



Research article

Aberrant intrinsic functional brain topology in methamphetamine-dependent individuals after six-months of abstinence

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Abstract: Our aim was to explore the aberrant intrinsic functional topology in methamphetamine-dependent individuals after six months of abstinence using resting-state functional magnetic imaging (rs-fMRI). Eleven methamphetamines (MA) abstainers who have abstained for six months and eleven healthy controls (HC) were recruited for rs-fMRI examination. The graph theory and functional connectivity (FC) analysis were employed to investigate the aberrant intrinsic functional brain topology between the two groups at multiple levels. Compared with the HC group, the characteristic shortest path length (L_p) showed a significant decrease at the global level, while the global efficiency (E_{glob}) and local efficiency (E_{loc}) showed an increase considerably. After FDR correction, we found significant group differences in nodal degree and nodal efficiency at the regional level in the ventral attentional network (VAN), dorsal attentional network (DAN), somatosensory network (SMN), visual network (VN) and default mode network (DMN). In addition, the NBS method presented the aberrations in edge-based FC, including frontoparietal network (FPN), subcortical network (SCN), VAN, DAN, SMN, VN and DMN. Moreover, the FC of large-scale functional brain networks revealed a decrease within the VN and SCN and between the networks. These findings suggest that some functions, e.g., visual processing skills, object recognition and memory, may not fully recover after six months of withdrawal. This leads to the possibility of relapse behavior when confronted with MA-related cues, which may contribute to explaining the relapse mechanism. We also provide an imaging

basis for revealing the neural mechanism of MA-dependency after six months of abstinence.

Keywords: methamphetamine; abstainers; graph theory; functional connectivity; mechanism

1. Introduction

Methamphetamine (MA) is a new synthetic compound with strong central euphoric effects [1]. The prevalence of MA abuse is continuously increasing and ranking as the second most widely used illicit drug globally due to its ease of production, which has become a global public health problem [2]. As an amphetamine-type stimulant, MA abuse can cause various physical illnesses and psychiatric deficits [3]. Moreover, even after undergoing treatment for substance abuse, patients often relapse when they encounter high-risk environments that may trigger drug abuse, severely impacting families and society [4]. However, the exact relapse mechanisms of MA addiction have not been fully explored [5]. The main reason is that we have not clarified the altered functional brain imaging mechanism in MA withdrawal patients. Therefore, how to explore alterations in functional brain imaging and, thus, potential mechanisms of relapse in MA abstainers has become a hot topic.

Magnetic resonance imaging (MRI) is essential for clinical and brain science research [6,7]. Numerous studies have shown that long-term abuse causes irreversible damage to brain function and induces psychiatric symptoms or cognitive deficits [8–10]. After a period of withdrawal, the structure and function of the brain can be improved or restored to a certain extent [11,12]. For example, Zhang et al. [13] focus on the brain structure change in long-term abstinence of MA by voxel-based morphometry. They found differences in grey matter volumes in the cerebellum and right thalamus compared with normal subjects. In addition, they suggested that prolonged abstinence benefits cognitive function recovery. By the seed-based functional connectivity (FC) analysis, Li et al. [14] found disrupted FC between the cerebellum and several cerebral functional networks in addicts after six months of abstinence. Based on diffusion-tensor imaging, Fan et al. [15] revealed abnormal microstructures of several brain regions in MA individuals following prolonged withdrawal by voxel-based analysis.

Among various methods to explore brain changes, graph theory is an efficient tool for whole-brain analysis to detect functional differences [16,17]. This method not only measures the connectivity between brain regions, but also allows the detection and quantification of changes in the characteristic properties of each brain region to explore microscopic changes in the central system [18]. Hence, it is widely used in research to explore MA addiction or relapse mechanisms [19,20]. In a study of diffusion-tensor imaging, Zhou et al. [21] found that the brains of male addicts have small-world networks, with significantly efficient transmission and integration of information between brain regions. Mansoori et al. [22] adopted resting-state fMRI and graph theory to investigate the FC and local efficiency in the left and right hemispheres of the brain. They demonstrated the alterations in the default mode network (DMN), executive control network (ECN) and salience network (SN) in both hemispheres of the brain. Generally, these studies have shown structural or functional abnormalities in the brain of addicts in withdrawal at the topological level. This allows for a better understanding of abnormalities in information interaction between brain regions and coordination and cooperation between brain networks [23]. However, most relevant studies focus on an extended range of withdrawal, i.e., they did not focus on investigating a particular point in time. Studying brain

abnormalities at specific withdrawal time points, such as the six-month voluntary MA withdrawal period in China, may better explain the mechanisms of relapse [14].

To the best of our knowledge, few functional brain imaging studies have explored functional alterations after six months of abstinence at the topological level. Consider that the brain changes during short-term withdrawal, i.e., less than six months, may reflect a compensatory response to MA rather than an actual recovery pathway. In the current study, we employed graph theory and FC analysis to explore aberrant intrinsic functional topology in MA-dependent individuals after six months of abstinence. We hypothesize that different brain areas remain dysfunctional, possibly leading to dysregulation of executive control, emotional processing and sensorimotor functions in MA-dependent individuals after six months of abstinence, which may provide imaging evidence for abnormal brain topology and a better understanding of the underlying relapse mechanisms.

2. Materials and methods

To explore the aberrant intrinsic functional topology in MA-dependent individuals after six months of abstinence, we first acquired fMRI data from the subjects and preprocessed it. Then, we use the atlas to construct and analyze the individual brain networks according to three dimensions: Nodes, edges and large-scale brain networks. Finally, a two-sample t-test was performed on the different metrics of the two groups, corrected for multiple comparison methods, such as false discovery rate (FDR) or network-based statistic (NBS), to find the significant differences in brain regions. The flowchart of the study is shown in Figure 1.

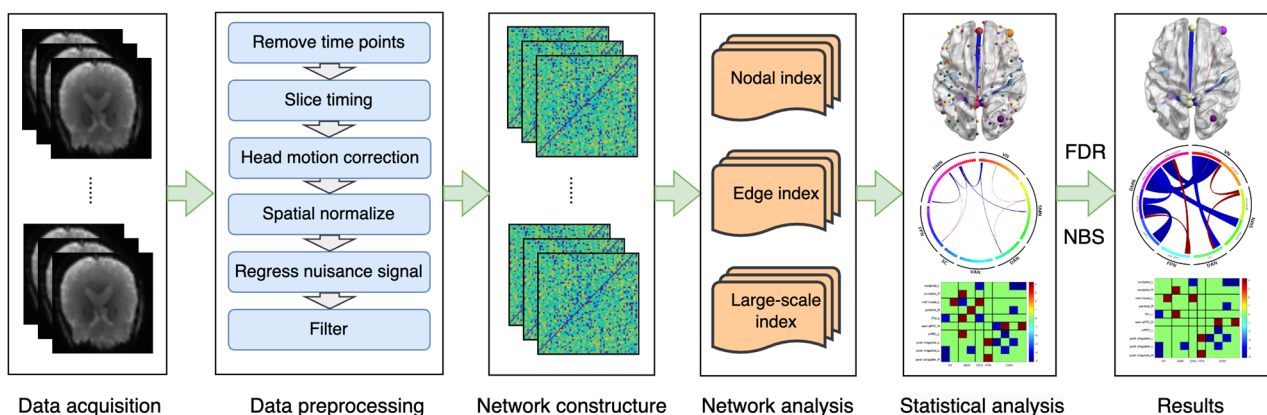


Figure 1. Flowchart of estimating aberrant intrinsic functional topology in the subjects.

2.1. Participants

We recruited 11 male MA abstainers and 11 age- and sex-matched HCs for the present study. The MA abstainers were recruited from the Shandong Detoxification Monitoring and Treatment Institute, while the HCs were recruited from the community. MA abstainers were diagnosed according to the Diagnostic and Statistical Manual on Mental Disorders (DSM-V) criteria and after that had received a six-month voluntary abstinence period. During withdrawal, addicts were given medication and physical exercises and showed no significant withdrawal symptoms by the end of the period.

The inclusion criteria included males, right-handed, ranging in age from 20 to 30 years old,

without a history of psychiatric or neurological disorders, not taking medication two weeks before data collection that could affect the experiment results and informed consent. The exclusion criteria included subjects with severe aphasia, cognitive loss or inability to communicate, a history of alcohol or drug abuse, previous or existing severe liver or kidney damage or cardiovascular disease or organic mental disorders, contraindications to MRI scannings, such as the presence of metal objects, pacemakers, metal dentures and claustrophobia. In addition, the abstainers abuse only one MA drug and escape the physiological detoxification period with a negative MA urine test after six months of abstinence.

This study was approved by the Ethics Committee of the Affiliated Hospital of Shandong University of Traditional Chinese Medicine (Ethics No. 2018-078). All volunteers were informed of the experimental procedure and wrote the informed consent before scanning.

2.2. Data acquisition and preprocessing

Brain imaging was performed on a Philips Ingenia 3.0T magnetic resonance scanner. The fMRI data were acquired using the T2 echo planar imaging sequence with the following parameters: TR = 2000 ms, TE = 30 ms, rotation angle = 90° , matrix = 64×64 , FOV = $220 \text{ mm} \times 220 \text{ mm}$, slice thickness = 3 mm, thickness interval = 1 mm. After resting-state scanning, structural images of each subject were also collected for subsequent spatial alignment using the fast acquisition gradient echo sequence with the following parameters: Rotation angle = 9° , TR = 2300 ms, TE = 3.01 ms, matrix = 256×256 , slice thickness = 1 mm. Foam padding and earplugs were used to reduce head motion and scanner noise. The scans were reviewed by two experienced neuroradiologists and showed no serious neuroanatomical abnormalities or image distortions associated with head movements in any of the participants.

The fMRI data were automatically preprocessed using the Data Processing and Analysis for Brain Imaging (DPABI) toolbox based on MATLAB with the standard process [24]. First, the first 10 scan time points data were removed to ensure scan signal stability. The remaining images were corrected by the slice-timing and head motion. They were then segmented and normalized to the standard Montreal Neurological Institute (MNI) space using the DARTEL tool. After alignment, the image is smoothed using a full-width half-height kernel to increase the signal-to-noise ratio. Then, the nuisance signals were regressed, including white matter signals and cerebrospinal fluid signals, linear trends and 24 Friston head movement parameters. The functional images were bandpass filtered (0.01–0.1 Hz).

The exclusion criteria were subjects with maximum head movement amplitude larger than 3 mm or rotation larger than 3° in the X, Y and Z axes. Furthermore, subjects with mean frame-wise displacement (mean FD) larger than 0.2 mm were also excluded from further analyses ($P > 0.05$) [25].

2.3. Network construction

We construct functional brain networks from node and edge perspectives via the Data Processing and Analysis for Brain Imaging Network analysis (DPABINet) toolbox, which has evolved from DPABI [26]. The functional brain networks of the node represent the different brain regions, while the edge represents the FC among the various brain regions. The Dosenbach atlas [27], which defines 160 ROIs across the whole brain, was used to segment the brain region. Since all scans did not fully cover the cerebellum, we excluded them from the analysis. After removing 18 brain regions from the

cerebellum, 142 ROIs were retained. For each ROI, the mean BOLD signal intensity of the ROI was extracted, and the Pearson correlation coefficient of the time series between any ROIs was calculated to construct a whole-brain FC matrix, which has a dimension of 142×142 and contains a total of 10,011 edges. Then, we converted the FC matrix to z-scores using Fisher's r-to-z, making the matrix more normal.

2.4. Network analysis

Global and nodal topological matrices were analyzed using the graph theory analysis module of the DPABINet toolbox. Specifically, global network matrices include clustering coefficient (C_p), characteristic shortest path length (L_p), small-world network (α), global efficiency (E_{glob}) and local efficiency (E_{loc}). The nodal network matrices mostly include the nodal degree and nodal efficiency. In the constructed functional brain networks, the strength of FC between different brain regions is defined as the edge. Then, the ratio of edges present in the network to the maximum number of possible edges is defined as the sparsity of the network [18]. For each metric, we analyzed the global and nodal topological matrices and calculated the area under the curve (AUC) at a sparsity of 0.1–0.34 with an interval of 0.01 to avoid the results being dependent on a particular sparsity value. The calculation formulas of network matrices are presented as follows [28]:

$$C_p = \frac{2C}{N(N-1)} \quad (1)$$

$$L_p = \frac{1}{N(N-1)} \sum_{i,j,i \neq j} D(i,j) \quad (2)$$

$$\alpha = \frac{C_p/C_{prand}}{L_p/L_{prand}} \quad (3)$$

$$E_{glob} = \frac{1}{N(N-1)} \sum_{i,j,i \neq j} \frac{1}{D(i,j)} \quad (4)$$

where N represents the number of nodes in the network and C denotes the number of statistically significant connected edges in the network after multiple comparisons. $D(i,j)$ represents the number of minimum steps required to connect the nodes i and j . C_{prand} and L_{prand} represent the clustering coefficient and characteristic shortest path length of the randomized network, respectively.

2.5. Statistical analysis

Demographic information was analyzed using SPSS statistical software (Version 26, IBM). Specifically, we adopted the two-sample t-test to compare the differences in clinical data between the two groups. In addition, we also conducted the two-sample t-tests to examine the differences in AUC of each network metric in global and nodal levels between the two groups. The results were corrected using the FDR method for multiple comparisons ($P < 0.05$).

Further, two-sample t-tests with the NBS method were used to assess the detailed differences in edge-based FC strength between the two groups. The primary threshold for each edge is set at $P < 0.001$ in the t-tests. The functional brain network is defined by the interconnectivity of suprathreshold edges in the topological space. The NBS method performs the statistical analysis on each edge to obtain the

statistic t-value, detects the network composed of suprathreshold edges based on the t-value and then calculates the number of edges the network contains. A permutation test with 5000 iterations was employed to generate the distributions of suprathreshold edge numbers in the network for non-parametric testing.

To better describe the significant node and edge obtained in the multiple comparisons, we also classified and reported the results according to their affiliation with the network defined by Yeo et al. [29]. Since the limbic network contains few ROIs, we used the subcortical network instead. The seven networks were the visual network (VN), the somatosensory network (SMN), the dorsal attentional network (DAN), the ventral attentional network (VAN), the subcortical network (SCN), the frontoparietal network (FPN) and default mode network (DMN).

Finally, we have investigated the FC conditions between large-scale intra- and inter-network. The results were calculated by averaging the z-score of FC across all involved edges and yielded seven intra-network and twenty-two inter-network values. Then, the two-sample t-tests were adopted to investigate the effect of long-term MA administration on FC changes by comparing the FC of large-scale networks in MA abstainers with those in the healthy group. The FDR was used to correct the results. We counted the number of edges that fell into the seven intra-network and inter-network classes.

3. Results

3.1. Demographic and clinical information

The demographic and clinical information of the two groups is shown in Table 1. A total of twenty-two subjects were included in this study. The mean duration and amount of drug use in the MA group are 48.95 ± 13.32 months and 0.36 ± 0.31 g, respectively. In addition, the two groups had no significant difference in age and education level ($P > 0.05$).

Table 1. Demographic and clinical information statistics for MA and HC groups.

Characteristics	MA Mean \pm SD	HC Mean \pm SD	P-value
Age (years)	29.43 ± 4.25	30.81 ± 3.22	0.375
Education level (years)	12.74 ± 2.41	13.35 ± 2.18	0.576
Duration of drug use (months)	48.95 ± 13.32	-	-
Amount of drug use (g/d)	0.36 ± 0.31	-	-

Note: MA: methamphetamine, HC: healthy controls, SD: standard deviation, -: no data.

3.2. Global topological networks

In the sparsity range of 0.1 to 0.34, we found significantly lower L_p and higher E_{glob} in the MA group, with statistically differences between the two groups ($P < 0.05$). Nevertheless, E_{loc} was only statistically different between the two groups within a sparsity range of 0.1 to 0.25 ($P < 0.05$). In addition, although the subjects in the two groups met the small-world networks, no statistical differences existed between groups for C_p and α ($P > 0.05$). The results are shown in Figures 2 and 3.

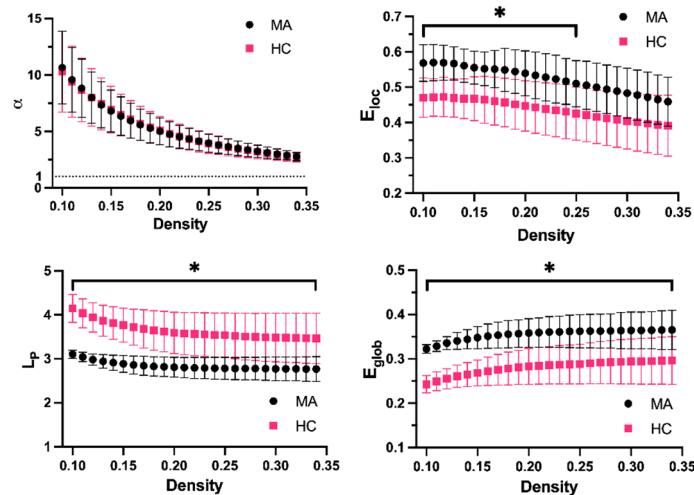


Figure 2. Differences global topological metrics of brain functional networks between two groups at the range of sparsity. *: $P < 0.05$, α : Small-world index, E_{loc} : Local efficiency, L_p : Characteristic shortest path length, E_{glob} : Global efficiency, MA: Methamphetamine and HC: Healthy controls.

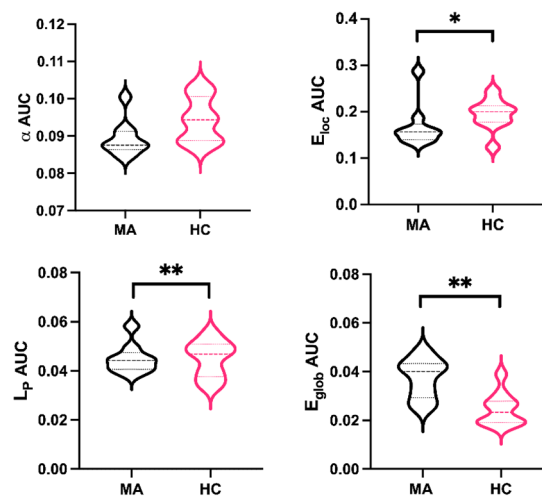


Figure 3. Differences AUC value of global topological metrics between two groups. *: $P < 0.05$ and **: $P < 0.01$, α : Small-world index, E_{loc} : Local efficiency, L_p : Characteristic shortest path length, E_{glob} : Global efficiency, MA: Methamphetamine and HC: healthy controls.

3.3. Regional topological networks

At the region level, there were significant group differences between nodal degree and nodal efficiency after FDR correction ($P < 0.05$). Compared with the HC group, the MA group showed significantly increased nodal degree in the FPN (e.g., left ventral anterior prefrontal cortex (vent aPFC_L) and left inferior parietal lobule (IPL_L)), VAN (e.g., right ventral prefrontal cortex (vPFC_R), right ventral prefrontal cortex (vFC_R) and left mid insula (Mid insula_L)), DAN (e.g., right dorsal frontal cortex (dFC_R)), SMN (e.g., left mid insula (Mid insula_L)), VN (e.g., right occipital lobe

(Occipital_R)) and DMN (e.g., right precuneus (Precuneus_R)). The results are shown in Table 2 and Figure 4(a). Similarly, in terms of nodal efficiency, we found that all seven large-scale brain networks in the MA group contained some brain regions that were significantly higher compared to the HC group. Differences were found mostly in VAN (e.g., right ventral prefrontal cortex (vPFC_R)), right ventral frontal cortex (vFC_R), left mid insula (Mid insula_L), as well as in FPN (right ventral inferior prefrontal cortex (vIPFC_R)), DAN (e.g., right dorsal frontal cortex (dFC_R) and left occipital lobe (Occipital_L)), SCN (e.g., bilateral basal ganglia (basal ganglia_L and basal ganglia_R), right thalamus (thalamus_R)), SMN (e.g., bilateral mid insula (Mid insula_L and Mid insula_R)), VN (left occipital lobe (Precuneus_L)) and DMN (left precuneus (Occipital_L)). The statistical and visualization results are shown in Table 3 and Figure 4(b).

Table 2. Group differences in node degree, with the MA group higher than HC.

ROIs	Coordinates (x, y, z)	Networks	MA Mean \pm SD	HA Mean \pm SD	P-value
vent aPFC_L	(-43, 47, 2)	FPN	5.260 \pm 1.377	3.263 \pm 0.787	0.002
IPL_L	(-48, -47, 49)	FPN	6.397 \pm 2.028	3.551 \pm 1.054	0.002
vPFC_R	(34, 32, 7)	VAN	2.428 \pm 0.951	0.594 \pm 0.705	0.001
vFC_R	(43, 1, 12)	VAN	4.501 \pm 1.305	0.976 \pm 0.824	0.001
Mid insula_L ¹	(-42, -3, 11)	VAN	6.255 \pm 3.215	1.164 \pm 0.477	0.001
dFC_R	(44, 8, 34)	DAN	5.611 \pm 1.565	2.329 \pm 1.640	0.001
Mid insula_L ²	(-36, -12, 15)	SMN	5.643 \pm 3.246	1.657 \pm 1.718	0.005
Occipital_R	(36, -60, -8)	VN	3.034 \pm 1.585	0.878 \pm 0.450	0.001
Precuneus_R	(11, -68, 42)	DMN	4.739 \pm 2.882	2.109 \pm 0.266	0.015

Note: MA: methamphetamine, HC: healthy controls, SD: standard deviation, L: left, R: right.

Table 3. Group differences in node efficiency, with the MA group higher than HC.

ROIs	Coordinates (x, y, z)	Networks	MA Mean \pm SD	HA Mean \pm SD	P-value
vIPFC_R	(39, 42, 16)	FPN	0.127 \pm 0.010	0.097 \pm 0.018	0.002
vPFC_R	(34, 32, 7)	VAN	0.112 \pm 0.008	0.045 \pm 0.034	0.001
vFC_R	(43, 1, 12)	VAN	0.130 \pm 0.018	0.052 \pm 0.036	0.001
Mid insula_L ¹	(-42, -3, 11)	VAN	0.134 \pm 0.027	0.085 \pm 0.019	0.002
dFC_R	(44, 8, 34)	DAN	0.137 \pm 0.020	0.097 \pm 0.017	0.001
Occipital_L ¹	(-29, -75, 28)	DAN	0.141 \pm 0.026	0.100 \pm 0.026	0.006
basal ganglia_L	(-20, 6, 7)	SCN	0.105 \pm 0.027	0.049 \pm 0.029	0.001
basal ganglia_R	(1, -26, 31)	SCN	0.118 \pm 0.012	0.085 \pm 0.017	0.002
thalamus_R	(11, -12, 6)	SCN	0.124 \pm 0.014	0.087 \pm 0.016	0.001
Mid insula_L ²	(33, -12, 16)	SMN	0.124 \pm 0.025	0.072 \pm 0.033	0.002
Mid insula_L ³	(-36, -12, 15)	SMN	0.137 \pm 0.026	0.072 \pm 0.034	0.001
Precuneus_L	(-3, -38, 45)	VN	0.123 \pm 0.023	0.096 \pm 0.012	0.006
Occipital_L ²	(-34, -60, -5)	DMN	0.122 \pm 0.023	0.038 \pm 0.043	0.001

Note: MA: methamphetamine, HC: healthy controls, SD: standard deviation, L: left, R: right.

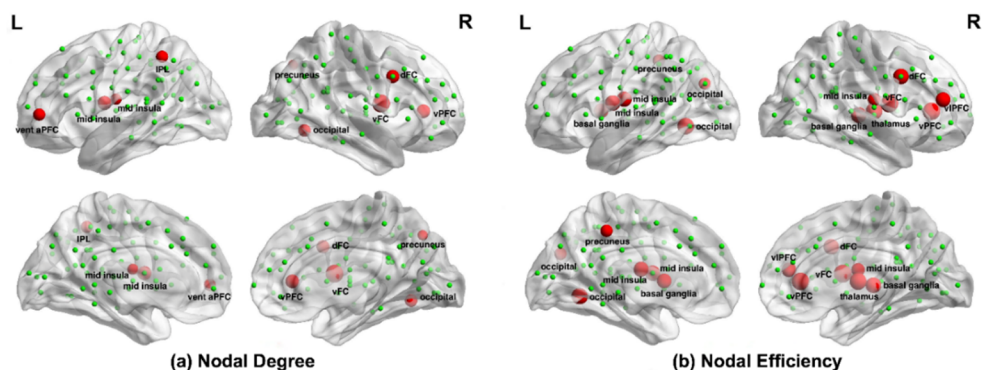


Figure 4. Brain regions with significant differences in node attribute changes in the MA group and HC group. The green balls indicate no statistical significance, while the red balls represent an increase in node attributes. vent aPFC: Ventral anterior prefrontal cortex, vPFC: Ventral prefrontal cortex, dFC: Dorsal frontal cortex, vFC: Ventral frontal cortex, IPL: Inferior parietal lobule, vIPFC: Ventral inferior prefrontal cortex.

3.4. Edge-based functional connectivity

The NBS-corrected results showed ten nodes and ten edges with significant FC differences between the MA and HC groups ($P < 0.05$). Among these results, four edges have significantly higher FC, while six are significantly lower than the HC group (Figure 5). The most significant increase edge is the FC between the left post cingulate (post cingulate_L) and right vent anterior prefrontal cortex (vent aPFC_R) brain regions (Figure 6(a),(b)). The most significant decrease edge is the FC between the left post cingulate and left medial prefrontal cortex (mPFC_L) (Figure 6(c),(d)). The edges with differences were mostly distributed in the VN, SMN, DAN, FPN and DMN (Figure 7).

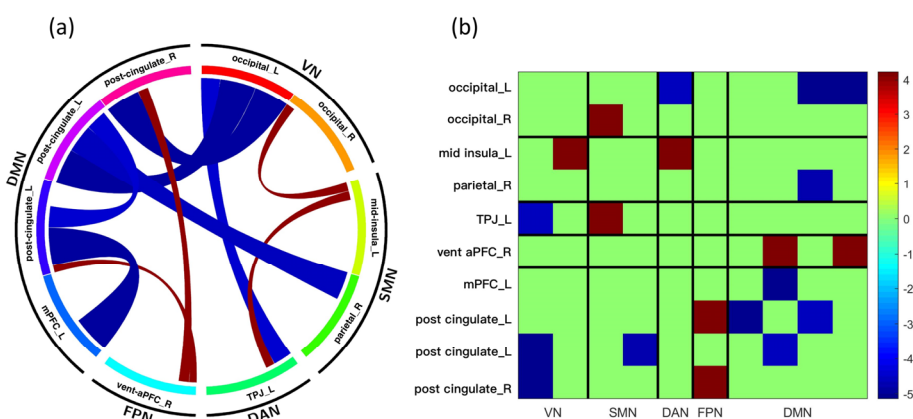


Figure 5. The NBS method shows significantly different nodes and edges between the MA and HC groups. (a, b) The 10 nodes defined by the Dosenbach atlas and the significant 10 unique Yeo network-pair edges in two groups are listed in the circus and heat map. For the heat map, the vertical axis represents the names of the brain regions, and the horizontal axis represents the large-scale brain networks to which the brain regions belong. For the color of the edges, red indicates that the FC has increased and while blue indicates that the FC has decreased after the NBS correction. L: Left, R: Right.

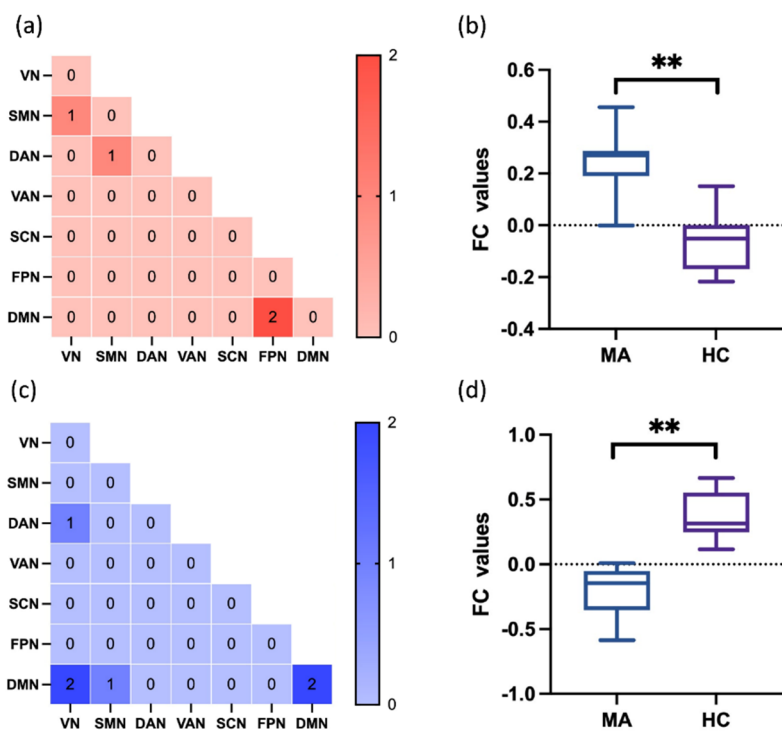


Figure 6. NBS results show the number of edges with significant differences. (a) and (c) Heatmaps show the number of significant edge changes in FC. (b) and (d) The box diagram shows the most significantly different edges in the increased or decreased FC. **: $P < 0.01$ after NBS correction.

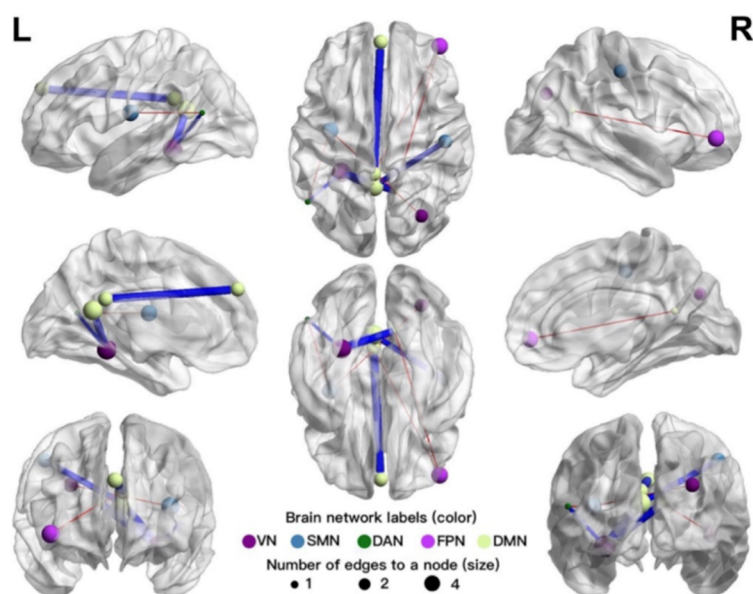


Figure 7. Visualization results of different edges after NBS correction. Red indicates that the FC is increased and blue means that the FC is decreased. VN: Visual network, SMN: Somatosensory network, DAN: Dorsal attentional network, FPN: Frontoparietal network, DMN: Default mode network.

3.5. Large-scale functional connectivity

Compared to the HC group, we found that the MA group significantly decreased within-network FC of the VN and SCN. In addition, the FC value between the VN and SCN of seven networks has reduced significantly ($P < 0.05$). No significant differences were found for other large-scale brain networks. The T-value diagram and visualization results are shown in Table 4 and Figure 8.

Table 4. Group differences in node efficiency, with the MA group higher than HC.

Networks	VN	SMN	DAN	VAN	SCN	FPN	DMN
VN	-2.7325*	-	-	-	-	-	-
SMN	0	0	-	-	-	-	-
DAN	0	0	0	-	-	-	-
VAN	0	0	0	0	-	-	-
SCN	-2.6490*	-	-	-	-3.1505*	-	-
FPN	0	0	0	0	0	0	-
DMN	0	0	0	0	0	0	0

Note: * represents a statistically significant difference ($P < 0.05$).

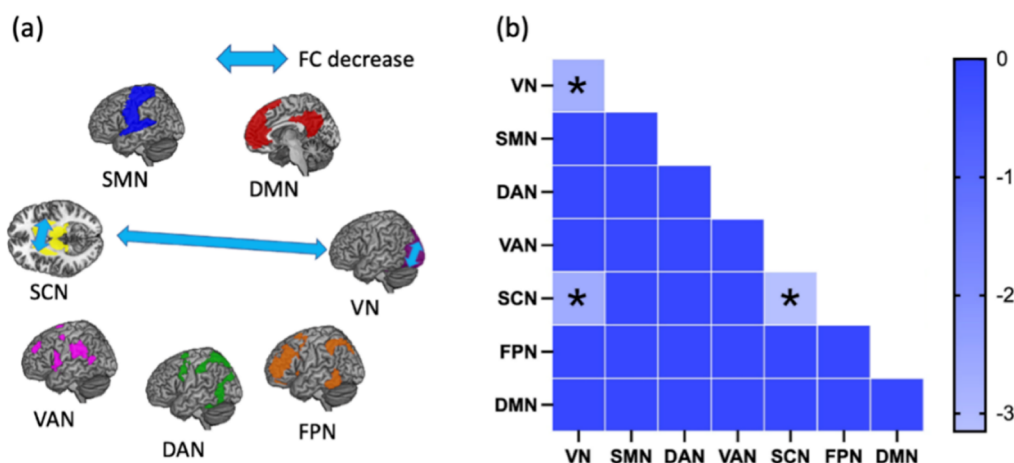


Figure 8. Visualization of functional connectivity differences between the two groups of large-scale brain networks. (a) Statistically significant decreases in functional connectivity between the SCN, VN and SCN-VN within the seven large-scale networks. (b) Heat map showing statistically significant T values between the VN and SCN and the VN and SCN within the seven networks ($P < 0.05$).

4. Discussion

In the present study, we compared the aberrant intrinsic functional brain topology at different levels between MA individuals after six months of abstinence and HC groups by graph theory and FC analysis methods. We found that the L_p showed a significantly decreased in global levels, while the E_{glob} and E_{loc} showed an increase considerably compared with the HC group. At the regional level,

we also found significant group differences in nodal degree and nodal efficiency in several brain networks after FDR correction. In addition, NBS methods extensively presented the anomalies in edge-based FC, involving almost all brain functional networks. Moreover, the FC of large-scale functional brain networks revealed a decrease within the VN and SCN and between the networks.

4.1. Discussion on aberrant brain topology at different levels

The human brain typically has a highly modular small-world network architecture, with different functional areas separated from each other and forming highly connected hub regions [30]. The properties of small-world networks can be reflected by well-defined metrics [31], such as C_p , L_p , E_{glob} and E_{loc} . Specifically, C_p represents the degree of aggregation between brain network nodes, and L_p can reflect the average value of information transfer pathways between different brain regions. E_{glob} and E_{loc} measure the brain's ability to transmit global and local information, respectively [32]. Small-world networks generally have high C_p and short L_p , ensuring efficient data transmission and processing while maintaining a balance between local specialization and global functional connectivity of the brain [33]. Liu et al. [34] found that MA disorder patients had economic small-world properties. We found that the brains of MA individuals after six months of abstinence also have small-world properties, suggesting a certain balance of integration and differentiation of brain functions. Moreover, compared to healthy individuals, MA withdrawal individuals showed lower L_p and higher E_{loc} and E_{glob} , indicating enhanced brain defenses and a tendency to convert the random networks. This is similar to the findings in patients with chronic MA addiction on EEG data [32,35]. These findings infer that long-term MA abuse disrupts the brain balance of small-world networks, and it still fails to recover to normal levels even after six months of abstinence.

In addition, we found a significant increase in nodal degree and efficiency in brain regions such as the frontal cortex, mid-insula, inferior parietal lobule, occipital lobe and precuneus, which are mostly distributed in the seven brain functional networks. These brain regions are closely associated with human cognition, emotional control, visual processing and dopamine secretion [36]. Previous studies have shown that long-term MA abuse can damage neurons, causing compensatory neuronal adaptations and eventually manifesting lasting cognitive or neurological deficits [37]. Li et al. [19] showed similar results in a population study of MA use disorders. Nodal degree and nodal efficiency reflect the efficiency of information interaction between different brain regions [32]. We hypothesize that long-term MA abuse leads to addiction in the brain, causing frequent communication between brain areas responsible for cognition, emotional control, visual processing and dopamine release. Addiction is the response of the brain to the environment, and withdrawing is challenging. Therefore, the findings may explain the high rate of MA relapse to some extent.

4.2. Discussion on aberrant FC at different scales

The edge-based FC analysis found several abnormal edges compared with the HC group, mostly distributed in the DMN, FPN, DAN, SMN and VN. It has been reported that MA addicts have clinical manifestations such as psychotic symptoms, anxiety, sleep disorder or craving after withdrawal, among which depression is the most apparent psychiatric symptom in the early stage of MA withdrawal [38]. The DMN is abnormal in many addictions and mental diseases, such as schizophrenia, depression, anxiety and heroin addiction [39]. Postcingulate is an essential component of the DMN. The emergence

of multiple psychiatric disorders is associated with functional abnormalities in the posterior cingulate gyrus [40]. The mPFC_L is implicated in self-processing decisions, such as personal information, autobiographical memories, future goals, or family decisions. The present study found significantly weakened FC between the mPFC_L and the post-cingulate, which may be related to cognitive deficits or psychiatric disorders in MA abstainers.

The FPN involves a wide range of exogenous cognitive processing and is critical for executive functions (e.g., working memory) and cognitive flexibility [41]. It can be divided into a left and a right part. The left FPN can be considered the language network, which also belongs to the cognitive category [42]. The vent aPFC_R is an essential component of the right FPN related to cognitive control [43]. In addition, the right FPN also has the main functions of activity inhibition, somatosensory perception, and pain processing, which is the core of the top-down cognitive control system [44]. Numerous studies have shown that the cognitive control system of the substance-addicted population is dysfunctional [9,10]. Previous research has shown that post-cingulate activity increases when heroin addicts are exposed to drug cues [45]. We found increased FC activity between the post-cingulate and vent aPFC_R, which may be related to the eagerness to seek drugs after methamphetamine withdrawals and ignore the harms of addictive behaviors. Unfortunately, after withdrawal, it still failed to improve the arousal of MA-independent individuals.

DAN allows one to focus attention and ignore external noise or changes in the environment [46]. As one of the core brain regions, the temporoparietal junction (TPJ) plays an essential role in attention and social cognition. TPJ is intensely active in visual detection, especially when the target is unexpected and needs to be redirected. For example, TPJ activity increases significantly during redirection but decreases during executive attention. The mid-insula is one of the components of SMN and is deeply involved in the brain reward-addicted system [10]. We found a significant increase in FC between TPJ and mid-insula and a substantial decrease between TPJ and occipital. It is hypothesized that it may be related to long-term abstinence from MA. After withdrawal, MA addicts redirect their attention and reshape the mapping of the reward system to the preoccupation system, thus gradually decreasing their concentration on drugs. In addition, Li et al. [19] studied the characteristics of the white matter structure network. They found that the connection strength between the reward and visual system in MA addicts was significantly enhanced. The present study also found that the FC between the occipital and mid-insula increased significantly, suggesting that it is related to long-term MA consumption. The long-term drug-seeking behaviors of MA addicts lead to long-term excitability on visual related functional regions. The long-term prominent visual stimulation leads to adaptive changes in the reward and visual systems [47]. The information transmitted between the visual and reward systems increases, leading to MA addicts being sensitive to drug-related clues. Although MA addicts reshape the mapping of the reward system to the preoccupation system to reduce preoccupation with the drug, the damage to the visual and the reward system from long-term MA abuse has not been restored, which may be one of the reasons why MA addicts relapse.

We also found significantly reduced FC between post-cingulate and parietal and between post-cingulate and occipital. The parietal is responsible for integrating sensory information from the outside and sensory feedback from the inside and merging this information into a coherent expression describing the connection between the body and the environment. Zhao et al. [48] studied the proactive inhibition ability of smokers. The findings showed that FC within the DMN and between the SMN-DMN was significantly decreased as the active inhibitory load increased. Moreover, it highlights that cigarette cues interfere with the effective dynamic interactions of active inhibition-related network

modules. We hypothesize that with more profound withdrawal, the active inhibition function of MA addicts continues to strengthen, and the active inhibition of the informational links between the DMN-SMN and the DMN-VN is conducive to reducing the interference of drug cues.

The large-scale FC analysis also found that the FC between VN-VN, SCN-SCN and VN-SCN was significantly decreased. SCN covers the dopamine neurotransmitter projection and transmission areas, such as the ventral dorsal tegmental area (VTA) of the midbrain, nucleus accumbens and substantia nigra [49]. The SCN is often associated with reward-seeking, encoding and impulsive behaviors related to the dopamine projection pathway [50]. Previous studies have pointed out that MA smoking primarily acts on the dopamine system, damaging neurons with long-term use, causing compensatory neuronal adaptations and eventually manifesting lasting cognitive or neurological deficits [51]. It is speculated that long-term MA intake will destroy the dopamine release pathway, thus leading to a significant decline in the internal FC of SCN. In addition, some studies have pointed out that the cingulate executive network can regulate SCN and affect individual performance in cognitive tasks, especially reward-related cognitive inhibition [52]. Moreover, Matteo et al. [53] found that early VN attention deficits can affect the SCN in Parkinson disease. Sun et al. [54] found that the lingual gyrus can regulate the human hormone level. The study found that the lingual gyrus region in the VN was activated. We hypothesized that long-term MA abuse would stimulate the above brain regions, altering hormone levels in the body and causing MA addicts to experience abnormal behavior, such as visual decline. However, after six months of withdrawal, MA addicts do not return to normal levels.

4.3. Limitations and future directions

Although our work revealed some abnormal brain networks, this study still has limitations that should be acknowledged. First, the sample size of this study was relatively small because of the strict exclusion criteria. This has an impact on the statistical power of the sample. Therefore, we will include more subjects to validate the experimental results in the future. Second, we focused only on alterations in topological properties of the brain between men with withdrawal and HC. It is necessary for future studies to include female abstainers. Third, the current study is cross-sectional and does not determine the causal effect of long-term interventional treatment on abnormal brain function. Further studies should explore the correlation between abnormal brain networks and clinical symptoms. In addition, the optical coherence tomography (OCT) imaging technique recently emerged as an alternative to fMRI in studying neural functional imaging of the brain and retina, which allows the functional response of neural circuits in the brain and retina [55–58]. Thus, we could adopt the OCT imaging technique to explore the aberrant intrinsic functional topology in methamphetamine-dependent individuals for future research.

5. Conclusions

In general, we investigate the aberrant intrinsic functional brain topology between MA individuals after six months of abstinence and HC groups via graph theory analysis. Results showed aberrant intrinsic functional brain topology at global and regional levels, and abnormal FC within or between several brain networks, including the FPN, VAN, DAN, SMN, VN, SCN and DMN. These findings suggest that MA-dependent individuals still have abnormal brain topology even when not exposed to drugs for long periods. After six months of withdrawal, some visual processing skills, object

recognition and memory deficits may not fully recover. This leads to the possibility of relapse behavior in MA-dependent individuals when confronted with MA-related cues, which may explain the relapse mechanism to a certain extent. Furthermore, the study provides an imaging basis for revealing the neural mechanism of MA-dependent after six months of abstinence. We hope that the results will be helpful in future neurobiological mechanisms and the treatment of MA addiction.

Use of AI tools declaration

The authors declare they have not used Artificial Intelligence (AI) tools in the creation of this article.

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Conflict of interest

The authors declare that there are no conflicts of interest.

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