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Human Equilibrative Nucleoside Transporter 1 Expression in Endoscopic Ultrasonography-Guided Fine-Needle Aspiration Biopsy Samples Is a Strong Predictor of Clinical Response and Survival in the Patients With Pancreatic Ductal Adenocarcinoma Undergoing Gemcitabine-Based Chemoradiotherapy

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(*Pancreas* 2016;45: 761–771)

Objectives: This study aimed to clarify whether pretreatment human equilibrative nucleoside transporter (hENT1) expressions in endoscopic ultrasonography-guided fine-needle aspiration biopsy (EUS-FNAB) specimens obtained from resectable, borderline resectable, and locally advanced unresectable pancreatic ductal adenocarcinoma (PDAC) are concordant with those in the resected specimen after gemcitabine-based chemoradiotherapy (Gem-CRT) and to validate the utility of hENT1 expression using EUS-FNAB samples as a prognostic marker.

Methods: We evaluated the relationship between hENT1 expressions assessed by immunohistochemical staining and clinical outcomes in 51 of 76 patients with PDAC who were diagnosed by EUS-FNAB and received preoperative Gem-CRT.

Results: The concordance rate of hENT1 expressions was 89.2% ($K = 0.681$). Median survival time (month) in the 51 whole patients and 37 patients with resection was significantly longer in hENT1 positive than in hENT1 negative: 25.0 and 30.0 versus 9.0 and 9.0, respectively. A multivariate analysis confirmed that hENT1 expression was an independent prognostic factor in both whole patients and those with resection. Regardless of T3 and T4, hENT1-positive patients with resection had significantly better prognosis than hENT1-negative patients, whose prognosis was similar to those without resection.

Conclusions: The assessment of hENT1 expression using EUS-FNAB samples before Gem-CRT provides important information on patients with PDAC who can benefit from curative-intent resection.

Key Words: EUS-FNAB, hENT1, chemoradiotherapy, gemcitabine, pancreatic ductal adenocarcinoma

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Received for publication March 23, 2015; accepted November 17, 2015.

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This study was supported by in part by Grants-in-Aid for scientific research from Japan Society for the Promotion of Science (25461033).

The authors declare no conflict of interest.

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Gemcitabine (Gem) therapy has been the standard treatment for pancreatic ductal adenocarcinoma (PDAC) since Burris et al¹ reported that Gem offered better overall survival (OS) than fluorouracil. However, its efficacy is limited; only 15% of patients with recurrent and metastatic PDAC² and up to 30% in general³ can be expected to respond to treatment. Because Gem is strongly hydrophilic, passive diffusion through hydrophobic cellular membranes is slow. Efficient permeation of Gem into cells requires specialized integral membrane transporter proteins to cross plasma membranes.⁴ Among these transporters, the major mediators of Gem uptake into human cells are the human equilibrative nucleoside transporter 1 (hENT1) and, to a lesser degree, the human concentrative nucleoside transporter 3.^{5–7}

The hENT1 has been reported as an important predictive marker of Gem-based therapy.⁸ In vitro studies indicated that hENT1 gene expression was positively associated with Gem-chemosensitivity.⁹ High hENT1 expression in resected specimen was also reported to be associated with increased OS in patients with PDAC who received postoperative Gem-based chemotherapy.^{8,10–16} These studies indicate that hENT1 expression is important in predicting the survival of patients with PDAC in the adjuvant setting. However, there have been a few reports describing the impact of hENT1 expression on the outcome after preoperative Gem-based chemoradiotherapy (Gem-CRT) in patients with PDAC. Our previous study showed that hENT1 expression was an independent predictor of OS after neoadjuvant Gem-CRT in the patients with Union Internationale Contrele Cancer (UICC) T3 to T4.¹⁷ We also reported that positive expression of hENT1 in the resected specimen was the significant prognostic factor especially for the treatment of locally unresectable (LUR) PDAC defined by the National Comprehensive Cancer Network (NCCN) guidelines (2010).^{18,19}

Based on these results, pretreatment/preoperative evaluation of hENT1 expression in PDAC tissue can be beneficial in predicting the efficacy of Gem-based therapy before initial treatment. The specimens obtained by endoscopic ultrasound-guided fine-needle aspiration biopsy (EUS-FNAB) might be suitable for evaluating hENT1 expression; however, the immunohistochemical (IHC) analysis of hENT1 expression in the pancreatic tumor tissue taken by EUS-FNAB has not been established. There have been several studies that examined gene expression including hENT1 in pretreated tissue biopsy samples obtained by EUS-FNAB in

patients with unresectable PDAC.²⁰⁻²² Based on genetic analysis of EUS-FNAB tissue samples, it is suggested that hENT1 messenger RNA expression levels might be biomarkers for predicting and monitoring Gem sensitivity in patients with unresectable PDAC.²² The examination of total RNA isolated from EUS-FNAB tissue samples without microdissection has a risk of contaminating cells, which could lead to false results. In contrast, IHC analysis using EUS-FNAB samples can examine cancer-specific expression of hENT1. However, there have been no previous reports performing IHC analysis of hENT1 expression in the pretreatment tissue taken by EUS-FNAB and comparing with posttreatment resected specimens of PDAC. One of the reasons why such studies were rare is the difficulty in obtaining sufficient quantity of cancer cells for IHC analysis because the materials aspirated for analysis are often bloody and contain contamination from gastrointestinal tract epithelium.²³⁻²⁶ Recently, Yamao et al²³ have revealed that EUS-FNAB with rapid on-site evaluation (ROSE) provides more accurate diagnosis than EUS-FNAB without it because a cytopathologist ensures that the samples taken by EUS-FNAB are adequate for assessment. Because the sampling rate of PDAC tissues in our institute has been high owing to the introduction of ROSE, we could retrospectively evaluate the stored cell block specimens for the IHC analysis of hENT1 expression.

The aims of our study were to clarify whether pretreatment hENT1 expressions in the EUS-FNAB specimens are concordant with those in the resected specimen after Gem-CRT and to validate the utility of hENT1 expression using EUS-FNAB samples as a prognostic marker in patients with locally advanced PDAC who underwent Gem-CRT.

MATERIALS AND METHODS

Between February 2005 and November 2011, we had enrolled 117 patients for our Gem-CRT protocol reported previously,^{17,18} who were cytologically or histologically diagnosed as having PDAC and having UICC T3 and T4 tumors determined

by using 64-slice multidetector computed tomography (MDCT). Computed tomography was performed according to a defined pancreas protocol as 4-phasic contrast-enhanced MDCT with thin slices at intervals of 1 mm. Patients were excluded when they showed evident distant metastatic lesions at the time of enrollment. They all gave their written informed consent for inclusion in the study. These patients were also retrospectively reclassified into the 3 resectability groups: resectable (R), borderline resectable (BR), or LUR, according to the NCCN guidelines (2010).¹⁹

Of the 117 patients, 76 were diagnosed with PDAC by cytology and/or histology using EUS-FNAB specimen (Fig. 1). Among 76 cases, 94.7% (n = 72) were diagnosed by cytology, 81.6% (n = 62) were diagnosed by histology, 100% (n = 76) were diagnosed by either of the 2 methods. We retrospectively reviewed the formalin-embedded specimens obtained by EUS-FNAB for these 76 patients, and the adequate amount of histological specimens required for the examination of hENT1 expression could be found in 52 patients (68.4%), all of which could have IHC staining successfully performed. Among these 52 patients, hENT1 positive was found in 34 (65.4%), of whom 29 (85.3%) could receive resection and 5 (14.7%) could not, whereas hENT1 negative was found in 18 (34.6%), of whom 1 was excluded because of refusal of treatment, 8 (47.0%) could receive resection, and 9 (53.0%) could not.

We evaluated the relation between hENT1 expressions and clinical courses in these 51 patients. The study measured intratumoral hENT1 expression, concordance rates of hENT1 expressions of EUS-FNAB specimens with those of resected tumors, and survival analysis based on hENT1 expression of EUS-FNAB specimen.

EUS-FNAB Procedure

Endoscopic ultrasonography was performed using a linear array endoscope (GF-UCP240; Olympus Medical Systems Co, Ltd, Tokyo, Japan), connected to a processor with a color Doppler function (SSD-α10; Hitachi-Aloka Medical, Ltd, Tokyo, Japan).

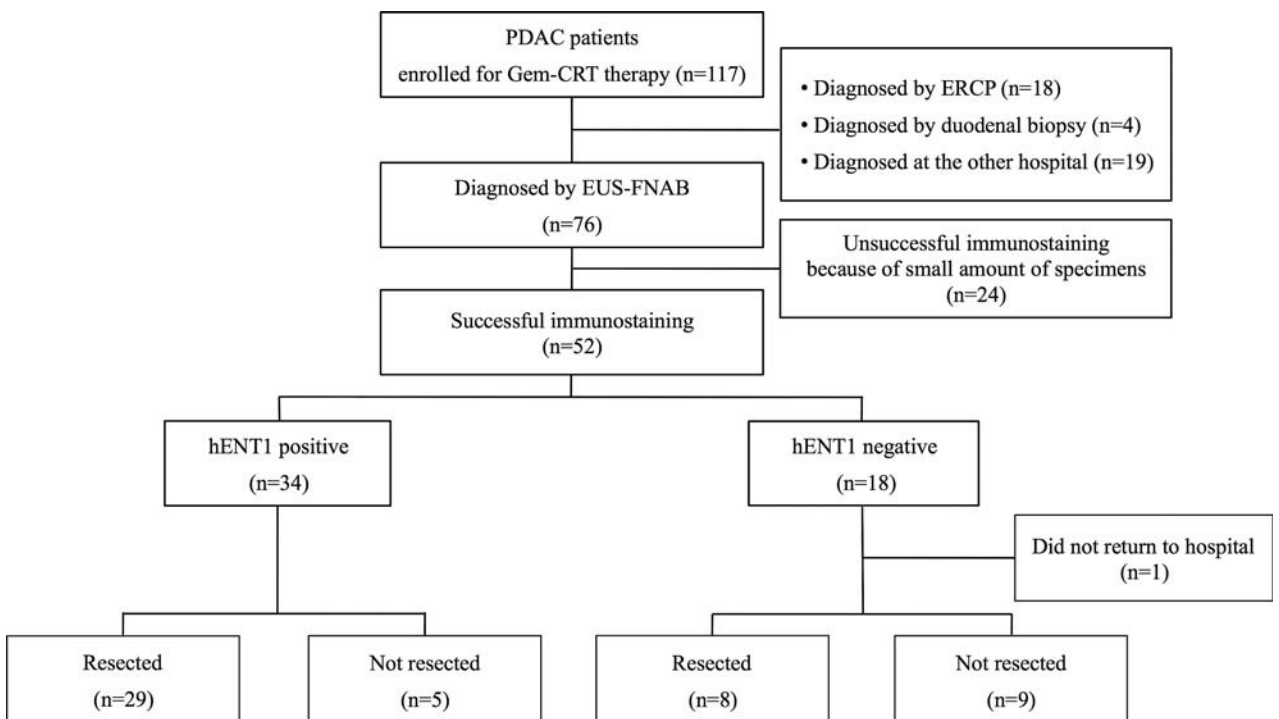


FIGURE 1. Flow diagram of the study participants. ERCP, endoscopic retrograde cholangiopancreatography.

After the tumor was identified using B-mode imaging, we confirmed the absence of vessels in the target area with the color Doppler mode. After we punctured an aspiration needle into the tumor under ultrasonographic guidance, the stylet was pulled out, and the specimen was aspirated with a 20-mL syringe, and then, the needle moved back and forth several times within the tumor. Negative pressure was released before the needle was removed from the tumor. A cytologist immediately examined the specimen with ROSE using rapid stain (Diff-Quik stain; International Reagents, Kobe, Japan) to verify that sufficient sample was obtained. When a tentative diagnosis of malignancy could be made by the on-site evaluation, we finished the EUS-FNAB procedure. If not, we performed an additional 1 to 2 punctures to obtain the diagnosis. The specimen from each EUS-FNAB pass was fixed in alcohol and then stained using the Papanicolaou multichromatic procedure. The remaining material was fixed in 10% formalin and then embedded in paraffin for the cell block analysis to obtain histological diagnosis (hematoxylin and eosin [H&E]).

Immunohistochemical Analysis and Evaluation of hENT1 Expression

After cytological and/or histological diagnosis of PDAC had been confirmed, we retrospectively evaluated 76 stored cell block specimens for the IHC analysis of hENT1 expression: IHC staining was able to be performed successfully on 52 specimens, whereas the remaining 24 failed. The causes of failure were as follows: blood clot alone in 4, normal pancreatic tissue in 9, and an insufficient quantity of malignant cells in 11. For the hENT1 IHC analysis, we used only cell block samples, neither core biopsy samples nor cytological smear.

The cell blocks were sliced into 2- μ m paraffin sections. The 2- μ m sections were used for the assessment of intratumoral hENT1 expressions with immunohistochemistry as well as being stained with H&E. Immunostaining procedure was performed

using the labeled streptavidin-biotin peroxidase complex method with the Benchmark XT auto-immunostaining system (Ventana Japan, Tokyo, Japan). The antigen retrieval step was performed at 90°C, 30 minutes, and then, the sections were incubated in rabbit-derived anti-hENT1 polyclonal antibody (Medical and Biological Laboratories Co, Ltd, Nagoya, Japan). The sections were labeled with an automated immunostaining system with I-View detection kit. Immunostained sections were lightly counterstained with Mayer's hematoxylin.

The resected specimens were fixed in a formalin solution, sliced into 5-mm sections, and embedded in paraffin blocks. A 3- μ m section was obtained from each block and stained with H&E. The sections were routinely examined for pathological differentiation and resection margin status. The histological response of Gem-CRT was evaluated according to histopathological criteria of Evans et al.²⁷ According to the result of H&E staining, the most appropriate one section that contained tumor cells rich enough for immunostaining was stained to assess intratumoral hENT1 expression in the same manner as the EUS-FNAB samples.

Two pathologists (T.S., K.U.) who were blinded to the clinical characteristics of the patients assessed EUS-FNAB samples and resected specimens. Scoring for hENT1 immunostaining was performed on the basis of the relative intensities of staining of the cancer cells, with reference to the normally strong hENT1 staining of cytoplasm within the lymphocytes in the EUS-FNAB samples and of cell membranes within the islets of Langerhans cells in the resected specimen as internal controls, respectively. The degree of hENT1 expression in the resected specimen was determined by the intensity as well as extent of positive staining according to our previous study.¹⁷ A revised scoring system expressing the degree of hENT1 expression in the EUS-FNAB samples was devised based on our previous study; the scoring system is represented as follows: a score ranging from 0 to 3 was assigned based on the intensity of staining, where 0 = no staining, 1 = weakly positive, 2 = moderately positive (same intensity as

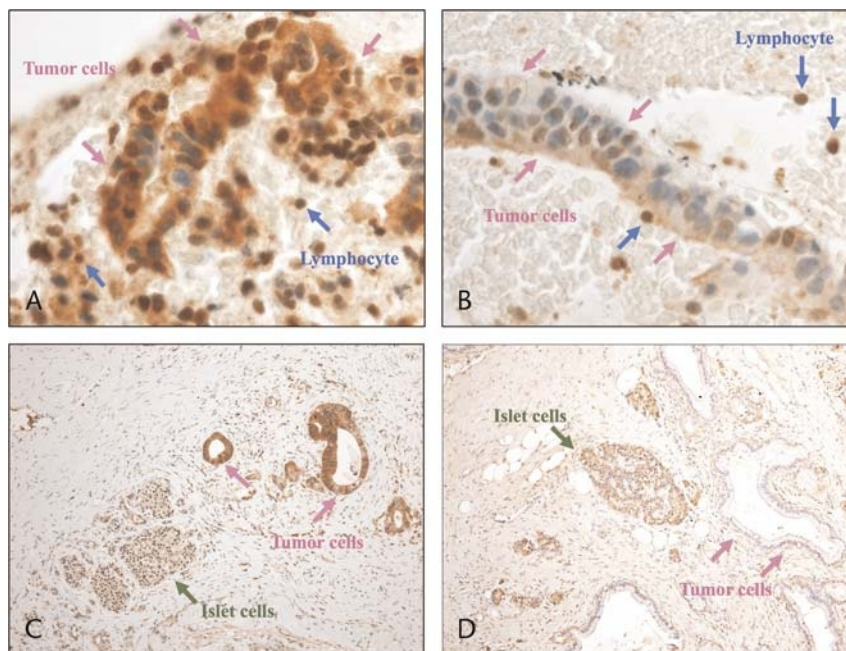


FIGURE 2. Immunohistochemical staining of PDAC for hENT1. A, EUS-FNAB sample showing high hENT1 expression relative to internal control (lymphocyte), "hENT1 positive." B, EUS-FNAB sample showing low hENT1 expression, "hENT1 negative." C, Resected specimen showing high hENT1 expression relative to internal control (islet cells), "hENT1 positive." D, Resected specimen showing low hENT1 expression, "hENT1 negative."

internal control), and 3 = strongly positive. The degree of hENT1 expression was defined as high (neoplastic cells with score 3 accounting for >50% of the total tumor cells), low (neoplastic cells with score 0 or 1 accounting for >50% of the total tumor cells), and intermediate (all other neoplastic cells). We defined high and intermediate staining as hENT1 positive and low staining as hENT1 negative in both EUS-FNAB samples and resected specimens (Fig. 2).

Treatment Protocol

The treatment protocol of Gem-CRT was described by our previous reports.^{17,18} Briefly, the total radiation dose was 45 Gy, delivered in 25 fractions (5 fractions per week), and the patients were administered an infusion of Gem at a dose of 800 mg/m² on days 1, 8, 22, and 29 for 1 cycle. The patients underwent reassessment at 4 to 6 weeks after the completion of Gem-CRT; when we determined that curative-intent resection was possible, they were scheduled to undergo pancreatectomy. At the time of reassessment, especially in the case of LUR patients, we determined that curative-intent resection was possible when the following findings on MDCT were observed: no stenosis or change of shape in the celiac trunk and superior mesenteric artery as well as the absence of metastatic lesions in other distant organs. Even after we decided that the tumor was inoperable, we continued chemotherapy mainly using Gem. Pancreaticoduodenectomy (PD) or distal pancreatectomy was performed as previously described.^{17,18} From 6 weeks after resection, we planned to start the postoperative chemotherapy regimen, consisting of Gem at a dose of 800 mg/m² biweekly for at least 6 months. After pancreatectomy, all patients were evaluated as follows: physical examination every month; laboratory tests including CEA serum levels and carbohydrate antigen (CA) 19-9 (CA19-9) levels every 2 or 3 months; and MDCT every 3 months within 2 years and thereafter every 6 months.^{17,18}

Analysis of Factors Contributing to Survival

We analyzed various clinicopathological factors in the whole patients and those with resection to clarify the significant prognostic factors, including (1) pretreatment factors such as tumor location, tumor size before Gem-CRT, UICC-T classification, resectability according to NCCN guideline 2010, and hENT1 expression of EUS-FNAB samples and (2) posttreatment clinical factors, such as response to Gem-CRT evaluated according to the Response Evaluation Criteria in Solid Tumors (RECIST),²⁸ reduction rate in serum CA19-9 level as previously described,¹⁸ presence of distant metastasis after Gem-CRT, and hENT1 expression of resected specimen.

Statistical Analyses

The results for continuous variables were expressed as mean or median. For the clinicopathological features of the patients, *P* values were calculated by χ^2 test or Fisher exact test, as appropriate. In the whole patients, the date of the initial treatment was chosen as the starting point for the measurement of survival time. The day of final follow-up was December 31, 2013, and there was no loss of follow-up. Survival time was calculated using the Kaplan-Meier method and was compared between the groups using the Wilcoxon test. The factors affecting survival time were analyzed using the multivariate Cox proportional hazard model. Individual variables with a significance of *P* < 0.05 in the univariate Cox proportional hazard model were selected for inclusion into the multivariate analysis. In the multivariate analysis, variables with a significance of *P* < 0.05 were selected. For all statistical tests, a *P* < 0.05 was considered statistically significant. All

statistical analyses were performed using SPSS version 20 software (IBM Inc., Chicago, Ill).

RESULTS

Immunostaining and Patient Background

Patient characteristics are summarized in Table 1. Comparing with the whole patients and those with resection, the resection rate according to tumor location, UICC-T classification, resectability classification, and hENT1 expression in EUS-FNAB samples differed significantly: head versus body/tail (85.3% vs 47.1%, *P* = 0.004), T3 versus T4 (89.2% vs 52.2%, *P* = 0.003), resectable versus borderline resectable versus LUR (40.0% vs 89.3% vs 55.6%, *P* = 0.01), and hENT1 positive versus hENT1 negative (85.3% vs 47.1%, *P* = 0.004). The positive rate of hENT1 expression in EUS-FNAB samples was 66.7% in the whole 51 patients and 78.4% in the 37 patients with resection.

We examined the homology of hENT1 expression between pretreatment samples obtained by EUS-FNAB and resected specimens after Gem-CRT in 37 resected specimens (Table 2). As the status of hENT1 expression in the resected specimens was determined as control, sensitivity, specificity, positive predictive value, negative predictive value, and accuracy of EUS-FNAB samples were 93.1%, 75.0%, 93.1%, 75.0%, and 89.2%, respectively. Therefore, the rate of concordance between EUS-FNAB samples and resected specimens was 89.2% (*K* = 0.681). We examined the characteristics of the 4 patients in whom hENT1 expression differed between EUS-FNAB samples and resected specimen (Table 3). In cases 1 and 2, hENT1 expression was found to be negative (low) in the EUS-FNAB samples, whereas positive (intermediate) in the resected specimen. On the other hand, in cases 3 and 4, it was positive (intermediate) in the EUS-FNAB sample, whereas negative (low) in the resected specimen. When we compared the intensity scores of hENT1 staining between the EUS-FNAB samples and the resected specimen (control) as shown in

TABLE 1. Background Characteristics of the Patients

Characteristic	Whole Patients	Patients With Resection	Resection Rate
	n = 51	n = 37	%
Age, mean (SD)	66.4 (9.6)	66.1 (8.8)	—
Sex			
Male	29 (56.9%)	22 (59.5%)	75.9
Female	22 (43.1%)	15 (40.5%)	68.2
Tumor location			
Head	34 (66.7%)	29 (78.4%)	85.3*
Body/tail	17 (33.3%)	8 (21.6%)	47.1*
UICC-T classification			
T3	28 (54.9%)	25 (67.6%)	89.2*
T4	23 (45.1%)	12 (32.4%)	52.2*
Resectability classification			
R	5 (9.8%)	2 (5.4%)	40.0*
BR	28 (54.9%)	25 (67.6%)	89.3*
LUR	18 (35.3%)	10 (27.0%)	55.6*
hENT1 expression in EUS-FNAB samples			
Positive	34 (66.7%)	29 (78.4%)	85.3*
Negative	17 (33.3%)	8 (21.6%)	47.1*

**P* < 0.05, χ^2 test.

TABLE 2. Homology of hENT1 Expression Between Pretreatment Samples Obtained by EUS-FNAB and Resected Specimens After Gem-CRT ($K = 0.681$)

	FNAB Sample Positive	FNAB Sample Negative
Resected specimen positive	27	2
Resected specimen negative	2	6

Sensitivity, 93.1%; specificity, 75.0%; positive predictive value, 93.1%; negative predictive value, 75.0%; and accuracy, 89.2%.

Table 3, the intensity scores in the resected specimen in all of 4 cases contained more than 2 kinds of intensity with various dominant area, and those in the EUS-FNAB samples contained one or more scores of the resected specimen with dominant area, which was different from the resected specimen. These findings suggested that the discrepancy between EUS-FNAB samples and resected specimen occurred because the intensity of staining and its area in the resected specimen varied widely in these 4 cases.

Patient Characteristics and Effect of Gem-CRT According to hENT1 Expression

Pretreatment clinical factors and the clinical response after Gem-CRT in the whole patients and those with resection are summarized in Table 4. Pretreatment clinical factors in the whole patients and in those with resection did not differ between hENT1-positive and hENT1-negative expression. As for RECIST after Gem-CRT in the whole patients, the percentage of the patients with partial response (PR) and stable disease (SD) was significantly higher in hENT1 positive than in negative: 82.4% versus 52.9% ($P = 0.047$). Distant metastasis after Gem-CRT occurred significantly less frequently in hENT1 positive than in negative: 11.8% versus 47.1% ($P = 0.005$). The incidence of the patients with CA19-9 reduction rate of 50% or more was significantly higher in hENT1 positive than in negative: 64.7% versus 24.5% ($P = 0.006$). In the patients with resection, the incidence of patients with CA19-9 reduction rate of 50% or more was significantly higher in hENT1 positive than in negative: 75.9% versus 37.5% ($P = 0.04$), whereas other factors did not differ between the 2 groups.

Univariable and Multivariable Analyses for Prognostic Factors

In the whole patients, UICC-T classification ($P = 0.002$), hENT1 expression of EUS-FNAB samples ($P < 0.001$), response of Gem-CRT ($P < 0.001$), CA19-9 reduction rate ($P = 0.001$), and distant metastasis after Gem-CRT ($P < 0.001$) were found to be significant in the univariate model; however, in the multivariate model, only hENT1 expression and UICC-T classification were found to be significant independent prognosis factors (Table 5). In the patients who underwent resection, UICC-T classification ($P = 0.015$), hENT1 expression of EUS-FNAB samples ($P < 0.001$), and hENT1 expression of resected specimen ($P < 0.001$) were found to be statistically significant in the univariable analyses; however, again, in the multivariate model, only hENT1 expression and UICC-T classification were found to be significant (Table 5).

In the 51 whole patients and 37 with resection, survival rates were significantly higher in hENT1 positive than in hENT1 negative as shown in Figure 3. Furthermore, we compared survival curves according to hENT1 expression in T3 (Fig. 4A, B) and

TABLE 3. The Characteristics of the 4 Patients in Whom hENT1 Expression Differed Between EUS-FNAB Samples and Resected Specimen

Age	Sex	hENT1 Expression of EUS-FNAB Sample		hENT1 Expression of Resected Specimen		Tumor Size, mm	UICC	Resectability	Tumor Location	Pretreatment CA19-9	Posttreatment CA19-9	Response	Histology	Evans Grade	Survival Time, mo	
		Judge	Score	Judge	Score											
1	49	M	Negative	Score: 1 Low	Positive	Score: 2 > 1 > 3 Intermediate	28	T3	BR	Head	458.0	249.3	SD	Well diff.	Ila	9
2	63	M	Negative	Score: 0 > 1 Low	Positive	Score: 2 > 1 > 0 Intermediate	22	T3	BR	Head	10093.0	940.9	SD	Well diff.	Ila	24
3	77	M	Positive	Score: 2 > 1 Intermediate	Negative	Score: 1 > 2 Low	32	T3	BR	Head	316.5	122.7	SD	Well diff.	Ila	16
4	65	M	Positive	Score: 2 > 1 Intermediate	Negative	Score: 1 > 2 > 0 Low	46	T4	UR	Head	1.0	1.0	SD	Moderate diff.	Ila	12

A score ranging from 0 to 3 for the intensity of hENT1 staining is shown in decreasing order according to the dominant area like intensity: 2 > 1 > 3 (intermediate), which means that score 2 (moderately positive) occupied the most predominant area followed by score 1 (weakly positive) and score 3 (strongly positive). UR indicates unresectable; diff., differentiated.

TABLE 4. Patient Characteristics and Effect of Gem-CRT According to hENT1 Expression

Values	Whole Patients			Patients With Resection		
	hENT1 Positive n = 34	hENT1 Negative n = 17	P	hENT1 Positive n = 29	hENT1 Negative n = 8	P
Pretreatment clinical factors						
Age, mean (SD), y	67.3 (8.5)	64.7 (11.6)	0.357	66.0 (8.1)	66.4 (11.7)	0.917
Sex (male/female)	18/16	11/6	0.552	17/12	5/3	0.221
Tumor size before Gem-CRT, mean (SD), cm	32.5 (9.3)	33.5 (13.0)	0.747	31.8 (9.2)	30.5 (7.5)	0.724
UICC-T classification			0.553			0.612
T3	20	8		19	6	
T4	14	9		10	2	
Resectability			0.373			0.722
R	3	2		2	0	
BR	21	7		19	6	
LUR	10	8		8	2	
Clinical response after Gem-CRT						
Response of Gem-CRT (RECIST)			0.047			0.450
Complete response	0	0		0	0	
PR	4 (11.8%)	0		4 (13.8%)	0	
SD	24 (70.6%)	9 (52.9%)		24 (82.8%)	8 (100%)	
PD	6 (17.6%)	8 (47.1%)		1 (4.4%)	0	
Distant metastasis after Gem-CRT	4 (11.8%)	8 (47.1%)	0.005	0	0	-
CA19-9 levels, median						
Pre-CA19-9, U/mL	313.15	218.6	0.839	309.9	202.75	0.928
Post-CA19-9, U/mL	82.4	249.3	0.100	40.4	134.5	0.346
Degree of reduction rate in CA19-9			0.006			0.040
≥50%	22 (64.7%)	4 (23.5%)		22 (75.9%)	3 (37.5%)	
<50%	12 (35.3%)	13 (76.5%)		7 (24.1%)	5 (62.5%)	

T4 patients (Fig. 5A, B). In T3 patients, the survival rates were significantly higher in hENT1 positive than in hENT1 negative in the whole patients and in those with resection. In T4 patients, the survival rates did not significantly differ between hENT1 positive and negative in the whole patients, whereas in the patients with resection, the survival rates were significantly higher in hENT1 positive than in negative. Interestingly, survival curves in the patients without resection (14 patients in Fig. 3B, 3 in Fig. 4B, and 11 in Fig. 5B) were very similar to those of hENT1 negative with resection (8 in Fig. 3B, 6 in Fig. 4B, 2 in Fig. 5B).

DISCUSSION

The hENT1 expression assessed immunohistochemically in the resected specimen has been proven to be a significant prognostic marker of patients with PDAC undergoing Gem-based adjuvant therapy,^{8,10-17} although the assessment method for grading of expression and reference cells (Langerhans cells or lymphocytes) differed among the studies. In our previous study on the 55 patients using Langerhans cells as a reference,¹⁷ staining intensity and extension of stained tumor cells (I-E) were graded as high (n = 14, 25.5%), intermediate (n = 25, 45.5%), and low (n = 16, 29.0%). High and intermediate were defined as positive (71.0%) and low was defined as negative (29.0%), and survival rate was significantly higher in the hENT1-positive group than in the hENT1-negative group. Using lymphocytes as a reference, Farrell et al¹⁰ reported that I-E was categorized as high (n = 34, 37.4%), low (n = 39, 42.8%), and no staining (n = 18, 19.8%), in which greater than 50% of cells showed no staining and that survival rate was significantly higher in the hENT1 high/low than in the no

staining. Using Langerhans cells as a reference, Nakagawa et al¹⁵ also reported that I-E was graded high (n = 78, 71.6%) and low (n = 31, 28.4%) and that survival rate was significantly higher in the hENT1 high than in the low. Therefore, the proportion of hENT1 expression was similar among these previous 3 studies, although the assessment method based on I-E for grading of expression slightly differed. In contrast, Kawada et al²⁹ revealed that hENT1 expression in the resected specimens was not associated with prognosis in the patients who underwent resection after preoperative Gem-CRT and immediately received postoperative liver perfusion chemotherapy using continuous infusion of 5-fluorouracil (for 28 days) into the hepatic artery and portal vein through a catheter inserted during the surgical procedure. They suggested that 5-FU liver perfusion had a negative impact on the role of hENT1 expression in prognosis.

If pretreatment evaluation of hENT1 expression in PDAC specimen obtained by EUS-FNAB becomes possible without difficulty, it is very useful to predict the efficacy of Gem-based therapy. The IHC analysis of hENT1 expression in the EUS-FNAB specimens has not been established, and thus, we first examined whether pretreatment hENT1 expressions in the EUS-FNAB specimens were concordant with those in the resected specimen after Gem-CRT. As a result, the rate of concordance between them was 89.2%, which is higher than the previous 2 reports concerning the other IHC studies: 86.5% in the study on SMAD4 protein and 73.9% in the study on ZIP4.^{30,31} The reason why the concordance rate in the 3 studies including ours did not reach 100% is unclear.

However, we could identify the features of the 4 patients in whom hENT1 expression differed between EUS-FNAB samples

TABLE 5. Univariate and Multivariate Analyses of Factors Affecting Cox Proportional Hazard Model

	Whole Patients			Patients With Resection		
	Univariate Analysis		Multivariate Analysis	Univariate Analysis		Multivariate Analysis
	HR (95% CI)	P	HR (95% CI)	HR (95% CI)	P	HR (95% CI)
Age, y						
<65	1	0.925	1	0.933 (0.440–1.976)	0.856	
≥65						
Sex						
Male	1	0.954	1	0.943 (0.445–2.002)	0.879	
Female						
Tumor location						
Head	1	0.072	1	0.909 (0.384–2.149)	0.828	
Body/tail						
Tumor size before Gem-CRT, cm						
<3.0	1	0.730	1	1.166 (0.542–2.507)	0.695	
≥3.0						
UICC-T classification						
T3	1	0.002	1	2.629 (1.211–5.707)	0.015	1
T4						
Resectability						
BR/LUR	1	0.936	1	0.396 (0.050–3.144)	0.396	
R						
hENT1 expression of EUS-FNAB samples						
Positive	1	<0.001	1	6.192 (2.439–15.715)	<0.001	1
Negative						
Response of Gem-CRT (RECIST)						
PD	1	<0.001	1	1.100 (0.148–8.171)	0.926	
PR/SD						
Reduction rate in serum CA19-9 level						
≥50%	1	0.001	1	2.137 (0.978–4.670)	0.057	
<50%						
Distant metastasis after Gem-CRT						
Metastasis	1	<0.001	1	—	—	
Nonmetastasis						
hENT1 expression of resected specimen						
Positive	—	—	1	7.791 (2.887–21.022)	<0.001	
Negative						

CI indicates confidence interval; HR, hazard ratio.

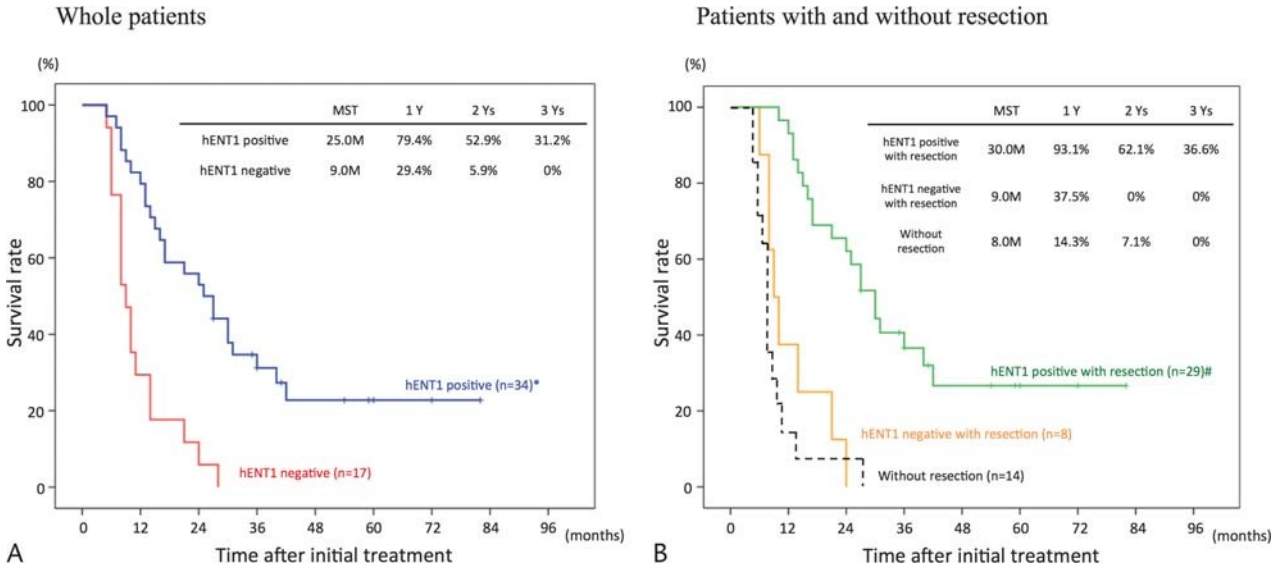


FIGURE 3. Cumulative survival curves according to hENT1 expression. A, Whole patients comparing hENT1 positive (n = 34) and negative (n = 17). B, Patients with resection comparing hENT1 positive (n = 29) and negative (n = 8), and those without resection (n = 14). *P < 0.001 versus hENT1 negative. #P < 0.001 hENT1 negative with resection.

and resected specimen by comparing the intensity scores and its area of hENT1 staining between the EUS-FNAB samples and the resected specimen: the intensity of staining and its area in the resected specimen varied widely, indicating the existence of tumor heterogeneity in these 4 cases. A recent study on the evaluation of Ki-67 index in pancreatic neuroendocrine tumors also demonstrated intratumoral heterogeneity by comparing its index in EUS-FNAB specimens and resected specimens as the criterion standard: concordance rate remained 74.0% using the mean Ki-67 index.³² It is however interesting to note that Gem-CRT did not seem to change the preoperative/postoperative correlation of hENT1 staining.

With the use of EUS-FNAB specimens, our hENT1 IHC analysis could be successfully performed in 68.4% (52/76) among

the cases diagnosed cytologically and/or histologically as PDAC, under the situations that the remaining materials followed by cytological/histological diagnosis were used and that some of adequate samples might be already consumed before the IHC analysis. These results suggested that EUS-FNAB specimens obtained from PDAC were appropriate for IHC analysis of hENT1. In the method similar to ours, which used the remaining samples after diagnosis to evaluate SMAD4 protein, only 44.4% (52/117) could be analyzed.³⁰ It is therefore considered that the success rate of IHC analysis using EUS-FNAB samples obtained from PDAC specimens remains not so high. Concerning the reason why the success rate remains low, Navina et al³³ recently evaluated the adequacy of EUS-FNAB samples of pancreatic masses for theranostic studies by assessing the cellularity of the cytology

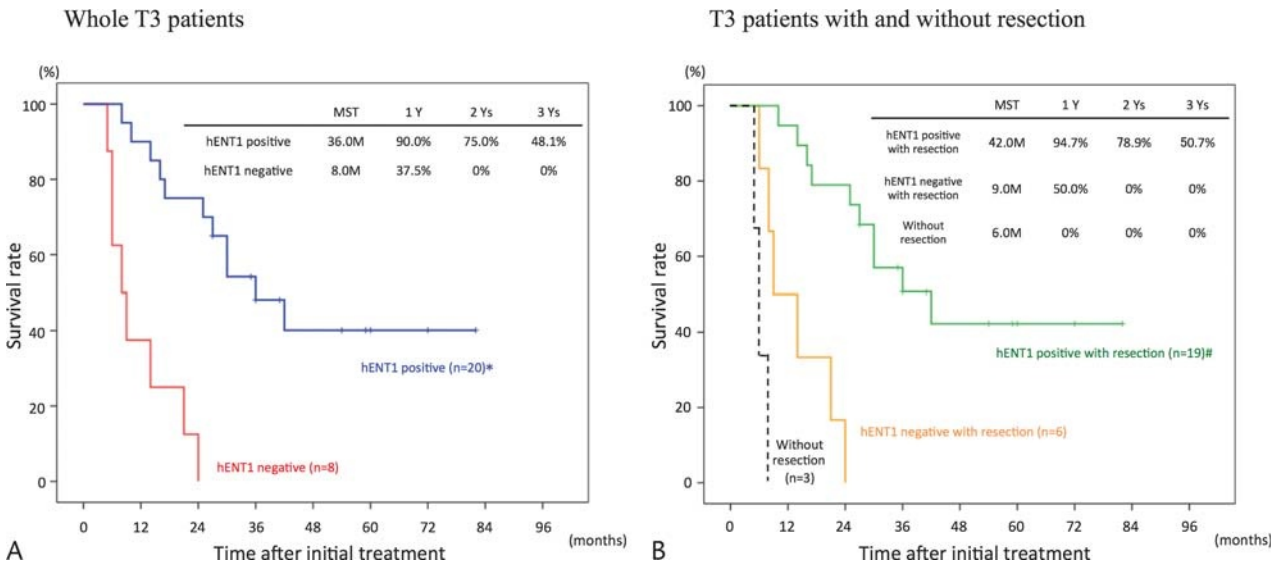


FIGURE 4. Cumulative survival curves in T3 patients according to hENT1 expression. A, Whole T3 patients comparing hENT1 positive (n = 20) and negative (n = 8). B, T3 patients with resection comparing hENT1 positive (n = 19) and negative (n = 6), and those without resection (n = 3). *P < 0.001 versus hENT1 negative. #P < 0.001 hENT1 negative with resection.

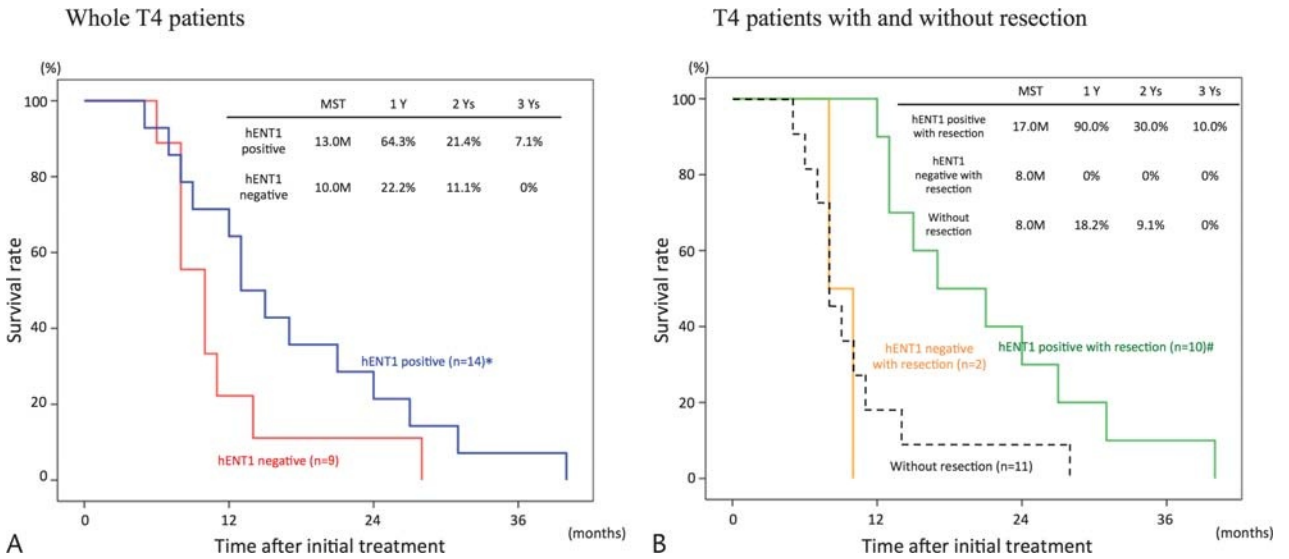


FIGURE 5. Cumulative survival curves in T4 patients according to hENT1 expression. A, Whole T4 patients comparing hENT1 positive (n = 14) and negative (n = 9). B, T4 patients with resection comparing hENT1 positive (n = 10) and negative (n = 2), and those without resection (n = 11). *P = 0.126 versus hENT1 negative. #P < 0.001 hENT1 negative with resection.

material. They retrospectively evaluated 169 EUS-FNAB specimens with positive diagnoses of solid epithelial pancreatic neoplasms (adenocarcinoma, 88%) for smear and cell block cellularity. Cellularity of cell blocks was scored on a scale of 1 to 4 (score 1 for <50 lesional cells; score 2 for 50–100, score 3 for 100–200, and score 4 for >200), and scores of 3 or 4 were deemed adequate for ancillary studies such as IHC analysis. As a result, only 12.4% of the positive cases had a cell block cellularity score that was adequate for theranostic studies. This score was not associated with ROSE, needle gauge, or number of passes. Tumor size and fibrosis score of resected tumors correlated with cellularity, but only larger size in pancreatic neuroendocrine tumors was significantly associated with adequacy. Furthermore, 75 PDAC cases were prospectively evaluated for cellularity score: score 0 in 39%, score 1 to 2 in 49%, and score 3 in 12%. When taking this cellularity score 1 to 3 of 61% and our result of yield 68.4% for hENT1 IHC analysis together, the cellularity score 1 or greater might be enough for hENT1 IHC analysis. Consequently, to enhance the clinical utility of pretreatment/preoperative IHC hENT1 examination, we have to develop a novel method to improve tumor cell yield, including modified cytological techniques and new needle designs.

Serum levels of CA19-9 have been accepted as a measure of pancreatic cancer burden, and the role of CA19-9 has been recently underscored for the evaluation of patients with pretreatment/preoperative therapy before planned surgical resection. Our previous 2 studies, which evaluated the clinical response after Gem-CRT for PDAC according to the hENT1 expression in the resected specimen, revealed that the hENT1-positive group had significantly higher reduction rate of CA19-9 than the hENT1-negative group, although RECIST did not differ between the 2 groups.^{17,18} In our present study using pretreatment/preoperative EUS-FNAB samples in the whole patients, the incidence of the patients with CA19-9 reduction rate of 50% or more was significantly higher in the hENT1-positive group than in the hENT1-negative group, and percentage of the patients with PR and SD in RECIST after Gem-CRT was significantly higher in the hENT1-positive than in the hENT1-negative group. In the patients with resection, however, RECIST did not differ between the 2 groups. Concerning the reason why RECIST results differed between the whole patients and those with resection, 47.1% (8/17) of hENT1 negative showed

PD after Gem-CRT, and all of them could not receive pancreatectomy, whereas only 18.6% (6/34) of hENT1 positive showed PD, and one of them could receive pancreatectomy. It was therefore considered that hENT1 expression was not associated with RECIST in the patients with resection. Other than our studies comparing the clinical response between hENT1 positive and negative, Poplin et al³⁴ evaluated clinical response using RECIST and survival in patients with metastatic PDAC, and hENT1 status had no influence on RECIST and survival (median survival time): the percentage of PR/complete response was 15.5% (9/58), and median survival time was 5.2 months in hENT1 high, whereas 26.3% (30/118) and 6.1 months, respectively, in hENT1 low. They considered that the role of hENT1 was less important in metastatic disease than after surgery with a presumed micrometastatic state. In contrast, our study included the patients with locally advanced (T3/T4) PDAC without distant metastasis at the time of enrollment, and at the time of reassessment (approximately 2–3 months after enrollment), distant metastasis became apparent in 23.5% (12/51) of the patients: hENT1 positive (n = 4) and negative (n = 8). These 12 patients died within 12 months regardless of hENT1 expression.

As for Gem-based pretreatment studies on hENT1 expression in PDAC, to the best of our knowledge, there have been 4 studies including our study: the clinical outcomes of patients undergoing Gem-CRT could be predicted by IHC analysis of hENT1 in EUS-FNAB samples obtained from T3/T4 (R/BR/LUR) PDAC. The one study, which evaluated messenger RNA expression levels of hENT1 using EUS-FNAB specimens obtained from patients with stage III/IV inoperable (LUR and metastatic) PDAC, did not show that its expression levels influenced survival.²² The remaining 2 studies on IHC hENT1 evaluation in PDAC, of which one used biopsy specimens of metastatic lesions³³ and the other used biopsy specimens from the primary and metastatic lesions in stage III/IV inoperable (LUR and metastatic) patients,³⁵ did not demonstrate any significant differences in prognosis between the high- and low-hENT1 subgroups either. The reason for the conflicting results between our study and the other 3 probably is that the other 3 studies included only inoperable patients who had basically poor prognosis in itself, whereas ours included the patients with locally advanced (T3/T4) PDAC without distant metastasis. It is considered that tumor progression

influences the role of hENT1 expression in clinical response as well as prognosis in patients with PDAC, and we therefore compared survival curves according to hENT1 expression in T3 (R/BR) and T4 (BR/LUR) patients. In T3, prognosis was significantly better in hENT1 positive in the whole patients and in those with resection. In T4, it did not significantly differ between hENT1 positive and negative in the whole patients, whereas it was significantly better in hENT1 positive in those with resection. These results indicate that the role of hENT1 expression in Gem-based treatment becomes less important as tumor progresses.

Our treatment protocol of Gem-CRT for patients with locally advanced (T3/T4) PDAC was conducted for aiming to achieve curative-intent resection after reassessment, even though it was determined initially LUR. Therefore, we have to clarify the significance of preoperative/pre-treatment assessment of hENT1 expression using EUS-FNAB specimens based on our results: its assessment identifies patients with PDAC who can benefit from curative-intent resection followed by Gem-based adjuvant therapy. Regardless of T3 and T4 tumors, hENT1-positive patients who underwent curative-intent resection had significantly better prognosis compared with hENT1-negative patients with resection, whose prognosis was similar to those without resection. To improve the prognosis in hENT1-negative patients, a novel regimen other than Gem-based treatment needs to be further investigated.

In conclusion, pre-treatment hENT1 expressions in the EUS-FNAB specimens are concordant with those in the resected specimen after Gem-CRT, and its assessment before Gem-CRT provides us the important information on patients with PDAC who can benefit from curative-intent resection followed by Gem-based adjuvant therapy.

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