

CASE REPORT

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Malignant Perivascular epithelioid cell tumour of the uterus without *TFE3* gene rearrangement: a case report

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Abstract

Background Perivascular epithelioid cell tumours (PEComas) are soft tissue tumours. These neoplasms belong to the family of mesenchymal tumours, which include angiomyolipomas, clear-cell sugar tumours of the lung, and PEComas not otherwise specified (NOS). The probability of a perivascular epithelioid cell tumour (PEComa) occurring in the uterus is low, and the incidence, diagnosis, treatment, and outcomes of such tumours are still unclear.

Case presentation A 51-year-old woman presented a 4-year history of natural menopause. An intrauterine mass was detected via ultrasound examination; the mass showed a tendency to increase but caused no symptoms. The levels of tumour markers were within the normal range. Pathological analysis of the curettage revealed perivascular epithelioid differentiation of the endometrial tumour. Consequently, a laparoscopic total hysterectomy with bilateral adnexectomy was performed. No distant metastasis was detected via whole-body positron emission computed tomography (PETCT) after the operation. Fluorescence in situ hybridization (FISH) revealed no *TFE3* gene rearrangement. Next-generation sequencing of bone and soft tissue revealed negative TSC1/2 and TP53 expression. No recurrence or metastasis was observed during the 18-month follow-up period.

Conclusion PEComa of the gynecologic tract is a rare and challenging entity. Diffuse HMB-45 expression, TSC alterations and TFE3 rearrangement are characteristic of uterine PEComas. Surgical resection is the first choice. Genetic testing is helpful for determining the nature of the mass and for choosing targeted therapy. Further research is needed to establish treatment protocols.

Keywords PEComa, Uterus, *TFE3*, Rearrangement, Case report

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Background

Perivascular epithelioid cell tumours (PEComas) were first proposed by Zamboni et al. in 1996 [1]. In 2003, the World Health Organization defined a PEComa as a mesenchymal tumour with perivascular epithelioid cell characteristics in terms of the histology and immunophenotype. PEComas of the female gynecological tract are rare, accounting for 25% of all PEComas, and they cause variable symptoms and yield different prognoses for each individual patient. The uterus is one of the most commonly involved sites (72%). Fewer cases of gynecological PEComas have also been reported in the cervix (11%), and even rarer cases have been reported in the vagina, broad ligament and ovary [2, 3]. Surgery is the main treatment. The rates of metastatic disease at diagnosis, recurrences, and/or death vary among larger studies, with percentages ranging from 35 to 64% of patients [4–6]. The etiology of PEComas remain unclear and may be related to estrogen levels [7]. Some studies have suggested that gene mutations in the tuberous sclerosis complex (TSC) [8] and the rearrangement of transcription Factor E3 (*TFE3*) [9] are associated with pathogenesis. PEComas with *TFE3* gene rearrangements are a group of subtypes with a unique morphology. Compared with PEComas without rearrangement, these subtypes are more aggressive and have malignant morphological characteristics. Here, a case of a malignant uterine PEComa without *TFE3* gene rearrangement is reviewed.

Case presentation

A 51-year-old woman with a uterine mass found 6 months prior was admitted to the gynecology department of Fujian Maternity and Child Health Hospital. The patient had undergone natural menopause 4 years prior. Ultrasonography revealed a 2 cm mass in the patient's uterus 6 months prior, and a regular review was performed. The presurgical ultrasound examination showed an intrauterine mass that was 4.0 cm × 3.2 cm × 5.1 cm

in size (uterine size, 4.9 cm × 4.5 cm × 5.3 cm; endometrial thickness, 0.3 cm). Colour Doppler revealed that the blood flow signal in the tumour had a pulsatility index of 0.52 and a resistive index of 0.40. The preoperative diagnostic hypothesis was uterine submucosal fibroids. There was no family history or clinical evidence of tuberous sclerosis. Pathological analysis of the curettage revealed perivascular epithelioid differentiation of the endometrial tumour. The results of laboratory tests, including carcinoembryonic antigen (CEA), cancer antigen (Ca) 125, Ca153, Ca199, alpha fetoprotein (AFP) and squamous cell carcinoma antigen (SCC) levels, were all within the normal range. As showed in Fig. 1, magnetic resonance imaging (MRI) revealed an abnormal signal indicating uterine mass (3.7 cm × 3.2 cm × 4.3 cm); and the internal strengthening was uneven after enhancement. No enlarged pelvic lymph nodes were observed.

A laparoscopic total hysterectomy with bilateral adnexectomy was performed. During the intraoperative exploration, no abnormalities were found in the abdominal pelvic cavity, and no obviously enlarged pelvic lymph nodes were observed. The uterus was enlarged, measuring 8.0 cm × 6.0 cm × 5.0 cm, whereas both the ovaries and fallopian tubes were unremarkable. Upon bivalving the uterus, a greyish tumour measuring 4 cm × 4 cm × 2.5 cm was observed in the endometrial cavity. Microscopically, the tumour had infiltrated into the superficial 1/2 of the muscle layer, resulting in necrosis and lymphovascular invasion. The mitotic count was >1/50 HPF. As shown by the immunohistochemistry results in Fig. 2A–D, the tumour cells were negative for ER, PR, ALK, and SOX-10. However, they showed diffuse strong positivity for HMB-45 and Melan-A and focal positive (1+) staining for smooth muscle actin (SMA), TFE-3, CD34, and myogenin. The Ki-67 index was 8%. The peritoneal lavage fluid tested negative for tumour cells. A final diagnosis of a malignant uterine PEComa was made.

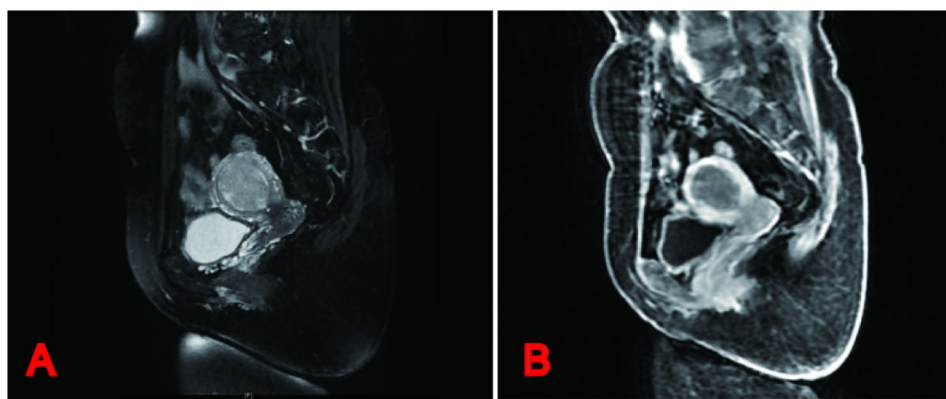


Fig. 1 Pelvis magnetic resonance imaging showing heterogeneously enhancing mass occupying the uterine cavity. (A) Sagittal T2-weighted image. (B) Sagittal contrast-enhanced T1-weighted image

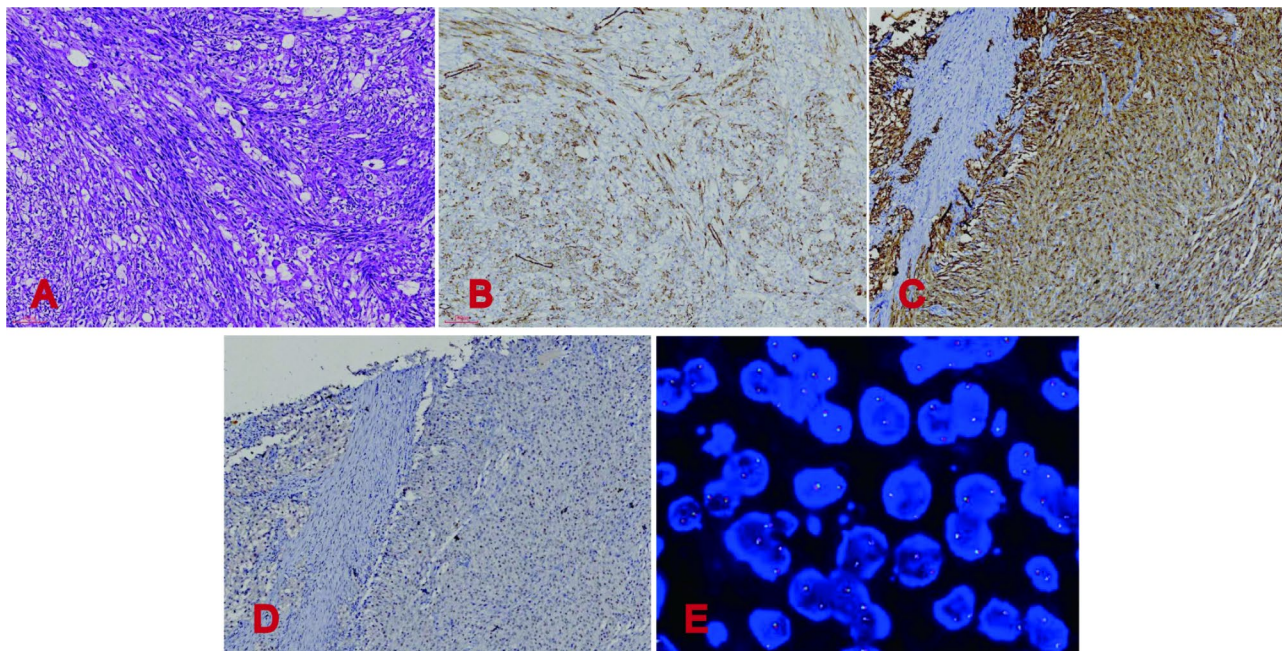


Fig. 2 (A) The tumor was composed of sheets of epithelioid cells and bundles of spindle cells, hematoxylin and eosin (H&E, $\times 100$). (B) Spindle cells were positive for smooth muscle actin (SMA) ($\times 100$). (C) Epithelioid cells were positive for HMB-45 ($\times 100$). (D) Epithelioid cells showed nuclear positivity of TFE3 ($\times 100$). (E) No *TFE3* gene rearrangement detected by FISH

The whole-body PET-CT scan was performed and revealed no abnormalities after the operation. Moreover, FISH showed no *TFE3* gene rearrangement (Fig. 2E). Next-generation sequencing for bone and soft tissue revealed negative *TSC1/2* and *TP53* expression.

A postoperative follow-up assessment was performed every 3 months for 2 years, every 6 months during years 3–5 and yearly thereafter. Routine follow-up appointments included a physical examination, vaginal examination, laboratory testing (including Ca125), chest radiography, and pelvic ultrasonography. Lung CT or pelvic MRI was performed when necessary. No recurrence or metastasis was observed during the 18-month follow-up period. Moreover, the gynecological ultrasound and tumour marker data remained normal.

Discussion and conclusions

Herein we report a case of uterine malignant PEComa without *TFE3* gene rearrangement detected by FISH. A combination of immunohistochemical and genetic testing is helpful for diagnosing PEComas. Genetically targeted therapy is more effective at improving overall survival.

PEComas in the uterus are rare. The clinical presentation is nonspecific and includes abnormal uterine bleeding, abdominopelvic pain, diagnosis of “fibroids,” or the identification of a mass on imaging. PEComas are easily misdiagnosed preoperatively as uterine fibroids [8]. Most cases are diagnosed accidentally or via quick-frozen pathology during surgery.

PEComas can be comprised of both epithelioid and spindle cells. PEComas uniquely show immunohistochemical positivity for both melanocytic (HMB-45, Melan-A) and myoid markers (SMA, desmin, caldesmon, and calponin), whereas cytokeratin and S-100 are generally negative [3]. PEComas can generally be considered present when HMB-45 is positively expressed because of its specificity. Currently, there is no unified standard for differentiating between benign and malignant uterine PEComas. According to Folpe’s criteria, PEComas are categorized as malignant if they demonstrate ≥ 2 of the following poor prognostic indicators: size ≥ 5 cm, significant nuclear atypia, invasive growth, mitosis $\geq 1/50$ HPF, necrosis, or evidence of lymphovascular invasion [7]. The pathologic features indicating a malignant PEComa in the index patient included invasive growth of the superficial 1/2 muscle layer, mitosis $\geq 1/50$ HPF, necrosis, and lymphovascular invasion.

Molecular and genomic profiling of endometrial cancer has increased in popularity in recent years. L1 cell adhesion molecule (L1CAM) is frequently mutated in endometrial cancer and is associated with a greater risk of distant recurrence, which provides a potentially useful tool for tailoring the need for adjuvant therapy [10, 11]. A few PEComas also show abnormal gene expression. Some PEComa patients have mutations in the *TSC1* and *TSC2* genes. A subset of PEComas has shown *TFE3* rearrangement [12]. *TFE3* is ubiquitously present at low levels in normal cells. When *TFE3* gene rearrangements occurs, *TFE3* protein overexpression is promoted, which

interferes with cell transcriptional regulation and leads to tumour formation [13]. Argani et al. [14] performed FISH analyses on *TFE3*-positive PEComas and confirmed that the *TFE3* gene rearrangement was accompanied by T (X; 1) (P11.2; P34) chromosome translocation, resulting in *PSF-TFE3* gene fusion, thereby promoting *TFE3* overexpression in tumour cells. *TFE3* protein is often strongly positive in immunohistochemistry, but this result does not indicate *TFE3* gene abnormalities in the FISH test. However, recent studies have shown that *TFE3* immunohistochemistry plays only a minor role in the diagnosis of *TFE3*-rearranged tumours. Thus, *TFE3* protein detected by immunohistochemistry alone is not sufficient to be used as a surrogate indicator of *TFE3* gene rearrangement; FISH analysis is recommended [15]. In the present case, the expression of *TFE3* was positive by immunohistochemistry but negative by FISH analysis, demonstrating the superior accuracy of FISH analysis.

Since PEComas with *TFE3* rearrangement are very rare, their exact biological behavior remains to be determined. Liu et al. [16] reported a more invasive case of malignant cervical PEComa accompanied by *TFE3* gene rearrangement. PEComas with *TFE3* gene rearrangements are considered more aggressive and should be considered independent subtypes of PEComas [7]. However, recent studies also revealed that PEComas with *TFE3* gene expression are benign [17]. In this case, the patient's tumour exhibited malignant biological behavior without *TFE3* rearrangement. Therefore, the relationship between *TFE3* gene rearrangement and the disease prognosis, as well as whether this factor should be considered a criterion for benign or malignant evaluation, needs to be proven with further research.

Standard treatment protocols are not yet available for these tumours owing to their rarity. Currently, surgical resection remains the preferred treatment. The choice of surgery depends on the patient's age and fertility requirements. Tumour resection alone may be considered only for patients who have fertility requirements and whose tumours are thought to be "benign". Shan et al. reported the case of a woman who had a natural pregnancy after tumour resection alone; she delivered a child and had a disease-free survival of 6 years [18]. According to the literature, total hysterectomy is the preferred treatment for patients without a fertility requirement [18]. The necessity for pelvic lymph node dissection is controversial because of the hematogenous metastasis of mesenchymal tumours in general, which needs to be verified by further studies. Whether postoperative adjuvant therapy is necessary for malignant PEComas has also been explored. The efficacy of chemoradiotherapy is uncertain [19]. Mutations in the *TSC1* and *TSC2* genes are driving factors in the development of some PEComas, resulting in activation of the mammalian target of rapamycin

(mTOR) pathway [8]. These alterations constitute the basis of mTOR inhibitor therapy. However, these findings need to be confirmed in additional clinical trials. The present patient's genetic test showed a negative TSC gene mutation, and she may have obtained limited benefit from mTOR inhibitors.

Currently, the diagnosis and treatment of uterine PEComa are predominantly based on case reports. Because of the lack of unified standards, surgery is the main treatment. However, genetically targeted therapy may be more effective. Further studies on the genomics, transcriptomics, proteomics, and epigenetics of PEComas are needed to identify criteria for accurately predicting outcomes and guiding disease-management decisions.

In conclusion, this case highlights the importance of a comprehensive approach for diagnosing PEComas, including genetic testing and immunochemical markers. Surgical resection is the first choice for treatment. Genetically targeted therapy is effective in improving the prognosis for patients with malignant PEComas. Long-term monitoring and follow-up are also needed.

Abbreviations

PEComa	Perivascular epithelioid cell tumour
TFE3	Transcription factor E3
TSC	Tuberous sclerosis complex
CEA	Carcinoembryonic antigen
AFP	Alpha fetoprotein
SCC	Squamous cell carcinoma antigen
PET-CT	Positron emission computed tomography
FISH	Fluorescence in situ hybridization
mTOR	The mammalian target of rapamycin

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Author contributions

MX participated in the acquisition of clinical data and drafted the manuscript. JHF carried out the pathological examination and interpretation. LZC revised the manuscript. All authors have read and approved the final manuscript.

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Data availability

The data supporting the conclusions of this article is available from corresponding author on reasonable request.

Declarations

Ethics approval and consent to participate

This study was approved by the Ethics Committee of the Fujian Provincial Maternity and Children's Hospital (No. 2024KY040).

Consent for publication

Written informed consent was obtained from the patient for publication of this case report and any accompanying images.

Competing interests

The authors declare no competing interests.

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