

Impaired Interhemispheric Synchrony in Bronchial Asthma

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Research Article

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Abstract

Objective: Converging evidence demonstrated that bronchial asthma (BA) individuals with intermittent hypoxia were associated with functional and morphological reorganization in the brain. However, the alterations of the interhemispheric functional connectivity in BA individuals remain unknown. The purpose of this study was to assess the interhemispheric functional connectivity changes in individuals with intermittent hypoxia due to BA using voxel-mirrored homotopic connectivity (VMHC) methods.

Methods: In total, 31 BA individuals (17 males and 14 females) and 30 healthy controls (HCs) (15 males and 15 females) closely matched in age, sex, and education underwent resting-state magnetic resonance imaging (MRI) scans. VMHC analysis was performed to investigate differences in interhemispheric functional connectivity between the two groups. Then, a seed-based resting-state functional connectivity (rsFC) analysis was conducted to further reveal the abnormal functional connectivity between the altered VMHC regions and the whole brain.

Results: Compared with HCs, BA individuals had significantly lower VMHC values in the bilateral basal ganglia/thalamus/insula, cuneus/calcarine/lingual gyrus [brodmann area (BA) 17/18/19], precentral and postcentral gyrus (BA 3/4/6). [voxel level $P < 0.01$, Gaussian random field (GRF) correction, cluster level $P < 0.05$]. Taking VMHC altered brain areas as seed points, the rsFC values of left insula/supramarginal/postcentral gyrus (PostCG) / inferior parietal lobule (IPL) brain areas in BA patients were increased, and rsFC values of right basic ganglia / thalamus, left caudate/antioxidant cingulate and bilateral cuneus/calcarine/lingual gyrus/precentral gyrus (PreCG)/PostCG were decreased in BA patients.

Conclusion: The abnormal resting-state functional connectivity of BA patients is altered in specific brain regions related to the basal ganglia network, visual network, and sensorimotor network, which may be related to the neuropathogenesis of asthma patients. Furthermore, these VMHC and FC values may be important clinical indicators for the diagnosis and treatment of asthma patients.

Introduction

Bronchial asthma (BA) is a severe respiratory disease characterized by chronic intermittent dyspnea. Asthma is a global health problem, nearly 300 million people suffer from asthma globally [1] According to a recent survey in China, the prevalence of asthma is 1.24%[2]. There are several risk factors for asthma such as obesity, respiratory infections [3], genetic [4], and environmental factors et al [5]. Chronic inflammation of the airways is the main pathological mechanism of asthma, subsequently results in narrowing of the airway and the classic symptoms of wheezing. The BA led to the impairment of lung function [6] and intermittent hypoxia. However, the BA patients are not only associated with respiratory symptoms, but also suffer from mental problems. The BA patients had a higher risk of developing mood and anxiety disorders relative to healthy individuals [7]. BA patients have also been shown to face higher risks of occurrence of depression and anxiety than healthy individuals [8, 9]. Moreover, BA patients suffer

from higher rates of cognitive impairment relative to healthy individuals [10]. Thus, the BA might lead to the dysfunction of the central nervous system.

To date, accumulated neuroimaging researches demonstrated that BA patients lead to significant brain functional and structural architecture changes. Xiong X et al reported that asthma without depression had decreased regional homogeneity (ReHo) in the right insula, whereas asthma with depression had decreased ReHo in the right insula [11]. Kline JN et al found that the insula and anterior cingulate cortex might play a critical role in inflammatory processes in BA patients [12, 13]. Meanwhile, the BA patients showed increased functional connectivity between the left ventral anterior insula and the right anterior cingulate cortex (ACC), decreased functional connectivity between the left ventral anterior insula and the bilateral parietal lobe relative to HCs [14]. Moreover, the asthma patients showed abnormal functional network centrality in the default mode network, the frontoparietal network, and the sensorimotor network [15]. In addition, BA patients lead to specific brain structural changes. Carlson SM et al revealed that smaller hippocampal volume was detected in asthma patients, which might have implications for impaired memory in BA patients [16]. Voxel-based morphometry (VBM) studies reported that gray matter volumes of the insular cortex and the brainstem periaqueductal grey were observed and it correlated with ratings of dyspnea unpleasantness in asthma patients [17]. Bian R et al reported that the asthma patients had abnormal white matter (WM) integrity in the left forceps major, cingulum, and right uncinate fasciculus, inferior longitudinal fasciculus relative to healthy individuals [18]. However, the abovementioned studies focused on the specific brain region functional and structural changes. The alternations of interhemispheric connectivity occurring in BA patients remain unknown. We hypothesized that BA patients might be associated with impaired interhemispheric connectivity changes implicated for impaired cognition and depression.

The human brain is two symmetric cerebral hemispheres with extensively anatomical and functional connectivity between homotopic locations. The synchrony of homotopic connectivity is an important feature of the functional structure of the brain. Interhemispheric coordination is involved in important physiological functions. Previous electroencephalogram (EEG) studies demonstrated that interhemispheric coupling was closely contributed to the processing of motor, auditory, and vision function [19–21]. Voxel-mirrored homotopic connectivity (VMHC) is an effective Resting-state-fMRI approach that can be used to investigate the homotopic connectivity between each hemisphere [22]. VMHC method has been shown the high test-retest reliability. The VMHC method had been successful to assess the neurophysiological mechanism of many diseases such as Obstructive Sleep Apnea-Hypopnea Syndrome [23] and chronic insomnia disorder [24]. However, the long-term effect of intermittent dyspnea due to BA on the homotopic connectivity is not well understood.

Here, the purpose of the study was to explore the interhemispheric connectivity changes in BA individuals. Furthermore, a seed-based resting-state functional connectivity (rsFC) analysis was applied to investigate the functional networks architecture seed as the altered VMHC regions in BA individuals. The relationship between the VMHC index and clinical variables was analyzed by Pearson correlation. These findings

might offer important insights into the understanding of abnormal hemispheric communication with BA individuals.

Materials And Methods

Participants: A total of 31 patients with BA (17 men and 14 women) were admitted to the Jiangxi Provincial People's Hospital (Nanchang, Jiangxi Province, China). The diagnostic criteria for bronchial asthma were as follows: (1) presence of intermittent hypoxia and wheezing symptoms; (2) more than 20% decrease in the rate of forced expiratory volume within 1 h; (3) absence of an upper respiratory tract infection; and (4) absence of psychiatric and cerebrovascular disorders.

30 HCs (15 men, 15 women) age, sex, and education status matched with patients in the BA group were enrolled. All healthy participants fulfilled the following criteria: (a) no psychiatric disorders (e.g. depression, bipolar disorder, sleep disorder, etc.); (b) without any contraindications of MRI. All research methods approved by the Ethics Committee of People's Hospital affiliated Nanchang University. Informed consent was obtained from all participants before MRI scanning, and the research methods complied with the principles of the Declaration of Helsinki.

MRI parameters

Our parameters were the same as our previous study [15]. MRI scanning was performed using a 3-Tesla MR scanner (Trio; Siemens, Munich, Germany). High-resolution T1-weighted anatomical images were acquired with a three-dimensional spoiled gradient-recalled sequence in a sagittal orientation: 176 images (repetition time= 1900ms, echo time= 2.26 ms, thickness= 1.0 mm, gap =0.5mm, acquisition matrix=256 × 256, the field of view =250mm × 250 mm, flip angle =9°) were obtained. Finally, a total of 240 functional images (repetition time =2000 ms, echo time=30 ms, thickness=4.0 mm, gap=1.2 mm, acquisition matrix=64×64, flip angle=90°, field of view=220 mm×220 mm, 29 axial slices with Gradient-Recalled Echo-Planar Imaging pulse sequence) covering the whole brain were obtained. All subjects underwent the MRI scanning with eyes closed without falling asleep.

fMRI data preprocessing

All preprocessing was performed using the toolbox for Data Processing & Analysis of Brain Imaging (DPABI, <http://www.rfmri.org/dpabi>) [25], which is based on Statistical Parametric Mapping (SPM8) (<http://www.fil.ion.ucl.ac.uk>) implemented in MATLAB 2013a (MathWorks, Natick, MA, USA) briefly the following steps: 1) DICOM format of the functional images converted to NIFTI format The first ten volumes of each subject were discarded due to the signal reaching equilibrium. 2) The remaining 230 volumes of functional BOLD images were corrected for slice timing effects, motion-corrected and realigned. Data from subjects whose head motion more than 2 mm or for whom rotation exceeded 2° during scanning were excluded according to the criteria of Van Dijk et al [26]. 3) Individual T1-weighted MPRAGE structural images were registered to the mean fMRI data, then resulting aligned T1-weighted images were segmented using the Diffeomorphic Anatomical Registration Through Exponentiated Lie

Algebra (DARTEL) toolbox for improving spatial precise in the normalization of fMRI data [27]. Normalized data (in Montreal Neurological Institute [MNI] 152 space) were re-sliced at a resolution of $3 \times 3 \times 3 \text{ mm}^3$. 4) smoothing with a 6 mm full-width-half-maximum Gaussian kernel; 5) linear regression analysis was used to regress out several covariate (six head motion parameters, mean framewise displacement (FD), global brain signal, and averaged signal from white matter signal and cerebrospinal fluid). 6) Data with linear trend were removed, and temporal band-pass was filtered (0.01–0.1 Hz).

Voxel-mirrored homotopic connectivity analysis

According to a previous study [22], the VMHC was calculated with REST software (<http://www.resting-fmri.sourceforge.net>) [28]. Briefly, the homotopic FC for each subject was computed as the Pearson correlation coefficient between each voxel's residual time series and that of its symmetrical interhemispheric counterpart. Correlation values were then Fisher z-transformed to improve normality.

Seed-based resting state functional connectivity (rsFC) analysis

To further detect the altered functional networks behind the impaired VMHC, the brain regions of decreased VMHC values were selected as the regions of interest (ROIs). Correlation analysis of the time course was performed between the spherical seed region and each voxel of the whole brain for each subject using REST software (<http://www.resting-fmri.Sourceforge.net>). Then, all functional connectivity maps were z-transformed with Fisher's r-to-z transformation to reduce the influence of individual variation for group statistical comparisons.

Statistical analysis

For demographic information

One-sample t-test was conducted to assess intra-group patterns of z-VMHC maps using SPM8 software. Two-sample t-tests including general linear model (GLM) for the groups of VMHC using SPM 8 (with age and gender treated as covariates; voxel-wise $P < 0.001$, Gaussian random field (GRF) theory connected cluster level, $P < 0.005$). Two-sample t-test was used to compare two group differences in the zFC maps including general linear model (GLM) for the groups of VMHC using SPM 8 (with age and gender treated as covariates; voxelwise $P < 0.001$, Gaussian random field (GRF) theory connected cluster level, $P < 0.005$).

Receiver operator characteristic (ROC) curve was used to assess the sensitivity of the z-VMHC values to distinguish the BA and HC groups. Pearson correlation was performed to investigate the relationship between the z-VMHC and zFC values and clinical variables in BA groups.

Results

Demographic and clinical characteristics of subjects

31 patients with BA (seventeen males and fourteen females, mean age: 51.62±5.20 years) and 30 HCs (seventeen males and fourteen females, mean age: 51.26±5.27 years) were included in this study, There were no significant differences in gender, age, weight and handedness between BA and HC groups ($P>0.05$). Details are presented in **Table 1**.

Table 1 Demographics and clinical measurements between BA and HC groups.

condition	BA	HC	t	P-value*
Male/female	17/14	15/15	N/A	>0.99
Age (years)	51.62±5.20	51.26±5.27	0.260	0.796
Weight (kg)	63.56±7.12	63.18±6.68	0.197	0.845
Handedness	31R	30R	N/A	>0.99
Duration of Asthma (years)	27±6.21	N/A	N/A	N/A

* $P < 0.05$ Independent t-tests comparing two groups, the data was showed in mean ±standard deviation.

Abbreviations: HC, healthy control; N/A, not applicable;

The difference of VMHC between BA group and HC group

The spatial pattern of VMHC at the average level of BA and HC groups. (**Figure 1**) The VMHC value of BA patients in bilateral ganglia/thalamus/ Insula, cuneus/ Calcarine/lingual gyrus, precentral gyrus (PreCG) and postcentral gyrus (PostCG) were lower than that of HCs. (voxel level $P < 0.01$, GRF-correction, cluster level $P < 0.05$) (**Table 2** and **Figure 2**)

Table 2. Differences in VMHC between two groups

Brain region	BA	Voxels size	Peak t-value	Peak coordinates in MNI space		
				X	Y	Z
BA <HC						
Basal ganglia/Thalamus/ Insula	-	557	-6.591	±42	-06	15
Cuneus/ Calcarine/ Lingual Gyrus	17,18,19	359	-6.423	±09	-93	0
PreCG /PostCG	3,4,6	211	-6.250	±66	0	9

Notes: t: statistical value of peak voxels showing significant differences between the two groups. X, Y, Z are the coordinates of primary peak locations in the MNI space. (voxel level $P < 0.01$, GRF-corrected for

multiple comparisons at a cluster level of $P < 0.05$).

Abbreviations: VMHC, voxel-mirrored homotopic connectivity; GRF, Gaussian random field; BA, bronchial asthma; HCs, healthy controls; MNI, Montreal Neurological Institute; BA's, Brodmann's area; PreCG, Precentral Gyrus; PostCG, Postcentral Gyrus;

Comparative analysis of the functional connectivity of VMHC altered brain regions in BA group and HC group

The altered VMHC of each brain region was used as the seed brain region, compared with the control group, the rsFC values of the left insula / supramarginal / PreCG / inferior parietal lobule (IPL) brain areas in BA group were significantly increased in those of control group, while the rsFC values in the right basal ganglia / thalamus, left caudate / anterior cingulate and bilateral cuneus / calcarine / lingual gyrus / PreCG / PostCG were significantly decreased in BA group. (voxel level $P < 0.01$, GRF-correction, cluster level $P < 0.05$) (**Table 3** and **Figure 3-5**)

Table 3. Differences in heterotopic connectivity of the regions with altered VMHC

Brain region	BA	Voxels size	Peak t-value	Peak coordinates in MNI space		
				X	Y	Z
ROI1: L-Basal ganglia/Thalamus/ Insula						
L- Insula/ SupraMarginal/ PreCG /IPL	13,6,40	1082	5.923	-42	-39	24
R- Basal ganglia/Thalamus	-	356	-6.269	-27	-30	3
ROI2: R-Basal ganglia/Thalamus/ Insula						
R- Basal ganglia/Thalamus	-	271	-5.474	-24	-30	6
L- Insula/ SupraMarginal/ PreCG /IPL	-	293	4.801	-27	24	9
L-Caudate/Anterior Cingulate	33	229	-6.059	-12	12	27
ROI3: L- Cuneus/ Calcarine/ Lingual Gyrus						
B-Cuneus/ Calcarine/ Lingual Gyrus	17,18,19	1506	-6.644	3	-69	6
ROI4: R-Cuneus/ Calcarine/ Lingual Gyrus						
B- Cuneus/ Calcarine/ Lingual Gyrus	17,18,19	1863	-6.628	15	-93	-9
ROI5: L-PreCG/PostCG						
R- PreCG/PostCG	3,4,6	490	-6.136	39	-18	33
L-PreCG/PostCG	3,4,6	275	-5.486	-42	-21	30
ROI6: R- PreCG/PostCG						
R- PreCG/PostCG	3,4,6	358	-6.104	42	-18	33
L-PreCG/PostCG	3,4,6	105	-5.114	-42	-21	30

Notes: t: statistical value of peak voxels showing significant differences between the two groups (negative values: BA<HC; positive values: BA>HC). X, Y, Z are the coordinates of primary peak locations in the MNI space. (voxel level P<0.01, GRF-corrected for multiple comparisons at a cluster level of P < 0.05).

Abbreviations: VMHC, voxel-mirrored homotopic connectivity; GRF, Gaussian random field; BA's, Brodmann's area; MNI, Montreal Neurological Institute Coordinate System; BA, bronchial asthma; HCs, healthy controls; MNI, Montreal Neurological Institute; BA's, brodmann's area; IPL, Inferior Parietal Lobule; PreCG, Precentral Gyrus; PostCG, Postcentral Gyrus; L, left; R, right; B, bilateral;

Receiver operating characteristic (ROC) curve analysis

The receiver operating Characteristic curve (ROC) tested the different combinations of abnormal mean VMHC values of 8 bilateral brain regions in the two groups, and we used ROC curve to analyze the diagnostic value of different brain regions. The areas under the curve (AUC) were as follows: bilateral-basal ganglia(BG)/thalamus (Tha)/insula (Ins) was 0.943; bilateral-cuneus (Cun)/ calcarine (Cal)/lingual Gyrus(LG) was 0.852; bilateral- precentral Gyrus (M1)/ postcentral Gyrus(S1) was 0.867, B-BG/Tha/Ins showed the highest AUG, which may mean that these brain regions are most important for the diagnosis of BA (Figure 6).

Correlation analysis

At the level of GRF correction $P < 0.05$, VMHC anomalies of BA patients were not correlated with clinical variables.

Discussion

In this study, rsMRI technology was used to detect BA patients and HC controls, and the VMHC method was used to evaluate the changes in the functional connections between cerebral hemispheres. We have two major findings, first, it was mainly found that the VMHC values of these brain areas (bilateral basal ganglia/thalamus/ insula, cuneus/ calcarine/ lingual gyrus, PreCG, and PostCG) in BA patients were significantly lower than those in the normal controls. Second, functional connectivity was studied using VMHC abnormal brain regions as seed points, compared with the HC group. As far as we know, this is the first report that VMHC combined with seed-based rsFC method which has been used to study the changes of functional connectivity in the whole brain of BA patients. We believe that these findings will contribute to the development of an imaging biomarker for the diagnosis of BA, which is of great significance in improving the clinical efficacy and quality of life of patients with asthma.

Functional homotopy refers to the high similarity in endogenous spontaneous activities of neurons of the same origin in the left and right hemispheres of the brain, and it is related to brain stratification [22]. The VMHC method measures interhemispheric coordination and is a rsMRI method used to study the functional connections between hemispheres, also, it is an important method to study brain information integration by measuring the correlation between cerebral hemisphere blood oxygen level-dependent time series and reflecting the information exchange and integration mode between cerebral hemispheres [29]. At present, this method has been widely used to assess the functional connectivity of the cerebral hemisphere in various neuropsychiatric diseases. Meanwhile, asthma is considered to be a respiratory disease that seriously affects the central nervous system [30]. Combined with our study, the VMHC values of eight brain regions, namely bilateral basal ganglia/thalamus/ insula, cuneus/calcarine/lingual gyrus, PreCG, and PostCG decreased in BA patients, suggesting that the functional connectivity in these brain regions is impaired in BA patients.

Both thalamus and basal ganglia are important to brain regions for the basal ganglia network [31], Xiaodan Yan et al. found that BOLD signal activation in the middle gyrus, occipital gyrus, lingual gyrus, and thalamus was reduced in a high-altitude group compared with those in normal altitude group,

moreover, these high altitudes subjects had longer response time and lower response accuracy, they believe that this may be caused by the long-term hypoxia concentration and low pressure of high altitude people, and the function and development of the brain in the hypoxia environment will be affected, which will lead to the decline in cognitive ability [32]. In this instance, we can explain the abnormality of the basal ganglia network of asthmatic patients, because asthmatics are characterized by blocked airflow and bronchospasm, and severe asthma attacks can lead to hypoxia [33]. Besides, many studies have shown that the insula is an important brain region involved in the neuropathology of asthma, for example, according to the research of Rosenkranz et al, patients with stress-related diseases such as asthma, regions of the brain such as the insula may lead to inflammatory responses by affecting disease-specific emotions and incoming physiological signals [13]. Also, Yuqun Zhang et al found that the FC value of the insular brain area of asthmatic patients decreased compared with that of normal patients, but after group cognitive behavior therapy treatment, the FC value of the insular subregion increased significantly [34]. Additionally, the insula receives afferent information about the cord cortex of the thalamus, carries information related to respiration, and is closely related to the nerve center that processes emotions [35]. Moreover, the insula is involved in the effective assessment of sensory stimuli, regulation of homeostasis responses, and visceral perception, moreover, the degree of insula activation can predict individual differences in the evaluation of stimulus intensity and visceral consciousness. All of these studies indicate that emotional stimulation and insular area are closely related to asthma. In brief, we found a decrease in VMHC in basal ganglia, thalamus and, the insular lobes, which may be used as a clinical indicator for evaluating BA patients.

Many pieces of evidence showed that asthma was a serious respiratory disease that can affect the nervous system [36, 7], furthermore, there has been functional MRI study with degree centrality (DC) that have confirmed that the sensorimotor network and visual network of asthmatic patients are abnormal [37]. However, relevant studies on VMHC methods are still vacant. Brodmann Area (BA) 17/18/19 is the visual cortex, these brain regions as the main area of the visual network, its VMHC value is altered in asthmatic patients, the abnormal change in the visual brain network may imply that asthma can affect the spontaneous changes of the central nervous system. These important brain regions of the visual network play an important role in the body's visual function. Calcarine (BA19) as the visual contact area, which is responsible for visual processing, and forms visual joint cortex with BA18, and an rs-fMRI study on asthma confirmed that the FC value of BA19 on the left side of asthma patients was significantly lower than that of the normal control group [34]. Besides, cuneus (BA17) mainly belongs to the primary visual center(V1), as the core of visual pathways [38], it is responsible for the preliminary processing of visual information, receiving information about the lateral geniculate body, and transmitting the information on other visual areas [39], and when the body is not paying attention, visual information will be difficult to transmit from V1 to other regions [40]. Also, the lingual gyrus (BA18) from the secondary visual cortex (V2) is the visual contact area, which is mainly responsible for visual processing. Moreover, it receives the pre feedback connection between V1 and plays an important role in object shape vision and stereo vision [41, 42]. Thus, we demonstrated that the neural activity of the visual network between

the two hemispheres of asthmatic patients changed synchronously, moreover, this change of visual brain network in BA patients may lead to abnormal visual function.

PreCG (BA4/6, M1) and PostCG (BA3, S1) belong to the motor and sensory centers respectively, they are also sensorimotor areas [43]. In addition, the PostCG is the core component of the sensorimotor network [44], the motor center mainly controls the movement behavior of the body, while the sensory center has the basic somatosensory function, which is responsible for encoding touch and pain [45], also, receiving the sensory input from the limbs. Furthermore, Liu et al found that the sensorimotor network of asthmatic patients would change, and this change was related to the altered respiratory amplitude, in addition, Li used MRI to contrast the brain structure of asthma patients and normal controls, found that patients with asthma sensorimotor network is abnormal, they through the network abnormal airway obstruction and the degree of correlation analysis reflect the brain damage associated with asthma caused by respiratory damage, they believe that brain network potential parameters can be used as an understanding of the pathophysiology of asthma [37]. Also, Zhang et al found the M1/S1 brain area abnormality in asthmatic patients, and the FC value of these two brain regions was lower in asthmatic patients than that of the normal control group [34]. Therefore, we hypothesized that sensorimotor function might change in BA patients, and this change was associated with abnormalities in key brain areas of the sensorimotor network.

Our further seed-based rsFC analyses revealed that FC values of these brain regions (B- PreCG/PostCG, B- Cuneus/ Calcarine/ Lingual Gyrus, and R- Basal ganglia/Thalamus) were lower in BA patients, furthermore, the VMHC values corresponding to these brain regions in BA patients are also lower than those in normal subjects, which suggests that there are abnormal changes in resting-state functional connectivity between these brain regions in asthmatic patients and corresponding ataxia between cerebral hemispheres. Moreover, the inferior parietal lobule (IPL) is involved in the control of body perception, motor orientation, memory retrieval, language understanding, digital processing, and social cognition, this brain area is one of the main brain areas of the frontoparietal network, which plays an important role in attention and executive ability [46–48]. Kuo et al found that the FC value of the IPL brain region in untreated heroin addicts was significantly lower than that in normal patients, and they speculated that the IPL brain region might be a neural target for treatment and intervention [49]. Besides, an MRI studied on asthma found that the degree centrality value of asthmatic patients in the IPL brain area was lower than that of normal controls, this study suggested that the IPL brain region can be used as one of the clinical indicators for the diagnosis of asthma [15]. The supramarginal gyrus is involved in verbal divergent thinking, and this brain region is activated during divergent thinking tasks [50]. The cingulate gyrus belongs to the limbic system, which is mainly involved in emotion formation, processing, learning, and memory [51], the interaction between the cingulate cortex and other brain networks may be very important to the body's consciousness [52]. Furthermore, the caudate nucleus plays an important role in cognitive function [53], Zhang et al not only found the abnormality of FC value in asthma patients, but also found the change of FC values in caudate and cingulate brain regions [34], which were similar to our findings in BA patients. Hence, combined with our rsFC results, we speculate that asthma may involve changes in functional connectivity of multiple brain regions, which are closely related to the body's

emotional, motor, cognitive and other functions, and this may mean that asthma patients are dysfunctional in the above abilities.

Conclusion

In conclusion, VMHC and rsFC based on seed points were used to compare the resting-state functional connectivity between BA patients and healthy controls. We found that the functional connectivity between the hemispheres of bilateral basal ganglia, thalamus, insula, cuneus, calcarine, lingual gyrus, precentral and postcentral gyrus was abnormal in asthmatic patients, these results may help to understand the intrinsic neural mechanism of BA patients, and provide some new reference to the formulation of treatment plans for asthma patients.

Declarations

Data availability

MRI data used to support the results of this study are available on request from the corresponding authors.

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Author contributions

Ya-Jun Wu: responsible for writing manuscript; Jie Rao: responsible for proofreading and revising the language of the manuscript; Xiao-Rong Wu and Jun Wang: guide the writing and revision of the

manuscript; Xin Huang was responsible for the production of all images and tables in the manuscript; Na Wu, Ling Shi, Hui Huang, Si-Yu Li, Xiao-Lin Chen, Shui-Qin Huang, Pei-Pei Zhong: responsible for clinical trials and data processing.

Conflict of Interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Figures

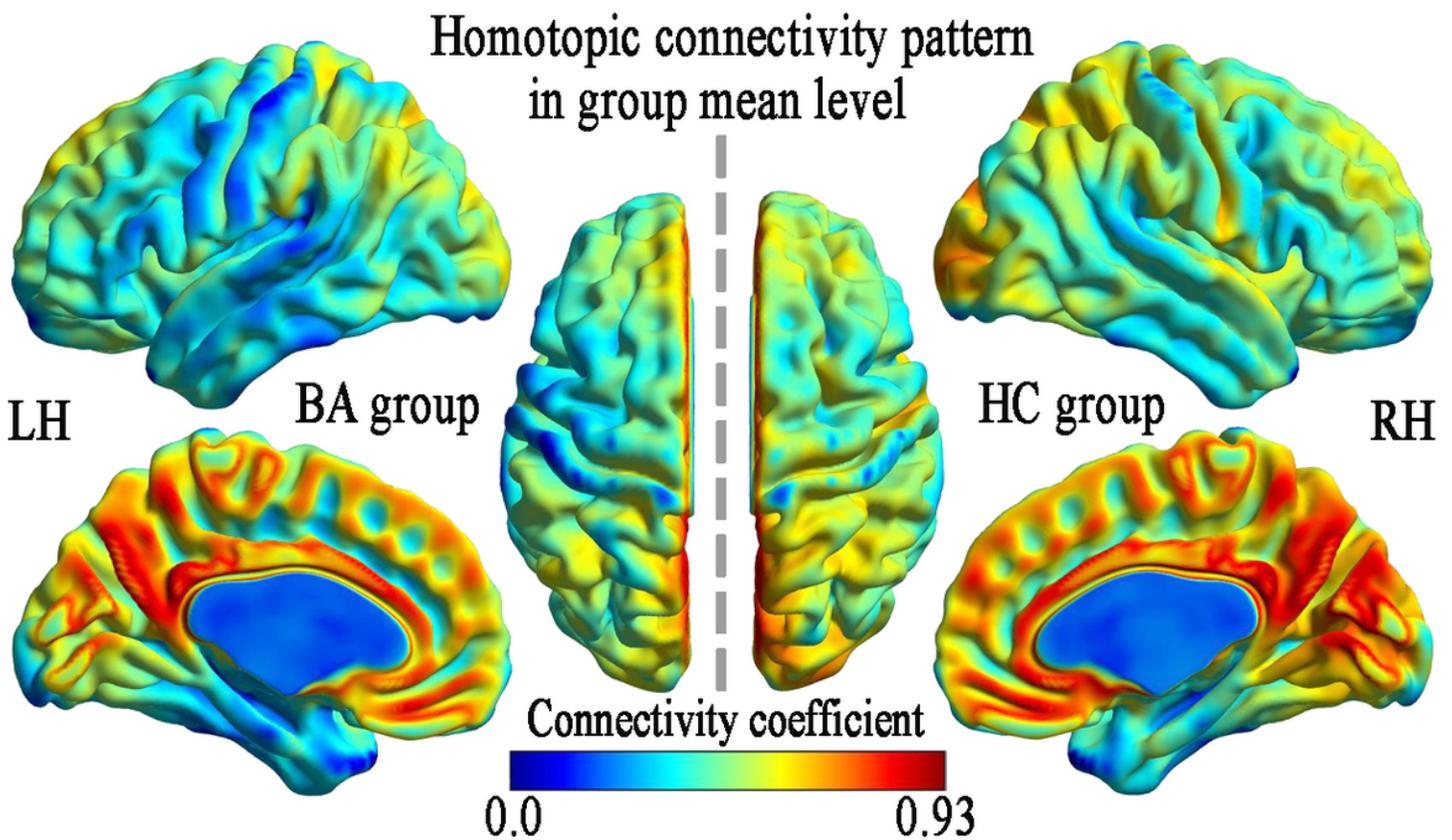


Figure 1

VMHC spatial patterns at the group mean level of the BA and HC groups. Note: Within-group mean VMHC maps within the BA (left) and HCs (right). Abbreviations: VMHC, voxel-mirrored homotopic connectivity; BA, bronchial asthma; HC, health control; LH, left hemisphere; RH, right hemisphere;

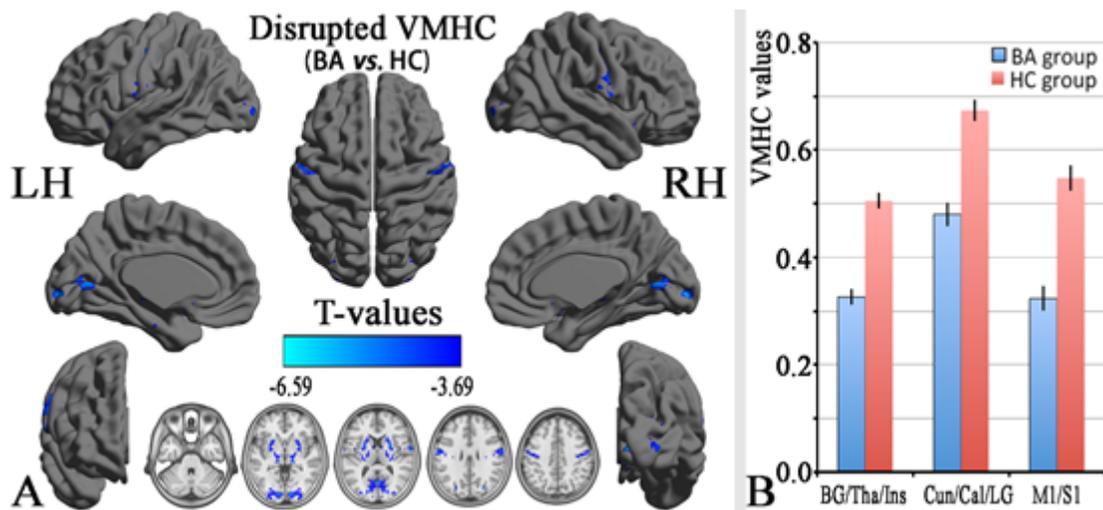


Figure 2

Group comparisons of the VMHC between the BA and HCs. Note: Significant VMHC differences were found between two groups. (A) The blue areas indicate lower VMHC values, respectively (two-tailed, voxel-level $P < 0.01$, GRF correction, cluster-level $P < 0.05$). The mean values of altered VMHC values between the BA and HC groups was shown in histogram. (B) Abbreviations: VMHC, voxel-mirrored homotopic connectivity; BA, bronchial asthma; HC, health control; LH, left hemisphere; RH, right hemisphere; BG, Basal ganglia; Tha, Thalamus; Ins, Insula; Cun, Cuneus; Cal, Calcarine; LG, Lingual Gyrus; M1, Precentral Gyrus; S1, Postcentral Gyrus;

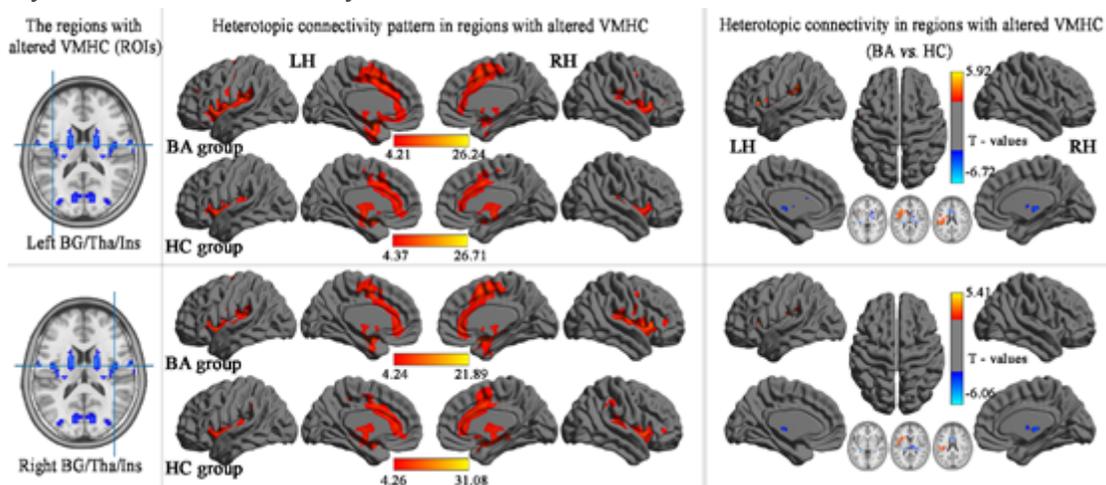


Figure 3

note: Group comparisons seed-based FC of the altered VMHC between the BA and HC seeded regions in bilateral BG/Tha/Ins. ($P < 0.01$, GRF-corrected at a cluster level of $P < 0.05$) Abbreviations: VMHC, voxel-mirrored homotopic connectivity; GRF, Gaussian random field; BA's, Brodmann's area; BA, bronchial asthma; HCs, healthy controls; FC, functional connectivity; ROI, region of interest; BG, Basal ganglia; Tha, Thalamus; Ins, Insula;

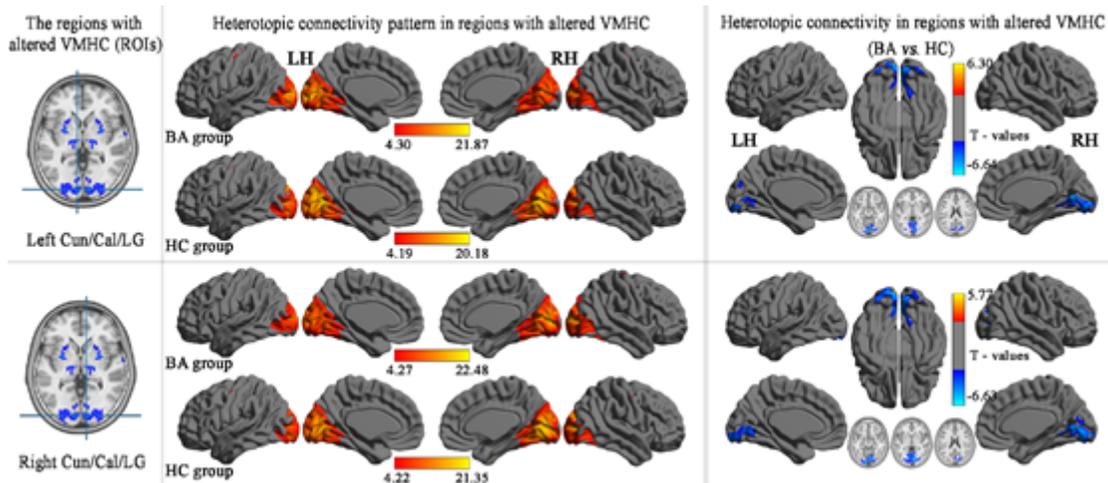


Figure 4

note: Group comparisons seed-based FC of the altered VMHC between the BA and HC seed regions in bilateral Cun/Cal/LG . ($P < 0.01$, GRF-corrected at a cluster level of $P < 0.05$) Abbreviations: VMHC, voxel-mirrored homotopic connectivity; GRF, Gaussian random field; BA's , Brodmann's area; BA, bronchial asthma; HCs, healthy controls; FC, functional connectivity; ROI, region of interest; Cun, Cuneus; Cal, Calcarine; LG, Lingual Gyrus;

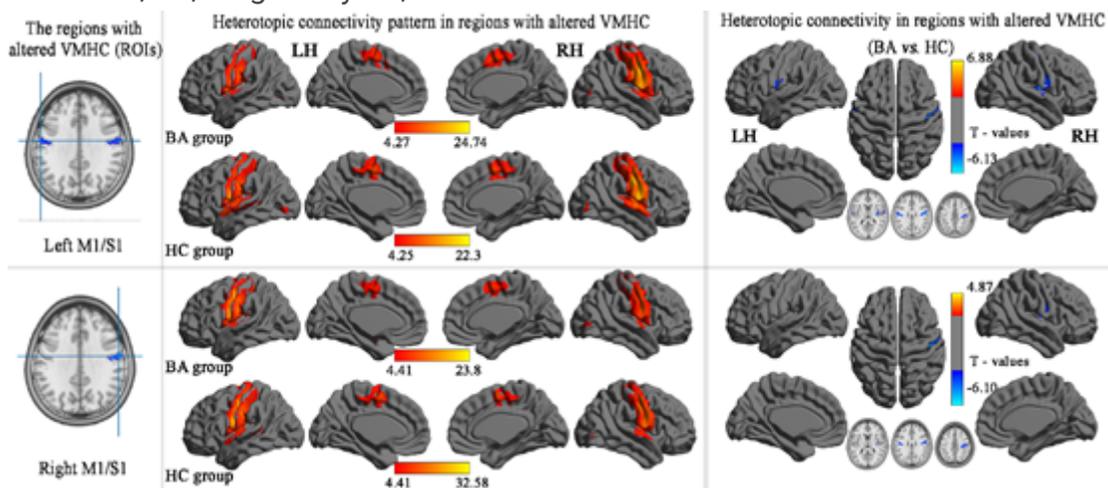


Figure 5

note: Group comparisons seed-based FC of the altered VMHC between the BA and HC seed regions in bilateral M1/S1. ($P < 0.01$, GRF-corrected at a cluster level of $P < 0.05$) Abbreviations: VMHC, voxel-mirrored homotopic connectivity; GRF, Gaussian random field; BA's, Brodmann's area; BA, bronchial asthma; HCs, healthy controls; FC, functional connectivity; ROI, region of interest; M1, Precentral Gyrus; S1, Postcentral Gyrus;

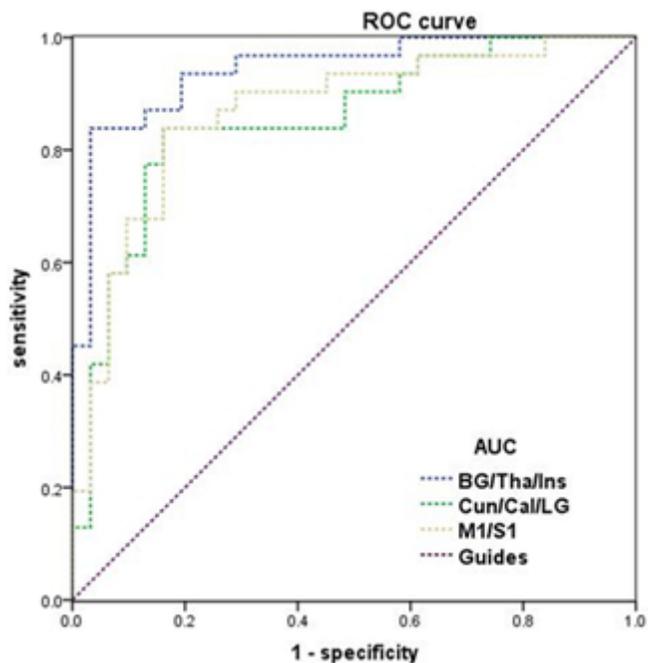


Figure 6

High sensitivity in mean VMHC values of different brain regions within two groups. Note: ROC curve in VMHC: BA < HC, for B-BG/Tha/Ins 0.943 ($P < 0.001$; 95% CI: 0.887–0.999); BA < HC, for B-Cun/Cal/LG 0.852 ($P < 0.001$; 95% CI: 0.755–0.950); BA < HC, for B-M1/S1 0.867 ($P < 0.001$; 95% CI: 0.774–0.960); Abbreviations: ROC, receiver operating characteristic; AUC, area under the curve; VMHC, voxel-mirrored homotopic connectivity; BA, bronchial asthma; HC, health control; BG, Basal ganglia; Tha, Thalamus; Ins, Insula; Cun, Cuneus; Cal, Calcarine; LG, Lingual Gyrus; M1, Precentral Gyrus; S1, Postcentral Gyrus; B, bilateral;