

# Distinct subcellular localization of E-cadherin between epithelioid angiomyolipoma and triphasic angiomyolipoma: A preliminary case-control study

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**Abstract.** Epithelioid angiomyolipoma (EAML) is a rare variant of angiomyolipoma (AML). Previous studies have demonstrated that epithelial (E-)cadherin is expressed in two subtypes of AML, EAML and triphasic AML; however, the expression pattern of E-cadherin remains unclear. In the present study, a preliminary case-control study was conducted to determine the expression pattern of E-cadherin between EAML and triphasic AML, the control, focusing on the subcellular localization and expression category of E-cadherin. No significant difference was identified in the age, sex, history of tuberous sclerosis, smoking and alcohol consumption between the two groups ( $P > 0.05$ ). In EAML, 9 patients were categorized as exhibiting a low risk of malignant behavior and the other two were categorized as exhibiting an intermediate or high risk of malignant behavior. The proportion of cases expressing E-cadherin, human melanoma black-45 (HMB45), melanoma antigen recognized by T cells 1 (Mart1/Melan A), smooth muscle actin and progesterone receptor were 95.5 (21/22),

95.5 (21/22), 86.4 (19/22), 77.3 (17/22) and 86.4% (19/22), respectively. E-cadherin was identified to be localized, using staining techniques, in the cell membrane and/or cytoplasm. The subcellular localization of E-cadherin was significantly different between EAML and triphasic AML; the majority of EAML cases revealed membranous and cytoplasmic staining, whereas triphasic AML cases demonstrated cytoplasmic staining ( $P = 0.0093$ ). The expression of E-cadherin may be positively associated with HMB45 ( $P = 0.0044$ ) and Mart1/Melan A ( $P = 0.0049$ ). The results of the present study identified that the subcellular localization of E-cadherin may be different between EAML and the control group of triphasic AML. Additionally, E-cadherin and melanocytic markers may be co-expressed in distinct subtypes of AML. A follow-up study with a large sample size to validate the results of the present study, followed by a mechanistic study based on cell lines to determine any significance, are warranted.

## Introduction

Angiomyolipoma (AML) is the most common type of kidney mesenchymal tumor and accounts for between 1 and 3% of kidney tumors (1,2). There are two clinical subtypes of AML: Sporadic AML and tuberous sclerosis complex (TSC)-associated AML (in patients with TSC) (3). The allelic loss of the TSC2 gene is a common genetic event in AML (4,5). There are three histological subtypes of AML: Triphasic, monophasic and epithelioid (3). Triphasic AML, the most common subtype of AML, is a benign tumor containing various proportions of mature fat, thick-walled and poorly organized blood vessels and smooth muscle (3,5). Monophasic and epithelioid AML (EAML) are classified as atypical AML (3). EAML is a rare subtype of AML which was first described by Mai *et al* (6) and is characterized by the proliferation of atypical epithelioid cells, with abundant eosinophilic and granular cytoplasm (5). Triphasic and EAML co-express melanocytic markers [human melanoma black-45 (HMB45), melanoma antigen recognized by T cells 1 (Mart1/Melan A), etc.] with variable intensity in smooth muscle markers, including smooth muscle actin

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**Abbreviations:** AML, angiomyolipoma; CK, cytokeratin; EAML, epithelioid angiomyolipoma; EMA, epithelial membrane antigen; EMT, epithelial-to-mesenchymal transition; ECs, epithelioid cells; IHC, immunohistochemistry; IRS, immunoreactive score; MET, mesenchymal-to-epithelial transition; OR, odds ratio; PR, progesterone receptor; SD, standard deviation; SMA, smooth muscle actin; TSC, tuberous sclerosis complex

**Key words:** kidney, angiomyolipoma, epithelioid, epithelial cadherin, immunohistochemistry, case-control

(SMA), and progesterone receptor (PR) (5). In contrast with triphasic AML, ~33% of EAML cases exhibit malignant potential with metastasis to the lymph nodes, liver, lungs or spine (5). However, there are no definite histopathological features to determine the prognosis of EAML. A number of genitourinary pathologists previously agreed that EAMLs should be divided into low, intermediate and high risk of malignant behavior (7-9). Currently, the majority of studies on malignant EAML have focused on the p53 gene and the protein Ki-67 (10-15).

One of the characteristic differences between EAML and triphasic AML is in the cellular morphology. The former is characterized by the appearance of epithelioid cells, which may exhibit the potency of metastasis, whereas the latter presents with spindle cells, which do not get the potency of metastasis (5). This difference may be similar to the process of epithelial-to-mesenchymal transition (EMT) and mesenchymal-to-epithelial transition (MET) to a certain extent (16,17). Epithelial (E-)cadherin was reported as one of the key factors in EMT (16-18). A previous study conducted on renal epithelial tumor cell lines demonstrated that loss-of-function mutations of the TSC2 gene resulted in cytoplasmic localization of E-cadherin, which led to a decrease in cell-cell adhesion and the development of EMT (19). This may indicate that E-cadherin is an important factor in the progression of kidney tumors. In addition, the expression of E-cadherin is reported as a common event in AML, and, generally, the membrane immunoreactivity is more marked in epithelioid tumor cells compared with that in spindle tumor cells (20,21). These results raise the question of whether EAML expresses E-cadherin with distinct subcellular localization, compared with triphasic AML.

In the present study, a preliminary case-control study was conducted to detect the expression of E-cadherin in EAML compared with triphasic AML (the most common AML subtype), as a control. The results of the present study demonstrated that the subcellular localization of E-cadherin in EAML was distinct from that in triphasic AML. It was additionally revealed that E-cadherin and melanocytic markers may be co-expressed in distinct subtypes of AML.

## Materials and methods

**Patients and tissues.** The present study was part of a series of retrospective studies which was approved by the National Cancer Center Ethics Committee/Institutional Review Board (IRB; Chinese Academy of Medical Sciences and Peking Union Medical College, Beijing, China). All identifying information of patients was anonymized, and there were no risks anticipated to the participants. It was difficult to acquire patient consent from each participant in a retrospective study and, with the permission of the National Cancer Center Ethics Committee/IRB, patient consent was waived in this series of studies. There were 9 males and 13 females recruited into the present study. The median age was 42.8, and ranged from 23-58 years old.

In the present study, 11 cases of EAML from the National Cancer Center/Cancer Hospital, Chinese Academy of Medical Sciences and Peking Union Medical College, between January 2005 and December 2011. Another 11 cases of triphasic

AML, during the same period, were selected as the control. The World Health Organization classification of renal tumors (2004) criteria were applied to all EAML and classic AML cases examined in the present study (5). EAML followed these diagnostic criteria, as follows: i) Polygonal larger cells with abundant granular cytoplasm; ii) enlarged vesicular nuclei, prominent nucleoli, multinucleated or enlarged ganglion-like cells; iii) nuclear atypia; iv) expression of one or both melanocytic markers (HMB-45, Mart1/Melan A); v) positive or negative for SMA or PR; and vi) negative immunoreaction for the epithelial markers pan-cytokeratin (CK), CK8, CK18 and epithelial membrane antigen (EMA). The cases of triphasic AML followed these diagnostic criteria, as follows: i) A variable mixture of fat, blood vessels and smooth muscle; ii) the smooth cells emanated from blood vessel walls in a radial fashion with or without atypia; iii) mature adipose tissue and thick-walled poorly organized blood vessels were observed in the tumor; and iv) immunoprofile was similar to that of EAML (as aforementioned).

In the EAML group, the epithelioid component was required to be  $\geq 5\%$  in the tumors, which used the same criteria as a previous study (7). The nuclear atypia was also required to be at least moderate in EAML. The nuclear atypia criteria followed the criteria of Brimo *et al* (7). The nuclear atypia was defined as vesicular nuclei, prominent nucleoli and nuclear size, which was at least twice the size of adjacent nuclei. Moderate atypia described epithelioid cells that were intermediate in size and exhibited enlarged nuclei with moderate pleomorphism and prominent nucleoli. Identifying carcinoma-like growth patterns was conducted following the definition of Nese *et al* (8): Tumor cells arranged as cohesive nests, broad alveoli and compartmentalized sheets. All EAMLs were divided into three groups of malignant behaviors (low, intermediate and high risk) on the basis of five adverse prognostic parameters: Associated TSC of concurrent AML, necrosis, tumor size  $>7$  cm, extrarenal extension and/or renal vein involvement and a carcinoma-like growth pattern (8,9). Tumors with  $<2$  adverse prognostic parameters were classified as low risk, tumors with between 2 and 3 adverse prognostic parameters were classified as intermediate risk and tumors with  $\geq 4$  adverse prognostic parameters were classified as high risk of exhibiting malignant behavior.

Three pathologists (Y.Z., H.T.Z. and S.Z.) from the Department of Pathology, National Cancer Center/Cancer Hospital, Chinese Academy of Medical Sciences and Peking Union Medical College (Beijing, China) reviewed all the hematoxylin and eosin slices of every case to evaluate the proportion of epithelioid cells (ECs), nuclear atypia and growth pattern. The demographic and clinical information including age, sex, history of TSC, smoking and alcohol consumption was obtained from medical records. Tumor size was obtained according to the gross description of the specimen following formalin fixation.

**Immunohistochemistry (IHC).** IHC staining was performed in all 22 cases using 9 antibodies (Table I) according to the manufacturer's protocol for each antibody. Tissue blocks with tumor and normal tissue were selected for the IHC staining. Prior to IHC, sections of 4  $\mu\text{m}$  from each block were deparaffinized by xylene, rehydrated by gradient alcohol and then steamed

Table I. Antibody information.

Antibody	Clone	Catalogue numbers	Dilution	Incubation time, min	Source
CK8	TS1	Kit-0034	Working solution	30	Fuzhou Maixin Biotech Co., Ltd., Fuzhou, China
CK18	DC10	MAB-0182	Working solution	20	Fuzhou Maixin Biotech Co., Ltd.
E-cadherin	4A2C7	MAB-0589	Working solution	30	Fuzhou Maixin Biotech Co., Ltd.
EMA	E29	IR62961	Working solution	30	Dako; Agilent Technologies, Inc., Santa Clara, CA, USA
Pan-keratin	AE1/AE3	Z2061	1:120	30	Zeta Corporation, Arcadia, CA, USA
Mart1/Melan A	A103	IR63361	Working solution	20	Dako; Agilent Technologies, Inc.
Melanoma	HMB45	MAB-0098	Working solution	30	Fuzhou Maixin Biotech Co., Ltd.
PR	1E2	790-4296	Working solution	30	Ventana Medical Systems, Inc., Tucson, AZ, USA
SMA	1A4	202M-97	Working solution	30	Cell Marque, Austin, TX, USA

CK, cytokeratin; E-cadherin, epithelial cadherin; EMA, epithelial membrane antigen; Mart1/Melan A, melanoma antigen recognized by T cells 1; PR, progesterone receptor; SMA, smooth muscle actin.

in 10 mM sodium citrate buffer, pH 6.0 (except SMA), for 2 min in a pressure cooker. SMA did not require any antigen retrieval according to the manufacturer's protocol. Endogenous peroxidase was blocked by incubating the sections at room temperature with ultraView Universal diaminobenzidine (DAB) inhibitor (3% H<sub>2</sub>O<sub>2</sub>; Ventana Medical Systems, Inc., Tucson, AZ, USA) for 5 min. Sections were subsequently incubated with primary antibodies for between 20 and 30 min according to the antibody (Table I). Following this, sections were incubated in the ultraView Universal horseradish peroxidase Multimer for 15 min and then visualized using ultraView Universal DAB Chromogen (0.2% DAB), ultraView Universal DAB H<sub>2</sub>O<sub>2</sub> and ultraView Universal DAB Copper (all Ventana Medical Systems, Inc.).

The results of staining were initially assigned an immunoreactive score (IRS) (22,23). Values for the proportion of positive cells were determined as follows: Negative, 0; 1-10%, 1; 11-50%, 2; 51-80%, 3; and 81-100%, 4. Staining intensity was qualitatively evaluated as follows: Negative, 0; weak, 1; moderate, 2; and strong, 3. The final IRS was calculated by multiplying the value for the proportion of positive cells by the value for staining intensity. The total range of values was 0-12. Subsequently, the expression of each biomarker was divided into four categories based on the IRS: IRS 0, negative (-); IRS 1-4, weak staining (+); IRS 5-8, intermediate staining (++); IRS 9-12, strong staining (+++). The slices were evaluated independently by three pathologists (X.L.W., Q.W. and S.Z.). In ambiguous cases, these three pathologists evaluated together using a Leica Multiview system (Leica Microsystems, Inc., Buffalo Grove, IL, USA) to obtain the final result.

*Statistical analysis.* The mean and standard deviation (SD) were calculated for continuous data and a Student's t-test was carried out. Frequency and proportion values were reported for the categorical data, including sex, surgical type, history of TSC, smoking and alcohol history. Fisher's exact test was conducted. Spearman's rank correlation was used to determine the association between E-cadherin and other markers of interest, including HMB45, Mart1/Melan A, SMA and PR. Odds ratios (ORs) and 95% confidence intervals for risk were calculated. To estimate the adjusted OR, the logistic regression models including age, sex and tumor size as factors were used. All the statistical tests carried out were two-tailed and P<0.05 was considered to indicate a statistically significant difference. No adjustment for multiplicity was applied since all analyses were to be exploratory. In EAML, only the staining result of epithelial components was compared with that in the control (triphasic AML) components. The data were analyzed using SAS (version 9.2; SAS Institute Inc., Cary, NC, USA).

## Results

*Baseline characteristics between EAML and control group of triphasic AML.* There were a total of 22 cases in the EAML group and the control (triphasic AML) group, some of which were reported previously (24) (Table II). Only 1 case with EAML exhibited metastases (24). In total, there were 13 females and 9 males with a mean  $\pm$  SD age of 42.8 $\pm$ 9.8 years. Tumor size ranged between 2.0 and 17.5 cm and the majority of cases underwent radical nephrectomy. In the EAML group, 9 cases exhibited  $\geq$ 90% ECs in tumors; the other two cases exhibited 10% ECs in tumors. None of the patients had a

Table II. Detailed baseline characteristics of EAML and triphasic AML.

Case no.	Pathological diagnosis	Age/sex	Tumor size, cm	ECs, %	Carcinoma-like growth pattern	Necrosis	Extrarenal extension and/or renal vein involvement	Surgical type	TSC	Smoking history	Alcohol consumption
1	EAML	40/F	5.5	100	-	-	-	Partial nephrectomy	-	Never	Never
2	EAML	41/F	6.0	90	-	+	-	Radical nephrectomy	-	Never	Never
3	EAML	49/M	4.0	10	-	+	-	Radical nephrectomy	-	Current	Current
4	EAML	23/M	3.5	100	-	-	+	Radical nephrectomy	-	Current	Never
5	EAML	41/M	2.0	10	-	-	-	Partial nephrectomy	-	Current	Never
6	EAML	46/M	8.5	100	-	-	-	Radical nephrectomy	-	Current	Former
7	EAML	25/F	12.5	100	+	+	+	Radical nephrectomy	-	Never	Never
8	EAML	51/F	4.5	100	-	+	-	Radical nephrectomy	-	Former	Former
9	EAML	56/F	6.0	100	+	-	-	Partial nephrectomy	-	Never	Never
10	EAML	58/F	17.5	95	+	-	-	Radical nephrectomy	-	Never	Never
11	EAML	47/M	2.0	100	+	-	-	Radical nephrectomy	-	Never	Never
12	Triphasic AML	57/M	5.0	-	-	-	-	Partial nephrectomy	-	Current	Former
13	Triphasic AML	47/F	2.5	-	-	-	-	Partial nephrectomy	-	Never	Never
14	Triphasic AML	33/F	13.0	-	-	-	-	Radical nephrectomy	-	Never	Never
15	Triphasic AML	55/M	5.5	-	-	-	-	Partial nephrectomy	-	Current	Current
16	Triphasic AML	39/F	15.0	-	-	-	-	Radical nephrectomy	-	Never	Never
17	Triphasic AML	31/F	4.0	-	-	-	-	Partial nephrectomy	-	Never	Never
18	Triphasic AML	35/F	15.0	-	-	+	-	Radical nephrectomy	-	Never	Never
19	Triphasic AML	41/F	2.0	-	-	-	-	Partial nephrectomy	-	Never	Never
20	Triphasic AML	49/M	3.0	-	-	-	-	Radical nephrectomy	-	Current	Former
21	Triphasic AML	38/M	6.5	-	-	+	-	Radical nephrectomy	-	Current	Never
22	Triphasic AML	40/F	13.0	-	-	-	-	Radical nephrectomy	-	Never	Never

AML, angiomylipoma; EAML, epithelioid angiomylipoma; ECs, epithelioid cells; F, female; M, male; TSC, tuberosus sclerosis complex; +, positive; -, negative.

Table III. Comparison of baseline characteristics between EAML and triphasic AML.

Characteristic	EAML	Triphasic AML	Total
Age, years			
n	11	11	22
Mean (SD)	43.4 (11.2)	42.3 (8.7)	42.8 (9.8)
Range	23-58	31-57	23-58
Sex			
Female, n (%)	6 (54.5)	7 (63.6)	13 (59.1)
Male, n (%)	5 (45.5)	4 (36.4)	9 (40.9)
Tumor size, cm			
Mean (SD)	6.5 (4.7)	7.7 (5.2)	7.0 (4.9)
Range	2.0-17.5	2.0-15.0	2.0-17.5
Surgical type			
Radical nephrectomy, n (%)	8 (72.7)	6 (54.5)	14 (63.6)
Partial nephrectomy, n (%)	3 (27.3)	5 (45.5)	8 (36.4)
History of TSC			
Yes, n (%)	0 (0)	0 (0)	0 (0)
No, n (%)	11 (100)	11 (100)	22 (100)
Smoking history			
Never, n (%)	6 (54.5)	7 (63.6)	13 (59.1)
Former or current, n (%)	5 (45.5)	4 (36.4)	9 (40.9)
Alcohol consumption			
Never, n (%)	8 (72.7)	8 (72.7)	16 (72.7)
Former or current, n (%)	3 (27.3)	3 (27.3)	6 (27.3)

AML, angiomyolipoma; EAML, epithelioid angiomyolipoma; SD, standard deviation; TSC, tumor sclerosis complex.

history of TSC; however, 15 of the patients had a history of smoking and/or alcohol consumption. The important baseline characteristics between EAML and control groups were comparable ( $P>0.05$ ; Table III). In the EAML group, 9 cases were classified as possessing a low risk of malignant behavior, 1 case was classified as possessing an intermediate risk of malignant behavior and 1 case was classified as possessing a high risk of malignant behavior (Table IV).

#### Expression of E-cadherin and other biomarkers in all AML.

First, the expression of 9 biomarkers in the two AML groups were identified (Figs. 1 and 2). In all 22 cases, the proportion of cases expressing E-cadherin, HMB45, Mart1/Melan A, SMA and PR was 95.5 (21/22), 95.5 (21/22), 86.4 (19/22), 77.3 (17/22) and 86.4% (19/22), respectively. All biomarkers, with the exception of PR, exhibited positive subcellular localization in the cell membrane and/or plasma. The latter exhibited nuclear staining. The majority of EAML cases (8/11, 72.7%) demonstrated intermediate E-cadherin expression, whereas that of the control cases (8/11, 72.7%) demonstrated weak E-cadherin expression. In all cases, results demonstrated that E-cadherin and HMB45 were expressed; however, E-cadherin and Mart1/Melan A were expressed in 86.4% (19/22) of cases (Table IV). AE1/AE3, CK18, CK8 and EMA were not expressed in any of the cases (Table IV).

The association between E-cadherin and other biomarkers was subsequently identified. The results from the present

study revealed that the expression of E-cadherin demonstrated a positive association with the expression of HMB45 ( $P=0.0044$ ; Table V) and Mart1/Melan A ( $P=0.0049$ ; Table V). Additionally, the results of the present study revealed negative associations between E-cadherin and SMA or PR. However, none of these results were considered to indicate a statistical significance.

#### Distinct subcellular localization of E-cadherin in EAML and control group of triphasic AML.

The subcellular localization and staining categories of E-cadherin were investigated between EAML and the control group of triphasic AML to observe any differences. The majority of cases in the EAML group demonstrated cell membrane and cell plasma staining of E-cadherin (8/11), whereas the majority of cases in the triphasic AML group demonstrated cell plasma staining only (9/11). The subcellular localization of E-cadherin was identified to demonstrate differences between the EAML group and the control group ( $P=0.0089$ ). The OR was calculated to be 20.0, indicating that cases with cell membrane E-cadherin staining (with or without cell plasma staining) had a 20-fold increased likelihood of exhibiting EAML rather than triphasic AML (Table VI). However, no statistically significant difference in staining categories of E-cadherin between these two groups was identified ( $P=0.0950$ ) following the adjustment for age, sex and tumor size (data not shown).

Table IV. Pathological characters and expression of biomarkers in EAML and triphasic AML.

Case no.	Pathological diagnosis	Risk of malignant behavior	E-cadherin									
			Expression category	Subcellular localization	HMB45	Mart1/ Melan A	SMA	PR	Pan-cytokeratin	CK18	CK8	EMA
1	EAML	Low	++	M+P	+	+	++	-	-	-	-	-
2	EAML	Low	+++	M+P	+	+	++	+	-	-	-	-
3	EAML	Low	++	M+P	+	+	++	+	-	-	-	-
4	EAML	Low	++	M+P	+	+	+	+	-	-	-	-
5	EAML	Low	+++	M+P	+	+	++	++	-	-	-	-
6	EAML	Low	++	M+P	+	+	-	+	-	-	-	-
7	EAML	High	++	M+P	+	+	+	+++	-	-	-	-
8	EAML	Low	+	P	+	+	+++	++	-	-	-	-
9	EAML	Low	++	M+P	+	+	-	+	-	-	-	-
10	EAML	Intermediate	-	-	-	+	-	+	-	-	-	-
11	EAML	Low	+	P	+	+	+	+	-	-	-	-
12	Triphasic AML	-	+	P	+	+	+++	++	-	-	-	-
13	Triphasic AML	-	+	M+P	+	+	++	++	-	-	-	-
14	Triphasic AML	-	++	P	+	+	-	+	-	-	-	-
15	Triphasic AML	-	+	P	+	+	++	+	-	-	-	-
16	Triphasic AML	-	+	P	+	+	++	+	-	-	-	-
17	Triphasic AML	-	+	P	+	+	+	++	-	-	-	-
18	Triphasic AML	-	++	P	+	+	-	+	-	-	-	-
19	Triphasic AML	-	++	M	+	+	+	+	-	-	-	-
20	Triphasic AML	-	+	P	+	+	+++	-	-	-	-	-
21	Triphasic AML	-	+	P	+	+	+	-	-	-	-	-
22	Triphasic AML	-	+	P	+	+	+++	+	-	-	-	-

AML, angiomylipoma; CK, cytokeratin; EAML, epithelioid angiomylipoma; EMA, epithelial membrane antigen; HMB45, Human Melanoma Black-45; M, membrane; Mart1/Melan A, melanoma antigen recognized by T cells 1; P, plasma; +, weak positive; ++, intermediate positive; +++, strong positive; SMA, smooth muscle actin; -, negative.

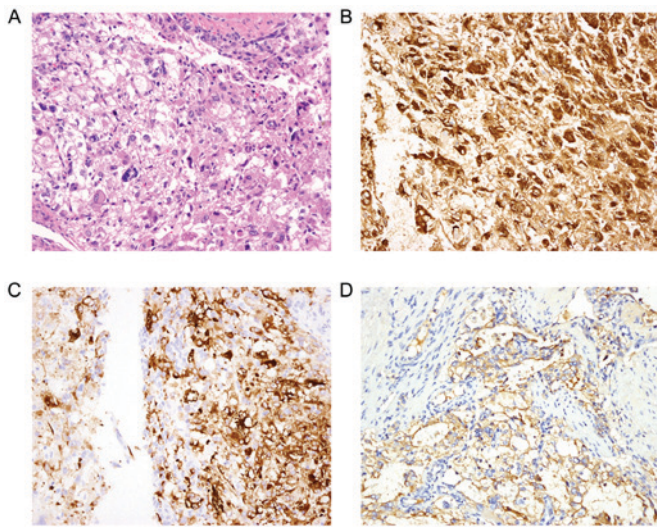


Figure 1. Hematoxylin and eosin staining demonstrating the histopathological features and the expression of biomarkers (magnification, x200) in EAML. (A) Histopathological features of EAML. Polygonal larger cells with abundant granular cytoplasm and nuclear atypia, prominent nucleoli and multinucleated cells. (B) Human Melanoma Black-45 staining for EAML (strong plasma staining). (C) Melanoma antigen recognized by T cells 1 for EAML (strong plasma staining). (D) Epithelial cadherin staining for EAML (middle membrane combined with plasma staining). EAML, epithelioid angiomyolipoma.

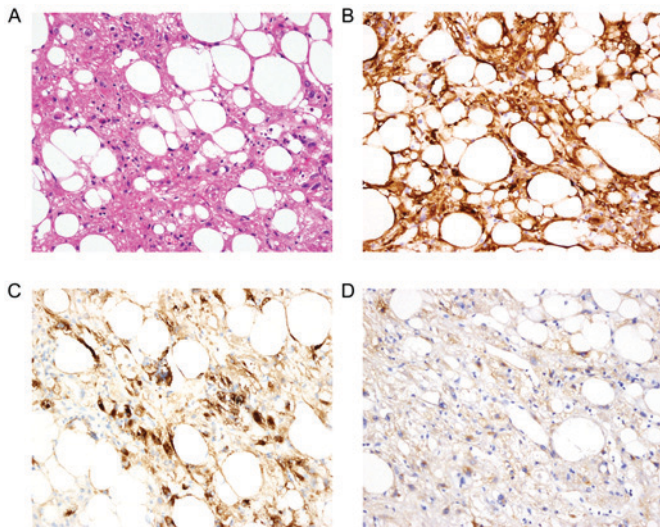


Figure 2. Hematoxylin and eosin staining demonstrating the histopathological features and the expression of biomarkers in AML (magnification, x200). (A) Histopathological features of AML, fat and smooth muscle cells. (B) Human Melanoma Black-45 staining for AML (strong plasma staining). (C) Melanoma antigen recognized by T cells 1 staining for AML (moderate plasma staining). (D) Epithelial cadherin staining for AML (weak plasma staining). AML, angiomyolipoma.

The majority of cases classified as possessing a low risk of malignant behavior exhibited at least intermediate E-cadherin staining (7/9, 77.8%), whereas the case classified as possessing an intermediate risk of exhibiting malignant behavior exhibited negative E-cadherin staining. The case classified as exhibiting a high risk of malignant behavior group also exhibited intermediate E-cadherin staining; however, due to the limited number of cases, the significance of this remains unclear.

Table V. Spearman's rank correlation between E-cadherin and other biomarkers.

Variable	Biomarker	n	Sample correlation	P-value
E-cadherin	HMB45	21	0.5956	0.0044
	Mart1/Melan A	22	0.5777	0.0049
	SMA	22	-0.0904	0.6893
	PR	22	-0.1879	0.4023

E-cadherin, epithelial cadherin; HMB45, human melanoma black-45; Mart1/Melan A, melanoma antigen recognized by T cells 1; SMA, smooth muscle actin; PR, progesterone receptor.

## Discussion

E-cadherin belongs to the classical cadherin superfamily; it has five repetitive extracellular cadherin domains and a conserved cytoplasmic domain (25,26). E-cadherin was first reported as a cell adhesion protein in chicken liver cells and mice (27,28), and Takeichi and colleagues named this  $Ca^{2+}$ -dependent cell-cell adhesion molecule E-cadherin in the early 1980s (29,30). Since then, the role E-cadherin serves in cell architecture and tissue formation of normal epithelium has been extensively investigated (25,26,31,32). It has been identified previously that E-cadherin is a key factor in EMT and MET (33). EAML exhibits epithelioid cells with metastatic potential, while triphasic AML presents with spindle cells without metastatic potential (5). These differences between EAML and triphasic AML were similar to the EMT process to a certain extent (16,17). Hence, the factors associated with the EMT process, including E-cadherin (33), may be a reason for these differences. Previous studies have revealed E-cadherin to be expressed in tumors of the urinary system (34,35), soft tissue sarcoma (36,37) and AML (20,21). E-cadherin may be associated with the differences in cellular morphology between EAML and triphasic AML. The present preliminary case-control study was conducted to detect the expression pattern of E-cadherin in EAML compared with the common subtype of AML, triphasic AML.

The results of the present study indicated that E-cadherin was expressed commonly in EAML and triphasic AML. All but one case expressed E-cadherin and half of the cases demonstrated at least intermediately stained E-cadherin. The proportion of cases demonstrating E-cadherin expression in EAML was slightly decreased compared with that in triphasic AML. The results of the present study were similar to those from a previous study by Wang *et al* (20) which revealed a slight increase in the proportion of cases demonstrating E-cadherin expression (98%), and the majority of cases (71%) demonstrated moderate or strong E-cadherin staining. The slight difference between the two studies may be due to the differences in case selection. In the present study, renal EAML and control (triphasic AML) cases were selected at the constituent ratio 1:1. Conversely, the study by Wang *et al* (20) collected a variety of AML types (including renal, hepatic and retroperitoneal), with only 2 cases of EAML (5%). The

Table VI. Fisher's exact test, OR for E-cadherin score and localization between EAML and triphasic AML.

E-cadherin expression	EAML, n (%)	Triphasic AML, n (%)	P-value (Fisher's exact test)	OR (95% CI), P-value
E-cadherin localization				
Membrane staining with or without plasma	8 (80.0)	2 (18.2)	0.0089	20.0 (2.0, 159.1), 0.0093
Cytoplasm only	2 (20.0)	9 (81.8)		
E-cadherin expression category				
++/+++	8 (72.7)	4 (36.4)	0.1984	4.7 (0.77, 28.5), 0.0950
-/+	3 (27.3)	7 (63.6)		

E-cadherin, epithelial cadherin; AML, angiomyolipoma; CI, confidence intervals; EAML, epithelioid angiomyolipoma; OR, odds ratio; +, positive; -, negative.

differences in the proportion of epithelioid cell and antibody clone of E-cadherin may be another reason for the differences between the results of the two studies. Additionally, the present study investigated the association between E-cadherin and other biomarkers which have been identified in AML (5,38). The results of the present study demonstrated that the expression of E-cadherin may be positively associated with the expression of HMB45 and Mart1/Melan A which are common biomarkers of melanomas ( $P < 0.05$ ). Tumors which expressed E-cadherin exhibited a decreased level of SMA and PR; however, no statistically significant difference was identified ( $P > 0.05$ ). In a previous study, Barnes *et al* (19) revealed that loss of E-cadherin expression may upregulate the expression of SMA in TSC2 gene (-/-) cell lines and is the only study, to the best of our knowledge, which has focused on the association between E-cadherin and other biomarkers. These cell lines were confirmed to exhibit a loss of the wild-type allele for the Tsc2 locus. In the present study, the association between E-cadherin and other biomarkers was observed at the tissue level; however, the underlying molecular mechanism and significance remains unknown. To elucidate this, a mechanistic study based on cell lines is required.

Secondly, the subcellular localization and staining category of E-cadherin in EAML and its control (triphasic AML) were investigated in the present study. The subcellular localization of E-cadherin was demonstrated to be significantly different between EAML and triphasic AML. Staining of the former demonstrated membranous (with or without cytoplasmic) E-cadherin, whereas triphasic AML revealed cytoplasmic E-cadherin staining only. In previous studies, E-cadherin exhibited a different staining intensity, a distinct localization between spindle and epithelioid cells in AML and staining identified cell membrane E-cadherin (20,21), which is consistent with the results of the present study. In addition, the results of the study by Barnes *et al* (19) were consistent with those of the present study at the cell line level. The study by Barnes *et al* (19) demonstrated that wild-type rat renal epithelial cells exhibited plasma membrane staining of E-cadherin which aided cells to maintain the epithelioid characteristics. In contrast, the TSC2 (-/-) renal epithelial tumor-derived cells demonstrated a paucity of membrane E-cadherin and cells lost

epithelioid morphology. In a previous study, Yang *et al* (39) hypothesized that increased expression of E-cadherin may be a critical step in MET. The results of the present study indicated that the various subcellular localizations of E-cadherin observed between EAML and triphasic AML may be one reason for the differences in cellular morphology between these two types of tumor.

Finally, the results of the present study identified differences between EAML cases possessing a low risk of exhibiting malignant behavior and EAML cases possessing intermediate and high risk of exhibiting malignant behavior. The cases categorized as low risk exhibited at least intermediate staining of E-cadherin, whereas the cases categorized as intermediate and high risk exhibited the opposite. Despite this, due to the limited number of cases, the significance of the association between expression of E-cadherin and malignant behavior remains unclear. Further studies are required to investigate this potential association.

The present case-control study was conducted to obtain more EAML cases retrospectively in a relatively short period. On the basis of current knowledge, it is understood that there are a number of risk factors for EAML; however, the only accepted risk factor was TSC (4,5). All the EAML cases had no history of TSC so only cases of triphasic AML without TSC were selected as a control. Additionally, the present study selected cases where patients also possessed common risk factors for cancer including a history of smoking and alcohol consumption. Although the cases with a strict condition were matched to guarantee the comparability between EAML and triphasic AML, there may be some confounding factors not identified in the present study which cannot be resolved using a case-control study. The small sample size and the heterogeneity in the proportion of ECs in EAML group were other limitations of the present study. However, EAML was a rare variant of AML and the present study established relative strict criteria of EAML and triphasic AML, on the basis of World Health Organization classification and the literature. The present study aimed for more suitable cases of EAML and triphasic AML to be recruited which resulted in all but 2 cases of EAML demonstrating  $\geq 90\%$  of ECs, which led to minimal heterogeneity. A study with a larger sample size from multiple centers and a relative homogeneity of the proportion



of epithelioid cells in EAML, is required to increase the reliability of the results.

E-cadherin may an important biomarker in AML. The mechanisms of E-cadherin may be distinct between EAML and triphasic AML, since the subcellular localization of E-cadherin may be distinct between EAML and triphasic AML. A study with a large sample size is required to validate the results of the present study, followed by a mechanistic study based on cell lines to determine any significance.

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