

Kinetics of plasma microRNA-499 expression in acute myocardial infarction

Xi Chen^{1*}, Lizhu Zhang^{1*}, Tong Su¹, Heng Li¹, Qiang Huang¹, Dan Wu¹, Chengjian Yang¹, Zhijun Han²

¹Department of Cardiology, ²Department of Laboratory Medicine, Wuxi Second People's Hospital of Nanjing Medical University, Wuxi 214002, China

*These authors contributed equally to this work.

Correspondence to: Chengjian Yang. Department of Cardiology, Wuxi Second People's Hospital of Nanjing Medical University, Wuxi 214002, China. Email: doctory2071@sina.com; Zhijun Han. Department of Laboratory Medicine, Wuxi Second People's Hospital of Nanjing Medical University, Wuxi 214002, China. Email: zjhan1125@163.com.

Background: MicroRNA (miRNA) is reported to be present in human plasma and has been increasingly suggested as a biomarker for diseases. Our study aimed to investigate the kinetics of cardiac-specific microR-499 (miR-499) in acute myocardial infarction (AMI).

Methods: Circulating concentrations of cardiac enriched miR-499 were measured by quantitative PCR in 73 patients with acute coronary syndrome (ACS), including 53 with AMI and 20 with unstable angina (UA). Thirty healthy subjects were used as controls. Plasma samples in AMI group were obtained immediately after admission and at 12 h, 24 h, 3 d and 7 d after onset of symptoms. Plasma samples in UA and healthy control groups were collected immediately after admission. The severity and extent of coronary stenotic lesions were evaluated on the basis of coronary angiography using Gensini score.

Results: miR-499 expression levels were significantly higher in the 53 AMI patients than in the 20 UA patients and 30 healthy controls immediately after admission ($P < 0.01$). A measurable increase in miR-499 levels was observed in AMI patients within 24 h of the last onset of chest pain and the levels returned to the baseline after 7 d. Plasma miR-499 levels in the patients with AMI were positively-correlated with cTnI ($r = 0.384$, $P < 0.01$) and CK-MB ($r = 0.402$, $P < 0.01$). In addition, miR-499 levels in AMI patients with two- and three-vessel coronary artery disease (CAD) were significantly higher than those in patients with single-vessel CAD ($P < 0.05$). Gensini scores were used to evaluate the severity of coronary stenosis. miR-499 were positively correlated with Gensini scores ($r = 0.52$, $P < 0.01$). miR-499 levels at admission were significantly higher than those 24 h after percutaneous coronary intervention (PCI) in AMI patients ($P < 0.01$) and were negatively correlated with LVEF ($r = 0.36$, $P = 0.008$).

Conclusions: Cardiac-specific miRNA-499 levels were found to be linearly proportional to myocardial damage. MiRNA-499 might prove to be a new biomarker for AMI and a predictor of the risk of myocardial ischemia.

Keywords: MicroRNA (miRNA); acute myocardial infarction (AMI); biomarker

Submitted Jul 15, 2014. Accepted for publication Oct 09, 2014.

doi: 10.3978/j.issn.2072-1439.2014.11.32

View this article at: <http://dx.doi.org/10.3978/j.issn.2072-1439.2014.11.32>

Introduction

Acute myocardial infarction (AMI) is the one of the leading causes of morbidity and mortality worldwide. At present, the diagnosis of AMI mainly depends on clinical symptoms, electrocardiographic (ECG) changes and circulating

biomarkers. As clinical presentations and echocardiographic findings are often nonspecific in patients with chest pain, cardiac biomarkers are more important for the diagnosis of AMI. Current biomarkers including CK-MB and troponins have been widely applied to the clinical diagnosis of AMI, but they have several shortcomings such as slow-

release patterns or limitations in specificity (1-3). It is therefore necessary to seek new more sensitive and specific biomarkers of AMI.

MicroRNA (miRNA) is a type of small molecular non-coding RNA with the length about 21-25 nucleotides (4). Accumulating evidence has indicated that miRNAs play a key role in various diseases such as leukemia, progressive liver disease, and neurodegenerative diseases (5-7). In addition, miRNAs were shown to be very stable in circulating blood (8), and the levels of individual miRNA and specific miRNA signatures were reported to be associated with the diagnosis and prognosis of cardiovascular diseases (9). Some recent studies (10,11) have demonstrated that miRNAs, including miR-1, miR-133a and miR-208a, are cardio- or skeletal- muscle specific, and therefore have been proposed as candidate biomarkers for myocardial infarction.

Our prophase study showed that microR-499 (miR-499) released into the circulation during AMI and could be used to detect and monitor the myocardial injury (12). Most studies have focused on the acute phase of myocardial infarction. The present study intended to detect miR-499 levels in AMI patients, and see whether miR-499 is associated with the severity and extent of coronary stenotic lesions and observe changes in plasma miR-499 during emergency percutaneous coronary intervention (PCI).

Materials and methods

Study population

In this study, 73 patients with acute coronary syndrome (ACS), including 53 patients with AMI and 20 patients with unstable angina (UA), who were admitted to emergency and cardiology departments in Wuxi Second People's Hospital (Wuxi, China) between January 2013 and December 2013. All of these patients were consecutively included from the admitted patients. The inclusion criteria for patients with AMI were based on the newly developed universal definition of MI (13). Briefly, patients with AMI were clinically diagnosed by biochemical markers (cTnI >0.1 ng/mL), acute ischemic-type chest pain, ECG changes, and coronary angiography. The diagnostic criteria for UA were as follows: chest pain with an accelerating pattern or a prolonged duration (>20 min) or recurrent episodes at rest or with minimal effort and ischemic ECG changes such as ST-segment elevation, ST-segment depression of 0.1 mV, or T-wave inversion in at least two contiguous ECG leads.

In addition, 30 healthy adult volunteers who underwent routine physical examinations in our hospital were enrolled as the control. Patient characteristics and clinical features are described in *Table 1*.

The research protocol was recognized by the Ethics Committee in Wuxi Second People's Hospital. Informed consent was obtained from all subjects before initiation of the study.

Plasma collection and storage

Plasma samples in AMI group were obtained immediately after admission and at 12 h, 24 h, 72 h and 7 d after onset of the symptoms. The duration between the onset chest pain and arrival at the emergency room was 4.46 ± 3.36 h. Blood samples in UA group were collected immediately after admission. All plasma samples (3-5 mL) were extracted from citrated tubes and stored at -80°C .

Coronary angiography and diagnostic criteria for coronary artery disease (CAD)

All coronary angiograms were performed by experienced investigators who were blinded to the study, using transradial or transfemoral approaches. At least four projections of the left anterior descending artery and the circumflex artery and two projections of the right coronary artery were taken into consideration. CAD was defined as the presence of one or more coronary stenoses with lumen narrowing more than 50% in a given patient. The extent of CAD was coded as 0, 1, 2, or 3 or more depending on the number of major coronary vessels with luminal stenosis more than 50%. Left main coronary artery stenosis 50% was considered as a two vessel disease.

Gensini score

Gensini score (14) was used to assess the severity of coronary artery stenosis. When the severity of coronary artery stenosis was 0-25%, 26-50%, 51-75%, 76-90%, 91-99% and 100%, the score was 1, 2, 4, 8, 16 and 32 points, respectively. The coefficient was determined in accordance with the locations of coronary artery lesions: 5 points for the left trunk; 2.5, 1.5 and 1 point for the proximal segment, middle segment and distal segment of the left anterior descending branch respectively; 2.5 and 1 point for the proximal segment and distal segment of the circumflex branch respectively; 1 point for the right coronary artery,

Table 1 Demographic and clinical characteristics of the patients and healthy control

Group	AMI	UA	Control	P (AMI vs. UA vs. control)
Total population	53	20	30	–
Age (mean ± SD)	68.8±7.3	66.2±5.1	70.3±7.9	0.131
Sex (male/female)	44/9	15/5	24/6	0.739
Hypertension (n, %)	37 (69.8)	13 (65.0)	15 (50.0)	0.195
Diabetes (n, %)	12 (22.6)	9 (45.0)	7 (23.0)	0.136
Tobacco (n, %)	33 (62.3)	12 (60.0)	15 (50.0)	0.544
TC (mmol/L)	4.16±0.94	3.85±1.13	3.78±0.83	0.153
TG (mmol/L)	1.76±0.92	1.63±1.01	1.37±0.67	0.149
HDL (mmol/L)	1.12±1.28	1.13±0.37	1.23±0.36	0.290
LDL (mmol/L)	2.69±0.91	2.35±0.94	2.32±0.71	0.119
Cr (μmol/L)	70.7±13.7	69.9±12.2	68.1±11.0	0.661
Medications, n [%]				
β-blockers	20 [38]	6 [30]	–	0.595
CCB	10 [17]	3 [15]	–	0.751
ACEI	16 [31]	2 [10]	–	0.710
ARB	14 [26]	6 [30]	–	0.775
Nitrates	21 [40]	11 [55]	–	0.294
Statins	48 [91]	15 [75]	–	0.124

All AMI patients with a large thrombus burden received aspirin, clopidogrel, heparin and abciximab. TC, total cholesterol; TG, total glyceride; HDL, high-density lipoprotein; LDL, low-density lipoprotein; Cr, creatinine; CCB, calcium channel blocker; ACEI, angiotensin-converting enzyme inhibitor; ARB, angiotensin inhibitors.

the first diagonal branch, the second diagonal branch and the posterior branch of the left ventricle; and 0.5 point for the others.

Detection of miR-499

Total RNA was extracted from the plasma samples using the mirVana PARIS kit (Ambion, Applied Biosystem) with enrichment for small RNAs. Reverse transcription of miRNA was performed with the miScript reverse transcription kit (Applied Biosystem). miR-499 was quantitated by using TaqMan miRNA quantitative reverse transcriptase-polymerase chain reaction (qRT-PCR) assay according to the protocol of the manufacturer (Applied BioSystems). U6RNA was performed as a miRNA internal control.

The data were analyzed with automatic setting for assigning baseline; the threshold cycle (Ct) was defined as the fractional cycle number at which the fluorescence exceeds the given threshold. The Ct values from real-time

PCR assays greater than 40 were treated as 40. The plasma levels of miRNA were detected and analyzed by two investigators who were blinded to the clinical data of the patients. Expression values were normalized by using the mean Ct, and the data obtained by real-time PCR were translated in log₂ (relative level) (15,16).

Statistical analyses

Experimental data are expressed as the mean ± standard deviation ($\bar{x} \pm s$). Before the analyses, all data were subjected to a normality test (Shapiro-Wilk). The Kruskal-Wallis H test or one-way ANOVA were used to compare data between the three groups, and the LSD *t*-test was used for inter-group comparisons. To assess whether the time courses of miR-499 expression differed, we compared them by repeated measures ANOVA. Correlations between two variables were determined by the Spearman tests. Statistical analyses were performed with the SPSS statistical package. P values <0.05 were considered statistically significant.

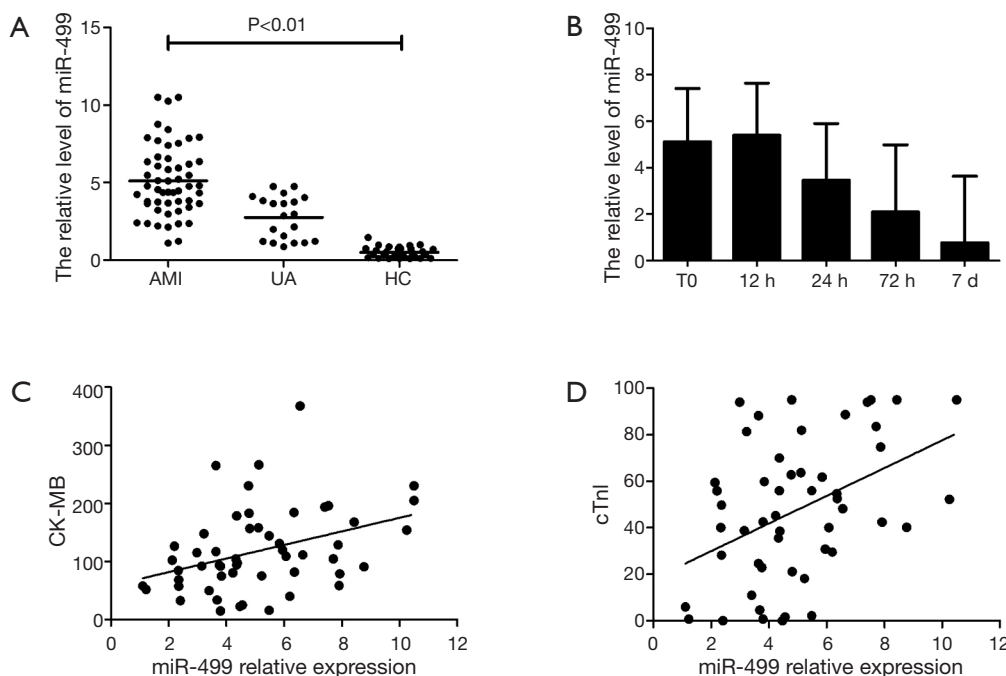


Figure 1 The pattern of plasma miR-499 levels in acute myocardial infarction. (A) miR-499 levels in AMI and UA patients and healthy controls (dark lines represent the mean values); (B) time course of miR-499 plasma levels in AMI patients after symptom onset; (C,D) correlations between miRNA-499 levels and CK-MB and cTnI concentrations. The relative levels of miR-499 in the patients with AMI were positively correlated with serum concentrations of cTnI ($r=0.384$, $P<0.01$) or CK-MB ($r=0.402$, $P<0.01$). AMI, acute myocardial infarction; UA, unstable angina.

Results

The pattern of plasma miR-499 levels in acute myocardial infarction (AMI)

This study included 53 AMI patients, 20 UA patients, and 30 healthy controls. The baseline clinical characteristics of the groups are shown in *Table 1*. There were no significant differences in the clinical characteristics of the patients between the three groups. Blood samples were obtained from each AMI patient at various time points (T0 h, 12 h, 24 h, 72 h, and 7 d) after the onset of AMI. The first plasma sample was obtained immediately after admission (T0 h). The average duration between the onset of chest pain and arrival at the emergency room was 4.46 ± 3.36 h. As shown in *Figure 1A*, plasma samples in AMI, UA and healthy groups were drawn immediately after admission, the relative level of plasma miR-499 in 53 patients with AMI (5.12 ± 2.29) was significantly higher than that in UA group (2.75 ± 1.39) and healthy control group (0.50 ± 0.35), and the differences were statistically significant ($P < 0.01$). We also investigate the time course of plasma miR-499 levels in AMI patients (*Figure 1B*). The relative level of

plasma miR-499 was 5.40 ± 2.24 , 3.47 ± 2.43 , 2.11 ± 2.88 and 0.77 ± 2.86 at 12 h, 24 h, 72 h and 7 d after onset of symptoms respectively, and returned to the baseline level after 7 d, when it was not significantly different from that in healthy control group ($P > 0.05$). Furthermore, correlation analysis showed a positive correlation between circulating levels of miR-499 and cTnI concentrations in AMI patients (*Figure 1C*, $r=0.384$, $P < 0.01$), as well as between miR-499 and CK-MB (*Figure 1D*, $r=0.402$, $P < 0.01$). These data suggested that circulating miR-499 might be regarded as a novel biomarker of AMI.

MicroRNA (miRNA)-499 and the severity of coronary atherosclerosis

Of the 53 AMI patients, 42 patients underwent coronary angiography, who were divided into three subgroups according to the number of affected branches ($n=1, 2$ and 3). The relative level of miR-499 was 3.82 ± 1.98 , 5.53 ± 2.62 and 5.92 ± 1.97 , respectively. It was higher in two- and three-vessel CAD than that in single-vessel CAD ($P < 0.05$) (*Figure 2A*). The Gensini scoring system was used to evaluate the

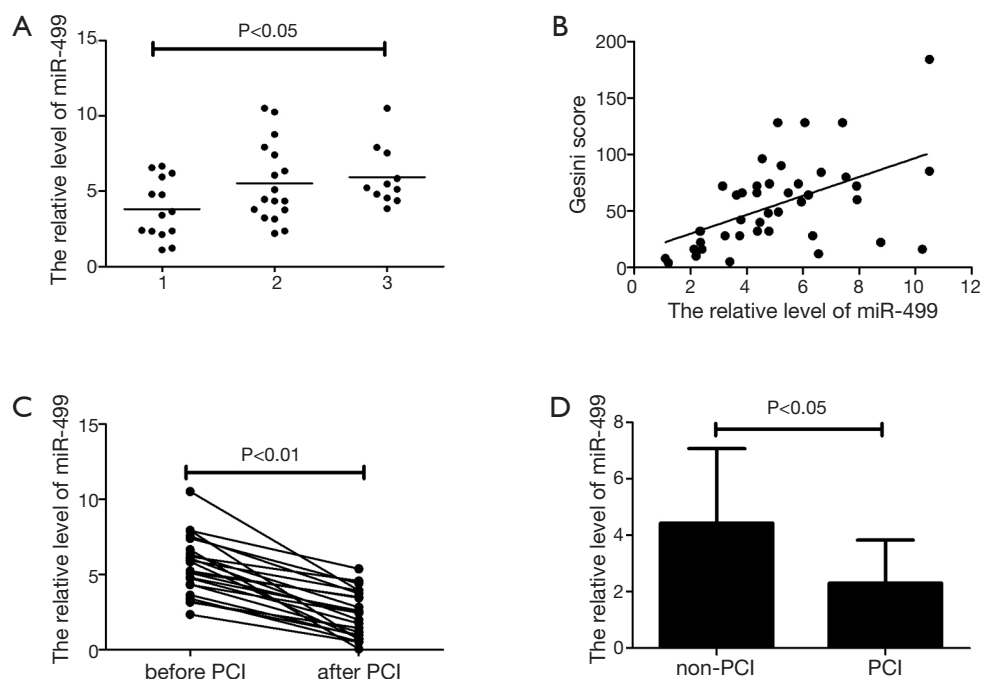


Figure 2 miRNA-499 expression levels in AMI patients. (A,B) miRNA-499 and the severity of coronary atherosclerosis: relative levels of miR-499 in patients with different numbers of affected branches ($n=1, 2$ and 3); dark lines represent the mean values. Correlation between plasma miR-499 levels and Gensini scores in AMI patients. Spearman's correlation analysis showed that plasma miR-499 levels were positively correlated with the severity of coronary stenosis ($r=0.52, P<0.01$); (C,D) miRNA-499 expression levels in AMI patients after emergency PCI: plasma miR-499 level at 24 h after PCI were significantly lower than those at admission in the emergency PCI group ($P<0.01$); plasma miR-499 levels at 24 h after PCI was significantly lower than those in the non-PCI group ($P<0.01$). AMI, acute myocardial infarction; UA, unstable angina.

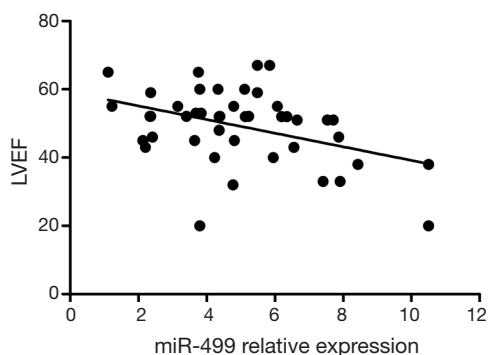


Figure 3 Correlation between miRNA-499 levels and long-term myocardial systolic function: miRNA-499 levels were negatively correlated with LVEF ($r=-0.36, P=0.008$).

severity and extent of coronary stenotic lesions in the 42 AMI patients by coronary angiography. Spearman's correlation analysis showed that plasma miR-499 was positively correlated with the severity of coronary stenosis ($r=0.52, P<0.01$) (Figure 2B).

miRNA-499 expression in AMI patients after emergency PCI

Of the 53 AMI patients, 29 patients received emergency PCI successfully. As shown in Figure 2C, circulating miR-499 level at 24 h after PCI was significantly lower than that at admission in emergency PCI group (2.31 ± 1.52 vs. $5.51 \pm 0.38, P<0.01$). Meanwhile, plasma miR-499 at 24 h after PCI was significantly lower than that in non-PCI group ($4.43 \pm 2.64, P<0.05$) (Figure 2D).

Correlation between miRNA-499 and myocardial systolic function

We investigated whether circulating cardiac-specific miRNA-499 levels correlated with long-term myocardial systolic function, as measured by LVEF. As shown in Figure 3, we found that miRNA-499 levels were negatively correlated with LVEF ($r=-0.36, P=0.008$). Scatter plots for these dates are shown in additional file 1: Figure 3.

Discussion

Recent studies (16-18) have shown that miR-499 may be a clinically practicable biomarker for AMI. Adachi *et al.* (17) proposed that miR-499 level was increased significantly at 6 and 12 h after myocardial infarction. Devaux *et al.* (16) found that miR-499 level was elevated significantly in AMI patients within 3 h after chest pain and was positively correlated with hs-cTnT, the positive rate of miR-499 being 93%. Olivieri *et al.* (18) found that plasma miR-499-5p in elderly NSTEMI patients was nearly 80-fold that in healthy control group, and its sensitivity and specificity were significantly higher than hs-TnT. However, these studies mostly focused on the acute phase of myocardial infarction. An ideal biomarker should not only have sensitivity and specificity but be measurable with respect to myocardial damage (19). Our study aimed to investigate the kinetics of plasma cardiac-specific miR-499 in AMI and see whether it is associated with the severity and extent of coronary stenotic lesions.

In our study, we found that plasma miR-499 levels significantly increased in the AMI group within 12 h after the onset of symptoms, and eventually returned to baseline levels, which were not significantly different from those in the healthy control group, suggesting that cardiac-specific miR-499 might be released into the blood circulation from the necrotic myocardium in the early stage of AMI and then change along with the progression of AMI. Correlation analysis showed a positive correlation between circulating levels of miR-499 and cTnI, as well as CK-MB, which is consistent with their cardiac expression and release from the injured heart. In addition, plasma miR-499 levels in UA patients were higher than those in the control group, which may be due to increased release from cardiomyocytes in the ischemic myocardial tissue. However, Widera *et al.* (11) proposed that there was no difference in the levels of miR-499 between UA and AMI patients. miR-499 levels are considered to be dynamic in the plasma, thus, we were concerned that the blood samples were taken too early to observe the change in miR-499 levels in their study.

Additionally, we investigated the correlation between the severity of CAD and miR-499 and found that the level of miR-499 in two- and three-vessel CAD was significantly higher than that in single-vessel CAD. In addition, we used Gensini score to assess the severity of CAD in AMI patients and found that plasma miR-499 was positively correlated with the severity of coronary stenosis. Meanwhile, we found

that higher severity of the coronary artery of AMI patient represents a higher risk of myocardial ischemia. This positive correlation between Gensini score and miR-499 level further suggest that miR-499 release from myocardial cells may reflect the risk of the occurrence of myocardial ischemia. Therefore, miR-499 may prove to be a useful marker for predicting the risk of myocardial ischemia and serve as a warning sign for the prevention and early intervention of AMI.

Finally, we found that plasma miR-499 levels in AMI patients 24 h after emergency PCI were significantly lower than those at admission, further, plasma miR-499 levels were significantly lower 24 h after the onset of symptoms in the emergency PCI group than those in the non-PCI group. Emergency PCI is said to be associated with a high rate of patency of the infarct-related artery, and effective reperfusion of the occluded coronary artery could reduce myocardial necrosis (20,21). These finding seem to suggest that miR-499 may be a good biomarker to be measurable in linear proportion to myocardial damage and has an important predictive effect in evaluating myocardial ischemia reperfusion in AMI patients. We also found that miRNA-499 levels were negatively correlated with LVEF, which might indicate poor prognosis and increase the risk of death or the development of heart failure. It remains possible that measuring miRNA-499 levels at later time points would ensure their prognostic value. However, our study did not assess the conditions of the AMI patients at discharge or their follow-up survival. Therefore, further research is needed.

Conclusions

In conclusion, miR-499 may prove to be a new biomarker for the diagnosis of AMI and clinical evaluation of the risk of myocardial ischemia, thus improving the sensitivity and specificity of screening patients with myocardial infarction and evaluation of effective reperfusion for myocardial ischemia.

Acknowledgements

Funding: This work was funded by Clinical Technology Foundation of Jiangsu Province (BL2012042), National Natural Science Foundation of China (81301503).

Disclosure: The authors declare no conflict of interest.

References

- de Lemos JA, Drazner MH, Omland T, et al. Association of troponin T detected with a highly sensitive assay and cardiac structure and mortality risk in the general population. *JAMA* 2010;304:2503-12.
- Alpert JS, Thygesen K, Antman E, et al. Myocardial infarction redefined--a consensus document of The Joint European Society of Cardiology/American College of Cardiology Committee for the redefinition of myocardial infarction. *J Am Coll Cardiol* 2000;36:959-69.
- de Winter RJ, Koster RW, Sturk A, et al. Value of myoglobin, troponin T, and CK-MBmass in ruling out an acute myocardial infarction in the emergency room. *Circulation* 1995;92:3401-7.
- Ambros V. The functions of animal microRNAs. *Nature* 2004;431:350-5.
- Calin GA, Dumitru CD, Shimizu M, et al. Frequent deletions and down-regulation of micro- RNA genes miR15 and miR16 at 13q14 in chronic lymphocytic leukemia. *Proc Natl Acad Sci U S A* 2002;99:15524-9.
- Murakami Y, Toyoda H, Tanaka M, et al. The progression of liver fibrosis is related with overexpression of the miR-199 and 200 families. *PLoS One* 2011;6:e16081.
- Cogswell JP, Ward J, Taylor IA, et al. Identification of miRNA changes in Alzheimer's disease brain and CSF yields putative biomarkers and insights into disease pathways. *J Alzheimers Dis* 2008;14:27-41.
- Lawrie CH, Gal S, Dunlop HM, et al. Detection of elevated levels of tumour-associated microRNAs in serum of patients with diffuse large B-cell lymphoma. *Br J Haematol* 2008;141:672-5.
- Gupta SK, Bang C, Thum T. Circulating microRNAs as biomarkers and potential paracrine mediators of cardiovascular disease. *Circ Cardiovasc Genet* 2010;3:484-8.
- Wang F, Long G, Zhao C, et al. Plasma microRNA-133a is a new marker for both acute myocardial infarction and underlying coronary artery stenosis. *J Transl Med* 2013;11:222.
- Widera C, Gupta SK, Lorenzen JM, et al. Diagnostic and prognostic impact of six circulating microRNAs in acute coronary syndrome. *J Mol Cell Cardiol* 2011;51:872-5.
- Han ZJ, Shi WQ, Shen HY, et al. Diagnostic performance of plasma miR-499 for acute myocardial infarction. *J chin J Lab Med* 2013;36:1096-9.
- Thygesen K, Alpert JS, White HD. Universal definition of myocardial infarction. *Eur Heart J* 2007;28:2525-38.
- Gensini GG. A more meaningful scoring system for determining the severity of coronary heart disease. *Am J Cardiol* 1983;51:606.
- Wang GK, Zhu JQ, Zhang JT, et al. Circulating microRNA: a novel potential biomarker for early diagnosis of acute myocardial infarction in humans. *Eur Heart J* 2010;31:659-66.
- Devaux Y, Vausort M, Goretti E, et al. Use of circulating microRNAs to diagnose acute myocardial infarction. *Clin Chem* 2012;58:559-67.
- Adachi T, Nakanishi M, Otsuka Y, et al. Plasma microRNA 499 as a biomarker of acute myocardial infarction. *Clin Chem* 2010;56:1183-5.
- Olivieri F, Antonicelli R, Lorenzi M, et al. Diagnostic potential of circulating miR-499-5p in elderly patients with acute non ST-elevation myocardial infarction. *Int J Cardiol* 2013;167:531-6.
- Lindahl B. Acute coronary syndrome - the present and future role of biomarkers. *Clin Chem Lab Med* 2013;51:1699-706.
- Sluijter JP, van Mil A, van Vliet P, et al. MicroRNA-1 and -499 regulate differentiation and proliferation in human-derived cardiomyocyte progenitor cells. *Arterioscler Thromb Vasc Biol* 2010;30:859-68.
- Araszkievicz A, Grygier M, Lesiak M, et al. The impact of ischemia-reperfusion injury on the effectiveness of primary angioplasty in ST-segment elevation myocardial infarction. *Postepy Kardiol Interwencyjne* 2013;9:275-81.

Cite this article as: Chen X, Zhang L, Su T, Li H, Huang Q, Wu D, Yang C, Han Z. Kinetics of plasma microRNA-499 expression in acute myocardial infarction. *J Thorac Dis* 2015;7(5):890-896. doi: 10.3978/j.issn.2072-1439.2014.11.32