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Misdiagnosis of synovial sarcoma: Cellular myofibroma with *SRF-RELA* gene fusion:

A case report

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Zhou Y *et al.* Cellular myofibroma

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Abstract

BACKGROUND

Cellular myofibroma is a rare subtype of myofibroma that was first described in 2017. Its diagnosis is often challenging because of its relative rarity, lack of known genetic abnormalities, and expression of muscle markers that can be confused with sarcomas that have myogenic differentiation. Currently, scholars have limited knowledge of this disease, and published cases are few. Further accumulation of diagnostic and treatment experiences is required.

CASE SUMMARY

A 16-year-old girl experienced left upper limb swelling for 3 years. She sought medical attention at a local hospital 10 months ago, where magnetic resonance imaging revealed a 5-cm soft tissue mass. Needle biopsy performed at a local hospital resulted in the diagnosis of a spindle cell soft tissue sarcoma. The patient was referred to our hospital for limb salvage surgery with endoprosthetic replacement. She was initially diagnosed with a synovial sarcoma. Consequently, clinical management with chemotherapy was continued for the malignant sarcoma. Our pathology department also performed fluorescence in situ hybridization for result validation, which returned negative for SS18

gene breaks, indicating that it was not a synovial sarcoma. Next-generation sequencing was used to identify the *SRF-RELA* rearrangement. The final pathological diagnosis was a cellular/myofibroblastic neoplasm with an *SRF-RELA* gene fusion. The patient had initially received two courses of chemotherapy; however, chemotherapy was discontinued after the final diagnosis.

CONCLUSION

This case was misdiagnosed because of its rare occurrence, benign biological behavior, and pathological similarity to soft tissue sarcoma.

Key Words: Cellular myofibroma; *SRF*; *RELA*; Fusion; Pathology; Case report

¹ Zhou Y, Sun YW, Liu XY, Shen DH. Misdiagnosis of synovial sarcoma: Cellular myofibroma with *SRF-RELA* gene fusion: A case report. *World J Clin Cases* 2024; In press

Core Tip: Cellular myofibroma (CMF) is a rare subtype of myofibroma that exhibits a broad age distribution, and manifests in diverse anatomical locations, including deep soft tissues and skeletal muscles. Owing to the absence of distinct features, CMF is frequently misdiagnosed. Microscopic examination revealed active mitotic figures (non-pathological), focal areas of necrosis, and occasional infiltrative growth. Next-generation sequencing aids in the accurate pathological diagnosis of CMF by detecting the *SRF-RELA* gene fusion.

INTRODUCTION

Myofibromas are common benign mesenchymal tumors that predominantly affect infants and young children. This disease was first reported by Williams and Sohm in 1951, and was initially referred to as congenital fibrosarcoma. In 1954, Stout renamed juvenile fibromatosis as congenital generalized fibromatosis. However, in 1981, Chung and Enzinger demonstrated that this tumor is actually a myofibroblastic proliferation, leading to its reclassification as infantile myofibromatosis^[1]. The 2002 World Health Organization (WHO) classification categorizes it as one of the most common fibroblastic and myofibroblastic proliferations in infants and children. Moreover, morphologically, myofibromas share some continuity with myopericytomas; thus, they were included in the category of myopericytic tumors in the 2013 WHO classification.

² In 2017, a subset of cellular variants of myofibroma and myopericytoma with a smooth muscle-like immunophenotype, harboring a recurrent *SRF-RELA* gene fusion was reported. Diagnosing cellular myofibroma (CMF) presents certain challenges as it is rare, and its histological features resemble those of other sarcomas^[2]. Furthermore, CMF is characterized by high cell density and visible nuclear divisions. Some CMF tumors exhibit a distinct fascicular growth pattern that morphologically resembles that of spindle cell sarcomas such as rhabdomyosarcoma, leiomyosarcoma, and synovial sarcoma. This type of tumor lacks specific immunohistochemical markers and exhibits significant overlap with low-grade myofibroblastic sarcomas and fibroblastic tumors within the immunophenotypic spectrum. This necessitates the use of next-generation sequencing (NGS) to detect characteristic genetic aberrations. Therefore, the differential diagnosis of CMF requires consideration of other sarcomas with similar morphologies or similar expression of immunohistochemical markers. Additionally, CMF is even rarer in young or adult patients and is more likely misdiagnosed as "sarcoma" in older patients.

Herein, we report a case of CMF and discuss its clinical, pathological, and molecular features to elucidate the differential diagnosis of this rare tumor subtype.

CASE PRESENTATION

Chief complaints

The patient had a palpable lump in the left arm for > 3 years and numbness in the fingers for > 1 month.

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History of present illness

A 16-year-old girl was admitted to the Bone Tumor Department of our hospital with left upper limb pain that had started 1 wk prior to presentation. The patient first noticed symptoms of left upper limb pain 3 years prior when she accidentally discovered a painless "walnut-sized" lump while bathing. However, she did not seek medical attention at that time. Over time, the lump size gradually increased. She had recently started to experience numbness in her fingers and decided to seek medical care at a local hospital. Magnetic resonance imaging (MRI) of the left humerus revealed a 5-cm soft tissue mass surrounding the bone. A fine-needle aspiration biopsy was performed at the local hospital, and the pathological report indicated the possibility of a malignant soft tissue tumor.

History of past illness

The patient had been previously healthy.

Expert consultation

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The patient was transferred to our hospital for expert consultation and further treatment on June 23, 2023. Considering the immunohistochemical staining results from the original institution, along with the imaging findings, the diagnosis was consistent with spindle cell mesenchymal sarcoma. The differential diagnoses included synovial sarcoma, malignant peripheral nerve sheath tumor, and other tumors with similar presentations.

Personal and family history

No evident abnormality was identified in the personal and family history.

Physical examination

A firm lump was palpated on the anterolateral aspect of the middle segment of the left upper arm. It was immobile and approximately 5 cm × 3 cm in size. Mild tenderness and scarring from the previous punch biopsy were observed.

Laboratory examinations

No evident abnormality in the serum tumor markers was detected.

Imaging examinations

Left humerus radiography (Figure 1A), computed tomography (Figure 1B), and MRI (Figure 1C) examinations revealed bone destruction in the middle segment of the left humerus, accompanied with the formation of a soft tissue mass.

Treatment

On July 7, 2023, the patient underwent segmental resection of the left humerus with prosthetic replacement (Figure 1D).

FURTHER DIAGNOSTIC WORK-UP

Postoperative pathological examination

Macroscopic examination: The specimen consisted of muscle and adipose tissues with visible tumors. The tumor appeared as multiple nodules with a total size of 5 cm × 4.5 cm × 2 cm. The diameter of each nodule ranged from 1.5 to 2 cm. The cut surface of the tumor was solid, slightly tough in texture, and grayish-yellow in color. No clear areas of bleeding were observed (Figure 2A).

Microscopic examination: Nodular tumors were observed in the fibromuscular tissue (Figure 2B). The cells were oval- or spindle-shaped (Figure 2C), densely arranged in bundles or swirling patterns. Focal necrosis and epithelioid cell-like regions were

observed (Figure 2D). ⁷ The average mitotic count was approximately 10/10 high-power fields (HPF), with visible focal mitosis (approximately 20/10 HPF) (Figure 2E).

Immunohistochemical staining: SATB2 (+), INI1 (+), S-100 (partially +), Sox-10 (-), ALK (-), CD99 (focally +), TLE1 (diffusely strong +) (Figure 2F), Desmin (focally +), SMA (+), Myogenin (-), Bcl-2 (focal +), CD34 (-), Ki-67 (30%+), H3K27Me3 (+), EMA (-), CD56 (+), CK19 (-), MUC4 (-), and Calretinin (-).

Preliminary pathological diagnosis

The preliminary pathological diagnosis was considered as synovial sarcoma (spindle cell type).

Genetic testing

¹⁰ Fluorescence in situ hybridization (FISH) analysis revealed a separation rate of 6/100 = 6% for SS18, indicating a negative result (Figure 2G). Additionally, NGS revealed an *SRF-RELA* rearrangement (Figure 2H).

FINAL DIAGNOSIS

The final pathological diagnosis was a cellular fibroma with *SRF-RELA* fusion.

TREATMENT

The patient received two cycles of chemotherapy with pegylated liposomal doxorubicin 70 mg/d, combined with ifosfamide 9 g/d postoperatively. Chemotherapy was discontinued based on the final diagnosis.

OUTCOME AND FOLLOW-UP

The patient was followed-up for 3 months, and no recurrence was noted.

DISCUSSION

Cellular fibromas, also known as atypical fibromas, are benign with no metastasis and limited growth^[3-4]. Surgical removal typically results in a good prognosis, with a local recurrence rate of approximately 5%. However, CMF may be misdiagnosed owing to its histological similarity to soft tissue sarcomas^[5]. Limited research has been conducted on CMF, domestically and internationally, and pathologists have not paid sufficient attention to diagnosing and distinguishing spindle cell tumors of soft tissues. Previous case reports have highlighted the misdiagnosis of CMF as leiomyomas, low-grade malignant fibrous histiocytomas, fibrosarcomas, and rhabdomyosarcomas. However, subsequent confirmation through NGS revealed the presence of the *SRF-RELA* gene fusion^[6-7]. Positive outcomes were observed during the follow-up.

Furthermore, CMFs can occur in children, adolescents, and adults of varying ages and manifests in different anatomical sites, including deep soft tissues and skeletal muscles^[8]. They typically present as nodular or multicystic masses with well-defined borders. Histologically, CMF tumor tissues display characteristics such as high cell density and solid or focal infiltrative growth. The cellular-rich fibromatous lineage closely resembles pediatric malignancies such as infantile fibrosarcoma or synovial sarcoma. Oudijk *et al*^[9] identified a subset of 9 out of 114 conventional fibromatous tumors that resembled fibrosarcoma, which they referred to as the cellular-rich subtype. Moreover, immunohistochemistry revealed diffuse SMA positivity and desmin expression in these tumors. Tumors with *SRF* fusion have been observed to exhibit a consistent smooth muscle-like immunophenotype, which leads to misdiagnosis as myogenically-differentiated sarcomas^[2,10].

The diagnostic journey of the patient described in this article was intricate and demanding. The patient was initially diagnosed with a leiomyosarcoma based on a biopsy conducted at a local hospital. However, upon referral to our hospital for consultation, malignant mesenchymal tumors were considered. The patient received surgical intervention and postoperative chemotherapy based on the suspicion of a malignant mesenchymal tumor. Immunohistochemical staining of the postoperative specimen revealed a strong and diffuse expression of TLE1 in the tumor cells, suggesting

a preliminary diagnosis of synovial sarcoma. However, further examination using the SS18 gene probe dual-color break-apart FISH yielded a negative result. Nevertheless, NGS gene testing identified the *SRF-RELA* gene fusion. Based on the onset location, histological morphology, and immunohistochemical findings, a final diagnosis of cellular fibroma was established.

In this case, the diagnostic challenge arose from the unusual location, age of onset, presence of active mitotic figures, abundant cellular content, and diffusely strong expression of the TLE-1 protein in tumor cells. After reviewing the histopathological slides and existing literature, we identified several clinical and pathological features of CMF. Generally, CMFs have well-defined borders that distinguish them from classic sarcomas^[11]. They do not exhibit the typical "fish-flesh" appearance upon cross-section, and extensive areas of necrosis or hemorrhage are rarely observed under microscopic examination. Additionally, CMFs tend to grow at a slower pace than classic sarcomas and do not undergo sudden enlargement^[12]. In this particular case, the tumor had a long duration and demonstrated slow growth with focal infiltrative growth or limited stromal wedge-shaped infiltration, all of which are not characteristic of malignancy^[13]. Upon reviewing the literature, we identified that this is the first reported case of CMF expressing TLE-1, which further complicated the diagnosis. Moreover, TLE-1 is commonly observed in synovial sarcoma tumor cells; however, fibromas have not been reported to express TLE-1. Both tumors typically present as well-defined nodules and share similarities in tissue morphology, such as a bundled cell arrangement, abundant cellularity, and short spindle-shaped cytology. This poses significant diagnostic challenges. This case highlights the need for future studies to differentiate between synovial sarcoma and fibroma, and to enhance our understanding of this rare tumor subtype. Although active mitotic figures were present in this case, the absence of pathological or atypical mitotic figures did not support the diagnosis^[14]. Our literature review did not reveal any relevant reports on TLE-1 expression in fibromas or cellular fibromas. Through a retrospective analysis, we observed that cellular or atypical fibromas

may exist. However, owing to the limited number of evidence-based cases of CMF, the biological behavior of this tumor remains unknown, necessitating close follow-up.

⁵ The 5th edition of the WHO classification of soft tissue and bone tumors in 2020 identified the *SRF-RELA* rearrangement as an aid in the pathological diagnosis of cellular/atypical fibroma^[9]. Additionally, *SRF* is a protein-coding gene that stimulates cell proliferation and differentiation. Similarly, ⁹ *RELA* (*RELA* proto-oncogene, NF-κB subunit) is another protein-coding gene^[15]. In this case, the *SRF-RELA* gene rearrangement was consistent with previous findings. The fusion gene retained the 5' end of the *SRF* gene promoter up to exon 4 and the 3' end of the *RELA* gene from exon 11 to the termination codon, without any predicted frameshift mutations.

CONCLUSION

With the rapid development of NGS technology, breakthroughs in the classification of classic tumors have been achieved. Currently, with the application of this technology, various tumors can be diagnosed more accurately, including those that were previously misdiagnosed as low-grade sarcomas or synovial sarcomas. Using second-generation sequencing technology, we discovered that these tumors were benign. This article describes a rare case of cellular fibroma with the *SRF-RELA* gene fusion. The tumor exhibits atypical and intriguing cytological features that may be mistaken for malignant sarcomas with smooth or striated muscle differentiation. The current follow-up results demonstrate a good prognosis for the patient. This case further suggests that the accuracy of pathological diagnosis relies on the assistance of new technologies, and that diagnosis based on emerging technologies must be mutually complementary with traditional pathology

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