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HER3 overexpression is a prognostic indicator of extrahepatic cholangiocarcinoma

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Abstract

Members of the HER (ERBB) receptor protein tyrosine kinase family play an important role in regulating cellular division, proliferation, differentiation, and migration and have prognostic significance in a number of cancers. Here, we sought to define their role in extrahepatic cholangiocarcinoma (EHCC). HER2 and HER3 protein expression was studied in 230 EHCC cases using a tissue microarray and compared with clinicopathological variables, including the survival of EHCC patients. HER3 was predominantly localized to the cytoplasm, whereas HER2 exhibited a membranous expression pattern. Overexpression of HER2 and HER3 was observed in 6 % (13/224) and 39 % (90/230) of EHCCs, respectively. Membranous HER2 overexpression occurred more frequently in intraductal papillary neoplasms with an associated invasive carcinoma than in tubular adenocarcinomas ($P=0.02$). HER3 protein was more commonly overexpressed in nodular and infiltrative than in papillary tumors ($P=0.03$). HER3 overexpression was associated with decreased survival in both univariate ($P=0.01$) and multivariate ($P=0.008$) analyses, whereas HER2 overexpression was not associated with survival. HER2 and HER3 are overexpressed in subsets of EHCC patients. Notably, HER3 overexpression is correlated with decreased patient survival, suggesting that HER3 constitutes a prognostic factor as well as a potential candidate for targeted therapy.

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Conflict of interest The authors declare that they have no conflict of interest.

Keywords

Extrahepatic; Bile duct; Cholangiocarcinoma; HER2; HER3; Immunohistochemistry; Prognosis

Introduction

Extrahepatic cholangiocarcinoma (EHCC) is a malignant epithelial neoplasm of the biliary tract, from the hepatic hilum to the distal bile duct, that accounts for 70–90 % of all cholangiocarcinomas [1]. EHCC is a relatively uncommon cancer in Western countries but is more prevalent in Eastern Asian countries, including Korea [1–3]. Surgical resection of the tumor is the only curative therapeutic modality for patients with EHCC but can be applied only in a limited number of patients with localized or locally advanced disease [4]. The 5-year survival rate of EHCC patients following surgical resection is approximately 20 % [5]. Several neoadjuvant therapies, including chemotherapy, radiation therapy, and photodynamic therapy, have been extensively studied, but none have yielded significant survival advantages for EHCC patients [1, 6]. Therefore, identification of new biomarkers for early detection and/or development of new therapeutic regimens based on a better understanding of the biological mechanisms of this deadly disease are essential for decreasing mortality among EHCC patients.

Epidermal growth factor receptors (EGFRs) are members of a membrane receptor protein tyrosine kinase family that includes HER1 (ErbB1/EGFR), HER2 (ErbB2), HER3 (ErbB3), and HER4 (ErbB4). After binding ligands, EGFRs are activated by receptor homo- or heterodimerization, resulting in the phosphorylation of various downstream substrates, which mediate intracellular signal transduction. EGFR signaling pathways play an important role in regulating cell division, proliferation, differentiation, and migration [7]. The EGFR family member HER3 lacks intrinsic protein tyrosine kinase activity [8]. However, binding of neuregulin family ligands enables HER3 to form functional heterodimers with EGFR or HER2 that act through the cytoplasmic domain of HER3 to exert tyrosine phosphorylation activity [9]. Thus, by heterodimerizing with other EGFR proteins, HER3 can participate in diverse signal transduction cascades and regulate various cellular processes [7, 9]. The development of tyrosine kinase inhibitors of EGFR or HER2 was initially considered a promising strategy for blocking EGFR or HER2 pathways in several solid cancers, including breast and lung cancers. However, evidence suggests that HER3 may be responsible for the resistance to these therapeutic regimens, which often develops [10]. Overexpression of HER3 has been reported in several malignant neoplasms in gastrointestinal tract organs, including gastric [11–13], colorectal [14–19], and ampulla of Vater [18] cancers. Although a few studies have demonstrated HER2 expression in cholangiocarcinoma, showing that it is relatively infrequent in these tumors [20–23], there have been no comprehensive studies of HER2 and HER3 expression in EHCCs. In the current study, we analyzed HER2 and HER3 protein expression in 230 EHCC cases using tissue microarray immunohistochemistry. We demonstrate that HER3 expression in EHCC constitutes a prognostic factor.

Materials and methods

Patients and tumor samples

The study population consisted of 230 patients with surgically resected EHCCs treated at Asan Medical Center, University of Ulsan College of Medicine, in Seoul, South Korea, between 1991 and 2005. Carcinomas with an epicenter in the extrahepatic bile duct were included in this study, whereas carcinomas from the ampulla of Vater or pancreas were not. Carcinomas arising in the gallbladder or intrahepatic bile duct with extension to the extrahepatic bile duct were also excluded. Information regarding the age and gender of patients, surgical procedure, survival time, and survival status was obtained by reviewing EHCC patients' medical records. Data pertinent to location, size, and growth pattern of the tumor were obtained from surgical pathology reports. Materials were obtained with appropriate human protection approval from the Institutional Review Board of the Asan Medical Center (Project Number 2011–0734). Information regarding post-operative radiation, chemotherapy, and performance status of patients was not analyzed in this study.

Tissue microarray construction

Tissue microarrays were constructed from archival formalin-fixed, paraffin-embedded tissue blocks as previously described [24]. Briefly, 230 extrahepatic cholangiocarcinoma cases and 12 normal extrahepatic bile ducts were included. A representative area was carefully selected for each tumor or normal biliary epithelia from a hematoxylin-and-eosin-stained section of a donor block. Each case was represented by two to four 1.5-mm-diameter cores.

Immunohistochemistry and scoring

Immunohistochemistry was performed on 4- μ m-thick tissue microarray sections as previously described [25]. Briefly, tissue sections were deparaffinized and hydrated in xylene and serially diluted ethanol, respectively. Endogenous peroxidase was blocked by incubation in 3 % H₂O₂ for 10 min. Antigen retrieval was performed in a steam pressure cooker using preheated antigen retrieval buffer, pH 6 (Dako, Glostrup, Denmark), at 95 °C for 10 min. Non-specific binding of antibodies was minimized by incubating sections with Protein Block (Dako) for 15 min. Microarrays were incubated at room temperature for 30 min with rabbit polyclonal anti-HER2 (c-erbB2, A0485; 1:750 dilution; Dako) or overnight at 4 °C with mouse monoclonal anti-HER3 (RTJ.2, 1:500 dilution; Santa Cruz, CA, USA). Antigen–antibody reactions were detected with an LSAB+peroxidase kit (Dako) and 3,3'-diaminobenzidine (Dako). Negative controls were composed of identically treated histologic sections with the omission of primary antibodies. Immunostained sections were lightly counterstained with hematoxylin, dehydrated in ethanol, and cleared in xylene. We did not perform immunohistochemistry for Her2 and Her3 on full sections. Tissue core representativity might be a concern, but certainly for 1.5-mm cores previous studies have demonstrated that two to four tissue cores are representative with 95–97 % concordance rate [62].

HER2 protein expression was scored on a scale of 0 to 3 using the following gastric cancer staging system [26]: 0, no reactivity or membranous reactivity in less than 10 % of cells; 1+, faint/barely perceptible membranous reactivity in 10 % of cells or reactivity in only part of

the cell membrane; 2+, weak to moderate complete or basolateral membranous reactivity in 10 % of tumor cells; 3+, strong complete or basolateral membranous reactivity in 10 % of tumor cells. Cases receiving a HER2 score of 3+ were considered positive for HER2 expression.

The results of immunohistochemical staining for HER3 were scored based on the intensity of staining as 0 (negative), 1 (weak), or 2 (strong), and centage of positive epithelial cells as 0 (<5 %), 1 (6–25 %), 2 (26–50 %), 3 (51–75 %), or 4 (>76 %). A Histo-score was generated as the product of intensity and area. The Histo-score was then dichotomized into no/lower expression (Histo-score, 0–6) and overexpression (Histo-score, 8).

Statistical analysis

Statistical analyses were performed using SPSS version 17 (SPSS Inc., Chicago, IL, USA). Associations between categorical variables were examined using Pearson's chi-square and Fisher's exact tests. Survival curves were calculated by the Kaplan–Meier method, and statistical significance was evaluated using the log-rank test and the Cox proportional hazards regression model. $P < 0.05$ was considered as statistically significant.

Results

Clinicopathological characteristics of cases

The clinicopathological characteristics of cases are summarized in Table 1. Patient age ranged from 30 to 84 years (mean, 60.9 years). Of the 230 patients, 164 were men and 66 were women. Tumor size ranged from 0.5 to 6 cm (mean, 2.5 cm). Thirty-one cases were pT1 tumors, 84 were pT2, 91 were pT3, and 24 were pT4. The length of patient follow-up time ranged from 1 to 127 months, and median survival time at last follow-up was 22 months.

HER2 and HER3 expression

HER2 protein expression patterns were analyzed in tumors from a total of 224 EHCC patients using tissue microarray immunohistochemistry. Information regarding six patients was excluded from the HER2 analysis because their tissue microarray cores were either detached or folded during the HER2 immunohistochemical staining procedure. Representative images of the expression of HER2 and HER3 in EHCC tumors and normal biliary epithelial cells are shown in Fig. 1. All 12 samples of normal biliary epithelial cells did not show immunoreactivity for either HER2 or HER3. HER3 was predominantly distributed in a cytoplasmic pattern, whereas HER2 was mainly localized to the cell membrane in EHCC cases. HER2 and HER3 overexpression was observed in 6 % (13/224 cases) and 39 % (90/230 cases) of EHCCs, respectively. Only the histological subtype was significantly associated with HER2 overexpression ($P = 0.02$). Specifically, intraductal papillary neoplasms with associated invasive carcinoma cases showed significantly more HER2 overexpression than other histological subtypes, including tubular adenocarcinoma, adenosquamous carcinoma, intestinal type adenocarcinoma, and mucinous carcinoma, which did not express HER2 (Table 2). HER3 protein was more commonly overexpressed in

nodular and infiltrative tumors than in papillary tumors ($P=0.03$). HER3 overexpression was more frequent in cases without than in cases with invasion of the pancreas ($P=0.02$).

Among 13 HER2-overexpressing EHCCs, seven (54 %) showed HER3 overexpression. There was a significant correlation between cases with HER2 overexpression and those with HER3 overexpression ($R=0.24$, $P<0.001$) but not between HER2 and HER3 co-overexpression and clinicopathological parameters (Table 3).

Survival analysis

Median survival of patients with HER3 overexpression (24.0 months) was significantly worse than that of patients with no/lower HER3 expression (31.9 months; $P=0.01$, log-rank test; Fig. 2a). The 1-, 3-, and 5-year survival rates in the HER3-overexpression group were 82.2, 22.6, and 13.9 %, respectively, whereas the corresponding rates in the no/lower HER3-expression group were 82.6, 40.5, and 31.2 %.

There was no significant difference in survival between patients with HER2 overexpression (median survival, 25.9 months) and those without HER2 expression (median survival, 27.8 months; $P=0.17$, log-rank test; Fig. 2b).

Median survival of EHCC patients with both HER2 and HER3 overexpression (24.8 months) tended to be lower than that in patients without HER2 and HER3 co-overexpression (28.0 months; $P=0.08$; Fig. 2c), but this did not reach statistical significance.

Association between survival and other clinicopathological factors

The relationships between other clinicopathological variables and survival are summarized in Table 4. Of these additional clinicopathological variables, tumor differentiation status ($P<0.001$), pT classification ($P=0.01$), lymph node metastasis ($P<0.001$), liver invasion ($P=0.01$), and vascular invasion ($P=0.02$) were also significantly associated with survival. In contrast, survival was not associated with gender ($P=0.38$), tumor location ($P=0.47$), growth pattern ($P=0.09$), pancreatic invasion ($P=0.24$), duodenal invasion ($P=0.07$), perineural invasion ($P=0.36$), or resection margin status ($P=0.10$).

Multivariate analysis of clinicopathological factors

The independent prognostic significance of HER3 expression as well as of other clinicopathological parameters was determined by applying the Cox proportional hazards model (Table 5). HER3 overexpression ($P=0.007$), differentiation ($P=0.002$), liver invasion ($P=0.03$), and lymph node metastasis ($P=0.002$) remained as prognostic factors.

Discussion

In this study, we observed overexpression of HER2 and HER3 protein in a subset of EHCCs, with about 6 % of EHCC cases overexpressing HER2 (13/224) and 39 % (90/230) overexpressing HER3. In addition, we found HER3 overexpression to be an independent prognostic factor in EHCC patients. HER3 is unique among members of the HER receptor tyrosine kinase family in that it lacks intrinsic tyrosine kinase activity but it does contain six consensus phosphotyrosine sites, which bind the SH2 domain of the three regulatory

subunits of phosphoinositide-3-kinase (PI3K) [27, 28]. Binding of HER3 can activate the PI3K/Akt signaling pathway, which is a critical regulator of many cellular processes. HER2-mediated transformation of epithelial cells is associated with the activation of the PI3K/Akt signaling pathway in breast cancer. Several targeted therapeutics, including trastuzumab, a monoclonal antibody against the extracellular domain of HER2, and lapatinib, an EGFR/HER2 tyrosine kinase inhibitor, have been approved for use in the treatment of HER2-overexpressing breast cancers. It has been proposed that inhibition of HER3 phosphorylation and blockade of the PI3K/Akt signaling pathway are required in order to obtain a tumor suppression effect despite differences in the operating mechanisms of these therapeutic agents [29]. HER2–HER3 heterodimerization plays a key role in HER2-mediated transformation, tumor progression, and drug resistance [29]. Previous studies on loss of HER3 expression in HER2-dependent cells have demonstrated reduced PI3K signaling and decreased cell proliferation, suggesting that HER3 is essential for HER2-driven carcinogenesis [30, 31]. Treatment of HER2-amplified tumors with tyrosine kinase inhibitors results in a compensatory increase in HER3 expression, membranous HER3 localization, and decreased HER3 dephosphorylation [32]. MET-dependent phosphorylation of HER3 has been identified as one mechanism of resistance to EGFR inhibitors in lung cancer [33]. A recent study demonstrated that blocking both HER2 and HER3 inhibits the PI3K/Akt pathway more effectively than blocking HER2 or HER3 alone, suggesting that also HER3 should be inhibited in patients with HER2- and PI3K-dependent cancers in order to completely block the PI3K/Akt pathway [29]. Our group previously demonstrated that the Akt pathway is activated in a large proportion of EHCCs (84 %) [24]. In the present study, we demonstrate that HER3 is overexpressed in 39 % of EHCC cases. Considering the current study and the results of previous studies of breast cancers and EHCCs, we hypothesize that application of a therapeutic regimen for blocking HER3, such as AMG888, together with HER2 inhibitors and PI3K inhibitors might be effective in a subset of EHCC patients with activated Akt and HER3.

HER3 overexpression has been identified in malignant neoplasms from several organs, including melanoma, and prostate, colorectal, lung, ovarian, and gastric cancers [17, 30,34–38]. In addition, HER3 overexpression has been reported to be associated with outcome in melanomas and lung and gastric cancers [34, 38, 39]. Although HER3 is specifically required for HER2-driven tumorigenesis and has been extensively studied in the HER2-amplified subtype of breast cancers, other types of cancer also show higher HER3 expression without accompanying HER2 overexpression [30, 40]. In these tumors, HER3 may function as an allosteric activator of other members of the HER receptor tyrosine kinase family, such as EGFR [41–43]. EGFR expression has been reported in 8–57 % of cholangiocarcinomas, and patients with EGFR-positive tumors showed worse clinical outcome than those with EGFR-negative cancers [20, 23, 44, 45]. The absence of a survival difference in patients with HER2 overexpression and the significantly shorter survival in patients with HER3 overexpression, as demonstrated in this study, suggest a possible link between shorter survival time in the HER3-overexpression group and EGFR–HER3 heterodimerization in a subset of these patients. These patients may benefit from a therapeutic regimen that includes other types of HER inhibitors, including erlotinib,

cetuximab, and lapatinib, which have been investigated recently in cholangiocarcinomas [46–50].

HER2 overexpression in cholangiocarcinomas has been explored in various studies and has shown to vary widely (between 5 and 76 %) [20, 51–57]. These discrepancies might be explained by the lack of a standardized methodology, different standards of interpretation, or differences in tumor location. In recent reports, the HER2 expression rate in resected EHCCs was reported to be 5 to 10 % [20, 23, 45, 52]. The approximately 6 % HER2 expression rate identified in the current study is in agreement with these previous studies. Consistent with some previous reports, we found a higher HER2 overexpression rate in intraductal papillary neoplasm with an associated invasive carcinoma than in other histological types [20, 45, 54]. Although the HER2 overexpression rate is not exceptionally high in EHCCs, a subset of patients with HER2 overexpression might benefit from HER2-targeted monoclonal therapy.

The recently updated (7th edition) American Joint Committee on Cancer Staging Manual divided EHCCs into perihilar and distal bile duct cancers [58]. The term “distal bile duct cancers” was applied to cancers distal to the insertion of the cystic duct. This division reflects the growing evidence for distinct biological patterns in perihilar and distal EHCCs [59–61]. However, in our study, the proportions of HER2 and HER3 overexpression were similar in perihilar and distal bile duct cancer groups.

In conclusion, HER2 and HER3 proteins are overexpressed in subsets of EHCCs. HER3 overexpression is correlated with decreased patient survival and therefore constitutes a prognostic factor and a potential therapeutic target in patients with EHCCs.

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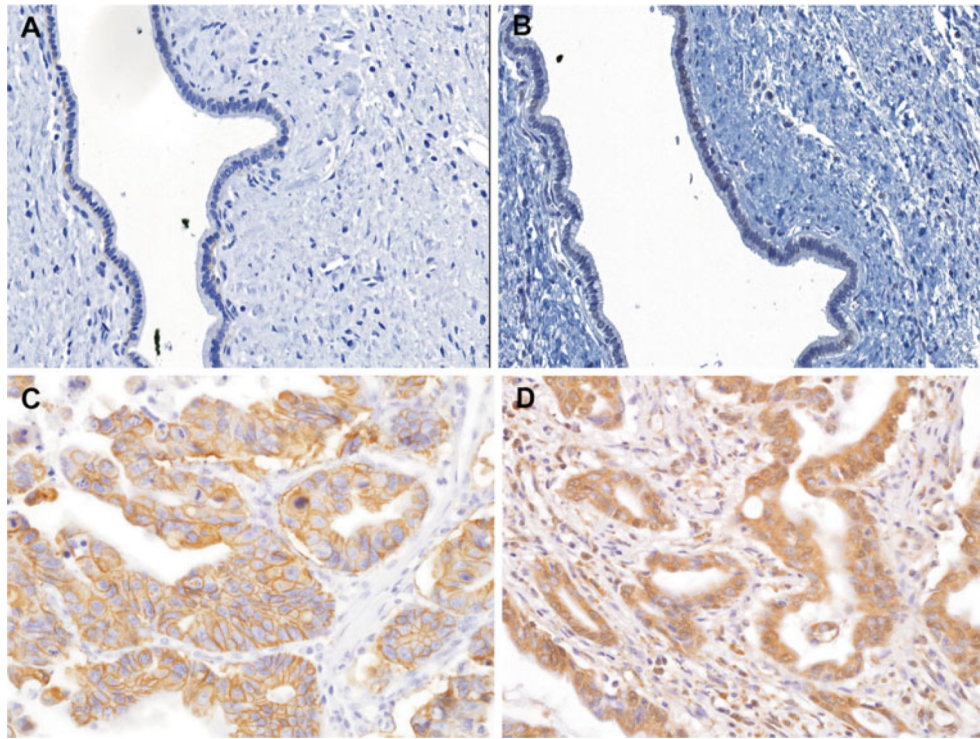


Fig. 1. Representative immunohistochemical staining of HER2 and HER3 in EHCC. Normal bile duct epithelia show negativity for HER2 (a) and HER3 (b). Cholangiocarcinoma show strong and complete membranous expression of HER2 (c) and diffuse and strong cytoplasmic HER3 immunopositivity (d)

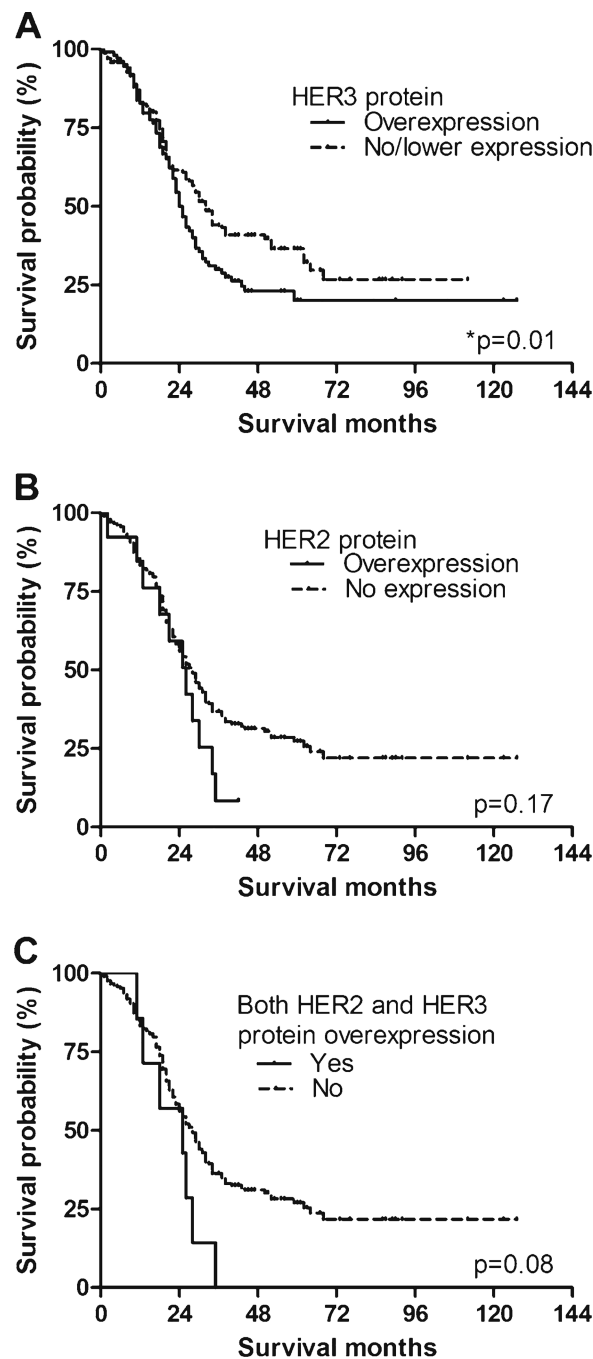


Fig. 2. Survival curves stratified by HER2 and HER3 expression in EHCC patients. **a** Median survival in patients in the HER3-overexpression group (24.0 months) was significantly shorter than that for patients in the no/ lower HER3-expression group (31.9 months; $P=0.01$, log-rank test). **b** Survival was not significantly different between patients in the HER2-overexpression group (25.9 months) and the no/lower HER2-expression group (27.8 months; $P=0.17$, log-rank test). **c** Median survival in EHCC patients with both HER2 and HER3 co-

overexpression (24.8 months) was marginally different from that in patients without HER2 and HER3 co-overexpression (28.0 months; $P=0.08$)

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Table 1

Clinicopathologic characteristics of patients with EHCC

Variables	No. of patients
Mean age	230
60.9 years	
Gender	
Male	164
Female	66
Mean tumor size	230
2.5 cm	
Location	
Perihilar	106
Distal	118
Diffuse	6
Histologic subtype	
Tubular adenocarcinoma	193
Intraductal papillary neoplasm with an associated invasive carcinoma	15
Intestinal type adenocarcinoma	9
Mucinous carcinoma	4
Adenosquamous carcinoma	6
Clear cell carcinoma	1
Signet ring cell carcinoma	1
Sarcomatoid carcinoma	1
pT classification	
pT1	31
pT2	84
pT3	91
pT4	24
Lymph node metastasis	
Present	76
Absent	154
Hepatic invasion	
Present	10
Absent	220
Pancreatic invasion	
Present	104
Absent	126
Duodenal invasion	
Present	24
Absent	206
Perineural invasion	
Present	167

Variables	No. of patients
Absent	63
Vascular invasion	
Present	71
Absent	159

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Table 2
Comparison between HER2 and HER3 immunohistochemical staining results and clinicopathologic factors in EHCC

Variable	Total no. of cases	HER2 expression (%)	P-value	Total no. of cases	HER3 expression (%)	P-value
Gender			0.36			0.46
Male	161	11 (6.8)		164	67 (40.9)	
Female	63	2 (3.2)		66	23 (34.8)	
Location			0.18			0.21
Perihilar	101	9 (8.9)		106	48 (45.3)	
Distal	117	4 (3.4)		118	40 (33.9)	
Diffuse	6	0 (0.0)		6	2 (33.3)	
Growth pattern			0.39			0.03*
Papillary	22	1 (4.5)		23	4 (17.4)	
Nodular	14	2 (14.3)		14	8 (57.1)	
Infiltrative	188	10 (5.3)		193	78 (40.4)	
Differentiation			1			0.13
Well differentiated	63	4 (6.3)		66	32 (48.5)	
Moderately differentiated	123	7 (5.7)		125	46 (36.8)	
Poorly differentiated	38	2 (5.3)		39	12 (30.8)	
Histologic subtype			0.02*			0.64
Tubular adenocarcinoma	188	8 (4.3)		193	73 (37.8)	
Intraductal papillary neoplasm with an associated invasive carcinoma	14	5 (35.7)		15	7 (46.7)	
Intestinal type adenocarcinoma	9	0 (0.0)		9	5 (55.6)	
Mucinous carcinoma	4	0 (0.0)		4	1 (25.0)	
Adenosquamous carcinoma	6	0 (0.0)		6	2 (33.3)	
Clear cell carcinoma	1	0 (0.0)		1	1 (100.0)	
Signet ring cell carcinoma	1	0 (0.0)		1	0 (0.0)	
Sarcomatoid carcinoma	1	0 (0.0)		1	1 (100.0)	
pT classification			0.35			0.08
pT1	29	3 (10.3)		31	16 (51.6)	
pT2	82	6 (7.3)		84	38 (45.2)	
pT3	89	4 (4.5)		91	27 (29.7)	

Variable	Total no. of cases	HER2 expression (%)	P-value	Total no. of cases	HER3 expression (%)	P-value
pT4	24	0 (0.0)		24	9 (37.5)	
Duodenal invasion						
Absent	200	13 (6.5)	0.37	206	81 (39.3)	1
Present	24	0 (0.0)		24	9 (37.5)	
Hepatic invasion						
Absent	215	12 (5.6)	1	220	87 (39.5)	0.74
Present	9	1 (11.1)		10	3 (30.0)	
Pancreatic invasion						
Absent	121	10 (8.3)	0.15	126	58 (46.0)	0.02*
Present	103	3 (2.9)		104	32 (30.8)	
Perineural invasion						
Absent	62	6 (9.7)	0.2	63	30 (47.6)	0.13
Present	162	7 (4.3)		167	60 (35.9)	
Vascular invasion						
Absent	154	10 (6.5)	0.56	159	60 (37.7)	0.56
Present	70	3 (4.3)		71	30 (42.3)	
Lymph node metastasis						
Absent	153	9 (5.9)	1	154	62 (40.3)	0.67
Present	71	4 (5.6)		76	28 (36.8)	
Resection marginal status						
Negative	173	10 (5.8)	1	176	64 (36.4)	0.15
Positive	51	3 (5.9)		54	26 (48.1)	
Stage grouping						
Stage IA	24	3 (12.5)	0.46	25	12 (48.0)	0.4
Stage IB	60	3 (5.0)		60	28 (46.7)	
Stage IIA	58	3 (5.2)		58	18 (31.0)	
Stage IIB	58	4 (6.9)		63	23 (36.5)	
Stage III	24	0 (0.0)		24	9 (37.5)	

* $P < 0.05$ (significant)

Table 3

Comparison of results of immunohistochemical staining for HER2 and HER3 co-expression with clinicopathologic factors in EHCC

Variable	Total no. of cases	HER2 and HER3 co-overexpression (%)	P-value
Gender			0.2
Male	161	7 (4.3)	
Female	63	0 (0.0)	
Location			0.09
Perihilar	101	6 (5.9)	
Distal	117	1 (0.9)	
Diffuse	6	0 (0.0)	
Growth pattern			0.36
Papillary	22	0 (0.0)	
Nodular	14	1 (7.1)	
Infiltrative	188	6 (3.2)	
Differentiation			1
Well differentiated	63	2 (3.2)	
Moderately differentiated	123	4 (3.3)	
Poorly differentiated	38	1 (2.6)	
Histologic subtype			0.12
Tubular adenocarcinoma	188	4 (2.1)	
Intraductal papillary neoplasm with an associated invasive carcinoma	14	3 (21.4)	
Intestinal type adenocarcinoma	9	0 (0.0)	
Mucinous carcinoma	4	0 (0.0)	
Adenosquamous carcinoma	6	0 (0.0)	
Clear cell carcinoma	1	0 (0.0)	
Signet ring cell carcinoma	1	0 (0.0)	
Sarcomatoid carcinoma	1	0 (0.0)	
pT classification			0.66
pT1	29	1 (3.4)	
pT2	82	4 (4.9)	
pT3	89	2 (2.2)	
pT4	24	0 (0.0)	
Duodenal invasion			0.61
Absent	200	7 (3.5)	
Present	24	0 (0.0)	
Hepatic invasion			1
Absent	215	7 (3.3)	
Present	9	0 (0.0)	
Pancreatic invasion			0.46
Absent	121	5 (4.1)	
Present	103	2 (1.9)	

Variable	Total no. of cases	HER2 and HER3 co-overexpression (%)	P-value
Perineural invasion			0.4
Absent	62	3 (4.8)	
Present	162	4 (2.5)	
Vascular invasion			0.44
Absent	154	6 (3.9)	
Present	70	1 (1.4)	
Lymph node metastasis			1
Absent	153	5 (3.3)	
Present	71	2 (2.8)	
Resection marginal status			1
Absent	173	5 (2.9)	
Present	51	2 (3.9)	
Stage grouping			0.94
Stage IA	24	1 (4.2)	
Stage IB	60	2 (3.3)	
Stage IIA	58	2 (3.4)	
Stage IIB	58	2 (3.4)	
Stage III	24	0 (0.0)	

Table 4

Univariate analysis of pathologic features affecting survival in EHCC patients

Variable	Characteristics	Median survival time (month)	P-value	95 % confidence interval	
				Lower	Upper
HER2 expression	Low	27.8	0.17	24.2	31.4
	High	25.9		18.6	33.2
HER3 expression	Low	31.9	0.01*	27.3	36.5
	High	24.0		22.0	26.0
Gender	Male	28.4	0.38	24.8	32.0
	Female	21.8		15.0	28.6
Location	Perihilar	24.8	0.47	21.1	28.4
	Distal	30.0		25.5	34.5
	Diffuse	12.0		0	42.7
Growth pattern	Papillary	50.0	0.09		
	Nodular	30.0		10.8	49.2
	Infiltrative	25.9		22.2	29.6
Differentiation	Well	34.0	<0.001*	24.0	44.0
	Moderate	26.8		22.5	31.2
	Poor	17.5		11.0	24.0
pT classification	pT1	44.0	0.01*		
	pT2	24.9		22.2	27.6
	pT3	25.9		19.0	32.9
	pT4	27.0		14.5	39.5
Perineural invasion	Absent	28.9	0.36	21.8	36.0
	Present	26.1		21.8	30.4
Vascular invasion	Absent	28.9	0.02*	22.9	34.9
	Present	26.1		20.9	31.2
Duodenal invasion	Absent	26.9	0.07	23.6	30.2
	Present	27.0		14.5	39.5
Liver invasion	Absent	28.0	0.01*	24.9	31.1
	Present	17.5		0.0	35.7
Pancreas invasion	Absent	26.1	0.24	21.6	30.6
	Present	27.0		22.2	31.8
Resection margin status	Negative	28.8	0.1	24.9	32.7
	Positive	23.1		17.5	28.7
Lymph node metastasis	Absent	30.0	<0.001*	27.0	33.0
	Present	18.2		15.5	20.9

* $P < 0.05$ (significant)

Table 5

Multivariate analysis for prognosis in EHCC

Variable	P-value	Relative risk	95 % confidence interval
HER3 expression	0.007*	1.6	1.14–2.26
Differentiation	0.002*	1.52	1.16–2.00
pT classification	0.066	1.21	0.99–1.48
Vascular invasion	0.646	1.1	0.75–1.62
Liver invasion	0.025*	2.47	1.12–5.45
Lymph node metastasis	0.002*	1.81	1.25–2.65

Duodenal invasion was not included because it was a covariate with pT classification

* $P < 0.05$ (significant)

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