level seed to voxel FC maps were generated with the preprocessed BOLD time course data. The three DMN-subgenual ROIs FC were averaged and then correlated with voxels throughout the remainder of the brain. Whereas our hypothesis based on rTMS therapeutic effects was that the ROI would correlate with a frontal subregion, a whole brain analysis enabled more conservative corrections in case alternative FC was detected (Beynel et al., 2020). This analysis produced subject-level correlation maps that were converted to z scores using Fisher's r-to-z transformation.

Group seed to voxel analyses

Group-based analyses were completed by correlating subject-level maps with reflection or brooding. Additionally, while our interests are primarily in the reflection and brooding subscales, to enable comparison to the previous literature we also evaluated the RRS total score, correlating the RRS total score with subject-level maps. Cluster thresholds were defined with a $p_{\text{uncorrected}} < .001$ and corrected for multiple comparisons with a two-sided $p_{\rm FDR}$ < .05. These analyses were repeated with the same statistical threshold while controlling for variables of no interest including demographic (age and sex) and clinical variables (PCL-M and BDI-II). Analyses with the addition of covariates provided additional information that the relationships between brain regions and rumination subtypes were selective and not better accounted for by depression severity or other comorbid and otherwise confounding variables. All seed to voxel analyses controlled for the effect of fMRI scanner site with four, individual by site, dummy coded variables (i.e., each variable contained 1 s for subjects scanned at a site and 0 s for subjects from other sites, repeated for four of five sites; these variables were controlled for in the general linear model). Additionally, to demonstrate that fMRI scanner site did not significantly alter results, significant results were analyzed without the fMRI scanner dummy coded variables while employing the same cluster threshold ($p_{\text{uncorrected}} < .001$) and multiple comparison controls (two-sided $p_{\rm FDR} < .05$).

Results

Patient characteristics

Table 1 shows descriptive statistics for the demographic and questionnaire data. On average, participants reported moderate depression severity ($\bar{x} = 23.3$; SD = 10.26) with approximately 60% of the participants experiencing depression severity in the moderate to severe range at the time of fMRI data acquisition. Seven participants received active rTMS treatment on average 21.07 (SD = 9.34) months prior to their fMRI data acquisition. To confirm our findings were not explained by these participants, significant seed to voxel analyses were repeated while controlling for the effect of these seven post-active rTMS participants (i.e., an additional dummy coded variable for these seven participants). The number of participants per fMRI scanner site are found in Supplementary Material Table 1 and correlations between the questionnaire results are found in Supplementary Material Table 2.



Regions correlated with DMN-subgenual and rumination subtypes and total score

Higher reflection scores were associated (two-sided $p_{\rm FDR}$ < .05) with FC between a subregion within the left lateral orbitofrontal cortex (peak -44, 26, -8; k = 172) and the DMN-subgenual seed. Seed to voxel analyses comparing brooding and the RRS total score with FC between the DMN-subgenual and voxels in the remainder of the brain were not significant using the same statistical threshold (two-sided $p_{\rm FDR} > .05$), hence no further analyses associated with brooding or the RRS total score are reported. Figure 1 depicts the spatial extent of the reflection-related cluster in the frontal lobe that extended beyond the left lateral orbitofrontal cortex peak to include the inferior frontal gyrus-pars triangularis, frontal operculum, and inferior frontal gyruspars opercularis. Individual participant FC values between the DMN-subgenual seed with the left lateral orbitofrontal cortex are shown in Fig. 2, which average to a positive correlation between these regions. Correlations between reflection and FC between the left lateral orbitofrontal cortex and the DMN-subgenual are represented in Fig. 3. This figure is displayed for descriptive purposes (Vul & Pashler, 2012). Overall, there is a negative correlation between the self-reported mental process of reflection and FC between the left lateral orbitofrontal with the DMN-subgenual. Further details are displayed in Table 2. The addition of demographic and clinical variable to the analyses continued to produce significant results (Table 2). Seed to voxel analyses with the addition of a control variable for the effect of the 7 participants with active rTMS prior to fMRI data acquisition produced similar results (Supplemental Material Table 3). Removal of the fMRI scanner site dummy coded variables from the significant seed to voxel analyses produced similar results, except for one analysis in which an additional right frontal pole cluster was significant (Supplemental Material Tables 4 and 5). The additional right frontal pole cluster (peak -46, 26, -8; k = 96) was identified in the seed to voxel

Fig. 1 Subject-level ROI-ROI analyses results. Significant seed to voxel analyses ($p_{\text{uncorrected}} < .001$ and two-sided $p_{\text{FDR}} < .05$) in which higher reflection scores are associated with a stronger anticorrelation between the seeded DMN-subgenual with the left lateral orbitofrontal cortex

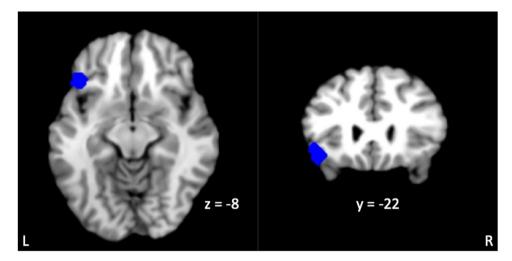


Fig. 2 Subject-level FC between the DMN-subgenual and the left orbitofrontal cortex. Subject-level FC between the DMN-subgenual and the left orbitofrontal cortex. The left lateral orbitofrontal cortex was identified with the seed-voxel analyses of the relationship between reflection and the DMN-subgenual seed. These z-transformed values have a group mean of .014 and standard deviation of .095, demonstrating that there is, on average, positive FC between the DMNsubgenual with the left lateral orbitofrontal cortex

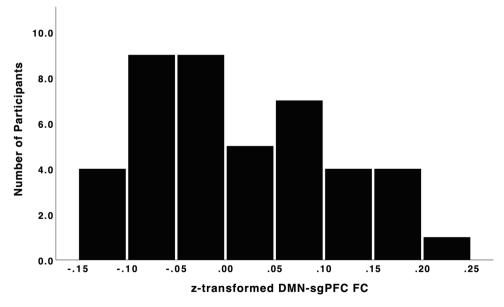




Fig. 3 Scatterplot of correlation between reflection with FC (cluster means) between the DMN-subgenual and the left lateral orbitofrontal cortex. Scatterplot depicting the Pearson correlation between the z-transformed reflection subscale of the Rumination Response Scale and the strength of FC (cluster means) between the DMN-subgenual seed with the left lateral orbitofrontal cortex. The FC values were derived from DMN-subgenual seed to voxel analysis with $p_{\rm uncorrected}$ < .001 and two-sided $p_{\rm FDR} < .05$

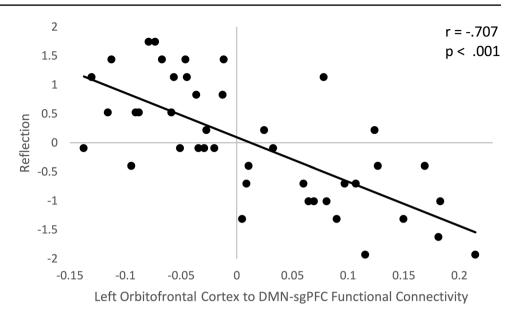


Table 2 Significant seed to voxel results

Label	Brod- mann Area	Peak Coordinates			Two-sided $p_{\rm FDR}$	t	Number of	Mean (SD)
		x	у	Z			voxels (k)	
Reflection								
Left lateral orbito- frontal cortex	47	-44	26	-8	.001	-6.25	172	.014 (.095)

Cluster subregions: orbitofrontal cortex = 107 voxels; inferior frontal gyrus pars triangularis = 28 voxels; frontal operculum = 11 voxels; inferior frontal gyrus pars opercularis = 2; not-labeled = 24 voxels

Reflection w/ Covariates

Cluster subregions: orbitofrontal cortex = 52 voxels; inferior frontal gyrus pars triangularis = 35 voxels; frontal operculum = 4 voxels; not-labeled = 17 voxels

All analyses controlled for the effects of fMRI scanner site. Covariates = age, sex, posttraumatic stress severity (PCL-M), and depression severity (BDI-II). Brodmann Area = area associated with peak MNI coordinates; t = 37 dof for primary analyses and 33 dof for analyses with covariates; Mean (SD) are Fisher r-to-z transformed functional connectivity values

analysis that evaluated the relationship between reflection while controlling for demographic and clinical variable.

Discussion

In this study, results indicate that increased reflection was associated with a stronger anticorrelation between a focal region with in the left lateral orbitofrontal cortex and the DMN-subgenual area, but no significant relationships were found between brooding or the RRS total score with FC

between the DMN-subgenual and the remainder of the brain. These results extend the literature relating symptoms of treatment resistant depression and FC. Notably, the association between FC and reflection suggests that modulating this network of regions may alter a potentially compensatory process rather than a process that is destructive such as brooding.

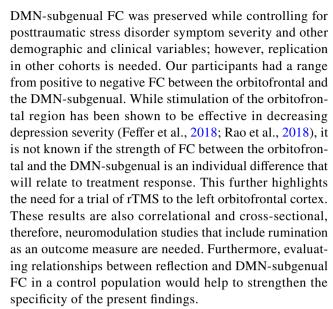
These results are consistent with previous rTMS studies suggesting depression can be treated through directly stimulating orbital frontal regions and those frontal regions with anticorrelations with the subgenual region. Whereas



modulating dorsolateral prefrontal cortical regions anticorrelated with the subgenual has been associated with more effective rTMS response (Cash et al., 2017; Fox et al., 2012; Mir-Moghtadaei et al., 2015; Weigand et al., 2018; Williams et al., 2018), other frontal regions have been suggested as rTMS targets such as a lateral orbitofrontal cortex (Downar & Daskalakis, 2013). Stimulation of the right orbitofrontal region in patients undergoing rTMS (Feffer et al., 2018) and invasive brain stimulation to left or right orbitofrontal regions (Rao et al., 2018) have been shown to decrease depression severity. Our findings demonstrating the longitudinally adaptive process of reflection is associated an anticorrelation between the orbitofrontal and the DMN-subgenual is consistent with other studies that have demonstrated that an increased anticorrelation between the orbitofrontal region and the DMN is adaptive for depression (Cheng et al., 2018; Jacobs et al., 2016), although our findings are in the left and the other findings are in the right hemisphere. Together, all three studies support that strengthening the anticorrelation between the DMN with the orbitofrontal cortex is associated with positive prognostic factors.

Our findings of an anticorrelation between the orbitofrontal cortex and the DMN-subgenual region is of particular interest because the orbitofrontal cortex is one of two prefrontal regions associated with rumination in major depressive disorder (Cooney et al., 2010). Specifically, during a rumination induction task, individuals diagnosed with major depressive disorder have greater activity than controls in the subgenual cingulate, orbitofrontal cortex, and dorsolateral prefrontal cortex (Cooney et al., 2010). While there is abundant evidence that targeting the dorsolateral prefrontal cortex with rTMS is effective for treating depression (Cash et al., 2017; Fox et al., 2012; Mir-Moghtadaei et al., 2015; Philip et al., 2018; Weigand et al., 2018; Williams et al., 2018), the orbitofrontal cortex is also demonstrating promise as a rTMS target for depression (Feffer et al., 2018). Our findings lend additional evidence that the orbitofrontal cortex may prove to be an effective target with rTMS for depression.

There are limitations to this study. Our participants were exclusively Veterans, many of whom were comorbid for posttraumatic stress disorder (Yesavage et al., 2018). Whereas clinician administered measures were used for the original diagnosis of posttraumatic stress disorder and major depressive disorder, symptom severity of posttraumatic stress disorder (PCL-M) and depression (BDI-II) were also collected at the time of MRI data acquisition in case there was change from the time of diagnosis. Whereas self-report measures are less burdensome and robust against variations in clinical rater biases, self-report measures may have been vulnerable to patient under or over-reporting. The relationship between reflection and



This study addressed a focused question about the nature of rumination and a focal abnormality in highly treatment resistant depressed patients (Hamilton et al., 2015); however, the study of perseverative cognition is a rich and growing literature (for a review see Makovac et al., 2020). Though resting-state fMRI is increasingly used to identify rTMS targets, future work should investigate the dynamics of networks (Chang & Glover, 2010) and further fMRI task-defined and resting-state regions identified by brooding (Kaiser et al., 2018; Vanderhasselt et al., 2013; Vanderhasselt et al., 2011) could be targeted with rTMS.

Conclusions

We demonstrated that increased reflection and not brooding or the RRS total-score is associated with an anticorrelation between the left lateral orbitofrontal region and the rumination-associated DMN-subgenual network of regions. This association suggests that modulating this system would alter a process believed to be compensatory.

Supplementary Information The online version contains supplementary material available at https://doi.org/10.1007/s11682-021-00596-4.

Author contributions Ehrlich and Rosen developed the study concept and design. Bhat provided integral assistance with data management. All other authors contributed data to the analyses from a larger project (Yesavage et al., 2018) and reviewed the manuscript. Ehrlich performed the data analyses and drafted the manuscript, which was revised by Rosen

Funding This work was supported by Cooperative Studies Grant # 556 and Merit Award # CX000604 from the United States (U.S.) Department of Veterans Affairs, Office of Research and Development.



Declarations

Ethical approval This study was approved by the Veterans Administration Central Institutional Review Board and the Research and Development Committee of the Veterans Affairs Palo Alto Health Care System.

Informed consent Informed consent was collected from all participants, who were paid for their participation.

Conflict of interest Dr. Mathalon is a consultant for Boehringer Ingelheim and Cadent Therapeutics. The remaining authors have no conflicts of interest to declare.

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