Original Article

HOTAIR is a predictive and prognostic biomarker for patients with advanced gastric adenocarcinoma receiving fluorouracil and platinum combination chemotherapy

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Abstract: Accumulating evidence suggests that long non-coding RNA (IncRNA) HOTAIR participates in many types of cancer such as gastric cancer and may confer malignant phenotype to tumor cells. Fluorouracil and platinum combination chemotherapy is the first line therapy for gastric cancer. However, it is still unknown whether HOTAIR influences the outcome of cancer patients treated with chemotherapy. This study aimed to evaluate the association of HOTAIR expression with the prognosis of patients with advanced gastric adenocarcinoma (GA) receiving fluorouracil and platinum based chemotherapy. We examined the levels of HOTAIR in 168 GA samples using quantitative real-time PCR and analyzed its relationship with clinical features and prognosis of patients with advanced GA treated with fluorouracil and platinum based chemotherapy. Compared with paracancerous tissues, HOTAIR was significantly upregulated in GA tissues, especially in more advanced cases. High HOTAIR expression was an independent poor prognostic factor for patients with advanced GA. Further stratification analyses revealed that the association between HOTAIR expression and survival in patients with advanced GA remained significant in the subgroup of patients with TNM stages IIIA and IIIB, poorly differentiated, and smaller tumors. In conclusion, our results provide first evidence that HOTAIR may be served as a biomarker that predicts which patient with advanced GA will benefit from fluorouracil and platinum combination chemotherapy.

Keywords: Gastric cancer, long noncoding RNA, HOTAIR, chemotherapy, prognosis

Introduction

Gastric cancer is the fifth most common cancer in the world, accounting for 7% of the total new cancer cases and 9% of the total cancer deaths [1]. The incidence rate of gastric cancer is highest in Eastern Asia [2]. Gastric adenocarcinoma (GA) is the majority of gastric cancer, which can be further divided into intestinal and diffuse types according to Lauren classification. Although multimodality therapy is available to gastric cancer, approximately half of patients with advanced gastric cancer who undergo surgical resection still develop local or distant metastases and die from the cancer [3, 4]. Therefore, it is urgent to identify biomarkers and explore the effective and personalized treatment.

Long noncoding RNAs (IncRNAs) are defined as transcripts > 200 nucleotides in length and are transcribed but non-translated noncoding RNAs in human genome [5, 6]. There are convincing evidence that IncRNA are differentially expressed in various diseases including cancer and play a vital role in cancer progression and metastasis [7, 8]. Functional studies of HOTAIR lifts IncRNAs to a new level with so many documents elucidated that HOTAIR is involved in the development and progression of various types of cancer, including ovarian [9], breast [10], pancreatic [11], and non-small cell lung cancers [12]. HOTAIR is overexpressed in gastric cancer, which promotes its carcinogenesis, invasion and metastasis [13-18]. Recent study revealed that when laryngeal squamous cell carcinoma was treated with increasing concen-

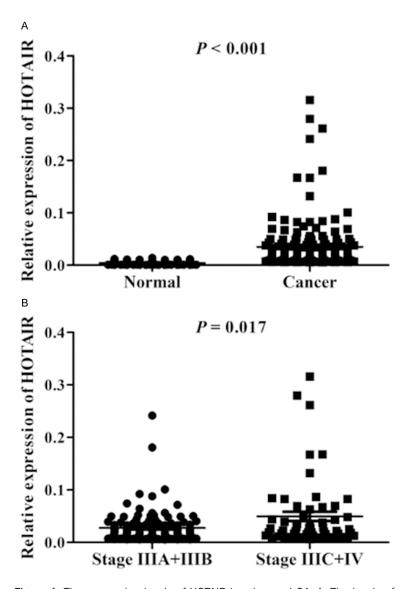


Figure 1. The expression levels of HOTAIR in advanced GA. A. The levels of HOTAIR in GA tissues were significantly upregulated compared paracancerous tissues. B. The levels of HOTAIR in cases with TNM stages IIIA and IIIB were higher than those with TNM stages IIIC and IV.

trations of drugs and extending the duration of treatment, the levels of HOTAIR were dramatically reduced. Furthermore, HOTAIR overexpression confers cisplatin resistance to lung adenocarcinoma cells and ovarian cancer cells [9, 19]. However, it remains largely unknown whether HOTAIR overexpression influences the clinical outcome of cancer patients receiving chemotherapy. In this study, we examined the levels of HOTAIR in 168 patients with advanced GA receiving fluorouracil and platinum based chemotherapy, and evaluated the association of its expression with patient prognosis.

Materials and methods

Patients

A total of 168 patients with advanced GA were collected from Subei People's Hospital and the Biobank of National engineering center for biochip at Shanghai. All patients were histologically confirmed and had TNM stage III-IV tumor. No patient had other cancer history. Samples were obtained before patients were treated with any anti-cancer therapy. This study was approved by the Ethics Committees of Subei People's Hospital and National engineering center for biochip at Shanghai, and all subjects gave written informed consent before enrolling in this study.

Patients were treated with fluorouraciland platinum combination chemotherapy. There were 111 men and 57 women, ranging in age from 28 to 75 years. Clinical information such as date of birth, sex, histologic grade, tumor size, TNM stage, therapeutic regimen, and other necessary information were extracted from computerized clinical database or follow-up records.

Quantitative RT-PCR (qRT-PCR)

Total RNA was extracted from tissue samples using Trizol reagent (Invitrogen, CA, USA) according to the manufacturer's protocols. The concentration of total RNA was quantified by Nanodrop ND-1000 spectrophotometer (Nanodrop, DE, USA). The first-strand cDNA was synthesized using a PrimeScript 1st Strand cDNA synthesis kit (TaKaRa, Dalian, China). Subsequently the cDNA template was amplified by qRT-PCR using SYBR Premix Ex Taq II (TliRNaseH Plus) (TaKaRa, Dalian, China) on an Applied Biosystems® 7900 Real-Time PCR Systems (Applied Biosystems, CA, USA). The relative

Table 1. Association of HOTAIR expression with clinicopathologic parameters

	HOTAIR		Р			
Clinicopathologic parameters	expression		value			
	High	Low	value			
Age (years)						
> 65	29	29	1.000			
≤ 65	55	55				
Sex						
Male	54	57	0.625			
Female	30	27				
Histologic grade						
1+2	16	13	0.540			
3	68	71				
Size (cm)						
> 5	47	52	0.433			
≤ 5	37	32				
TNM stage						
III	66	75	0.092			
IV	18	9				

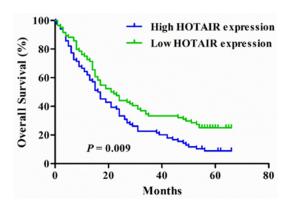


Figure 2. Kaplan-Meier curves of overall survival in 168 patients with advanced gastric adenocarcinoma according to HOTAIR expression.

expression level of HOTAIR was normalized to GAPDH and fold change was calculated using the equation $2^{-\Delta\Delta CT}$. All experiments were performed in triplicate.

Statistical analyses

The level of HOTAIR was compared using Wilcoxon nonparametric test within patient groups. Patients were divided into high and low HOTAIR expression groups according to the median expression level of HOTAIR. Chi-square test or Fisher's exact test were performed to determine the relationship between HOTAIR

expression and clinical pathological parameters, as appropriate. Kaplan-Meier and log-rank analyses were carried out to evaluate the effect of HOTAIR expression on survival. Independent prognostic indicators were assessed in the multivariate analysis using Cox's proportional-hazard model. All statistical analyses were performed using SPSS v20.0 (SPSS, IL, USA) and GraphPad Prism 5.0 (GraphPad Software, CA, USA). A *P* value less than 0.05 was considered statistically significant.

Results

The expression levels of HOTAIR in advanced GA

To investigate the levels of HOTAIR in advanced GA tissues, qRT-PCR was used to determine the levels of HOTAIR in 168 advanced GA. The levels of HOTAIR in GA tissues were significantly higher than those in paracancerous tissues (P < 0.001) (Figure 1A), and was positively related to tumor size (P = 0.034). Furthermore, when patients were subdivided into TNM stages IIIA+IIIB and IIIC+IV groups, the levels of HOTAIR in TNM stage IIIC+IV group were higher than those in TNM stage IIIA+IIIB group (P = 0.017, Figure 1B).

Association of HOTAIR expression with clinicopathologic features of patients with advanced GA

The association of HOTAIR expression with clinicopathologic parameters in patients with advanced GA was shown in **Table 1**. No difference was observed between HOTAIR expression and clinicopathologic parameters. However, there was a weak but not statistically significant association between HOTAIR expression and TNM stage (P = 0.092).

Association of HOTAIR expression level with prognosis of patients with advanced GA

To investigate the effect of HOTAIR expression on survival, Kaplan-Meier curves were adopted in patients with advanced GA receiving fluorouracil and platinum based chemotherapy. As shown in **Figure 2**, median survival time (MST) of patients with high HOTAIR expression was 16.5 months, while MST in those with low HOTAIR expression was 21.5 months (P = 0.009). Multivariate Cox proportional hazard

HOTAIR and gastric adenocarcinoma

Table 2. Univariate and multivariate Cox regression analysis of overall survival in 168 patients with advanced GA

Navialela a	Univariate analysis		Multivariate analysis	
Variables	HR (95% CI)	P value	HR (95% CI)	P value
Age (≤ 65 years vs > 65 years)	0.818 (0.578-1.157)	0.255		
Sex (male vs female)	0.938 (0.654-1.344)	0.726		
Histologic grade (3 vs 1, 2)	1.169 (0.752-1.814)	0.488		
Size (> 5 cm vs ≤ 5 cm)	1.627 (1.150-2.303)	0.006	1.470 (1.029-2.100)	0.034
TNM (IIIC, IV vs IIIA, IIIB)	2.014 (1.424-2.847)	< 0.001	1.755 (1.227-2.511)	0.002
HOTAIR (high vs low)	1.547 (1.105-2.167)	0.011	1.466 (1.044-2.059)	0.027

Table 3. Stratification analysis of HOTAIR associated with survival of patients with advanced GA

	Univariate analysis		Multivariate analysis			
Variables	HR (95% CI)	P value	HR (95% CI)	P value		
Age, years						
≤ 65	1.489 (0.979-2.265)	0.063	1.249 (0.818-1.908)	0.304		
> 65	1.669 (0.944-2.950)	0.078	1.588 (0.894-2.821)	0.115		
Sex						
Male	1.487 (0.984-2.248)	0.060	1.350 (0.890-2.048)	0.159		
Female	1.696 (0.938-3.065)	0.080	1.542 (0.849-2.799)	0.155		
Histologic grade						
1+2	0.769 (0.343-1.725)	0.524	0.780 (0.347-1.751)	0.546		
3	1.802 (1.241-2.617)	0.002	1.679 (1.148-2.456)	0.008		
Tumor size						
≤ 5 cm	1.881 (1.066-3.319)	0.029	1.856 (1.024-3.363)	0.042		
> 5 cm	1.405 (0.922-2.141)	0.113	1.302 (0.848-1.999)	0.227		
TNM						
IIIA+IIIB	1.594 (1.038-2.448)	0.033	1.718 (1.113-2.651)	0.015		
IIIC+IV	1.255 (0.728-2.162)	0.414	1.263 (0.732-2.178)	0.401		

regression analysis that larger tumor size [adjusted hazard ratio [HR] = 1.470, 95% CI: 1.029-2.100, P = 0.034], TNM stages IIIC and IV (adjusted HR = 1.755, 95% CI: 1.227-2.511, P = 0.002), and high HOTAIR expression (adjusted HR = 1.466, 95% CI: 1.044-2.059, P = 0.027) were significant risk factors related to death status in patients with advanced GA (**Table 2**).

Given that HOTAIR expression was correlated with clinical features, further stratification analysis was undertaken to elucidate the exact role of HOTAIR expression in patients with different clinicopathologic parameter. As shown in **Table 3**, among patients with poorly differentiated tumors, high HOTAIR expression conferred an increased risk of mortality (adjusted HR = 1.679, 95% CI: 1.148-2.456, *P* = 0.008) (**Figure**

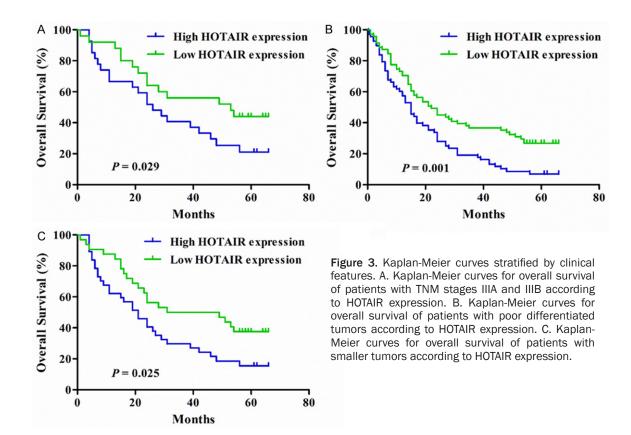
3). Among patients with smaller tumors, those with high HOTAIR expression had a 1.856-fold (95% CI: 1.024-3.363) higher risk of mortality compared with those with low HOTAIR expression. In addition, among patients with TNM stages IIIA and IIIB, high HOTAIR expression was associated with an elevated risk of mortality (adjusted HR = 1.718, 95% CI: 1.113-2.651, P = 0.015).

Discussion

Gastric cancer is a leading cause responsible for cancer deaths [1], of which the vast majority are GA and associated with infection of Helicobacter pylori and Epstein-

Barr virus [20, 21]. The management for gastric cancer is complicated and constantly evolving. Although surgery resection is regarded as the only option for cure, most patients exhibit an advanced disease at diagnosis and usually have a poor prognosis [22]. Furthermore, fluorouracil and platinum combination chemotherapy is the first line therapy for gastric cancer. Therefore, a potential biomarker is urgent for identifying the outcome of patients with advanced gastric cancer who received fluorouracil and platinum based chemotherapy.

Dysregulation of HOTAIR is correlated with multiple types of cancer [9-12, 23-25]. Recent studies have revealed that HOTAIR can act as a scaffold and interact with at least two distinct histone modification complexes to regulate the expression of multiple genes involved in vari-



ous cellular functions [10, 26-28]. HOTAIR binds polycomb repressive complex 2 (PRC2) and LSD1/CoREST/REST complexes via its 5' and 3' domains, respectively, leading to H3K27 methylation and H3K4 demethylation and ultimately gene silencing [29-31]. Previous study have shown that the level of HOTAIR positively correlates with TNM stage and metastasis of gastric cancer [13, 17]. In this study, although high HOTAIR expression was predominant in cases with distant metastasis, the difference did not reach statistical significance. This may be explained that all cases are advanced gastric adenocarcinoma in our study. HOTAIR promotes gastric cancer metastasis by inhibiting Poly r(C)-Binding Protein 1 [32]. Inhibition of HOTAIR impairs gastric cancer cell viability, induces apoptosis, and suppresses the growth of gastric cancer xenograft [15, 16, 33]. Furthermore, HOTAIR acts as a competing endogenous RNA to regulate HER2 expression by sponging miR-331-3p [34]. Given that trastuzumab is used as first line therapy for HER2positive gastric cancer, HOTAIR is potential therapeutic target for gastric cancer.

It is well known that drug resistance is a main obstacle and challenge in cancer treatment.

Previous studies have revealed that HOTAIR overexpression is related to cisplatin resistance [9, 19]. Upregulation of HOTAIR contributes to cisplatin resistance via downregulation ofp21 (WAF1/CIP1) expression in lung adenocarcinoma and exhibits downregulation in cisplatin-responding lung adenocarcinoma tissues [19]. Downregulation of HOTAIR restores cisplatin sensitivity in cisplatin-resistant ovarian cancer cells [9]. In addition, decreased expression of HOTAIR is associated with good response to treatment in sarcoma patient [35]. It has been demonstrated that HOTAIR expression affect the prognosis of patients with many types of cancer [12, 18, 23, 27, 33]. In the present study, we found that high HOTAIR expression was associated with worse survival in patients with advanced GA receiving fluorouracil and platinum combination chemotherapy, especially those with TNM stages IIIA and IIIB, poorly differentiated, and smaller tumors. Therefore, drug resistance caused by HOTAIR overexpression is one plausible reason for the association of HOTAIR overexpression with poor prognosis of cancer patients.

In summary, our findings provide the first evidence that HOTAIR influences the prognosis of

patients with advanced GA treated with fluorouracil and platinum combination chemotherapy. HOTAIR may be served as a biomarker classifier for identifying the subset of patients who will benefit from fluorouracil and platinum combination chemotherapy.

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Disclosure of conflict of interest

None.

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