

# Small intestinal SMARCA4-deficient undifferentiated carcinoma with intussusception: A case report

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**Abstract.** Switch/sucrose non-fermentable (SWI/SNF) related matrix associated actin-dependent regulator of chromatin sub-family A member 4-deficient undifferentiated carcinoma (SMARCA4-DUC) of the small intestine is a rare and highly aggressive subtype of gastrointestinal malignancy. The present study reported the case of SMARCA4-DUC in a patient who presented with intestinal intussusception and multi-organ metastasis to the lungs, liver, spleen and skin, which led to rapid progression and death in 43 days. The present case highlighted the unique histological and molecular features of SMARCA4-DUC, characterized by the absence of SMARCA4 expression and undifferentiated cellular morphology. The present case report aimed to describe the clinical presentation, imaging findings, pathological features and treatment course, to further the current understanding of SMARCA4-associated malignancies and improve the recognition of SMARCA4-DUC among clinicians and pathologists. Next-generation sequencing of the primary small intestinal tumor identified pathogenic mutations in PTEN, TP53, ataxia telangiectasia mutated, ephrin type-A receptor 5 and EGFR p.F997L, suggesting a DNA-repair-deficient genomic

background and providing potential targets for future precision therapy. Further research is warranted to determine the molecular mechanisms underlying SMARCA4-DUC, for the development of targeted therapeutic strategies and to improve the understanding of the aggressiveness of SMARCA4-DUC.

## Introduction

Gastrointestinal malignancies account for >25% of the global incidence of all tumors and 35% of tumor-related mortality, which poses a notable threat to the health of individuals worldwide (1,2). Based on histological features, gastrointestinal malignancies can be classified into adenocarcinoma, squamous cell carcinoma, sarcomatoid carcinoma and undifferentiated carcinoma (3,4). Among them, switch/sucrose non-fermentable (SWI/SNF) related matrix associated actin-dependent regulator of chromatin sub-family A member 4 (SMARCA4)-deficient undifferentiated carcinoma (DUC) of the gastrointestinal tract is a subtype of gastrointestinal undifferentiated carcinoma (4,5).

SMARCA4 deficiency is more commonly observed in non-small cell lung cancer, with an incidence of 5-10%. However, it is relatively rare in gastrointestinal cancer types, with limited reports available (6). The absence of SMARCA4 in primary tumors is associated with higher tumor invasiveness and a poorer prognosis. Compared with common tumors, there may be differences in clinical features, imaging manifestations, pathological characteristics and treatment responses (7,8). A previous study validated that in 436 cancer cell lines, tumors with low SMARCA4 expression demonstrated markedly higher IC<sub>50</sub> values for multiple chemotherapeutic drugs compared with those with high SMARCA4 expression, which implied that SMARCA4 deficiency may be a key reason for patient drug resistance (9). Another previous study indicated that SMARCA4 deficiency induces chemoresistance in tumor cells and also activates the EMT process, which thereby enhances tumor invasion (10). Therefore, additional studies and reports of cases of SMARCA4-deficient gastrointestinal undifferentiated carcinoma may enhance the current understanding of its biological behavior and potentially provide key insights for clinical diagnosis, the formulation of treatment strategies and prognosis assessment in the future.

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The present study aimed to report a rare case of small intestinal SMARCA4-DUC. The patient presented with intestinal intussusception and progression to metastasis in the lungs, liver, spleen and skin. The condition of the patient deteriorated rapidly, with death occurring 42 days following the initial detection of the lesion. The present study aimed to provide a detailed description of the clinical features, imaging manifestations, pathological characteristics and molecular biological features of the present case, to aid in the diagnosis and treatment of SMARCA4-deficient gastrointestinal undifferentiated carcinoma.

### Case report

A 69-year-old man with gastrointestinal discomfort was admitted to Weifang No. 2 People's Hospital (Weifang, China) on April, 2024 (day 1). The patient reported that they had suffered from chronic gastritis and gastric ulcers for >10 years. The mental health of the patient was poor at the time of admission, with dyspnea and no other abnormalities.

After the patient was admitted to the hospital, a biochemical examination was conducted (Table I). Results of the examination demonstrated carcinoembryonic antigen levels of 22.81 ng/ml (reference value, 0-4.5 ng/ml), indicative of a malignant neoplastic lesion. The medical team conducted a chest CT (Optima CT660, GE HealthCare; 5-mm-layer-thickness; helical scanning; Table II) with plain scanning and enhancement examination. Results of the CT examination revealed multiple nodules and masses in both lungs, with the largest mass located in the right lower lobe of the lung in the outer basal segment. The mass exhibited a diameter of ~31 mm, with an average CT value of 19 Hounsfield unit (HU), clear borders and smooth edges (Fig. 1A). There was no notable enhancement in the central area of the lesion on enhanced scanning (double-phase enhancement CT values of 20 HU and 21 HU, respectively), with circumferential enhancement around the edges (Fig. 1B). Furthermore, the lymph node in the right hilar region of the right lung was enlarged (Fig. 1C), with a maximal short diameter of ~15 mm. An abdominal CT scan (Optima CT660, Cytiva; 5-mm layer thickness; spiral scan) and enhanced examination revealed multiple hypodense nodules in the liver, with the largest nodule observed in the hepatic segment VIII (Fig. 2A). This nodule exhibited a diameter of ~12 mm, a CT value of 20 HU on scanning and enhanced scans of 30 HU, 35 HU and 39 HU in hepatic arterial, portal and delayed phases, respectively. A brain CT was also performed and indicated no evidence of metastasis, hemorrhage or infarction.

The clinical history of the patient, imaging and laboratory findings were suggestive of a malignant neoplastic lesion. On day 6, the patient underwent a puncture biopsy of a mass in the extramedullary segment of the lower lobe of the right lung. The puncture specimen was a greyish-white tissue of two strips with a length of 0.8-1.0 cm and a diameter of 0.1 cm. The puncture specimen was further examined using H&E staining. The tissue sections were fixed with 4% paraformaldehyde at room temperature for 30 min. The thickness of each section was 5  $\mu$ m. Hematoxylin staining was performed at room temperature for 7 min, followed by eosin staining at room temperature for 1 min. The sections were then observed under a light microscope. The

fresh specimen was dehydrated using alcohol in a room-temperature gradient, soaked in xylene at room temperature and embedded in paraffin at 65°C. Subsequently, the pathological tissue was cut into 5- $\mu$ m slices and soaked in room temperature xylene, anhydrous ethanol, 95% ethanol, 80% ethanol and H<sub>2</sub>O. The tissue sample was stained using hematoxylin for 5 min and eosin for 1 min, followed by gradient alcohol dehydration and neutral resin sealing. Tissue samples were observed using a light microscope (scale bar, 200  $\mu$ m). The tumor cells in the punctured tissue of the patient were diffusely distributed in the form of sheet nests. Furthermore, the tumor cells exhibited unclear borders, the cytoplasm was weakly basophilic or translucent, the nuclei were ovoid or short spindle-shaped, the frequent mitotic figures was irregular and small nucleoli were partially visible. Mitotic figures and a high amount of necrosis were observed (Fig. 3A). Histopathological staining of the patient tissue sample revealed negative expression of tumor markers for colorectal or lung cancer, though there is focal expression of squamous cell carcinoma markers. The tissue samples were fixed with 4% paraformaldehyde at room temperature for 30 min, and serial sections were cut at a thickness of 5  $\mu$ m. After blocking, the sections were incubated with the primary antibody at 37°C for 30 min, followed by incubation with the secondary antibody at room temperature for 8 min. To visualize the nuclei, the sections were treated with 3,3'-diaminobenzidine chromogenic solution for 8 min at room temperature and then counterstained with hematoxylin for 7 min at room temperature. Finally, the stained sections were examined under a light microscope. However, the specific origin of the tumor cannot be determined based solely on these findings. Therefore, clinical manifestations must be taken into account. The patient was admitted due to hematochezia and was found to have a positive fecal occult blood test. No pulmonary symptoms were observed. Imaging studies demonstrated tumors in the lungs and liver, which were peripherally located and well-defined, which suggested a higher possibility of metastatic tumors. Additionally, the mass causing intussusception had already invaded the muscular layer of the small intestine. Combining these clinical findings, it was concluded that the systemic tumors in the patient most likely originated from the small intestine.

Immunohistochemical staining was performed using Benchmark GX (Roche Diagnostics GmbH). The tissue samples were fixed with 4% paraformaldehyde at room temperature for 30 min, after which they were embedded in paraffin wax. Permeabilization was performed by treating the sections with 0.1% Triton X-100 at room temperature for 15 min. Each section was cut to a thickness of 5  $\mu$ m. EDTA (pH, 8) was used for antigen repair and repair parameters were set at 100°C for 30 min, according to the manufacturer's protocol. Endogenous catalase was blocked using 3% hydrogen peroxide for 5 min at room temperature. Tumor tissues were incubated with monoclonal antibodies against thyroid transcription factor-1 (TTF-1), NapsinA, cytokeratin (CK)5/6, synaptophysin (Syn), CK7, p40 and SMARCA4 at 37°C for 30 min. Following primary incubation, samples were incubated with the HRP-labeled secondary antibodies (Roche Diagnostics GmbH) for 8 min at room temperature. Subsequently, the nuclei were visualized using 3,3'-diaminobenzidine chromogenic solution for 8 min at room temperature and stained using hematoxylin for 7 min at room temperature (Table III).

Table I. General examination and laboratory test results on admission.

A, General clinical examination		
Parameter	Value	Reference range
Body temperature, °C	36.6	36.3-37.2
Heart rate, beats per min	82	60-100
Respiratory rate, breaths per min	18	15-20
Blood pressure, mmHg	130/82	90/60-120/80
B, ABG analysis		
Parameter	Value	Reference range
pCO <sub>2</sub> , mmHg	36.00	35-45
pO <sub>2</sub> , mmHg	86.00	80-100
HCO <sub>3</sub> , mmol/l	24.50	18-23
TCO <sub>2</sub> , mmol/l	25.60	22-28
SO <sub>2</sub> C, %	97.00	95-98
A-aDO <sub>2</sub> , mmHg	20.00	15
BE, mmol/l	0.30	-2-3
C, Tumor marker assays		
Parameter	Value	Reference range
CEA, µg/l	22.81	0-10
CA125, U/ml	28.62	0-35
CA19-9, U/ml	26.88	0-27
CA72-4, U/ml	2.97	0-6.9
AFP, ng/ml	1.85	0-20
D, Pathogen detection		
G test, pg/ml	Value	Reference range
G test, pg/ml	32.53	<60
CP-DNA test	Negative	Negative
MP-DNA test	Negative	Negative
GM ratio	0.21	<0.5
Fungal smear test	Negative	Negative

ABG, arterial blood gas; BE, base excess; CP, chloroplast; MP, *Mycoplasma pneumoniae*; G, 1,3-β-D-Glucan; GM test, Galactomannan test.

Results of the present study demonstrated that the tumor tissues obtained from the patient were negative for TTF-1, NapsinA, CK5/6 and Syn, positive for CK7, partially positive for p40 and negative for SMARCA4 (Fig. 3B-D; Table IV). The specimen was pathologically examined and used the 7th edition American Joint Committee on Cancer staging system for small bowel cancer (11). The tumor of the patient was stage IV, T2NXM1. The NX staging was due to insufficient

peritumoral lymph nodes sampled during surgery. However, with distant metastasis (M1), all T and N categories default to stage IV (12). Results of the pathological analysis demonstrated a SMARCA4-DUC with localized squamous carcinoma expression (Table IV). Given the low incidence of small intestine cancer and the lack of specific clinical guidelines, the guidelines for gastrointestinal tumors were referred to, which indicated that programmed cell death protein-1 (PD-1) therapy as the first-line treatment for advanced patients with SMARCA4-DUC (13). Additionally, previous studies have reported that SMARCA4 deficiency can lead to resistance to platinum-based and other traditional chemotherapeutic drugs (9,10,14). Therefore, PD-1 therapy was chosen as the primary treatment without employing a combination therapy regimen. Tislelizumab (200 mg, intravenous; cat. no. BGB-A317; BeOne Medicines) was intravenously administered to the patient in the clinic on day 11. Paclitaxel and nedaplatin were suggested for the patient on day 13 but not administered due to bleeding that occurred in the patient. The patient was passing tarry stools; thus, a follow-up blood test was performed. The results demonstrated hemoglobin levels of 57 g/l (healthy range, 130-175 g/l). Gastrointestinal bleeding was considered, chemotherapy was stopped and fasting, blood transfusion (4 units) and nutritional support were advised. However, the patient did not agree to the test because the results of the immunotherapy did not meet the expectations of the patient and they were not willing to continue with the immunotherapy.

On day 18, the patient had not passed gas for 1 day. A complete abdominal CT examination and enhanced examination (Optima CT660, Cytiva; 5-mm layer thickness; helical scan) revealed increased and enlarged liver metastases, novel metastases in the spleen, intestinal intussusception in the jejunal segment of the small intestine and intestinal obstruction (Fig. 2B-E). On day 20, a laparoscopic partial resection of the small intestine was performed and observed a resected small intestinal specimen of 8 cm in length in the lumen of the lassoed area. A red elevated area was observed in the intestinal lumen of the intestinal trocar, measuring 3.5x2.5x1.5 cm. This area exhibited an ulcerated vesiculated surface and the cut surface was grayish-white and grayish-red with a moderate solid texture. H&E staining was performed using the previously described protocol and light microscopy revealed that the tumor cells were diffusely distributed in nests of sheets. In addition, the tumor cells were large with unclear borders, weakly basophilic or translucent cytoplasm and large, rounded or oval nuclei with pronounced nuclear membranes. Multiple small nucleoli were observed and nuclear schizophrasia was readily visible. There were numerous hemorrhagic necroses within the lesion and small blood vessels in the tumor interstitium were hyperplastic and partially dilated with bruising (Fig. 4A). Immunohistochemistry demonstrated lack of expression for TTF-1, CK5/6, Syn, PSA, villin, CK20, caudal type homeobox 2 (CDX-2), CD34, Syn, CgA, insulinoma-associated protein 1 (INSM1), melanoma antigen (MelanA), S100, myogenin, leukocyte common antigen and CD30, and positive expression for CK7 and CK8. There were small foci of p40 positive expression, partial positive expression of glypican-3 and CD56, and lack of expression of SMARCA4 and nuclear protein in the testis (Fig. 4B-D; Table V). Small intestinal

Table II. Summary of patient radiological examination results.

Day	Radiological examination	Results
Day 1	Chest CT	Multiple nodules and masses found in both lungs, enlargement of right hilar lymph node in right lung
Day 18	Abdominal CT	Multiple low-density nodules in the liver, with larger nodules located in segment VIII of the liver
	Abdominal CT	Enlarged liver metastases, new metastases in the spleen, intussusception in the jejunal segment of the small intestine, intestinal obstruction
Day 31	Chest CT	Enlarged lesions in both lungs with new ground-glass shadows in the dorsal aspect of both lungs

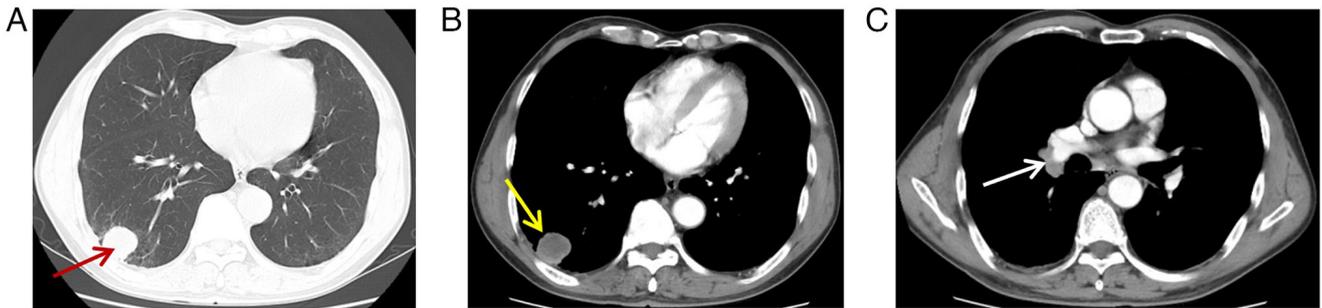


Figure 1. Thoracic CT examination (plain and enhanced) dated on day 1. (A) Lung window. A well-defined, smooth-edged mass measuring 31 mm in diameter was observed in the outer basal segment of the right lower lobe (red arrow). (B) Mediastinal window (enhanced). The central region of the mass in the outer basal segment of the right lower lobe revealed no enhancement, while the peripheral rim exhibited ring-like enhancement (yellow arrow). (C) Enlarged lymph node in the right hilar region (station 11), with a maximum short-axis diameter of ~15 mm (white arrow).

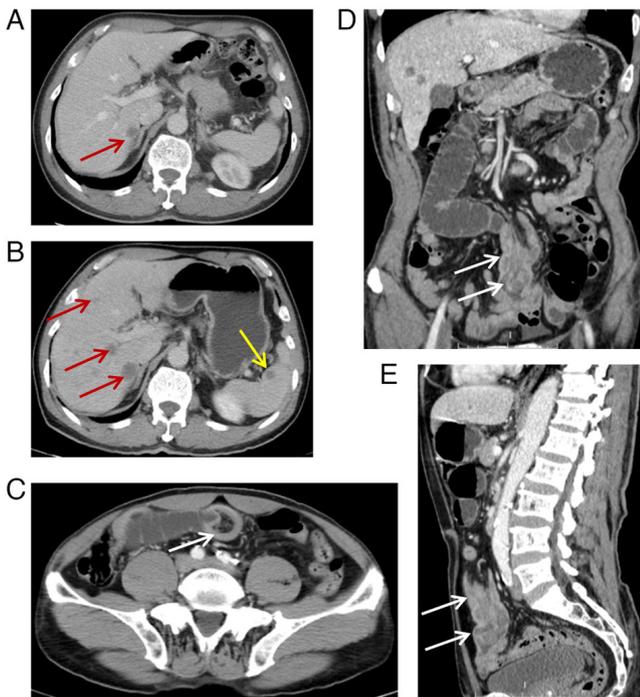


Figure 2. Plain and enhanced abdominal CT examination. (A) Multiple hypodense nodules in the liver, with the largest nodule located in segment VIII, measuring 12 mm in diameter with moderate enhancement (red arrow), on Day 1. (B) Increased and enlarged liver metastases (indicated by the red arrows), novel metastases in the spleen (indicated by the yellow arrows), on Day 18. (C) Transverse plane, (D) coronal plane and (E) sagittal plane views demonstrated localized intussusception in the small intestinal jejunum segment of the patient, with notable thickening and edema of the bowel wall (white arrows), on Day 18.

tumor samples obtained from the patient were investigated for 50 genes associated with lung cancer and 23 genes associated with intestinal cancer using next-generation sequencing (NGS). High-throughput sequencing was conducted on the MGISEQ-2000 platform. DNA libraries were constructed with the EGFR-KRAS-ALK Gene Mutation Combination Test kit (Cat. No. RM0914, Shenzhen Huada Gene Technology Co., Ltd.) to generate 150-bp paired-end reads. Library integrity was validated using the Qsep1™ capillary electrophoresis system (Product No. 100001, BiOptic Inc.) with the Standard Cartridge kit (Cat. No. C105201, BiOptic Inc.). Final library concentrations were quantified with the Equalbit 1x dsDNA HS Assay Kit (Cat. No. EQ121-02, Vazyme) and adjusted to 10 pM for cluster generation. Raw sequencing data were processed and analyzed using HALOS software (v4.1.3, Shenzhen Huada Gene Technology Co., Ltd.), and mutations in PTEN, TP53, ataxia telangiectasia mutated (ATM), ephrin type-A receptor 5 (EPHA5) and EGFR p.F997L were observed (Table SI). The raw sequence data reported in the present case report have been deposited in the Genome Sequence Archive (GSA) in the National Genomics Data Center, China National Center for Bioinformatics/Beijing Institute of Genomics, Chinese Academy of Sciences (GSA-Human database; accession no. HRA011751) that are publicly accessible at GSA for Human ([ngdc.cncb.ac.cn/gsa-human/browse/HRA011751](http://ngdc.cncb.ac.cn/gsa-human/browse/HRA011751)) (15,16). The genes tested include both those recommended for testing by Food and Drug Administration, National Medical Products Administration and National Comprehensive Cancer Network authoritative guidelines and key genes associated with drug targets during clinical

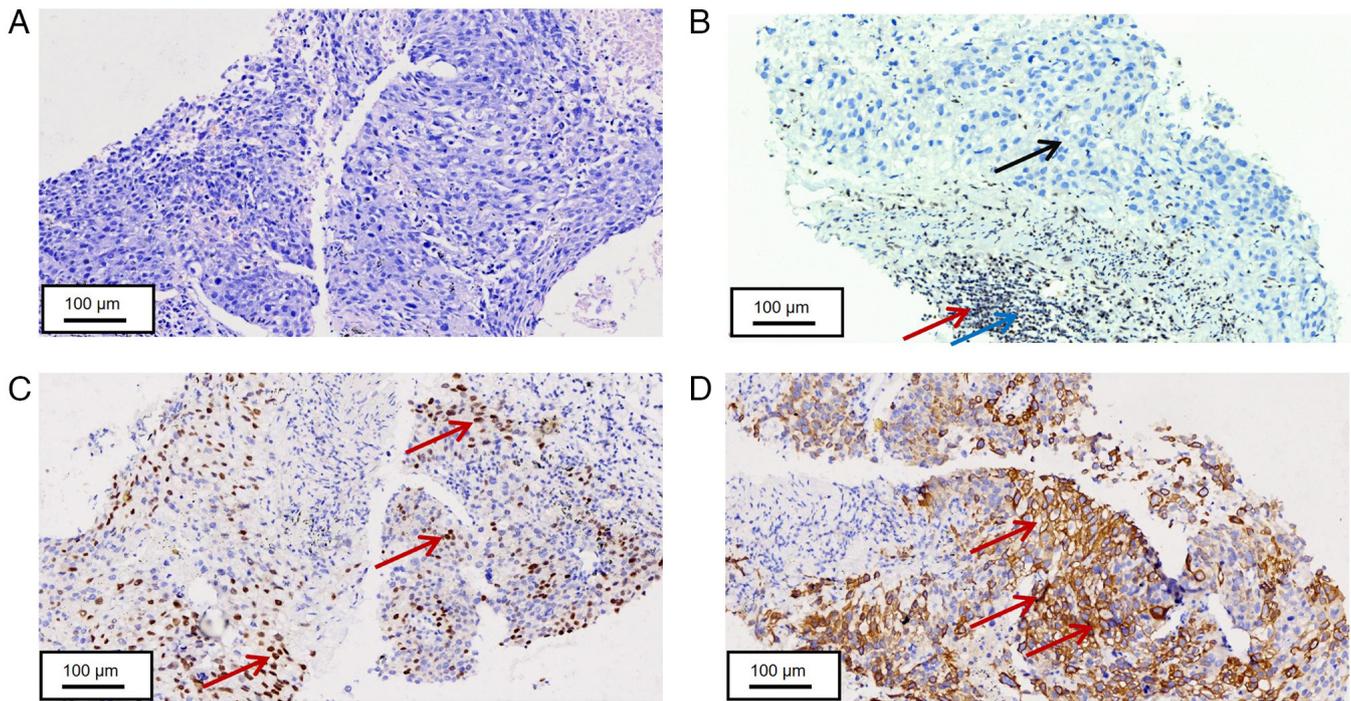


Figure 3. Histopathological staining of lung biopsy tissue. (A) Tumor cells in lung biopsy tissue were arranged in solid sheets, with some cells exhibiting visible nucleoli following H&E staining (scale bar, 100  $\mu$ m). (B) Tumor cells obtained from the lung biopsy tissue revealed negative expression of SMARCA4 (Red arrows indicate SMARCA4-positive sites, blue arrows indicate lymphoid regions, black arrows indicate tumor regions, scale bar, 100  $\mu$ m). (C) Partial positive expression of p40 in tumor cells obtained from the lung biopsy tissue (Red arrows indicate p40-positive sites, scale bar, 100  $\mu$ m). (D) Positive expression of CK7 in tumor cells obtained from the lung biopsy tissue (red arrows indicate CK7-positive sites, scale bar, 100  $\mu$ m). SMARCA4, switch/sucrose non-fermentable (SWI/SNF) related matrix associated actin-dependent regulator of chromatin sub-family A member 4; CK, cytokeratin.

trials and the development of diseases (17-27). Pan-cancer studies have linked SMARCA4/PTEN loss to DNA repair defects that may underlie the TP53, ATM, EPHA5 and EGFR mutations, as unrepaired DNA damage can trigger oncogenic mutations (28-30). Thus, the clinical features, imaging manifestations and pathological features were indicative of undifferentiated carcinoma with localized squamous carcinoma expression and intussusception in SMARCA4-deficient small intestine, with metastasis to the lung, liver and spleen. Although these mutations were detected, the treatment plan could not be adjusted and optimized based on them due to the rapid progression of the condition of the patient.

On day 31, a CT examination of the chest was performed. The patient exhibited enlarged and increased lesions in both lungs (Fig. 5A-C) and new ground-glass shadows in the dorsal aspect of both lungs, with no evidence of infection, the Response Evaluation Criteria in Solid Tumors (RECIST) score was disease progression (31). Furthermore, immune-associated pneumonia-non-specific interstitial pneumonitis was considered (Fig. 5B) (32). Multiple new lesions were observed on the skin and gingiva, indicative of metastasis (Fig. 6A-C). The patient presented with persistent symptoms of gastrointestinal bleeding and a decrease in hemoglobin levels (69 g/l) and was therefore administered 9 units of erythrocyte transfusions and posterior pituitary hormones (a dilute acetic acid sterilized aqueous solution of posterior pituitary powder containing posterior pituitary hormone) for hemostasis (2 ml intravenously once daily), and was provided with nutritional support. The family of the patient requested that the patient be discharged from the hospital on day 42 and the patient was discharged on

the same day. Within 24 h of discharge, the patient succumbed to the disease. The complete timeline is shown in Fig. 7.

## Discussion

The SMARCA4 gene subgroup belongs to the SWI/SNF family, which serve a key role in chromatin remodeling and repair. Results from previous studies demonstrated that SMARCA4 serves a key role in developmental processes, transcriptional regulation, DNA repair, cell cycle regulation, tumorigenesis and tumor development. Furthermore, inactivating mutations in SMARCA4 lead to a loss of protein expression in the nucleus, which result in malignancies associated with SMARCA defects (33,34). In non-small cell lung cancer, mutations in the SMARCA4 gene occur in ~10% of cases. Notably, these mutations have been reported in malignant tumors in alternative sites, including the ovaries, esophagus, stomach, uterus and liver; however, the majority of previous studies were case reports (35-39).

SMARCA4-DUC is a rare yet highly malignant tumor that is aggressive and associated with poor prognosis (Table VI). Cases of SMARCA4-DUC in the gastrointestinal tract predominantly affects elderly men and the common sites of development include the stomach, colon, small intestine and lower esophageal cardia, with few cases affecting the ileum, rectum and pancreas (40,41). In a study by Wang *et al* (42), it was noted that lung cancer cells undergo SMARCA4 loss, which results in a reduced response to anti-PD-1 immunotherapy, which may be associated with a notable reduction in the infiltration of dendritic cells and CD4<sup>+</sup> T cells into

Table III. Sources of reagents for patient testing.

Reagent or medicine	Manufacturer	Cat. no.	Dilution
EGFR-KRAS-ALK gene mutation combination test kit	Shenzhen Huada Gene Technology Co Ltd	RM0914	Not applicable
Anti-Rabbit IgG H&L (HRP)	Roche Diagnostics GmbH	760-500	1:1
Anti-human NUT	Guangzhou Anbiping Medical Technology	IR429	1:1
Anti-human TTF-1	Fuzhou Maixin Biotechnology Development Co., Ltd.	MAB-0677	1:1
Anti-human CK5/6	Fuzhou Maixin Biotechnology Development Co., Ltd.	MAB-0774	1:1
Anti-human Syn	Fuzhou Maixin Biotechnology Development Co., Ltd.	MAB-0742	1:1
Anti-human Villin	Fuzhou Maixin Biotechnology Development Co., Ltd.	MAB-0540	1:1
Anti-human CK20	Fuzhou Maixin Biotechnology Development Co., Ltd.	MAB-0834	1:1
Anti-human CDX2	Fuzhou Maixin Biotechnology Development Co., Ltd.	MAB-1056	1:1
Anti-human CD34	Fuzhou Maixin Biotechnology Development Co., Ltd.	MAB-1076	1:1
Anti-human CgA	Fuzhou Maixin Biotechnology Development Co., Ltd.	MAB-0707	1:1
Anti-human INSMI	Fuzhou Maixin Biotechnology Development Co., Ltd.	MAB-1017	1:1
Anti-human MelanA	Fuzhou Maixin Biotechnology Development Co., Ltd.	MAB-1033	1:1
Anti-human S-100	Fuzhou Maixin Biotechnology Development Co., Ltd.	Kit-0007	1:1
Anti-human Myogenin	Fuzhou Maixin Biotechnology Development Co., Ltd.	MAB-0362	1:1
Anti-human CD30	Fuzhou Maixin Biotechnology Development Co., Ltd.	MAB-0023	1:1
Anti-human CK7	Fuzhou Maixin Biotechnology Development Co., Ltd.	MAB-0828	1:1
Anti-human CK8	Fuzhou Maixin Biotechnology Development Co., Ltd.	MAB-1002	1:1
Anti-human p40	Fuzhou Maixin Biotechnology Development Co., Ltd.	RMA-0815	1:1
Anti-human CD56	Fuzhou Maixin Biotechnology Development Co., Ltd.	MAB-0743	1:1
Anti-human SMARCA4	Fuzhou Maixin Biotechnology Development Co., Ltd.	MAB-1016	1:1
Anti-human NapsinA	Fuzhou Maixin Biotechnology Development Co., Ltd.	MAB-0704	1:1

NUT, nuclear protein in testis; TTF-1, thyroid transcription factor-1; Syn, synaptophysin; CK; CDX2, caudal type homeobox 2; INSMI, insulinoma-associated protein 1; MelanA, melanoma antigen; SMARCA4; switch/sucrose non-fermentable-related matrix associated actin-dependent regulator of chromatin sub-family A member 4.

the tumor microenvironment (TME); however, this study focused on cellular and animal experiments. The results of previous studies examining the impact of SMARCA4 loss on the response to PD-1 immunotherapy in lung cancer are conflicting and these discrepancies may be associated with differences in study models. As noted in a study of advanced undifferentiated carcinoma of the duodenum, PD-1 therapy is effective in some patients with SMARCA4 deficiency, which potentially provides an option for the treatment of these highly malignant tumors (43). A clinical study reported that partial SMARCA4 deletion may make patients with lung cancer more sensitive to PD-1 monoclonal antibody drugs (27). However, this study primarily included an animal model for its experiments and the underlying mechanisms may be more complex in humans. Tumors are often located within the muscular wall of the digestive tract and involve the mucosa and may also form masses outside of the wall of the digestive tract (38,44-47). Few characteristic imaging features of SMARCA4-DUC lead to complexities in diagnosis; however, the majority of these tumors exhibit notable signs of malignancy, such as large foci that are lobulated, which encircle blood vessels, invade adjacent tissues and metastasis to mediastinal lymph nodes and distant organs (5). In the present case, a chest CT examination demonstrated multiple nodules and masses with clear borders and smooth edges and enhanced CT examination demonstrated

that the majority of areas in the lesions did not exhibit notable enhancement. Furthermore, the edges of the lesions were notably enhanced, which was consistent with the manifestation of bloodstream metastasis in both lungs. Enhanced abdominal CT examination revealed localized condensation of the jejunal segment of the small intestine, with edema and thickening of the tube wall. In addition, multiple nodules in the liver and spleen were observed, with moderate bulls-eye enhancement observed through enhanced examination. These findings were consistent with the manifestation of hematogenous metastasis in the liver and spleen. The imaging manifestations of bilateral lung, liver and spleen metastases were consistent with those reported in previous literature (48-50); however, to the best of our knowledge, the present study is the first to report the case of intestinal intussusception caused by SMARCA4-DUC in the small intestine.

The histology of SMARCA4-DUC is characterized by rounded cells and epithelioid cell structures arranged in sheets, nests or beams within the tumor tissue, which may be accompanied by a loss of adherent growth. Furthermore, rhabdoid morphology, which is defined as the presence of cells with large, eosinophilic cytoplasm, prominent nucleoli and a vesicular chromatin pattern is seen in focal areas in ~2/3 of the cases (34). Results from previous studies revealed that tumor cells are often uniform. However, tumor cells that

Table IV. Summary of patient IHC results.

Name	pH	Incubation time, min	IHC result
NUT	8	32	-
TTF-1	8	32	-
CD5/6	8	32	-
Syn	8	32	-
Villin	8	32	-
CDX2	8	32	-
CD34	8	32	-
CgA	8	32	-
INSM1	8	32	-
Myogenin	8	32	-
CD30	8	32	-
CK7	8	32	+
p40	8	32	+
CD56	8	32	+
SMARCA4	8	32	-
NapsinA	8	32	-
MelanA	8	28	-
S-100	8	24	-
CK8	8	24	+
CD20	8	24	-

IHC, immunohistochemistry; -, negative; +, positive; NUT, nuclear protein in testis; TTF-1, thyroid transcription factor-1; Syn, synaptophysin; CK, cytokeratin; CDX2, caudal type homeobox 2; INSM1, insulinoma-associated protein 1; MelanA, melanoma antigen; SMARCA4; switch/sucrose non-fermentable (SWI/SNF) related matrix associated actin-dependent regulator of chromatin sub-family A member 4.

were pleomorphic, contained spindle cell components or exhibited pseudoadenoids or pseudoglandular structures were observed in some cases (5,51). A previous study demonstrated that undifferentiated carcinomas of the gastrointestinal tract often possess multifarious rhabdomyolysis features, accompanied by deletion of SMARCB1, SMARCA2, SMARCA4 and AT-rich interaction domain-containing protein 1A (ARID1A) (41). Notably, a deletion in SMARCA4 is often present in conjunction with a deletion in SMARCA2, which affects ~39.9% of cases. These deletions are often accompanied by negative or reduced expression (<10% positivity) of epithelial markers, including CKpan, CK7, CK20 and CDX2. In addition to epithelial markers, tumor cell markers, such as Sox2, spalt-like protein 4, p53, ARID1A and neuroendocrine markers, such as CgA, Syn, CD56 and INSM1, may also be expressed to varying degrees in SMARCA4-DUC (46,52-54). In a previous study (36), it was noted that SMARCA4 deletion and TP53 have similar mutation rates in gastrointestinal tumors and that this mutation seems to occur more often in the esophagus compared with the stomach or elsewhere, which suggests a possible location of origin for gastrointestinal tumors. The patient from the present case also demonstrated focal expression of p40; however, SMARCA4 deletion was detected in small bowel cancer in a study on SMARCA4

Table V. Immunohistochemistry of tumor tissues from different sites in patients.

	Lung tumor	Small bowel tumor
CK5/6	-	-
p40	+	+
CK7	+	+
TTF-1	-	-
NapsinA	-	Untested
Syn	-	Untested
CK8	Untested	+
Villin	Untested	-

-, negative; +, positive; TTF-1, thyroid transcription factor-1; Syn, synaptophysin; CK, cytokeratin.

deletion in lung cancer, which indicated that tumors with deletion of SMARCA4 were mostly non-squamous cell carcinomas and the expression of p40 was mostly negative or weakly positive at the focal level (34). This is similar to the expression of p40 detected in the patient in the present case. Considering the DNA repair function of SMARCA4, SMARCA4 deletion may lead to a variety of morphological and immunophenotypic features of tumor cells during differentiation and the focal expression of p40 may reflect the partial squamous differentiation tendency of the associated tumor cells (55). According to previous studies, diffuse strong positivity of p40 expression in tumor tissue can be used to diagnose squamous cell carcinoma. However, the present tumor tissue showed focal p40 expression, which did not meet the diagnostic criteria. Therefore, the patient's treatment plan was based on the criteria for undifferentiated carcinoma (56,57).

During clinical diagnosis, a combination of histological and molecular pathological analyses is required to distinguish SMARCA4-DUC from large-cell neuroendocrine carcinomas with rhabdomyosiform morphology, often accompanied by diffuse or strong positive expression of neuroendocrine markers (58,59). Notably, melanomas with rhabdomyosiform morphology often express melanocyte markers, such as S100, MelanA and human melanoma black-45 (60,61). Rhabdomyosarcoma with distinctive rhabdomyosarcoma-like features often express muscle-specific markers, such as myogenic differentiation antigen 1 and myogenin (62,63). Cases of primary interstitial large-cell lymphoma of the gastrointestinal tract often include large cells with a pronounced nucleolus and these may exhibit high levels of expression of lymphocyte markers, such as CD30 and ALK (64,65). Furthermore, proximal-type epithelioid sarcoma with epithelioid morphology often express epithelioid markers, such as CK and epithelial membrane antigen (66,67), which aid in identification and differentiation.

Genomic analysis of SMARCA4-DUC demonstrated that common co-mutations included TP53, KRAS, serine/threonine kinase 11 (STK11) and Kelch-like ECH-associated protein 1 (KEAP1), with less common co-mutations with common genes, such as EGFR, ROS proto-oncogene 1, receptor tyrosine kinase

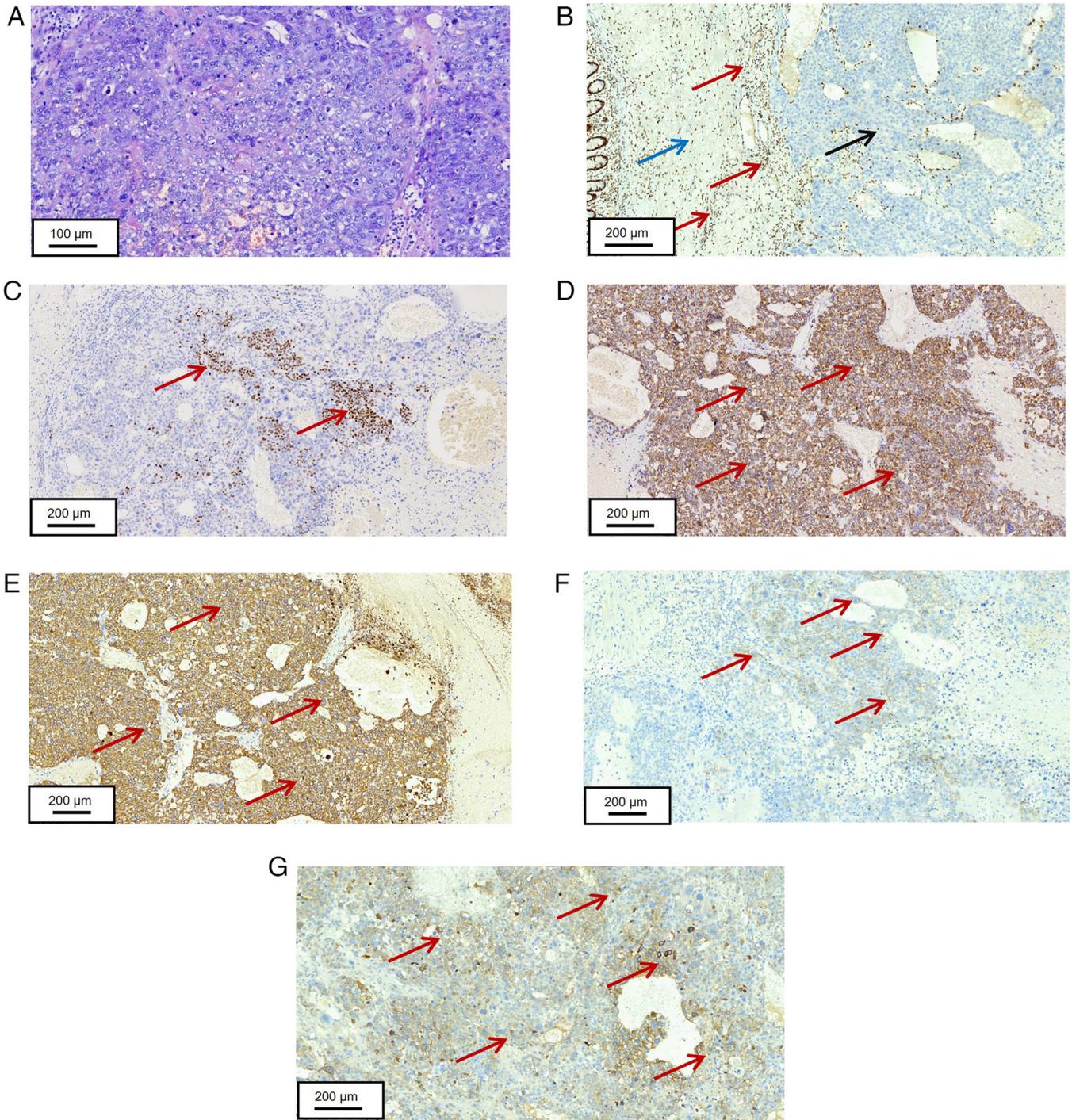


Figure 4. Histopathological staining of tumor tissue in the small intestine. (A) Tumor cells in the small intestine were arranged in solid nests, with large, lightly stained nuclei and one or more visible small nucleoli (scale bar, 100  $\mu\text{m}$ ). (B) Negative SMARCA4 expression in tumor cells of the small intestine (red arrows indicate SMARCA4-positive sites, blue arrows indicate lymphoid regions, black arrows indicate tumor regions, scale bar, 200  $\mu\text{m}$ ). (C) Partial positive expression of p40 in tumor cells of the small intestine (red arrows indicate p40-positive sites, scale bar, 200  $\mu\text{m}$ ). (D) Positive expression of CK7 in tumor cells of the small intestine (Red arrows indicate CK7-positive sites, scale bar, 200  $\mu\text{m}$ ). (E) Positive expression of CK8 in tumor cells of lung biopsy tissue (Red arrows indicate CK8-positive sites, scale bar, 200  $\mu\text{m}$ ). (F) Positive expression of CD56 in tumor cells of lung biopsy tissue (red arrows indicate CD56-positive sites, scale bar, 200  $\mu\text{m}$ ). (G) Positive expression of GPC-3 in tumor cells of lung biopsy tissue (red arrows indicate GPC-3-positive sites, scale bar, 200  $\mu\text{m}$ ). SMARCA4, switch/sucrose non-fermentable (SWI/SNF) related matrix associated actin-dependent regulator of chromatin sub-family A member 4; CK, cytokeratin; GPC-3, glypican-3.

and ALK (68,69). The Dana-Farber Cancer Institute (Boston, USA) reported that the KRAS gene is the most commonly co-mutated gene in SMARCA4-DUC, which accounts for 5-11% of gene mutations (70). Patients with SMARCA4-DUC often present with a poor prognosis, with markedly shorter median

progression-free survival (4.1 vs. 1.4 months) and median overall survival (15.1 vs. 3.0 months) compared with patients exhibiting the SMARCA4/KRAS wild-type (71,72).

Treatment of SMARCA4-DUC is complex, due to its unique physiological characteristics and low prevalence rates. In

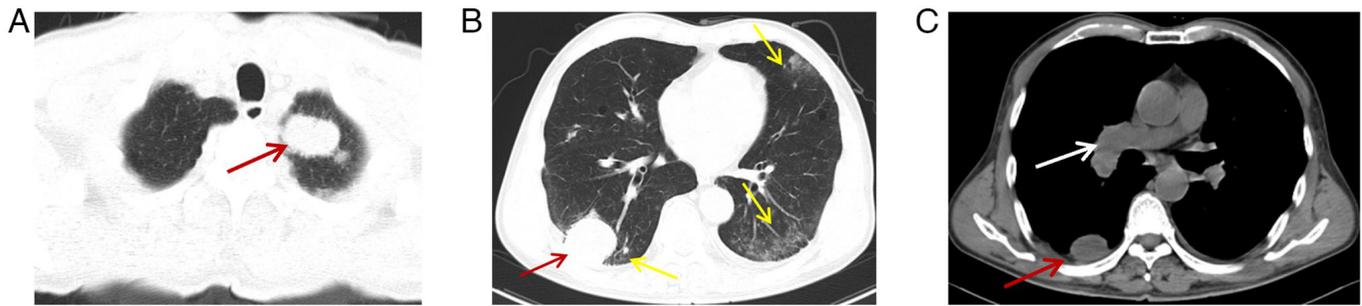


Figure 5. Thoracic CT examination (plain) on Day 31. (A) Lung window. Multiple nodules and masses in the left upper lobe, indicative of novel lesions compared with the thoracic CT examination (indicated by the red arrow), dated on Day 31. (B) Lung window. Mass in the extra-basal segment of the lower lobe of the right lung, enlarged on contrast with chest CT examination, on Day 31 (red arrow). Subpleural ground-glass shadows in both lungs were indicative of a novel lesion consistent with the CT pattern of immune-associated pneumonia-non-specific interstitial pneumonitis (yellow arrow). (C) Mediastinal window on Day 31 (red arrow). Enlarged lymph nodes in region 11 of the right hilar (white arrow).

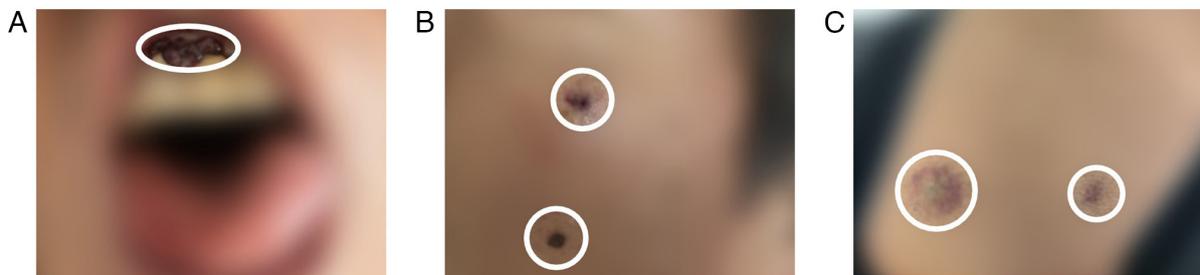


Figure 6. Multiple cutaneous metastases. Images of multiple cutaneous metastases in the (A) gingiva, (B) left face and (C) anterior chest (white circles).

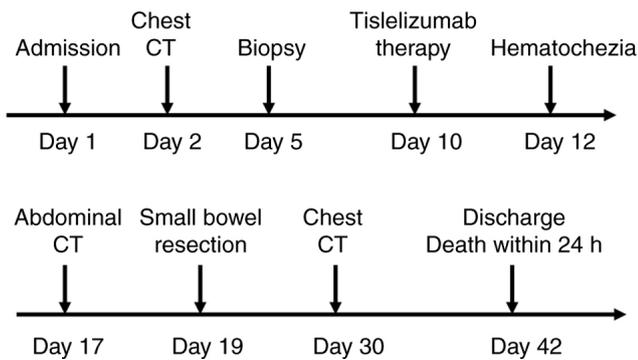


Figure 7. Timeline of major tests and interventions received by the patient from admission to discharge.

addition, there are no unified treatment guidelines to follow, with a lack of effective therapies used in clinical practice (73). Partial SMARCA4 deletion may make patients with lung cancer more sensitive to PD-1 monoclonal antibody drugs, which was noted in a clinical study (44). A previous case report demonstrated that surgical resection may be an effective initial treatment option for patients with SMARCA4-DUC with clinical staging (74). However, the present case report also demonstrated that 4 months following surgery, patients with SMARCA4-DUC developed tumor recurrence with invasion to the adjacent ribs. Results from a previous study revealed that patients with SMARCA4-DUC may benefit from platinum-based chemotherapy and the systemic effects of chemotherapy may control distal tumor metastasis to some extent, due to the high aggressiveness and metastatic

nature of the disease (75). Furthermore, targeted drug therapies, including anilotinib, sindilizumab and bevacizumab, may aid in controlling the progression of SMARCA4-DUC (54,76). In a previous case report, a patient with SMARCA4-DUC who developed genetic mutations in STK11/KEAP1, KEAP1, PBRM1, SMARCA4 and STK11 received tirilizumab immunotherapy (77). However, rapid disease progression was observed, with hepatic and pulmonary metastases within 1 month, which ultimately led to the death of the patient. Findings from previous studies revealed that STK11/KEAP1, KEAP1, PBRM1 and STK11 gene mutations may affect the tumor microenvironment of patients, which weakens the tumor-killing effect of immune cells and increases the risk of disease progression following immunotherapy (78-80). Due to the high degree of heterogeneity of tumor cells in patients with SMARCA4-DUC, NGS technology is required for the development of personalized treatment plans (81). The rapid disease progression of the present patient after immunotherapy suggested that there may be hyperprogression, although there is no direct evidence that SMARCA4 may be an independent influence on hyperprogression of immunotherapy in gastrointestinal patients. It was noted in Han *et al* (82) that environmental factors, especially the gut microbiome, can have an impact on immunotherapy and it is possible that the intussusception and gastrointestinal bleeding that the patient suffered at a later stage would have led to an imbalance in the gut microbiome, which could be an important cause of hyperprogression.

In addition, the present patient tumor exhibited partial squamous carcinoma and in a case report by Misra *et al* (83), it was noted that patients with SMARCA4 in the gastrointestinal

Table VI. Summary of key findings on SMARCA4 loss in various malignancies.

First author, year	Type of cancer	Key finding	(Refs.)
Witkowski <i>et al</i> , 2023	Ovarian	Associated with tumor aggressiveness and poor prognosis.	(35)
Yu <i>et al</i> , 2024	Esophageal	Led to reduced sensitivity to chemotherapeutic drugs.	(55)
Yu <i>et al</i> , 2024	Gastric	Associated with tumor aggressiveness and poor prognosis.	(55)
Affandi <i>et al</i> , 2018	Endometrial	Associated with tumor aggressiveness and poor prognosis.	(57)
Garcia-Porrero <i>et al</i> , 2024	Liver	Associated with tumor aggressiveness and poor prognosis.	(58)
Xue <i>et al</i> , 2021;	Non-small cell	Reduced sensitivity to chemotherapeutic drugs; activated	(9)
Kim <i>et al</i> , 2021	lung cancer	EMT process and enhanced tumor invasiveness	(10)

EMT, epithelial-mesenchymal transition.

tract exhibiting focal squamous carcinoma differentiation can be resistant to a variety of chemotherapeutic agents (cisplatin and docetaxel) and lead to an increase in their invasiveness, which may also explain the rapid progression of the patient in the present case (84,85). Due to the highly heterogeneous nature of tumor cells in SMARCA4-DUC patients, deep gene sequencing of SMARCA4-DUC patients using NGS technology is of great importance for the development of personalized treatment protocols, taking into account the impact of different gene mutations on patient prognosis and the toxicity and side effects of traditional chemotherapeutic agents (81,86). In addition, some non-pharmacological interventions, such as magnetic fields, exosomes, may also provide options for the treatment of SMARCA4-DUC (87-89). In the present case, the patient was administered tislelizumab once following diagnosis and the disease was reassessed at 22-day intervals. After 1 month of treatment, the symptoms of the patient had worsened, with an increase in the number of metastatic lesions in the lungs and liver, enlargement, novel metastases in the spleen, the development of a CT pattern indicative of immune-associated pneumonia-nonspecific interstitial pneumonitis in both lungs and a Response Evaluation Criteria in Solid Tumors score of progressive disease (31). Following intussusception surgery, gastrointestinal bleeding and treatment for aggressive symptoms, the death of the patient occurred after 12 days. Although the specific cause of death was not determined via autopsy, it was considered that the highly aggressive nature of SMARCA4-DUC, which led to multiple organ metastasis was the main cause of death and the gastrointestinal hemorrhage and immune-associated pneumonia caused by the intussusception may have accelerated the deterioration of the disease.

The present study exhibits numerous limitations. For example, the present study included a retrospective analysis of rare cases, with a lack of guideline support and inexperience. In the present case, diagnosis and treatment were delayed, as it remained unclear whether immunotherapy caused hyperprogression of the disease. In addition, the specific cause of death in the patient was not determined due to the absence of autopsy confirmation.

In conclusion, the present study reported a rare case of SMARCA4-DUC in the gastrointestinal tract, which provides a theoretical basis for the clinical management of this disease. SMARCA4-DUC often affects elderly male patients and imaging often reveals large lesions, invasion of surrounding

tissues and metastasis to lymph nodes and distant organs. Definitive diagnosis of the disease relies on pathological examination, immunohistochemistry and molecular testing. A diagnosis of SMARCA4-DUC should be considered following the observation of solid high-grade undifferentiated morphology of cells and negative expression of genes used for conventional staging. Thus, appropriate immunohistochemical and molecular tests should be carried out to aid in early diagnosis and effective treatment.

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#### Availability of data and materials

The data generated in the present study may be requested from the corresponding author. The raw sequence data generated in the present study may be found in the Genome Sequence Archive (GSA) (14) in National Genomics Data Center (15), China National Center for Bioinformatics/Beijing Institute of Genomics, Chinese Academy of Sciences, under accession number (accession no. HRA011751) or at the following URL: [ngdc.cncb.ac.cn/gsa-human/browse/HRA011751](http://ngdc.cncb.ac.cn/gsa-human/browse/HRA011751).

#### Authors' contributions

MQ, LL, BY, and YZ analyzed data. BY and XY designed the study protocol. XC and SL were the primary care physicians of the patient and developed and implemented the treatment plan. YW, QG and ZC performed the literature review and analyzed and interpreted the data in the paper. XY drafted the work and critically revised intellectual content. BY and XY confirm the authenticity of all the raw data. All authors commented on the manuscript and agreed with the conclusions of the manuscript. All authors read and approved the final manuscript.

### Ethics approval and consent to participate

The present case report was approved by the Ethics Committee of the Weifang No. 2 People's Hospital (approval no. KY2023-055-01; Weifang, China) and was performed in accordance with the Declaration of Helsinki.

### Patient consent for publication

The patient's family member provided written informed consent for the publication of the manuscript including any identifying images or data.

### Competing interests

The authors declare that they have no competing interests.

### Use of artificial intelligence tools

During the preparation of this work, artificial intelligence tools were used to improve the readability and language of the manuscript or to generate images and subsequently, the authors revised and edited the content produced by the artificial intelligence tools as necessary, taking full responsibility for the ultimate content of the present manuscript.

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