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ORIGINAL RESEARCH

Long-term total sleep deprivation decreases the default spontaneous activity and connectivity pattern in healthy male subjects: a resting-state fMRI study

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Correspondence: Ren-Lai Zhou Beijing Key Lab of Applied Experimental Psychology, School of Psychology, and National Key Laboratory of Cognitive Neuroscience and Learning, Beijing Normal University, Beijing 100875, People's Republic of China Tel +86 010 58802021 Email rlzhou@bnu.edu.cn

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Department of Radiology, The First Affiliated Hospital of Nanchang University, Number 17, YongWai Zheng Street, DongHu District, Nanchang 330006, Jiangxi, People's Republic of China Tel +86 791 8869 2582 Email honghan_gong@sina.com **Objective:** The aim of this study is to use resting-state functional connectivity (rsFC) and amplitude of low-frequency fluctuation (ALFF) methods to explore intrinsic default-mode network (DMN) impairment after sleep deprivation (SD) and its relationships with clinical features.

Methods: Twelve healthy male subjects underwent resting-state functional magnetic resonance imaging twice: once following rested wakefulness (RW) and the other following 72 hours of total SD. Before the scans, all subjects underwent the attention network test (ANT). The independent component analysis (ICA), rsFC, and ALFF methods were used to examine intrinsic DMN impairment. Receiver operating characteristic (ROC) curve was used to distinguish SD status from RW status.

Results: Compared with RW subjects, SD subjects showed a lower accuracy rate (RW =96.83%, SD = 77.67%; P < 0.001), a slower reaction time (RW = 695.92 ms; SD = 799.18 ms; P = 0.003), a higher lapse rate (RW =0.69%, SD =19.29%; P < 0.001), and a higher intraindividual coefficient of variability in reaction time (RW =0.26, SD =0.33; P=0.021). The ICA method showed that, compared with RW subjects, SD subjects had decreased rsFC in the right inferior parietal lobule (IPL, BA40) and in the left precuneus (PrC)/posterior cingulate cortex (PCC) (BA30, 31). The two different areas were selected as regions of interest (ROIs) for future rsFC analysis. Compared with the same in RW subjects, in SD subjects, the right IPL showed decreased rsFC with the left PrC (BA7) and increased rsFC with the left fusiform gyrus (BA37) and the left cluster of middle temporal gyrus and inferior temporal gyrus (BA37). However, the left PrC/ PCC did not show any connectivity differences. Compared with RW subjects, SD subjects showed lower ALFF area in the left IPL (BA39, 40). The left IPL, as an ROI, showed decreased rsFC with the right cluster of IPL and superior temporal gyrus (BA39, 40). ROC curve analysis showed that the area under the curve (AUC) value of the left IPL was 0.75, with a cutoff point of 0.834 (mean ALFF signal value). Further diagnostic analysis exhibited that the AUC alone discriminated SD status from RW status, with 75% sensitivity and 91.7% specificity.

Conclusion: Long-term SD disturbed the spontaneous activity and connectivity pattern of DMN.

Keywords: sleep deprivation, amplitude of low-frequency fluctuation, default-mode network, functional magnetic resonance imaging, functional connectivity, independent component analysis, receiver operating characteristic curve

Introduction

Sleep deprivation (SD) is common and can be a serious problem negatively affecting health. Sleep has been reported to allow the removal of free radicals accumulated

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in the brain during wakefulness,¹ while SD can induce oxidative damage and disturb the removal of free radicals in the brain.² Long-term SD is harmful to health and can lead to multisystemic and multiorgan dysfunction, causing maladaptive emotional regulation, exaggerated neural reactivity, aversive experiences, and negative metabolic, psychological, physiological, or even behavioral reactivity.³⁻⁶ However, their mechanisms remain unclear.

Anatomically the "default-mode network" (DMN) spans the bilateral inferior parietal lobule (IPL), posterior cingulate cortex (PCC), precuneus (PrC), medial prefrontal cortex (MPFC), retrosplenial cortex, and parts of the hippocampal formation and medial temporal lobe.7,8 Recent research has suggested that the DMN might be associated with collection and evaluation of information,9 self-referential mental activity,¹⁰ extraction of episodic memory,¹¹ emotion and anxiety,12,13 and mind wandering or daydreaming.14 More and more researches have been devoted to exploring the DMN changes in various kinds of diseases, including cognitive impairment,^{15,16} chronic pain,¹⁷ hepatic encephalopathy,¹⁸ and autism.¹⁹ Furthermore, the focus of many researches has been on delineating the regional effects of SD on DMN. Gujar et al²⁰ found that deactivation within DMN regions alone discriminated SD from sleep-control subjects with 93% sensitivity and 92% specificity. In addition, the relative balance of deactivation within these default nodes significantly correlated with the amount of prior sleep in the control group. Bosch et al²¹ found that SD reduced resting-state functional connectivity (rsFC) between PCC and bilateral anterior cingulate cortex, in addition to enhancing rsFC between the dorsal nexus and distinct areas in the right dorsolateral PFC. Chee and Chuah²² found SD-related reduced IPL deactivation during a visual short-term memory task. Horovitz et al^{23,24} revealed reductions in the IPL-MPFC rsFC during deep sleep and after partial SD. De Havas et al²⁵ also found that the IPL node of the DMN was consistently impaired and might represent an early marker for the effects of 24-hour SD, as well as serving as an indicator of hitherto-unexplored behavioral impairments. Although these studies provide insights into the neural events occurring during SD, far less is still known about the effect of long-term SD on intrinsic DMN node impairment.

It is proposed that resting-state functional magnetic resonance imaging (rfMRI), one of the hot areas in neuroimaging and one that is suitable for the mechanism research of central nervous system, can detect the spontaneous neuronal activity of the human brain and provide new insights into the pathophysiology of disease, because of its advantages in not requiring exposure to radioactive tracers, accurate positioning, and ease of combining functional imaging with structural imaging. Amplitude of low-frequency fluctuation (ALFF) is an index in which the square root of the power spectrum is integrated in a low-frequency range for detecting the regional spontaneous neuronal activity in blood oxygenation-level dependent (BOLD) signal.²⁶ Previous studies have demonstrated that the ALFF showed good-to-moderate test-retest reliability.27,28 The simple calculation and reliable characterization of the ALFF measurement makes it a suitable and useful tool for rfMRI data analysis to investigate a disease trait. The ALFF measurement has been successfully applied in the exploration of functional modulations and characterization of the pathophysiological changes during different sleep conditions, such as the condition of being awake²⁹ and light sleep.³⁰ However, little attention has been given to the possibility that SD may be equally well described on the basis of differences in resting-state modes of spontaneous brain activation. The rsFC indicates temporal correlations among neural or hemodynamic signals arising from distinct brain regions, and it measures the signal synchrony of ALFF activity among different brain areas, while it still cannot ascertain which brain area is abnormal when abnormal rsFC is seen between two or more areas. Thus, it is desirable that the ALFF and the rsFC methods are used together.

In this study, we hypothesized that long-term SD disturbs spontaneous default activity and connectivity pattern. For a more comprehensive understanding of the effect from long-term total SD on DMN, we examined the statistical differences between long-term total SD condition and rested wakefulness (RW) condition, both in regional spontaneous activities using the ALFF method from regional perspectives and in default connectivity pattern using the rsFC method from integrative perspectives.

Materials and methods Subjects

The sleeping habit and sleep quality of all participants were monitored throughout 1-week duration by wearing a Fitbit Flex tracker (http://help.fitbit.com/), and only those whose data indicated habitual good sleep were recruited for this study. Twelve healthy male participants, aged 20–32 years (mean: 24.83 \pm 2.88 years) with an education duration of 15.6 \pm 1.5 years and who were university students responding to a Web-based questionnaire, were recruited. All subjects met the following criteria as in our previous studies^{3,31}: 1) had no symptoms associated with sleep disorders and no history of any psychiatric or neurologic disorders; 2) right-handed; 3) had a good sleeping habit, ie, >6.5 hours of sleep per night, falling asleep no later than 1.00 am and waking up no later than 9.00 am; 4) had a good sleep onset and/or maintenance, and has no history of swing shift, shift work, or sleep complaints; 5) had a regular dietary habit and did not consume any stimulants, alcohol, tea, cigarette, medications, and caffeine for at least 3 months before the study; 6) had no foreign implants in the body and no inborn or other acquired diseases; 7) moderate body shape and weight; and 8) Pittsburgh Sleep Quality Index score <5, and Hamilton Depression Rating Scale and Hamilton Anxiety Rating Scale scores <7. The protocols conformed to the Declaration of Helsinki and were approved by the First Affiliated Hospital of Nanchang University. All participants participated voluntarily and gave their written informed consent.

Sleep deprivation

The experiment was conducted in a closed narrow room, which was about 8 m² large and was equipped with monitors and ventilation. Two doors were used to isolate the room and the supervisory room. The researchers did not interact directly with the participants during the SD experiment. Thus, food and other necessities were passed through the doors. The SD was from 9.00 am of the first day to 9.00 am on the third day. During the 72 hours of total SD, the participants were required to stay awake all the time. The staffs of the research team took charge of monitoring in turns through video, preventing participants from falling asleep. If there were any signs of falling asleep, they were awakened by an alarm clock immediately.

Research design and process

Each of the subjects underwent rfMRI scans twice at 9:15 am; one followed RW and the other followed 72 hours of total SD. During the rfMRI scans, the subjects were asked to look at the black screen in the MRI scanner, where a "+" still stay on the screen. A video monitor was installed to ensure that the participants were awake with their eyes open during the rfMRI scans. A simple questionnaire was administered immediately to inquire whether the subjects were awake after the scans. Before the scans, all subjects underwent the attention network test (ANT).^{32,33}

Image acquisition

MRI scanning was performed on a 3-Tesla MRI scanner (Trio; Siemens, Erlangen, Germany). Each subject underwent a 7-minute fMRI scan immediately after T1-weighted images were obtained. High-resolution T1-weighted images were acquired with a three-dimensional spoiled gradient-recalled sequence in a sagittal orientation: 176 images (repetition time =2,530 ms, echo time =3.39 ms, thickness =1.33 mm, gap =0 mm, matrix =256×256, field of view =240 mm ×240 mm, flip angle =7°) were obtained. Finally, the 210 functional images (repetition time =2,000 ms, echo time =30 ms, thickness =3.5 mm, gap =0.7 mm, matrix =64×64, flip angle =90°, field of view =220 mm ×220 mm, 33 axial slices with gradient-recalled echo-planar imaging pulse sequence) covering the whole brain were obtained.

Data analysis

Behavioral data analysis

On the basis of the ANT, the behavioral accuracy rate was measured using corrected recognition; the reaction time (RT) was determined using only trials with correct responses (correct RT), and the lapse rate was measured using missing recognition. The intraindividual coefficient of variation was calculated for each participant by dividing the standard deviation of the correct RT by the mean correct RT.

fMRI data analysis

Data preprocessing

Functional data were checked by MRIcro software (www.MRIcro.com) to exclude defective data. The first ten time points of the functional images were discarded due to the possible instability of the initial MRI signal and the participants' adaptation to the scanning environment. Using MATLAB2010a (Mathworks, Natick, MA, USA), the remainder of the data preprocessing was performed by DPARSFA (http://rfmri.org/DPARSF) software, including Digital Imaging and Communications in Medicine standards for form transformation, slice timing, head motion correction, spatial normalization, and smoothing with a Gaussian kernel of 6×6×6 mm³ full-width at half-maximum. Motion time courses were obtained by estimating the values for translation (millimeters) and rotation (degrees) for each subject. Participants who had >2.0-mm maximum displacement in the x, y, or z directions and 2.0° of angular motion during the whole rfMRI scans were rejected. The Friston six head motion parameters were used to regress out head motion effects based on recent work showing that higher-order models were more effective in removing head motion effects.34,35 Linear regression was also applied to remove other sources of spurious covariates along with their temporal derivatives, including the signal from a ventricular region of interest (ROI) and the signal from a region centered in the white matter.36 Of note, the global signal was not regressed out in the present data, as in our previous study,³¹ for the reason that it is still controversial to remove the global signal in the preprocessing of resting-state data.^{36,37} After head motion correction, the fMRI images were spatially normalized to the Montreal Neurological Institute space using the standard echo-planar imaging template and resampling the images at a resolution of 3 mm \times 3 mm \times 3 mm.

Independent component analysis

After data preprocessing, the smoothed images of each subject were inserted into an overall independent component analysis (ICA) to decompose fMRI data into spatially independent component (IC) patterns and time courses by using the group ICA of fMRI toolbox (GIFT).³⁸ The fMRI data were split into a set of spatially independent functional networks (components). Each component was presented as a spatial map along with an associated time course. The optimal number of ICs was estimated using a modified minimum description length algorithm,³⁹ and this number was found to be 22 in this study. According to the templates provided by GIFT, the corresponding ICs of the DMNs for each subject were individualized and selected. The details of ICA analysis were reported in a previous study.⁴⁰

ALFF differences

After preprocessing, the time series for each voxel were temporally bandpass filtered (0.01–0.08 Hz) and linearly detrended to reduce low-frequency drift and physiological high-frequency respiratory and cardiac noise. The details of ALFF calculation were reported in a previous study.⁴¹ To reduce the global effects of variability across the participants, the ALFF of each voxel was divided by the global mean value for each participant.

rsFC differences

After pre-processing, temporal filtering settings were applied using a band-pass filtered (0.01~0.08 Hz) to reduce lowfrequency drift and physiological high frequency respiratory and cardiac noise. The influence of covariant (including head motion parameters, global mean signal, white matter signal and cerebrospinal fluid signal) should be eliminated. The different areas from ALFF method and ICA method were selected as ROIs for rsFC analysis. The average time series from the ROIs were extracted from the residual image. To make the data fit the normal distribution, we calculated the coefficient of Pearson correlation between ROIs and other voxels of whole brain, the coefficient was participated in Fisher's Z transformation With Z value represents function connection coefficient. To reduce the global effects of variability across the participants, the rsFC of each voxel was divided by the global mean value for each participant.

Receiver operating characteristic curve

Receiver operating characteristic (ROC) curve analysis is a widely accepted method for identifying and comparing the diagnostic accuracy of biomarkers. Our previous study found that the ALFF method showed high sensitivity and specificity and may be a useful noninvasive imaging tool and an early biomarker for the detection of cerebral changes of obstructive sleep apnea patients.⁴² In this study, the ROC curve was used to investigate whether these specific ALFF differences have the sensitivity and specificity to distinguish SD status from RW status.

Statistics

The behavioral deficits were evaluated using two paired *t*-tests with the IBM Statistical Package for the Social Sciences version 21.0 (SPSS 21.0), and the statistical threshold was set at P < 0.05.

For each selected component, individual maps were entered into one sample *t*-test (P < 0.0001 with a minimum continuous cluster volume, V, of 1,080 mm³, corrected by false discovery rate). To reveal the between-condition differences of regional activities and connectivity patterns, paired *t*-tests were performed with Statistical Parametric Mapping 8 (SPM8, http://www.fil.ion.ucl.ac.uk/spm), with age and years of education as nuisance covariates of no interest. A corrected significance level of individual voxel P < 0.005 and a continuous $V \ge 1,080$ mm³, using an AlphaSim-corrected cluster threshold of P < 0.05, were used to determine statistical significance.

Results Behavioral results

Compared with RW subjects, SD subjects showed a lower response accuracy rate (RW mean accuracy rate =96.83%, SD mean accuracy rate =77.67%; t=-5.123; P<0.001), a slower response (RW mean RT =695.92 ms; SD mean RT =799.18 ms; t=3.892; P=0.003), a significantly higher lapse rate (RW mean lapse rate =0.69%, SD mean lapse rate =19.29%; t=5.762; P<0.001), and a higher intraindividual coefficient of variability in RT (RW coefficient =0.26, SD coefficient =0.33; t=2.736; P=0.021). The details are presented in Figure 1.

DMN maps determined by group independent component analysis

Figure 2 shows the DMN maps in the RW group and in the SD group respectively, as determined by the group independent



Figure 1 ANT findings of RW subjects and SD subjects.

Notes: Compared with RW subjects, SD subjects showed (A) lower accuracy rate, and (B) higher RT, (C) lapse rates, and (D) intraindividual variability. Abbreviations: ANT, attention network test; RW, rested wakefulness; SD, sleep deprivation; RT, reaction time.

component analysis (GICA) approach when performing the one-sample *t*-test. The areas within the DMN in the SD group were significantly smaller than those in the RW group.

The GICA method showed that SD subjects, compared with RW subjects, displayed decreased rsFC between whole brain region and right IPL (BA40) or left PrC/PCC (BA30, 31). The details are presented in Figure 3 and Table 1. The two different areas were selected as ROIs, namely, the IPL1-FC and PrC/PCC-FC maps, respectively.

ALFF differences within the DMN

Compared with RW subjects, SD subjects showed lower ALFF area in the left IPL (angular gyrus, supramarginal gyrus; BA39, 40). To further understand the intrinsic DMN impairment, a different region was selected as ROI for the subsequent rsFC analysis, namely, the IPL2-FC map. The details are presented in Figure 4 and Table 1.

ROC curve

Since the lower ALFF area was found in the left IPL in the SD group relative to the RW group, the brain region might be utilized as a marker to distinguish the SD status from the RW status. To test this possibility, the mean ALFF signal value of the region was extracted. The ROC curve analysis showed that the area under the curve (AUC) value of the left IPL was 0.75, with a cutoff point of 0.834 (mean ALFF signal value). Further diagnostic analysis showed that the left IPL alone discriminated the SD status from the RW status with 75% sensitivity and 91.7% specificity. The details are presented in Figure 5.



Figure 2 The DMN maps in (A) the RW and (B) the SD groups respectively.

Notes: The covered DMN areas were smaller in the SD group than in the RW group. These maps are the results of the within-group comparisons by the one-sample t-test with the GICA method, corrected by FDR.

Abbreviations: DMN, default-mode network; RW, rested wakefulness; SD, sleep deprivation; GICA, group independent component analysis; FDR, false discovery rate; L, left; R, right.

Differences in rsFC

The IPL1-FC map demonstrated that the right IPL showed decreased rsFC with the left Prc (BA7), in addition to increased rsFC with left fusiform gyrus (BA37) and left cluster of middle temporal gyrus and inferior temporal gyrus (BA37). The details are presented in Figure 3 and Table 1. No connectivity difference was found in the PrC/PCC-FC map. The IPL2-FC map

showed that the left IPL had decreased rsFC with right cluster of IPL and superior temporal gyrus (BA39, 40). The details are presented in Figure 6 and Table 1.

Discussion

Our previous study had demonstrated that a few brain areas had obvious sex differences in both RW and SD subjects.³



Figure 3 The rsFC differences between the SD group and the RW group.

Notes: (A) The three maps represent the results of the ICA method, which showed decreased rsFC in right IPL and left PrC/PCC in the SD group relative to the RW group. (B) The three maps reflected the result of the IPL1-FC map, which showed decreased rsFC between the right IPL and left Prc and increased rsFC between the right IPL and left fusiform gyrus and left middle/inferior temporal gyrus.

Abbreviations: rsFC, resting-state functional connectivity; SD, sleep deprivation; RW, rested wakefulness; ICA, independent component analysis; IPL, inferior parietal lobule; PrC, precuneus; PCC, posterior cingulate cortex; L, left; R, right.

To avoid the influence of sex differences, only healthy males were recruited in this study. This study demonstrated that both rsFC method and ALFF method could reflect the intrinsic DMN impairment, and both showed equivalent results. SD resulted in a decline in response accuracy rate, as well as significant increase in RT and lapse rate. Furthermore, compared with RW subjects, SD subjects showed significantly higher value of intraindividual coefficient of variability of RT. Sleep debt had been shown to reportedly augment emotional instability (eg, anxiety and confusion), dilatation of the pupils, sleepiness, and feeling of fatigue, receding psychomotor performance, and changes in heart rate and blood pressure.^{43–49} In this study, SD significantly reduced all the behavioral measures and increased the risk of the variability, because of which SD may increase emotional instability and have a tendency to increase the risk of developing a disease.

ROC curve has been previously used to discriminate disease from health in subjects.^{42,50} The discrimination result is considered excellent for AUC values between

Methods	Conditions	Brain regions of peak coordinates	R/L	BA	V (mm³)	t-score of peak voxel	MNI coordinates of peak voxel (x, y, z)
ICA	SD <rw< td=""><td>PrC/PCC</td><td>L</td><td>31</td><td>1,890</td><td>-4.92</td><td>-12, -57, 30</td></rw<>	PrC/PCC	L	31	1,890	-4.92	-12, -57, 30
		IPL	R	40	1,755	-5.94	51, -42, 27
ALFF	SD <rw< td=""><td>IPL (angular gyrus, supramarginal gyrus)</td><td>L</td><td>39, 40</td><td>2,592</td><td>-6.81</td><td>-48, -57, 39</td></rw<>	IPL (angular gyrus, supramarginal gyrus)	L	39, 40	2,592	-6.81	-48, -57, 39
IPLI-FC	SD <rw< td=""><td>Prc</td><td>L</td><td>7</td><td>1,188</td><td>-4.3 I</td><td>-12, -78, 33</td></rw<>	Prc	L	7	1,188	-4.3 I	-12, -78 , 33
	SD>RW	Fusiform gyrus	L	37	86	5.99	-42, -63, -18
	SD>RW	Middle temporal gyrus, inferior temporal gyrus	L	37	43	7.10	-60, -60, -9
IPL2-FC	SD <rw< td=""><td>IPL, superior temporal gyrus</td><td>R</td><td>39, 40</td><td>2,484</td><td>-12.12</td><td>60, -57, 18</td></rw<>	IPL, superior temporal gyrus	R	39, 40	2,484	-12.12	60, -57, 18

Table I Two-sample paired t-test differences within the DMN areas between SD subjects and RW subjects

Notes: The between-condition statistical threshold was set at cluster size with P < 0.05, voxel with P < 0.050 ($|t| \ge 4.437$), and $V \ge 1,080$ mm³, corrected for AlphaSim. **Abbreviations:** DMN, default-mode network; SD, sleep deprivation; RW, rested wakefulness; R, right; L, left; BA, Brodmann's area; V, volume; MNI, Montreal Neurological Institute; ICA, independent component analysis; PrC, precuneus; PCC, posterior cingulate cortex; IPL, inferior parietal lobule; ALFF, amplitude of low-frequency fluctuation; FC, functional connectivity.



Figure 4 The ALFF differences within the DMN between the SD group and the RW group.

Notes: (A) Compared with RW group, SD group displayed lower ALFF area in the left IPL. (B) The right stapeldiagram presents the mean ALFF signal value of the left IPL area. The mean ALFF signal value of the left IPL was 0.90 ± 0.57 in the RW group, and that of the SD group was 0.44 ± 0.44 , and the difference was statistically significant (P=0.04).

Abbreviations: ALFF, amplitude of low-frequency fluctuation; DMN, default-mode network; RW, rested wakefulness; SD, sleep deprivation; IPL, inferior parietal lobule.



Figure 5 ROC curve of the mean ALFF value of the left IPL area. Notes: The arrow represents the best cutoff point. The ALFF value represents the mean ALFF value of the corresponding time point. Abbreviations: ROC, receiver operating characteristic; ALFF, amplitude of low-frequency fluctuation; IPL, inferior parietal lobule.

0.9 and 1.0, good for values between 0.8 and 0.9, fair for values between 0.7 and 0.8, poor for AUC values between 0.6 and 0.7, and failed for AUC values between 0.5 and 0.6.⁵¹ In this study, ROC curve analysis revealed that the mean ALFF signal decline value in the left IPL had a fair AUC value (AUC =0.75). Further diagnostic analysis exhibited that it alone discriminated SD subjects from RW subjects, with 75% sensitivity and 91.7% specificity. In short, our findings suggest that the ALFF method might be an effective measurement method of the spontaneous neuronal activity in brain regions associated with the low-frequency BOLD fluctuations obtained by rfMRI, suggesting that the IPL might be applied as a trait-related node to identify SD status from RW status.

Interestingly, the GICA method showed that SD subjects, compared with RW subjects, displayed decreased rsFC between whole brain region and the right IPL or the left PrC/PCC. Furthermore, the right IPL area showed decreased rsFC with left Prc, heterolateral IPL, and small area in temporal lobe, as well as increased rsFC with left temporal lobe. These findings support intrinsic dysfunctional connectivity of the DMN, as shown by previous deep sleep and partial SD studies.^{23,24,52} The IPL region is obligatorily or unintentionally engaged in the recall, consolidation, and retrieval of episodic memory information,^{9,53,54} as well as being implicated in diverse cognitive operations⁵⁵ that include bodily

awareness,⁵⁶ generating a sense of personal responsibility,⁵⁷ and moral decision making.⁵⁸ In this study, the IPL, as a core node, showed abnormal spontaneous activity and dys-functional connectivity. We therefore speculate that the IPL might be an early and important hub node of default brain impairment and might be associated with abnormal emotional and cognitive symptoms.

PrC is a deactivated area with the highest metabolic rate during the resting state.9 Many diseases, such as obstructive sleep apnea,⁵⁹ hyperactivity disorder,⁶⁰ and amnestic type mild cognitive impairment,⁶¹ can lead to PrC impairment with the adjacent PCC being involved. The abnormally high metabolic rate and the distribution of arterial anatomy could explain the susceptibility of the PrC/PCC. Previous studies found that SD increased the RT and lapse rate, which was attributed to the less-efficient communication among brain areas^{62,63} and to the changes in activity in the cingulate cortex.^{20,64} Sämann et al^{24,65} found that the PCC was associated with the regulation of consciousness, and the reduced integration of the PrC/PCC may be either an indicator of reduced awareness during wakefulness with increased sleep propensity or an indicator of sleep propensity itself. These findings suggest that the PrC/PCC may be involved in the worsening of behavioral measures (eg, RT, lapse rate) and sleep propensity by decreasing the rsFC among brain areas.

Conclusion

In this study, we demonstrated that long-term SD resulted in the intrinsic DMN dysfunction from both regional perspectives and integrative perspectives and that the DMN may be involved in the mechanism of cerebral and behavioral changes by decreasing efficient communication among brain areas. Furthermore, the IPL region might be an early and important default node of cerebral changes during SD status. Our findings provide insights into the pathophysiological mechanism of SD and may be helpful for understanding the pathophysiology of long-term SD and the development of symbolic early-imaging biomarkers for the detection of cerebral changes.⁶⁶

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Figure 6 The IPL2-FC differences between SD and RW groups.

Notes: The color in the map represents the differences. The left IPL showed decreased rsFC with the right cluster of IPL and superior temporal gyrus (*P*<0.005, corrected for AlphaSim).

Abbreviations: IPL, inferior parietal lobule; FC, functional connectivity; SD, sleep deprivation; RW, rested wakefulness; rsFC, resting-state functional connectivity.

Disclosure

The authors report no conflicts of interest in this work.

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