Self-dependent neural variability predicts recovery from depressive symptoms

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Abstract

Researchers have increasingly paid attention to the neural dynamics of depression. This study examined whether self-dependent neural variability predicts recovery from depressive symptoms. Sixty adults with depressive symptoms who were not officially diagnosed with major depressive disorder participated in this study. Participants completed functional magnetic resonance imaging (fMRI) scanning, including a resting-state and a self-reflection task. The fMRI data were used to estimate neural variability, which refers to the temporal variability in regional functional connectivity patterns. Participants then completed the Self-Construal Scale and the Beck Depression Inventory (BDI). The change in BDI scores over 3 months indicated the degree of recovery from depressive symptoms. Self-construal moderated the effects of general neural variability on predicting recovery from depressive symptoms. Interdependent individuals became less depressive with higher general neural variability, but the relationship was not significant in independent individuals. The differences in neural variability between self-related and other-related conditions also predicted recovery from depressive symptoms. The regions contributing to the prediction were mainly distributed in the default-mode network. Based on these results, the harmony between individuals’ neural dynamics and self-concept is important for recovery from depressive symptoms, which might be a foundation for individualized treatment and counseling.

Key words: self; fMRI; neural variability; recovery; depressive symptoms

Introduction

Depression has affected over 264 million people worldwide (GBD 2017 Disease and Injury Incidence and Prevalence Collaborators, 2018). Based on studies examining the neurological mechanisms that underlie depression, various treatments were proposed to intervene in major depressive disorder (MDD), such as medications and cognitive therapies (Beck, 1979; van der Velden et al., 2015; Trivedi et al., 2016; Cohen and DeRubeis, 2018; Tozzi et al., 2019). However, studies also found that only approximately one-quarter of individuals with temporary depressive symptoms develop chronic depression, whereas others recover without treatment (DeRubeis et al., 2008; Houle et al., 2013; van Grieken et al., 2015). To date, the process of recovering from depressive symptoms without treatment has rarely been studied.

Based on accumulating evidence, self-related processing is crucial to depression. MDD is associated with an increased
self-focus by which one engages in self-referential processing (Derry and Kuiper, 1981; Watkins and Teasdale, 2004; Grimm et al., 2009; Nejad et al., 2013). Negative self-referential bias predicts a deterioration of depressive symptoms (Disner et al., 2017; LeMoult et al., 2017). Self-related processing was found to be associated with specific neural activity in midline regions of the brain, such as the default-mode network (DMN), the anterior cingulate cortex (ACC) and the medial prefrontal cortex (mPFC) (Northoff and Bermpohl, 2004; Northoff et al., 2006; Qin et al., 2013, 2016). Consistent with behavioral findings, individuals with MDD show abnormalities in the DMN originating from global sources, both within- and intra-DMN sources (Sheline et al., 2009; Scalabrini et al., 2020). Abnormally increased self-focus in patients with MDD is related to neural activity in subcortical–cortical midline structures (Grimm et al., 2009). Both behavioral and neural evidence show that self-related processing plays an important role in depressive symptoms.

Moreover, social relationships and cultural circumstances are associated with depression (Cohen, 1996). The social network (Cohen et al., 1995) and self-focused processing (Yim et al., 2010) are associated with depression. In addition, self-focused processing is related to depression (Bae (1999) found that self-focused processing is related to depression. Bae (1999) reported a positive correlation between interdependent self-focused processing and depressive symptoms, and a negative correlation between independent self-focused processing and depressive symptoms. In addition, Lam (2005) found that interdependent self-focused processing indirectly affects depression by increasing family cohesion, which enhances self-esteem among Vietnamese-American adolescents. Therefore, self-focused processing may modulate the process of depression.

Neural studies have found that brain networks, including the DMN, and the salience and emotion network (SEN), are closely related to depression (Yeo et al., 2011; Wang et al., 2012; Kaiser et al., 2015; Jacobs et al., 2016). Neural dynamics have recently received increased attention in psychiatric and neurological disorders. The temporal variability in regions of the DMN differs between patients with distinct mental disorders (Zhang et al., 2016). Patients with MDD present higher temporal variability in the right inferior frontal gyrus, the left inferior occipital gyrus, the bilateral fusiform gyri and the left Heschl's gyrus than healthy controls (Hou et al., 2018). Researchers have also suggested that temporal variability might be a promising indicator for individualized therapy of MDD (Hou et al., 2018). Based on these findings, researchers speculated that temporal variability might be an indicator of brain flexibility and adaptability or a predictor of neural rehabilitation (Zhang et al., 2016), which in turn might affect recovery from depression.

This study aimed to investigate the neural dynamics of recovery from depressive symptoms. We hypothesized that self-dependent neural variability predicts recovery from depressive symptoms. Brain regions associated with self-related processing, including the DMN, contribute to the prediction.

Materials and methods

Participants

Two hundred thirty-nine individuals participated in depressive symptom screening. These participants were asked to complete the Beck Depression Inventory-II (BDI-II; Beck et al., 1996). The BDI included 21 sets of statements with four statements in each set. Sixty individuals (42 males and 18 females; age = 20.11 ± 2.33 years) participated in the functional magnetic resonance imaging (fMRI) studies. Among the 60 individuals, the 28 participants with a BDI score higher than 14 (range: 14–28, mean ± s.d. = 19.21 ± 2.19) were assigned to the high-depression group and the other 32 participants with a BDI score lower than 4 (range: 0–4, mean ± s.d. = 2.25 ± 1.44) were assigned to the low-depression group. The participants completed the BDI again 3 months after the first screen. Recovery from depressive symptoms was assessed by subtracting the BDI score recorded at the second screen from the BDI score reported at the first screen. Higher scores indicated that the participants had become less depressive. None of the participants took psychotropic medicine or treatment between the two screens. Written informed consent was obtained from all participants before starting the experiments. All studies were approved by the ethics committee of the Department of Psychology at Sun Yat-sen University.

Procedure and stimuli

During fMRI scanning, the participants completed a 5-min resting-state and a self-reflection task (Figure 1). During the task, the participants were asked to judge whether a particular item described themselves (self-related conditions) or a public figure (other-related conditions; i.e. Liu Xiang, a famous athlete). The items were divided into three categories: mental attributes (i.e. personality characteristics), physical attributes (i.e. physical appearances) and social attributes (i.e. social identities). Each category contained 48 items. A font judgment (bold vs light) was used as the control condition. Six scans with seven blocks in each scan were performed. Each block presented a type of judgment. The blocks were presented in a random order and with a 10-s interval. Each item was presented for 2 s and followed by a 1-s fixation.

After scanning, participants completed the Self-Construal Scale (Singelis, 1994). The Self-Construal Scale was divided into two dimensions: the interdependent subscale and the independent subscale. Each subscale included 12 items that were rated on a 7-point Likert scale (1 = strongly disagree and 7 = strongly agree). Interdependence was assessed by subtracting the mean score for the independent subscale from the mean score for the interdependent subscale (Figure 1). Higher scores indicated greater interdependence on social contexts and others.

Imaging acquisition and preprocessing

We used a GE Signa MR750 3.0T scanner with a standard head coil to acquire the fMRI data. The data were acquired using T2-weighted, gradient-echo, echo-planar imaging (EPI) sequences with the following parameters: repetition time (TR) = 2000 ms, echo time (TE) = 30 ms, flip angle (FA) = 90°, field of view (FOV) = 240 × 240 mm, matrix = 64 × 64 × 32 and spatial resolution = 3.75 × 3.75 × 5 mm3. During resting-state scanning, the participants were asked to keep their eyes open.
A standard preprocessing procedure was performed on the fMRI data using the Data Processing Assistant for Resting-State fMRI (DPARSF) toolbox (Yan and Zang, 2010). The data from the first five volumes were removed. The remaining data underwent slice timing and realignment to correct the time delay of scans and head motion. None of the participants was excluded during realignment because their maximum head motion was all within the criterion of 3.0 mm and 3.0°. The corrected data were registered to the Montreal Neurological Institute (MNI) space with the EPI template. The normalized data were Gaussian smoothed [full-width at half-maximum (FWHM) = 4 mm], detrended, and bandpass filtered (0.01–0.08 Hz). Nuisance covariates were removed by multiple regressions, including six rigid-body head motion parameters and the mean time courses of the white matter and cerebrospinal fluid. In addition, we used different nuisance covariate removal strategies (removal of the global signal vs no removal) to test the robustness of the prediction effects across nuisance covariate removal strategies (see supplementary results).

A 264-node atlas (Power et al., 2011) was used to define nodes and networks. The atlas included the cerebral cortex, subcortical structures and cerebellum. The nodes were 264 spheres (diameter = 10 mm) divided into 14 networks (see supplementary methods).

Estimation of neural variability

We used the method proposed by Zhang et al. (2016) to estimate neural variability. The neural variability of each node was defined as the temporal variability of functional connectivity in the node.

Estimation of general neural variability. We used the fMRI data collected during the resting state to estimate general neural variability. Time series were extracted and split into nine non-overlapping windows with a length of 30 s. In space, we calculated the functional connectivity matrix within each window using Pearson’s correlation analysis. Each row (or each column) of the matrix represented the functional connectivity between a specific node and the remaining nodes across the whole brain. For each node, we compared the functional connectivity matrices across different windows of time using Pearson’s correlation analysis. General neural variability of the node was calculated by subtracting the mean of correlation coefficients from one. The following formula was used:

\[ V_k = 1 - \text{corrcoef}(F_{ik}, F_{jk}), i, j = 1, 2, 3, \ldots, 9, i \neq j, \]

where \( k \) indicated the node and \( i \) and \( j \) indicated windows.

Estimation of the difference in neural variability between the self-related conditions and other-related conditions. We used the fMRI data collected during the self-reflection task to estimate the difference in neural variability between the self-related conditions and other-related conditions. The neural variability of each condition was calculated by comparing regional functional connectivity matrices across different scans under the same condition. The procedure was similar to the estimation of general neural variability. The difference in neural variability was calculated by subtracting the mean neural variability in the three other-related conditions from the mean neural variability in the three self-related conditions.

Prediction analysis

We used the leave-one-out cross-validation method to study whether self-dependent neural variability predicted recovery from depressive symptoms in a novel individual (Figure 1). In each iteration, one participant was excluded as the test set and the remaining participants were the training set. The training set established the predictive models, and the test set evaluated the predictive models. Because each of the 60 participants was excluded once, 60 iterations were performed.

Self-construal-dependent neural variability predictive models. Before the prediction analysis, the dot product of normalized general neural variability and normalized interdependence was defined as the interaction between general neural variability and interdependence. We used the interaction between general neural variability and interdependence to study whether self-construal moderated the effects of general neural variability on predicting the recovery from depressive symptoms in a novel individual.
In the feature-selection stage, based on the training set, we performed Pearson’s correlation analysis between the interactions in each node and recovery scores (feature-selection threshold = 0.05). The nodes whose interactions were significantly positively correlated with recovery were selected as positive features, and the significantly negatively correlated nodes were negative features. In addition, we used different feature-selection thresholds (thresholds = 0.05 and 0.01) to test the robustness of the prediction effects across feature-selection thresholds (see supplementary methods and results).

In the model establishment stage, positive features or the opposite number of negative features were averaged, resulting in the positive feature interaction strength or the negative feature interaction strength, respectively, and they were averaged together, resulting in the total feature interaction strength. Simple linear regression analyses were conducted to construct the relationships between the three feature interaction strengths and recovery, resulting in three models: the total model, positive model and negative model.

In the prediction stage, the same features as the training set were extracted from test set, and the three interaction strengths were substituted in the corresponding model, resulting in the predicted recovery for the participant in each of the three models.

In the model evaluation stage, the correlation coefficient between the predicted recovery calculated using the model and the observed recovery assessed using the scale was defined as the predictive power of the model. Only significant positive predictive power indicated that the prediction was successful. The nodes that were selected as features in more than 95% of the iterations were regarded as important nodes. We performed permutations tests to further confirm the significance of the predictions (see supplementary methods).

Self-referential-related neural variability predictive models. The procedure was similar to the prediction analyses of the interaction between general neural variability and interdependence except that the interaction was replaced with the differences in neural variability between self-related and other-related conditions. Moreover, for the domain-specific prediction effect, we used the differential neural variability between self-related conditions and other-related conditions in which participants made judgments about mental attributes, physical attributes or social attributes to perform the prediction analyses (see supplementary methods).

Results
Self-construal-dependent neural variability predictive models
The interaction between general neural variability and interdependence successfully predicted recovery from depressive symptoms in the total model ($r = 0.31$, $P = 0.016$, Figure 2A) and in the positive model ($r = 0.32$, $P = 0.012$). In the negative model, no feature was selected in at least one iteration. The prediction effects were robust across nuisance covariate removal strategies (no removal of the global signal vs removal) and feature selection thresholds (thresholds = 0.05, 0.01) (see supplementary results). However, general neural variability itself failed to predict recovery from depressive symptoms in the three models (total model: $r = 0.13$, $P = 0.323$; positive model: $r = 0.23$, $P = 0.070$; negative model: $r = -0.23$, $P = 0.081$). The results indicated that self-construal moderated the effects of general neural variability on predicting recovery from depressive symptoms in individuals. Permutation tests further confirmed that the observed predictive power significantly differed from the predictive power in the null distribution (total model: $P = 0.015$; positive model: $P = 0.009$, Figure 2B). Thirty-four important nodes were identified, all of which were positive features (Figure 2C). The important nodes were mainly distributed in the visual network ($n = 17$, Figure 2D), sensory–somatomotor network ($n = 6$) and DMN ($n = 5$).

We conducted a simple effects analysis with the mean general neural variability of the important nodes to examine the relationship between general neural variability and recovery from depressive symptoms in individuals with high interdependence ($M + s.d.$) and in individuals with low interdependence ($M − s.d.$). General neural variability was positively correlated with recovery from depressive symptoms ($b = 4.24$, $SE = 1.35$, $P = 0.003$, Figure 2E) in interdependent individuals, but was not significant in independent individuals ($b = −1.95$, $SE = 1.03$, $P = 0.063$). The simple effects were consistent on the high-depression group (interdependent individuals: $b = 5.58$, $SE = 2.13$, $P = 0.015$; independent individuals: $b = 0.12$, $SE = 1.41$, $P = 0.935$) and the low-depression group (interdependent individuals: $b = 3.10$, $SE = 0.028$, independent individuals: $b = −1.08$, $SE = 1.48$, $P = 0.471$). Based on the results, interdependent individuals became less depressive if they had higher general neural variability.

Self-referential-related neural variability predictive models
The differences in neural variability between the self-related conditions and other-related conditions during the self-reflection task successfully predicted recovery from depressive symptoms in the total model ($r = 0.33$, $P = 0.010$, Figure 3A) and in the negative model ($r = 0.32$, $P = 0.014$). In the positive model, no feature was selected in at least one iteration. The prediction effects were robust across feature selection thresholds (thresholds = 0.05, 0.01; see supplementary results). However, the differences in neural variability between the self-related conditions and font condition (total model: $r = 0.15$, $P = 0.255$; negative model: $r = 0.15$, $P = 0.255$; positive model: no feature) or the differences in neural variability between the other-related conditions and font condition (total model: $r = −0.19$, $P = 0.145$; negative model: $r = −0.15$, $P = 0.244$; positive model: $r = 0.08$, $P = 0.525$) failed to predict recovery from depressive symptoms. Thus, the difference between self-related neural variability and other-related neural variability predicted recovery from depressive symptoms in individuals. Permutation tests further confirmed that the observed predictive power significantly differed from the predictive power in the null distribution (total model: $P = 0.012$; negative model: $P = 0.004$, Figure 3B). Twenty-seven important nodes were identified, among which 26 nodes were negative features and one node was a positive feature (Figure 3C). The important nodes were mainly distributed in the DMN ($n = 10$, Figure 3D) and visual network ($n = 7$).

Moreover, the difference in neural variability between the self-related conditions and other-related conditions in which participants made judgments about mental attributes marginally significantly predicted recovery from depressive symptoms in the total model ($r = 0.25$, $P = 0.056$, Figure 4A) and in the negative model ($r = 0.25$, $P = 0.055$). The prediction
Fig. 2. Results of the self-construal-dependent neural variability predictive models. (a) The predictive power (correlation between the predicted recovery and the observed recovery) of the total model (left panel, dark gray) and positive model (right panel, red). (b) The results of permutation tests in the total model (left panel, dark gray) and positive model (right panel, red). The black arrow indicates the observed predictive power in the two models. The light gray bar indicates the permutations in which no feature was selected in at least one iteration. (c) The locations of the important nodes (nodes that were selected as features in more than 95% of the iterations). (d) The distribution of the important nodes in the visual network (blue), sensory–somatomotor network (cyan), DMN (red), uncertain network (striped), subcortical network (brown), cingulo-opercular task control network (purple), dorsal attention network (green), frontoparietal task control network (yellow), salience network (black), auditory network (pink), hand ventral attention network (teal), mouth sensory–somatomotor network (orange), memory retrieval network (gray) and cerebellar network (pale blue). The dark colors indicate the important nodes, and the light colors indicate the remaining nodes. The networks were ranked by the number of the important nodes. (e) The simple effects on all participants (left panel), high-depression group (middle panel) and low-depression group (right panel).
effects were robust across feature selection thresholds (thresholds = 0.05, 0.01; see supplementary results). Thirty-four important nodes were identified, all of which were negative features (Figure 4B). The important nodes were mainly distributed in the DMN (n = 11, Figure 4C) and visual network (n = 11). However, the differences in neural variability in the physical condition or in the social condition failed to predict recovery from depressive symptoms (physical attribute—total model: \( r = -0.01, P = 0.967 \); negative model: \( r = -0.01, P = 0.923 \); positive model: no feature. Social attribute—total model: \( r = -0.17, P = 0.205 \); negative model: \( r = -0.03, P = 0.834 \); positive model: \( r = -0.10, P = 0.431 \)). The results indicated the importance of the mental aspect of the self.

We examined the robustness of the predictive models in the high-depression group and in the low-depression group using the important nodes as features. The differences in neural variability within the important nodes successfully predicted recovery from depressive symptoms in both the high-depression group \( (r = 0.61, P < 0.001, \text{Figure 3E}) \) and the low-depression group \( (r = 0.36, P = 0.044) \). In the mental condition, the prediction effect was significant in the high-depression group \( (r = 0.70, P < 0.001, \text{Figure 4D}) \), but not in the low-depression group \( (r = -0.06, P = 0.762) \).

**Discussion**

The present study provided the first evidence for the relationship between neural dynamics and mental dynamics (recovery from depression). Interdependent individuals with higher general neural variability in the resting state recovered better from depression, while independent individuals with higher neural variability showed no recovery. Interdependent individuals interpret their thoughts, feelings and actions in association with connectedness and integration with the external social environments, and independent individuals comprehend their thoughts, feelings and actions concerning separateness and uniqueness from within rather than reference to others (Markus and Kitayama, 1991). Interdependent individuals must consider and integrate more external information to link to various people and maintain this interconnectedness (Yeh and Hwang, 2000). In the process of adapting to various external environmental demands, the brain continues to modify its function and structure by strengthening, weakening, pruning or adding synaptic connections and by promoting neurogenesis (Faucal-Leone et al., 2005, 2011). This finding is consistent with our results mentioned above; neural variability may be an indicator of brain flexibility and adaptability (Zhang et al., 2016). Taken together, these results suggest that the harmony between self-construal and neural variability plays an important role in individuals’ mental health.

Moreover, the differences in neural variability between self-related conditions and other-related conditions successfully predicted recovery from depressive symptoms. A greater difference in neural variability between self-related conditions and other-related conditions was related to better recovery from depression symptoms. Analogously, previous research has emphasized the important role of self-related processing in depression, which has previously been reported in several studies (Sheline et al., 2009; Disner et al., 2017). This result is also consistent with a previous study showing that neural variability in the left pallidum is positively correlated with recovery from depressive symptoms (Hou et al., 2018). Neural variability in the left pallidum predicts implicit self-esteem (Izuma et al., 2018), which is closely related to depression (Battle, 1978; Orth et al., 2016; Rieger et al., 2016).

Furthermore, the DMN contributed to the prediction. The important nodes in our task-based predictive models were mainly distributed in the DMN. The DMN is involved in various self-related processes, including self-referential thoughts (Gusnard et al., 2001), internal thought (Andrews-Hanna, 2012),

![Figure 3. Results of the self-referential-related neural variability predictive models. (a) The predictive power of the total model (left panel, dark gray) and negative model (right panel, blue). (b) The results of permutation tests in the total model (left panel, dark gray) and negative model (right panel, blue). The black arrow indicates the observed predictive power in the two models. The light gray bar indicates the permutations in which no feature was selected in at least one iteration. (c) The locations of the important nodes. (d) The distribution of the important nodes. The colors of the networks are the same as those in Figure 2. The dark colors indicate the important nodes, and the light colors indicate the remaining nodes. The networks were ranked by the number of the important nodes. (e) The predictive power in the high-depression group (left panel) and low-depression group (right panel).](image-url)
Fig. 4. Results of the self-referential-related neural variability predictive models in the mental condition. (a) The predictive power of the total model (left panel, dark gray) and negative model (right panel, blue). (b) The locations of the important nodes. (c) The distribution of the important nodes. The colors of the networks are the same as those in Figure 2. The dark colors indicate the important nodes, and the light colors indicate the remaining nodes. The networks were ranked by the number of the important nodes. (d) The predictive power in the high-depression group (left panel) and low-depression group (right panel).
autobiographical memory and theory of mind (Spreng and Grady, 2010). Furthermore, several studies found that the DMN is involved in self-related processing in individuals with depression (Lemogne et al., 2009; Sheline et al., 2009; Grimm et al., 2011). For instance, negative blood-oxygen-level-dependent (BOLD) responses decreased in the DMN when patients with MDD made judgments on self-relatedness (Grimm et al., 2011). The present study extended these findings from a dynamic perspective by showing that self-related neural variability in the DMN contributes to recovery from depression.

Moreover, the prediction effects of self-related neural variability were significant only when participants made judgments about mental attributes. Thus, different aspects of self may not be related to depression in a similar manner. Patients with MDD exhibit increased neural activity in the DLPFC and dorsal MFG and functional connectivity between the DLPFC, MFG and dorsal anterior cingulate cortex (dACC) when they made judgments on personality characteristics describing themselves rather than general desirable characteristics (Lemogne et al., 2009). Meanwhile, these brain regions have been shown to correspond to important parts of the neural mechanisms underlying depression (Sheline et al., 2009; Kaiser et al., 2015; Jacobs et al., 2016). The neural variability in the mental aspect of self-related processing appears to be particularly important for recovery from depression.

We used the leave-one-out cross-validation method to study whether self-dependent neural variability predicted recovery from depressive symptoms. Leave-one-out cross-validation was evidenced to provide an almost unbiased estimate of the probability of test error (Cawley and Talbot, 2003) from limited data. The method enabled us to test whether the predictive models established using the training data were generalizable to a novel observation. The results supported the generalization of the prediction effects.

One limitation of our study was that all the participants were Chinese. As previously shown, interdependent self-construal is adaptive and dominant in Eastern Asian culture, but independent self-construal is adaptive and dominant in Western culture (Singelis, 1994). Danish participants show greater activation in the mPFC when making judgments of self as a public figure than Chinese participants, regardless of the attribute dimensions for the judgments, while greater activity in the temporoparietal junction (TPJ) was induced in Chinese participants than in Danish participants when making self-judgments of social attributes (Ma et al., 2014). Researchers have suggested that people in different social and cultural contexts might adopt different self-reflection strategies by changing the neural activity of the mPFC and TPJ in social brain networks (Ma et al., 2014). The present study illustrated that only interdependent Chinese participants became less depressive with higher general neural variability. Based on previous research, this relationship might be different in Western culture. Further studies should explore whether self-dependent neural variability remains a predictor of recovery from depressive symptoms in samples from a Western culture where independent self-construal is dominant. Moreover, future research should also examine the neural dynamics of the process from depressive symptom onset to the diagnosis of depression to test whether self-dependent neural variability also adequately predicts the development of depression.

Conclusions

Taken together, our findings suggest that self-dependent neural variability predicts recovery from depressive symptoms in individuals. The effects of general neural variability on predicting recovery from depressive symptoms may be moderated by self-construal. Interdependent individuals with higher general neural variability in the resting state recover better from depressive symptoms. The difference between self-related neural variability and other-related neural variability predicted the recovery from depressive symptoms, especially in the mental domain. Our results provide the inspiration to understand and improve individuals’ mental health, which might be a foundation for individualized treatment and counseling in the future.

Funding

This work was supported by the National Natural Science Foundation of China [grant numbers 32071081 and 31800916] and the Guangdong Basic and Applied Basic Research Foundation [grant number 2020A1515010975].

Conflict of interest

The authors have no conflicts of interest to declare.

Authors’ contributions

L.S.Y. designed the research; F.L.Y. and L.S.Y. collected and analyzed the data; and F.L.Y., D.Q. and L.S.Y. wrote the manuscript. All authors commented on the manuscript.

Code availability

The custom computer code that was used in the main analysis of this study is available from the corresponding author.

Data availability

The de-identified data that support the main findings of this study are available from the corresponding author upon reasonable request.

Ethical approval

All experimental procedures conformed to the Declaration of Helsinki and were approved by the local Research Ethics Committee of the Department of Psychology, Sun Yat-sen University. Informed consent was obtained from participants prior to inclusion in the study.

Supplementary data

Supplementary data are available at SCAN online.

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