The functional and structural characteristics of the emotion network in alexithymia

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Background: Alexithymia is a multifaceted personality trait characterized by emotional dysfunction.

Methods: In this study, the functional and structural features of the emotion network in alexithymia were investigated using resting-state functional MRI (rsfMRI), voxel-based morphometry (VBM), functional connectivity (FC) analysis, and diffusion tensor imaging (DTI). Alexithymic and non-alexithymic students were recruited from the local university. The intrinsic neural activity and gray matter density of the brain regions in the emotion network were measured using rsfMRI and VBM; the FC and structural connectivity of the brain regions in the emotion network were measured using FC analysis and DTI.

Results: The altered intrinsic neural activity in V1, rostral dorsal anterior cingulate cortex, and left amygdala, and the weak FC between V1 and left superior temporal gyrus and V1 and left paracentral lobule in alexithymia subjects were identified. However, no alteration of the structure and structural connectivity of the emotion network was identified.

Conclusion: The results indicated that the development of alexithymia might have been caused only by slight alteration of the neural activity. Furthermore, the results suggest that noninvasive treatment technologies for improving the brain activity are suitable for alexithymic individuals.

Keywords: emotion network, rsfMRI, VBM, functional connectivity, structural connectivity

Introduction

Alexithymia is a personality trait characterized by difficulties in identifying and describing one's own emotional feelings and an external thinking style.¹ Alexithymia is associated with increased risks for psychosomatic complaints, autism, anxiety, and depression disorder.²⁻⁵ Moreover, impaired identification and expression of emotions (the core symptoms of alexithymia) may play a mediating role in these disorders.^{2-4,6,7} Clarifying the neural mechanism of emotional dysfunction in alexithymic individuals can improve the understanding of this personality trait and its relationships with psychosomatic disorders.

The neurobiological basis of the emotion function has been a research focus for a long time. Through a meta-analysis study, an emotion network covering 21 brain areas was identified (Table 1).⁸ Notably, the distinct structural or functional abnormalities of the brain regions in the emotion network have been linked to the specific emotional symptoms of different mental disorders.⁹ For instance, in the case of schizophrenic individuals, unusual activities are seen in the amygdala, insula, and ventral striatum in response to emotional stimulus, and a reduced gray matter (GM) volume is observed in the frontal cortex.^{10,11} In the case of depressed patients, abnormal activities are seen in the amygdala, anterior insula, ventral striatum, and prefrontal cortex (PFC) in response to emotional stimuli, and reduced volume is observed in the amygdala,

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Table I The MNI coordinates of the selected ROIs in the emotion network

Name	X	Y	Z	Radius (mm)
Visual area 8/bilateral superior cerebellum (V8/sCB, Bi)	-11	-62	-20	12
Bilateral middle temporal complex (MT+, Bi)	-28	-69	9	12
Right visual area 4 (V4, R)	44	-79	-4	12
Right lateral occipital cortex/temporal lobe (latOCC/Temp, R)	49	-56	-2	12
PCC	1	-54	25	12
Primary visual cortex (VI)	-7	-89	0	12
Dorsal medial prefrontal cortex (dmPFC)	-2	51	29	12
Pregenual anterior cingulate cortex (pgACC)	1	39	2	12
Rostral dorsal anterior cingulate cortex (rdACC)	0	29	25	12
Ventral anterior insula/frontal operculum/right temporal pole, bilateral posterior Orbitofrontal cortex (valns/frOP/TP, RpOFC, Bi)	38	19	-11	12
Anterior insula, posterior orbitofrontal cortex (alns, pOFC, Bi),	-40	23	-6	12
Bilateral ventral anterior insula/temporal cortex/orbitofrontal cortex (valns/TC/OFC, Bi)	-7	12	-23	12
Bilateral middle insula, right dorsal putamen (midIns, Bi dPut, R)	21	0	2	12
Left ventral Striatum, middle insula, hippocampus (vStr, midIns, HCMP, L)	-36	0	-8	12
Left Basolateral Amygdala (Amy_L)	-23	-3	-21	12
Hypothalamus (Hy)	-7	-5	-13	12
Periaqueductal gray/thalamus (PAG/Thal)	-1	-23	2	12
Bilateral amygdala, ventral striatum, ventral globus pallidus, thalamus (Amy vStr vGP Thal, Bi)	11	-6	-10	12
Right frontal operculum (frOP, R)	46	24	6	12
Right inferior frontal gyrus (IFG, R)	49	20	19	12
Left pre-supplementary motor area, inferior frontal gyrus (preSMA, IFG, L)	-23	13	37	12

Abbreviations: MNI, Montreal Neurological Institute; PCC, posterior cingulate cortex; ROI, region of interest.

ventral striatum, anterior cingulate gyrus (ACC), PFC, and hippocampus.^{12–14} Moreover, abnormal neural activity in the amygdala, insula, ACC, medial PFC, and posterior cingulate cortex (PCC) is observed in alexithymic individuals.¹⁵ Two voxel-based morphometry (VBM) studies revealed decreased GM density in the ACC and left amygdala and increased GM density in the right posterior insula and right cingulate cortex in alexithymic individuals.^{16,17}

It is notable that alexithymia is related to abnormal activity in multiple brain regions instead of only a single region. Multiple brain regions have to cooperate with each other to maintain the brain functions. Therefore, it is reasonable to hypothesize that the occurrence of alexithymia is associated with not only functional or structural abnormalities of some brain regions in the emotion network but also some abnormal functional connectivity (FC) or structural connectivity among these brain regions. Resting-state functional MRI (rsfMRI) analyses on FC provide a powerful method to investigate the FC among brain regions. Liemburg found abnormal FC levels among several brain regions (such as the medial PFC, occipital areas, insula cortex, and ACC) in alexithymic individuals.¹⁸ Diffusion tensor imaging (DTI) is a highly sensitive method for measuring the structural integrity of the white matter tracts that connect brain regions.¹⁹ Structural connectivity is also an important index to investigate the

cooperative effects among the brain regions in the emotion network. A DTI study conducted by Ho et al suggested that the structural connectivity of the superior longitudinal fasciculus was significantly correlated with alexithymia.²⁰ These studies partly supported the hypothesis presented earlier. Unfortunately, Liemburg's study investigated only the brain regions in the default network. Ho et al focused only on the corpus callosum and superior longitudinal fasciculus. In order to verify the hypothesis proposed earlier, it is necessary to investigate the functional and structural features of the emotion network in alexithymic individuals.

In order to investigate the structural and functional features of the emotion network in alexithymia, the present study included four programs: first, examining the neural intrinsic activity of the brain regions in the emotion network using rsfMRI; second, examining the structural features of these brain regions using VBM analysis; third, examining the FCs among the brain regions showing abnormal neural intrinsic activity or abnormal structural features in the emotion network via FC analysis; and fourth, examining the structural connections among the brain regions showing abnormal FC in the emotion network via DTI analysis. Holistic knowledge on the emotion network will contribute to a deeper understanding of the neural nature of alexithymia.

Methods

Subjects

All the participants were recruited from among the freshmen students of a local university. All the students underwent a mental health checkup (including alexithymia measurement, see the “Psychological measurements” section) during the university entrance examination in 2015. The participants satisfied the following inclusive criteria: right-handed and physically healthy individuals without histories of traumatic brain injuries, substance abuse, or mental disorders. Participants were excluded if they had contraindications to MRI scans or had consumed any psychotropic medications within 6 months before the study. The eligible participants with alexithymia were randomly chosen, telephonically contacted, and assigned to the alexithymia group (Alex group). The eligible non-alexithymia participants, who matched the alexithymia subjects with regard to age and sex, were randomly chosen, contacted by telephone, and assigned to the healthy control group (HC group). All the protocols were approved by the Research Ethics Review Board of Hangzhou Normal University. Thus, 42 participants (21 each in the Alex and HC groups) were recruited in the present study. The participants were informed of the study protocol and signed the informed consent form.

Psychological measurements

The characteristics of alexithymia were evaluated by the Toronto Alexithymia Scale-20 (TAS-20), which is a 20-item self-report scale, rating from 1 (strongly disagree) to 5 (strongly agree). The participants with TAS-20 total scores >60 and <52 were assigned to Alex group and HC group, respectively.²¹

MRI measurements

All MRI scan processes were performed with a 3T whole-body MR imager (GE MR750). The rsfMRI scans were acquired: sequence = gradient recalled echo-echo planar images, axial slices, 43 slices, repetition time (TR)/echo time (TE)=2,000/30 ms, field of view (FOV)=220×220 mm², resolution=64×64, flip angle (FA)=90°, total scan time=480 s. Participants were instructed to simply rest with their eyes closed, not to think of anything in particular, and not to fall asleep. The 3D T1-weighted images were acquired: sagittal slices, 180 slices, TR/TE=8,100/3 ms, resolution=256×256, FOV=256×256 mm², thickness/gap=1/0 mm, FA=8°, total scan time=305 s. Lastly, DTI was acquired: spin-echo echo-planar imaging sequence, 67 axial slices, TR/TE=8,600/minimum ms, FOV=192×192 mm², matrix=128×128, number of shots=1, slice thickness=1.5 mm. Diffusion direction: TENSOR, 30 directions, b-value=1,000 s/mm²,

NEX (number of excitations)=2, number of T2 images=4, voxel size=1.5×1.5×1.5 mm³, total scan time=593 s.

Data processing

rsfMRI data analysis and statistical analysis

Based on the MatLab software, data preprocessing was performed by the Data Processing Assistant for Resting-State fMRI (DPARSF, <http://www.restfmri.net>) and the rsfMRI data analysis toolkit (REST, <http://www.restfmri.net>).^{22,23} The images were corrected for slice-time differences, realigned for head motion correction, co-registered to T1, and spatially smoothed, according to a previous study's suggestion.²⁴ The amplitude of low-frequency fluctuation (ALFF) map was calculated and the ALFF values of 21 regions of interest (ROI) in the emotion network were extracted. The ROI selection was based on the previous studies (Table 1).^{8,25} The ALFF values of the two groups were compared using an independent *t*-test by SPSS19.0. Differences were considered significant at a two-sided *p*-value of 0.05.

VBM data analysis and statistical analysis

Data were processed using the VBM toolbox (VBM8, <http://dbm.neuro.uni-jena.de/vbm.html>) in a statistical software package (SPM8, <http://www.fil.ion.ucl.ac.uk/spm/>). Then preprocessing was undertaken: the T1-weighted images were bias-corrected and segmented into GM, white matter (WM), and cerebrospinal fluid (CSF); Diffeomorphic Anatomical Registration Through Exponentiated Lie Algebra (DARTEL) create template; DARTEL existing template; normalize to Montreal Neurological Institute space; spatial smoothing with 8 mm full-width at half-maximum Gaussian kernel; the GM density of 21 ROIs in the emotion network was extracted. The GM density of the two groups was compared using independent *t*-test by SPSS19.0. Differences were considered significant at two-sided *p*-value of 0.05.

Functional connectivity analysis and statistical analysis

The FC between each ROI in the emotion network and other brain regions in the whole brain was determined using REST software. According to the previous studies, the preprocessing was undertaken: time series extraction, Pearson's correlation analysis of ROIs and whole-brain voxel time series (WM, CSF, and six head motion parameters were covariates), Fisher's Z-transformation, voxel-by-voxel based comparisons of FC of 21 ROIs between two groups by two-sample *t*-test using REST, and AlphaSim correction ($p<0.001$, AlphaSim correction, cluster size >20).^{26,27} The result was visualized by BrainNet Viewer (<https://www.nitrc.org/projects/bnv/>).²⁸

Table 2 Comparison of the demographic variables and psychological measurements between the Alex and the HC groups

Variables	Alex group, n=21	HC group, n=21	p-value
Demographic data			
Males, n (%)	11 (52.4)	11 (52.4)	
Age (years)	18.33 (1.06)	18.62 (1.07)	p=0.39
TAS-20 scores*	58.10 (3.2)	41.00 (7.75)	p<0.001

Notes: Data are presented as percentage/ratio or mean $\pm$ SD; *p<0.05.

Abbreviations: Alex, alexithymia; HC, healthy control; TAS-20, Toronto Alexithymia Scale-20.

Structural connectivity analysis and statistical analysis

The DTI data were processed using the Pipeline for Analyzing Brain Diffusion Images (PANDA, <http://www.nitrc.org/projects/panda/>) and DTI Studio (<https://www.mristudio.org/>). The DTI data procession included transferring Digital Imaging and Communications in Medicine to Neuroimaging Informatics Technology Initiative, image registration, normalizing, and smoothing. Maps of FA were computed from the DTI data. Based on the FC analysis results in the present study, the FA values were extracted using the Fiber Assignment by Continuous Tracking (FACT) algorithm.²⁹ The FA values of the two groups were compared using independent *t*-test by SPSS.

Differences were considered significant at two-sided *p*-value of 0.05. The result was visualized by BrainNet Viewer.

Results

Demographic characteristics and psychological measurements of the students

The results indicated no significant difference in the demographic variables between the two groups. TAS-20's global and three subscales' scores in the Alex group were significantly higher than those in the HC group (Table 2).

Comparison analysis of rsfMRI between two groups

The comparative analysis revealed significant differences in ALFF values of V1, rostral dorsal anterior cingulate cortex (rdACC), and Amy_L between the Alex and the HC groups (Table 3, Figure 1).

Comparison analysis of gray matter density between two groups

The results revealed no significant difference in GM density between the Alex and the HC groups (Table 4).

Table 3 Brain regions in the emotion network showing significant differences on ALFF between the Alex and the HC groups

	ALFF value		t-value	p-value
	Alex group	HC group		
V8/sCB, Bi	0.996 $\pm$ 0.015	0.991 $\pm$ 0.021	0.832	0.410
MT+, Bi	1.001 $\pm$ 0.016	1.000 $\pm$ 0.040	0.025	0.980
V4, R	0.886 $\pm$ 0.014	0.893 $\pm$ 0.012	-1.574	0.123
latOCC/Temp, R	0.992 $\pm$ 0.017	0.997 $\pm$ 0.029	-0.703	0.486
PCC	1.003 $\pm$ 0.017	0.993 $\pm$ 0.033	1.189	0.242
V1*	1.001 $\pm$ 0.019	1.019 $\pm$ 0.028	-2.477	0.018
dmPFC	1.004 $\pm$ 0.019	0.999 $\pm$ 0.022	0.744	0.461
pgACC	0.993 $\pm$ 0.027	0.995 $\pm$ 0.031	-0.254	0.801
rdACC*	1.002 $\pm$ 0.022	0.984 $\pm$ 0.022	2.587	0.013
valns/frOP/TP, RpOFC, Bi	0.998 $\pm$ 0.010	0.989 $\pm$ 0.019	1.784	0.082
alns, pOFC, Bi	0.998 $\pm$ 0.015	0.997 $\pm$ 0.022	0.125	0.901
valns/TC/OFC, Bi	0.990 $\pm$ 0.015	0.994 $\pm$ 0.021	-0.769	0.446
midlns, Bi dPut, R	0.993 $\pm$ 0.024	0.993 $\pm$ 0.023	0.073	0.942
vStr, midlns, HCMP, L	0.995 $\pm$ 0.012	1.001 $\pm$ 0.021	-1.048	0.301
Amy_L*	0.989 $\pm$ 0.019	1.002 $\pm$ 0.023	-2.141	0.038
Hy	0.990 $\pm$ 0.020	0.999 $\pm$ 0.019	-1.498	0.142
PAG/Thal	0.993 $\pm$ 0.023	1.001 $\pm$ 0.017	-1.321	0.194
Amy vStr vGP Thal, Bi	0.995 $\pm$ 0.015	1.000 $\pm$ 0.014	-1.065	0.293
frOP, R	1.004 $\pm$ 0.014	0.993 $\pm$ 0.026	1.702	0.097
IFG, R	1.004 $\pm$ 0.011	0.996 $\pm$ 0.021	1.451	0.155
preSMA, IFG, L	1.004 $\pm$ 0.012	1.004 $\pm$ 0.010	0.000	1.000

Notes: Data are presented as percentage/ratio or mean $\pm$ SD; *p<0.05.

Abbreviations: Alex, alexithymia; alns, pOFC, anterior insula, posterior orbitofrontal cortex; ALFF, amplitude of low-frequency fluctuation; Amy vStr vGP Thal, Bi, Bilateral amygdala, ventral striatum, ventral globus pallidus, thalamus; Amy_L, Left basolateral amygdala; dmPFC, dorsal medial prefrontal cortex; frOP, R, right frontal operculum; HC, healthy control; Hy, hypothalamus; IFG, R, right inferior frontal gyrus; latOCC/Temp, R, right lateral occipital cortex/temporal lobe; midlns, Bi dPut, R, Bilateral middle insula, right dorsal putamen; MT+, Bi, bilateral middle temporal complex; PAG/thal, periaqueductal gray/thalamus; PCC, posterior cingulate cortex; pgACC, pregenual anterior cingulate cortex; preSMA, IFG, L, left pre-supplementary motor area, inferior frontal gyrus; rdACC, rostral dorsal anterior cingulate cortex; V4, R, right visual area 4; V8/sCB, Bi, visual area 8/bilateral superior cerebellum; valns/frOP/TP, RpOFC, Bi, ventral anterior insula/frontal operculum/right temporal pole, bilateral posterior orbitofrontal cortex; valns/TC/OFC, Bi, bilateral ventral anterior insula/temporal cortex/orbitofrontal cortex; vStr, midlns, HCMP, L, left ventral striatum, middle insula, hippocampus.

Abbreviations: Alex, alexithymia; alns, pOFC, anterior insula, posterior orbitofrontal cortex; ALFF, amplitude of low-frequency fluctuation; Amy vStr vGP Thal, Bi, bilateral amygdala, ventral striatum, ventral globus pallidus, thalamus; Amy_L, left basolateral amygdala; dmPFC, dorsal medial prefrontal cortex; frOP, R, right frontal operculum; HC, healthy control; Hy, hypothalamus; IFG, R, right inferior frontal gyrus; latOCC/Temp, R, right lateral occipital cortex/temporal lobe; midlns, Bi, bilateral middle insula, right dorsal putamen; MT+, Bi, bilateral middle temporal complex; PAG/thal, periaqueductal gray/thalamus; pgACC, pregenual anterior cingulate cortex; preSMA, IFG, L, left pre-supplementary motor area, inferior frontal gyrus; rdACC, rostral dorsal anterior cingulate cortex; V4, R, right visual area 4; V8/SCB, Bi, visual area 8/bilateral superior cerebellum; valns/frOP/TP, RpOFC, Bi, ventral anterior insula/frontal operculum/right temporal pole, bilateral posterior orbitofrontal cortex; valns/TC/OFC, Bi, bilateral ventral anterior insula/temporal cortex/orbitofrontal cortex; vStr, midlns, HCMP, L, left ventral striatum, middle insula, hippocampus.

area (SMA) in the alexithymia group, compared with the HC group (Table 5, Figure 2). The results revealed no significant difference in the FA value between the Alex and the HC groups (Table 5).

	GM density value		t-value	p-value
	Alex group	HC group		
V8/sCB, Bi	0.718 ± 0.045	0.734 ± 0.033	-1.345	0.186
MT+, Bi	0.2298 ± 0.027	0.235 ± 0.029	-0.602	0.551
V4, R	0.378 ± 0.039	0.358 ± 0.047	1.449	0.155
latOCC/Temp, R	0.438 ± 0.076	0.446 ± 0.048	-0.429	0.670
PCC	0.651 ± 0.055	0.668 ± 0.043	-1.107	0.275
VI	0.601 ± 0.044	0.721 ± 0.033	0.991	0.328
dmPFC	0.529 ± 0.048	0.529 ± 0.039	-0.028	0.978
pgACC	0.546 ± 0.077	0.547 ± 0.068	-0.072	0.943
rdACC	0.560 ± 0.095	0.574 ± 0.064	-0.542	0.591
valns/frOP/TP, RpOFC, Bi	0.557 ± 0.078	0.571 ± 0.046	-0.735	0.467
alns, pOFC, Bi	0.540 ± 0.051	0.528 ± 0.046	0.805	0.426
valns/TC/OFC, Bi	0.431 ± 0.025	0.429 ± 0.035	0.238	0.813
midlns, Bi dPut, R	0.426 ± 0.047	0.431 ± 0.032	-0.415	0.681
vStr, midlns, HCMP, L	0.655 ± 0.039	0.654 ± 0.042	0.103	0.919
Amy_L	0.700 ± 0.027	0.706 ± 0.028	-0.663	0.511
Hy	0.408 ± 0.025	0.406 ± 0.028	0.299	0.767
PAG/Thal	0.517 ± 0.048	0.503 ± 0.043	1.015	0.316
Amy vStr vGP Thal, Bi	0.348 ± 0.021	0.345 ± 0.027	0.499	0.621
frOP, R	0.422 ± 0.087	0.439 ± 0.044	-0.809	0.423
IFG, R	0.409 ± 0.075	0.426 ± 0.036	-0.946	0.350
preSMA, IFG, L	0.183 ± 0.035	0.182 ± 0.028	0.143	0.887

Abbreviations: Alex, alexithymia; alns, pOFC, anterior insula, posterior orbitofrontal cortex; Amy vStr vGP Thal, Bi, Bilateral amygdala, ventral striatum, ventral globus pallidus, thalamus; Amy_L, left basolateral amygdala; dmPFC, dorsal medial prefrontal cortex; frOP, R, right frontal operculum; GM, gray matter; HC, healthy control; Hy, hypothalamus; IFG, R, right inferior frontal gyrus; latOCC/Temp, R, right lateral occipital cortex/temporal lobe; midlns, Bi dPut, R, Bilateral middle insula, right dorsal putamen; MT+, Bi, bilateral middle temporal complex; PAG/thal, periaqueductal gray/thalamus; pgACC, pregenual anterior cingulate cortex; preSMA, IFG, L, left pre-supplementary motor area, inferior frontal gyrus; rdACC, rostral dorsal anterior cingulate cortex; V4, R, right visual area 4; V8/SCB, Bi, visual area 8/bilateral superior cerebellum; valns/frOP/TP, RpOFC, Bi, ventral anterior insula/frontal operculum/right temporal pole, bilateral posterior orbitofrontal cortex; valns/TC/OFC, Bi, bilateral ventral anterior insula/temporal cortex/orbitofrontal cortex; vStr, midlns, HCMP, L, left ventral striatum, middle insula, hippocampus.

Table 5 Comparison of the functional and structural connectivity between the Alex and the HC groups

	FC/FA value		t-value	p-value
	HC group	Alex group		
Functional connectivity (FC value)				
VI and Temporal_Sup_L (MNI: -57, -12, 15)*	0.12 ± 0.041	0.024 ± 0.011	-4.0639	0.002
VI and Paracentral_Lobule_L (MNI: -6, -18, 72)*	0.427 ± 0.075	0.337 ± 0.065	-4.140	0.001
Amy_L and Temporal_Sup_L (MNI: -57, -12, 15)*	0.314 ± 0.047	0.283 ± 0.044	-3.9274	0.003
Amy_L and SMA (MNI: 12, -9, 51)*	0.105 ± 0.035	0.038 ± 0.016	-4.771	<0.001
Structural connectivity (FA value)				
VI and Temporal_Sup_L	0.476 ± 0.115	0.411 ± 0.174	1.422	0.163
VI and Paracentral_Lobule_L	0.328 ± 0.214	0.280 ± 0.226	0.707	0.484
Amy_L and Temporal_Sup_L	0.315 ± 0.255	0.275 ± 0.245	0.507	0.615
Amy_L and SMA	0.390 ± 0.196	0.294 ± 0.237	1.439	0.158

Notes: Data are presented as percentage/ratio or mean ± SD; * $p < 0.05$.

Abbreviations: Alex, alexithymia; Amy_L, left basolateral amygdala; HC, healthy control; MNI, Montreal Neurological Institute; Paracentral_Lobule_L, left paracentral lobule; SMA, supplementary motor area; Temporal_Sup_L, left superior temporal gyrus.

Discussion

Alexithymia is a personality trait characterized by emotional dysfunction. This study examined the emotion network in people with alexithymia. Compared with the HC group, the Alex group displayed higher intrinsic neural activity in rdACC and lower activity in V1 and the Amy_L. The ACC includes two parts: the dorsal ACC (dACC) and the ventral ACC. The dACC is in charge of the detection and appraisal of emotional and social information.³⁰ Furthermore, alexithymia is linked to the blunted response of dACC to social rejection.³¹ The rdACC plays a role in regulating the startle reaction and mediating the anxiety emotion.³² In the present study, the part of dACC associated with alexithymia is more precisely located in the rdACC. The V1 is in charge of processing visual information and regulating pattern

recognition.³³ It has been reported that people with high-level alexithymia have difficulty in processing visual information from emotional body postures.³⁴ Therefore, a low activity in V1 is closely related to alexithymia. The Amy_L belongs to the limbic system and plays a key role in memory and emotional reactions.³⁵ Alexithymia is characterized by emotional dysfunction and contributes to social function impairment.^{36–38} A previous study reported that Amy_L was associated with alexithymia.³⁶ The present study also supports this finding.

Based on the fact that three brain regions show abnormal intrinsic neural activity in people with alexithymia, the present study further investigated the roles of these brain regions in the occurrence of alexithymia. As a multifaceted personality trait, alexithymia should be associated with the

Figure 2 The functional connectivity showing significant differences between the Alex and the HC groups.

Notes: Red balls indicate Alex group > HC group; blue balls indicate Alex group < HC group and the brain regions with weak functional connectivity; gray balls represent other brain regions in the emotion network. Blue lines indicate weak functional connectivity in Alex group, compared with HC group.

Abbreviations: Alex, alexithymia; dmPFC, dorsal medial prefrontal cortex; frOP, alns, anterior insula, frontal operculum; HC, healthy control; Hy, hypothalamus; IFG, inferior frontal gyrus; latOCC, lateral occipital cortex; MT+, middle temporal complex; PAG, periaqueductal gray; PCC, posterior cingulate cortex; PARA, paracentral lobule; pgACC, pregenual anterior cingulate cortex; pOFC, anterior insula, posterior orbitofrontal cortex; preSMA, L, left pre-supplementary motor area; STG, superior temporal gyrus; V8/sCB, visual area 8/ superior cerebellum; valns, ventral anterior insula; vGP, ventral globus pallidus; vStr, ventral striatum.

cooperative action of numerous brain regions. This study identified weak FC between V1 and Temporal_Sup_L and V1 and Paracentral_Lobule_L in alexithymia subjects. The Temporal_Sup_L is the primary auditory cortex, and it plays a key role in language comprehension.³⁹ The Paracentral_Lobule is the continuation of the precentral and postcentral gyrus and regulates motor and sensory innervations.^{40,41} Identifying and describing one's own feelings (including visual and auditory senses) are involved in language comprehension, motor controls and sensory information processing. Moreover, the present study presented a weak FC between Amy_L and Temporal_Sup_L and Amy_L and SMA in the alexithymia group. The SMA plays a key role in movement control, such as bimanual coordination and postural stability during standing or walking, which are all involved in identifying and describing emotional feelings.^{42–44} Therefore, the result of this study implies that the difficulties in identifying emotion and expression (the core symptoms of alexithymia) could be associated with weak FC between V1 and Temporal_Sup_L; V1 and Paracentral_Lobule_L; Amy_L and Temporal_Sup_L; and Amy_L and SMA. In the present study, no abnormal functional connection was identified between rdACC and other brain regions. This indicates that the dysfunction of rdACC alone is involved in the occurrence of alexithymia, and thus rdACC itself should be a therapeutic target for alexithymia. However, in terms of the V1 and left amygdala, the FC between V1 and Temporal_Sup_L; V1 and Paracentral_Lobule_L; Amy_L and Temporal_Sup_L; and Amy_L and SMA should be considered a therapeutic target.

According to the VBM and DTI analysis results, it is notable that no significant difference was identified in the gray matter or structural connectivity in the emotion networks of people with and without alexithymia. This suggests that the occurrence of alexithymia might only be due to a slight alteration in the neural activity of V1, rdACC, and left amygdala, and that it is not associated with any structural brain alteration. This is why alexithymia is considered just a personality trait rather than an emotional disorder. Most emotional disorders are caused not only by neural activity alteration but also by structural alteration. Furthermore, it implies that some noninvasive treatment technologies focusing on improving neural activity (such as transcranial magnetic stimulation, psychotherapy, and physical therapy) may be effective in helping people with alexithymia.

In conclusion, this study focused on the functional and structural characteristics of the emotion network in people with alexithymia. There were several limitations to this

study. The emotion network's functional characteristics should be investigated during emotionally motivated situations. Moreover, depression and anxiety, which are closely related to alexithymia, should be measured and analyzed. This study identified the altered intrinsic neural activity in V1, rdACC, and the left amygdala, as well as the weak functional connectivity between V1 and Temporal_Sup_L; V1 and Paracentral_Lobule_L; Amy_L and Temporal_Sup_L; and Amy_L and SMA in people with alexithymia. This study did not identify any structural alteration of the emotion network in people with alexithymia. However, the results of this study can help identify treatment options and may have heuristic value. To further investigate the features of the brain regions in people with alexithymia, a series of studies using different neural imaging methods is necessary.

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Author contributions

Han Dai designed the study, managed the imaging data selection and analysis, and wrote the first draft of the manuscript. Mei Li, Minjun Mei and Xiaofei Sun undertook the psychology measurement and statistical analysis. All authors contributed toward data analysis, drafting and critically revising the paper, gave final approval of the version to be published, and agree to be accountable for all aspects of the work.

Disclosure

The authors report no conflicts of interest in this work.

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