

The angiogenesis regulator vasohibin-1 inhibits ovarian cancer growth and peritoneal dissemination and prolongs host survival

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Abstract. Vasohibin-1 (VASH1) is expressed in vascular endothelial cells stimulated by several angiogenic growth factors and displays autocrine activity to regulate angiogenesis via a negative feedback mechanism. In this study, we investigated the effect of VASH1 on ovarian cancer progression using VASH1-expressing ovarian cancer cells *in vitro* and *in vivo*. The growth ability of ovarian cancer cells engineered to express the VASH1 gene remained unchanged *in vitro*. However, we showed that VASH1 secretion by tumor cells inhibited the growth of human umbilical vein endothelial cells. Further, animal experiments showed that VASH1 expression inhibited tumor angiogenesis and growth. In a murine model of peritoneal dissemination of ovarian cancer cells, VASH1 inhibited peritoneal dissemination and ascites, resulting in significantly prolonged survival in mice. This indicates that VASH1 exerts an antitumor effect on ovarian cancer by inhibiting angiogenesis in the tumor environment. These findings suggest that a novel therapy based on VASH1 could be a useful therapeutic strategy for ovarian cancer.

Introduction

Ovarian cancer is the second most common malignant tumor in gynecology and is the fifth leading cause of cancer-related mortality in women (1). Since early ovarian cancer is often asymptomatic, more than half of patients are initially diagnosed at advanced disease stages with peritoneal dissemination and ascites (2). The standard therapy for advanced ovarian cancer is a combination of cytoreductive surgery and platinum/taxane-

based chemotherapy. Ovarian cancer is relatively sensitive to chemotherapy and many patients achieve remission with this combinatorial therapy (3,4). However, the benefit is temporary, and more than half of these patients develop recurrence and die of the disease. Thus, there are limitations to current treatment options, necessitating the development of novel treatment strategies.

Angiogenesis is a phenomenon in which new blood vessels develop from the existing vasculature. Normal angiogenesis is a vital reaction observed only in strictly controlled situations, such as organogenesis during fetal development, the endometrial proliferative stage, and the wound healing process. However, pathological angiogenesis is closely associated with the progression of proliferative diseases (such as diabetic retinopathy and rheumatoid arthritis) and solid tumors. In 1971, Folkman proposed the theory that the growth of tumor cells depends on angiogenesis and that inhibition of angiogenesis is a useful antitumor strategy (5). Subsequently, many studies have been published describing the regulatory mechanisms of tumor angiogenesis and related processes.

Angiogenesis is regulated by a balance between angiogenic growth factors, such as VEGF (6) and FGF (7), and angiogenesis inhibitors, such as angiostatin (8), endostatin (9), pigment epithelium-derived factor (10), 16 kDa prolactin (11), and thrombospondin-1 (12). In 2004, the novel angiogenesis inhibitor vasohibin-1 (VASH1) produced by human umbilical vein endothelial cells (HUVEC) stimulated by VEGF and FGF was identified (13). VASH1 is expressed in VEGF-stimulated vascular endothelial cells and acts on vascular endothelial cells themselves via a negative feedback mechanism that regulates angiogenesis (13,14). Mouse corneal assays have also confirmed that VASH1 has inhibitory effects on pathological angiogenesis caused by several angiogenic factors, including VEGF, FGF, and PDGF (13,15), suggesting that the targets of the anti-angiogenic action are wide-ranging. VASH1 lacks a classical secretory signal and its secretion mechanism was previously unknown. However, it was recently discovered that VASH1 binds small vasohibin-binding protein (SVBP), thereby enabling its secretion (16). VASH1 has been associated with diseases involving abnormal angiogenesis, including malignant tumors (17-21), atherosclerosis (22), diabetic retinopathy

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(23), diabetic nephropathy (24,25), and age-related macular degeneration (26,27). It has also been shown that VASH1 inhibits lymphangiogenesis and lymph node metastasis as well as angiogenesis (15). VASH1 could be effective for the treatment of various diseases, including malignant tumors. Animal experiments using murine lung cancer cell lines have demonstrated the anti-angiogenic and antitumor activity of VASH1 (13). In this study, we investigated the previously unreported antitumor effect of VASH1 on ovarian cancer in an experimental animal model of ovarian cancer.

Materials and methods

Cell lines and culture. The human ovarian cancer cell line SKOV-3 (28) was purchased from the American Type Culture Collection (Manassas, VA, USA). The SKOV-3 cell line, which has previously been reported to produce VEGF (29,30), was cultured in Dulbecco's modified Eagle's medium/F12 (DMEM/F12) medium (Life Technologies, Carlsbad, CA, USA) containing 10% fetal calf serum (Sigma-Aldrich, St. Louis, MO, USA) and 1% penicillin/streptomycin (Life Technologies). HUVECs were obtained from Kurabo Industries, Ltd. (Osaka, Japan) and cultured in a 10-cm, type 1 collagen-coated culture dish (Asahi Glass Co., Ltd., Tokyo, Japan) containing EBM-2 medium (Lonza, Walkersville, MD, USA) supplemented with EGM-2-MV-SingleQuots (Lonza), VEGF, FGF, insulin-like growth factor-1, epidermal growth factor, and 2% fetal calf serum. All cells were cultured at 37°C in a 5% carbon dioxide atmosphere.

Establishment of a VASH1-expressing cell line. Human VASH1 cDNA (NCBI database; Gene ID 22846) was inserted into the cloning site (*Sma*I, *Xba*I) of pCMV-IRES-bsr (31) to generate a VASH1 expression vector. The VASH1 expression vector or the control plasmid vector expressing luciferase [pCMV-LUC-IRES-bsr (31)] was transfected into SKOV-3 cells using Lipofectamine LTX Plus reagent (Invitrogen, Carlsbad, CA, USA) according to the manufacturer's instructions. Subsequently, the transfected cells were selectively cultured in medium containing blasticidin S HCl (Funakoshi Co., Ltd., Tokyo, Japan) and blasticidin-resistant cells were obtained. These cells were cloned by limiting dilution and VASH1-expressing and control clones were established.

Quantitative reverse transcriptase-polymerase chain reaction (RT-PCR). Extraction of mRNA from cultured cells was performed using an RNeasy Mini kit (Qiagen, Valencia, CA, USA) according to the manufacturer's instructions. The mRNA was subjected to quantitative RT-PCR using a One Step SYBR PrimeScript PLUS RT-PCR kit (Perfect Real-Time, Takara, Otsu, Japan) and Thermal Cycler Dice Real Time System II (Takara) according to the manufacturer's protocol. The amount of each mRNA was expressed as a ratio of the glyceraldehyde 3-phosphate dehydrogenase (GAPDH)-corrected fluorescent signal intensity. The primer sequences were as follows: GAPDH forward, ACCACAGTCCATGCC ATCAC; reverse, GGGCCTCTTTGGTCATTTC; VASH1 forward, AGATCCCATAACCGAGTGTG; reverse, GGGCCT CTTTGGTCATTTC.

In vitro cell growth. A total of 1×10^3 cancer cells were seeded in each well of a 96-well plate and an XTT assay kit (Roche Diagnostics, Mannheim, Germany) was used every 24 h. The absorbance was measured 24 h later at 490 nm, and cell growth curves were generated.

Endothelial cell growth in vitro. A total of 1×10^6 cancer cells were seeded in each well of a 6-well plate and cultured in EBM-2 medium containing no growth factors. The culture supernatant was collected 24 h later. A total of 2×10^3 HUVEC were seeded in each well of a 96-well plate and cultured using the culture supernatant. An XTT assay kit was used after 48 h and the absorbance was measured 24 h later at 490 nm.

Western blotting. Cells were lysed using a lysis buffer (1% NP-40, 150 mM NaCl, and 50 mM Tris-HCl, pH 8.0), and proteins were extracted. These samples were mixed with an SDS sample buffer [10 mM Tris-HCl (pH 7.5), 150 mM NaCl, 1% SDS, and EDTA-free proteinase inhibitor cocktail (Roche)]; the mixtures were electrophoresed on 10% polyacrylamide gels, followed by transfer to polyvinylidene fluoride membranes (Merck Millipore, Ltd., Billerica, MA, USA). The membranes were incubated at room temperature for 1 h in Tris buffer (pH 7.6) containing 5% skimmed milk (Wako Pure Chemical Industries, Ltd., Tokyo, Japan), and were incubated at 4°C overnight with 1 µg/ml of mouse monoclonal anti-VASH1 (13) or rabbit anti-actin antibody (Sigma-Aldrich). After washing three times with phosphate-buffered saline-Tween-20 (PBS-T), the membranes were incubated with peroxidase-conjugated anti-mouse or anti-rabbit IgG antibody (GE Healthcare, Buckinghamshire, UK) at room temperature for 1 h. After washing with PBS-T three times, the membranes were subjected to chemiluminescence using an ECL kit (American Biosciences, Piscataway, NJ, USA) and chemiluminescence was detected using a cooled CCD camera system (LAS-4000 mini, GE Healthcare).

Animal experiments. Four- to six-week-old BALB/c nude mice (Clea Japan, Tokyo, Japan) were kept in a specific pathogen-free environment. All animal experiments followed the Guidelines for Animal Experiments of Tohoku University and Jichi Medical University.

Subcutaneous tumor transplant model. Nude mice were subcutaneously transplanted with 5×10^6 tumor cells. The tumors formed were measured with a caliper and tumor volumes [long diameter x (short diameter)² x 1/2] were calculated.

Peritoneal dissemination model and survival time. A total of 5×10^6 cancer cells were used to inoculate the peritoneal cavity of nude mice, and the amount of ascitic fluid and number of tumor nodules on the surface of the intestine and mesentery were measured. Mouse survival was determined twice per day and survival curves were constructed using the Kaplan-Meier method.

Immunohistochemical staining. Subcutaneous tumor transplant model mice were sacrificed using isoflurane inhalation. The tumor was removed, embedded in OCT compound (Sakura Finetek Japan Co., Ltd., Tokyo, Japan) and frozen.

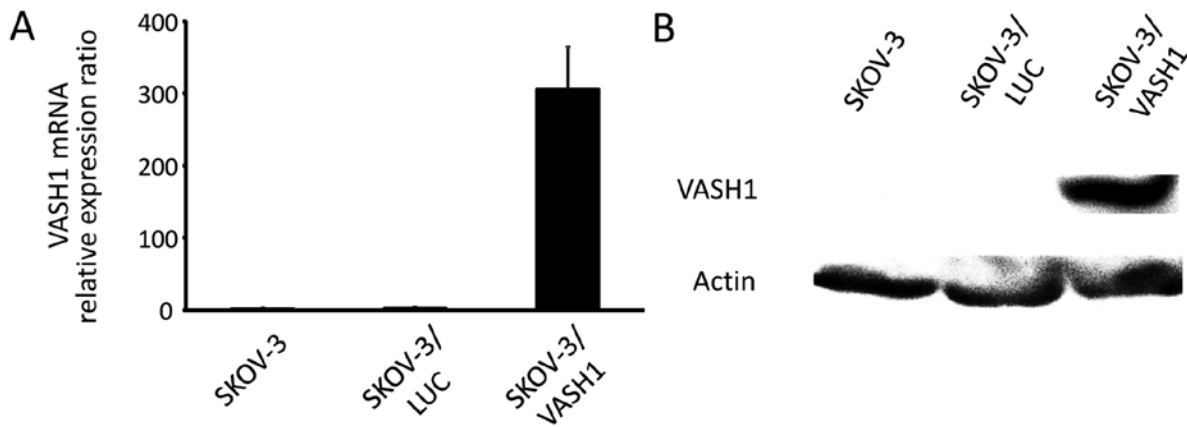


Figure 1. Establishment of a VASH1-expressing cell line. (A) Quantitative RT-PCR of VASH1. (B) Western blotting results. Expression of VASH1 mRNA and protein was only observed in SKOV-3/VASH1.

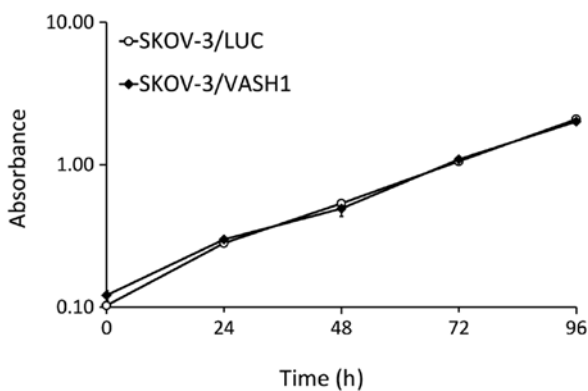


Figure 2. Cell growth curves of SKOV-3/LUC and SKOV-3/VASH1 *in vitro*. No significant difference was observed between groups.

Sections 7-mm thick were cut, fixed in methanol at -20°C for 20 min, blocked with 1% BSA at room temperature for 30 min, and incubated with a 1:500 dilution of anti-mouse CD31 antibody (Research Diagnostics, Flanders, NJ, USA) overnight at 4°C . After washing three times with PBS, sections were incubated with a 1:500 dilution of Alexa 488-conjugated anti-rat IgG antibody (Molecular Probes, Eugene, OR, USA) at room temperature for 1 h, washed three times with PBS and embedded in fluorescent mounting medium. Specimens were analyzed using a BZ-9000 fluorescence microscope (Keyence, Osaka, Japan). The vascular area was calculated as the average blood vessel area in five microscopic fields. Quantitative analysis was performed using BZ-HIC software (Keyence).

Statistics. Student's t-test was used for comparison between two groups. The log-rank test was used to compare Kaplan-Meier survival curves. p-values <0.05 were considered significant.

Results

Establishment of a VASH1-expressing cell line. Fig. 1 shows the results for VASH1 expression analyzed using quantitative RT-PCR and western blotting, respectively, for SKOV-3, SKOV-3/LUC, and SKOV-3/VASH1. Expression of VASH1 mRNA and protein was only observed in SKOV-3/VASH1,

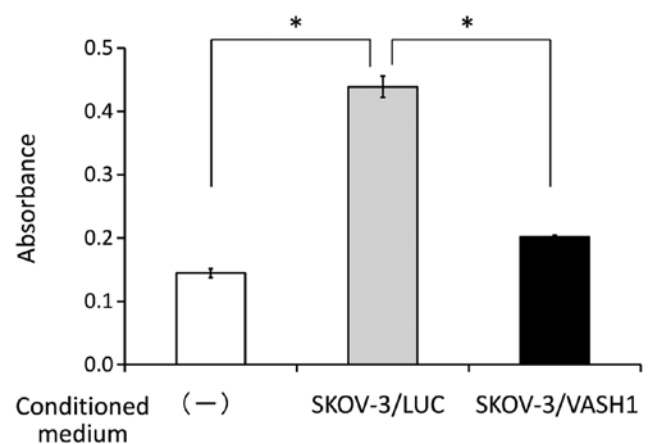


Figure 3. HUVEC number after supplementation with culture supernatant. The absorbance was significantly higher in the group supplemented with the SKOV-3/LUC culture supernatant than in the non-supplemented group but significantly lower in the group supplemented with the SKOV-3/VASH1 culture supernatant compared with the group supplemented with the SKOV-3/LUC supernatant. * $p < 0.01$.

indicating the establishment of the VASH1-expressing ovarian cancer cell line SKOV-3.

Effect of VASH1 on cell growth *in vitro*. The effects of SKOV-3/LUC and SKOV-3/VASH1 on cell growth *in vitro* were compared. As shown in Fig. 2, there was no significant difference in the cell growth curve, indicating that VASH1 expression did not influence the growth of SKOV-3 cells *in vitro*.

The effect of VASH1 secretion by SKOV-3/VASH1 on HUVEC. The effect of VASH1 secretion by SKOV-3/VASH1 on HUVEC was examined. As shown in Fig. 3, the absorbance indicating the number of HUVEC 48 h later was significantly higher in the group supplemented with the culture supernatant of SKOV-3/LUC ($n=3$) than in the non-supplemented group ($n=3$) (0.44 ± 0.02 vs. 0.15 ± 0.01 , respectively; $p < 0.01$), but was significantly lower in the group supplemented with the culture supernatant of SKOV-3/VASH1 ($n=3$) than in the group supplemented with the supernatant of SKOV-3/LUC

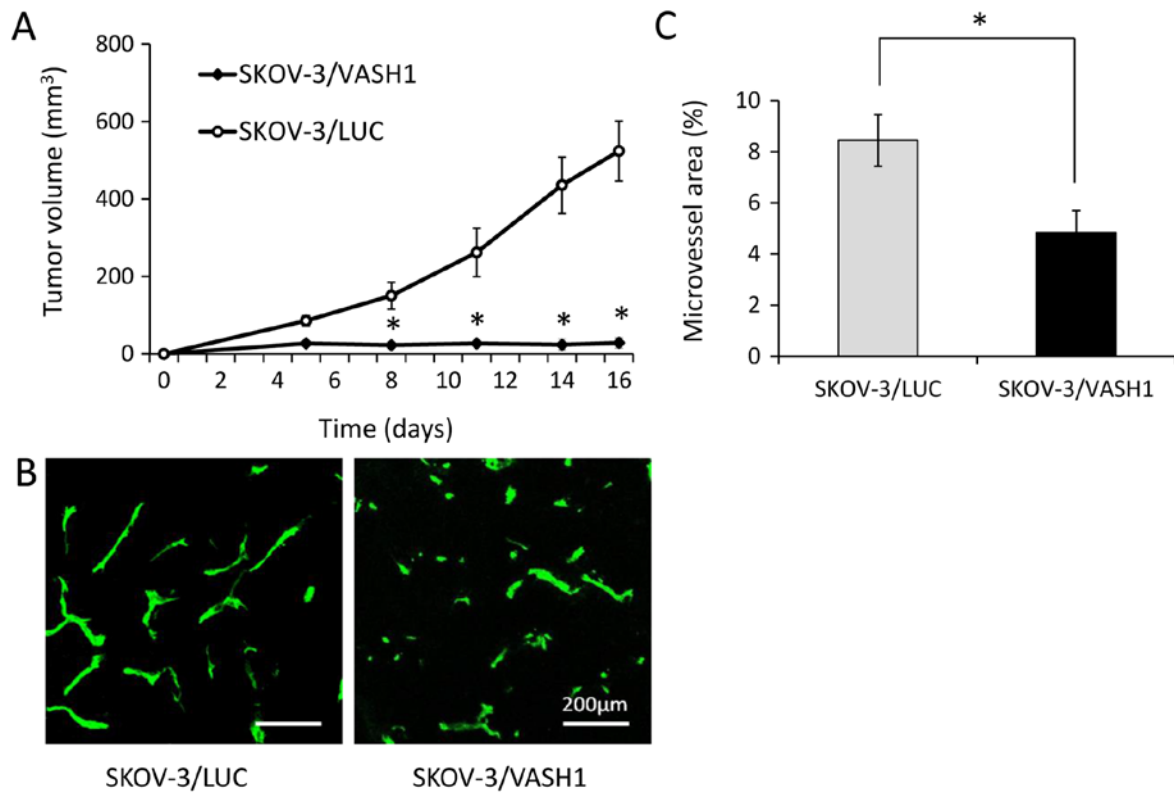


Figure 4. Anti-angiogenic and antitumor activity of VASH1. (A) Growth curves of SKOV-3/LUC and SKOV-3/VASH1 subcutaneous tumors. (B) Immunohistochemical staining of new blood vessels in tumor tissue. (C) The microvessel area in SKOV-3/VASH1 tumors per high-power field was significantly lower than that in SKOV-3/LUC tumors. * $p < 0.01$.

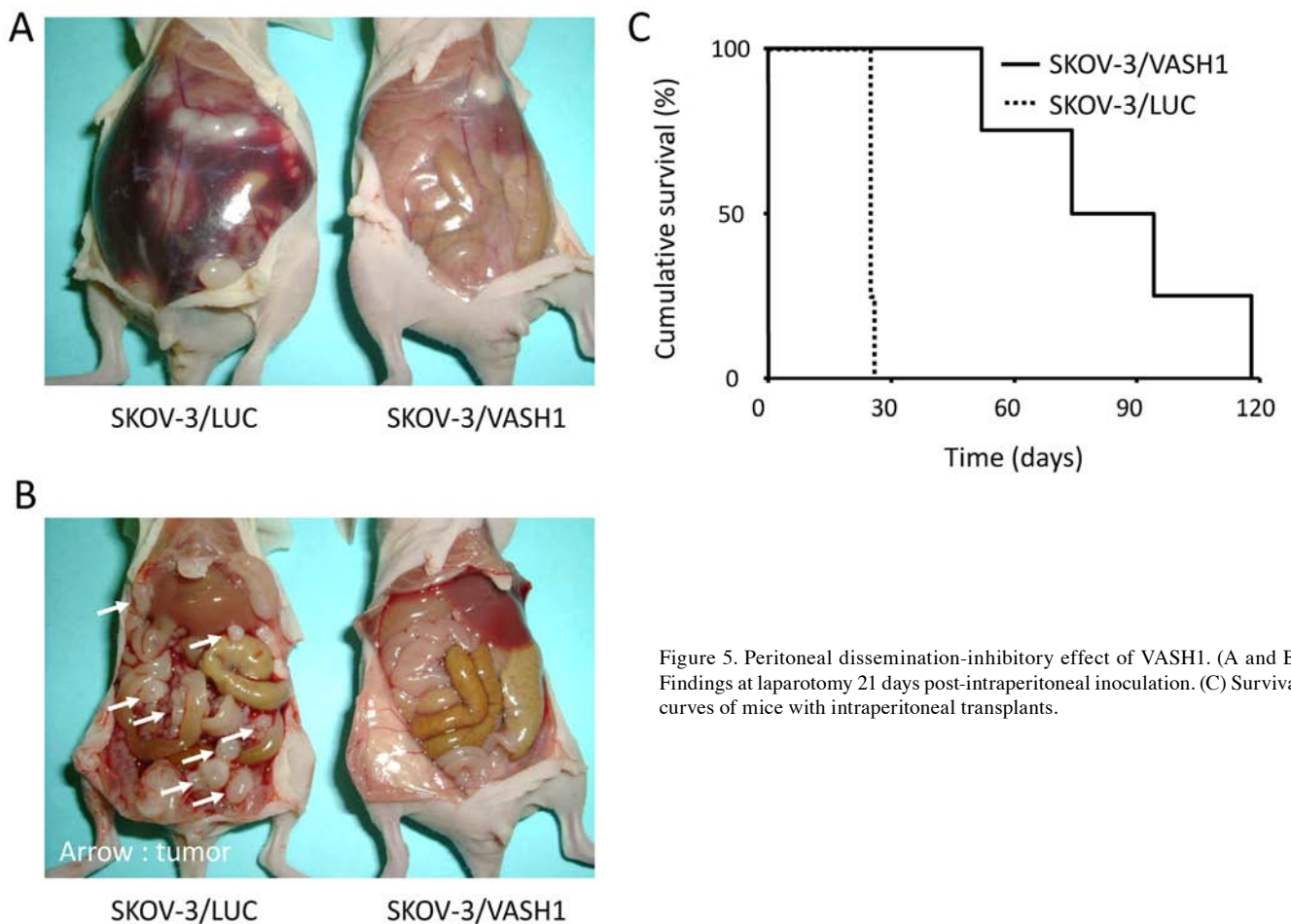


Figure 5. Peritoneal dissemination-inhibitory effect of VASH1. (A and B) Findings at laparotomy 21 days post-intraperitoneal inoculation. (C) Survival curves of mice with intraperitoneal transplants.

(n=3) (0.20 ± 0.002 vs. 0.44 ± 0.02 , respectively; $p<0.01$). This indicates that VASH1 secreted by SKOV-3/VASH1 inhibited the growth of HUVEC *in vitro*.

Antitumor and anti-angiogenic effects of VASH1. Fig. 4A shows the growth curves of SKOV-3/LUC and SKOV-3/VASH1 subcutaneous tumors. Both SKOV-3/LUC and SKOV-3/VASH1 groups formed tumor nodules 1 week after inoculation. Subsequently, tumor enlargement was observed in the SKOV-3/LUC group but not in the SKOV-3/VASH1 group. Thus, VASH1 expression inhibited the subcutaneous growth of SKOV-3 tumors *in vivo*. Next, new blood vessel formation in resected SKOV-3/LUC and SKOV-3/VASH1 tumors was evaluated using immunohistochemical staining (Fig. 4B). The microvessel density in SKOV-3/VASH1 tumors (n=3) per high-power field ($4.9\pm 0.9\%$) was significantly lower than that in SKOV-3/LUC tumors (n=3) ($8.5\pm 1.0\%$, $p<0.01$) (Fig. 4C). Thus, VASH1 expression inhibited angiogenesis and growth in SKOV-3 tumors.

Peritoneal dissemination-inhibiting effect of VASH1. The effect of VASH1 expression on ascites and peritoneal dissemination was studied in murine models of peritoneally disseminated ovarian cancer. Fig. 5A and B show findings at laparotomy 21 days after the inoculation of the peritoneal cavity of nude mice with SKOV-3/LUC or SKOV-3/VASH1 cells. Marked ascites (Fig. 5A, left) and many tumor nodules disseminated in the peritoneum (Fig. 5B, left) were observed in SKOV-3/LUC-inoculated mice but little or no ascites or peritoneal dissemination was observed in SKOV-3/VASH1-inoculated mice (Fig. 5A, right and B, right). The number of disseminated tumor nodules on the surface of the intestine and mesentery was significantly lower in SKOV-3/VASH1-inoculated (n=4) than in SKOV-3/LUC-inoculated (n=4) mice (3 ± 4 vs. 87 ± 48 nodules, respectively; $p<0.01$). Thus, VASH1 expression inhibited peritoneal dissemination of SKOV-3 cells.

Next, the survival of SKOV-3/LUC- and SKOV-3/VASH1-inoculated mice was examined. Survival curves are shown in Fig. 5C. All mice in the SKOV-3/LUC group (n=4) died by day 28 post-inoculation, whereas survival of SKOV-3/VASH1 mice (n=4) was significantly increased ($p<0.01$). Thus, VASH1 expression inhibited the peritoneal dissemination of SKOV-3 cells, significantly prolonging mouse survival.

Discussion

Animal studies using lung or breast cancer cell lines have reported the anti-angiogenic and antitumor activity of VASH1 (13,15,17). However, no studies have reported its effects on ovarian cancer, which spreads by peritoneal dissemination, a unique pattern of tumor progression. Therefore, we investigated the antitumor effect of VASH1 on ovarian cancer in murine models of peritoneally disseminated ovarian cancer.

In this study, the growth ability of ovarian cancer cells engineered to express the VASH1 gene remained unchanged. An experiment using HUVECs showed that VASH1 secreted by tumor cells inhibited the growth of vascular endothelial cells. Next, animal experiments showed that VASH1 expression inhibited angiogenesis in tumors. In addition, in a murine model of peritoneal dissemination of cancer cells, VASH1

inhibited peritoneal dissemination and ascites, thus significantly prolonging survival. This indicates that VASH1 exerts an antitumor effect on ovarian cancer by inhibiting angiogenesis in the tumor environment.

In recent years, molecular-targeted therapy against VEGF has been clinically applied and has demonstrated some effectiveness. Bevacizumab, an anti-VEGF monoclonal antibody, has been widely used in the clinical setting in the treatment of colorectal cancer, non-small cell lung cancer, and breast cancer. In addition, a phase III clinical trial of bevacizumab for ovarian cancer showed that the progression-free survival time was significantly higher in a group of patients treated with bevacizumab and chemotherapy followed by continued bevacizumab monotherapy than in a group of patients treated with chemotherapy alone (32). Furthermore, large-scale clinical studies reported similar midterm results (33,34), supporting the efficacy of bevacizumab therapy for ovarian cancer. However, a phenomenon called 'evasive resistance', which is observed in tumors after anti-VEGF therapy, has attracted attention (35). Proposed mechanisms for the acquisition of this resistance include the induction of angiogenic factors other than VEGF, mobilization of bone marrow-derived endothelial progenitor cells, covering of tumor vascular endothelial cells by pericytes, and an increase in the invasive capacity of tumor cells. An animal experiment using an antibody targeting the VEGF receptor (VEGFR) 2 showed that vascular regression and tumor reduction occurred first, followed by the induction of angiogenesis, leading to tumor regrowth (36). Analysis of the mRNA of regrown tumors revealed increased expression of angiogenic factors other than VEGF, such as FGF, ephrin, and angiotensin (36). In addition, a clinical study reported that the treatment of glioblastoma patients with VEGFR inhibitors resulted in increased serum FGF levels (37). Since various angiogenic factors are involved in tumor angiogenesis, there are limitations to treatment targeting a single angiogenic factor. Therefore, to achieve sufficient therapeutic benefit, it may be necessary to simultaneously target multiple angiogenic factors. In the above-cited study, the combined use of anti-VEGFR2 antibodies and soluble FGF receptor (an FGF antagonist) successfully delayed tumor regrowth (36). In our study, VASH1 inhibited angiogenesis mediated by various angiogenic factors other than VEGF (13,15). These observations suggest that VASH1, which acts alone to inhibit multiple angiogenic factors, is a more effective therapeutic agent compared with VEGF inhibitors in terms of overcoming evasive resistance.

Physiologically, VASH1 is induced in vascular endothelial cells to prevent excessive angiogenesis associated with VEGF stimulation. The application of VASH1 to the treatment of diseases caused by abnormal angiogenesis requires doses that far exceed the physiological VASH1 expression (38). To date, the local administration of purified VASH1 protein preparations (27,38) and local (38,39) or systemic (15,17,22,24,25) administration of an adenovirus vector encoding VASH1 have been attempted. We previously reported the efficacy of therapeutic strategies using adeno-associated virus (AAV) vectors carrying various therapeutic factors (40,41). AAV vectors have advantages, in that they are derived from non-pathogenic viruses with a strong safety profile; also, non-dividing cells, such as neurons and muscle cells, can also be efficiently

transfected, with transfected genes expressed over a prolonged period of time (42). If these advantages were exploited, it would be possible to transfect skeletal muscle cells with therapeutic genes using AAV vectors carrying therapeutic factors to induce muscle cells expression and secretion of these factors into the blood and to deliver them to target lesions. Using an animal model, we demonstrated the feasibility of inhibiting ovarian cancer progression by constructing an AAV vector containing soluble VEGFR-1 (sFlt-1), a VEGF antagonist, and injecting this vector into skeletal muscle (40). Using a similar strategy, we showed that an AAV vector carrying soluble VEGFR-3 (sFlt-4), an antagonist of VEGF-C that promotes tumor lymphangiogenesis and lymphatic metastasis, successfully inhibited lymph node metastasis of endometrial carcinoma (41). A gene therapy strategy using vectors such as AAV is a promising application of VASH1 therapy. In this case, since SVBP is required for the secretion of VASH1, co-transfection of VASH1 and SVBP expression vectors could be effective in increasing VASH1 secretion into the blood.

Unlike sFlt-1 and sFlt-4, the underlying inhibitory mechanism of VASH1 is still unknown. We are currently trying to elucidate this mechanism. If this mechanism is understood in future, it might be possible to identify patients who will benefit from VASH1.

Anti-angiogenic therapy is now approved for several types of cancers and drugs targeting the VEGF signals are in clinical use. However, such drugs can have side effects that include hypertension and proteinuria due to the impairment of normal quiescent vessels. It was recently reported that VASH1 protects endothelial cells from premature senescence and stress-induced cell death via the induction of superoxide dismutase 2 and sirtuin 1 (43). Animal experiments demonstrated that VASH1 did not increase mean blood pressure and urinary albumin excretion (44). It has also been reported that VASH1 does not affect any morphological changes in normal blood vessels (15), wound healing, body weight, and peripheral blood flow (45) in adenoviral VASH1 gene-treated mice. These findings suggest that VASH1 can be a potential candidate for anti-angiogenic treatment.

In conclusion, this study showed that VASH1, a negative feedback regulator of angiogenesis, inhibited tumor angiogenesis and peritoneal dissemination in ovarian tumors, thereby prolonging survival in mouse models of ovarian cancer. These results suggest that a novel therapy based on VASH1 could be a very useful therapeutic strategy for ovarian cancer.

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