

Original Article

Serum miR-34a is a potential diagnostic and prognostic marker for osteosarcoma

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Abstract: Background and Aims: MicroRNA-34a (miR-34a) has been shown to be a potential diagnostic and prognostic marker for several cancers. In addition, miR-34a has been reported to suppress osteosarcoma cell proliferation in vitro. However, the clinical value of miR-34a is still unknown. In the present study, we evaluated serum miR-34a level in osteosarcoma patients to explore its diagnostic and prognostic value for this particular malignancy. Methods: Serum from 120 patients with primary osteosarcoma, and 87 sex- and age-matched healthy individuals were obtained. Serum miR-34a level was measured with by a real-time quantitative reverse transcription-polymerase chain reaction assay (qRT-PCR) and correlation with clinicopathological characteristics was further analyzed using t test. Receiver operating curves (ROC), Kaplan-Meier curves, and log-rank analyses as well as Cox proportional hazard models were used to evaluate diagnostic and prognostic significance. Results: The serum miR-34a levels was significantly decreased in the serum of osteosarcoma patients compared to healthy controls ($P < 0.001$). Low miR-34a had significant association with clinical stage ($P = 0.006$), distant metastasis ($P = 0.002$), tumor grade ($P = 0.038$) and response to chemotherapy ($P = 0.017$). The Kaplan-Meier curve showed that patients with high miR-34a level survived significantly longer than patients with low miR-34a levels ($P = 0.036$). Multivariate analysis demonstrated that miR-34a level ($P = 0.001$) was an independent prognostic biomarker for overall survival. To distinguish osteosarcoma patients from healthy controls, ROC/AUC analysis indicated an AUC of 0.83 (sensitivity 0.68; specificity 0.92). Conclusions: Decreased miR-34a might be related to the metastasis of osteosarcoma and might be a novel diagnostic and prognostic biomarker in osteosarcoma.

Keywords: Osteosarcoma, serum, marker, microRNA-34a

Introduction

Osteosarcoma (OS) is the most common malignancy of bone and among the deadliest cancers in adolescents [1, 2]. OS patients are commonly treated with multiagent neoadjuvant chemotherapy, combined with surgery to remove the primary tumor mass and subsequent adjuvant chemotherapy. Introduction of chemotherapy has increased the mean 5-year survival rates of patients with localized disease from 20% in the early 1970s to above 60% at present [3]. However, there has been no impressive progress in improving the survival rate of those with recurrence or metastasis over the last three decades [4]. Unfortunately, most of the current strategies have limited efficacy in the treatment of metastatic and recurrent OS, which remains a major challenge in bone can-

cer fields. Therefore, there is an urgent need to develop novel early molecular markers of diagnostic and therapeutic targets for osteosarcoma. Detection of cancer biomarkers in human serum is a novel method in diagnosis and prognosis of cancer. The serum level of bone-specific ALP reflects the cellular activity of osteoblasts, and elevated serum level of ALP has been found during bone formation or increased bone turnover [5]. However, there is still no effective serum indicator for diagnosis and prognosis of patients with OS.

MicroRNAs (miRs), a class of small non-coding RNA molecules, play critical roles in a variety of biological events, including development, cell proliferation and cell differentiation [6, 7]. MiRNAs negatively regulate gene expression by binding to the 3'-untranslated regions (UTRs) of

Prognostic value of serum miR-34a in patients with osteosarcoma

the corresponding target mRNAs of protein-coding genes, thereby leading to mRNA degradation or translation inhibition [8-10]. Multiple miRs are involved in the invasion and metastasis of different types of cancers, including gastric cancer, breast cancer, hepatocellular carcinoma, colorectal cancer and OS [11-17].

Serum miR-34a is potential biomarker of pancreatic ductal adenocarcinoma [18]. Xiang et al. has reported that reduced serum and intratumoral miR-34a expression levels were independent risk factors for developing BM. Migration and invasion experiments indicated that a reverse correlation existed between miR-34a and HCC tumor migration and invasion [19]. Tian et al. has reported that reduced expression of miR-34a was associated with vascular invasion, and advanced TNM stage. Kaplan-Meier revealed that reduced expression of miR-34a was associated with poor overall survival [20]. Recently, Wu et al. demonstrated that miR-34a was significantly downregulated in osteosarcoma tissues and cell lines, and overexpression of miR-34a inhibited the proliferation of MG-63 and Saos-2 cells. Furthermore, xenograft nude mice model showed that miR-34a inhibited osteosarcoma growth in vivo [21]. Zou et al. demonstrated that miR-34a has a negative regulatory effect on osteosarcoma cell proliferation, migration and invasion [22]. However, the role of serum miR-34a levels in the diagnosis and prognosis of OS has not been reported.

In the present study, we detected the expression levels of miR-34a in serum samples of osteosarcoma patients and healthy individuals. Then, the correlations between serum miR-34a level and clinicopathological factors or overall survival of osteosarcoma patients were evaluated. The prognostic value of miR-34a expression level was demonstrated for overall survival of patients with osteosarcoma.

Materials and methods

Patients and specimens

120 patients diagnosed with osteosarcoma and 87 sex- and age-matched healthy individuals from the affiliated hospital of Qingdao University between March 2007 and July 2012 were recruited in this study. Neither chemother-

apy nor radiotherapy had been used in all of the patients before surgery treatment. The clinical stage of the osteosarcoma patients was classified according to the Tumor Node Metastasis (TNM) Classification of Malignant Tumors (Sixth edition) from the Union for International Cancer Control (UICC). Clinical information of patients was obtained from medical records and pathology reports. All of the osteosarcoma patients received regular followed-up. Overall survival time was defined as the time interval from primary surgery to the date of death or last follow-up. Prior patients' written consent and approval were obtained from the affiliated hospital of Qingdao University according to institutional regulations. We have obtained consent to publish from the participant to report individual patient data.

Total RNA isolation

For each subject, 3 mL venous blood was collected from 207 cases and placed in a test tube. The whole blood samples were incubated at 37°C for 1 h and then centrifuged immediately at 1500 g for 15 min at 4°C. The supernatant serum was stored at -20°C until analysis. Total RNA was isolated from 400 µl serum sample by using mirVana miRNA isolation kit (Applied Biosystems, Foster City, CA) according to the manufacturer's instructions. The concentration and quality of total RNA were monitored by NanoDrop ND-2000 spectrophotometer (Thermo Fisher Scientific, Waltham, MA) and Agilent's 2100 Bioanalyzer (Agilent Technologies, Santa Clara, CA).

Quantitative real-time polymerase chain reaction (qRT-PCR)

The total RNA was reversely transcribed into cDNA (20 µl solution), and qRT-PCR was conducted for each sample using NCode™ EXPRESS SYBR® GreenER™ miRNA qRT-PCR Kit Universal (Invitrogen Corporation, USA) and Light Cycler 96® Real-Time PCR System (Roche, Germany) in a final 20 µl reaction volume according to the protocol of manufacturer. At the end of PCR cycles, melting curve analyses were performed to validate the specific generation of the expected PCR products. The Cq uniformity (SD < 0.2) of Real-Time PCR System had been detected by measuring the samples in triplicates. In order to avoid test errors that

Prognostic value of serum miR-34a in patients with osteosarcoma

Table 1. Osteosarcoma patient characteristics

Characteristic	Osteosarcoma (n = 120)	miR-34a level	p-value
Age (Years)			0.142
≤ 20	72	4.08 ± 0.31	
> 20	48	4.21 ± 0.36	
Gender (n, %)			0.263
Male	69	4.17 ± 0.36	
Female	51	4.13 ± 0.32	
Tumor location			0.562
Femur	73	4.18 ± 0.36	
Tibia	47	4.12 ± 0.33	
Tumor size (cm)			0.094
< 6	78	4.68 ± 0.42	
≥ 6	42	3.87 ± 0.29	
Clinical stage			0.006
IIA	28	5.12 ± 0.64	
IIB	69	3.94 ± 0.25	
III	23	2.83 ± 0.16	
Tumor grade			0.038
Low	52	3.43 ± 0.31	
High	68	5.04 ± 0.38	
Distant Metastasis			0.004
Yes	37	2.34 ± 0.13	
No	83	5.79 ± 0.68	
Response to chemotherapy			0.017
Poor	89	3.66 ± 0.24	
Good	31	5.37 ± 0.32	

Table 2. Multivariate survival analysis of overall survival in 120 osteosarcoma patients

Variables	Hazard ratio	95% CI	P
Age	0.58	0.42-1.17	0.286
Gender	1.29	0.93-2.77	0.573
Tumor location	0.87	0.64-1.54	0.742
Tumor size	1.23	0.89-2.67	0.193
Clinical stage	2.84	2.16-7.45	0.026
Tumor grade	1.47	1.03-2.48	0.12
Distant Metastasis	4.72	3.86-10.15	0.017
Response to chemotherapy	0.93	0.67-2.34	0.163
miR-34a level	3.18	2.87-9.56	0.001

were caused by pollution we set up two negative controls per test. miR-34a primers were purchased from GENEWIZ (Suzhou, China). The relative expression levels of miR-34a for qRT-PCR were normalized by the mean Cq value of U6 snRNA, and were calculated utilizing the $2^{-\Delta\Delta Ct}$ method.

Statistical analysis

Continuous variables were expressed as mean ± standard deviation (SD). Associations between serum levels of miR-34a and clinicopathological characteristics or survival of osteosarcoma patients were evaluated using t test. Nonparametric receiver operating characteristic (ROC) curves were generated to assess diagnostic efficiency. Disease-free survival time was measured as the time from the surgery day until the date of cancer reoccurrence or patient death or the day of the last live follow-up to represent disease progression. Survival probabilities were estimated with a Kaplan-Meier analysis, and significant differences were analyzed with a log-rank test. Multivariate analysis of prognostic factors was performed using a COX regression analysis. All P values were two-sided and $P < 0.05$ was considered statistically significant. The statistical analyses of all experimental data were conducted using SPSS 18.0 statistical software (SPSS Inc., Chicago, IL, USA).

Results

Patient characteristics

In this study, it was shown that the demographic and clinicopathological data of the cases in **Table 1**. Our study comprised of 120 osteosarcoma patients. The mean age was 28.9 ± 10.4 years. Of the 87 healthy controls, male:female was 49:38; the age ≤ 20:> 20 were 51:36. All patients featured osteosarcoma with I-III Enneking stage and received the same curative resection treatment. During the follow-up period, 49 patients (40.8%) died of disease.

Distant metastases developed in 37 patients after the original diagnosis. Of these patients, 7 had bone metastases and 30 had lung metastases (2 patients had both bone and lung metastases). The median overall and disease-free survival of patients was 32 months (95% confidence interval [CI], 31.3-45.3 months) and 26

Prognostic value of serum miR-34a in patients with osteosarcoma

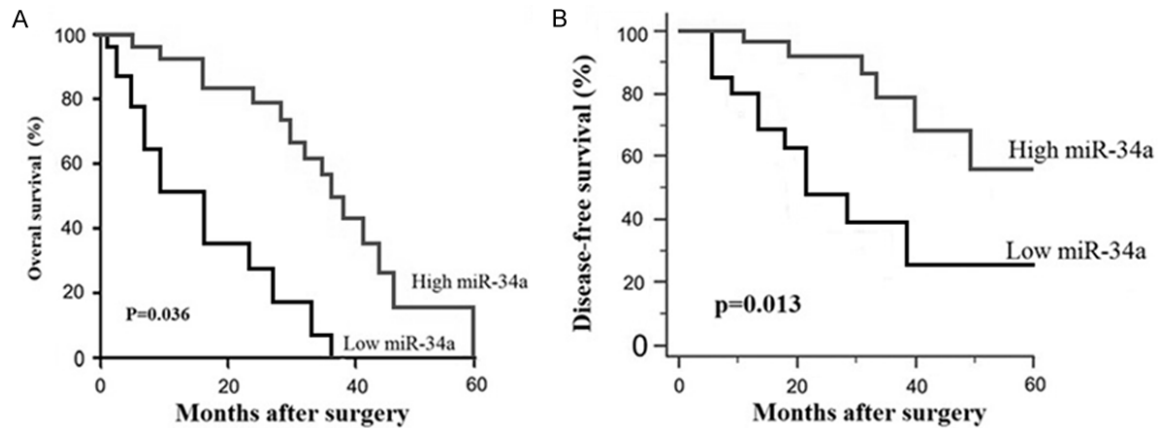


Figure 1. Kaplan-Meier survival curves for osteosarcoma patients with high or low expression of miR-34a. The overall survival curves and disease free survival curves for two groups of osteosarcoma patients with low and high expression of serum miR-34a. A. The overall survival rate of osteosarcoma patients with high miR-34a expression were significantly higher than those with low miR-34a; B. The disease-free survival rate of osteosarcoma patients with high miR-34a level were significantly higher than those with low miR-34a level.

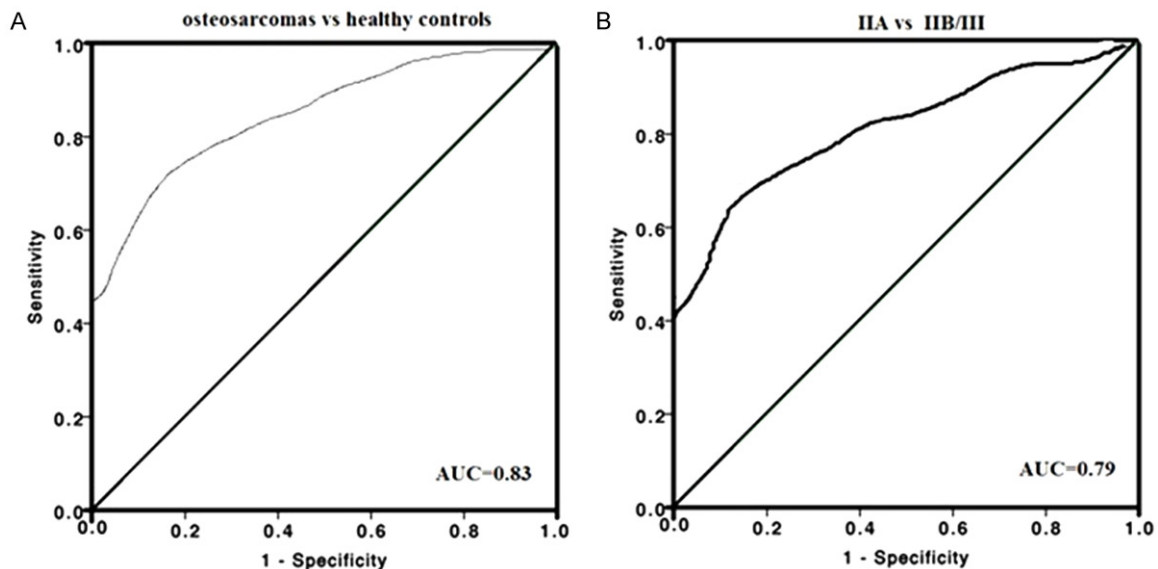


Figure 2. ROC analysis for serum miR-34a for distinguishing osteosarcoma patients from healthy controls. A. ROC for serum miR-34a differentiating osteosarcoma patients from controls; B. Distinguishing osteosarcoma patients with IIB/III stages from those with IIA stages.

months (95% CI: 24.6-37.1 months), respectively.

Serum miR-34a in osteosarcoma patients

qRT-PCR was conducted to determine the levels of serum miR-34 in 120 patients with osteosarcoma and 87 sex- and age-matched healthy controls. The relative serum miR-34a level was 0.72 ± 0.13 patients with osteosarcoma, which was significantly decreased in comparison

with those of healthy controls (4.16 ± 0.34 , $P < 0.001$).

Serum miR-34a and association with clinicopathologic features

The correlations between serum miR-34a and clinicopathologic characteristics of tumors are shown in **Table 2**. Serum miR-34a level was significantly lower in stage III, distant metastases, low stage and poor response to chemotherapy

Prognostic value of serum miR-34a in patients with osteosarcoma

than those with IIA or IIB stages, no metastases, high stage and good response to chemotherapy. However, no significant difference was observed between the serum miR-34a and patient gender, age, tumor location, tumor size and histological grade.

Decreased expression of miR-34a in serum level associates with poor prognosis in osteosarcoma patients

We used the 4.16 as the cutoff value. Serum miR-34a value > 4.16 was the high levels and serum miR-34a value ≤ 4.16 was the low levels. The overall survival ($P = 0.036$, **Figure 1A**) and disease-free survival ($P = 0.013$, **Figure 1B**) of the patients with osteosarcoma was assessed using Kaplan-Meier survival analysis. The Kaplan-Meier curves for overall survival showed that osteosarcoma patients with high serum miR-34a level survived significantly longer than those with low miR-34a levels. Multivariate Cox proportional hazards model analysis suggested that expression level of serum miR-34a was a significant independent prognostic factor of overall survival for patients with osteosarcoma ($P = 0.001$, shown in **Table 2**).

ROC analysis of serum miR-34a level in osteosarcoma patients

ROC/AUC analysis revealed sensitivity and specificity of different serum miR-34a levels. ROC curve analysis illustrated that serum miR-34a level was a potential biomarker for screening OS patients from controls (AUC of 0.83 (sensitivity 0.68; specificity 0.92) (**Figure 2A**)). To assess the potential of miR-34a as a diagnostic biomarker to distinguish advanced cancer patients (IIB/III) from early clinical stages (IIA), we got an AUC of 0.79 (95% CI 0.73-0.95) with a sensitivity of 0.83 and a specificity of 0.70 (**Figure 2B**).

Discussion

Osteosarcoma derives from primitive bone-forming mesenchymal cells and is the most common type of primary bone malignancy [23]. Over the past few decades, the introduction of combinatorial chemotherapy has improved the 5-year overall survival rate of patients with osteosarcoma to approximately 50-60% [24]. However, osteosarcoma-related morbidity re-

mains high due to the difficulty of early diagnosis and the lack of efficient therapeutic approaches for osteosarcoma. Therefore, it is necessary to identify highly sensitive and specific diagnostic and prognostic biomarkers to diagnose osteosarcoma at an early stage and initiate aggressive therapy.

Overexpression of miR-34a reportedly suppresses tumor progression and leads to improved prognoses, whereas reduced miR-34a expression is associated with poor overall survival in several cancers [25-27]. Gao *et al.* found that lower miR-34a expression was correlated with reduced progression-free survival and overall survival in glioma patients [28], but Genovese *et al.* found that lower miR-34a expression led to improved overall survival in glioblastoma patients [29]. However, the role of miR-34a as a serum diagnostic and prognostic biomarker has not been previously explored in osteosarcoma patients. In the present study, we quantified the serum expression levels of miR-34a in osteosarcoma patients and healthy controls and then assessed the potential value of miR-34a as a serum diagnostic and prognostic marker in osteosarcoma patients.

The results from our study show for the first time that the expression of miR-34a was remarkably decreased in the serum of osteosarcoma patients. Furthermore, we showed that the expression of miR-34a could be used to discriminate osteosarcoma from healthy controls, with a specificity of 68% and sensitivity 92%. Notably, decreased serum miR-34a levels were found to be significantly associated with distant cancer metastasis and advanced clinical stage in osteosarcoma patients, which suggests that miR-34a might act as a tumor suppressor in the development of osteosarcoma. In addition, we proved that low levels of miR-34a significantly correlated with poor prognosis in OS patients. Therefore, we suggested that serum miR-34a levels could be a diagnostic biomarker and prognostic factor in the progression of OS.

Chemotherapy is an important treatment modality for osteosarcoma. However, it often fails because of chemoresistance. The underlying mechanisms of chemoresistance are still poorly understood. In recent years, growing evidence demonstrate *miR-34a* has a key role in tumor cell responses to chemotherapeutic ag-

ents and may serve as an effective antitumor therapeutic target [30-34]. Li et al. has reported that increased expression of a panel of tumor suppressive microRNAs (miRNAs), including miR-34a, miR-143, miR-145, and miR-200b/c that were typically lost in osteosarcoma, was observed during diallyl trisulphide treatment, which suggested that miRNAs may be involved in the chemotherapeutic response of osteosarcoma [35]. In the present study, we found that serum miR-34a was higher in OS patients with good response to chemotherapy than those in patients with poor response to chemotherapy, suggesting that miR-34a could be as a marker to predict chemosensitivity in OS patients.

In summary, our findings provide the first hints that serum miR-34a level may be a useful diagnostic and prognostic biomarker that could be used for risk stratification and selection of osteosarcoma patients. In addition, the potential role of miR-34a in osteosarcoma warrants further investigation.

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Disclosure of conflict of interest

None.

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Prognostic value of serum miR-34a in patients with osteosarcoma

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