

Expression of Integrin $\beta 6$ and HAX-1 Correlates with Aggressive Features and Poor Prognosis in Esophageal Squamous Cell Carcinoma

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Purpose: The development of esophageal squamous cell carcinoma (ESCC) is a complicated process in which cell adhesion and motility, mediated by integrins, are involved through connecting the cytoskeleton to extracellular matrix. Different mechanisms via which integrin $\beta 6$ participates in cancer invasion and metastasis have been described by numerous studies; however, the expression and clinical significance of integrin $\beta 6$ in ESCC remain unknown.

Methods: To investigate the differential expression of integrin $\beta 6$ in ESCC, qPCR and immunohistochemistry assays were performed in 10 paired human samples. A total of 137 ESCC samples were further enrolled to evaluate the expression levels of integrin $\beta 6$ and its endocytic trafficking regulator HS1-associated protein X-1 (HAX-1), followed by the evaluation of their correlation with clinicopathological parameters. The overall survival was analyzed using the Kaplan–Meier method, with significant variables further evaluated by multivariate Cox regression analyses.

Results: The expression of integrin $\beta 6$ was markedly increased in ESCC compared with matched adjacent normal tissues. Among the ESCC samples, positive expression of integrin $\beta 6$ was observed in 41.6% tumors, which was associated with histological differentiation, lymph node metastasis and TNM stage. High expression of HAX-1 was detected in 47.4% tumors, and there was a positive relationship between the expression levels of integrin $\beta 6$ and HAX-1. Furthermore, the expression of integrin $\beta 6$ and HAX-1 were independent unfavorable indicators for prognosis. Patients with positive integrin $\beta 6$ and high HAX-1 expression demonstrated worst outcomes.

Conclusion: The present findings suggested the predictive value of integrin $\beta 6$ and HAX-1 as independent indicators of poor prognosis for patients with ESCC, both of which may contribute to the tumor proliferation and metastasis, leading to ESCC progression. Therefore, combined targeting of integrin $\beta 6$ and HAX-1 may provide a potential novel approach for the treatment of ESCC.

Keywords: integrin, HAX-1, esophageal squamous cell carcinoma, prognosis

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Introduction

As a common aggressive malignancy, esophageal cancer (EC) is characterized by a high incidence rate and lethality, exacerbated by distant metastasis and therapeutic resistance. Esophageal squamous cell carcinoma (ESCC) is the predominant EC subtype in East Asia in terms of histological classification, accounting for up to 90% of EC cases.¹ While ongoing progress has been achieved with regards to surgical techniques and adjuvant therapies, the outcomes of ESCC remain

unsatisfactory, with a 5-year overall survival rate $\leq 20\%$, and numerous patients are at a late stage at initial diagnosis.² Therefore, it is important to further clarify the underlying mechanisms of ESCC progression, as well as identify novel biomarkers and potential therapeutic targets, which could improve the prognosis of patients with ESCC.

Cell motility and invasion are important steps in tumor progression and metastasis, in which the adhesion of the cellular cytoskeleton to the extracellular matrix (ECM) occurs. As a large family of transmembrane receptors on the cell surface, integrins act as anchoring and bidirectional signaling molecules that serve as a physical and functional bridge between cells and ECM.³ Currently, 24 distinct types of integrins have been discovered, composed of an α and a β subunit. Integrin $\beta 6$ only partners with αv , giving rise to a single heterodimer $\alpha v\beta 6$ that is unique, with an exclusive and specific expression in epithelial cells. Integrin $\beta 6$ is barely detected in normal epithelia but becomes upregulated during carcinogenesis.⁴ By binding to corresponding ECM ligands, such as fibronectin and vitronectin, integrin $\beta 6$ can activate downstream pathways so as to regulate a vast array of cellular behaviors including, but not limited to, epithelial-mesenchymal transition (EMT), transforming growth factor- β (TGF- β) activation and anoikis-resistance in a tumor context.^{5–7} Thus, integrin $\beta 6$ can facilitate cell survival, proliferation, migration and tumorigenesis.

With regards to the mechanism of dysregulated integrin activity, endocytic trafficking is important for the regulation of integrin-mediated adhesion and migration in a dynamic manner by controlling the availability of integrin at the cell surface.⁸ Therefore, the assembly and disassembly of cellular adhesion contacts may be dynamically regulated via the trafficking of integrins.^{9,10} Moreover, the process of endocytosis simultaneously induces the ability of integrins to facilitate co-trafficking of growth factor receptors, referred to as intracytoplasmic “inside-in” signaling.¹¹ The clathrin-mediated endocytosis of integrin $\beta 6$, which could be regulated by HS1-associated protein X-1 (HAX-1), participates in cancer progression. The interaction of HAX-1 with integrin $\beta 6$ could determine the net signaling output of integrin $\beta 6$ on the cell surface, acting as a tight and elaborate means to regulate endocytosis of integrin $\beta 6$ and thus consequent cell motility.¹² Based on the role of endocytic trafficking in integrin $\beta 6$ -mediated migration,¹³ it is important to take endocytosis into consideration for the mechanistic study.

It has been reported that upregulation of integrin $\beta 6$ occurs in a variety of carcinomas. However, the expression and clinical significance of integrin $\beta 6$ in ESCC remain unknown. As an important regulator of endocytosis of integrin $\beta 6$, HAX-1 has been revealed to be upregulated in ESCC, which is a risk factor of lymph node metastasis.¹⁴ However, it is yet to be elucidated whether HAX-1 is involved in integrin $\beta 6$ -mediated ESCC progression; thus, the present study aimed to investigate the expression of integrin $\beta 6$ and HAX-1 in ESCC, followed by the evaluation of their correlation with various clinicopathological parameters and the prognosis of patients. The current study evaluated the prognostic value of integrin $\beta 6$ and HAX-1, thus providing a theoretically potential basis for molecular studies and possible targets for the future therapeutic interventions of ESCC.

Patients and Methods

Patients and Tissue Samples

A total of 10 pair matched frozen ESCC and adjacent non-tumor tissues were collected from patients who received radical resection at The First Affiliated Hospital of Xi'an Jiaotong University. In addition, a series of 137 paraffin-embedded samples from patients with ESCC were selected between March 2011 and September 2014, with the enrollment criteria to exclude the influence of neoadjuvant therapy on the prognosis of patients as follows: a) R0 curative surgery, b) no neoadjuvant chemotherapy or radiotherapy, c) no perioperative death, and d) no death due to other diseases rather than ESCC. The median follow-up period was 52 months (range, 8–60 months). The pathologic TNM (pTNM) staging was based on the 8th edition of the American Joint Committee on Cancer.¹⁵ The protocol was approved (approval no. 2018-G-162) by the Institutional Medical Ethics Committee of The First Affiliated Hospital of Xi'an Jiaotong University, and was in agreement with the Declaration of Helsinki. Written informed consent was obtained from the patients prior to study commencement.

Quantitative Real-Time PCR (qPCR) and Immunohistochemistry (IHC)

RNA extraction, reverse transcription and qPCR assays were performed according to our previous study.¹⁶ The mRNA expression of integrin $\beta 6$ was normalized to housekeeper β -actin for the calculation of fold change using the $2^{-\Delta\Delta Ct}$ method. For IHC staining, the slide sections were

blocked with normal goat serum after optimal epitope retrieval. Subsequently, the slides were incubated with primary monoclonal antibody against integrin $\beta 6$ (1:200; Clone 442.5C4; Sigma-Aldrich) or HAX-1 (1:500; cat. no. 166845; Santa Cruz Biotechnology, Inc.) at 4°C overnight. After incubation with horseradish peroxidase (HRP)-conjugated secondary antibody, the staining was developed using a DAB peroxidase substrate (Beyotime Institute of Biotechnology). Non-specific IgG was utilized as the negative control for primary antibody.

Assessment of Integrin $\beta 6$ and HAX-1 Immunostaining

The immunostaining of integrin $\beta 6$ and HAX-1 was mainly expressed in the cytoplasm of tumor cells. The staining levels were based on the extent and intensity assessed by two blinded observers. The staining intensity was graded as 0 (no staining), 1 (weak staining), 2 (moderate staining) or 3 (strong staining), and the staining extent was scored as 0 (0%), 1 (1–25%), 2 (26–50%), 3 (51–75%) or 4 (76–100%). The IHC score for each ESCC sample was determined by the product of staining intensity and extent grade. Given the heterogeneity of IHC scores, the optimal cut-off value was finally determined by Log-rank tests with respect to the overall survival as previously reported.¹⁷ An IHC score of 8 was used as the cut-off value for positive or negative integrin $\beta 6$ staining. As for HAX-1 immunostaining, the patients were classified into high or low expression group, with IHC score 9 as the cut-off value.

Statistics

Statistical analyses were performed using GraphPad Prism 5 software (La Jolla, CA, USA). A 2-tailed Wilcoxon matched pairs test was utilized to compare the expression levels of integrin $\beta 6$ in paired ESCC and adjacent normal tissues. Chi-square tests were used to investigate the association of integrin $\beta 6$ and HAX-1 expression levels with different clinicopathological factors. Spearman correlation analysis was used to evaluate the relationship between integrin $\beta 6$ and HAX-1 expression levels. The overall survival was analyzed via the Kaplan–Meier method with Log rank tests, with significant variables further evaluated using multivariate Cox regression analyses. $P < 0.05$ was considered to indicate a statistically significant difference.

Results

Upregulation of Integrin $\beta 6$ in ESCC

To examine whether there was a differential expression of integrin $\beta 6$ in ESCC, qPCR analysis was performed in 10 paired human samples. The mRNA expression of integrin $\beta 6$ was markedly increased in ESCC compared with matched adjacent normal tissues (Figure 1A). Subsequent IHC analysis of the same samples also demonstrated stronger immunostaining of integrin $\beta 6$ in ESCC (Figure 1B–D).

A total of 137 ESCC samples were hence enrolled to further investigate the staining of integrin $\beta 6$ and HAX-1 via IHC, both of which were predominantly detected in the cytoplasm of tumor cells. Among the ESCC specimens, 57 cases (41.6%) demonstrated positive staining for integrin $\beta 6$ (Figure 2A and B), and 65 cases (47.4%) showed high expression of HAX-1 (Figure 2C and D).

Relationship Between Integrin $\beta 6$ and Clinicopathological Factors in ESCC

The relationship between integrin $\beta 6$ and various clinicopathological factors is summarized in Table 1. There was a positive correlation between integrin $\beta 6$ and lymph node metastasis, TNM stage and histological differentiation. The clinical cases with positive integrin $\beta 6$ expression demonstrated a higher probability of lymph node metastasis compared with those with negative expression ($P < 0.001$). In addition, there was an increased number of advanced-TNM-staged ESCC samples in integrin $\beta 6$ -positive cases compared with integrin $\beta 6$ negative cases ($P < 0.001$). Specimens with positive integrin $\beta 6$ expression were more likely to exhibit poor differentiation ($P = 0.006$). More importantly, patients with positive integrin $\beta 6$ expression had worse outcomes compared with those with negative integrin $\beta 6$ expression (Figure 3A; $P < 0.001$).

Relationship Between HAX-1 and Clinicopathological Factors in ESCC

There was a positive correlation between HAX-1 expression and tumor size, pT stage, lymph node metastasis and TNM stage (Table 2). ESCC specimens with high HAX-1 expression were more likely to demonstrate late pT stage ($P < 0.001$) and lymph node metastasis ($P = 0.024$) compared with those with low HAX-1 expression. Furthermore, there was an increased number

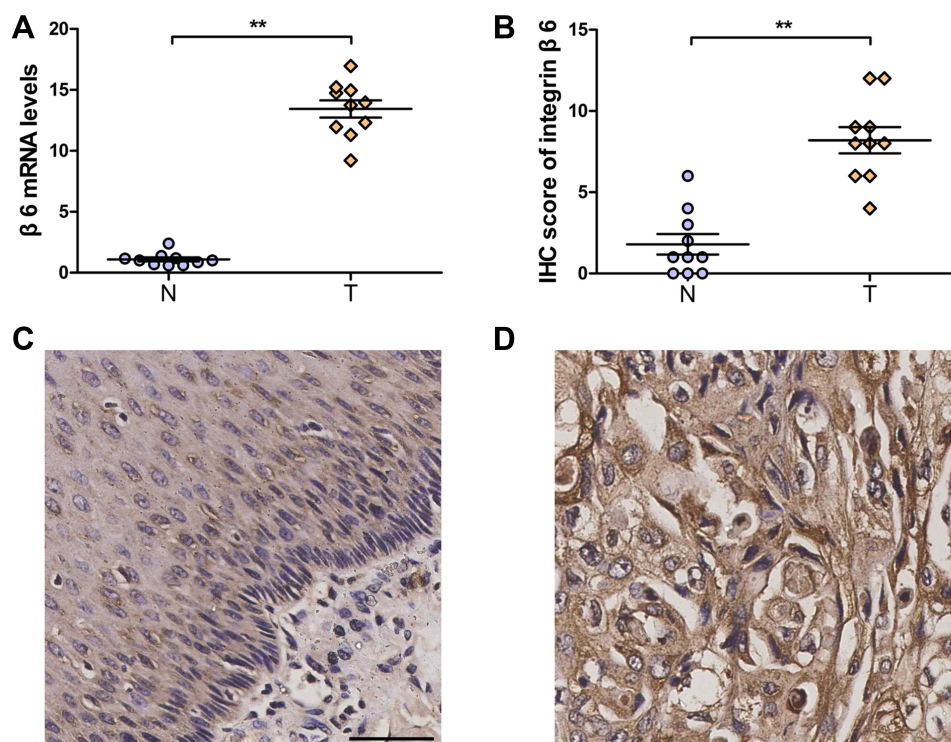


Figure 1 Upregulation of integrin $\beta 6$ in ESCC. **(A)** mRNA expression of integrin $\beta 6$ was determined via qPCR in 10 pairs of ESCC and matched adjacent healthy tissues. β -actin was used as the housekeeper gene. **(B)** IHC score demonstrated stronger immunostaining of integrin $\beta 6$ in ESCC than in adjacent normal tissues. **(A and B)** N, adjacent normal tissue; T, tumor. Statistical analysis was performed by a 2-tailed Wilcoxon matched pairs test. $P=0.002$ **(A)**; $P=0.009$ **(B)**. Data represent the mean \pm SEM. $^{**}P<0.01$. **(C and D)** Representative immunostaining of integrin $\beta 6$ in paired normal **(C)** and ESCC **(D)** tissues. Scale bar, 100 μ m.

Abbreviations: ESCC, esophageal squamous cell carcinoma; HAX-1, HSI-associated protein X-1; IHC, immunohistochemistry; qPCR, quantitative real-time PCR; SEM, standard error of mean.

of advanced-TNM-staged ESCC samples in high HAX-1 cases compared with low HAX-1 cases ($P=0.016$). It was also found that specimens with high HAX-1 staining were positively associated with tumor size ($P=0.001$). Additionally, patients with high HAX-1 expression had worse outcomes compared with those with low HAX-1 expression (Figure 3B; $P=0.004$).

Correlation of Integrin $\beta 6$ with HAX-1 in ESCC

The positive expression of integrin $\beta 6$ was observed in 64.6% of samples highly expressing HAX-1, while the ratio of samples with low expression of HAX-1 was 20.8%. Spearman analysis further demonstrated that the expression of integrin $\beta 6$ was positively correlated with HAX-1 expression (Table 2; $r=0.444$; $P<0.001$). The 137 ESCC cases enrolled in the present study were classified into four different groups based on integrin $\beta 6$ and HAX-1 expression levels for Kaplan–Meier analysis, and the results indicated that patients with positive integrin $\beta 6$ and high HAX-1 expression had

worst outcomes compared with the other groups (Figure 4; $P=0.003$).

Univariate and Multivariate Analyses for Prognosis of Patients with ESCC

To investigate the potential value of integrin $\beta 6$ and HAX-1 expression levels as predictors of poor prognosis for patient with ESCC, univariate and multivariate Cox regression analyses were performed. Univariate analysis suggested the ability of integrin $\beta 6$ ($P<0.001$) and HAX-1 ($P=0.004$) expression levels to indicate poor outcomes, as well as clinicopathological parameters such as tumor size ($P=0.032$), histological differentiation ($P=0.014$), pT stage ($P=0.025$), lymph node metastasis ($P=0.003$) and TNM stage ($P=0.017$; Table 3). Subsequent multivariate analysis demonstrated that positive integrin $\beta 6$ ($P=0.004$) and high HAX-1 expression ($P=0.011$) were independent unfavorable indicators for prognosis. In addition, pT stage ($P=0.009$) and lymph node metastasis ($P=0.001$) remained independent unfavorable prognostic indicators (Table 3).

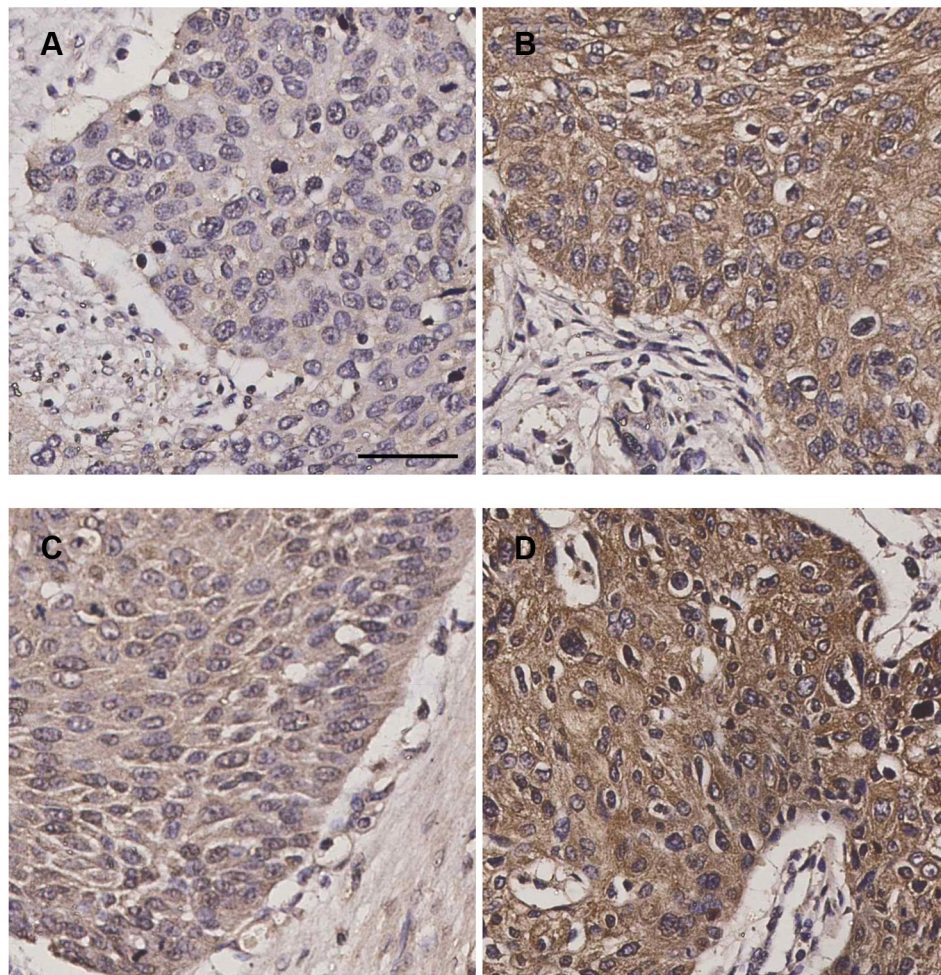


Figure 2 Immunostaining of integrin $\beta 6$ and HAX-1 in ESCC samples. **(A)** Negative expression of integrin $\beta 6$. **(B)** Positive expression of integrin $\beta 6$. **(C)** Low expression of HAX-1. **(D)** High expression of HAX-1. Scale bar, 100 μ m.

Abbreviations: ESCC, esophageal squamous cell carcinoma; HAX-1, HSI-associated protein X-1.

Discussion

Metastasis may occur at the early stage and initial diagnosis of ESCC, and along with the high incidence of post-operational recurrence, there are poor outcomes for patients with ESCC. Although pTNM stage classification is mostly utilized for survival reference, there is a great interest to discover alternative molecular biomarkers to supplement the pTNM staging system for early diagnosis and prognostic evaluation,¹⁸ which will facilitate the development of target therapy to treat highly lethal malignancy, such as ESCC. The development and progression of ESCC is a complicated process in which cancer cell adhesion, proliferation and motility mediated by integrins are involved by connecting the cytoskeleton to ECM. Different mechanisms via which integrin $\beta 6$ participates in cancer invasion and metastasis have been described in

previous studies,⁴ but a preliminary understanding of the role of integrin $\beta 6$ in ESCC is lacking.

The present study for the first time characterized the expression of integrin $\beta 6$ in ESCC, and demonstrated the mRNA expression of integrin $\beta 6$ was significantly elevated in ESCC compared with corresponding normal tissues, which was also found at the protein level using IHC. In line with its oncogenic function, integrin $\beta 6$ expression is elevated in a variety of malignancies, including human oral squamous cell carcinoma, suggesting its diagnostic value in colorectal cancer and cholangiocarcinoma.^{19–21} Furthermore, epithelial-specific integrin $\beta 6$ has been reported to be a promising target for molecular imaging of cutaneous squamous cell carcinoma, head and neck squamous cell carcinoma, lung cancer and pancreatic ductal adenocarcinoma.^{22–24} Besides epigenetic regulation,²⁵

Table 1 Correlation of the Expression of Integrin $\beta 6$ and HAX-I with Clinicopathological Features in Human ESCC

Clinicopathological Factors	n	Integrin $\beta 6$ Expression		P	HAX-I Expression		P
		Negative	Positive		Low	High	
Gender				0.982			0.563
Male	96	56	40		52	44	
Female	41	24	17		20	21	
Age (years)				0.941			0.247
<60	38	22	16		23	15	
≥ 60	99	58	41		49	50	
Tumor size (cm)				0.111			0.001
<3	59	39	20		41	18	
≥ 3	78	41	37		31	47	
Histological differentiation				0.006			0.256
Well/Moderate	101	66	35		56	45	
Poor	36	14	22		16	20	
Tumor location				0.161			0.466
Upper	28	12	16		15	13	
Middle	80	49	31		39	41	
Lower	29	19	10		18	11	
pT stage				0.790			<0.001
T1 - T2	92	53	39		64	28	
T3 - T4	45	27	18		16	29	
Lymph node metastasis				<0.001			0.024
Absent	77	65	12		47	30	
Present	60	15	45		25	35	
TNM stage				<0.001			0.016
I-II	84	61	23		51	33	
III-IV	53	19	34		21	32	

Abbreviations: ESCC, esophageal squamous cell carcinoma; HAX-I, HSI-associated protein X-I.

the marked increase of integrin $\beta 6$ expression during carcinogenesis could be induced by lysophosphatidic acid (LPA) via LPA receptor-1 coupling to $G\alpha_i$, with

subsequent activation of transcription factor Smad3 and Ets-1.¹⁹ Moreover, STAT3 is involved in the transcription of integrin $\beta 6$.²⁶ Cancer-associated fibroblasts (CAFs) in

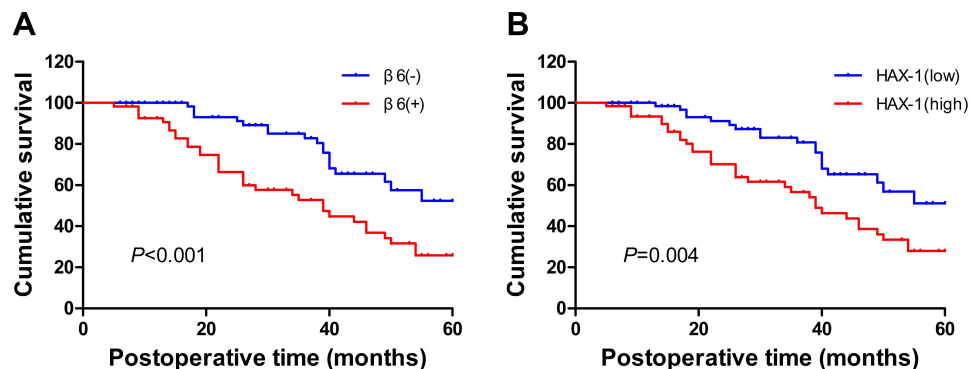


Figure 3 Association of integrin $\beta 6$ (A) and HAX-I (B) expression levels with overall survival of patients with ESCC. Statistical analysis was performed by Kaplan-Meier analytical method and compared by Log rank test. $P < 0.001$ (A); $P = 0.004$ (B).

Abbreviations: ESCC, esophageal squamous cell carcinoma; HAX-I, HSI-associated protein X-I.

Table 2 Relationship Between Integrin $\beta 6$ and HAX-1 in Human ESCC

HAX-1	Integrin $\beta 6$		Total
	Negative	Positive	
Low	57	15	72
High	23	42	65
Total	80	57	137

Abbreviations: ESCC, esophageal squamous cell carcinoma; HAX-1, HSI-associated protein X-1.

the tumor microenvironment (TME) may upregulate the expression of integrin $\beta 6$ in colon cancer via the secretion of stromal cell-derived factor-1 (SDF-1) via the SDF-1/CXCR4 axis.²⁷ Our previous study reported that interleukin-8 could increase integrin $\beta 6$ expression via the ERK/Ets-1 pathway.²⁸ In addition to the crosstalk between cancer cells and TME, integrin $\beta 6$ could also be translationally regulated by eukaryotic initiation factor 4E (eIF4E), which contributes to tumor malignancy.²⁹

The present study demonstrated that there were significant differences in histological differentiation, lymph node metastasis and TNM stage between positive and negative integrin $\beta 6$ expression groups. It was identified that integrin $\beta 6$ was an independent unfavorable indicator of prognosis for patients with ESCC. Integrin $\beta 6$ serves as a poor prognostic indicator in multiple tumors, such as hilar cholangiocarcinoma¹⁶ and breast cancer.³⁰ Integrin $\beta 6$ can also initiate a cascade of downstream signaling pathways, as oppose to a single mechanism, to promote metastasis, which may explain why it sets the tumor stage

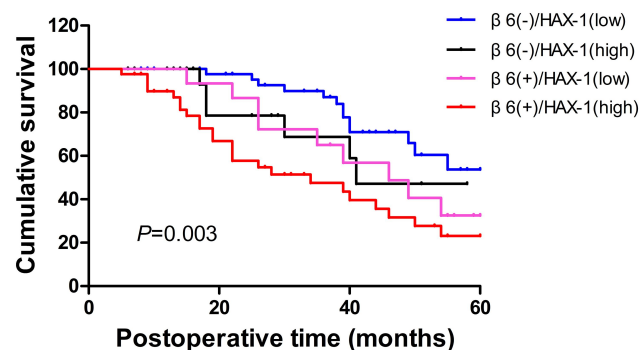


Figure 4 Association between integrin $\beta 6$ expression, HAX-1 expression and overall survival of patients with ESCC. Statistical analysis was performed by Kaplan–Meier analytical method and compared by Log rank test. $P=0.251$ ($\beta 6^{-}/HAX-1^{low}$ vs $\beta 6^{-}/HAX-1^{high}$); $P=0.094$ ($\beta 6^{-}/HAX-1^{low}$ vs $\beta 6^{+}/HAX-1^{low}$); $p<0.001$ ($\beta 6^{-}/HAX-1^{low}$ vs $\beta 6^{+}/HAX-1^{high}$); $P=0.797$ ($\beta 6^{-}/HAX-1^{high}$ vs $\beta 6^{+}/HAX-1^{low}$); $P=0.108$ ($\beta 6^{-}/HAX-1^{high}$ vs $\beta 6^{+}/HAX-1^{high}$); $P=0.215$ ($\beta 6^{+}/HAX-1^{low}$ vs $\beta 6^{+}/HAX-1^{high}$).

Abbreviations: ESCC, esophageal squamous cell carcinoma; HAX-1, HSI-associated protein X-1.

and correlates with shortened survival.³¹ In particular, integrin $\beta 6$ can induce the expression of MMP-3/9 in colon cancer cells via the ERK/Ets-1 pathway,³² contributing to invasion³³ and chemo-resistance.³⁴ Furthermore, integrin $\beta 6$ promotes EMT via the TGF- $\beta 1$ /Smad2/3 signaling pathway,³⁵ and the canonical Rho-Rac pathway may be downstream of integrin $\beta 6$ via increased expression of MMP-9/15 in HER2 amplified breast cancer.³⁶ In terms of TME, integrin $\beta 6$ on cancer cells can activate fibroblasts via the TGF- β pathway, thus forming a positive feedback and mutually promotive crosstalk with CAF.²⁷ In addition to intracellular pathway, integrin $\beta 6$ can be transferred intercellularly via prostate cancer cell-derived small extracellular vesicles to microvascular endothelial cells, which enhances angiogenesis and progression.³⁷

Disassembly of focal adhesions mediated by integrin via endocytosis is a key process for the facilitation of migration. Given the participation of abnormal integrin endocytosis in tumor progression, insights into the endocytic mechanisms by which integrin exerts its functions to promote invasiveness are vital to gain functional understanding of carcinogenesis.³⁸ Ramsay et al provided the first evidence that the interaction with HAX-1 was crucial to the endocytosis of integrin $\beta 6$ in the context of cancer.¹² As such, the present study also investigated the expression of HAX-1 in ESCC samples. Indeed, high expression of HAX-1 was observed in ESCC cells, which was associated with tumor size, lymph node metastasis, advanced pT and TNM stage, consistent with a previous report.¹⁴ HAX-1 can modulate actomyosin contractility through RhoA/septin signaling, thus promoting cell adhesion and collective migration associated with breast cancer progression.³⁹ Moreover, HAX-1 regulates proliferation, metastasis and angiogenesis of hepatocellular carcinoma cells by affecting Akt phosphorylation and FOXO3A expression.⁴⁰ Endocytic trafficking is vital for the regulation of integrin-mediated malignant behaviors of tumor cells. Endocytosis participates in the dynamic process to redistribute the pools of integrin $\beta 6$ from one part of the cell to another, allowing cancer cells to react more rapidly to adverse environmental stress.¹³ In the present study, it was found that there was a positive relationship between the expression levels of integrin $\beta 6$ and HAX-1. Furthermore, patients with positive integrin $\beta 6$ and high HAX-1 expression demonstrated worst outcomes, which is in accordance with the involvement of integrin $\beta 6$ and HAX-1 in tumor progression. It should be noted that although 7 patients enrolled in the present study received chemotherapy or

Table 3 Univariate and Multivariate Analyses of Clinicopathological Features for Overall Survival of 137 Patients with ESCC

Clinicopathological Factors	Univariate		Multivariate	
	HR (95% CI)	P value	HR (95% CI)	P value
Gender Male vs Female	0.629 (0.347–1.204)	0.125	–	–
Age (years) <60 vs ≥60	0.852 (0.451–1.524)	0.077	–	–
Tumor size (cm) <3 vs ≥3	2.824 (1.206–4.865)	0.032	1.924 (0.663–3.657)	0.279
Histological differentiation Well/Moderate vs Poor	1.455 (1.022–3.445)	0.014	1.307 (0.657–2.651)	0.310
Tumor location Upper vs Middle vs Lower	0.813 (0.322–1.730)	0.243	–	–
pT stage T1 - T2 vs T3 - T4	2.541 (1.162–5.795)	0.025	3.225 (1.426–6.245)	0.009
Lymph node metastasis Absent vs Present	4.451 (1.976–9.221)	0.003	3.347 (2.425–7.400)	0.001
TNM stage I–II vs III–IV	3.161 (1.257–7.145)	0.017	–	–
Integrin β6 expression Negative vs Positive	3.752 (1.012–6.624)	<0.001	4.133 (2.881–7.265)	0.004
HAX-1 expression Low vs High	2.315 (1.704–6.866)	0.004	1.951 (1.145–4.986)	0.011

Abbreviations: CI, confidence interval; ESCC, esophageal squamous cell carcinoma; HAX-1, HSI-associated protein X-1; HR, hazard ratio.

radiotherapy following radical surgery, adjuvant therapies were reported to fail to improve survival when compared with surgery alone and appeared less effective compared with neoadjuvant therapies.⁴¹ Hence, it is likely that adjuvant therapies would not impact the final analyses and results to a large degree. Still, additional multi-center studies with larger sample size are needed to validate the results obtained from the present study.

The discovery that integrin β6 promotes pathological processes marks it as a potential target. Indeed, integrin β6 has been investigated in therapeutic therapies, based on its role in cancer progression. The combination of integrin β6-targeted photodynamic therapy with immune check-point inhibition demonstrates enhanced anti-tumor efficacy with improved anti-tumor immunity and suppressed tumor growth.⁴² Additionally, integrin β6-targeted immunoliposomes show tumor-specific 5-fluorouracil delivery in colon cancer, offering a potential anticancer strategy to improve therapeutic efficacy.⁴³ Integrin β6-targeted adenovirus vectors also provide a promising platform for local

intraperitoneal treatment of ovarian cancer metastasis.⁴⁴ Moreover, integrin β6-specific therapy developed from foot-and-mouth-disease virus could selectively eliminate β6-positive pancreatic cancer.⁴⁵ Whilding et al reported the clinical evaluation of integrin β6 re-targeted chimeric antigen receptor T cell immunotherapy for tumors that express this integrin, providing a promising new molecular-specific therapy.⁴⁶ Therefore, in the future mechanistic studies would be performed to further examine the role of HAX-1 in integrin β6-mediated ESCC progression, and evaluate the potential of HAX-1 and/or integrin β6-targeted therapy in the translation into clinical benefits.

In conclusion, the present study identified the upregulation of integrin β6 and its relevance to HAX-1, as well as unfavorable clinicopathological factors in ESCC. The current results further defined the predictive value of integrin β6 and HAX-1 as independent indicators of poor prognosis for patients with ESCC, both of which may contribute to tumor proliferation and metastasis, thus promoting ESCC progression. Therefore, combined targeting of integrin β6

and HAX-1 are required for future investigation, which may provide a potential novel approach for the treatment of ESCC.

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Disclosure

The authors declare no conflict of interest in this work.

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