Abnormal whole-brain functional connectivity in patients with primary insomnia

Chao Li
Mengshi Dong
Yi Yin
Kelei Hua
Shishun Fu
Guihua Jiang

Department of Medical Imaging,
The Affiliated Guangdong No 2
Provincial People's Hospital of
Southern Medical University,
The Third Clinical Medical College
of Southern Medical University,
Guangzhou, People's Republic of China

Abstract: The investigation of the mechanism of insomnia could provide the basis for improved understanding and treatment of insomnia. The aim of this study is to investigate the abnormal functional connectivity throughout the entire brain of insomnia patients, and analyze the global distribution of these abnormalities. Whole brains of 50 patients with insomnia and 40 healthy controls were divided into 116 regions and abnormal connectivities were identified by comparing the Pearson's correlation coefficients of each pair using general linear model analyses with covariates of age, sex, and duration of education. In patients with insomnia, regions that relate to wakefulness, emotion, worry/rumination, saliency/attention, and sensory-motor showed increased positive connectivity with each other; however, regions that often restrain each other, such as regions in salience network with regions in default mode network, showed decreased positive connectivity. Correlation analysis indicated that some increased positive functional connectivity was associated with the Self-Rating Depression Scale, Insomnia Severity Index, and Pittsburgh Sleep Quality Index scores. According to our findings, increased and decreased positive connectivities suggest function strengthening and function disinhibition, respectively, which offers a parsimonious explanation for the hyperarousal hypothesis in the level of the whole-brain functional connectivity in patients with insomnia.

Keywords: primary insomnia, hyperarousal hypothesis, resting-state functional magnetic resonance imaging, functional connectivity, whole brain

Introduction

Sleep is as vital to human as water and food. However, many sleep disorders' neural mechanisms are still elusive. Among them, insomnia has received relatively more attention as one of the most prevalent sleep disorders.¹ Insomnia² is an independent psychiatric syndrome that results in difficulties in falling asleep or maintaining sleep for at least 1 month and is associated with a sequela of daytime impairment or clinically significant distress unrelated to any other medical, physical, or psychiatric disorder. Insomnia confers an increased risk for physiological and psychiatric disorders, such as increased cardiovascular morbidity and mortality, and especially depression and anxiety, causing pronounced direct and indirect socioeconomic burden.³

In the past decade, neuroimaging techniques have made significant contributions to the study of insomnia. A series of studies⁴⁻⁹ found that patients with insomnia showed abnormal metabolism, function, and structure related to widespread cortical and subcortical areas, including the ascending reticular activating system, amygdala, hippocampus, cingulate cortex, insula, parietal, frontal cortex, and caudate nucleus. From the functional connectivity network's point of view,¹⁰⁻¹³ these regions mainly belong to the default mode network (DMN), salience network (SN), affective network (AN), central-executive network (CEN), and subcortical regions (SUB).

Correspondence: Guihua Jiang
Department of Medical Imaging,
The Affiliated Guangdong No 2
Provincial People's Hospital of Southern
Medical University, The Third Clinical
Medical College of Southern Medical
University, No 466, Xingang Road,
Haizhu District, Guangzhou 510317,
People's Republic of China
Tel/fax +86 20 8916 8071
Email jiangguihua1970@163.com

However, little is known about the functional connectivities among these regions.

Indeed, a few studies^{14–18} have emerged to investigate functional connectivity in patients with insomnia. However, results of some of these studies are inconsistent.^{17,18} Most importantly, most of these studies only focused on single or a few seed regions. It was difficult to fully clarify the neural mechanism of insomnia because the brain interconnects among widespread regions.¹⁰ Given the possibility that distributed abnormality of functional connectivity may exist in the rest of the brain of insomnia patients, we think it will be helpful to study functional connectivity from the perspective of whole brain for a better understanding of the neural mechanism of insomnia.

Thus, the aim of this study is to investigate the abnormal functional connectivities on the whole-brain level and analyze the global distribution of these abnormalities. Considering the overarching neural mechanism of insomnia, namely hyperarousal hypothesis,¹⁹ we hypothesized that global distribution of these abnormal functional connectivities throughout the entire brain of insomnia patients supports the hyperarousal hypothesis.

Patients and methods

Subjects

All patients with insomnia were recruited from Guangdong No 2 Provincial People's Hospital from April 2014 to May 2016. Each study participant gave written informed consent, and the guidelines of the Declaration of Helsinki were followed in the study protocol. The study protocol was approved by the Ethics Committee of Guangdong No 2 Provincial People's Hospital. The inclusion criteria for patients with insomnia were as follows: 1) conformity to the definition of primary insomnia by the *Diagnostic and Statistical Manual of Mental Disorders*, Fourth Edition;²⁰ 2) insomnia lasting at least 1 month, with a complaint of difficulty falling asleep, maintaining sleep, or early awakening; 3) no other sleep disorders (such as sleep-related movement disorders, hypersomnia, or parasomnia) or other psychiatric disorders determined by a semistandardized psychiatric and sleep-related interview conducted by an experienced psychiatrist; 4) insomnia not due to the effects of any medications or substance abuse, such as caffeine, nicotine, or alcohol; 5) right-hand dominance; 6) no serious organic disease; 7) no foreign implants in the body; 8) age 25–60 years; 9) no abnormal signal as verified by conventional T1-weighted or T2 fluid-attenuated inversion recovery magnetic resonance (MR) imaging; and

10) head motion less than 1.5 mm or 1.5° during MR imaging. The exclusion criterion for recruitment of patients was that patients must meet all of the above inclusion criteria; if not, the patient would be excluded. Five patients were excluded because of head motion greater than 1.5 mm or 1.5° and 5 patients were excluded because of brain infarcts and abnormal signal intensity. Finally, 50 patients with insomnia (21 men, 29 women; mean age, 39.64±10.42 years) were included in this study.

We also recruited 40 healthy control subjects matching in levels age, sex, and education (10 men and 30 women; mean age 39.20±8.47 years) from the local community by using advertisements. All control subjects met the following criteria: 1) good sleep quality and an Insomnia Severity Index (ISI) score of less than 7; 2) no history of swing shifts, shift work, or sleep complaints; 3) no medication or substance abuse, such as caffeine, nicotine, or alcohol; and 4) fulfillment of inclusion criteria 5–10 for the patients with insomnia.

Questionnaires

All volunteers were asked to complete a number of questionnaires, including the Pittsburgh Sleep Quality Index (PSQI),²¹ the ISI,²² the Self-Rating Anxiety Scale (SAS),²³ and the Self-Rating Depression Scale (SDS).²⁴

MR imaging

MR imaging data were obtained by using a 1.5T MR scanner (Achieva Nova-Dual; Philips, Best, the Netherlands) in the *BLINDED*. Head motion was restricted by a belt and foam pads. During the resting state of the functional MR imaging scanning, subjects were instructed to rest with their eyes closed and heads still. The resting-state functional MR images were obtained using a gradient-echo planar imaging sequence (interleaved scanning, repetition time/echo time =2,500 ms/50 ms, matrix =64×64, field of view =224×224 mm, flip angle =90°, section thickness =4 mm, gap =0.8 mm, 27 sections covering the whole brain were positioned along the anterior commissure-posterior commissure line). A total of 240 volumes were acquired in approximately 10 minutes.

Data preprocessing

An overview of the data processing framework is summarized in Figure 1. Data Processing Assistant for Resting-State Functional MR Imaging toolbox (<http://www.restfmri.net/forum/DPARSF>) was used to preprocess the

Figure 1 Overall data processing framework.

Abbreviations: AAL, anatomical automatic labeling; GLM, general linear model; FC, functional connectivity.

imaging data. Volumes at the first 10 time points were discarded so that magnetization reached a steady state and subjects had adapted to the MR scanning noise. The slice timing and realignment for head motion correction and spatial normalization were conducted according to the standard Montreal Neurologic Institute template, resampled into a voxel size of $3 \times 3 \times 3$ mm³. In order to avoid introducing artificial local spatial correlations, we did not apply spatial smoothing.^{25,26} To reduce the effects of low-frequency drift and high-frequency noise, the data were processed to remove linear trends and be filtered temporally (band pass, 0.01–0.08 Hz). Nuisance covariates including cerebrospinal fluid signals, white matter signals, and 6 head motion parameters were regressed from the functional MR imaging data. Of note, we did not regress out the global signal because of an ongoing controversy over removing the global signal during the preprocessing of resting-state functional MR imaging data.²⁷ The residuals of these regressions were used for the following analysis. Then, mean time series of 116 regions of interests (ROI) defined by Automated Anatomic Labeling atlas²⁸ were extracted for each individual by averaging the functional

MR imaging time series throughout all voxels in each ROI. Functional connectivity between each pair of ROI was evaluated using Pearson's correlation coefficients, producing $(116 \times 115)/2 = 6,670$ functional connectivities for each subject. Fisher *r*-to-*z* transformation was performed for all functional connectivities to improve the normality of the correlation coefficients.

Statistical analysis

Two-sample 2-tailed *t*-tests were used to compare differences in age, duration of education, PSQI, ISI, SAS, and SDS scores between patients with insomnia and healthy controls. Two-tailed Pearson's chi-square test was performed to determine differences of sex between the 2 groups.

General linear model analyses with covariates of age, duration of education, and sex were used to compare the differences of functional connectivities between the 2 groups. Significantly different connectivities were chosen as connectivities of interest (COIs).

Correlations between clinical characteristics and COIs were computed using Pearson's correlation analyses ($P < 0.05$).

Table 1 Demographics and clinical characteristics of all participants

Parameter	Patients with insomnia	Healthy controls	P-value
Sex			0.09
No of men	21	10	
No of women	29	30	
Age (years)	39.64±10.42	39.20±8.47	0.83
Duration (months)	40.31±44.09		
Education (years)	7.38±3.65	8.78±3.96	0.07
PSQI	12.32±2.93	5.88±2.19	<0.001
ISI	12.56±2.96	5.78±2.34	<0.001
SAS	51.92±10.87	42.73±5.73	<0.001
SDS	55.06±8.93	43.95±5.34	<0.001

Note: Unless otherwise noted, data are mean ± standard deviation.

Abbreviations: PSQI, Pittsburgh Sleep Quality Index; ISI, Insomnia Severity Index; SAS, Self-Rating Anxiety Scale; SDS, Self-Rating Depression Scale.

Results

Demographic and clinical characteristics

As shown in Table 1, the patients with insomnia and the healthy controls showed no significant differences regarding age ($P=0.83$), sex ($P=0.09$), and education level ($P=0.07$).

Patients with insomnia had higher PSQI, ISI, SAS, and SDS scores than the healthy controls (all $P<0.001$).

Functional connectivity of the whole brain

Overall, there were 36 functional connectivities that showed significant differences between the patients with insomnia and the healthy controls (all $P<0.05$, uncorrected). These included 28 increased functional connectivities (Table 2 and Figure 2) and 8 decreased functional connectivities (Table 3 and Figure 3). Regions (nodes) that constituted the 36 functional connectivities are shown in Table S1.

Compared with the healthy controls, patients with insomnia showed increased functional connectivities among wide-spread regions, and these regions belong to some functional connectivity networks as follows: between DMN and sensory-motor network (SMN) and orbital part of frontal lobe (ORB), between SN and SMN and ORB, between AN and SMN and cerebellum (CER), between ORB and SMN and CER, between SUB and SMN and ORB, between SMN and SMN

Table 2 Increased positive functional connectivities in patients with insomnia

COI	Network	z value in patients with insomnia	z value in healthy controls	Two-sample 2-tailed t-tests	P-value
R precuneus-L supramarginal gyrus	DMN-SMN	0.43±0.21	0.30±0.26	2.64	0.0096
R precuneus-R olfactory cortex	DMN-SMN	0.29±0.18	0.19±0.17	2.75	0.0070
L parahippocampal gyrus-L orbital part of middle frontal gyrus	DMN-ORB	0.22±0.17	0.12±0.22	3.22	0.0017
L hippocampus-L orbital part of superior frontal gyrus	DMN-ORB	0.17±0.19	0.06±0.19	3.82	0.0002
L insula lobe-L orbital part of middle frontal gyrus	SN-ORB	0.18±0.23	0.06±0.23	2.72	0.0076
L insula lobe-R rectus gyrus	SN-ORB	0.16±0.23	0.04±0.21	2.68	0.0086
L anterior cingulate gyrus-L olfactory cortex	SN-SMN	0.36±0.16	0.26±0.20	2.83	0.0057
R insula lobe-L olfactory cortex	SN-SMN	0.18±0.18	0.08±0.20	2.98	0.0036
R anterior cingulate gyrus-L olfactory cortex	SN-SMN	0.31±0.17	0.21±0.21	2.80	0.0061
R amygdala-R cerebellum 10	AN-CER	0.15±0.21	0.03±0.23	2.66	0.0091
R amygdala-L supramarginal gyrus	AN-SMN	0.30±0.22	0.19±0.19	2.66	0.0090
R orbital part of median superior frontal gyrus-R cerebellum 10	ORB-CER	0.21±0.19	0.09±0.23	2.70	0.0082
L orbital part of superior frontal gyrus-R cerebellum 10	ORB-CER	0.21±0.21	0.07±0.20	3.27	0.0014
L orbital part of superior frontal gyrus-R cerebellum 7b	ORB-CER	0.28±0.24	0.12±0.29	3.36	0.0011
L rectus gyrus-R cerebellum 10	ORB-CER	0.23±0.21	0.10±0.20	2.69	0.0083
R rectus gyrus-R cerebellum 10	ORB-CER	0.26±0.22	0.10±0.21	2.67	0.0088
R rectus gyrus-R cerebellum 7b	ORB-CER	0.43±0.18	0.30±0.20	3.03	0.0030
R rectus gyrus-L olfactory cortex	ORB-SMN	0.27±0.22	0.13±0.28	2.67	0.0088
L putamen-R rectus gyrus	SUB-ORB	0.23±0.18	0.13±0.16	2.67	0.0088
R putamen-L olfactory cortex	SUB-SMN	0.21±0.16	0.11±0.18	2.77	0.0067
L putamen-R olfactory cortex	SUB-SMN	0.26±0.17	0.18±0.15	2.73	0.0075
L supramarginal gyrus-R cerebellum 4.5	SMN-CER	0.39±0.23	0.27±0.25	2.63	0.0098
L olfactory cortex-L cerebellum 7b	SMN-CER	0.17±0.17	0.07±0.21	2.77	0.0067
R supplementary motor area-R olfactory cortex	SMN-SMN	0.28±0.16	0.19±0.17	3.13	0.0023
R olfactory cortex-L superior temporal gyrus	SMN-SMN	0.29±0.17	0.18±0.18	2.73	0.0075
R olfactory cortex-L supramarginal gyrus	SMN-SMN	0.17±0.19	0.07±0.16	2.75	0.0070
R olfactory cortex-L paracentral lobule	SMN-SMN	0.27±0.17	0.18±0.16	2.73	0.0075
R superior parietal gyrus-R cerebellum 7b	CEN-CER	0.36±0.35	0.17±0.35	2.65	0.0094

Abbreviations: COI, connectivity of interest; R, right; L, left; DMN, default mode network; SMN, sensory-motor network; ORB, orbital part of frontal lobe; SN, salience network; AN, affective network; CER, cerebellum; SUB, subcortical regions; CEN, central-executive network.

Figure 2 Increased positive functional connectivities in patients with insomnia.

Notes: (A) From the network point of view, ribbon represents connectivity. (B and C) From the brain region point of view. Visualization (A) was created using Circos (<http://circos.ca/>). (B and C) were visualized with the BrainNet Viewer.⁴⁰

Abbreviations: AN, affective network; CEN, central-executive network; CER, cerebellum; DMN, default mode network; ORB, orbital part of frontal lobe; SMN, sensory-motor network; SN, salience network; SUB, subcortical regions.

and CER, and between CEN and CER. Details can be seen in Table 2 and Figure 2.

Compared with the healthy controls, patients with insomnia showed decreased functional connectivities among several regions, and these regions belong to some functional connectivity networks as follows: between DMN and SN, between ORB and SMN, and between SUB and CER. Details can be seen in Table 3 and Figure 3.

Relationship between aberrant functional connectivities and clinical characteristics

Table 4 shows the results of correlation analysis. We found that only regions with increased functional connectivity correlated with PSQI, ISI, and SDS (all $P < 0.05$). Functional connectivities between some CER regions and 3 ORB regions, 1 AN region, 1 CEN region, and 1 SMN region were significantly correlated with PSQI ($r = -0.34, -0.34, -0.33, -0.29, -0.40, -0.38$). In addition, functional connectivities

between a DMN region and an ORB region and between 2 SMN regions were significantly positively correlated with ISI ($r = 0.29, 0.31$). Besides, functional connectivities between a DMN region and an SMN region and between an ORB region and an SN region were significantly negatively correlated with SDS ($r = -0.28, -0.30$).

Discussion

The current study investigated the whole-brain functional connectivity in patients with insomnia and healthy controls. The most interesting finding is global distribution of these abnormal functional connectivities throughout the entire brain of insomnia patients. Regions that relate to wakefulness, emotion, worry/rumination, saliency/attention, and sensory-motor showed increased positive connectivity with each other; however, regions that often restrain each other, such as SN with DMN and pallidum with thalamus, showed decreased positive connectivity in patients with insomnia.

Table 3 Decreased positive functional connectivities in patients with insomnia

COI	Network	z value in patients with insomnia	z value in healthy controls	Two-sample 2-tailed t-tests	P-value
L anterior cingulate gyrus-L angular gyrus	SN-DMN	0.32±0.22	0.45±0.20	-2.88	0.0049
R anterior cingulate gyrus-L angular gyrus	SN-DMN	0.31±0.19	0.43±0.20	-2.77	0.0066
L pallidum-R thalamus	SUB-SUB	0.25±0.19	0.35±0.15	-2.70	0.0081
L orbital part of superior frontal gyrus-L inferior occipital gyrus	ORB-SMN	0.18±0.25	0.34±0.24	-3.37	0.0011
R orbital part of superior frontal gyrus-R cuneus	ORB-SMN	0.28±0.21	0.40±0.23	-3.08	0.0026
L orbital part of middle frontal gyrus-L inferior occipital gyrus	ORB-SMN	0.18±0.27	0.33±0.25	-3.03	0.0031
R orbital part of median superior frontal gyrus-R cuneus	ORB-SMN	0.31±0.22	0.43±0.23	-3.00	0.0033
Vermis 9-Vermis 10	CER-CER	0.18±0.24	0.29±0.18	-2.63	0.0099

Abbreviations: COI, connectivity of interest; L, left; SN, salience network; DMN, default mode network; R, right; SUB, subcortical regions; ORB, orbital part of frontal lobe; SMN, sensory-motor network; CER, cerebellum.

Figure 3 Decreased positive functional connectivities in patients with insomnia.

Notes: (A) From the network point of view, ribbon represents connectivity. (B) From the brain region point of view. Visualization (A) was created using Circos (<http://circos.ca/>). (A and B) were visualized with the BrainNet Viewer.⁴⁰

Abbreviations: CER, cerebellum; DMN, default mode network; ORB, orbital part of frontal lobe; SMN, sensory-motor network; SN, salience network; SUB, subcortical regions.

These increased or decreased positive connectivity may contribute to hyperarousal in patients with insomnia.

We should note that the results of our study should be regarded as descriptive. Since there are so many comparisons ($116 \times 115 / 2 = 6,670$), correction for multiple comparisons may make most of the differences not statistically significant. However, from the perspective of the whole brain, this study offers a descriptive representation of the global distribution of the abnormal connectivity in patients with insomnia.

Although the neural mechanisms of insomnia are largely unknown, the most compelling hypothesis that explains insomnia is hyperarousal hypothesis.¹⁹ According to this hypothesis, insomnia patients commonly show increased glucose metabolism in multiple brain areas²⁹ and elevated

energy requirements in the whole gray matter.³⁰ Consistent with previous studies and as complementary, our study explains the hyperarousal hypothesis in the level of the whole-brain functional connectivity.

In the current study, we tended to interpret increased positive connectivity among regions in patients with insomnia as function strengthening, while interpreting decreased positive connectivity among regions in patients with insomnia as function disinhibition. Both function strengthening and function disinhibiting may cause hyperarousal in patients with insomnia. To be specific, increased positive connectivity was mainly between DMN, SN, and SUB (bilateral putamen) and SMN and ORB, between AN (right amygdala) and ORB and SMN and CER, and between SMN and CER. It is not difficult to see that the aforementioned

Table 4 Associations of aberrant functional connectivities and clinical characteristics

COI	Network	PSQI		ISI		SDS	
		r	P-value	r	P-value	r	P-value
L olfactory cortex-L cerebellum 7b	SMN-CER	-0.38	0.007				
R orbital part of middle frontal gyrus-R cerebellum 10	ORB-CER	-0.34	0.017				
L rectus gyrus-R cerebellum 10	ORB-CER	-0.33	0.019				
R rectus gyrus-R cerebellum 7b	ORB-CER	-0.34	0.015				
R amygdala-R cerebellum 10	AN-CER	-0.29	0.040				
R superior parietal lobule-R cerebellum 7b	CEN-CER	-0.40	0.004				
L orbital part of middle frontal gyrus-L parahippocampal gyrus	ORB-DMN			0.29	0.042		
R olfactory cortex-L paracentral lobule	SMN-SMN			0.31	0.027		
R rectus gyrus-L Insula lobe	ORB-SN					-0.30	0.033
L supramarginal gyrus-R precuneus	SMN-DMN					-0.28	0.049

Abbreviations: COI, connectivity of interest; L, left; SMN, sensory-motor network; CER, cerebellum; R, right; ORB, orbital part of frontal lobe; AN, affective network; CEN, central-executive network; DMN, default mode network; SN, salience network; PSQI, Pittsburgh Sleep Quality Index; ISI, Insomnia Severity Index; SDS, Self-Rating Depression Scale.

networks and SUB are closely associated with wakefulness (putamen),³¹ self-referential processing, introspection, worry and rumination (DMN),^{32,33} saliency, switching, attention and control (SN),³⁴ emotion (amygdala, ORB),^{35,36} and sensory-motor (SMN). Consequently, it is very possible that increased positive connectivity indicates that these networks promote each other, resulting in hyperemotion, hypersensorimotor, and finally hyperarousal, which is in line with hyperarousal hypothesis.¹⁹ Our speculations were also partially supported by previous studies that found emotion impairment in insomnia patients^{37,38} as well as our correlation analyses that several increased connectivities were associated with depressive emotion, severity of insomnia, and sleep quality.

However, temporal coupling between left pallidum and right thalamus and between SN and DMN showed decreased positive connectivity. An important function of the external pallidum was promoting sleep³¹ through inhibiting pyramidal cells. Animal studies have shown that rats with cell body-specific lesioning of the external pallidum lead to a dramatic increase in total wakefulness and pronounced fragmentation of non-rapid eye movement sleep and wakefulness. Interestingly, however, the thalamus is responsible for promoting wakefulness and is a component of the ascending reticular activating system.³⁹ Thus, decreased positive connectivity between these 2 SUB may indicate that the mutual inhibited effect between the 2 SUB has been compromised, which may result in decreased suppression of the thalamus during sleep, causing insomnia. We also detected decreased positive connectivity between SN and DMN. One key function of SN was switching between DMN and CEN by deactivating DMN and activating CEN.³⁴ Weakened association between SN and DMN in patients with insomnia indicated that the deactivating effect of SN on DMN is diminished, resulting in hyperactivity of DMN. Taking into account the functions of DMN,^{32,33} weakened association between SN and DMN may underpin the high levels of self-referential processing, introspection, worry, and rumination in patients with insomnia.¹⁸

Limitations

Our study had some limitations. First, in addition to correlation coefficient, there are many indexes to measure functional connectivity and network properties, including degree, clustering coefficient, lengths of paths, and so on. Further studies could use more indexes to investigate neural mechanism in insomnia. Second, although we found aberrant functional connectivity in patients with insomnia, we did not know the direction of the functional connectivity, which

needs further study with Granger causality analysis or other methods. Third, this was a cross-sectional study, and the causal relationship between insomnia and aberrant functional connectivity remains unclear. Longitudinal studies may help to resolve the question.

Conclusion

Our findings provide evidence for aberrant functional connectivity in widespread regions in insomnia patients. Furthermore, this offers a parsimonious explanation for the hyperarousal hypothesis in the level of the whole-brain functional connectivity.

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Disclosure

The authors report no conflicts of interest in this work.

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Supplementary material

Table S1 Brain regions and their abbreviations used in this paper

Region	Abbreviation	Network	Region	Abbreviation	Network
L orbital part of superior frontal gyrus	ORBsup.L	ORB	R precuneus	PCUN.R	DMN
L orbital part of middle frontal gyrus	ORBmid.L	ORB	L paracentral lobule	PCL.L	SMN
R supplementary motor area	SMA.R	SMN	L putamen	PUT.L	SUB
L olfactory cortex	OLF.L	SMN	R putamen	PUT.R	SUB
R olfactory cortex	OLF.R	SMN	L superior temporal gyrus	STG.L	SMN
R orbital part of median superior frontal gyrus	ORBsupmed.R	ORB	R cerebellum 4.5	CRBL45.R	CER
L rectus gyrus	REC.L	ORB	L cerebellum 7b	CRBL7b.L	CER
R rectus gyrus	REC.R	ORB	R cerebellum 7b	CRBL7b.R	CER
L insula lobe	INS.L	SN	R cerebellum 10	CRBL10.R	CER
R insula lobe	INS.R	SN	R orbital part of superior frontal gyrus	ORBsup.R	ORB
L anterior cingulate gyrus	ACG.L	SN	R cuneus	CUN.R	SMN
R anterior cingulate gyrus	ACG.R	SN	L inferior occipital gyrus	IOG.L	SMN
L hippocampus	HIP.L	DMN	L angular gyrus	ANG.L	DMN
L parahippocampal gyrus	PHG.L	DMN	L pallidum	PALL	SUB
R amygdala	AMYG.R	AN	R thalamus	THA.R	SUB
R superior parietal gyrus	SPG.R	CEN	Vermis9	Vermis9	CER
L supramarginal gyrus	SMG.L	SMN	Vermis10	Vermis10	CER

Abbreviations: L, left; ORB, orbital part of frontal lobe; R, right; DMN, default mode network; SMN, sensory-motor network; SUB, subcortical regions; CER, cerebellum; SN, salience network; AN, affective network; CEN, central-executive network.

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