

Increased miR-132 level is associated with visual memory dysfunction in patients with depression

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Background: Impaired visual memory seems to be a core feature of depression, while increased microRNA-132 (miR-132) levels have been widely reported in depression patients. The authors aimed to explore the relationship between miR-132 changes and visual memory deficits in unmedicated patients with major depressive disorder (MDD).

Patients and methods: A total of 62 medication-free MDD patients and 73 matched healthy controls (HCs) were tested for miR-132 expression level in peripheral blood using quantitative real-time polymerase chain reaction. We used a computerized neurocognitive task from the Cambridge Neuropsychological Test Automated Battery (CANTAB) – pattern recognition memory (PRM) task – as a measurement of visual memory. The relationship between visual memory, miR-132 expression level, and clinical symptoms was explored in patients with MDD.

Results: Upregulated miR-132 expression levels were seen in MDD patients but not in HCs. Two-sample *t*-tests showed that MDD patients had decreased visual memory, mainly memory delayed compared to that of HCs. Correlation analyses revealed that in MDD patients, increased miR-132 expression levels were significantly correlated with visual memory as measured by the CANTABPRM. Hamilton Rating Scale for Anxiety scores were negatively correlated with PRM – number correct (immediate) and PRM – percent correct (immediate).

Limitations: The main limitations were missing data and lack of follow-up studies.

Conclusion: Our study suggests that increased miR-132 expression levels were associated with visual memory deficits, which may underlie the pathophysiology of MDD. In individuals with depression, immediate visual memory defects were positively correlated with anxiety symptoms.

Keywords: major depressive disorder, miR-132, cognition, memory, CANTAB

Introduction

Depressive disorders, affecting 121 million people worldwide, are a major public health concern. The prevalence of depression is known to be between 5% and 19%.¹ Regarding course of recovery, major depressive disorder (MDD) was historically seen as an episodic disorder, but recent findings have indicated that 50% of depressed patients experience a relapse within 2 years after their first episode and 80% experience more than one depressive episode during their life course.² Although depression traditionally is seen as affective disorder in nature, research conducted during the last decades has shown that depression is associated with important disturbance in cognitive functioning, such as memory loss, attention deficit, and executive function impairment.^{3,4} Among them, memory decline was reported by most patients with depression, which seriously affected their social functioning and quality of life.³ Numerous studies showed that visual memory impairment is one of the important features of memory defects in depression patients (DPs).^{5–7}

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Despite considerable efforts to underpin the molecular and cellular changes associated with depressive states, a global understanding of the pathophysiology of MDD is still lacking. Recently, some studies particularly focused on the interplay between genetic and environmental factors. Through epigenetic mechanisms, environmental factors can induce changes in gene expression levels that might mediate the onset of a disease without altering the DNA sequence.^{8,9} These mechanisms include histone modification, DNA methylation, and post-transcriptional regulation by non-coding RNAs such as microRNAs (miRNAs). It is now generally accepted that these epigenetic factors may mediate susceptibility to disorders like MDD.¹⁰ miRNAs are short (~22 nucleotides) RNA molecules that negatively regulate gene expression at the post-transcriptional level¹¹ and trigger mRNA degradation and/or translational repression in turn.^{12,13} Each miRNA may target several mRNAs, often in specific locations on their 3'-untranslated region (3'-UTR). More than 1,000 reported human miRNAs (miRbase database¹⁴) are estimated to control the expression of more than half of human genes.¹⁵ Within the central nervous system (CNS), a recent work has indicated that miRNAs play key roles in a wide range of physiological processes, including neurogenesis, neuronal proliferation, and synaptic plasticity.⁵ The disruption of miRNA expression through genetic deletion of Dicer results in aberrant neuronal development and differentiation, as well as altered morphogenesis and neuronal signaling.⁵ For example, Konopka et al¹⁶ demonstrated that conditional deletion of Dicer1 enhanced learning and memory capacity in mice.

A growing number of studies on psychiatric illness have focused on miRNAs in peripheral blood.^{17–20} First, the CNS may exert its influence on the gene expression of peripheral lymphocytes via cytokines, neurotransmitters, or hormones,⁵ which may explain the comparable gene expression levels between the peripheral blood and some CNS tissues.⁵ Meanwhile, this also provides the explanation of alterations in the dopamine transporter in the lymphocytes of psychotic patients.⁵ Second, the peripheral expression levels of some dysregulated genes were also found to be correlated with certain clinical symptoms and neurocognitive functions in patients with depression.⁵ In fact, blood-based miRNAs are easier to access and less intrusive than brain tissues, which is becoming an increasingly important approach to identify clinically applicable biomarkers for MDD.

Altered expressions of microRNA-132 (miR-132) are associated with a series of neurological disorders, such as Alzheimer's disease,²¹ Huntington disease,^{22,23} drug

addiction,²⁴ and schizophrenia.⁵ Previous studies have also shown that miR-132 is related to proteins or genes, which are important for depression.²⁵ For example, a study revealed that miR-132 induced by light in the suprachiasmatic nucleus regulates the expression of period genes, which may play a role in circadian rhythms.²⁶ We know that impaired neurogenesis and alteration of circadian rhythms as potential dysregulated pathways may contribute to depressive disorders. Another study revealed that miR-132 targets p250GAP, a Rac1 GTPase-activating protein.²⁷ By repressing p250GAP levels, miR-132 increases the activity of the actin depolymerizing protein n-cofilin via Rac1-PAK signaling.²⁸ This modulation of actin turnover in dendritic spines has been shown to regulate both spine size and spine density.^{29,30} Inhibition of miR-132 in primary hippocampal cultures decreased spine density and size, whereas in vivo overexpression increased spine density. We know that memory function is closely related to the number and density of spine.

Despite these findings, the mechanism of cognitive function defects in MDD is not fully understood till now. In this study, we aimed to explore the possible association between peripheral miR-132 expression level and visual memory defects in patients with MDD.

Patients and methods

Participants and blood samples

Although we included 72 drug naïve patients and 74 healthy controls (HCs), 11 individuals failed to fully complete the CANTAB test for personal reasons and were excluded. The final sample size was 62 patients and 73 HCs. All participants were recruited from the Mental Health Center of West China Hospital of Sichuan University. All patients were independently diagnosed by two licensed psychiatrists according to Structured Clinical Interview for DSM-IV Axis I Disorders (SCID), thus excluding patients with concurrent other axis I disorders except major depression.³¹ All patients were assessed immediately after the baseline interview. Disease severity was evaluated using the Hamilton Rating Scale for Depression (HAMD; 17-item version, 1960) and Hamilton Rating Scale for Anxiety (HAMA; 14-item version, 1959). The exclusion criteria included age younger than 18 years or older than 60 years, severe medical diseases, mental retardation, structural brain disorders, movement disorders, and drug abuse. Meanwhile, demographically similar (age-, gender-, and education-matched) healthy volunteers were recruited via advertisements within the local community. The control subjects were also interviewed according to

the SCID, non-patient edition,³² to assure that no one had a current or past history of depression or other axis I disorders. The ethical committee of Sichuan University approved the study and the sample collection procedures. Written informed consent was obtained from all participants.

RNA isolation

Whole blood (3 mL) was collected from each participant using Tempus Blood RNA Tubes (Thermo Fisher Scientific, Waltham, MA, USA; PN: 4342792). Blood was mixed vigorously for at least 10 seconds after collection and immediately frozen at -80°C until use. Total RNAs were isolated using the MagMAX for Stabilized Blood Tubes RNA Isolation Kit (PN: 4451893; Ambion, Carlsbad, CA, USA) according to the manufacturer's protocol. The RNA quality was assayed using NanoDrop ND-2000 (Thermo Fisher Scientific), and integrity was assessed using Bioanalyzer 2100 (Agilent Technologies, Santa Clara, CA, USA).

cDNA synthesis and quantitative real-time polymerase chain reaction (qPCR)

Portions (100 ng) of each of the total isolated RNA preparations were reverse transcribed to cDNA using a TUREscript 1st Strand cDNA SYNTHESIS Kit (PC1802; Aidlab, Beijing, People's Republic of China) according to the manufacturer's instructions. The reaction mixture consisted of 2 μL 5 \times RT Reaction Mix, 0.8 μL gene special primer (GSP) (miR-132 reverse transcription primer), and 100 ng total RNA template in a total volume of 10 μL . Reverse transcription was performed in GeneAmp PCR System 9700 (Thermo Fisher Scientific) at 42°C for 60 min and 85°C for 5 min. The resulting reverse transcription reaction product was immediately stored at -20°C for further analysis. The qPCR reactions were done in real-time cyclers (qTOWER 2.2; Analytik Jena, Jena, Thuringia, Germany). The reaction mixture consisted of 5 μL 2 \times SYBR Green Supermix (Aidlab; PC3302), 200 nM miRNA specific primers, 200 nM universal primers, and 0.8 μL reverse transcription product; the volume was adjusted to 10 μL by H_2O . PCR reactions were initiated with 3-min incubation at 95°C , followed by 40 cycles at 95°C for 10 s and at 60°C for 60 s. A melting curve was performed at the end of the PCR run over a range of 66°C – 99°C , increasing the temperature stepwise by 0.5°C every 2 s. Experiments were performed, as described earlier, in triplicate (Table 1). The relative level of miR-132 was calculated with the comparative ΔC_t ($\Delta\Delta\text{C}_t$) method using small RNA U6 as a normalization control. Fold change was calculated using $2^{-\Delta\Delta\text{C}_t}$.

Table 1 Sequence of primers in RT-qPCR

Name	Sequence 5'-3'
U6-F	CTCGCTTCGGCAGCAC
U6-R	AACGCTTCACGAATTTGCGT
miR-132-RT	GTCGTATCCAGTGCAGGGTCCGAGG TATTCGACTGGATACGACCGACCA
miR-132-F	GCCCCTAACAGTCTACAGCCAT
miR-132-R	GCAGGGTCCGAGGTATTC

Abbreviations: RT-qPCR, reverse transcription quantitative polymerase chain reaction; miR-132-RT, microRNA-132 reverse transcription primer; miR-132-F, microRNA-132 forward primer; miR-132-R, microRNA-132 reverse primer.

Visual memory test

We used pattern recognition memory (PRM) in the Cambridge Neuropsychological Test Automated Battery (CANTAB) to assess visual memory processing.³³ CANTAB is a standard, computerized, nonlinguistic, and culturally blind test to assess cognitive dysfunction in various psychiatric disorders⁵ in experimental and clinical settings.³⁴

The PRM test is designed to measure the capacity of visual processing in a two-choice forced discrimination paradigm.³⁵ The participants were presented with a series of visual geometric patterns, one at a time, in the center of the screen. These patterns were designed so that they cannot be easily given verbal labels. In the recognition phase, the participants were required to choose the correct pattern between an already seen pattern and a novel pattern. In this phase, the test patterns were presented in the reverse order to the original order of presentation. This was then repeated with new patterns. The second recognition was given either immediately or after a delay. The immediate mode always ran ~ 20 min before the delayed mode.

Statistical analysis

All statistical analyses were performed using SPSS 21.0 software (IBM Corporation, Armonk, NY, USA) for Mac. We used two-sample *t*-test or chi-square test to compare the demographic data. Student's *t*-test was used to identify significant differences in the relative expression of miR-132 and PRM test scores, respectively, between the healthy participants and patients. The relationship between the expression of miR-132 and visual memory function was examined using Pearson's correlation coefficients. $P \leq 0.05$ was considered statistically significant for all tests.

Results

Demographic and clinical characteristics

No significant differences were observed between MDD patients and HC groups in the demographics, including age

Table 2 Comparison of general characteristics of patients with depression and HCs

Characteristics	MDD	HC	t	P-value
Gender (male/female)	22/40	21/52	–	0.460 ^a
Age (years)	28.35±8.54	27.67±7.80	–0.486	0.628 ^b
Years of education (years)	15.56±3.08	14.58±3.84	1.688	0.098 ^b
HAMD score	22.74±4.877	–	–	–
HAMA score	17.51±5.778	–	–	–

Notes: ^aP-value for gender distribution was obtained by chi-square test. ^bP-values were obtained by Student's t-test.

Abbreviations: MDD, major depressive disorder; HC, healthy control; HAMD, Hamilton Depression Rating Scale; HAMA, Hamilton Anxiety Rating Scale.

(MDD: 28.35±8.54; HC: 27.67±7.80; $t=-0.486$; $P=0.628$), gender (male/female; MDD: 22/40; HC: 21/52; $P=0.460$), or education years (MDD: 15.56±3.08; HC: 14.58±3.84; $t=1.688$; $P=0.098$). The mean severity of clinical symptoms in MDD patients was 22.74±4.877 in HAMD and 17.51±5.778 in HAMA (Table 2).

MicroR-132 expression level

We found that miR-132 expression levels were significantly upregulated in MDD patients compared with HCs ($t=-2.613$; $P=0.01$). The relative expression of miR-132 between patients and control samples is shown in Figure 1.

Visual memory task results

Altogether 62 patients and 73 HCs completed all the neurocognitive tests. Between the two groups, we found that MDD patients performed poorer than HCs in PRM – number correct

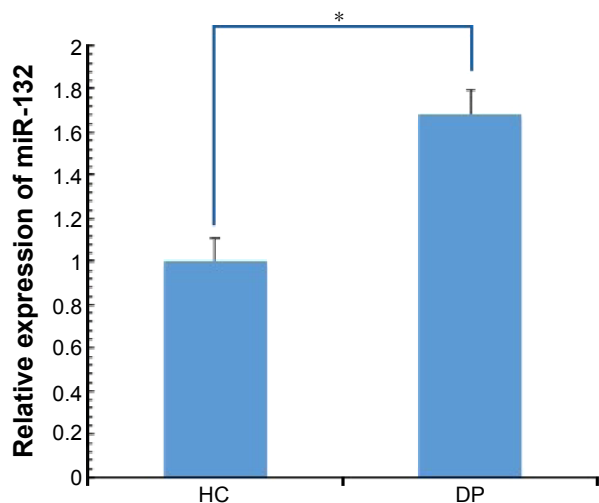


Figure 1 Blood-based miR-132 levels (indicated by $2^{-\Delta\Delta C_t}$): drug-free DPs vs HCs. **Notes:** ^a $P=0.010$. Relative expression of miR-132 in the two groups is expressed as mean ± SE in the bar chart. Differences in RQ values between the two groups were statistically measured by Student's t-test.

Abbreviations: miR-132, microRNA-132; HC, healthy control; DP, depression patient; RQ, relative quantification.

(delayed) (MDD: 9.10±1.87, HC: 9.95±1.75, $t=2.722$, $P=0.007$) and PRM – percent correct (delayed) (MDD: 75.84±15.56, HC: 82.86±14.61, $t=2.702$, $P=0.008$; Table 3).

Correlations among miR-132 expression level, visual memory function, and clinical symptoms

In the MDD group, the increased miR-132 expression levels were negatively correlated with PRM – number correct (delayed) ($r=-0.273$, $P=0.032$) and PRM – percent correct (delayed) ($r=-0.274$, $P=0.031$). However, there was no correlation between miR-132 levels and performance in the HC group. In the MDD group, HAMA scores were negatively correlated with PRM – number correct (immediate) ($r=-0.301$, $P=0.017$) and PRM – percent correct (immediate) ($r=-0.305$, $P=0.016$; Table 4).

No significant relationship between miR-132 level and HAMD ($r=0.019$, $P=0.883$) or HAMA ($r=0.175$, $P=0.172$) scores in the MDD group was observed.

Discussion

In our study, we found that visual memory function in MDD patients was significantly decreased compared to that in HCs. MDD patients showed a lower correct number and correct percent than HCs in the delayed modes of PRM. Numerous studies have reported an association between MDD and memory impairments.^{36,37} Using a variety of cognitive tasks, Airaksinen et al¹ analyzed the data from patients with major depression, dysthymia, mixed anxiety–depressive disorder, and minor depression. Their results showed that major depression and mixed anxiety–depression disorder groups exhibited significant memory dysfunction, whereas minor depression did not affect cognitive performance. Landrø et al³⁸ reported that patients with MDD performed significantly worse than HCs in working memory and verbal long-term memory. Some studies have found MDD patients were impaired in both verbal memory and visual memory^{39,40} and delayed verbal memory and verbal percent retention, but not in immediate verbal memory or visual memory.³⁷ Another study has found that immediate and delayed visual–spatial memories were impaired while immediate and verbal long-term memories were preserved.⁴¹ Our results showed that delayed visual memory decreased in patients with MDD. This finding was consistent with previous research outcomes.

Studies have revealed that miR-132 expression has tissue specificity, which is highly expressed in nervous tissues and involved in axonal growth, proliferation and differentiation of synapses, and neural tumor formation process.⁴² In addition,

Table 3 Comparison of the expression of cognition scores between patients with depression and HCs

Test item	Measurement	HC	MDD	t	P-value
PRM_NCi	PRM – number correct (immediate)	11.05±1.39	11.06±1.42	-0.040	0.971
PRM_PCi	PRM – number correct (immediate)	92.14±11.68	92.31±11.90	-0.083	0.934
PRM_MCLi	PRM – mean correct latency (immediate)	2,095.48±896.50	2,317.31±1,510.91	-1.167	0.293
PRM_NCd	PRM – number correct (delayed)	9.95±1.75	9.10±1.87	2.722	0.007*
PRM_PCd	PRM – percent correct (delayed)	82.86±14.61	75.84±15.56	2.702	0.008*
PRM_MCLd	PRM – mean correct latency (delayed)	2,047.58±621.16	2,195.68±757.48	-1.255	0.245

Notes: Lower scores represent better neuropsychological performance. * $P < 0.05$.

Abbreviations: HC, healthy control; MDD, major depressive disorder; PRM, pattern recognition memory.

miR-132 expression levels were the highest among upregulated miRNAs in response to neuronal activity in the dentate gyrus, which was implicated in mood disorders.⁴³ One study demonstrated that miR-132 expression in neurons and glia mediated the brain-derived neurotrophic factor (BDNF) action whose low levels are implicated in depression.⁴⁴ Suppression of miR-132 by 2-*O*'-methyl oligoribonucleotide can increase BDNF transcript levels.⁵ Studies reported that miR-132 expression was increased in the peripheral blood samples of patients with MDD, while BDNF and MeCP2 were decreased, which suggested that miR-132 may be a key factor in controlling stress-induced hippocampal neuroplasticity and neuronal survival in depression.^{44,45} In line with previous findings, our results confirmed that MDD patients had increased miR-132 expression level.

Numerous behavioral studies in animal models have identified behavioral changes in rodents accompanied with the disruption in the miR-132 expression level.⁵ Hansen et al²⁹ demonstrated impaired recognition memory in transgenic mice with overexpressed miR-132 throughout the forebrain. Scott et al⁴⁶ reported that specific overexpression of miR-132

selectively in the perirhinal cortex of rat was associated with impairment of short-term recognition memory. The primary function of miR-132 in response to neural activity is to serve as a proteome switch, broadly modulating the translation of proteins critical for the establishment or maintenance of memory.⁴⁷ Our study found that delayed visual memory deficits were positively correlated with overexpression of miR-132 in DPs. To our knowledge, this is the first study to explore the possible association between alterations in visual memory function of patients with depression and the miR-132 expression level.

Correlation analyses also suggested the association between memory deficits and clinical symptoms in patients with depression (Table 4). In the PRM test, anxiety score and the correct number and the correct percent of pattern recognition were negatively correlated in patients with depression. A number of investigations have indicated that only some depressed individuals have memory dysfunction, but these studies still cannot clearly identify who those individuals are.^{48,49} However, another study suggested that anxious individuals perform less well than non-anxious individuals on

Table 4 Relationship between PRM scores and miR-132 expression and HAMD or HAMA scores in 62 patients and 73 HCs

	miR-132	HAMD	HAMA
MDD			
PRM_NCi	$r = -0.107, P = 0.407$	$r = -0.135, P = 0.297$	$r = -0.301, P = 0.017^*$
PRM_PCi	$r = -0.103, P = 0.424$	$r = -0.135, P = 0.295$	$r = -0.305, P = 0.016^*$
PRM_MCLi	$r = 0.126, P = 0.328$	$r = 0.164, P = 0.203$	$r = 0.026, P = 0.108$
PRM_NCd	$r = -0.273, P = 0.032^*$	$r = -0.097, P = 0.453$	$r = -0.101, P = 0.437$
PRM_PCd	$r = -0.274, P = 0.031^*$	$r = -0.100, P = 0.440$	$r = -0.100, P = 0.440$
PRM_MCLd	$r = 0.139, P = 0.283$	$r = 0.288, P = 0.230$	$r = 0.113, P = 0.383$
HC			
PRM_NCi	$r = -0.104, P = 0.382$		
PRM_PCi	$r = -0.102, P = 0.391$		
PRM_MCLi	$r = 0.006, P = 0.959$		
PRM_NCd	$r = -0.035, P = 0.768$		
PRM_PCd	$r = -0.036, P = 0.760$		
PRM_MCLd	$r = 0.042, P = 0.721$		

Note: * $P < 0.05$.

Abbreviations: PRM, pattern recognition memory; miR-132, microRNA-132; HAMD, Hamilton Depression Rating Scale; HAMA, Hamilton Anxiety Rating Scale; HC, healthy control; MDD, major depressive disorder.

memory tasks.⁵⁰ Epidemiological studies from 1972 to 1985 have revealed comorbidity rates of 21%–91% between anxiety and depression.^{51,52} This linkage indicated that depressed individuals may differ and therefore potentially add to the variability of memory functioning. A previous study suggested that among individuals with symptoms of depression and/or anxiety, those experiencing high levels of coexisting depression and anxiety are more likely to have immediate and delayed memory difficulties than individuals reporting either depressive or anxious symptoms alone or those with low levels of both.⁵³ In our study, we found that depressed individuals with anxiety symptoms were positively correlated with immediate visual memory defects.

The current study has three limitations. First, the sample size was relatively small. Some patients did not finish the PRM task, which may affect the results of cognition analysis. Second, our MDD patients were not followed over time, and for this reason, it was difficult for us to determine the alteration of miR-132 expression in depression symptoms' recovery or memory improvement. We yet cannot determine whether the reported relation between miRNA-132 alteration and visual memory impairment was restricted to periods of acute episode or persisted during recovery. Finally, we need further experiments to explore the causal relationship between upregulated miR-132 expression level and visual memory changes.

Conclusion

We found that: 1) miR-132 expression levels were upregulated in major depression; 2) peripheral miR-132 expression levels were associated with visual memory defects in patients with depression; and 3) visual memory dysfunction was positively correlated with clinical symptoms.

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Disclosure

The authors report no conflicts of interest in this work.

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