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Case report: Primary pleural low-grade fibromyxoid sarcoma in a 4-year-old boy with molecular confirmation

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Low-grade fibromyxoid sarcoma (LGFMS) is a rare malignant fibroblastic tumor, principally affecting the deep tissues of the proximal trunk and extremities in young adults. However, primary pleural LGFMS is extremely rare, and only three cases have been reported in the previous English literature without genetic confirmation. Furthermore, the historical pleural LGFMS cases were all adults, and the primary pleural LGFMS in children has never been reported to date. Here, we presented a primary pleural LGFMS in a 4-year-old boy with detailed clinical, pathological, and molecular results. Histologically, the current tumor showed typical alternating collagenous and myxoid areas, containing spindled or oval tumor cells arranged in a whorled and short fascicular pattern. In some areas, the tumor cells exhibited moderate atypia, and mitotic figures were identified but without the identification of giant collagen rosettes. Immunohistochemically, all the neoplastic cells showed strong and diffuse positivity for MUC4. Genetically, FUS gene rearrangement was revealed by fluorescence in-situ hybridization (FISH), and subsequently, next-generation sequencing (NGS) and polymerase chain reaction (PCR) further demonstrated the FUS::CREB3L2 fusion transcript. To the best of our knowledge, this is the first case of primary pleural LGFMS with the identification of FUS gene rearrangement and FUS::CREB3L2 fusion in a 4year-old child. Our study expands the age range of pleural LGFMS and highlights the combination of morphological, immunohistochemical, and molecular analyses in such challenging cases.

KEYWORDS

low-grade fibromyxoid sarcoma, pleural tumor, pediatric sarcoma, *FUS::CREB3L2* gene fusion, molecular analysis

Introduction

Low-grade fibromyxoid sarcoma (LGFMS) is a rare malignant fibroblastic neoplasm, mainly occurring in the deep tissues of the proximal trunk and extremities. The tumors typically arise in young adults with a slight male predilection, and approximately 20% of cases are <18 years of age (1–3). Morphologically, classic LGFMS is composed of fibrous and myxoid areas, with bland, spindled cells in a whorled or fascicular growth pattern, and easy to be misdiagnosed as a benign tumor by histology only. Immunohistochemically, MUC4 has been reported as a sensitive and specific marker for the diagnosis of LGFMS. Genetically, more than 90% of LGFMS cases harbor t(7;16)(q33; p11) translocation, resulting in *FUS::CREB3L2* fusion gene, and a minority of the cases contain *FUS::CREB3L1* or *EWSR1::CREB3L1* fusions. The identification of *FUS::CREB3L2* or other rare fusions by molecular analysis could aid in the diagnosis of LGFMS (4–6).

In recent years, it has been described that LGFMS can also arise from other unusual sites, including the head/neck, abdominopelvic, retroperitoneal, gastrointestinal viscera, and mediastinum. It is worth highlighting that primary pleural LGFMS is extremely rare. To the best of our knowledge, only a total of three cases of primary pleural low-grade fibromyxoid sarcoma have been reported to date, and all lacked genetic validation (7–9).

We presented a primary pleural LGFMS case with the identification of *FUS* gene rearrangement by fluorescence *in-situ* hybridization (FISH) and the *FUS::CREB3L2* fusion gene by next-generation sequencing (NGS) and polymerase chain reaction (PCR). Moreover, although 20% of LGFMS cases aged <18 years, primary pleural LGFMS in children has never been reported before. Here, we reported the first genetically confirmed pleural LGFMS case in a 4-year-old boy, with detailed clinical, pathological, and molecular information.

Case presentation

A 4-year-old boy was admitted to a local hospital with a 1month history of cough and fever and diagnosed with "pneumonia." Contrast-enhanced computed tomography (CT) revealed a well-defined, heterogeneous soft tissue mass measuring 2.2 cm \times 1.9 cm, occupying the left side of the posterior mediastinal paravertebral region (Figure 1).

The patient was admitted to the department of pediatric surgery, and surgery was performed in August 2022. During the operation, a well-circumscribed mass was identified under the visceral pleura and clearly demarcated from the lung parenchyma. The patient underwent complete resection of the mass. Grossly, this lesion revealed a well-demarcated mass measuring 5 cm at its greatest dimension. On the cut surface, the mass demonstrated a white color and a brittle texture without obvious necrosis. The pathologists of the peripheral hospital first considered the diagnosis of the current case as a spindle-cell proliferative lesion, which was inclined to the tumor, and they suggested consultation to further clarify the diagnosis.

Our department received consultation slides from the peripheral hospital. Microscopically, the lesion was relatively well delineated and composed of fibrous stroma and myxoid areas with an abrupt transition (Figures 2A, B). The tumor cells were spindle and oval-shaped, arranged in a whorled and short fascicular pattern (Figure 2C). In some areas, the lesion had increased cellularity, and the neoplastic cells had spindled hyperchromatic nuclei showing moderate atypia, with indistinct pale eosinophilic cytoplasm and a few visible nucleoli (Figure 2D), and mitotic figures were identified without atypical forms (2/10 high power fields) (Figure 2E). Immunohistochemically, all the tumor components stained diffusely and strongly positive for MUC4 (Figure 2F). The neoplastic cells were negative for desmin, S-100, STAT-6, myogenin, CD34, ERG, TLE1, β-catenin, EMA, myoD1, SOX10, ALK, TRK, SMA, and loss of H3K27me3. The Ki-67 (MIB-1) index was 5% of these cells.

FISH analyses for *FUS*, *EWSR1*, and *MDM2* were performed using the GSP *FUS* gene probe (Anbiping, Guangzhou, China), GSP *EWSR1* gene probe (Anbiping, Guangzhou, China), and GSP *MDM2* (12q15) Gene Amplification probe (Anbiping, Guangzhou, China). The FISH results demonstrated the presence of *FUS* gene rearrangement (Figure 3A). In addition, the tumor was negative for *EWSR1* gene rearrangement and *MDM2* gene amplification.

Subsequently, next-generation sequencing (NGS)-based technology (1,084 cancer-relevant genes, hybrid DNA/RNA panels, Genetron Health, Beijing, China) was applied to the formalin-fixed paraffin-embedded tissue of the lesion. Notably, a *FUS::CREB3L2* fusion was identified at both the DNA and RNA levels. A fusion transcript of *FUS* (exon 7, transcript NM_004960.3) and *CREB3L2* (exon 5, NM_194071.4) was discovered (Figure 3B). Then, we further validated the presence of the *FUS::CREB3L2* fusion transcript by PCR (polymerase chain reaction) and Sanger sequencing using primers (*FUS*-forward: 5'-GTCTGATTGTTCA TTTGCAGA-3' and *CREB3L2*-reverse: 5'-GACTCAGAG GGCAGCCTGAGT-3') (Figure 3C). Based on the histological,



FIGURE 1

Imaging findings. The contrast-enhanced computed tomography showed a hypodense mass occupying the left side of the posterior mediastinal paravertebral region (arrow).



FIGURE 2

The histological and immunohistochemical results of the tumor. (A) The lesion was composed of fibrous stroma and myxoid areas with an abrupt transition [hematoxylin and eosin staining (H β E); magnification: ×40]. (B) The classic myxoid areas and fibrous stroma were shown in the tumor (H β E; magnification: ×100). (C) Neoplastic cells were spindle-shaped and arranged in a whorled and short fascicular pattern (H β E; magnification: ×200). (D) Some areas had increased cellularity with moderate atypia (H β E; magnification: ×400) with the identification of mitotic figure (arrow) [(E) H β E; magnification: ×400]. (F) The tumor cells showing diffuse and strong positivity for MUC4 (magnification: ×400).

immunohistochemical, and molecular results, the pleural tumor was diagnosed as LGFMS.

Considering that LGFMS mainly occurs in the deep tissues of the proximal trunk and extremities, a complete physical examination was performed again in this case to exclude the possibility that the pleural LGFMS was a metastasis from the trunk or extremities and revealed no palpable mass. PET-CT was carried out to further rule out the possibility that the pleural tumor was a metastasis and revealed no other lesions. Finally, combining the clinical, radiological, and pathological findings and molecular results, the diagnosis was primary LGFMS of the pleura. At the most recent follow-up, 11 months following the surgery, the patient was in a good status with no evidence of disease.

This study was approved by the West China Hospital Institutional Review Board.

Discussion

LGFMS mainly occurs in young adults, and approximately 20% of cases are <18 years of age. However, LGFMS arising in children aged <5 years is extremely rare. A SNOMED search of the West China Hospital surgical pathology files from July 2008 to January 2023 has identified 36 LGFMS cases, while only 2 cases younger than 5 years old were detected. Subsequently, we carefully reviewed the previously reported LGFMSs in English literature, and 122 cases (before 18 years old) have been described. Moreover, only 22 cases (22/122, 18.0%) under 5 years old were reported (10–21). Among the 22 cases, the tumors mainly occurred in the extremities (8/22, 36.5%), followed by the trunk (7/22, 31.8%), head and neck regions (5/22, 22.7%), liver (1/22, 4.5%), and abdominal cavity (1/22, 4.5%). To the best of our knowledge, the current case is the first primary



LGFMS case that occurred in the pleura of this age group (under 18 years old) in English literature.

In fact, primary pleural LGFMS is exceedingly scarce and only three cases have been reported before (7–9). In 2005, Kim et al. reported the first pleural LGFMS in a 37-year-old man, and the second case was described in a 42-year-old woman. However, MUC4 immunostaining and molecular testing were not performed in these cases. Recently, Perez et al. reported pleural LGFMS in a 32-year-old man, with MUC4 positivity. Our case is the first pleural LGFMS in a 4-year-old child that was immunohistochemically and genetically confirmed. Furthermore, primary intrathoracic LGFMS is also very rare, and only 27 cases have been reported before (including the current case) (Table 1) (7–9, 22–42). The historical intrathoracic cases were composed of 13 male and 14 female patients, aged 4 to 77 years old (median 35 years). These tumors involved the mediastinum (10/27, 37.0%), lung parenchyma (10/27, 37.0%), pleura (4/27, 14.8%), heart (1/27, 3.7%), intrathoracic/pericardium (1/27, 3.7%), and right heart/epicardium (1/27, 3.7%). It needs to be pointed out that intrathoracic LGFMS could sometimes be extremely large, and it may be difficult to determine where the lesion arises from. These results indicated that primary intrathoracic LGFMS cases, especially

TABLE 1 Clinicopathologic and molecular features of previously reported primary intrathoracic LGFMS cases.

Case	References	Age	Sex	Symptoms	Location	Size (cm)	Histological features	IHC		Genetic results				Treatment	Outcome
no.									Ki-	FISH			Sequencing		
								MUC4	67	FUS	ESWR1	Others			
1	Current case	4	М	Cough	Pleura	5	Classic LGFMS	(+)	5%	(+)	(-)	MDM2 amplification (–)	FUS::CREB3L2 (NGS and PCR)	Complete resection	11 mo/NED
2	Perez, D. et al. (7)	32	М	Cough	Pleura	11	LGFMS with collagen rosettes	(+)	ND	ND	ND	SYT rearrangement (–)	ND	Surgical resection	29 mo/NED
3	Liang, W. et al. (8)	42	F	Cough, shortness of breath	Pleura	11	Classic LGFMS	ND	ND	ND	ND	ND	ND	Marginal excision, radiotherapy	NA
4	Kim, S. Y. et al. (9)	37	М	Chest pain, dyspnea	Pleura	NA	Classic LGFMS	ND	ND	ND	ND	ND	ND	Surgical resection	NA
5	Ershadi, R. et al. (22)	26	F	Chest pain, shortness of breath	Left lung	30	LGFMS with collagen rosettes and epithelioid cell	ND	Low	ND	ND	ND	ND	Surgical resection	6 mo/NED
6	Yoshimura, R. et al. (23)	22	М	Asymptomatic	Right upper lung	4.5	LGFMS with collagen rosettes	(+)	ND	(-)	ND	ND	ND	Surgical resection	NA
7	Bartuma, H. et al. (24)	77	М	NA	Right upper lung	5	LGFMS with collagen rosettes	ND	ND	(+)	ND	ND	FUS:: CREB3L2 (PCR)	Surgical resection	NA
8	Sargar, K. et al. (25)	12	М	NA	Left lower lung	NA	NA	ND	ND	ND	ND	ND	ND	NA	NA
9	Oramas, D. M. et al. (26)	37	F	Cough, dyspnea, chest pain	Left lower lung	2.4	LGFMS with collagen rosettes	ND	ND	ND	ND	ND	ND	Surgical resection	6 mo/NED
10	Oramas, D. M. et al. (26)	42	М	Cough, dyspnea, chest pain	Right upper lung	3	LGFMS with collagen rosettes	ND	ND	ND	ND	ND	ND	Surgical resection	14 mo/NED
11	Magro, G. et al. (27)	20	F	Asymptomatic	Lung	2	LGFMS with collagen rosettes	ND	ND	ND	ND	ND	ND	Biopsy	12 mo/AWD
12	Kim, L. et al. (28)	50	F	Chest pain	Left lower lung	7.5	LGFMS with collagen rosettes	ND	ND	ND	ND	ND	FUS:: CREB3L2 (PCR)	Surgical resection	NA
13	Whale, K. et al. (29)	66	F	Asymptomatic	Right lung	2.6	LGFMS with collagen rosettes and SEF area	ND	2%	(+)	ND	ND	ND	Surgical resection	NA

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TABLE 1 Continued

Case	References	Age	Sex	Symptoms	Location	Size	Histological features	IHC		Genetic results				Treatment	Outcome
no.						(cm)			1/2	FISH			Sequencing		
								MUC4	Кі- 67	FUS	ESWR1	Others			
14	Kurul, İ. C. et al. (30)	16	F	Pain in the arms and legs	Lung	10	Classic LGFMS	ND	ND	ND	ND	ND	ND	Surgical resection	24 mo/NED
15	Mustafa, S. et al. (31)	44	F	NA	Mediastinum	12	LGFMS with collagen rosettes	ND	ND	(+)	ND	ND	ND	Surgical resection	NED
16	Xie, Y. et al. (32)	32	F	Shortness of breath, cough	Mediastinum	14	Classic LGFMS	ND	ND	ND	ND	ND	ND	Surgical resection	36 mo/NED
17	Sajid, M. I. et al. (33)	26	М	Fever, vomiting	mediastinum	17	LGFMS with collagen rosettes	(+)	ND	ND	ND	ND	ND	Surgical resection	22 mo/NED
18	Maeda, E. et al. (34)	50	М	Asymptomatic	Superior mediastinum	13	Classic LGFMS	ND	ND	ND	ND	ND	ND	Surgical resection	60 mo/NED
19	Maeda, E. et al. (34)	19	F	Asymptomatic	Anterior mediastinum	23.5	LGFMS with collagen rosettes	ND	ND	ND	ND	ND	ND	Surgical resection	Recurrence at 60 mo, resected; 60 mo/AWD
20	Takanami, I. et al. (35)	35	М	Asymptomatic	Anterior mediastinum	9	Classic LGFMS	ND	ND	ND	ND	ND	ND	Surgical resection	Recurrence at 108 mo, resected; 110 mo/NED
21	Aissaoui, R. et al. (36)	29	М	Dyspnea	Mediastinum	18.9	Classic LGFMS	ND	ND	ND	ND	FUS::DDIT3 (-) ESWR1:: DDIT3 (-)	ND	Biopsy	NA
22	Williams, C. M. et al. (37)	50	М	Shortness of breath	Mediastinum	NA	LGFMS with epithelioid, round cell	ND	50%	ND	ND	ND	FUS:: CREB3L2 (NGS)	Biopsy, chemotherapy	5 mo/died of pericardial effusion and tamponade
23	Galetta, D. et al. (38)	41	М	Asymptomatic	Anterior mediastinum	8	LGFMS with collagen rosettes	ND	ND	ND	ND	ND	ND	Surgical resection radiotherapy	35 mo/NED
24	Gülhan, S. Ş. E. et al. (39)	25	F	Back pain and dyspnea	Posterior mediastinum	17	LGFMS with collagen rosettes	ND	ND	ND	ND	ND	ND	Surgical resection	NA

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Outcome			12 mo/NED	7 mo/NED	Recurrence, died of cardiac failure at 84 mo aration sequencing;	ò -
Treatment			Chemotherapy, surgical resection	Surgical resection	Surgical resection zation; NGS, next gen	о
	Sequencing		FUS:: CREB3L2 (NGS)	QN	<i>FUS</i> :: <i>CREB3L2</i> (PCR) rescence <i>in situ</i> hybridi	
enetic results	Ŧ	Others	ND	ND	ND reaction; FISH, fluo	in reaction; FISH, flu
Ū	FISH	ESWR1	QN	QN	ND olymerase chaii	
		FUS	(+)	(+)	ND V; PCR, po	T
	r, K	67	ŊŊ	Ŋ	ND tochemistr	
HI		MUC4	(+)	ŊŊ	ND	ĸ
Histological	reatures		Classic LGFMS	LGFMS with collagen rosettes	Classic LGFMS (primary); LGFMS with pleomorphic areas (recurrent) oositive: "-", negative; IHG	sease: mo, month.
Size	(cm)		27	12	5 coma: "+", ;	live with di
Location			Intrathoracic/ pericardium	Right heart/ epicardium	Right ventricle epithelioid fibrosar	of disease: AWD. a
Symptoms			Cough and weight loss	Cough, retrosternal discomfort	Right cardiac failure oma: SEF, sclerosing	disease: DOD. dead
Sex			щ	щ	F	ridence of
Age			6	44	57 ade fibrom	JED, no ev
References			Chan, Y. C. et al. (40)	Jakowski, J. D. et al. (41)	Ferlosio, A. et al. (42) emale: LGFMS, low-gre	ilable: ND, not done: N
Case	OL		25	26	27 M, male: F, fi	NA, not avai

the pleural lesions, are exceedingly rare, and the diagnoses of LGFMS in such sites are extremely challenging.

Histologically, classic LGFMS is composed of alternating fibrous and myxoid areas with the proliferation of bland, spindled cells in a whorling or short fascicular pattern. Almost 30% of LGFMS cases exhibit collagenous rosettes and a subset of cases harbor some unusual features, including the presence of sclerosing epithelioid fibrosarcoma (SEF)-like areas and epithelioid or round tumor cells with increased pleomorphism and atypia (3). In the reported 27 intrathoracic LGFMS cases, 26 cases had available histology information (including the current case). The morphologic results showed that 9 (9/26, 34.6%) cases were classic LGFMS and 13 (13/26, 50.0%) cases were LGFMS with a collagen rosette structure. In addition, 4 (4/26, 15.4%) cases had unusual features, consisting of 1 case with collagen rosettes and epithelioid component, 1 case with collagen rosettes and SEF-like areas, 1 lesion with round and epithelioid cell areas, and 1 case with classic morphology in the primary tumor and pleomorphic areas in the recurrent tumor (22, 29, 37, 42). The frequency of these unusual features in the intrathoracic LGFMS (15.4%) was slightly higher than that of the overall LGFMS cases (<10%) (1). Furthermore, the frequency of collagen rosettes in pulmonary LGFMS cases was particularly high, reaching 80% (8/9, 88.9%). These results revealed that giant collagen rosettes were more common in the intrathoracic cases, especially in the pulmonary lesions.

Immunohistochemically, MUC4 is a sensitive and specific marker for LGFMS, and 80% of LGFMS cases are positive for epithelial membrane antigen (EMA) and 30% show positivity for SMA (43). The current case was positive for MUC4 and negative for EMA and SMA. In the historical intrathoracic LGFMS cases, MUC4 immunohistochemistry was carried out in 4 cases and all of them displayed positivity, indicating that MUC4 immunostaining is a useful ancillary diagnostic tool (7, 23, 33, 40). Notably, MUC4 could also show positivity in SEF and fusion-positive alveolar rhabdomyosarcomas (44). In such difficult cases, molecular testing for *FUS* gene rearrangement could be helpful.

Cytogenetically, over 90% of LGFMSs harbor FUS gene rearrangement, with FUS::CREB3L2 or FUS::CREB3L1 fusion gene. Additionally, a subset of LGFMSs have EWSR1 gene rearrangement, with EWSR1::CREB3L1 fusion (5, 6). Our present case was the first pleural case with confirmed FUS gene rearrangement and FUS::CREB3L2 fusion transcript. Eleven historical intrathoracic cases had molecular results, including 3 cases with FUS rearrangement and 5 cases with FUS::CREB3L2 fusion gene. The genetic analysis of 2 cases was performed for differential diagnosis, and another 1 case was proved to be negative for FUS rearrangement but positive for MUC4 immunostaining (24, 28, 29, 31, 37, 40-42). MUC4 could also be positive in other lesions, such as in alveolar rhabdomyosarcomas (44). In such cases, FISH for EWSR1 rearrangement and NGS would be helpful to confirm the diagnosis. Moreover, we endorsed genetic analysis for cases with ambiguous morphology in such rare locations.

It is worth noting that the current case exhibited extraordinarily rare clinicopathological features. First, the pleura is an extremely rare location for LGFMSs. Second, LGFMS is exceedingly rare in this age group (under 5 years old). Third, collagen rosettes were

FABLE 1 Continued

absent in the present case, which seemed to be common in the intrathoracic LGFMS cases. Hence, the differential diagnosis of this case is extremely challenging, and this tumor must be distinguished from other fibrous or myxoid lesions.

Inflammatory myofibroblastic tumor (IMT) often occurs in the pleuropulmonary area, outnumbering LGFMS. The current tumor was diffusely positive for MUC4 and negative for ALK, which is usually positive in IMT tumors. More importantly, IMT can be excluded because of the presence of the *FUS::CREB3L2* fusion gene in the current case, as IMT cases mainly harbor the rearrangement of the *ALK* gene (45).

Solitary fibrous tumor (SFT) is another common soft tissue tumor in this area and shares some morphologic features with LGFMS. However, the identification of MUC4 positivity could exclude the diagnosis of SFT. Furthermore, the genetic hallmark of SFT is the *NAB2::STAT6*, and LGFMS is featured by the *FUS* gene rearrangement (46).

Synovial sarcoma (SS) can also arise in the pleuropulmonary sites. Nevertheless, SSs harbor more conspicuous cellularity and positivity with EMA, TLE1, and cytokeratins. Cytogenetically, SSs are characterized by the t(X;18) (p11.2;q11.2) translocation, which leads to *SS18-SSX* gene fusion (47).

Desmoid fibromatosis with myxoid change could be also confused with LGFMS. However, almost 80% of desmoid fibromatosis was positive for β -catenin, which was negative in the present case (48). The majority of desmoid fibromatosis harbors somatic *CTNNB1* gene mutation, and a subset of cases arising in Gardner syndrome patients have *APC* gene mutation, which was different from the genetic hallmark of LGFMS.

Although dedifferentiated liposarcoma (DDL) rarely occurs in the thorax, it is necessary to differentiate LGFMS from DDL, as the two entities have different behaviors. DDL is usually positive for MDM2 and CDK4 immunostaining and characterized by the amplification of the *MDM2* gene, while the current case was negative for *MDM2* amplification (49).

The histology of malignant peripheral nerve sheath tumor (MPNST) and LGFMS can overlap significantly. However, our case was positive for MUC4 immunostaining but negative for S100 protein and without loss of H3K27me3, which are useful diagnostic biomarkers for MPNST (50). The identification of *FUS* rearrangement further demonstrated the diagnosis as LGFMS.

The current case also needed to be discriminated from SEF. However, classic SEF was composed of bland, monomorphic epithelioid cells, arranged in cords or nests within a densely collagenous stroma, while such morphologic changes were not identified in our case. Furthermore, the majority of SEFs harbor the *EWSR1::CREB3L1* fusion gene, while the present case exhibited *FUS::CREB3L2* fusion (51). Moreover, the recent study found that LGFMS and SEF could be classified by the different methylation profiles of the two entities (52).

Surgical resection is the main therapy for LGFMS, and our case showed no evidence of the disease after 11 months of the surgery. In 27 previously reported intrathoracic LGFMS cases, follow-up information was available in 16 cases who received surgical resection (including the current case), with a median follow-up of 23 months (range 6-110 months) (7, 22, 26, 30, 32-34, 38, 40-42). Three patients (3/16, 18.8%) developed recurrence, and none of the patients developed metastasis. The outcome results showed that 14 patients (87.5%) were alive without disease, 1 patient was alive with disease, and 1 patient died of cardiac failure. Long-term follow-up is still needed, as Evans et al. found that recurrence, metastasis rates, and mortality were 64%, 45%, and 42% of LGFMS with long-term follow-up (2). Additionally, LGFMSs with SEF-like areas and round-cell morphologic changes were reported to have more aggressive behavior (2). Four previous intrathoracic LGFMS cases had such morphologic change, and three of them had available follow-up information: one case was alive without disease, one case died of cardiac failure, and one case died of pericardial effusion and tamponade (22, 29, 37, 42). Hence, more cases are needed to verify the relation between the SEF-like and round-cell components with the behavior and prognosis of LGFMS cases.

In summary, we presented an extremely rare primary pleural LGFMS in a child with detailed clinicopathological and genetic results and carefully reviewed the literature on intrathoracic cases. To the best of our knowledge, the present case is the first genetically confirmed primary pleural LGFMS in a 4-year-old child with the identification of the *FUS::CREB3L2* fusion gene. Our study expands the age range of pleural LGFMS and highlights the use of immunohistochemical and molecular analyses in such challenging cases.

Data availability statement

The original contributions presented in the study are included in the article/supplementary material. Further inquiries can be directed to the corresponding author.

Ethics statement

Written informed consent was obtained from the minor(s)' legal guardian/next of kin for the publication of any potentially identifiable images or data included in this article.

Author contributions

XNH: Writing – original draft. WJ: Writing – original draft. XH: Writing – review & editing. MC: Writing – original draft. HZ: Writing – review & editing, Conceptualization, Funding acquisition, Project administration, Supervision.

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