

Intracellular localization of mesothelin predicts patient prognosis of extrahepatic bile duct cancer

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Abstract. Mesothelin is expressed in various types of malignant tumors, and we recently reported that the expression of mesothelin was related to unfavorable patient outcome in pancreatic ductal adenocarcinoma and gastric adenocarcinoma. In this study, we examined the clinicopathological significance of mesothelin expression in extrahepatic bile duct cancer (EHBDC), especially in terms of its association with the staining pattern. Tissue samples from 61 EHBDC (16 hilar cholangiocarcinoma, 17 upper bile duct adenocarcinoma, 20 middle bile duct adenocarcinoma and 8 distal bile duct adenocarcinoma) were immunohistochemically examined. The expression levels of mesothelin in tumor cells was classified into the localization of mesothelin in luminal membrane and/or cytoplasm, in addition to high and low according to the staining intensity and proportion as a conventional analysis. 'High-level expression' of mesothelin (47.5%) was statistically correlated with liver metastasis ($P=0.013$) and poorer patient outcome ($P=0.022$), while 'luminal membrane positive' of mesothelin (52.5%) was more significantly correlated with liver metastasis ($P=0.006$), peritoneal metastasis ($P=0.024$) and unfavorable patient outcome ($P=0.017$). Moreover, we found that 'cytoplasmic expression' isolated from 'luminal membrane negative' of mesothelin represented the best patient prognosis throughout this study. We describe the expression pattern level of mesothelin, i.e., in luminal membrane or cytoplasm both high and low level, evidently indicate the patient prognosis of EHBDC, suggesting the pivotal role of mesothelin in cancer promotion depending on its intracellular localization.

Introduction

Extrahepatic bile duct cancer (EHBDC), consisting of hilar cholangiocarcinoma and distal bile duct adenocarcinoma (excluding gallbladder cancer), is a rare disease in the United States with an incidence of 1-2/100,000/year (1). It occurs with great frequency in Asian countries, and is one of the common causes of cancer death in Japan, with near to 17,000 deaths annually (2). The 5-year survival rate of EHBDC, even after the surgical resection is poor, ranging from 20 to 45% (3-5). The incidence of EHBDC is increasing throughout the world with a high fatality rate; therefore, new prognostic markers and treatment for EHBDC patients are urgently needed.

Mesothelin is expressed on normal mesothelial cells lining the pleura, pericardium and peritoneum (6,7). In addition, the overexpression of mesothelin has been found in several cancer types, including malignant mesothelioma, ovarian cancer and pancreatic cancer (8-11,12). The full length of human *mesothelin* gene codes the primary product, which is a 71-kDa precursor protein. This protein can be physiologically cleaved by certain furin-like proteases into a 40-kDa C-terminal fragment that remains membrane-bound and a 31-kDa N-terminal fragment, which is secreted into the blood (6). The C-terminal 40-kDa fragment is named mesothelin and is attached to the cell membrane through a glycosyl-phosphatidylinositol (GPI) anchor (13). The biological functions of mesothelin are not clearly understood, although recent studies have suggested that enforced expression of mesothelin increases cell proliferation and migration (14). In ovarian cancers, higher mesothelin expression was found to be associated with chemoresistance and shorter patient survival (15). In pancreatic cancer, mesothelin expression was immunohistochemically observed in all cases, while its absence was noted in non-cancerous pancreatic ductal epithelium, with or without pancreatitis (8,12,16,17). We recently found that the expression of mesothelin was related to an unfavorable patient outcome in pancreatic ductal adenocarcinoma (12), while the opposite result was reported in gastric cancer, in which the mesothelin expression was correlated with prolonged patients' survival (18). However, our consecutive investigation for mesothelin expression patterns in gastric cancer recently discovered that luminal membrane expression, not cytoplasmic expression

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of mesothelin is a prominent negative prognostic factor for gastric cancer (19), suggesting the significance of expression pattern of mesothelin in clinicopathological analysis of cancer. In EHBDCA, Zhao *et al*, who first studied mesothelin expression in dysplasia and carcinoma of external bile duct, reported that mesothelin was expressed in 5 of 10 adenocarcinomas (50%) in cell membranes and cytoplasm (20); however, the detailed clinicopathological analysis of mesothelin expression in EHBDCA, especially with large number of the cases, has not yet been performed.

In this study, we investigated the mesothelin expression in 61 EHBDCA cases by immunohistochemistry, and its clinicopathological significance associated with patients' outcome was analyzed. Moreover, we focused on the intracellular localization of mesothelin, i.e., in luminal membrane and/or cytoplasm, and its clinicopathological significance associated with the patients' outcome.

Materials and methods

Patients' demography and tumor specimens. This study was performed with the approval of the Internal Review Board on Ethical Issues of Hokkaido University Hospital, Sapporo, Japan. The samples and the patient information were obtained under a blanket written informed consent. The subjects of this study were 61 patients who underwent radical surgery for bile duct adenocarcinoma between the years 2000 and 2008 at Hokkaido University Hospital by the Department of General Surgery, Hokkaido University, Graduate School of Medicine, Sapporo, Japan. The clinicopathological characteristics of these cases are summarized in Table I.

Mean age of patients was 67.5 years [± 9.0 standard deviation (SD)]; 47 patients (77.0%) were male and 14 patients (23.0%) were female. The predominant sites of the cancer were the hilar bile duct in 16 cases (26.2%), upper bile duct in 17 cases (27.9%), middle bile duct in 20 cases (32.8%) and distal bile duct in 8 cases (13.1%). The surgical procedures consisted of the standard pancreatoduodenectomy in 21 (34.4%) cases, the pylorus-preserving pancreatoduodenectomy in 5 cases (8.2%), the extended right or left hemihepatectomy with extrahepatic bile duct resection in 28 cases (45.9%), and the extrahepatic bile duct resection in 7 cases (11.5%). Intraoperative diagnosis of the ductal resection margins was performed using frozen sections. When a positive margin was found, additional resection of marginal bile duct was performed to the maximum extent possible. R0 curative resection was achieved in 39 cases (63.9%), and R1 resection was achieved in 22 cases (36.1%). T-factor, N-factor, M-factor and clinical stage were assigned according to the TNM classification of the Union Internationale Contre le Cancer (UICC) (21). The median survival time of patients was 29.8 months (± 3.5 SD).

Formalin-fixed paraffin-embedded tissue blocks were prepared from surgical specimens and sections were sliced and stained with hematoxylin and eosin (H&E) for routine histopathological examination. All specimens were diagnosed as EHBDCA.

Immunohistochemical evaluation. Immunohistochemical staining against mesothelin was performed as described

Table I. Clinicopathological characteristics of 61 patients with EHBDCA in this study.

Parameter	No. of cases
Age (years)	
<60	11
≥ 60	50
Mean \pm SD	67.5 \pm 9.0
Gender	
Male	47
Female	14
Location	
Hilar	16
Upper	17
Middle	20
Distal	8
Surgical procedure	
Pancreatoduodenectomy	21
Pylorus-preserving pancreatoduodenectomy	5
Extended right or left hemihepatectomy with bile duct resection	28
Extrahepatic bile duct resection	7
Resection status	
R0	39
R1	22
T-factor	
T1	5
T2	27
T3	19
T4	10
N-factor	
N0	25
N1	36
M-factor	
M0	58
M1	3
Stage	
IA	4
IB	14
IIA	4
IIB	28
III	8
IV	3
Median survival (months)	29.8 \pm 3.5

SD, standard deviation.

previously (12). In brief, the tissue sections were incubated with a mouse monoclonal antibody against mesothelin (clone 5B2 diluted 1:50; Novocastra, Newcastle Upon Tyne, UK) at a 1:50 dilution, and reacted with a dextran polymer reagent combined with secondary antibodies and peroxidase (Envision/HRP; Dako). All assessments were made

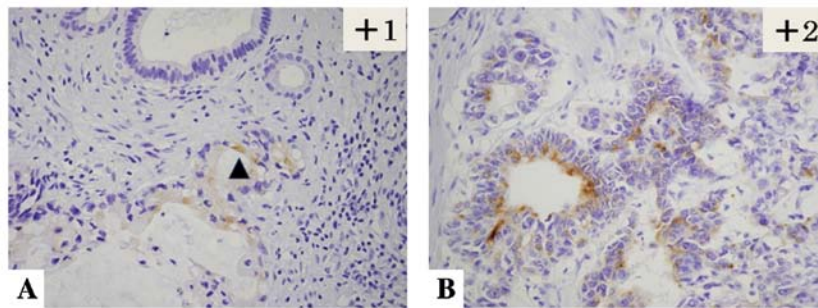


Figure 1. Representative cases of 'low-level expression' (A) and 'high-level expression' (B) of mesothelin in EHBDCa specimens by immunohistochemistry. (A) Partial luminal membrane staining (arrowhead; intensity, +1) and the weak cytoplasmic staining were observed in <50% area (proportion, +2). (B) Entire circumference of the luminal membrane was strongly positive in >50% tumor cells (intensity, +2; proportion, +3). (Magnification, x200).

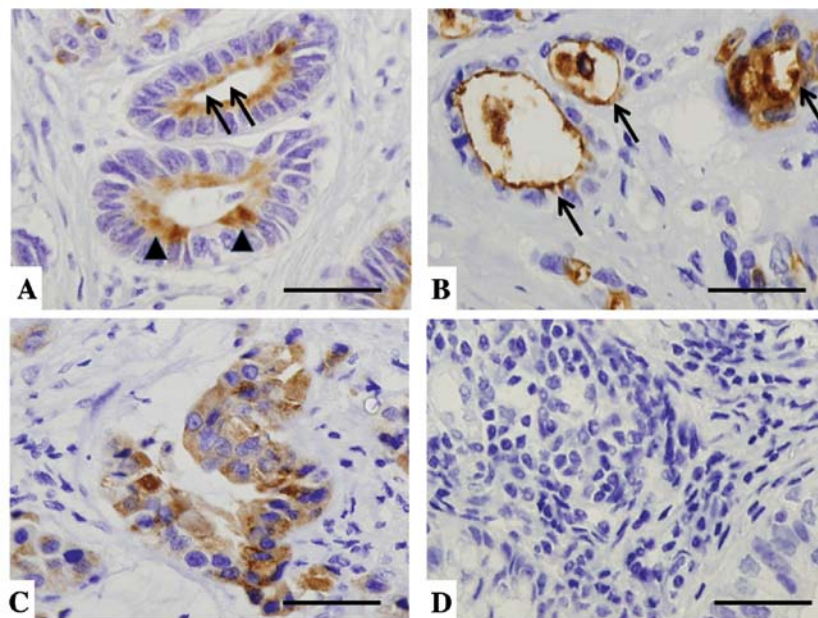


Figure 2. Representative cases of 'luminal membrane positive' (A, B) and 'luminal membrane negative' (C, D) of mesothelin in EHBDCa specimens by immunohistochemistry. (A) Granular cytoplasmic staining was observed (arrowheads; intensity, +2) and luminal membrane was also stained partially (arrows). (B) Entire circumference of the luminal membrane was explicitly stained (arrows). (C) Granular cytoplasmic, but no membranous staining in cancer cells was observed. (D) No expression of mesothelin was found in tumor cells, also designated 'mesothelin negative'. (Magnification, x400; scale bars, 50 μ m).

on the tumor region of the specimen (x400). Each slide was evaluated independently by three pathologists (F. Kawamata, M. Miyazaki and H. Nishihara) who did not know the clinical outcomes. Immunostaining for mesothelin was evaluated for both the proportion and staining intensity of tumor cells in each case. The proportion of mesothelin expression was assessed according to the percentage of mesothelin-positive cells as follows: 0, 0%; +1, 1-10%; +2, 10-50%; and +3, >50%. The staining intensity of mesothelin was evaluated as weak (+1) and moderate to strong (+2) (Table II). The final evaluation of mesothelin expression was assessed using the following scoring system: 'high-level expression' of mesothelin was defined as $\geq +3$ of the proportion score and/or +2 of the intensity score, while a 'low-level expression' of mesothelin was given when the total score was $\leq +3$ except in cases when the proportion score was +1 and the intensity score was +2 (Fig. 1).

Furthermore, among the 61 cases of EHBDCa, the staining localization of mesothelin was evaluated in luminal membrane

Table II. Immunohistochemical findings of mesothelin expression.

Staining intensity on tumor cells	No. of cases (%)			
	Percentage of mesothelin-positive cells			
	0	1-10%	10-50%	>50%
Score 0	17 (27.9)	0 (0.0)	0 (0.0)	0 (0.0)
Score 1	0 (0.0)	13 (21.3)	2 (3.3)	1 (1.6)
Score 2	0 (0.0)	6 (9.8)	12 (19.7)	10 (16.4)

or cytoplasm. Cases in which the luminal membrane was stained even partially or faintly (Fig. 2A), or the entire circumference of the luminal membrane was explicitly stained

Table III. Correlation between mesothelin expression levels and clinicopathological features.

Parameter	Total	Mesothelin		P-value	Luminal membrane expression		
		High-level (n=29)	Low-level (n=32)		Positive (n=32)	Negative (n=29)	P-value
Histopathological grade							
1 or 2	54	26	28	1.000	28	26	1.000
3	7	3	4		4	3	
pT-factor							
pT1-2	32	13	19	0.310	19	13	0.310
pT3-4	29	16	13		13	16	
pN-factor							
Negative	25	11	14	0.795	16	9	0.193
Positive	36	18	18		16	20	
pStage							
I-IIIB	50	24	26	1.000	26	24	1.000
III-IV	11	5	6		6	5	
Lymphatic permeation							
Negative	23	10	13	0.792	12	11	1.000
Positive	38	19	19		20	18	
Blood vessel permeation							
Negative	26	11	15	0.606	11	15	0.200
Positive	35	18	17		21	14	
Perineural invasion							
Negative	9	3	6	0.478	3	6	0.287
Positive	52	26	26		29	23	
Resection margin							
pR0	39	20	19	0.594	24	15	0.069
pR1	22	9	13		8	14	
Recurrence							
No	18	6	12	0.172	6	12	0.090
Yes	43	23	20		26	17	
Liver metastasis							
No	47	18	29	0.013	20	27	0.006
Yes	14	11	3		12	2	
Local recurrence							
No	46	22	24	1.000	25	21	0.767
Yes	15	7	8		7	8	
Peritoneal metastasis							
No	49	20	29	0.052	22	27	0.024
Yes	12	9	3		10	2	

(Fig. 2B) were judged as 'luminal membrane positive'. In cases with no membrane staining (Fig. 2D) and those in which only cytoplasmic staining (Fig. 2C) was observed in any intensity level, the term 'luminal membrane negative' was given.

Statistical analysis. We used the χ^2 test or Fisher's exact test to determine the correlation between mesothelin and clinicopathologic data. Survival curves for patients were drawn by the Kaplan-Meier method. Differences in survival curves were analyzed by the log-rank test. Prognostic implications of mesothelin expression and clinicopathologic parameters were

analyzed by Cox univariate and multivariate proportional hazards models. All differences were considered significant at a P-value of <0.05. All statistical analyses were performed using the Ekuseru-Toukei 2010 software for Windows (Social Survey Research Information Co., Ltd., Tokyo, Japan).

Results

High-level expression of mesothelin was correlated with liver metastasis and poor patient outcome. The overexpression of mesothelin has been found in several cancer types, including

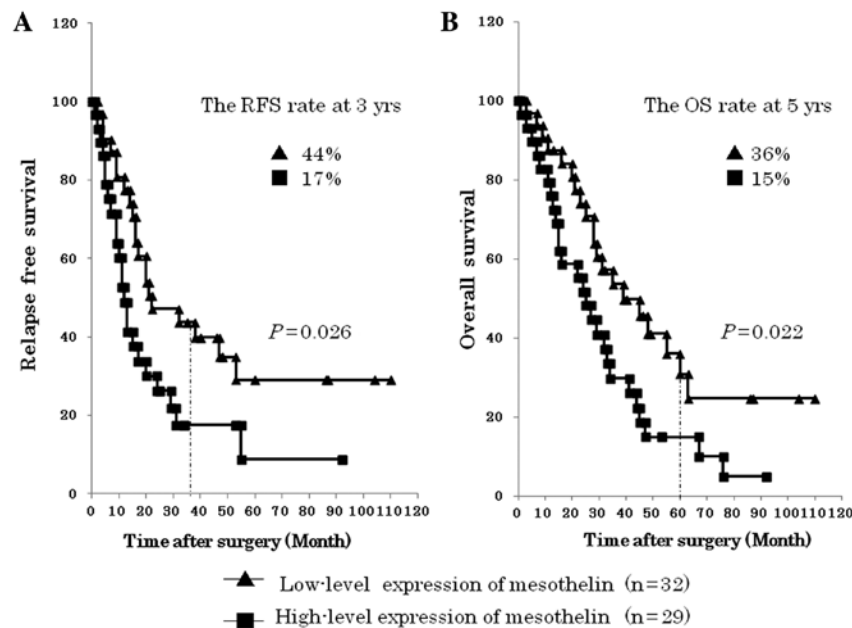


Figure 3. Relapse-free survival (RFS) and overall survival (OS) curves of EHBDCa patients according to the expression levels of mesothelin. The group of 'high-level expression' of mesothelin represented a statistically significantly unfavorable outcome compared to the group of 'low-level expression' ($P=0.026$ and 0.022 , respectively).

malignant mesothelioma, ovarian cancer, and pancreatic cancer (8-11,12); thus, we first evaluated the comprehensive expression of mesothelin in EHBDCa. As described in Materials and methods, 'high-level expression' and 'low-level expression' of mesothelin was attributed to all 61 cases of EHBDCa (Fig. 1). As summarized in Table II, 'high-level expression' was detected in 29 cases (47.5%), whereas 'low-level expression' was detected in 32 cases (52.5%). The statistical analysis for the clinicopathological parameters such as histological grade, T-factor and metastasis revealed that 'high-level expression' of mesothelin was significantly correlated with liver metastasis ($P=0.013$, Table III). Furthermore, recent studies reported that higher mesothelin expression was found to be associated with shorter patient survival; therefore, we examined the correlation of mesothelin overexpression with relapse-free survival (RFS) and overall survival (OS) in the EHBDCa patients. The group of 'high-level expression' of mesothelin had a significantly poorer RFS than the group of 'low-level expression' of mesothelin ($P=0.026$). In addition, the group of 'high-level expression' of mesothelin had a significantly poorer OS than the group of 'low-level expression' of mesothelin ($P=0.022$) (Fig. 3).

Luminal membrane expression of mesothelin is a prominent negative prognostic factor for the patients with EHBDCa. During our previous studies on pancreatic adenocarcinoma and gastric adenocarcinoma, we already noted that expression of mesothelin was found in the luminal membrane as well as in the cytoplasm (19). Mesothelin was reported to attach to the cell membrane through a glycosyl-phosphatidylinositol (GPI) anchor after being physiologically cleaved by some furin-like proteases (22), which are involved in the translocation of mesothelin, although the biological functions of mesothelin associated with its intracellular localization are not fully understood. Thus, we analyzed the intracellular localization

of mesothelin by immunostaining to explore the clinicopathological significance of its translocation.

As shown in Table III, the group 'luminal membrane positive', which consisted of the cases with luminal membrane staining even partially, was 32 (52.5%) cases, while the group 'luminal membrane negative', which contained 17 cases which were completely mesothelin negative was comprised of 29 (47.5%) cases. The statistical analysis revealed that the incidence of luminal membrane positivity was significantly correlated with peritoneal metastasis ($P=0.024$) in addition to liver metastasis ($P=0.006$) (Table III). The analysis of the patients' overall survival showed that 'luminal membrane positive' of mesothelin indicated a significantly unfavorable RFS ($P=0.012$) and OS ($P=0.017$) compared to 'luminal membrane negative' of mesothelin (Fig. 4).

To clarify the mesothelin expression as an independent prognostic factor, we performed a univariate analysis of the 61 EHBDCa using the Cox proportional hazards model, the result indicated that resection margin, 'high-level expression' and 'luminal membrane positive' of mesothelin were significantly correlated with risks of cancer mortality. Multivariate analysis also confirmed that resection margin (RR 3.361, 95% CI, 1.670-6.763, $P=0.0007$) and 'luminal membrane positive' of mesothelin (RR 2.964, 95% CI, 1.401-6.296, $P=0.0045$) were independent predictors of the overall patient survival (Table IV).

Isolation of 'cytoplasmic expression' of mesothelin potentiates more exquisite prediction of prognosis in EHBDCa. To explore the clinicopathological value of the cytoplasmic expression of mesothelin, we performed a sub-analysis in 'luminal membrane negative', dividing the group into 17 cases of 'mesothelin negative' and 12 cases of 'cytoplasmic expression'. The P-value (OS, $P=0.0085$) between 'luminal membrane positive' and 'cytoplasmic expression' was minimum in these

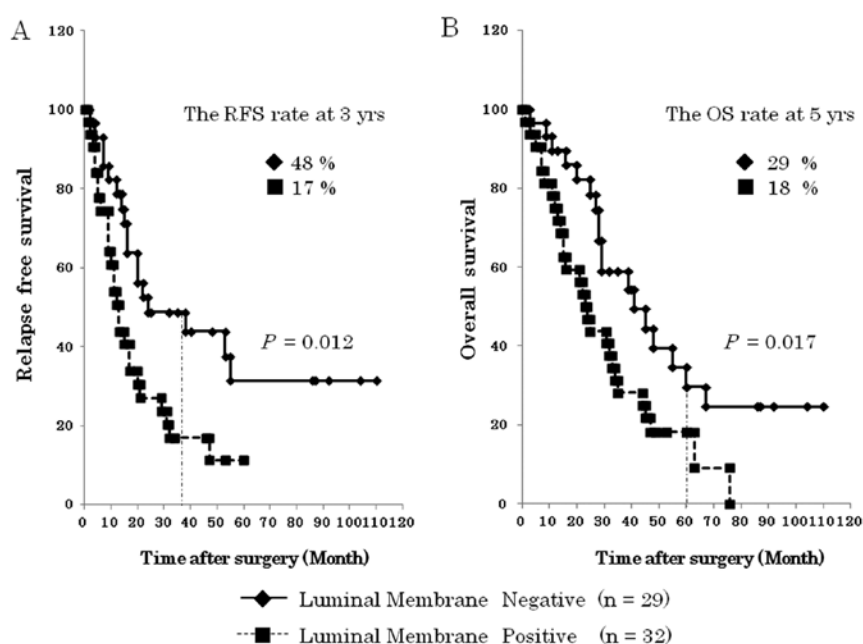


Figure 4. Relapse-free survival (RFS) and overall survival (OS) curves of EHBDCa patients according to the expression pattern of mesothelin. The group of 'luminal membrane positive' represented a statistically significantly unfavorable outcome compared to the group of 'luminal membrane negative' ($P=0.012$ and 0.017 , respectively).

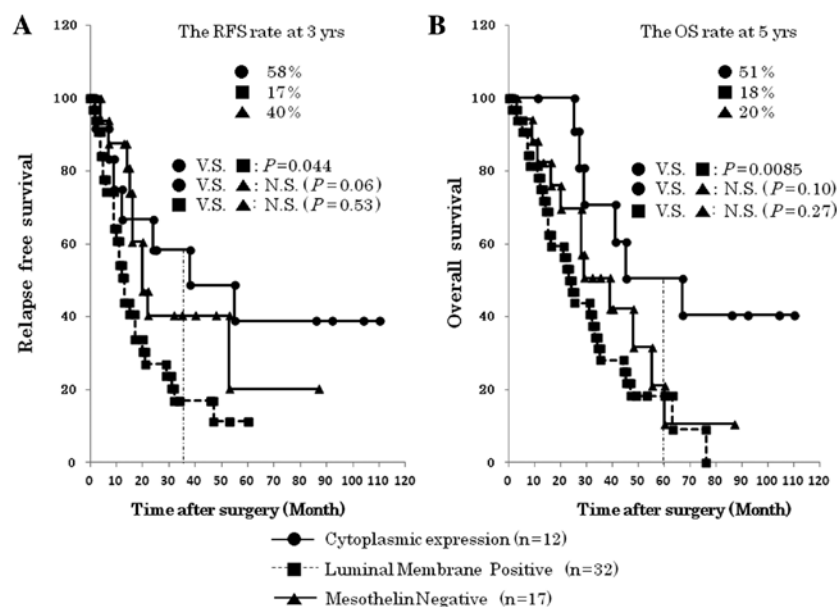


Figure 5. Relapse-free survival (RFS) and overall survival (OS) curves of EHBDCa patients among three groups of detailed expression patterns of mesothelin. 'Cytoplasmic expression' of mesothelin represented the best prognosis among the 3 groups.

survival analyses, suggesting the clinical benefit of isolation of 'cytoplasmic expression' of mesothelin (Fig. 5). Interestingly, 'cytoplasmic expression' of mesothelin represented relatively favorable patients' prognosis compared to 'mesothelin negative', although it was statistically not significant (RFS, $P=0.06$; OS, $P=0.10$).

Discussion

In this study, we confirmed that mesothelin expression is a prominent prognostic factor for EHBDCa patients as well

as for other tumors such as pancreatic cancer and ovarian carcinoma described previously (12,15,23). Furthermore, we revealed that the expression pattern of mesothelin, in luminal membrane or cytoplasm, could be a more evident prediction factor for these patients. These results evidently support our recent report of mesothelin expression patterns in gastric cancer in which luminal membrane expression, not cytoplasmic expression of mesothelin is a prominent negative prognostic factor for gastric cancer (19).

The mechanism for the membranous localization of mesothelin should be explained as follows: the full length of the

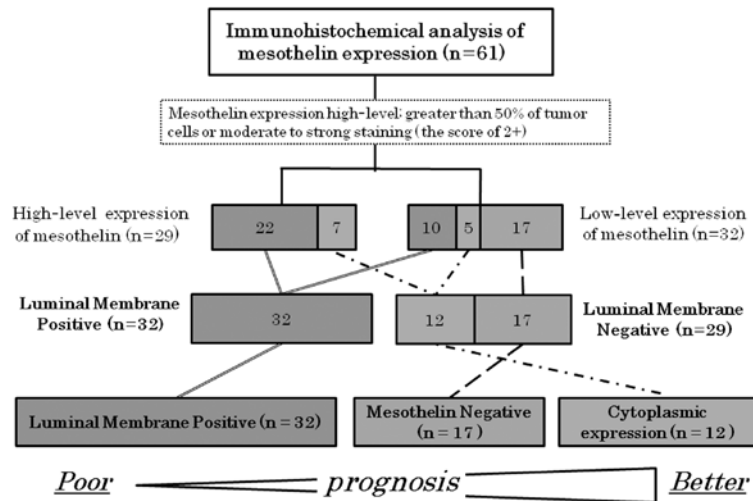


Figure 6. Flow chart of immunohistochemical evaluation of mesothelin expression and the prognostic aspect. The P-value (OS, P=0.0085) between 'luminal membrane positive' and 'cytoplasmic expression' was minimum in our survival analyses, suggesting the clinical benefit of isolation of 'cytoplasmic expression' of mesothelin.

Table IV. Univariate and multivariate analysis of patients' survival in EHBDCa.

Factor	n=61	Univariate analysis		Multivariate analysis		
		P-value	RR (95% CI)	RR (95% CI)	Hazard ratio	P-value
Histopathological grade						
1 or 2	54	0.3931	1		NC	
3	7		1.508 (0.588-3.871)			
pT-factor						
pT1-2	32	0.4264	1		NC	
pT3-4	29		1.266 (0.708-2.262)			
pN-factor						
Negative	25	0.3639	1		NC	
Positive	36		1.314 (0.729-2.368)			
pStage						
I-IIIB	50	0.2026	1		NC	
III-IV	11		1.608 (0.774-3.339)			
Lymphatic permeation						
Negative	23	0.1908	1		NC	
Positive	38		1.537 (0.807-2.924)			
Blood vessel permeation						
Negative	26	0.2999	1		NC	
Positive	35		1.370 (0.756-2.482)			
Perineural invasion						
Negative	9	0.4733	1		NC	
Positive	52		0.728 (0.306-1.732)			
Resection margin						
pR0	39	0.0398	1	1.670-6.763	1	0.0007
pR1	22		1.859 (1.029-3.356)		3.361	
Mesothelin expression						
Low-level	32	0.0236	1	0.864-3.067	1	0.1317
High-level	29		1.968 (1.095-3.538)		1.621	
Luminal membrane expression of mesothelin						
Negative	29	0.0175	1	1.401-6.296	1	0.0045
Positive	32		2.078 (1.137-3.798)		2.964	

RR indicates relative risk/hazard ratio; CI, confidence interval. NC, not calculable.

Table V. Sub-analysis among three groups according to the intracellular expression pattern of mesothelin.

Parameter	Total (n=44)	Luminal membrane positive (n=32)	Cytoplasmic expression (n=12)	P-value	Total (n=49)	Luminal membrane positive (n=32)	Negative expression (n=17)	P-value	Total (n=29)	Cytoplasmic expression (n=12)	Negative expression (n=17)	P-value
Histopathological grade												
1 or 2	39	28	11	1.000	43	28	15	1.000	26	11	15	1.000
3	5	4	1		6	4	2		3	1	2	
pT-factor												
pT1-2	23	19	4	0.179	28	19	9	0.765	13	4	9	0.452
pT3-4	21	13	8		21	13	8		16	8	8	
pN-factor												
Negative	18	16	2	0.083	23	16	7	0.764	9	2	7	0.234
Positive	26	16	10		26	16	10		20	10	10	
pStage												
I-IIIB	37	26	11	0.653	39	26	13	0.722	24	11	13	0.370
III-IV	7	6	1		10	6	4		5	1	4	
Lymphatic permeation												
Negative	14	12	2	0.282	21	12	9	0.370	11	2	9	0.064
Positive	30	20	10		28	20	8		18	10	8	
Blood vessel permeation												
Negative	16	11	5	0.732	21	11	10	0.134	15	5	10	0.462
Positive	28	21	7		28	21	7		14	7	7	
Perineural invasion												
Negative	3	3	0	0.551	9	3	6	0.049	6	0	6	0.028
Positive	41	29	12		40	29	11		23	12	11	
Resection margin												
pR0	30	24	6	0.152	32	24	8	0.065	14	6	8	1.000
pR1	14	8	6		17	8	9		15	6	9	
Recurrence												
No	11	6	5	0.139	13	6	7	0.172	12	5	7	1.000
Yes	33	26	7		36	26	10		17	7	10	
Liver metastasis												
No	30	20	10	0.282	36	20	16	0.020	26	10	16	0.553
Yes	14	12	2		13	12	1		3	2	1	
Local recurrence												
No	34	25	9	1.000	37	25	12	0.729	21	9	12	1.000
Yes	10	7	3		12	7	5		8	3	5	
Peritoneal metastasis												
No	34	22	12	0.041	37	22	15	0.175	27	12	15	0.498
Yes	10	10	0		12	10	2		2	0	2	

human *mesothelin* gene encodes a 71-kDa precursor protein that is proteolytically cleaved by some furin-like proteases into an *N*-terminal secreted form and a *C*-terminal fragment, the 40-kDa mesothelin, which is a glycosyl-phosphatidylinositol (GPI)-linked glycoprotein (6,13,15). Many researchers have investigated the role of the mesothelin expression in tumor biology and demonstrated the importance of mesothelin expression for tumor progression *in vitro* (14,24-26) and *in vivo* (27,28); however, the clinicopathological significance of the membrane localization of mesothelin has not been clarified. The 5B2 anti-mesothelin antibody, which we employed here for IHC, can detect both the 71-kDa precursor protein and the 40-kDa *C*-terminal fragment, but not the 30-kDa *N*-terminal fragment. According to the reported molecular processing mechanism of mesothelin and specificity of antibody, luminal membrane staining probably indicates the 40-kDa membrane-bound form of mesothelin, while cytoplasmic staining would mean the 71-kDa precursor form of mesothelin. Our results support the idea that the 40-kDa membrane-bound form of mesothelin is an active form and promotes the aggressive features including increased cell motility, invasion or migration capabilities and growth of metastatic tumors (24,25,29).

The fact that 'cytoplasmic expression' of mesothelin paradoxically resulted in better OS than mesothelin with 'mesothelin negative' took us by surprise (Fig. 5B). The RFS rate at 3 years (58 and 40%, respectively) and OS at 5 years (51 and 20%, respectively) were demonstrably better in 'cytoplasmic expression' compared to 'mesothelin negative', although the final RFS and OS were not statistically significant (RFS, $P=0.06$; OS, $P=0.10$). As indicated above, the majority of mesothelin in cytoplasm must be the 71-kDa precursor form and might behave like a dominant negative form of mesothelin as a tumor suppressor. The conflicting results in some previous reports in which mesothelin expression was correlated with prolonged patient survival in gastric cancer (18) and in ovarian serous carcinoma (30), may be explained by confusing the luminal membrane and cytoplasmic expression of mesothelin. Isolation of 'mesothelin negative' might give us another disease entity, mesothelin-independent EHBDCa. The tumor cells in such a type of EHBDCa would obtain invasive ability without the association of mesothelin; therefore, this could indicate an alternative gene expression profiling. In fact, additional sub-analysis for clinicopathological parameters among the three groups showed interesting results. Frequent perineural invasion was observed in 'mesothelin negative' rather than in mesothelin positive cases even in luminal membrane or cytoplasm ($P=0.049$ and 0.028 , respectively), while liver metastasis was abundantly found in 'luminal membrane positive' (Table V). Such conflicting results may suggest the distinct oncogenic process between mesothelin-associated and mesothelin-independent EHBDCa.

In terms of discovering the clinicopathological parameters, there are many previous studies demonstrating the prognostic significance of various molecules, such as epidermal growth factor receptor (EGFR) and c-erbB-2 (HER-2) in colorectal, breast and lung cancer (31). There are some other case reports describing a series of promising results targeting EGFR in patients with advanced biliary tract cancer (32-34); however, identification of useful prognostic markers for

EHBDCa still needs investigation. In addition, lack of effective adjuvant therapy against advanced EHBDCa requires establishing new therapeutic methods based on reliable molecular targeting markers; thus, mesothelin could be one of the potential targets for cancer molecular targeting therapy. Recombinant anti-mesothelin immunotoxin SS1P (CAT-5001) and a high affinity chimeric anti-mesothelin monoclonal antibody MORAb-009 recently entered phase II clinical trials (35,36). To evaluate the therapeutic effect of such antibody-based medicine, pathological verification of membranous expression of the target molecule must be performed, because antibody-based drugs can usually access the molecules located on the cell membrane. We believe that luminal membrane expression of mesothelin in EHBDCa would be of clinical benefit not only as a prognostic factor but also as a predictive factor for the eligibility to mesothelin-targeting therapies (13,14,27,37,38).

In conclusion, we demonstrated the clinicopathological significance of the mesothelin expression as an independent prognostic factor. Moreover, identification of luminal membrane or cytoplasmic expression of mesothelin could be a reliable prognostic factor for EHBDCa and might offer a novel therapeutic strategy for patients with EHBDCa, including immunotherapy using peptide vaccine or monoclonal antibody therapy.

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