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Gastroblastoma mimics the embryonic mesenchyme of the foregut: a case report

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Abstract

Background Gastroblastoma is a rare gastric tumor composed of epithelial and spindle cell components. The characteristic *MALAT–GL11* fusion gene has only been identified in 5 reported cases. We report the morphological characterization of gastroblastoma with the *MALAT1–GL11* fusion gene in a young Japanese woman.

Case presentation A 29-year-old Japanese woman visited Iwate Medical University Hospital with upper abdominal pain. Computed tomography revealed a tumor in expansive lesions involving the gastric antrum. Histologically, we observed a biphasic morphology composed of epithelial and spindle cell components. The epithelial components appeared as slit-like glandular structures with tubular or rosette-like differentiation. The spindle cell components consisted of short spindle-shaped oval cells. Immunohistochemical (IHC) analysis revealed that the spindle cell component was positive for vimentin, CD10, CD56, GL11, and HDAC2, and focally positive for PD-L1. The epithelial component was positive for CK AE1/AE3, CAM5.2, and CK7, and negative for CK20 and EMA. Both components were negative for KIT, CD34, DOG1, SMA, desmin, S100 protein, chromogranin A, synaptophysin, CDX2, and SS18-SSX. The *MALAT-GL11* fusion gene was detected molecularly.

Conclusions We report the following new findings with this case: (i) gastric tumors mimic the gastrointestinal mesenchyme in the embryonic period; (ii) nuclear expression of PD-L1 and HDAC2 were observed in the spindle cell component of a gastroblastoma. We speculate that histone deacetylase (HDAC) inhibitors may offer a promising treatment option for gastroblastoma.

Keywords Gastroblastoma, MALAT1-GLI fusion gene, PD-L1

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Background

Gastroblastoma, first described by Miettinen and colleagues in 2009, is a rare gastric tumor characterized by epithelial and spindle cell components [1]. To our knowledge, 16 cases have been reported in the medical literature [1–11]. However, the characteristic MALAT1–GLI1 fusion gene was identified in only 5 of those cases. Due to the rarity of this disease, its pathogenesis remains unknown. We report a case of gastroblastoma containing a MALAT1–GLI1 fusion gene in a young Japanese woman.

Case presentation

A 29-year-old Japanese woman visited Iwate Medical University Hospital with upper abdominal pain for over a week. The patient had no surgical history, drug allergies, or family history of malignancy. Her laboratory examination was unremarkable. A contrast-enhanced computed tomography (CT) scan showed a 70-mm antral expansive mass (Figs. 1A, B). A biopsy of the tumor was not performed, and she was diagnosed clinically with a gastrointestinal stromal tumor (GIST). Although a preoperative histological diagnosis was not made, a laparoscopic partial gastrectomy was performed. Eight months have passed since the surgery; however, no recurrence or metastasis has been found.

Grossly, the surgical specimen was revealed to be a nodular and well-circumscribed mass measuring 7 x 7 x 6 cm in the gastric antrum. The tumor grew as an expansive mass and involved the gastric wall structures. The cut surface showed a greyish-yellow and tan solid



Fig. 1 A 29-year-old Japanese woman with gastroblastoma. **A** Axial plane enhanced computed tomography (CT) image of the abdomen. The gastroblastoma appears as a well-circumscribed mass (arrows) showing a multi-cystic, heterogeneously enhanced mass in the lower part of the stomach; **B** Coronal plane enhanced CT image of the abdomen. Multi-cystic nature and heterogeneity of the gastroblastoma (arrows) arising from the bottom of the gastric wall in the upper-left quadrant



Fig. 2 Cut surface and histology of the gastroblastoma. A Cut surface of the gastroblastoma. B Most of the tumor cells were spindle cells, which appeared oval-shaped without atypia (x200). C Tubular or rosette-like differentiation (x200). D Glandular and slit-like structure (x100)

mass with cystic and hemorrhagic components (Fig. 2-A). Histologically, a biphasic morphology of epithelial and spindle cell components was observed. The epithelial components were slit-like glandular structures composed of low cuboidal epithelium with tubular or rosette-like differentiation with eosinophilic secretions. The spindle cell components consisted of short spindle-shaped oval cells infiltrating the smooth muscle. These cells had small round nucleoli and well-defined cell borders (Figs. 2-B, C, D). Mitotic activity was low in both components. Lymph-vascular invasion was not found. Immunohistochemical (IHC) analysis revealed that the spindle cell component was positive for vimentin, CD10 (Fig. 3-A), CD56 (Fig. 3-B), and glioma-associated oncogene homolog 1 (GLI1) (Fig. 3-C), focally positive for PD-L1 (Fig. 3-D), and histone deacetylase 2 (HDAC2) (Fig. 3-E). The epithelial component was positive for pan-cytokeratin (CK AE1/AE3), CAM5.2, and CK7, but negative for CK20 and epithelial membrane antigen (EMA). Both components were negative for KIT (Fig. 3-F), CD34 (Fig. 3-G), DOG1, smooth muscle actin (SMA), desmin, S100 protein, chromogranin A, synaptophysin, CDX2, and SS18-SSX (Fig. 3-H). Antibodies used for the IHC analysis are shown in Table 1, while results of the analysis are shown in Table 2.

We performed reverse transcriptase-polymerase chain reaction (RT-PCR) analysis [6], which revealed that the tumor harbored the *MALAT1-GLI1* fusion gene (Fig. 4-A, B). In addition, we developed a customized next-generation sequencing (NGS) gene panel for use in this case. The panel consisted of 28 genes (*APC, TP53, CDKN2A, MET, ATM, MLH-1, PMS2, HRAS, AXIN2, BAX, DCC, MSH2, POLE, RNF43, PTEN, BRAF, EPCAM, MSH6, BUB1B, RhoA, KRAS, NRAS, SMAD4, CDK4, PIK3CA, STK11, TGFBR2, and EGFR*) for exploring the genetic causes of colorectal cancer. This panel was employed for gastroblastoma in the present case to detect gene mutations. However,



Fig. 3 Immunohistochemical findings of the gastroblastoma. A Expression of CD10 by tumor cells (x100). B Expression of CD56 by tumor cells (x100). C Expression of GL11 by tumor cells (x100). D Expression of PD-L1 by tumor cells (x200). E Expression of HDAC2 by tumor cells (x200). F Lack of KIT expression by tumor cells (x100). G Lack of CD34 expression by tumor cells (x100). H Lack of SS18-SSX expression by tumor cells (x100).

Primary antibody	Source	Clone	Dilution	Treatment
KIT (CD117)	DAKO	Polyclonal	Ready to use	Heat retrieval (pH6.0)
CD56	DAKO	1B6	Ready to use	Heat retrieval (pH9.0)
CD10	DAKO	56C6	Ready to use	Heat retrieval (pH9.0)
PD-L1	DAKO	22C3	Ready to use	Heat retrieval (pH6.0)
Anti-HDAC2	abcam	HDAC2-62	1:1000	Heat retrieval (pH9.0)
SMA	DAKO	1A4	Ready to use	Heat retrieval (pH9.0)
CDX-2	DAKO	DAKO-CDX2	Ready to use	Heat retrieval (pH9.0)
CAM5.2	BectonDickinson	CAM5.2	1:20	Heat retrieval (pH9.0)
Cytokeratin	DAKO	AE1/AE3	Ready to use	Heat retrieval (pH9.0)
CD34	DAKO	NU-4A1	Ready to use	Heat retrieval (pH9.0)
Desmin	DAKO	D33	Ready to use	Heat retrieval (pH9.0)
Vimentin	DAKO	Vim 3B4	Ready to use	Heat retrieval (pH9.0)
SS18-SSX	Cell signaling Technology	SS18-SSX	1:500	Heat retrieval (pH6.0)
GLI1	Santa Cruz Biotechnology	C-1	1:500	Heat retrieval (pH6.0)
CK7	DAKO	OV-TL 12/30	Ready to use	Heat retrieval (pH9.0)
CK20	DAKO	Ks20.8	Ready to use	Heat retrieval (pH9.0)
EMA	DAKO	E29	Ready to use	Heat retrieval (pH9.0)
S100	DAKO	Polyclonal	Ready to use	Heat retrieval (pH6.0)
Chromogranin A	abcam	Polyclonal	1:100	Heat retrieval (pH9.0)
Synaptophysin	DAKO	DAK-SYNAP	Ready to use	Heat retrieval (pH9.0)
DOG1	Leica	K9	1:50	Heat retrieval (pH9.0)

Table 1 Summary of primary antibodies used in this report

positive pathogenic / likely pathogenic variants were not detected with this NGS panel.

Discussion and conclusions

To our knowledge, 16 cases of gastroblastoma have been reported in the medical literature. Table 2 summarizes the clinicopathological features of these cases [1-11], as well as the clinicopathological findings associated with the present case. Nuclear PD-L1 and HDAC2 expression was observed in the spindle cell component by IHC analysis. PD-L1 is transported from the cell membrane into the nucleus and activates other checkpoint inhibition-related genes. PD-L1 transport into the nucleus was shown to be regulated by HDAC2 [12]. In the present case, PD-L1 and HDAC2 were co-expressed in tumoral nuclei. We suggest that PD-L1 migrated into the nucleus via intranuclear HDAC2 activation. As a result, we speculate that HDAC inhibitors may offer a promising treatment option for gastroblastoma. Surgical resection is the standard treatment for gastroblastoma; however, in a small number of cases, chemotherapy or radiotherapy was used [1, 10]. The postoperative course of the disease is generally favorable. However, a few cases of postoperative local recurrence, distant metastases and death have been reported [3, 6, 7, 10]. Therefore, it is valuable to mention feasible treatment options; in this case, we showed nuclear migration of PD-L1 and overexpression of HDAC2, suggesting that HDAC2 inhibitors may be helpful. However, as this was a single case report, further studies are needed to confirm this result.

The tumorigenesis of gastroblastoma is not completely understood. Although Toumi et al. reported that gastroblastoma is believed to develop from a totipotent stem cell, the relationship between gastroblastoma and stem cells is still unclear [10]. Histologically, the embryonic gastrointestinal mesenchyme is morphologically analogous to gastroblastoma (Fig. 5-A, B). In the development and differentiation of the gastrointestinal system, the epithelium and mesenchyme exhibit crosstalk via molecular signaling pathways, such as FGF, TGF-b, Wnt, Hippo, Notch and Hedgehog (Hh). In particular, the Hh signaling pathway is the common pathway for embryo and gastroblastoma development [13]. Gastroblastoma activates GLI1 transcription by the MALAT1-GLI1 fusion gene. We suggest that the morphological similarities between the tumor and the embryonic gastrointestinal mesenchyme might be due to the effect of GLI1 expression via the Hh pathway.

	dn _	⁸				
	Follow-i (month)	42, 60, 11	0	m	20	ω
	Outcome	ANED	ANED	ANED	ANED	ANED
	Metastases	° Z	<u>0</u>	LN, liver,	0 Z	O
	Treatment	SG, PG, antrectomy and radia- tion tion	œ	CHT fol- lowed by PG	Dd	SG and gastroduo- denostomy
	Fusion gene	Q	QN	QN	QN	QN
	IHC negative	CK20 (E); EMA (E); KIT (CD117) (E; S); CD34 (E, S); CD99 (E, S); collectinin (E, S); s); S100 (E, S); s); S100 (E, S); S); S100 (E, S); CDX2 (E, S); and TTF-1 (E, S)	Calretinin; CD34; CEA; CgA; CK (HMW); Des; inhibin; NSE; p63; SMA; Syn	CK20 (E); calretinin (E); CDX2 (E); Des (E); EMA (E); inhibin; p63 (E); S100 (E); SMA	Calretinin; CD34; KIT (CD117); CgA; Des; DOG1; S100; SMA; Syn	ALK; CD34; KIT (CD117); CK5/6; Des; DOG1; PLAP; S100; SMA
	IHC positive	Vim (S), CD10 (S); CK (pan) (E); CK 18 (2/2, focally in one of the cases) (E); CK 7 (focally in E); DOG 1 (focally in E, dot-like posi- tivity in S)	CD10 (S); CD56; KIT (CD117) (E); CK (pan) (E); CK (LMW) (E); EMA (E); Vim (S)	CD10; CD56; KIT (CD117) (E); CK (pan) (E); CK (LMW) (E); CK7 (focally in E); Vim (S)	CD10 (S, focally in E); CD56; CK AE1/AE3 (E); CK CAM5.2 (E), Vim (S)	CD10 (S); CD56 (S); CK AE1/AE3 (E); CK CAM5.2 (E); Vim (S)
	Chief complication	N/A, AP, anemia and fatigue	AP and palpa- ble mass.	N/A	AP	Abdominal mass and blood in stool
	Size (mm)	50, 60. 150	6	38	105	45
	Locus	В, В, А	∢	∢	K	K
	Specimen type	~	٣	٣	£	۲
2 2 2 4	Sex	М, F, М	Σ	Σ	ш	Σ
	Age (year)	19, 27, 30	0	28	19	12
	Author (publication year)	Miettinen et al. (2009) [1]	Shin et al. (2010) [2]	Wey et al. (2012) [3]	Femandes et al. (2014) [4]	Ma et al. (2014) [5]
	Case	<u>~</u>	4	Ŋ	9	\sim

Table 2 Summary of previously published gastroblastoma cases

Table	2 (continue	d)												
Case	Author (publication year)	Age (year)	Sex	Specimen type	Locus	Size (mm)	Chief complication	IHC positive	IHC negative	Fusion gene	Treatment	Metastases	Outcome	Follow-up (month)
∞	Toumi et al. (2017) [10]	29	L L	æ	□	70	Epigastric pain	CD10 (focally); CD99	CK (pan); KIT (CD117); CgA; Syn	Q	æ	L	DEAD	Q
0	Graham et al. (2017) [6]	28	Σ	£	<	38	N/A	CK (pan) (E), CD56 (E), NSE (E), low Ki-67	CgA, Syn, CEA, TTF-1, PLAP, CD30, AFP, HCG	MALAT1– GLI1	R NOS	LN, liver, peritoneum	N/A	N/A
0	Graham et al. (2017 [6])	27	Σ	œ	R	Q	N/A	CK (pan) (E), SMA (patchy in S)	CgA, Syn, KIT (CD117), DOG1, Des, S100, mel- anA, SOX10, TLE-1, CD99, CK5/6	MALAT1 GLI1	R NOS	0 Z	ANED	12
=	Graham et al. (2017) [6]	0	Σ	£	R	8	N/A	CK (pan) (E), CD10 (focally in S), KIT (CD117) (E, S), CD56 (E), Vim (S)	CgA, Syn, NSE, Des, SMA, calretinin, inhibin, CK34βE12, CD34	MALAT 1– GLI 1	R NOS	° Z	ANED	33
1	Graham et al. (2017) [6]	20	ш	Needle biopsy	SZ	6	N/A	OSCAR (patchy in E), low Ki-67, Vim (S)	CK34βE12, CKT7, CK20, Syn, CD34, Syn, CD34, CD99, KIT (CD117), DOG1, wT1, SMA, wT1, SMA, WT1, SMA, MOC31, melanA, HMB45, De5, EMA, MD45, MCFA	GLI1 GLI1	Biopsy	Liver	A/A	A/N

Tablé	2 (continue	()												
Case	Author (publication year)	Age (year)	Sex	Specimen type	Locus	Size (mm)	Chief complication	IHC positive	IHC negative	Fusion gene	Treatment	Metastases	Outcome	Follow-up (month)
<u>ت</u>	Castri et al. (2019) [7]	62	Σ	<u>ح</u>	<	8	Weight loss and dysphagia	bcl2; CD10 (S> E), CD56; CK (pan) (E> S); Vim (S)	aFP, bHCG, h-calde- smon, calponin, CD31, CD99, KIT (CD117), CGA, Des, DOG1, ERG, GFAP, HMB45, melanA, PLAP, S100, STAT6, Syn	GLI1 GLI1	2	Local relapse	ANED	52
,	Giovanni et al. (2019)	6	ш	œ	<	23	bleeding	CD 10 (focally in S), EMA (focally in E), CK (pan) (E), CM 5.2 (E), CM 5.2 (E), in E), Vim (S), GL11 (E, S)	SMA (E, S), Calretinin (E, S), CgA (E, S), CgA (E, S), NSE (E, S), CD34 (E, S), CD34 (E, S), CD29 (E, S), CD29 (E, S), CD20 (E, S), CC20 (E, S), Inhibin (E, S), S100 (E, S), TTF1 (E, S) TTF1 (E, S) TTF1 (E, S)	Q	5	2	ANED	00
15	Koo et al. (2021) [8]	17	Σ	٣		63	Intestinal bleeding	Vim (E, S), CD56 (E, S), CD10 (E, S), CK (pan) (E, S), Syn (E, S)	CD34 (E, S), KIT (CD117) (E, S), DOG1 (E, S), S100 (E, S), Des (E, S), CgA (E, S)	EWSR1- CTBP1	Dd	0 Z	ANED	23

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Tablé	e 2 (continue	(d)												
Case	Author (publication year)	Age (year)	Sex	Specimen type	Locus	Size (mm)	Chief complication	IHC positive	IHC negative	Fusion gene	Treatment	Metastases	Outcome	Follow-up (month)
9	Chen C et al. (2022) [1 1]	ß	Σ	œ	Σ	53	A A	Vim (E, S), CD10 (E, S), bb-2 (E, S), CD56 (E), CD100 (E), EMA (focal), Ki-67 (low)	KIT (CD117) (E, S), DOG1 (E, S), CD34 (E, S), SD34 (E, S), SD34 (E, S), DG6 (E, S), D68 (E, S), D68 (E, S), SOX10 (E, S), S), SOM5.2 (E, S), CAM5.2 (E, S), CK (pan) (E, S).	PTCH-GU2	ESD	2	NVA	A/M
1	Present case / Sugimoto	28	ш	۲	<	2	AP	CK (pan) (E), CAM5.2 (E), CD10 (S), CD56 (S), VIm (S), (S), VIm (S), GL11 (S), CK GL11 (S), CK 7 (E), PD-L1 f(ocally in S), HDAC2 (E, S)	CK20 (E, S), EMA (E, S), CDX2 (E, S), KIT (CD117) (E, S), CD34 (E, S), SMA (E, S), Des (E, S), S100 protein (E, S), Syn (E, S), SS18-SSX (E, S).	MALAT1 GLI1	5 d	Ŷ	ANED	ω
<i>Abbrev</i> SMA Si gastrec	<i>viations: U</i> Gastric mooth muscle ac ctomy, <i>PG</i> Partial	cardia and fund ctin, <i>TTF-1</i> Thyro gastrectomy, <i>R</i>	Jus, B Ga: oid transc Resectio	stric body, A Ga ription factor 1, in, LN Lymph nc	astric antru <i>CgA</i> Chrc ode	um, AP Abdom omogranin A, S	inal pain, <i>IHC</i> Imm <i>yn</i> Synaptophysin,	unohistochemic , S Spindle cell cc	al stain <i>, ANED</i> Al omponent <i>, E</i> Epi	ive no evidence thelial cell comp	of disease, CK_C) oonent, N/A_Not a	ytokeratin, <i>Des</i> D available, <i>ND</i> Not	Jesmin, <i>Vim</i> Vi t done, SG Suk	mentin, itotal



Fig. 4 A MALAT1-GL11 fusion gene detected by molecular analysis. A Confirmation of the presence of a MALAT1-GL11 fusion transcript by RT-PCR analysis. B Sequencing of the cDNA-confirmed fusion of MALAT1 and GL11



Fig. 5 Histology of the fetal gastrointestinal tract (at 10 weeks). A, B Mesenchyme of the gastrointestinal tract, which resembles the spindle component of the gastroblastoma (A: x100, B: x200)

In conclusion, we report the following new findings associated with a case of gastroblastoma: (i) gastric tumors mimic the gastrointestinal mesenchyme in the embryonic period and (ii) nuclear expression of PD-L1 and HDAC2 were observed. We speculate that HDAC inhibitors may offer a promising treatment option for gastroblastoma.

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Authors' contributions

RS contributed to the preparation of the manuscript, Figures and Tables. TS contributed to the preparation of the manuscript, including all aspects of data collection. YA and AS provided clinical support during the preparation of the manuscript. NU, MO, NY, KI, and YO supported interpreting the pathological findings. WH helped carry out the fusion gene analysis. NY performed the immunohistochemical staining. The author(s) read and approved the final manuscript.

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Declarations

Ethics approval and consent to participate Not applicable.

Consent for publication

Written informed consent was obtained from the patient for publication of this case report.

Competing interests

The authors declare no competing interests.

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