

High DEPTOR expression correlates with poor prognosis in patients with esophageal squamous cell carcinoma

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Objective: The disheveled, Egl-10, and pleckstrin (DEP) domain containing mammalian target of rapamycin (mTOR)-interacting protein (DEPTOR) is a binding protein containing mTOR complex 1 (mTORC1), mTOR complex 2 (mTORC2), and an endogenous mTOR inhibitor. DEPTOR shows abnormal expressions in numerous types of solid tumors. However, how DEPTOR is expressed in esophageal squamous cell carcinoma (ESCC) remains elusive.

Methods: The expression of DEPTOR in 220 cases of ESCC and non-cancerous adjacent tissues was detected by immunohistochemistry. DEPTOR levels in ESCC and paired normal tissue were quantified using reverse transcription-polymerase chain reaction and Western blot analysis to verify the immunohistochemical results. The relationship between DEPTOR expression and the clinicopathological features of ESCC was analyzed based on the results of immunohistochemistry. Finally, we analyzed the relationship between DEPTOR expression and the prognosis of patients with ESCC.

Results: Immunohistochemical staining showed that the expression rate of DEPTOR in ESCC tissues was significantly increased. *DEPTOR* mRNA and protein expression was significantly higher in ESCC tissues than in normal adjacent esophageal squamous tissues. High DEPTOR expression was significantly correlated with regional lymph node status in the TNM stage of patients with ESCC. Kaplan–Meier survival curves showed that the rate of overall survival was significantly lower in patients with high DEPTOR expression than in those with low DEPTOR expression. Additionally, high DEPTOR expression was an independent prognostic predictor for ESCC patients.

Conclusion: High DEPTOR expression is an independent prognostic biomarker indicating a worse prognosis for patients with ESCC.

Keywords: DEPTOR, ESCC, prognosis

Introduction

Esophagectomy remains the primary treatment for patients with esophageal squamous cell carcinoma (ESCC). Although the survival rates have improved over the past decade, the 5-year survival rate is still only 30%.¹ Most tumor recurrences occur within the first year after surgery and are typically fatal. However, patients surviving for 5 years have approximately the same life expectancy as the general population.^{1,2} The post-operative recurrence, invasion, and metastasis of ESCC are affected by many closely linked factors such as proto-oncogene activation, tumor-suppressor gene inactivation, and abnormal protein expression.

Previous studies have shown that the mammalian target of rapamycin (mTOR) channel plays an important role in the regulation of protein synthesis and is activated in many human cancers.^{3,4} Therefore, the mTOR inhibitor, a rapamycin derivative, is

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used to inhibit the transcription of cancer gene(s), the growth of cancer, and the generation of the tumor vasculature caused by abnormal signal channels, and has been examined in recent studies.⁵⁻⁷ Some scholars have suggested that a coherent understanding of the mTOR network is imperative, as its dysregulation has been implicated in diverse pathologies including cancer, diabetes, autism, and aging.⁸ In addition, a study has shown that disheveled, Egl-10, and pleckstrin (DEP) domain containing mammalian target of rapamycin (mTOR)-interacting protein (DEPTOR) abnormal expression may be associated with resistance to chemotherapy in breast cancer patients; these authors also highlighted the dichotomous functions of DEPTOR in modulating the proliferation and survival of triple-negative breast cancers during metastasis.⁹ DEPTOR is a binding protein containing mTOR complex 1 (mTORC1), mTOR complex 2 (mTORC2), and an endogenous mTOR inhibitor. DEPTOR shows abnormal expression in numerous types of solid tumors.² However, how DEPTOR is expressed in ESCC remains unclear.

Materials and methods

Patients and specimens

ESCC was diagnosed in patients, who were enrolled in the current study, at the Nanfang Hospital and The Third Affiliated Hospital of the Southern Medical University, Guangzhou, People's Republic of China from 2004 to 2008. Patients included in the current study had undergone esophageal cancer resection. Pathological review confirmed ESCC in resected tumors and carcinoma-adjacent non-tumorous esophageal tissues. All samples were obtained from the two participating institutions, and were identified using anonymous codes that met the requirements of Nanfang Hospital of Southern Medical University and The Third Affiliated Hospital of Southern Medical University's guidelines, as formulated based on the Declaration of Helsinki. Patients were required to have provided written informed consent, and the Review Board of the Southern Medical University approved the protocol. The American Joint Committee on Cancer/Union for International Cancer Control (AJCC/UICC) classification guidelines were used as criteria to divide the ESCC specimens into different phases, while World Health Organization (WHO) criteria were applied for subtyping the grading and histopathology of ESCC specimens. ESCC biopsy samples and matching non-cancerous esophageal tissues were frozen in liquid nitrogen and stored.

Immunohistochemistry

Immunohistochemical analysis using streptomycin avidin-biotin peroxidase complex (SP) method was conducted to

evaluate paraffin sections. The immunohistochemical staining procedure was conducted according to the manufacturer's instructions, mainly including paraffin sectioning, regular dewaxing to water, antigen retrieval, 3% hydrogen peroxide blocking of endogenous peroxidase, goat serum closing, rabbit polyclonal antibody against human DEPTOR (1:200; BS-8255R; Beijing Biosynthesis Biotechnology Co, Ltd, Beijing, People's Republic of China), secondary antibodies, SP, 3,3'-Diaminobenzidine (DAB) staining, and a contrast dye seal. Biopsies of known normal tissues were used as positive controls, while phosphate-buffered saline (PBS) was substituted for primary antibodies in the negative controls.

A single-blind study was conducted to read the biopsies (pathologists were blinded to the sample information). Five medium views were randomly selected (200 times), and 200 cancer cells were counted in every view; thus, a total of 1,000 cells were counted and were divided into the following types according to the staining intensities: score 0, no staining; score 1, light yellow staining; score 2, brown yellow staining; and score 3, sepia staining. Samples were classified as 0%–100% according to the proportion of cells with positive cell nuclei. In the present study, the product >0 from the staining intensity multiplied by the positive proportion was defined as DEPTOR-positive; otherwise, the sample was considered to be negative. In addition, DEPTOR staining intensities with scores of 0 and 1 were defined as having low expression of DEPTOR, while those with scores of 2 and 3 were defined as having high expression.

Reverse transcription-polymerase chain reaction

Trizol reagent was used to extract total RNA from the cells, and absorbance was measured at 260 nm and 280 nm. RNA content was also determined. Photos were taken after 1% Sepharose gel electrophoresis. Reverse transcription of the mRNA was conducted according to the reverse transcription-polymerase chain reaction (RT-PCR) guidance manual to produce cDNA, which was then used as a template for PCR amplification for 30 cycles under the following PCR conditions: 94°C for 5 minutes; 94°C for 30 seconds; 56°C for 30 seconds; and 72°C for 40 seconds. Final extension was conducted at 72°C for 5 minutes. The primer sequences of DEPTOR were as follows: forward, 5'-TTTGTGGTGCGAGGAAGTAA-3'; reverse, 5'-CATTGCTTTGTGTCATTCTGG-3'. The amplified products were separated by electrophoresis, and the gel image was photographed under ultraviolet light. Quantity One® software was used to analyze the electrophoresis banding gray values

with glyceraldehyde 3-phosphate dehydrogenase (GAPDH) as the standardized internal reference.

Western blot analysis

Protein was extracted from the tissue lysate, and the bicinchoninic acid assay (BCA) method was used to measure protein concentration. Proteins were separated by 15% sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE), were transferred to a membrane, and were blocked with 5% skim milk for 1 hour. DEPTOR and GAPDH primary antibodies were incubated with the membranes overnight, after which the membranes were washed, and immunoglobulin G (IgG)/biotin IgG secondary antibodies were incubated with the membrane for 2 hours. After incubation, the membrane was washed three times with mixture of Tris-buffered saline and Tween 20 (TBST). Finally, the electrogenerated chemiluminescence (ECL) method was used to detect the protein bands. A Gel-Pro analyzer was used to analyze band absorbance, with GAPDH used as the standardized internal reference.

Statistical methods

To study the relationship between DEPTOR expression and patient characteristics including age, sex, tumor stage, TNM

classification, histological differentiation, and vital status, we applied Pearson's chi-square test and Spearman's correlation analysis. The Kaplan–Meier method was used to draw the survival curves for patients with both DEPTOR-high and DEPTOR-low expression, while the log-rank test was used to compare statistical differences. Overall survival (OS) was calculated from the date of surgery to the date of death or last follow-up. Cox regression analysis was conducted to analyze univariable and multivariable survival. *P*-values <0.05 were regarded as statistically significant. SPSS version 13.0 (SPSS Inc, Chicago, IL, USA) software was used for statistical analyses.

Results

DEPTOR expression in ESCC and adjacent tissues

DEPTOR expression was detected in 220 cases of ESCC and in corresponding adjacent tissue samples by immunohistochemistry. Immunohistochemical staining showed that DEPTOR was predominantly localized in the nucleus or cytoplasm of cancer and normal esophageal squamous cells (Figure 1). Compared with normal adjacent esophageal squamous tissues, the high expression rate of DEPTOR in ESCC tissues was significantly increased ($P=0.002$).

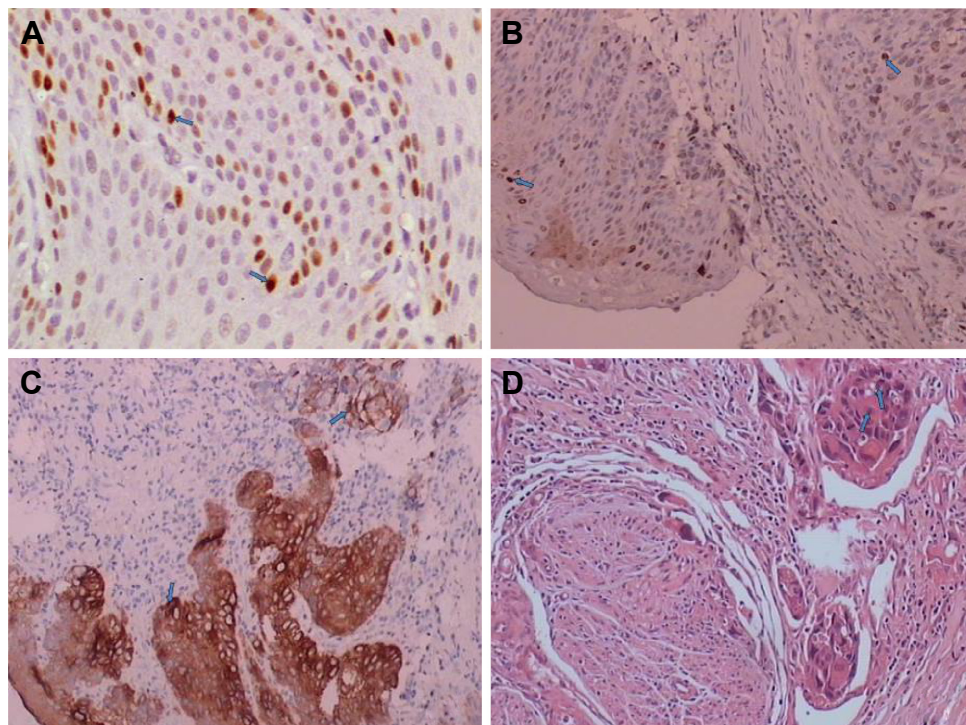


Figure 1 DEPTOR expression in ESCC and adjacent tissues detected by immunohistochemistry.

Notes: (A) Normal adjacent tissue, SP $\times 100$; (B) high differentiation, SP $\times 50$; (C) moderate differentiation SP $\times 50$; (D) poor differentiation, SP $\times 50$. Blue arrows indicate DEPTOR protein expression in ESCC or adjacent tissues.

Abbreviations: DEPTOR, the disheveled, Egl-10, and pleckstrin (DEP) domain containing mammalian target of rapamycin (mTOR)-interacting protein; ESCC, esophageal squamous cell carcinoma; SP, streptomycin avidin-biotin peroxidase complex.

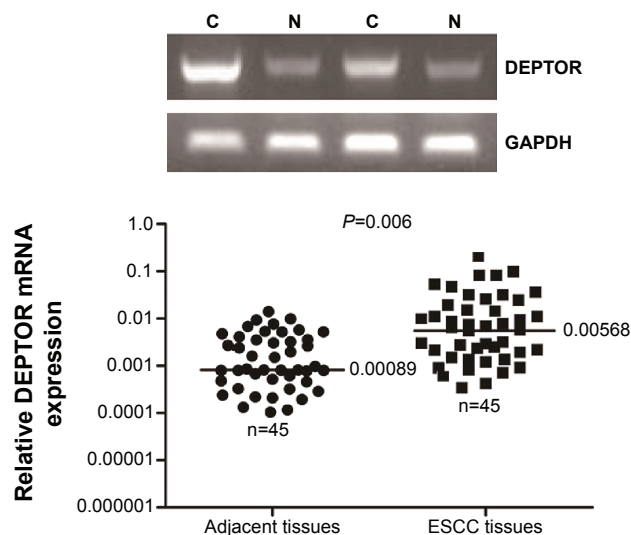


Figure 2 RT-PCR analyses of DEPTOR expression in 45 cases of ESCC and in corresponding normal adjacent tissues.

Notes: mRNA was detected in ESCC tissues (C columns) and adjacent tissues (N columns). mRNA expression was significantly higher in ESCC tissues than in normal adjacent esophageal squamous tissues.

Abbreviations: DEPTOR, the disheveled, Egl-10, and pleckstrin (DEP) domain containing mammalian target of rapamycin (mTOR)-interacting protein; ESCC, esophageal squamous cell carcinoma; mRNA, messenger ribonucleic acid; RT-PCR, reverse transcription-polymerase chain reaction; GAPDH, glyceraldehyde 3-phosphate dehydrogenase.

To confirm this finding, we investigated DEPTOR expression in 45 cases of ESCC and in corresponding normal adjacent tissues using RT-PCR and Western blot analysis. The results indicated that *DEPTOR* mRNA and protein expression was significantly higher in ESCC tissues than in normal adjacent esophageal squamous tissues ($P=0.0006$ and $P=0.009$, respectively; Figures 2 and 3, respectively).

DEPTOR expression and clinicopathological characteristics of ESCC

We analyzed the relationship between the expression of DEPTOR and clinicopathological factors of patients with ESCC based on immunohistochemical results. High DEPTOR expression was significantly correlated with regional lymph node status in the TNM stage of patients ($P<0.05$). However, high DEPTOR expression was not significantly correlated with other clinical parameters including age, sex, tumor size, degree of differentiation, location, smoking history, and radio-chemotherapy ($P>0.05$) (Table 1).

DEPTOR-high expression as a prognostic marker for ESCC

In the current study, the primary outcome was the OS of patients. In order to investigate the relationship between DEPTOR expression and patient prognosis, we plotted OS

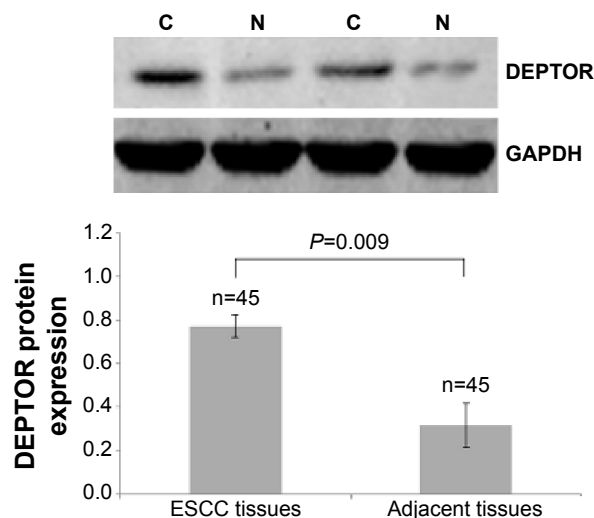


Figure 3 Western blot analyses of DEPTOR expression in 45 cases of ESCC and in corresponding normal adjacent tissues.

Notes: DEPTOR protein was detected in ESCC tissues (C columns) and adjacent tissues (N columns). Protein expression was significantly higher in ESCC tissues than in normal adjacent esophageal squamous tissues.

Abbreviations: DEPTOR, the disheveled, Egl-10, and pleckstrin (DEP) domain containing mammalian target of rapamycin (mTOR)-interacting protein; ESCC, esophageal squamous cell carcinoma; GAPDH, glyceraldehyde 3-phosphate dehydrogenase.

curves using the Kaplan–Meier method. Kaplan–Meier survival curves showed that the rate of OS was significantly lower in patients with high DEPTOR expression than in those with low DEPTOR expression ($P<0.001$; Figure 4). Subgroup analysis showed the same results (Figure 4). This indicates that patients with high DEPTOR expression had significantly shorter OS than those with low expression. We performed univariate and multivariate analysis using the Cox regression model. In addition to lymph node metastasis and TNM stage, high DEPTOR expression was also an independent prognostic predictor for ESCC patients (Table 2).

Discussion

mTOR is a protein kinase that interacts with rapamycin in mammalian cells.¹⁰ It is composed of 2,549 amino acids and has a relative molecular mass of 289 KDa.¹¹ mTOR consists of two isoforms, mTORC1 and mTORC2, which have different functions because of their different subunits.¹² DEPTOR is an important constituent of both of these isoforms.¹³ The mTOR signal transduction pathway can inhibit cancer cell apoptosis, maintain cell survival, and promote tumor neovascularization.^{7,14} It plays an important role in the generation and metastasis of malignant tumors, and thus is a well-studied signaling pathway for targeted therapy. For example, both Torisel® and everolimus are derivatives of rapamycin and can reduce the activity of mTOR while working in conjunction with a drug that prohibits the proliferation of

Table 1 Correlation between DEPTOR expression and clinicopathological factors of ESCC

Variable	Cases (n)	DEPTOR expression		χ^2	P-value
		Low	High		
Overall frequency				9.627	0.002
ESCC tissues	220	114	106		
Adjacent tissues	220	146	74		
Age (years)				0.672	0.412
≥ 60	112	55	57		
< 60	108	59	49		
Sex				0.669	0.792
Female	79	40	39		
Male	141	74	67		
Smoking history				0.530	0.466
Yes	78	43	35		
No	142	71	71		
Differentiation				2.330	0.312
Well	85	39	46		
Moderate	68	36	32		
Poor	67	39	28		
Location				0.240	0.887
Upper	46	23	23		
Middle	119	61	58		
Lower	55	30	25		
T stage				1.419	0.234
T1 + T2	105	50	55		
T3 + T4	115	64	51		
Lymph node metastasis				6.847	0.009
N0	101	62	39		
N1 + N2	119	52	67		
TNM				8.856	0.003
I + II	150	88	62		
III + IV	70	26	44		
Radio-chemotherapy				1.650	0.199
Yes	126	70	56		
No	94	44	50		

Abbreviations: DEPTOR, the disheveled, Egl-10, and pleckstrin (DEP) domain containing mammalian target of rapamycin (mTOR)-interacting protein; ESCC, esophageal squamous cell carcinoma.

breast cancer cells. This treatment is viable for patients with advanced and/or metastatic breast cancer.¹⁵

However, some studies have shown that DEPTOR was highly expressed in some solid tumors, possibly because DEPTOR not only inhibited the activity of mTORC1, but also reduced the negative feedback loop that inhibits the function of mTORC1 on phosphatidylinositol 3 kinase (PI3K) through inhibiting ribosomal protein S6 kinase 1 (S6K1) to activate protein kinase B (Akt).¹⁶⁻¹⁸ Activated Akt factors may regulate many molecular families involved in cell apoptosis, inhibit tumor cell apoptosis, and disrupt the balance between cell proliferation and apoptosis in tissues to maintain cell survival.¹⁹ Moreover, active PI3K and Akt may also upregulate HIF1 α through other channels to promote the expression of vascular endothelial growth factor, which stimulates the generation of tumor vessels, thus providing

sufficient nutrition to the tumor.^{17,20} However, the inhibition of DEPTOR expression may also promote apoptosis. Therefore, overexpression of DEPTOR may promote the generation and development of some malignant tumors. However, the role of DEPTOR in esophageal cancer or the relationship between its expression and the generation, development, and prognosis of esophageal cancer remains unclear.

In the present study, DEPTOR was identified in ESCC tissues and normal esophageal mucosa by immunohistochemistry. We found that the rate of DEPTOR-high expression was significantly higher in ESCC tissues than in corresponding adjacent tissues. To verify this result, DEPTOR was detected on the mRNA level by RT-PCR analysis. The mRNA levels of DEPTOR in ESCC tissues were significantly higher than those in normal esophageal tissues. Additionally, we investigated the relationship between DEPTOR expression

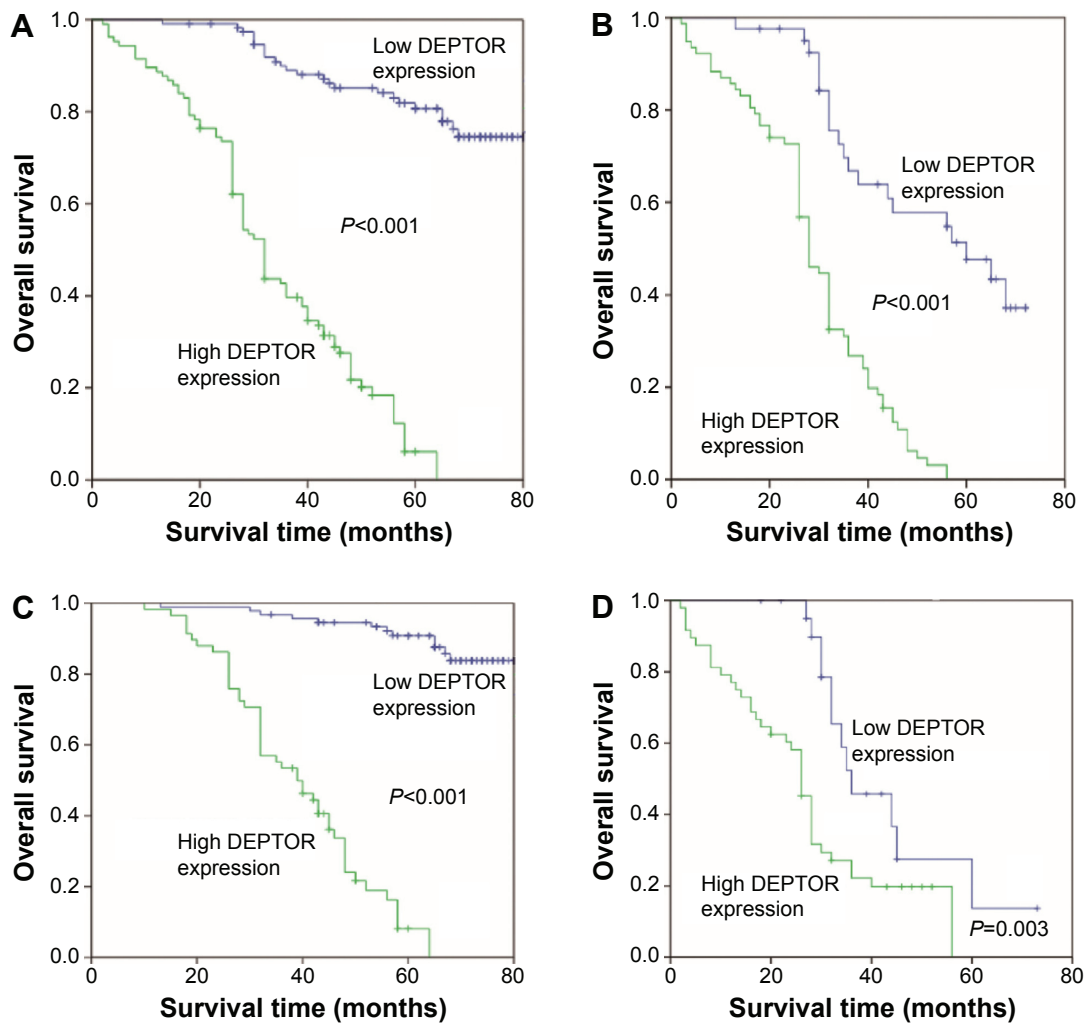


Figure 4 Kaplan-Meier analysis showing the OS of patients with ESCC.

Notes: (A) All patients; (B) patients with stages I-II; (C) patients with stages III-IV; (D) patients with lymph node metastasis.

Abbreviations: DEPTOR, the disheveled, Egl-10, and pleckstrin (DEP) domain containing mammalian target of rapamycin (mTOR)-interacting protein; OS, overall survival; ESCC, esophageal squamous cell carcinoma.

levels and clinicopathological parameters of ESCC patients. The results showed that DEPTOR expression was significantly correlated with lymph node status and TNM stage, but not with age, sex, differentiation, location, and T stage

of patients with ESCC. Through long-term follow-up, the survival rates of the ESCC patients with high DEPTOR expression were significantly lower than in patients with low DEPTOR expression. Univariate analysis showed

Table 2 Univariate and multivariate analysis of the prognosis for ESCC

Variable	Univariate analyses	P-value	Multivariate analyses	P-value
	HR (95% CI)		HR (95% CI)	
Age	0.982 (0.665–1.450)	0.925		
Sex	0.957 (0.653–1.403)	0.822		
Differentiation	0.895 (0.596–1.345)	0.595		
Location	0.910 (0.689–1.201)	0.505		
T stage	0.853 (0.565–1.287)	0.447		
Lymph node metastasis	6.229 (4.487–8.646)	<0.001	1.796 (1.271–2.538)	0.001
TNM	11.185 (6.660–18.783)	<0.001	4.846 (2.902–8.091)	<0.001
High DEPTOR expression	6.410 (4.156–9.888)	<0.001	1.381 (1.024–1.862)	0.034

Abbreviations: DEPTOR, the disheveled, Egl-10, and pleckstrin (DEP) domain containing mammalian target of rapamycin (mTOR)-interacting protein; ESCC, esophageal squamous cell carcinoma; HR, hazard ratio; CI, confidence interval.

that DEPTOR expression, lymph node status, and TNM stage were correlated with the OS of patients with ESCC. Multivariate analysis further revealed that high DEPTOR expression independently predicted unfavorable OS of ESCC patients.

Conclusion

In conclusion, high DEPTOR expression is an independent prognostic biomarker that indicates a worse prognosis for patients with ESCC, and the specific mechanisms require further analysis.

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Disclosure

The authors report no conflicts of interest in this work.

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