# AMPD3 is associated with the malignant characteristics of gastrointestinal stromal tumors

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Received June 1, 2015; Accepted September 1, 2016

DOI: 10.3892/ol.2016.5532

Abstract. Gastrointestinal stromal tumors (GISTs) are mesenchymal tumors of the gastrointestinal tract. It is well known that activating mutations in the receptor tyrosine kinases KIT and platelet-derived growth factor receptor-α have essential roles in the pathogenesis of GISTs. The activation of these receptor protein kinases triggers multiple signaling pathways that promote cell proliferation and survival; however, the exact mechanism by which the activation of these kinases promotes the progression of GISTs remains uncertain. The aim of the present was to search for genes that are associated with the progression of GIST. The present study used reverse transcription-quantitative polymerase chain reaction to demonstrate that adenosine monophosphate deaminase 3 (AMPD3) was highly expressed in GISTs. Furthermore, transfection of GIST-T1 cells with KIT-specific small interfering RNA (siRNA) demonstrated that the expression of AMPD3 was dependent on KIT expression, while the depletion of AMPD3 in human GIST-T1 cells using AMPD3-specific siRNA resulted in the suppression of cell migration and invasion. In addition, AMPD3 depletion sensitized GIST-T1 cells to the tyrosine kinase inhibitor imatinib. The results of the present suggested that the combined inhibition of tyrosine kinases and AMPD3 may be effective for the treatment of GISTs.

## Introduction

Gastrointestinal stromal tumors (GISTs) are the most common mesenchymal tumors of the gastrointestinal tract,

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*Key words:* gastrointestinal stromal tumors, KIT, adenosine monophosphate deaminase 3, migration, invasion, imatinib

and although the exact incidence of GISTs worldwide is not known, the approximate annual incidence of GISTs in Japan is 1500-2000 cases. GISTs are believed to originate from the interstitial cells of Cajal (ICC) (1). The ICC are electrical pacemakers that regulate the motility of the gastrointestinal tract, and are located around the myenteric plexus and the muscularis propria (1). Approximately 85% of GISTs harbor activating mutations in the KIT receptor tyrosine kinase or platelet-derived growth factor receptor-α (PDGFRA) (2,3). Constitutive activation of these receptor tyrosine kinases induces multiple signaling pathways that are essential for the pathogenesis of GISTs (2,3). For example, oncogenic mutations of KIT promote the activation of various signaling cascades, including the Ras/mitogen-activated protein kinase (MAPK), phosphoinositide 3-kinase (PI3K)/Akt and Janus kinase (JAK)/signal transducer and activator of transcription (STAT) signaling pathways, which have been shown to be activated in various tumors (4,5). Imatinib, a potent inhibitor of KIT, is effective for the treatment of unresectable or metastatic GISTs. However, patients with metastatic GISTs eventually develop resistance to imatinib (6); thus, it is necessary to further define the molecular pathogenesis of GISTs, in order to develop alternative therapeutic strategies.

Adenosine monophosphate (AMP) deaminases (AMPDs) catalyze the hydrolytic deamination of AMP to inosine monophosphate, which is a critical step in nucleotide metabolism (7). There are three members of the AMPD family: AMPD1, AMPD2 and AMPD3 (8,9). AMPD1 is exclusively expressed in skeletal muscle, whereas AMPD2 and AMPD3 are ubiquitously expressed. Deficiencies in all three AMPDs have been reported in humans (7), and mutations in AMPD1 were associated with muscle weakness or pain in certain patients (10,11). AMPD2 deficiency is associated with pontocerebellar hypoplasia, which is a rare inherited and progressive neurodegenerative disorder (12). Conversely, individuals with a complete AMPD3 deficiency did not exhibit any significant disorders (13), although elevated levels of adenosine triphosphate (ATP) were detected in the cells of AMPD3-deficient mice (14). Accumulating evidence has suggested that metabolic alterations are associated with the progression of tumor formation (15-19). A recent report suggested that AMPD2 and AMPD3 may be potential targets for cancer treatment (20). The present study aimed to determine the expression and role of AMPD3 in GISTs.

#### Materials and methods

Patients. A total of 16 GIST samples and 6 normal gastrointestinal tract tissues samples were obtained from 16 patients with GIST who underwent surgery at Nagoya University Hospital (Nagoya, Japan) between September 2008 and April 2014. Informed consent was obtained from all patients. Ethical approval was obtained from the Nagoya University Graduate School of Medicine (Nagoya, Japan). Patients were histologically diagnosed. The clinical characteristics of the patients with GIST are shown in Table I.

Cell lines. The human GIST cell line, GIST-T1, was purchased from Cosmo Bio, Co., Ltd. (Tokyo, Japan). GIST-T1 cells were cultured in Dulbecco's modified Eagle's medium (DMEM; Wako Pure Chemical Industries, Osaka, Japan) supplemented with 10% fetal bovine serum (FBS; Biowest, Nuaille, France), and penicillin and streptomycin (100 U/ml; Thermo Fisher Scientific, Inc., Waltham, MA, USA) at 37°C with 5% CO<sub>2</sub> for at least 1 month. SKOV3 (ovarian cancer) and HCT116 (colorectal cancer) cell lines were obtained from the American Type Culture Collection (Manassas VA USA), and KP4 (pancreatic cancer), TE1 (esophagus cancer), KASUMI1 (acute myeloid leukemia), MKN28 (stomach cancer) and QG90 (lung cancer) cell lines were obtained from RIKEN BioResource Center (Tsukuba, Japan). The cells were maintained in DMEM supplemented with 10% FBS and antibiotics.

Small interfering RNA (siRNA) transfection. siRNAs were obtained from Hokkaido System Science, Co., Ltd. (Sapporo, Japan). The sequences of the siRNAs targeting AMPD3 were 5'-CGGGACUUCUAUAACGUGAGA-3' (siAMPD3-1) and 5'-CCGGAUGGCAUUCCGAUAUGA-3' (siAMPD3-2). The sequences of the siRNAs used for KIT knockdown were 5'-GAC GAGAUUAGGCUGUUAUGC-3' (siKIT-1) and 5'-GCAUCA CGGUGACUUCAAUUA-3' (siKIT-2). The sequence of the control siRNA that targeted luciferase was 5'-CUUACGCUG AGUACUUCGATT-3'. GIST-T1 cells were transfected with 50 nM siRNAs using Lipofectamine RNAiMAX (Invitrogen; Thermo Fisher Scientific, Inc.), according to the manufacturer's protocol.

Reverse transcription-quantitative polymerase chain reaction (RT-qPCR). Total RNA was purified from the GIST and normal gastrointestinal tract tissue samples, the nontransfected cell lines and the siRNA-transfected cells using the RNeasy Mini kit (QIAGEN Benelux B. V., Venlo, The Netherlands). cDNA was synthesized using PrimeScript™ Reverse Transcriptase (Takara Bio Inc., Shiga, Japan). qPCR was performed on a LightCycler® Nano Instrument using a FastStart Essential DNA Green Master kit (Roche Diagnostics, Basel, Switzerland), according to the manufacturer's protocol. Thermal cycling conditions were 95°C for 10 sec and 60°C for 30 sec for 35 cycles. The relative mRNA expression levels were normalized to GAPDH using LightCyclerR® Nano Software 1.0 (Roche Life Science, Tokyo, Japan). The sequences of primers used to amplify each gene were as follows: GAPDH forward, 5'-AGGTGGAGGAGTGGGTGT CGCTGTT-3' and reverse, 5'-CCGGGAAACTGTGGCGTG ATGG-3'; AMPD3 forward, 5'-ACATCCTGGCTCTCATCA

CC-3' and reverse, 5'-CAGCAGATGCTTTTGGTTCA-3'; AMPD2 forward, 5'-CGTAGTGCCCCGTATGAGTT-3' and reverse, 5'-CGAGTCACTGTCCGTCTTCA-3'; and KIT forward, 5'-GAAGTCACCGTGATGCCAGC-3' and reverse, 5'-CTCTGTCTGCATTGTTCTGTG-3'. RT-minus was used for negative control and three independent experiments were performed.

Western blotting. Total protein was extracted from the non-transfected and siRNA-transfected cell lines using laemmli sample buffer [20% glycerol, 135 mM Tris-HCl (pH 6.8), 4% SDS, 10% 2-Mercaptoethanol and 0.003% BPB]. The protein concentrations of the lysates were measured using the RC-DC Protein assay (Bio-Rad Laboratories, Hercules, CA). Proteins were separated by 0.01% SDS-PAGE and transferred onto polyvinylidene difluoride membranes. Following blocking with 0.5% skimmed milk in phosphate-buffered saline (PBS), membranes were incubated with anti-KIT (catalog no., 3392; rabbit polyclonal; 1,000 dilution; Cell Signaling Technology, Inc., Danvers, MA, USA) and anti-βactin (catalog no., A1978; clone, AC15; mouse monoclonal; 1:5,000 dilution; Sigma-Aldrich; Merck Millipore, Darmstadt, Germany) antibodies overnight at 4°C. After washing with Tris-buffered saline plus Tween-20 [20 mM Tris (pH 7.5), 150 mM NaCl and 0.1% Tween 20], the membranes were incubated with horseradish peroxidase-conjugated secondary antibodies, followed by 3,3-diaminobenzidine.

*Imatinib sensitivity assay*. GIST-T1 cells were transfected with siRNAs, and 48 h later, the cells were treated with different concentrations of imatinib for 48 h. A cell proliferation assay was performed and the ratio of viable cells to control cells (not treated with imatinib) was calculated. Imatinib was obtained from Sigma-Aldrich (Merck Millipore).

Cell proliferation assay. The siRNA-transfected cells were plated onto a 96-well plate at a density of 1,000 cells/well. The day of transfection was set as day 0, and the number of viable cells on days 1, 2, 3 and 4 were assessed using the Cell Counting kit-8 (CCK-8) assay (Dojindo Molecular Technologies, Inc., Kumamoto, Japan).

EdU incorporation assay. EdU incorporation assays were performed using the Click-iT® Plus EdU Alexa Fluor® 594 Imaging kit (Thermo Fisher Scientific, Inc.). The siRNA-transfected cells were incubated with EdU for 24 h and fixed with 4% paraformaldehyde. The cells were permeabilized using 0.5% Triton X-100, and stained with a reaction cocktail (0.5% Triton X-100 in PBS) and Hoechst stain, according to the manufacturer's protocol. Images of the cells were captured using a fluorescence microscope, and the percentage of EdU-positive cells was evaluated.

Invasion assay. The invasion assay was performed using  $8-\mu m$  pore filters inserted into 24-well Boyden Chambers (Corning Incorporated, Corning, NY, USA). The filter was pre-coated with Matrigel (BD Bioscience, San Jose, CA, USA). GIST-T1 cells (1.5x10<sup>5</sup>) were seeded into the upper chamber and allowed to invade the lower surface of the filter. After 18 h, the cells were fixed with 100% ethanol and stained with 0.5%

Table I. Clinical characteristics of patients with GISTs.

| Variable               | Value      |
|------------------------|------------|
| Gender                 |            |
| Male                   | 6          |
| Female                 | 10         |
| Age (years)            | 63 (46-81) |
| Anatomic site of GIST  |            |
| Stomach                | 12         |
| Small intestine        | 4          |
| c-KIT exon 11 mutation |            |
| Deletion               | 6          |
| Insertion              | 5          |
| Point mutation         | 5          |
| Malignancy             |            |
| Benign                 | 8          |
| Malignant              | 8          |
| Imatinib treatment     |            |
| Yes/no                 | 4/9        |
| Unknown                | 3          |

Data are presented as n or median (range). GISTs, gastrointestinal stromal tumors.

crystal violet. The number of cells in five randomly selected fields was counted, and three independent experiments were performed.

Migration assay. Cell migration was determined using a wound healing assay and a Boyden chamber assay. For the wound healing assay, confluent monolayers of siRNA-transfected cells were scratched with a 200  $\mu$ l sterile pipette tip and imaged 24 h later. The distance between the leading edges of the wound was measured in five randomly selected fields, and three independent experiments were performed. To assess cell migration using the Boyden chamber (8  $\mu$ m pore size and 6.5 mm membrane diameter), siRNA-transfected cells were seeded onto the upper chamber and allowed to migrate to the lower surface of the filter that had been pre-coated with fibronectin. After 12 h, the cells were fixed in 100% ethanol and 0.5% crystal violet. The number of cells in five randomly selected fields was counted, and three independent experiments were performed.

Statistical analysis. Data are expressed as the mean ± standard error. An unpaired t-test was performed to evaluate P-values. The correlation of KIT expression with AMPD3 expression was analyzed by Pearson's correlation analysis using Analyse-it 3.0 (Analyse-it Software, Ltd., Leeds, UK). A difference was considered statistically significant when P<0.05.

### Results

AMPD3 is highly expressed in GISTs and is positively correlated with KIT. To obtain further insight into the

molecular basis of GISTs, genes whose expression levels were upregulated in GISTs were searched for using the Oncomine database (21). This search suggested that AMPD3 is upregulated in GISTs. The present study examined the expression levels of AMPD1, AMPD2 and AMPD3 in 16 GIST specimens and 6 normal gastrointestinal tissues (2 gastric and 4 colon tissues) using RT-qPCR. As is shown in Fig. 1A, the mRNA expression levels of AMPD3 were significantly increased in the GIST specimens, as compared with the normal tissue specimens (P=0.0002). In addition, there was a significant correlation between KIT and AMPD3 expression levels in the GIST samples (r=0.687; P=0.0047; Fig. 1B). The expression levels of AMPD1 and AMPD2 in the GIST samples were ~10-times lower than that of AMPD3. Similar to AMPD3, increased expression of AMPD2 in GIST specimens was observed by RT-qPCR (P=0.0045; Fig. 1C), whereas AMPD1 expression was not increased in GIST specimens (data not shown). AMPD2 expression was also significantly correlated with KIT expression (r=0.449; P=0.0806; Fig. 1D), although to a lesser extent than AMPD3 and KIT. The present study assessed whether AMPD3 expression was correlated with the malignancy of GISTs; however, there was no significant difference in AMPD3 expression between benign and malignant GISTs (P=0.455; Fig. 1E).

AMPD3 and KIT regulate the expression of each other's protein. The present study examined whether AMPD3 expression was regulated by KIT in GIST-T1 cells, which were originally established using GISTs from a Japanese woman (22). The GIST-T1 cell line contains heterozygous deletion mutations at 57 bases in exon 11 of the KIT gene (22). Western blot analysis demonstrated that the protein expression levels of KIT were markedly increased in GIST-T1 cells, as compared with the other cancer cell lines examined (Fig. 2A). Notably, the level of AMPD3 mRNA was significantly increased in GIST-T1 cells, as compared with the other cancer cell lines (Fig. 2B). The expression of AMPD2 was also upregulated in GIST-T1 cells, as compared with other cell lines, although its expression was ~10-times lower than AMPD3 expression (data not shown).

Subsequently, the effect of KIT depletion on AMPD3 expression was assessed. The expression of KIT was suppressed using two different siRNAs, and the level of AMPD3 mRNA was determined using RT-qPCR. Both KIT-specific siRNAs markedly reduced the KIT mRNA level (Fig. 2C). In addition, the level of AMPD3 mRNA was significantly reduced by KIT knockdown (Fig. 2C). Furthermore, whether AMPD3 knockdown using two AMPD3-specific siRNAs was able to affect the level of KIT mRNA in GIST-T1 cells was evaluated using RT-qPCR. As is shown in Fig. 2D and E, AMPD3 knockdown reduced KIT expression at the mRNA and protein level. Notably, depletion of AMPD3 did not affect the level of AMPD2 mRNA (data not shown). These results indicate that KIT and AMPD3 regulate the expression of each other's transcript.

Depletion of AMPD3 reduces the proliferation, migration and invasion of GIST cells. To determine the role of AMPD3 in the malignant characteristics of GIST-T1 cells, the effect of AMPD3 knockdown on cell proliferation was assessed using CCK-8 assays. GIST-T1 cells were transfected with

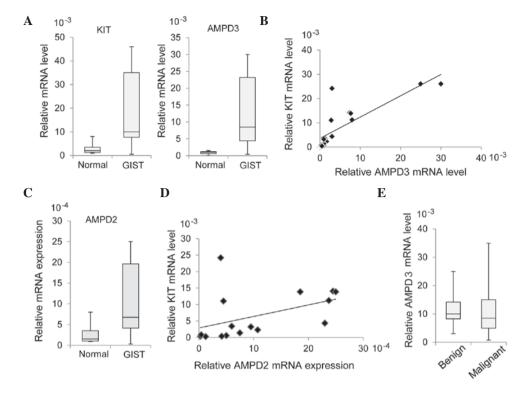


Figure 1. AMPD3 is highly expressed in GISTs. (A) Box plots show the relative mRNA expression levels of KIT and AMPD3 in normal and GIST specimens. Horizontal lines in the boxes indicate the medians and the boxes extend from the 25th to the 75th percentiles. (B) Pearson correlation analysis detected a positive correlation between the mRNA expression levels of AMPD3 and KIT in GIST specimens (r=0.687; P=0.0047). (C) Box plots show the relative mRNA expression level of AMPD2 in normal and GIST specimens. (D) Pearson correlation analysis detected a positive correlation between the mRNA expression levels of AMPD2 and KIT in GIST specimens (r=0.449; P=0.0806). (E) Box plots show the distribution of AMPD3 expression in benign (n=8) and malignant (n=8) GIST specimens. AMPD, adenosine monophosphate deaminase; GISTs, gastrointestinal stromal tumors.

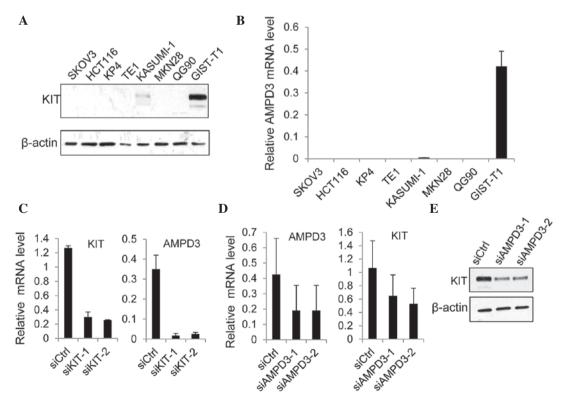


Figure 2. AMPD3 and KIT regulate the expression of each other. (A) The protein expression of KIT in various cancer cell lines was examined by western blotting. (B) The mRNA expression levels of AMPD3 in the different cell lines were evaluated by reverse transcription-quantitative polymerase chain reaction (RT-qPCR). (C) GIST-T1 cells were transfected with siCtrl, siKIT-1 or siKIT-2. After 72 h, the mRNA expression levels of KIT and AMPD3 were measured by RT-qPCR. (D) The mRNA expression levels of AMPD3 and KIT in GIST-T1 cells transfected with siCtrl, siAMPD3-1 or siAMPD3-2 were evaluated by RT-qPCR. (E) The protein expression of KIT in AMPD3-depleted GIST-T1 cells was confirmed by western blotting. AMPD, adenosine monophosphate deaminase; siCtrl, control-small interfering RNA (siRNA); siKIT, siRNA targeting KIT; siAMPD3, siRNA targeting AMPD3.

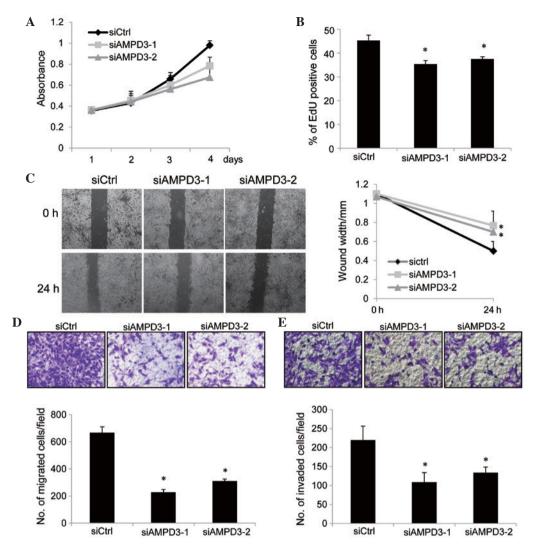


Figure 3. AMPD3 depletion suppressed the proliferation, migration and invasion of gastrointestinal stromal tumor (GIST) cells. (A) GIST-T1 cells were transfected with small interfering RNAs (siRNAs), and the number of viable cells was evaluated using the Cell Counting kit-8 assay. (B) GIST-T1 cells were transfected with siRNAs and, after 48 h, cell proliferation was assessed using the EdU incorporation assay. The graph shows the percentage of EdU-positive cells. Three independent experiments were performed, and the data are presented as the mean ± standard deviation (SD) (\*P<0.05). (C) Confluent monolayers of siRNA-transfected cells were scratched, and the distances between the leading edges were measured at 0 and 24 h. Representative images of the migrated cells are shown. The graph shows the average distance of the wound edge at the indicated time point (\*P<0.05). (D) siRNA-transfected GIST-T1 cells were subjected to a migration assay. Representative images of the migrated cells are shown (0.5% crystal violet stain), and the graph indicates the average number of migrated cells per field. Three independent experiments were performed, and the data are shown as the mean ± SD (\*P<0.05). (E) siRNA-transfected GIST-T1 cells were subjected to a cell invasion assay. Representative images of the invaded cells are shown (0.5% crystal violet stain), and the graph indicates the average number of invaded cells per field. Three independent experiments were performed and data are shown as the mean ± SD (\*P<0.05). Original magnification, x40. AMPD, adenosine monophosphate deaminase 3; siCtrl, control-siRNA; siAMPD3, siRNA targeting AMPD3.

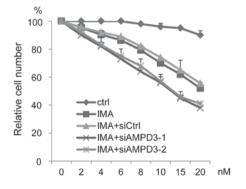


Figure 4. Depletion of AMPD3 sensitizes GIST-T1 cells to imatinib. The siRNA-transfected GIST-T1 cells were cultured in the presence of various concentrations of imatinib, and the proliferation ratio was evaluated. Data are presented as the percent growth of cells treated without imatinib (IMA; imatinib).

siRNAs, and the number of cells was evaluated at various time points. As is shown in Fig. 3A, the proliferation of AMPD3-knockdown cells was reduced compared with the control-siRNA-transfected cells. To examine whether the suppression of cell proliferation was mediated by the inhibition of cell cycle progression or by the induction of apoptosis, EdU incorporation and TUNEL assays (data not shown) were performed. EdU is a thymidine analog and is incorporated into newly synthesized DNA during the S phase. As is shown in Fig. 3B, depletion of AMPD3 significantly reduced the ratio of EdU-positive cells (siAMPD3-1, P=0.01; siAMPD3-2, P=0.02), whereas apoptosis was not induced by AMPD3 knockdown (data not shown). These results indicate that AMPD3 is associated with the progression of the GIST cell cycle.

The migration of AMPD3-knockdown cells was evaluated by performing a scratch assay. Confluent monolayers of siRNA-transfected GIST-T1 cells were scratched, and the migration of the cells into the scratch was observed after 24 h. The migration of AMPD3-knockdown cells was delayed compared with the control-siRNA-transfected cells (siAMPD3-1, P=0.008; siAMPD3-2, P=0.002; Fig. 3C). A modified Boyden chamber assay was performed to further confirm this result. The siRNA-transfected cells were placed on the upper surface of the filter and allowed to migrate to the bottom surface, which was coated with fibronectin. Cells that migrated to the bottom surface were counted to evaluate cell migration. Notably, the migration of AMPD3-depleted cells was suppressed compared with the control-siRNA-transfected cells (siAMPD3-1, P=0.008; siAMPD3-2, P=0.002; Fig. 3D). Finally, the invasion of AMPD3-depleted GIST-T1 cells was examined using a Matrigel-coated Boyden chamber. As is shown in Fig. 3E, AMPD3 suppression significantly delayed the invasion of GIST-T1 cells (siAMPD3-1, P=0.0004; siAMDP3-2, P=0.0007).

AMPD3 depletion increases GIST-T1 cell sensitivity to imatinib. The ability of AMPD3 depletion to affect the sensitivity of GIST-T1 cells to imatinib was evaluated. The siRNA-transfected GIST-T1 cells were treated with various concentrations of imatinib, and cell growth was assessed. As shown in Fig. 4, AMPD3-siRNA-transfected cells were more sensitive to imatinib than the control-siRNA-transfected cells.

#### Discussion

Previous studies have demonstrated that activating mutations in the receptor tyrosine kinases KIT and PDGFRA are critical for the pathogenesis of GISTs (23-25). Recent studies have shown the important function of ETS variant 1 (ETV1), an ETS family transcriptional factor, in the progression of GISTs, and suggested that targeting both KIT and ETV1 may be effective for the treatment of this type of tumor (26,27). KIT is known to activate multiple pathways, including the RAS/MAPK and PI3K/AKT signaling pathways (28,29); thus, activated KIT may promote the expression of various cancer-associated genes. The present study demonstrated that AMPD3 was significantly upregulated in GIST tissue samples, as compared with normal tissue samples, and that AMPD3 expression was correlated with KIT expression. In addition, KIT depletion was shown to suppress the expression of AMPD3 in GIST-T1 cells at the mRNA level. These results suggested that KIT regulates the expression of AMPD3 in GISTs. Notably, it was also demonstrated that AMPD3 knockdown suppressed KIT expression in GIST-T1 cells. Therefore, KIT and AMPD3 may form a positive feedback loop to promote their expression and the progression of GISTs.

AMPD3 is ubiquitously expressed, and mice with a total AMPD3 deficiency showed increased levels of ATP in their cells (14), indicating that AMPD3 contributes to the energy balance in cells. In the present study, the depletion of AMPD3 in GIST-T1 cells suppressed the cell proliferation, migration and invasion. These results indicated that AMPD3 is associated with the malignant characteristics of GIST-T1 cells. AMPD3 depletion reduced KIT expression; thus, the suppression of

malignant characteristics by AMPD3 may be partly mediated by KIT suppression. AMPD3 may also promote cancer progression by contributing to the energy-dependent activation of cancer-associated pathways. For example, AMP-activated protein kinase (AMPK) is known to suppress the progression of various tumors (30-32). Therefore, increases in the levels of AMP as a result of AMPD3 knockdown may promote the activation of AMPK, which can subsequently inhibit anabolic pathways that are essential for cancer cell growth and survival. AMPD3 depletion may promote the activation of protein kinases, such as AMPK, and suppress cancer cell migration and invasion.

In summary, we have shown that the expression levels of KIT and AMPD3 were correlated in GIST-T1 cells, and that the two proteins likely form a positive feedback loop. In addition, it was demonstrated that AMPD3 depletion suppressed the migration and invasion of GIST-T1 cells. Although imatinib is effective in GIST treatment, the drug cannot completely eradicate the tumor. The results of the present study suggested that the combined inhibition of KIT and AMPD3 may be effective for the treatment of GIST.

#### Acknowledgements

The authors would like to thank the members of the Division of Cancer Biology, Nagoya University Graduate School of Medicine for their helpful discussions and technical assistance. This research was funded by a grant from the Ministry of Education, Culture, Sports, Science and Technology of Japan (grant no. 24790689), and supported by the Naito Foundation.

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