

# Membrane mesothelin expression positivity is associated with poor clinical outcome of luminal-type breast cancer

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**Abstract.** Mesothelin is expressed in various types of malignant tumors. The present study immunohistochemically investigated mesothelin expression and its clinicopathological significance in each subtype of breast cancer, with special reference to its cellular localization, in particular, membrane mesothelin expression. Using tissue specimens from 482 patients with breast cancer, immunohistochemistry was used to study mesothelin expression and help classify its localization as membrane or cytoplasmic expression. Mesothelin expression was detected in 77 (16.0%) cases and was the highest in triple-negative breast cancer (31/75; 41.3%), followed by human epithelial growth factor receptor type 2 type (6/33, 18.2%) and luminal type (36/374; 9.6%). Among the 482 cases, membrane mesothelin expression was detected in 73 cases and was significantly associated with a negative hormone receptor status, higher Ki-67 labeling index, nuclear grade 3 and a lower relapse-free survival rate. Cytoplasmic mesothelin expression was not significantly associated with a lower relapse-free survival rate (P=0.058). In the 343 cases of luminal type, the membrane mesothelin expression-positive

group had significantly worse prognosis than the membrane mesothelin-expression-negative group (P=0.042). There was no significant difference in the relapse-free survival rate according to the membrane mesothelin expression status in the triple-negative type and other types. It was suggested that membrane mesothelin expression in luminal type breast cancer is associated with a lower rate of relapse-free survival.

## Introduction

Mesothelin is a 40-kDa cell surface glycoprotein that is expressed on normal mesothelial cells that line the pleura, pericardium, and peritoneum (1). Mesothelin is overexpressed in various types of malignant tumors, including malignant mesothelioma, ovarian cancer, and pancreatic cancer (2-4). The full-length human mesothelin gene encodes a 71-kDa precursor protein, which can be physiologically cleaved by some fern-like proteases into a membrane-bound 40-kDa C-terminal fragment and a 31-kDa N-terminal fragment, which is secreted into the blood (5). The 40-kDa C-terminal fragment, mesothelin, is attached to the cell membrane through a glycosyl-phosphatidylinositol anchor (5,6).

The biological functions of mesothelin are unclear, although recent studies have suggested that overexpression of mesothelin increases cell proliferation and migration (7). Furthermore, positive mesothelin expression (ME) is associated with an unfavorable prognosis in pancreatic cancer, gastric cancer, and colorectal cancer (4,8-10).

The number of studies on ME in breast cancer is limited. Wang *et al* (11) suggested that the ME level in invasive breast cancer tissue was inversely correlated with overall survival (OS). Furthermore, based on subtype classification, Parinyanitikul *et al* (12) and Bayoglu *et al* (13) showed that mesothelin was overexpressed in most triple-negative breast cancers (TNBCs), but was not correlated with survival outcomes in TNBC. Tchou *et al* (14) demonstrated that ME-positivity was rare and was not correlated with patient outcome in estrogen receptor (ER)-positive or human epithelial growth factor receptor type 2 (HER2)-positive breast cancers. Moreover, Einama *et al* (9) reported that the localization of ME was related to OS in gastric cancer. To the best of our knowledge, there have been no previous studies on the

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**Abbreviations:** CAR, chimeric antibody receptor; CME, cytoplasmic mesothelin expression; CTL, cytotoxic T lymphocytes; ER, estrogen receptor; FISH, fluorescence *in situ* hybridization; HER2, human epithelial growth factor receptor type 2; IHC, immunohistochemically; LI, labeling index; Ly, lymphatic invasion; ME, mesothelin expression; MME, membrane mesothelin expression; NG, nuclear grade; OS, overall survival; PgR, progesterone receptor; RFS, relapse-free survival; TNBCs, triple-negative breast cancers; V, vascular invasion

**Key words:** breast cancer, mesothelin, membrane, luminal type, triple-negative breast cancers

relationship between the cellular localization of mesothelin and the prognosis of breast cancer patients.

In this study, we investigated ME in each subtype of breast cancer using immunohistochemistry, with special reference to its cellular localization, and analyzed its clinicopathological significance, including patient outcome.

## Materials and methods

**Ethics approval and consent to participate.** This study was performed in accordance with the Declaration of Helsinki and was approved by the Institutional Review Board of the National Defense Medical College (registration no. 3003). All patients agreed to participate in this study, and written informed consent was obtained from all patients.

**Patients.** The subjects of this study comprised 482 patients who underwent radical surgery for primary breast cancer between 2002 and 2013. Patients with stage IV and non-primary breast cancer were excluded. The ER and progesterone receptor (PgR) statuses of the tumor were assessed using immunohistochemistry and defined as positive if 1% or more of the constituent carcinoma cells were immunoreactive. The HER2 status of the tumor was evaluated immunohistochemically (IHC), and fluorescence in situ hybridization (FISH) was performed in cases with a score of 2+. HER2 was considered positive when the IHC score was 3+ or FISH was positive according to the 2013 American Society of Clinical Oncology/College of American Pathologists guidelines. The nuclear grade (NG) was determined by the sum of the nuclear atypia score and the mitosis count score (15). Ki-67 was evaluated according to the recommendation of the Breast Cancer Working Group (16), and the Ki-67 labeling index (LI) was defined as high if 14% or more of the constituent carcinoma cells were immunoreactive (17). Breast cancers were classified into four subtypes according to the St. Gallen consensus as follows (18): Luminal type, ER+ and/or PgR+/HER2-; luminal HER2 type, ER+ and/or PgR+/HER2+; HER2 type, ER-/PgR-/HER2+; TNBC type, ER-/PgR-/HER2-.

**Immunohistochemistry.** Tissue microarrays of 482 breast cancer patients were used in this study. Sections with a thickness of 4  $\mu$ m were cut from formalin-fixed paraffin-embedded tissue blocks and mounted on charged glass slides, deparaffinized, and rehydrated through a graded ethanol series. For antigen retrieval, a target retrieval solution at pH 9.0 (catalogue number, S2368; Dako Japan) was used, and the slides were boiled in a pressure cooker (Pascal Pressure Cooker, Model: S2800; Dako) at 125°C for 3 min. Endogenous peroxidase was blocked with 0.3% hydrogen peroxidase. The slides were incubated with a 1:50 dilution of a mouse monoclonal antibody in mesothelin (clone 5B2, diluted 1:50; Novocastra) at room temperature for 30 min, and then reacted with a dextran polymer reagent combined with secondary antibodies and peroxidase (Envision/HRP; Dako) for 30 min at room temperature. Specific antigen-antibody reactions were visualized with 0.2% diaminobenzidine tetrahydrochloride and hydrogen peroxide. Slides were counterstained with hematoxylin for 10 min and then rinsed gently in reagent quality water.

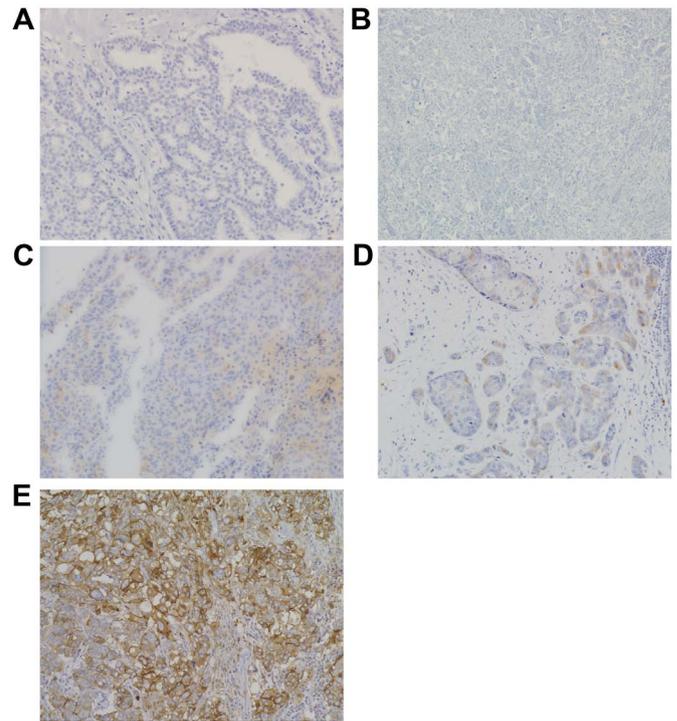


Figure 1. Variations in mesothelin expression and its cellular localization in breast cancer. Immunohistochemical staining for mesothelin. Magnification, x200. (A) No mesothelin staining in the normal breast tissue. (B) No mesothelin expression in breast cancer. (C) Scant cytoplasmic staining of mesothelin (1+). No mesothelin expression in the membrane. (D) Incomplete mesothelin expression in the membrane (1+) and in the cytoplasm with granular staining (2+). (E) Mesothelin expression in the membrane (2+) and cytoplasm (2+).

**Immunohistochemical evaluation.** All assessments were made on the tumor region of the specimen (x200). Each slide was independently evaluated by two observers who were blinded to the clinical outcomes (Y.Y. and T.E.). Immunostaining for mesothelin was evaluated for both the proportion and staining intensity of tumor cells in each case. The proportion of ME-positive cancer cells was assessed as 1-10%, >10-50%, and >50%. The staining intensity was evaluated as weak (+1) or moderate to strong (+2). The staining proportion and intensity were judged separately between the membrane and cytoplasm (Fig. 1). No mesothelin staining was observed in the normal breast tissue. ME was defined as positive when the percentage of positive cells was  $\geq$ 1% of the tumor cells, regardless of the intensity. Furthermore, among the ME-positive cases, the staining localization of mesothelin was evaluated as membrane, cytoplasm, or both. When the entire circumference of the membrane was evenly or partially stained throughout the whole section, 'membrane mesothelin expression (MME)' was defined as positive. When cytoplasmic staining, including cytoplasmic granular staining, was clearly observed in  $\geq$ 1% of the tumor cells, 'cytoplasmic mesothelin expression (CME)' was defined as positive.

**Statistical analysis.** The  $\chi^2$  test or Fisher's exact test were used to determine the correlation between ME and clinicopathological parameters. Survival curves were drawn using the Kaplan-Meier method, and compared using the log-rank test. The prognostic implications of these parameters were analyzed using Cox's univariate and multivariate proportional

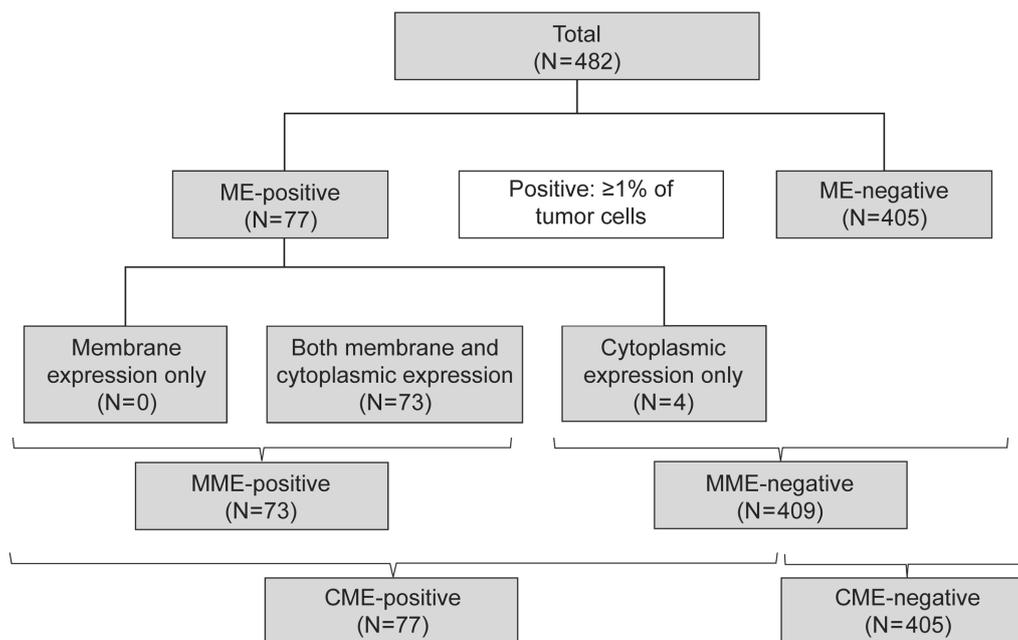


Figure 2. Flow chart for the evaluation of ME. CME, cytoplasmic mesothelin expression; ME, mesothelin expression; MME, membrane mesothelin expression.

hazards models. All differences were considered significant at a P-value <0.05. Statistical analyses were performed using JMP® 14 (SAS Institute Inc.).

## Results

**Clinicopathological analysis of mesothelin expression.** Representative membrane and cytoplasmic ME are shown in Fig. 1. Of the 482 breast cancer tissue samples, ME-positivity was detected in 77 cases (16.0%), MME-positivity in 73, and CME-positivity in 77, which included all 73 MME-positive cases (Fig. 2).

The total clinicopathological demography of patients is shown in Table I. The mean patient age was 59.1 years [ $\pm 11.3$  standard deviation (SD)]. Total mastectomies were performed on 226 patients (46.9%), and partial mastectomies were performed on 256 patients (53.1%). Luminal type breast cancer was the most common (71.2%), followed by TNBC (15.6%). Neoadjuvant chemotherapy was administered to 31 patients, while adjuvant chemotherapy was administered to 190 patients (including administration of trastuzumab to 15 patients). Adjuvant endocrine therapy was administered to 311 patients, and postoperative radiation therapy was administered to 209 patients. A total of 70 patients had a relapse of breast cancer, with a median relapse-free survival (RFS) of 7.5 years ( $\pm 3.9$  SD).

The correlation between ME and clinicopathological characteristics is summarized in Table II. ME was not correlated with age, pT, pN, pStage, lymphatic invasion (Ly), or vascular invasion (V). ME-positivity was more frequent in NG 3 cases (58/221, 26.2%) than in NG 1/2 cases (19/261, 7.3%) ( $P < 0.001$ ), more frequent in ER-negative cases (43/115, 37.4%) than in ER-positive cases (34/367, 9.3%) ( $P < 0.001$ ), and more frequent in PgR-negative cases (45/139, 32.4%) than in PgR-positive cases (32/343, 9.3%) ( $P < 0.001$ ). The mean Ki-67 LI was also higher in ME-positive cases ( $27.6\% \pm 26.5$  SD) than in negative

cases ( $13.4\% \pm 13.9$  SD) ( $P < 0.001$ ), while HER2 positivity was similar between the ME-positive and negative groups. Neoadjuvant chemotherapy had no significant impact on ME positivity ( $P = 0.629$ ) or other clinicopathological factors, such as Ki-67. Furthermore, in these 31 patients, ME was not associated with the pathological response after neoadjuvant chemotherapy. ME-positivity was the highest in TNBC (44.0%), followed by HER2 (21.2%), luminal (10.5%), and luminal HER2 (3.2%) subtypes.

**Survival analysis associated with mesothelin expression.** In analysis of RFS, the MME-positive group had a significantly worse outcome than the MME-negative group ( $P = 0.027$ ). In contrast, the CME-positive group had a relatively worse RFS than the CME-negative group ( $P = 0.058$ ) (Fig. 3). In the analysis of the luminal subtype, the MME-positive group had a significantly worse prognosis than the MME-negative group ( $P = 0.034$ ). The 5-year RFS rates of the MME-positive and negative groups with luminal subtype were 82.6 and 94.1%, respectively. There were no significant differences in RFS curves between the MME-positive and negative patients in the TNBC subtype and other subtypes. The 5-year RFS rates of the ME-positive and negative groups were 80.7 and 77.5% with TNBC, and 85.7 and 88.8% with the other types, respectively (Fig. 4).

In terms of OS, the MME-positive and CME-positive groups had a significantly worse outcome than the corresponding MME-negative and CME-negative groups, respectively ( $P = 0.037$  and  $P = 0.028$ , respectively) (Fig. 5). There was no significant difference in OS curves between the MME-positive and negative patients in any of the included subtypes. In the analysis of the luminal subtype, the MME-positive group had a relatively worse prognosis than the MME-negative group ( $P = 0.068$ ).

We performed univariate analyses of all 482 cases using the Cox proportional hazards model and found that pT, pN,

Table I. Demographics of 482 patients with breast cancer in the present study.

Parameter	Value
Age, years (mean ± SD)	59.1±11.3
Surgery, n (%)	
Mastectomy	226 (46.9)
Partial resection	256 (53.1)
pT, n (%)	
Tis	1 (0.2)
T1	267 (55.4)
T2	195 (40.5)
T3	19 (3.9)
pN, n (%)	
N0	304 (63.1)
N1	124 (25.7)
N2	34 (7.1)
N3	20 (4.1)
Subtype, n (%)	
Luminal HER2-	343 (71.2)
Luminal HER2+	33 (6.8)
HER2	31 (6.4)
TNBC	75 (15.6)
Neoadjuvant chemotherapy, n (%)	
Yes	31 (6.4)
No	451 (93.6)
Adjuvant therapy, n (%)	
Chemotherapy	185 (38.4)
Anti-HER2 therapy	15 (3.1)
Endocrine therapy	311 (64.5)
Radiation therapy	209 (43.4)
Postoperative radiation therapy, n (%)	
Yes	209 (43.4)
No	273 (56.6)
Recurrence, n (%)	
Yes	70 (14.5)
No	412 (85.5)
Death, n (%)	
Yes	51 (10.6)
No	431 (89.4)

HER2, human epidermal growth factor receptor type 2; SD, standard deviation; TNBC, triple-negative breast cancer.

Ly, V, ER, PgR, Ki-67 LI, NG and MME significantly correlated with the risk of relapse, whereas CME did not correlate with RFS. Furthermore, in Cox multivariate analysis, pT and pN were finally chosen as independent prognostic factors for RFS (Table III). In addition, Cox's univariate and multivariate analyses were performed for the 343 luminal type cases (Table IV). Univariate analyses showed that pT, pN, Ly, V and NG significantly correlated with the risk of relapse, whereas MME had no independent impact ( $P=0.062$ ). In

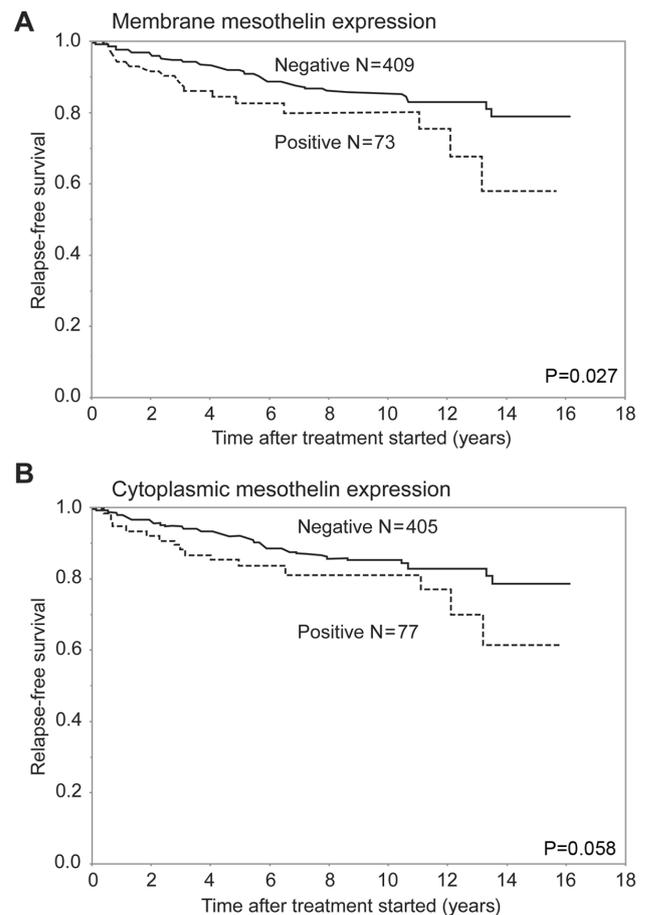


Figure 3. RFS curves for patients with breast cancer after surgical therapy, stratified by mesothelin expression status. (A) Curves for membrane mesothelin expression-positive and -negative groups, and (B) curves for cytoplasmic mesothelin expression-positive and -negative groups. The membrane expression-positive group had a less favorable outcome than the membrane expression-negative group ( $P=0.027$ ). However, cytoplasmic mesothelin expression was not significantly associated with RFS ( $P=0.058$ ). RFS, relapse-free survival.

multivariate analysis, pT and pN were independent prognostic factors for RFS.

## Discussion

In the current study, breast cancer cases were divided into ME-positive and ME-negative groups. In the ME-positive cases, the cellular localization was further divided into MME and CME, and the relationship between ME patterns and clinicopathological factors, including prognosis, was retrospectively investigated. Our results demonstrated that ME was related to conventional prognostic factors, including negative ER, negative PgR, higher Ki-67 LI, and higher NG. Furthermore, we revealed that MME-positivity was associated with lower RFS and OS rates. These results suggest that ME-positivity was an unfavorable prognostic factor in patients with breast cancer, as well as some other cancer types (4,8-10).

In the present study, we showed that MME in the luminal type, but not in other subtypes, was significantly associated with poor RFS. Several previous studies have reported that the ME-positive rates in TNBC were 30-40%, which was in

Table II. Association between mesothelin expression and clinicopathological parameters in 482 patients with breast cancer.

Clinicopathological parameters	Total (n=482)	MME			CME		
		Positive	Negative	P-value	Positive	Negative	P-value
Age, years (mean ± SD) <sup>a</sup>		59.5±11.5	59.1±11.3	0.766	60.1±11.5	58.9±11.3	0.429
pT, n (%)							
Tis	1	0 (0.0)	1 (100.0)	0.071	0 (0.0)	1 (100.0)	0.071
T1	267	32 (12.0)	235 (88.0)		34 (12.7)	233 (87.3)	
T2	195	35 (17.9)	160 (82.1)		37 (19.0)	158 (81.0)	
T3	19	6 (31.6)	13 (68.4)		6 (31.6)	13 (68.4)	
pN, n (%)							
N0	304	45 (14.8)	259 (85.2)	0.433	49 (16.1)	255 (83.9)	0.464
N1	124	16 (12.9)	108 (87.1)		16 (12.9)	108 (87.1)	
N2	34	8 (23.5)	26 (76.5)		8 (23.5)	26 (76.5)	
N3	20	4 (20.0)	16 (80.0)		4 (20.0)	16 (80.0)	
pStage, n (%)							
0	1	0 (0.0)	1 (100.0)	0.298	0 (0.0)	1 (100.0)	0.405
1	204	28 (13.7)	176 (86.3)		30 (14.7)	174 (85.3)	
2	222	32 (14.4)	190 (85.6)		34 (15.3)	188 (84.7)	
3	55	13 (23.6)	190 (76.4)		13 (23.6)	42 (76.4)	
NG, n (%)							
1	119	12 (10.1)	107 (89.9)	<0.001	13 (10.9)	106 (89.1)	<0.001
2	142	6 (4.2)	136 (95.8)		6 (4.2)	136 (95.8)	
3	221	55 (24.9)	166 (75.1)		58 (26.2)	163 (73.8)	
Ly, n (%)							
Negative	197	31 (15.7)	166 (84.3)	0.764	31 (15.7)	166 (84.3)	0.965
Positive	285	42 (14.7)	243 (85.3)		46 (16.1)	239 (83.9)	
V, n (%)							
Negative	312	48 (15.4)	264 (84.6)	0.843	51 (16.3)	261 (83.7)	0.763
Positive	170	25 (14.7)	145 (85.3)		26 (15.3)	144 (84.7)	
ER, n (%)							
Negative	115	40 (34.8)	75 (65.2)	<0.001	43 (37.4)	72 (62.6)	<0.001
Positive	367	33 (9.0)	334 (91.0)		34 (9.3)	333 (90.7)	
PgR, n (%)							
Negative	154	42 (27.3)	112 (72.7)	<0.001	45 (29.2)	109 (70.8)	<0.001
Positive	328	31 (9.5)	297 (90.5)		32 (9.8)	296 (90.2)	
HER2, n (%)							
Negative	418	66 (15.8)	352 (84.2)	0.313	69 (16.5)	349 (83.5)	0.415
Positive	64	7 (10.9)	57 (89.1)		8 (12.5)	56 (87.5)	
Ki-67 labeling index, % (mean ± SD) <sup>a</sup>		28.7±26.7	13.4±13.8	<0.001	27.6±26.5	13.4±13.9	<0.001
<14, n (%)	301	31 (10.3)	270 (89.7)	<0.001	34 (11.3)	267 (88.7)	<0.001
≥14, n (%)	181	42 (23.2)	139 (76.8)		43 (23.8)	138 (76.2)	
Neoadjuvant chemotherapy, n (%)							
Yes	31	4 (12.9)	27 (87.1)	0.719	4 (12.9)	27 (87.1)	0.629
No	451	69 (15.3)	382 (84.7)		73 (16.2)	378 (83.8)	
Subtype, n (%)							
Luminal type	343	35 (10.2)	308 (89.8)	<0.001	36 (10.5)	307 (89.5)	<0.001
Luminal HER2 type	31	1 (3.2)	30 (96.8)		1 (3.2)	30 (96.8)	
HER2 type	33	6 (18.2)	27 (81.8)		7 (21.2)	26 (78.8)	
TNBC type	75	31 (41.3)	44 (58.7)		33 (44.0)	42 (56.0)	

<sup>a</sup>Student's t-test. CME, cytoplasmic mesothelin expression; ER, estrogen receptor; HER2, human epidermal growth factor receptor type 2; Ly, lymphatic invasion; MME, membrane mesothelin expression; NG, nuclear grade; PgR, progesterone receptor; SD, standard deviation; TNBC, triple-negative breast cancer; V, vascular invasion.

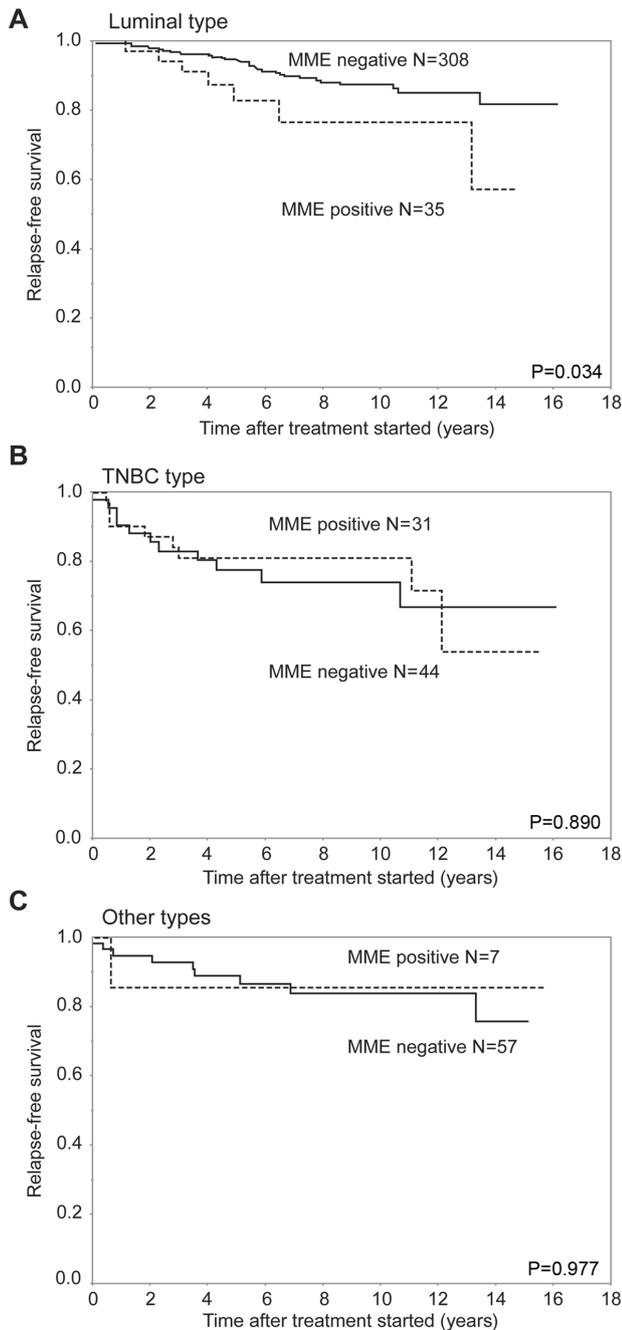


Figure 4. RFS for patients with each subtype of breast cancer after surgical therapy, stratified by the MME status. (A) Luminal type, (B) TNBC type and (C) other types. MME in the luminal subtype was associated with RFS ( $P=0.034$ ) but this was not the case for MME in the TNBC subtype and other types ( $P=0.890$  and  $P=0.977$ , respectively). MME, membrane mesothelin expression; RFS, relapse-free survival; TNBC, triple-negative breast cancer.

agreement with the results of our study (12-14,19). TNBC has a relatively higher relapse rate and worse OS rates than the other subtypes of breast cancer (20). Moreover, since there are few effective therapies that can improve the prognosis of TNBC patients, mesothelin has been researched as a potential candidate for targeted therapies. However, whether ME correlates with survival outcomes in TNBC remains unclear (12).

However, there were few studies on the clinicopathological implications of ME in the luminal type, which comprises 60-70% of all breast cancers. In the luminal type, the frequency

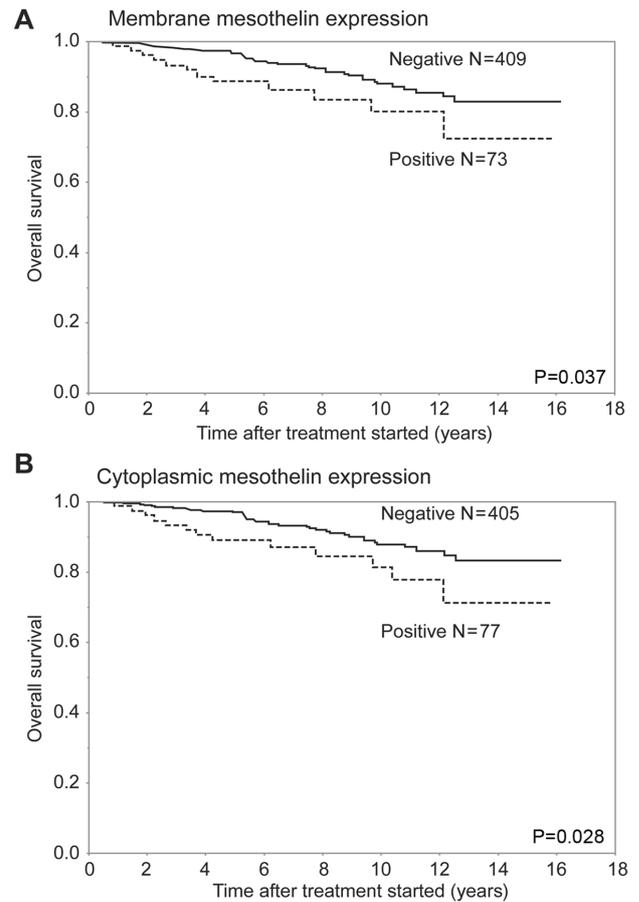


Figure 5. Overall survival curves for patients with breast cancer according to subtype after surgical therapy, stratified by the mesothelin expression status. (A) Membrane mesothelin expression. (B) Cytoplasmic mesothelin expression. Both membrane and cytoplasmic mesothelin expression-positive groups exhibited significantly less favorable outcomes than the membrane and cytoplasmic mesothelin expression-negative groups, respectively ( $P=0.037$  and  $P=0.028$ , respectively).

of ME-positivity was lower than that of TNBC (10.2 vs. 41.3%); however, MME-positivity was associated with a lower RFS rate, and consequently, MME might be useful as a biomarker to predict poor prognosis in luminal type breast cancer. In this group, Cox's multivariate analysis indicated that pT and pN were independent prognostic factors, whereas MME was not, likely because it was strongly related to biological factors, such as negative ER, negative PgR, a higher Ki-67 LI, and a higher NG, and not to anatomical factors, such as pT and pN. These results suggest that MME was not superior to pT and pN as a prognostic factor. ME was defined positive when the percentage of positive cells was  $\geq 1\%$  of tumor cells in this study. If we had defined ME-positivity more strictly, MME could have become an independent prognostic factor. Therefore, additional studies are necessary to determine whether ME can be used as a powerful prognostic factor.

The biological function of mesothelin remains unclear. Recent studies have suggested that overexpression of mesothelin increased cell proliferation and migration (7,21). Moreover, it has also been suggested that mesothelin can elicit cytotoxic T lymphocyte (CTL) responses, and could efficiently activate CTL to lyse human tumors. Previous studies have shown that response to chemotherapy was

Table III. Cox's univariate and multivariate analysis including clinicopathological parameters and mesothelin expression of recurrence-free survival in 482 patients with breast cancer.

Parameter	Univariate		Multivariate	
	HR (95% CI)	P-value	HR (95% CI)	P-value
pT				
Tis, T1	1	<0.001	1	0.003
T2, T3	3.64 (2.19-6.32)		2.27 (1.31-4.08)	
pN				
Negative	1	<0.001	1	0.002
Positive	3.76 (2.30-6.35)		2.78 (1.61-4.84)	
NG				
1,2	1	<0.001	1	0.471
3	2.26 (1.40-3.73)		1.04 (0.57-1.92)	
Ly				
Negative	1	0.002	1	0.646
Positive	2.32 (1.34-4.27)		1.16 (0.63-2.22)	
V				
Negative	1	0.003	1	0.066
Positive	2.05 (1.28-3.30)		1.59 (0.97-2.60)	
ER				
Negative	1	0.003	1	0.497
Positive	0.47 (0.29-0.77)		0.72 (0.35-1.68)	
PgR				
Negative	1	0.007	1	0.412
Positive	0.52 (0.32-0.84)		0.72 (0.35-1.61)	
Ki-67 labeling index, %				
<14	1	0.035	1	0.374
≥14	1.68 (1.04-2.69)		1.26 (0.75-2.10)	
MME				
Negative	1	0.040	1	0.659
Positive	1.86 (1.03-3.17)		1.11 (0.57-2.07)	
CME				
Negative	1	0.075		
Positive	1.71 (0.94-2.91)			

CI, confidence interval; CME, cytoplasmic mesothelin expression; ER, estrogen receptor; MME, membrane mesothelin expression; HR, hazard ratio; Ly, lymphatic invasion; NG, nuclear grade; PgR, progesterone receptor; V, vascular invasion.

improved in patients with pancreatic cancer and other solid cancers using amatuximab, a chimeric monoclonal antibody that targets mesothelin (22,23). TNBC is considered a good target of immunotherapy, and Tchou *et al* (14) reported that genetically modified T-cells expressing a chimeric antibody receptor (CAR) specific for mesothelin (mesoCAR T-cells) had high anti-tumor cytotoxicity. In the present study, we showed that the target for molecular target therapy could be expanded to the luminal type.

We demonstrated that MME-positivity was significantly associated with RFS and that CME-positive patients had relatively worse RFS. Both MME and CME are significantly related to OS, and previous studies have shown that MME-positivity was a poorer prognostic factor than CME in various cancer

types (4,8-10,21). Furthermore, Kawamata *et al* (21) presumed that mesothelin in the cytoplasm was in the 71 kDa precursor form and might behave in a dominant-negative manner as a tumor suppressor in extrahepatic bile duct cancer. In this study, there were only four cases of cytoplasmic-only ME, and additional studies in a greater number of cases are required to prove this hypothesis in breast cancer.

In conclusion, we suggest that MME-positivity in breast cancer, especially in luminal type, is associated with poorer clinical outcomes. The clinical utility of the immunohistochemical examination of ME in surgically resected tumor specimens is expected to be useful for prognostication and decision making with regards to further treatment procedures after surgical therapy in breast cancer patients.

Table IV. Cox's univariate and multivariate analysis including clinicopathological parameters and mesothelin expression of recurrence-free survival in 343 patients with luminal type breast cancer.

Parameter	Univariate		Multivariate	
	HR (95% CI)	P-value	HR (95% CI)	P-value
pT				
Tis, T1	1	<0.001	1	0.011
T2, T3	4.82 (2.45-10.35)		2.58 (1.24-5.79)	
pN				
Negative	1	<0.001	1	<0.001
Positive	6.58 (3.26-14.71)		4.74 (2.27-10.91)	
NG				
1, 2	1	0.003	1	0.185
3	2.62 (1.41-4.86)		1.55 (0.81-2.98)	
Ly				
Negative	1	0.024	1	0.923
Positive	2.18 (1.10-4.71)		1.04 (0.51-2.27)	
V				
Negative	1	0.005	1	0.113
Positive	2.45 (1.32-4.58)		1.68 (0.88-3.21)	
Ki-67 labeling index, %				
<14	1	0.113		
≥14	1.72 (0.87-3.25)			
MME				
Negative	1	0.062		
Positive	2.35 (0.95-5.02)			
CME				
Negative	1	0.067		
Positive	2.31 (0.94-4.93)			

CI, confidence interval; CME, cytoplasmic mesothelin expression; HER2, human epidermal growth factor receptor type 2; HR, hazard ratio; Ly, lymphatic invasion; NG, nuclear grade; MME, membrane mesothelin expression; V, vascular invasion.

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### Availability of data and materials

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

### Authors' contributions

TK, TY, MFK, YI, YT and TSh collected the data and assisted in data analysis. YM, TI, ES and KS contributed

to interpretation of data and manuscript preparation. TSu analyzed the data and wrote the original draft. YY, TE and HT conceived the study and reviewed and revised the manuscript. JY and YK made substantial contributions to the conception and design. HU made substantial contributions to analysis and interpretation of data. JY, YK and HU provided supervision of the manuscript. All authors read and approved the final manuscript.

### Ethics approval and consent to participate

The present study was performed in accordance with the Declaration of Helsinki, and was approved by the Institutional Review Board of the National Defense Medical College (registration no. 3003). All patients agreed to participate in the present study, and written informed consent was obtained from all patients.

### Patient consent for publication

Not applicable.

## Competing interests

The authors declare that they have no competing interests.

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