


RESEARCH ARTICLE

Neural changes following equine-assisted therapy for posttraumatic stress disorder: A longitudinal multimodal imaging study

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Abstract

Background: While effective treatments for posttraumatic stress disorder (PTSD) exist, many individuals, including military personnel and veterans fail to respond to them. Equine-assisted therapy (EAT), a novel PTSD treatment, may complement existing PTSD interventions. This study employs longitudinal neuro-imaging, including structural magnetic resonance imaging (sMRI), resting state-fMRI (rs-fMRI), and diffusion tensor imaging (DTI), to determine mechanisms and predictors of EAT outcomes for PTSD.

Method: Nineteen veterans with PTSD completed eight weekly group sessions of EAT undergoing multimodal MRI assessments before and after treatment. Clinical assessments were conducted at baseline, post-treatment and at 3-month follow-up.

Results: At post-treatment patients showed a significant increase in caudate functional connectivity (FC) and reduction in the gray matter density of the thalamus and the caudate. The increase of caudate FC was positively associated with clinical improvement seen immediately at post-treatment and at 3-month follow-up. In addition, higher baseline caudate FC was associated with greater PTSD symptom reduction post-treatment.

Conclusions: This exploratory study is the first to demonstrate that EAT can affect functional and structural changes in the brains of patients with PTSD. The findings suggest that EAT may target reward circuitry responsiveness and produce a caudate pruning effect from pre- to post-treatment.

KEYWORDS

equine assisted therapy, neuroimaging, PTSD, resting state fMRI

Trial Registration: ClinicalTrials.gov identifier: NCT03068325

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1 | INTRODUCTION

Posttraumatic stress disorder (PTSD) is a debilitating mental health disorder that occurs in the aftermath of trauma exposure. The prevalence of PTSD in the general population of the United States is estimated at nearly 9% (American Psychiatric Association, 2013), but ranges from 10 to 30% for US veterans (Kang, Natelson, Mahan, Lee, & Murphy, 2003). Active-duty military personnel and veterans suffering with PTSD are often reluctant to enroll or simply fail to respond to currently available psychotherapies (Holder, Holliday, Wiblin, LePage, & Suris, 2019; Lewis, Roberts, Gibson, & Bisson, 2020) and pharmacotherapies (Sullivan & Neria, 2009). One novel alternative and complementary treatment for PTSD is equine-assisted therapy (EAT) (Bachi, Terkel, & Teichman, 2012; Selby & Smith-Osborne, 2013). Specifically, EAT uses a horse (or horses) to facilitate communication and mindful awareness of thought and behaviors in an experientially oriented approach (Anestis, Anestis, Zawilinski, Hopkins, & Lilienfeld, 2014). A psychotherapist, assisted by an equine specialist, helps a patient understand and negotiate interactions with a horse, which should help foster regulation, reflection, and verbal and nonverbal communication (Selby & Smith-Osborne, 2013). We have recently developed and manualized EAT for PTSD and tested it in a pilot trial (Arnon et al., 2020) and in a large open trial (Fisher et al., 2021, under review) for military veterans with PTSD. Eight weekly sessions yielded reduction in PTSD and depressive symptoms, high acceptability (very few drop-outs), and tolerability. As part of the open trial, we conducted an exploratory neuroimaging study on a subset of participants. We aimed to examine whether observable neural changes can a) explain the mechanisms whereby EAT might ameliorate PTSD and b) predict those individuals most likely to successfully respond to this therapeutic intervention.

Assessment of neural mechanisms related to treatment response has increased in psychotherapy research (Nisar et al., 2020; Zhu et al., 2018). In PTSD, magnetic resonance imaging (MRI) has been also used to assess PTSD treatment-related changes, including structural alteration (Helpman et al., 2016), task-based functional activity (Fonzo et al., 2017a; Helpman et al., 2016), and resting-state functional connectivity (rsFC) (Zhu et al., 2018). Studies have explored these brain changes following prolonged exposure (PE), first line treatment for PTSD (Fonzo et al., 2017b; Helpman, Papini, et al., 2016; Suarez-Jimenez et al., 2020; Zhu et al., 2018), affect-focused interpersonal psychotherapy (IPT) (Suarez-Jimenez et al., 2020), eye movement desensitization and reprocessing (EMDR) (Laugharne et al., 2016), cognitive behavioral therapy (CBT) (Levy-Gigi, Szabo, Kelemen, & Keri, 2013), brief eclectic psychotherapy (BEP) (Lindauer et al., 2005), and trauma-focused therapy (van Rooij et al., 2015). The findings suggest that different PTSD treatments may target distinct mechanisms of action. To our knowledge, no study has utilized different neuroimaging modalities to investigate the CNS mechanisms by which EAT may affect the severity of PTSD.

The present study employs standardized longitudinal multimodal MRI, including structural MRI (sMRI), resting state-fMRI (rs-fMRI), and diffusion tensor imaging (DTI). Our first aim was to explore pre- to

post-treatment changes in: a) rsFC in reward, executive control, default mode, and salience networks; b) regional grey matter volume (GMV), measured by sMRI; and c) fractional anisotropy (FA) measured by DTI. The second aim was to clarify the association between changes of multimodal imaging biomarkers (sMRI, rsfMRI, DTI) and clinical response at post-treatment and at three-month follow-up (FU); and to examine whether these biomarkers might serve as baseline predictors of clinical improvement.

2 | METHODS

2.1 | Participants

The study sample comprised 20 treatment-seeking veterans with PTSD, a subset of participants taking part of a larger open trial ($n = 63$) of EAT for veterans with PTSD (Fisher, Lazarov et al. under review). Additional funding support allowed us to introduce MRI scans later in this larger open trial after 47.6% of participants ($n = 30$) had completed the treatment protocol. Due to the scheduling issues and MRI scanning requirements (e.g., no metal, tolerance of MRI procedure) 20 participants were recruited for the MRI portion of the study of which 19 underwent clinical evaluation at baseline, post treatment and three-month follow-up, and completed MRI assessment at baseline and post-treatment. One patient discontinued treatment prematurely. Table 1 illustrates the demographic information and the clinical outcomes of 19 treatment completers (mean age = 46.64, std = 12.34; 53% female), respectively. The New York State Psychiatric Institute Institutional Review Board approved the study. Participants provided written informed consent. ClinicalTrials.gov identifier: NCT03068325.

2.2 | Procedure

Participants were recruited through clinical referrals and print and online advertisements. After a telephone screen to determine probable eligibility, using the PTSD Checklist DSM-5 (PCL-5; Blevins, Weathers, Davis, Witte, & Domino, 2015), potential participants underwent clinical assessment by a master's or PhD-level clinician trained to $\geq 85\%$ reliability with a senior clinician on all interview-based measures. Psychiatric diagnoses were ascertained using the Structured Clinical Interview for DSM-5 (SCID-5; First, Williams, Karg, & Spitzer, 2015). Additional inclusion criteria were age 18–70, reported military experience, and English fluency. Exclusion criteria were: history of psychotic disorder or unstable bipolar disorder, determined by SCID-5 (First et al., 2015); 17-item Hamilton Rating Scale for Depression (HAM-D; Hamilton, 1960) score > 25 ; elevated suicide risk, determined by clinical interview; severe substance or alcohol use disorder within the past six months, or moderate use disorder within the past two months; and physical limitations impeding participation. Concurrent ongoing mental health treatment, including medication, was allowed. Initiating concurrent psychotherapy or pharmacotherapy

TABLE 1 Sample demographic and clinical characteristics

Variable	n	%
Gender, n (%)		
Female	10	53
Male	9	47
Ethnicity		
Hispanic, n (%)	3	16
Non-Hispanic	15	79
Prefer not to answer	1	5
Race, n (%)		
White	6	32
Black	7	37
Other	5	26
Prefer not to answer	1	5
Employment, n (%)		
Working full-time	1	5
Unemployed	4	21
Retired	2	11
Disabled	9	47
Student	3	16
Income, n (%)		
<30 K	12	64
30–40 K	3	16
>80 K	4	22
Medications, n (%)		
Yes	11	58
No	8	42
Psychotherapy, n (%)		
Yes	11	58
No	8	42
Any treatment, n (%)		
Yes	14	74
No	5	26
Co-morbid diagnoses		
Major depressive disorder	10	53
Persistent depressive disorder	4	21
Other depressive disorder	1	5
Alcohol use disorder	3	16

during the course of EAT was also deemed exclusionary. For the imaging protocol, additional exclusion criteria were having metal in the body, unable to tolerate the scanning procedures, severe medical disorder, or significant claustrophobia.

2.3 | Treatment

EAT for PTSD comprised eight weekly 90-minute group sessions held at an equestrian center. Each group was led by a licensed mental

health professional, an equine specialist, and an extra horse wrangler to enhance safety. All sessions began with a grounding exercise, focusing attention on current physical sensations. The first session has oriented patients to EAT (rationale, description, possible benefits), provided safety guidelines, psychoeducation (common reactions to trauma, development and maintenance of PTSD), and a barn tour, and ended with meeting two horses in a round-pen. Subsequent sessions were built upon previous ones, by reviewing previous session content and introducing increasingly complex encounters and interactions with horses, coupled with team feedback and direction. Each session ended with an opportunity for participants to review and discuss their experiences. The final session included a graduation ceremony celebrating patients' treatment progress and accomplishments (see Arnon et al., 2020; Fisher et al., under review).

2.4 | Clinical measures

Clinical interviewers assessed PTSD and depression symptoms at pre-, post-treatment, and 3-month follow-up. The CAPS-5 was the primary outcome assessment of DSM-5 PTSD criteria. The 17-item HAM-D (Hamilton, 1960) assessed depressive symptoms. We also administered two self-report measures, the PCL-5 (Blevins et al., 2015) for PTSD symptom severity and Beck Depression Inventory-II (BDI-II) for depressive symptoms (Beck, Steer, Ball, & Ranieri, 1996). These measures have well-documented psychometrics and validly, reliably assess their respective constructs (Beck et al., 1996; Bovin et al., 2016).

2.5 | Image procedures

2.5.1 | Image acquisition and analysis

Eleven participants were scanned twice using a 3T General Electric MR750, and nine participants were scanned twice using a 3T Siemens Prisma scanner due to scanner upgrade, equipped with a 32-channel receive-only head coil. Each participant was scanned twice with the same scanner. For each participant a high-resolution T1-weighted sequence was acquired using the following parameter: T1 = 450 ms (GE) or 900 ms (Siemens), Flip angle = 12° (GE) or 8° (Siemens), 256 × 256 matrix, slice thickness = 1 mm. T2*-weighted echo-planar images (EPis) depicting the blood-oxygen-level-dependent (BOLD) were acquired for each participant with TR = 1.3 s (GE) or 1.5 s (Siemens), TE = 28 ms, FA = 60° (GE) or 75° (Siemens), FOV = 19.2 cm, number of slices = 27, slice thickness = 4 mm. A head cushion limited head motion.

DTI data were obtained in the axial plane using a single-shot diffusion-weighted echo planar imaging sequence with the following parameters: TR/TE = 8500/77ms, FOV = 240mm, matrix size = 122×122, slice thickness = 2.5 mm, number of slices = 60, number of diffusion gradient directions = 64, b -value = 1000 s/mm², number of images at b -value of 0 s/mm² = 5, acceleration factor = 2. The scanner

factor was controlled as a covariate of no-interest in all three modalities analysis.

2.5.2 | Resting state fMRI processing

All rsFC images were preprocessed using MATLAB version R2018a (The MathWorks, Inc., Natick, Massachusetts) and statistical parametric mapping software (SPM12; Wellcome Trust Centre for Neuroimaging, UCL, London, UK). (a) Functional images were spatially realigned to the first image in the time series using a six-parameter rigid body transformation; (b) then slice-time correction was performed; (c) outlier detection was carried out using artifact removal tools (ART). The principal component-based noise-correction method, “CompCor,” implemented in this toolbox, was used to additionally control physiological noise and head motion effects. Outlier volumes in each participant were identified as having large spiking artifacts (i.e., volumes >3 standard deviations from the mean image intensity), or large motion (i.e., 0.5 mm for scan-to-scan head-motion composite changes in the x , y , or z direction); (d) each functional image was then spatially normalized to the standard T1 template included in SPM12; Functional images were then resliced to $2 \times 2 \times 2$ mm voxels, according to the resulting spatial realignment and normalization parameters; (e) anatomical images were segmented into grey matter, white matter, and cerebrospinal fluid (CSF) regions; (f) functional scans were smoothed with an 8mm full-width-at-half-maximum (FWHM) Gaussian kernel; (g) covariates corresponding to head motion (6 realignment parameters and their derivatives), outliers (one covariate per outlier consisting of 0s everywhere and a 1 for the outlier time point), and the BOLD time series from the subject-specific white matter and CSF masks were used in the GLM and connectivity analysis as predictors of no interest, and were removed from the BOLD functional time series using linear regression. Sum of root mean square (RMS) of 6 relative head motion parameters (movement from this time point to the next one) was calculated for each participant in pre- and post-treatment. No significant difference in head motion was found between pre- and post in PTSD ($p > .5$).

The Independent Component Analysis (ICA) was performed with the CONN-fMRI FC toolbox v18b in conjunction with SPM12 (Wellcome Department of Imaging Neuroscience, London, UK; <http://www.fil.ion.ucl.ac.uk/spm/>). ICA is a data-driven method that is based on a blind source separation algorithm. Unlike the seed-based method, which identifies the correlation between the seed region and other target brain regions, ICA examines multiple simultaneous voxel-to-voxel interactions of distinct networks in the brain, and separate independent patterns by maximizing the mutual spatial independence among components (Calhoun, Adali, Pearson, & Pekar, 2001). The resting state functional connectivity (rsFC) data were submitted to a subject-wise group ICA implemented in the CONN. A group-ICA methodology was used (Calhoun et al., 2001), with optional subject-level dimensionality reduction, concatenation across subjects, group-level Singular Value Decomposition (SVD) for dimensionality reduction, a fastICA algorithm for group-level IC, and a GICA3 for

subject-level back-projection. Group ICA was performed with 20 factors and a dimensionality reduction of 64. To compare these spatial patterns for one particular network of interest for two sessions, masks of each network were generated and only activations within each component network were examined.

To examine changes in brain functional connectivity between the two sessions, a paired sample t test was performed (whole brain FDR correction). The statistical results were used to determine the brain regions showing significant differences in each network.

2.6 | T1 image processing

Voxel-Based Morphometry (VBM) analysis was performed using SPM12 (<http://www.fil.ion.ucl.ac.uk/spm>) with MATLAB R2018a. Pairwise longitudinal registration was carried out using inverse-consistent rigid-body registrations to generate the subject average images. Each individual image was bias-corrected and then segmented into GM, WM, and cerebrospinal fluid (CSF) using standard Tissue Probability Maps (TPM) in SPM12. Images were normalized to an isotropic voxel size of 1.5 mm. This was followed by partial volume estimation, which provides a more accurate segmentation. Afterwards, tissue segments were spatially normalized to a common reference space using DARTEL for all subjects (Ashburner, 2007). Images were modulated to guarantee that relative GM/WM volumes were preserved following the spatial normalization procedure. Finally, images were smoothed with an 8 mm full width at half maximum isotropic Gaussian kernel. Spatial pre-processing, smoothed, modulated normalized GM and WM datasets were used to perform statistical analysis. Total gray matter volume (GMV) was estimated for each participant and used as a covariate in the second-level statistical test.

For statistical analysis of the longitudinal study, paired t tests were conducted. The activated regions of regional GMV were extracted using MarsBar (Matthew, Jean-Luc, Romain, & Jean-Baptiste, 2002).

2.7 | DTI image processing

All data analyses were performed in Functional MRI of the Brain Software Library (FSL) version 6.0 (Analysis Group, FMRIB, Oxford, UK, <http://fsl.fmrib.ox.ac.uk>). Diffusion weighted images were corrected for motion using Eddy Correct and non-brain tissue was removed using Brain Extraction Tool. Brain-extracted images were then visually inspected to confirm accurate results. Fractional anisotropy (FA) maps, the common DTI measure that describes the degree of anisotropy of a diffusion process, were created using DTIfit and input into Tract-Based Spatial Statistics (TBSS) to obtain a projection of all participants' FA data onto a mean FA skeleton. All participants' FA data were non-linearly aligned to a common space (FMRIB58_FA). Then, the mean FA image was created and thresholded ($FA > 0.2$) to create the mean FA skeleton. Next, each participant's FA data were projected onto the thresholded mean FA skeleton. Voxel-wise statistical analysis of the white matter skeleton was performed using *Randomise*,

FSL's nonparametric permutation inference tool. A one-sample, paired t test assessed pre- versus post-treatment differences. Threshold-free cluster enhancement was used to correct for multiple comparisons ($p < .05$). TBSS was performed for mean diffusivity (MD), which describes the rotationally invariant magnitude of water diffusion within brain tissue. Non-linear registration estimated from the FA images was applied to MD data and each participant's MD image was projected onto the mean FA skeleton before applying voxel-wise statistics. White matter regions were identified with Johns Hopkins University's white matter atlas available in FSL.

2.8 | Statistical analysis

To assess whether observed post-treatment brain changes were associated with the clinical changes, we first carried out correlation analysis between brain changes (rsFC and sMRI) and changes in clinical scores (CAPS-5, HAM-D, PCL, and BDI), controlled for scanners. Second, to examine whether these neural markers might serve as baseline predictors of clinical improvement, we carried out correlation analysis to examine whether the baseline brain findings (rsFC and sMRI) predicted changes in clinical scores (CAPS-5, HAM-D, PCL, and BDI). Bonferroni correction was used to correct for the four comparisons.

3 | RESULTS

3.1 | Examining pre- to post-treatment brain changes

3.1.1 | Resting state functional connectivity (rsFC) changes

Whole-brain baseline rsFC revealed six networks of interest using data-driven group ICA: left and right executive control network (ECN), anterior and posterior default mode network (a-DMN, p-DMN), salience network (SN), and the basal ganglia network (BGN) (Figure 1). However, only BGN connectivity showed changes from pre- to post-treatment. Specifically, within-BGN, one cluster (caudate) showed a significant increase in rsFC after treatment (Figure 2, Table 2). These changes of caudate rsFC were independent of whether subjects were taking medications.

3.1.2 | sMRI T1 VBM changes

Whole-brain sMRI showed reduction of regional GMV in the thalamus and caudate with $p < .05$ whole brain FWE correction (Table 3). There

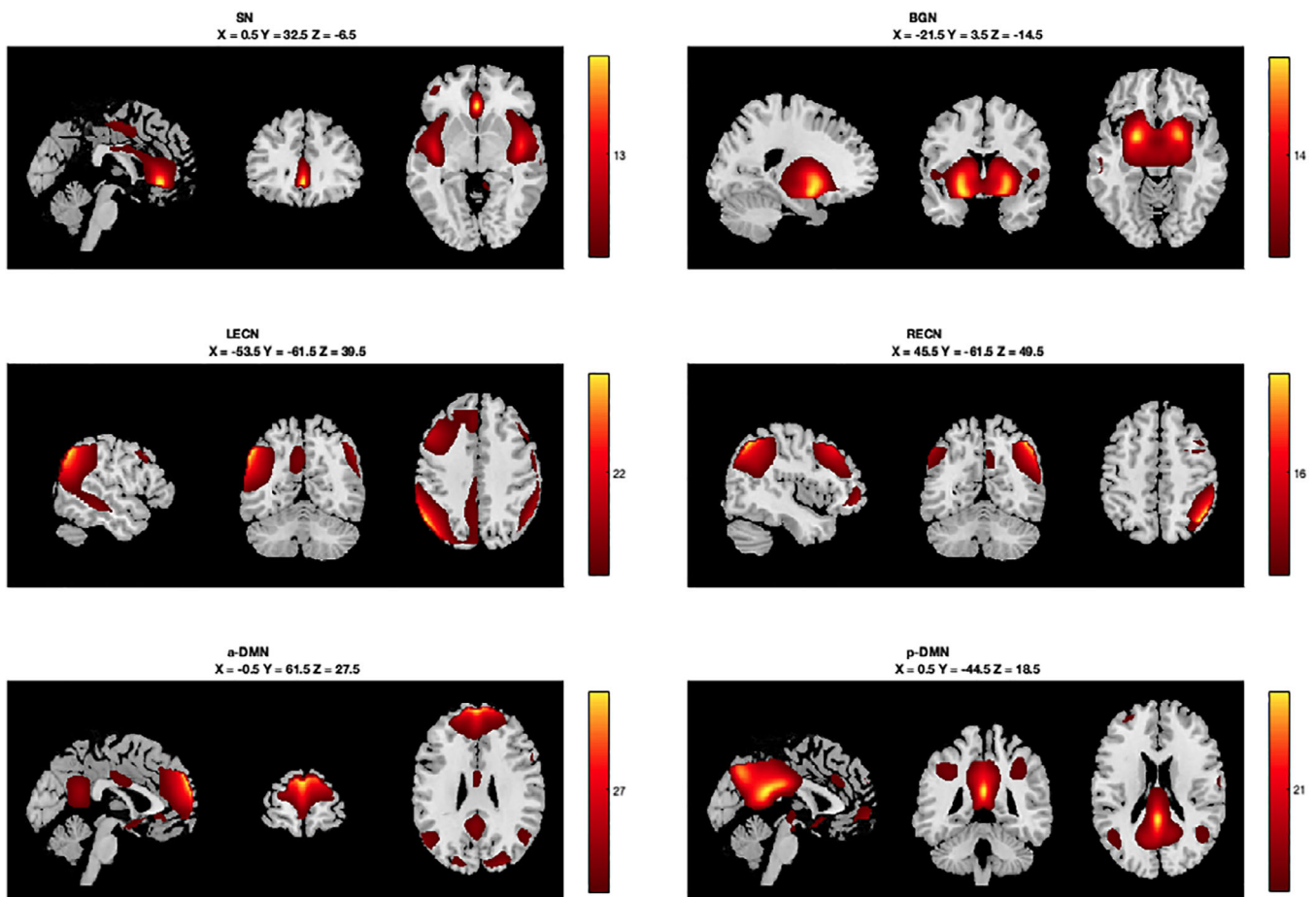


FIGURE 1 Networks identified by ICA

was no significant difference in regional GMV in the thalamus and caudate cluster between subgroups taking and not taking medication.

3.1.3 | DTI TBSS changes

Whole-brain DTI paired pre/post-treatment *t* test yielded no significant clusters in the FA and MD analyses.

3.2 | Examining associations between pre- to post-treatment brain changes and clinical change

3.2.1 | Clinical changes

Patients had significantly lower scores on the CAPS-5 ($p = .002$ post; $p = .005$ FU), PCL ($p < .001$ post; $p = .008$ FU), and BDI ($p = .007$ post;

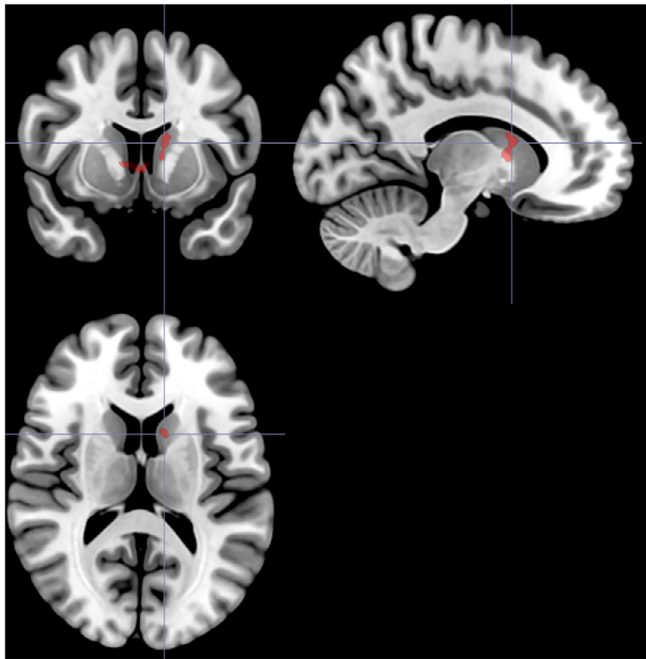


FIGURE 2 Within-BGN showed significant increased connectivity after treatment

TABLE 2 Resting state functional connectivity (rsFC) changes from pre- to post-treatment

Regions	$p(\text{FWE-corr})$	$p(\text{FDR-corr})$	Cluster size	<i>T</i>	<i>Z</i>	<i>x,y,z</i>
Caudate	0.245	0.044	37	5.33	4.03	[-12 8 16]
				4.8	3.77	[-10 6 6]
	0.048	0.028	56	5.03	3.88	[4 8 0]
				4.19	3.43	[8 18 2]

TABLE 3 Whole-brain sMRI showed reduction of regional GMV

Regions	$p(\text{FWE-corr})$	$p(\text{FDR-corr})$	Cluster size	<i>T</i>	<i>Z</i>	<i>x,y,z</i>
Caudate	0.009	0.022	959	6.58	4.64	[12 -3 18]
Thalamus				4.93	3.87	[6 -10 15]

$p < .001$ FU) at post treatment and follow-up compared with baseline (Table S1). There was no significant change on HAM-D at post treatment and FU compared with baseline ($p = .052$ post, $p = .14$ FU).

3.2.2 | Neural changes and correlation with clinical outcomes

Post-treatment: There was a significant positive correlation between increased caudate rsFC and improved PCL scores (pre-post; $r = .60$, $p = .015$; Figure 3), indicating that increased caudate rsFC was associated with reduction in PCL scores at post treatment. No significant correlation of changes in the caudate sMRI and clinical scores was found post treatment.

3 months follow-up: Changes in caudate rsFC were significantly correlated with the extent of changes seen in both PCL ($r = .58$, $p = .019$) and BDI ($r = .54$, $p = .03$) scores from FU, compared to pre-treatment baselines. Specifically, changes in caudate rsFC were positively associated with sustained improvement in BDI seen at 3 months after the conclusion of treatment.

No significant correlation of changes in the caudate sMRI and clinical scores were found at 3-months FU.

3.2.3 | Associations between baseline neural predictors and clinical improvement at post-treatment and 3-month FU

Post-treatment: baseline caudate rsFC was positively correlated with improvements in PCL scores (pre-post; $r = .69$, $p = .006$; Figure 4). Specifically, the baseline caudate rsFC was positively correlated with baseline PCL ($r = .34$, $p = .23$), but negative correlated with post-treatment PCL ($r = -.26$, $p = .38$). We also found a trending towards positive correlation with improved scores on both the HAM-D ($r = .57$, $p = .035$ before correction) and BDI instruments ($r = .61$, $p = .02$ before correction). Higher baseline rsFC seen in the caudate before beginning treatment was positively correlated with greater post-treatment symptom improvement. Baseline caudate sMRI was not significantly correlated with clinical improvement at post-treatment.

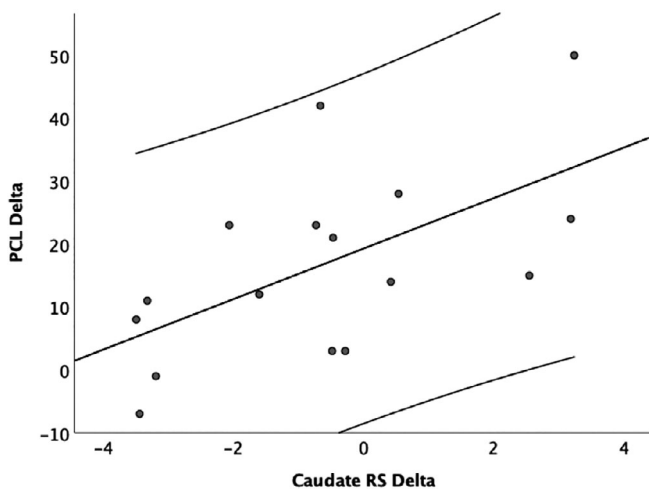


FIGURE 3 The association between changes of resting state caudate connectivity and PCL changes

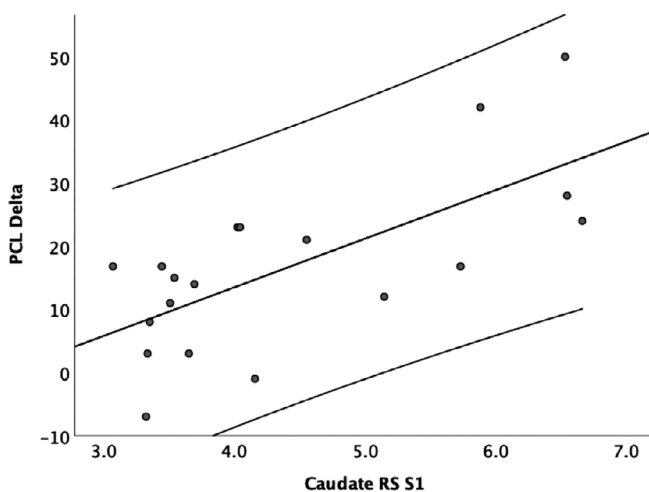


FIGURE 4 Baseline resting state caudate connectivity predicting the PCL changes

3 months follow-up: baseline caudate rsFC was positively correlated with improvements in CAPS-5 ($r = .658, p = .011$), HAM-D ($r = .73, p = .003$), PCL ($r = .82, p < .001$), and BDI scores (pre-FU; $r = .67, p = .008$). Specifically, for all 4 clinical measures, the baseline caudate rsFC was positively correlated with baseline clinical measures, but negative correlated with clinical measures at follow-up (see Table S2 for each correlation).

In summary, baseline caudate rsFC was positively associated with changes in these indices measured at 3 months after the conclusion of EAT. There was no correlation between baseline sMRI and clinical scores at 3-month follow-up evaluation.

4 | DISCUSSION

Our multimodal MRI study of neural mechanisms and predictors of treatment response to EAT for patients with PTSD revealed

significant increase in caudate connectivity within the BGN network after treatment, as well as a reduction in the regional GMV of the thalamus and caudate. Increased caudate connectivity following treatment was associated with greater symptom reduction. Higher levels of baseline caudate connectivity was predictive of greater symptomatic reduction at the end of EAT, and was even more strongly correlated with sustained improvement seen at three-month follow-up after the conclusion of EAT.

Overall, our findings suggest that the limbic-basal ganglionic reward system may be modified by the eight-week-long EAT employed for the treatment of PTSD. Previous resting-state fMRI studies have identified the basal ganglia/limbic network at rest subsumes areas of the striatum, thalamus, and amygdala (Cole et al., 2012; Moussa, Steen, Laurienti, & Hayasaka, 2012). These regions have been strongly implicated in dopaminergic reward and learning processes (Sesack & Grace, 2010). The caudate nucleus is involved in reward anticipation and response. Abnormal functioning within the caudate has been documented in depression (Eshel & Roiser, 2010), and substance abuse (Volkow, Baler, & Goldstein, 2011), and most relevant for the present study, in PTSD (Elman et al., 2009; Sailer et al., 2008). Increased bilateral nucleus accumbens to right dorsomedial prefrontal cortex connectivity was associated with higher levels of anhedonia in trauma-exposed individuals (Olson, Kaiser, Pizzagalli, Rauch, & Rosso, 2018). This evidence suggests that connectivity abnormalities within the basal ganglia/limbic network may contribute to aberrant reward processing (Schmidt et al., 2015) in PTSD. The current findings suggest that these abnormalities may be successfully targeted by EAT.

Results further show that baseline rsFC within the BGN predicted treatment outcomes. Specifically, baseline caudate rsFC was not correlated with PCL at independent time points, but rather was positively correlated with improvement in PCL scores after treatment. This PCL score improvement correlation was stronger three months later and correlated with CAPS-5, HAM-D, and BDI symptom improvement. These findings suggest that a stronger caudate connectivity before treatment predicted better outcomes post treatment, which strengthened with time. Clinical scores improvement also correlated with increase of caudate. Strengthening of caudate rsFC was significantly correlated with PCL score improvements post treatment and at three-month follow-up. Three-month follow-up caudate rsFC improvement was also correlated with BDI score improvements. The findings suggest that while baseline caudate connectivity predicts better treatment outcome overall, particularly at three-month follow-up, mainly PCL score improvement correlated with strengthening of caudate connectivity itself. This suggests that level of baseline striatal connectivity-and rsFC in the caudate nucleus, in particular-may be a stronger predictor of EAT outcome. Furthermore, EAT is changing the rsFC of the Caudate which in turns improves the symptoms of the patient.

T1 findings show a reduction in caudate regional GMV. Together with the pre- to post-treatment connectivity changes, these results appear to indicate a pruning effect in the caudate from pre- to post-treatment. In depression, lower caudate volume has been associated with anhedonia and greater depressive severity

(Pizzagalli et al., 2009). Felmingham et al. (2014) suggested that impaired recruitment of the striatum may partially mediate the anhedonia characteristic of depression, addictive disorders, and PTSD (Felmingham et al., 2014). Studies have reported significantly increased gray matter density in left caudate in individuals with PTSD (Yang, Wu, Bai, Hou, & Gao, 2009). In addition, we also found a reduction in thalamus regional GMV. The thalamus, along with the caudate and putamen, comprise an essential part of the limbic cortico-striatal-thalamocortical circuit which is involved in fear and anxiety. Dysfunction in this circuit appears to impair cognitive control, self-regulation of behavior, and emotion. Thalamus volume abnormalities appear to be a consistent finding across depression studies (Espinoza Oyarce, Shaw, Alateeq, & Cherbuin, 2020). Our findings of decreased GMV of the caudate and thalamus at post-treatment supports the idea that patients with PTSD might have a disrupted, dysfunctional reward circuitry, including cortico-striatal-thalamocortical circuits, that might be alleviated through EAT.

The present findings revealed no fractional anisotropy (FA) changes from pre- to post-treatment. Previous findings in adult and youth with PTSD have suggested structural connectivity changes in white matter (WM) tracts (Fani et al., 2012; Kennis et al., 2015). Both increased and reduced FA in the cingulum bundle has been reported in patients with PTSD (Sanjuan, Thoma, Claus, Mays, & Caprihan, 2013; Zhang et al., 2012). Decreased FA in the corpus callosum, most prominently in the anterior and posterior midbody, the isthmus, and splenium were reported in a meta-analysis summarizing findings from samples of children with PTSD and adult-onset PTSD with childhood trauma (Siehl, King, Burgess, Flor, & Nees, 2018). For adult-onset PTSD with traumatic experience in adulthood, changes were found in FA in the anterior and posterior part of the cingulum, the superior longitudinal fasciculus, and frontal regions. These findings suggest that changes in white matter microstructure vary depending on traumatic experience. Changes of FA have also been reported in PTSD following trauma-focused therapy, including increases in FA in both the dorsal cingulum and posterior cingulum bundle (Zhang et al., 2012) in patients with chronically symptomatic PTSD (Kennis et al., 2015). Our data showed no FA changes, perhaps suggesting that EAT targets different neural circuitry than trauma-focused therapy. Future studies with a larger sample size could help to investigate whether traumatic experience may moderate the effect of EAT treatment on DTI changes.

This study is the first to evaluate both structural and functional brain changes associated with EAT, in particular, with a specific manualized approach to the application of EAT to the treatment of PTSD. Still, several limitations must be acknowledged. The main limitation is that this study was part of an open trial design and lacks a control group. This design precludes comparison of neuroimaging results seen with EAT with a non-treatment control group or with subjects undergoing other interventional immersive treatment therapies to assess whether they activate similar or different circuits. Another important limitation is the small sample size, which lacked statistical power to compare patient subgroups based on adjunctive treatment, trauma type, etc. Future neuroimaging studies treating larger samples in controlled designs are needed to replicate these findings and allow to

examine to potential role of demographics, trauma history and concurrent treatments in the study of neural mechanisms and predictors of treatment outcome.

5 | CONCLUSION

This study demonstrated that an eight-week course of EAT in subject suffering with PTSD was associated with changes in the post-treatment resting-state reward networks in the caudate and volume changes in the thalamus and caudate on neuroimaging. Furthermore, baseline, pretreatment caudate connectivity significantly correlated with clinical improvement, suggesting enhanced reward responsiveness may underlie the improvements seen in subjects with PTSD after EAT treatment. PTSD is a heterogeneous disorder that has been shown to include symptoms related to diminished reward circuitry responsivity that can impact social and emotional responses. Improving reward circuitry might be a significant, initial step to finding new and enhanced methods to treat this complex disorder.

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CONFLICT OF INTEREST

The authors declared no potential conflicts of interest with respect to the research, authorship, and/or publication of this article.

AUTHOR CONTRIBUTIONS

Xi Zhu involved in formal MRI analysis, writing—original draft. Benjamin Suarez-Jimenez and Amit Lazarov involved in writing—review and editing. Sigal Zilcha-Mano involved in regression analysis, writing—review and editing. Shay Arnon, Maja Bergman, and Matthew Ryba involved in data curation and project administration. Ari Lowell involved in project administration and supervision. Allan J. Hamilton, Jane F. Hamilton, and John C. Markowitz involved in conceptualization, writing—review and editing. J. Blake Turner involved in writing—review and editing. Prudence W. Fisher involved in conceptualization, data curation, funding acquisition, investigation, project administration, and supervision. Yuval Neria involved in conceptualization, funding acquisition, investigation, resources, supervision, writing—review and editing.

DATA AVAILABILITY STATEMENT

The data that support the findings of this study are available on request from the corresponding author.

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