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HDAC inhibitor, MS-275, increases vascular permeability by suppressing Robo4 expression in endothelial cells

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ABSTRACT

Roundabout guidance receptor 4 (Robo4) is an endothelial-specific membrane protein that suppresses pathological angiogenesis and vascular hyperpermeability by stabilizing endothelial cells. Robo4 suppresses severe systemic inflammation induced by pathogens and endotoxins and inhibits tumor growth and metastasis, therefore serving as a potential therapeutic target. Although the regulation of Robo4 expression through transcription factors and epigenetic mechanisms has been studied, the role of histone deacetylases (HDACs) has not been explored. In the present study, we investigated the involvement of HDACs in the regulation of Robo4 expression. An HDAC inhibitor, MS-275, which inhibits HDAC1, HDAC2, and HDAC3, was found to suppress Robo4 expression in endothelial cells. Small interfering RNA (siRNA)-mediated knockdown of HDAC3, but not of HDAC1 and 2, also decreased its expression level. MS-275 downregulated the expression of the transcription factor complex GABP, in addition to suppressing Robo4 promoter activity. GABP expression was also downregulated by the siRNA against HDAC3. MS-275 decreased the transendothelial electrical resistance of a monolayer of mouse endothelial cells and increased the rate of leakage of Evans blue dye in the mouse lungs. In addition, MS-275 accelerated cell migration through the endothelial cell monolayer and augmented cell extravasation in the mouse lungs. Taken together, we demonstrated that MS-275 suppresses Robo4 expression by inhibiting HDAC3 in endothelial cells and enhances endothelial and vascular permeability. Thus, we demonstrated a novel mechanism regulating Robo4 expression and vascular permeability, which is anticipated to contribute to future therapies for infectious and inflammatory diseases.

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Introduction

Endothelial cells form a layer on the inner surface of the vasculature. The permeability of blood vessels is regulated by modulation of the VEcadherin-mediated adherens junction between endothelial cells. Endothelial permeability is regulated by various stimulators, such as vascular endothelial growth factor (VEGF) and tumor necrosis factor α (TNFα). Roundabout guidance receptor 4 (Robo4) is an endothelial-specific membrane protein that stabilizes endothelial cells and the vasculature.^{[1](#page-9-0),2} It suppresses vascular hyperpermeability through multiple mechanisms

by interacting with different proteins, including Slit2, Unc5B, and TRAF7. $3-5$ Robo4 improves the survival rate in mouse systemic inflammation models injected with pathogens and endotoxins by suppressing endothelial and vascular hyperpermeability.^{[3,](#page-9-2)6} Thus, Robo4-mediated regulation of vascular permeability could serve as a potential therapeutic target. It is, therefore, important to understand the mechanisms governing the regulation of Robo4 expression and to devise strategies to regulate its expression.

Robo4 expression is regulated by a 3 kb promoter.^{[7](#page-9-4)} SP1 and GA-binding protein

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(GABP), transcription factors that bind to the proximal region of this promoter, are essential for its activity, and thereby for Robo4 expression.[7](#page-9-4)[,8](#page-9-5) GABP is a heterodimeric complex of proteins consisting of two structurally unrelated subunits: GABPα and GABPβ/γ. GABPα binds to the DNA via the ETS domain, whereas GABPβ and its isoform, GABPγ, interact with GABPα. In addition to the transcription factors, epigenetic mechanisms also regulate Robo4 expression. For example, the methylation pattern of the proximal promoter regulates Robo4 expression. $9,10$ $9,10$ The components of the polycomb repressor complex 2 (PRC2) binding to the upstream promoter regulate DNA methylation and Robo4 expression.¹¹ These findings implied that other epigenetic factors, such as histone deacetylases (HDACs), may also be involved in the regulation of Robo4 expression in endothelial cells. However, the involvement of such factors has not been studied yet.

HDACs are enzymes that remove acetyl groups from histones and other proteins. Eighteen characterized members of the HDAC superfamily have been identified in humans and are divided into four classes.[12](#page-10-0) HDACs regulate the acetylation of histones in nucleosomes, which in turn modulates chromatin conformation and gene expression. HDACs also deacetylate non-histone proteins, including transcription factors, and regulate cellular processes under physiological and pathological conditions.

Several inhibitors targeting various combinations of HDAC subtypes have been generated. Certain inhibitors, including the pan-HDAC inhibitor trichostatin A (TSA), have been shown to regulate the expression of endothelial genes and endothelial hyperpermeability in inflammation.^{[13](#page-10-1)[,14](#page-10-2)} In the present study, we investigated whether Robo4 expression is regulated by HDACs by treating endothelial cells with HDAC inhibitors. We determined that MS-275 suppresses Robo4 expression by inhibiting HDAC3 and enhances vascular permeability. Herein, we demonstrated a hitherto unknown mechanism of HDACmediated regulation of Robo4 expression and vascular permeability, and proposed a novel strategy for regulating vascular permeability with small molecules.

Materials and methods

Cell culture

Human umbilical vein endothelial cells (HUVECs) were obtained from Lonza (Basel, Switzerland) and maintained in EGM-2-MV medium (Lonza). The mouse pancreatic endothelial cell line, Mile Sven 1 (MS1), and mouse melanoma cell line, B16-F10, were obtained from the American Type Culture Collection (ATCC; Manassas, VA); they were maintained in Dulbecco's modified Eagle's medium (Nacalai Tesque, Kyoto, Japan), supplemented with 10% fetal bovine serum, 100 U/mL penicillin, and 100 µg/mL streptomycin. All the cells were cultured under a humidified atmosphere with 5% $CO₂$ at 37°C.

Quantitative reverse transcription-polymerase chain reaction

For the assay using HDAC inhibitors, HUVECs were treated with 0.3 μM TSA (FUJIFILM, Osaka, Japan), 40 μM TC-H 106 (Tocris, Bristol, UK), 5–20 μM MS-275 (Santa Cruz Biotechnology, Dallas, TX; AdooQ BioScience, Irvine, CA; Carbosynth, Berkshire, UK), 10 μM MC1568 (MedChemexpress, Monmouth Junction, NJ), 100 μM bufexamac (FUJIFILM), or 0.5% dimethyl sulfoxide (DMSO; as a control). MS1 cells were treated with 20 μM MS-275. For the assay using small interfering RNAs (siRNAs), HUVECs $(2 \times 10^5 \text{ cells})$ were transfected with 9-30 pmol siRNAs against hHDAC1 (SI00070609; Qiagen, Hilden, Germany), HDAC2 (s6494; Thermo Fisher Scientific, Waltham, MA), hHDAC3 (SI03057901; Qiagen), or their control siRNA (SI03650318; Qiagen) (Table S1). Total RNA from the HDAC inhibitor-treated or siRNA-transfected cells was prepared using the RNeasy Mini Kit (Qiagen) and subsequently reverse-transcribed into complementary DNA (cDNA) using Superscript VILO Master Mix (Invitrogen, Carlsbad, CA). Real-time PCR was performed using cDNA as a template, QuantiTect SYBR Green PCR Kit (Qiagen), and specific primers (Table S1). The copy number of the target transcripts was calculated from standard curves prepared using known amounts of plasmids containing the target sequences. The expression levels of Robo4, SP1, GABP subunits, and HDAC1– 3 were normalized against glyceraldehyde 3-phosphate dehydrogenase (GAPDH) levels.

Western blot analysis

Total cell extracts from HUVECs treated with HDAC inhibitors were separated by sodium dodecyl sulfatepolyacrylamide gel electrophoresis, and transferred onto polyvinylidene fluoride membranes. The membranes were incubated with primary antibodies against Robo4 (AF2366; R&D Systems, Minneapolis, MN) or GAPDH (MAB374; Merck Millipore, Burlington, MA) and horseradish peroxidaseconjugated secondary antibodies. Immunoreactive bands were detected using the ImageQuant LAS4010 system (GE Healthcare, Chicago, IL).

Reporter assay

HUVECs (1×10^5 cells) were transfected with 1 µg of a luciferase reporter plasmid containing the Robo4 promoter (pGL3-Robo4⁷) using FuGENE6 (Promega, Madison, WI) and incubated for 24 h. The transfected cells were then treated with 20 μ M MS-275 for 24 h and analyzed by reporter assays using the Dual-Luciferase Reporter Assay system (Promega).

Measurement of transendothelial electrical resistance

MS1 cells (1×10^5 cells) were seeded in cell culture inserts with a pore size of 0.4 µm (BD Falcon, Franklin Lakes, NJ) and incubated for 24 h. The medium in the upper chamber was replaced with fresh medium containing 20 µM MS-275 or DMSO (for control) and transendothelial electrical resistance (TEER) was measured using the CellZscope system (NanoAnalytics, Münster, Germany). Data are expressed for each TEER value (Ω/cm^2) of MS-275-treated or DMSO-treated cells normalized to the values at time $(t) = 0$.

WST-8 assay

MS1 cells (2.5 \times 10⁴ cells) were treated with 20 μ M MS-275 for 48 h or were transfected with 2.5 pmol of siRNAs against mRobo4 or its control for 48 h and

then incubated in a medium containing 10% WST-8 reagent (Dojindo, Kumamoto, Japan) for 1 h. The reaction was terminated by adding 0.1 N HCl and optical density of the samples was measured at 450 nm.

Vascular permeability assay (Miles assay)

Female C57BL/6 N mice (7–8 weeks old) were administered MS-275 (30 mg/kg/day) by oral gavage for four consecutive days. One hour after the final dose, the mice were intravenously injected with PBS containing 1% Evans blue. One hour later, the mice were perfused with PBS containing 2 mM ethylenediaminetetraacetic acid (EDTA) and the lungs were harvested, minced, and incubated in formamide for 48 h. The eluted Evans blue was quantified by measuring the optical density at 620 nm.

Experimental metastasis model

A single-cell suspension of B16-F10 cells was prepared in phosphate-buffered saline (PBS) containing 0.02% EDTA. For the assay using MS-275–treated mice, female C57BL/6 N mice (7–8 weeks old) were administered 200 µL of water containing 1% DMSO and 30% polyethylene glycol 400 with or without MS-275 (30 mg/ kg/day) by oral gavage for four consecutive days. On the day of final injection of MS-275, the B16-F10 cell suspension $(3 \times 10^5 \text{ cells})$ was intravenously injected into the mice. For the assay using Robo4-deficient mice,^{[3](#page-9-2)} the same number of B16-F10 cells was intravenously injected into female Robo4-deficient and littermate control mice (7–8 weeks old). After 14 days of the injections, lungs were harvested from these mice, and fixed in Bouin's solution. The metastasized melanoma colonies on the lung surface were counted under an SZX7 stereo microscope (Olympus, Tokyo, Japan) equipped with an MAS-500 USB camera (Marutsuelec, Tokyo, Japan). All animal experiments were approved by the ethics committee of Osaka University and performed in accordance with the relevant guidelines and regulations.

Transmigration assay

MS1 cells $(4 \times 10^4$ cells) were seeded in FluoroBlok inserts with a pore size of 8.0 μm (Corning, Corning, NY) and transfected with 10 pmol of siRNA against mRobo4 or LacZ (as a control) (Table S1; Hokkaido System Science, Hokkaido, Japan) using Lipofectamine RNAiMAX (Invitrogen, Carlsbad, CA, USA) or were treated with 20 µM MS-275. Twenty-four hours after the transfection or treatment, B16- F10 cells $(2 \times 10^5$ cells) labeled with Cellstain Calcein AM solution (Dojindo) were added to the MS1 cells and incubated for 24 h. The transmigrated B16-F10 cells were then counted using a BZ-X700 fluorescence microscope with the BZ-X analyzer software (Keyence, Osaka, Japan).

Statistical analysis

Data are expressed as mean values ± standard error (s.e.m.) and graphically presented as scatter plots. Normality was tested using the Shapiro–Wilk test. For samples with a normal distribution, *p*-values were calculated using the

unpaired *t*-test and one-way analysis of variance (ANOVA) followed by Bonferroni correction or Dunnett's test. All statistical analyses were performed using the Prism 9 software (GraphPad Software Inc., San Diego, CA). The statistical significance of differences in the means was determined using the tests indicated in the figure legends. A p -value < 0.05 was considered statistically significant.

Results

HDAC inhibitors suppressed Robo4 expression in endothelial cells

To investigate whether HDACs regulate Robo4 expression in endothelial cells, HUVECs were treated with HDAC inhibitors targeting the different HDAC subtypes. Following this, Robo4 expression was analyzed at both the mRNA and protein levels ([Figure 1a,b](#page-3-0)). TSA, TC-H 106, and MS-275, which inhibit HDAC1, HDAC2, and HDAC3, respectively, strongly suppressed Robo4 expression; conversely, MC 1568 and Bufexamac, which target other HDACs, resulted in moderate or no effect on Robo4 expression

Figure 1. HDAC inhibitors targeting HDAC1-3 suppress Robo4 expression in endothelial cells. (A) HDAC inhibitors for various combinations of HDAC subclasses. (B) Effect of HDAC inhibitors on Robo4 expression. HUVECs were treated with the HDAC inhibitors and Robo4 expression was analyzed based on mRNA (n = 4) and protein levels (lower panel). (C) Robo4 expression in HVUECs treated with various concentrations of MS-275 (n = 4). Data are expressed as the mean \pm s.e.m. Statistical significance was determined by Dunnett's test (***p* < .01; N.S., not significant).

at the mRNA and protein levels. Moreover, MS-275-induced suppression of Robo4 was determined to be dose-dependent ([Figure 1c](#page-3-0)). These results indicated that Robo4 expression is regulated by HDAC1, HDAC2, or HDAC3. To identify the HDAC that primarily regulates Robo4 expression, HUVECs were transfected with siRNAs against HDAC1, HDAC2, and HDAC3. The knockdown of HDAC3, but not of HDAC1 and HDAC2, suppressed Robo4 expression ([Figure 2](#page-4-0)). This indicated that MS-275 suppresses Robo4 expression by inhibiting HDAC3.

MS-275 suppressed Robo4 promoter activity by downregulating GABP expression

The expression of human Robo4 is regulated by a 3 kb promoter, which is activated by the binding of transcription factors SP1 and GABP to its proximal region [\(Figure 3a\)](#page-5-0). To investigate whether MS-275 influences Robo4 promoter activity, we analyzed the promoter activity in HUVECs treated with or without MS-275 using a luciferase reporter assay. MS-275 hindered luciferase activity, which indicated that it suppresses Robo4 promoter activity [\(Figure 3b](#page-5-0)). We further investigated whether MS-275 regulates the expression of SP1 and GABP using quantitative PCR [\(Figure 3c\)](#page-5-0). MS-275 suppressed the expression of the GABP subunits but not that of SP1. Similarly, the siRNA against HDAC3 also suppressed the

expression of the GABP subunits but not that of SP1 [\(Figure 3d\)](#page-5-0). Taken together, these results indicated that MS-275 downregulates GABP expression by inhibiting HDAC3 activity and suppresses the activity of the Robo4 promoter.

MS-275 enhanced vascular and endothelial permeability

Because Robo4 has been shown to suppress vascular and endothelial permeability, we investigated whether MS-275 impacts vascular permeability. The mice were treated with MS-275 and analyzed for vascular leakage by injecting Evans blue intravenously. MS-275 treatment increased the leakage of Evans blue in the mouse lungs ([Figure 4a\)](#page-6-0). Consistent with this result, MS-275 suppressed Robo4 expression in mouse pancreatic endothelial cells (MS1 cells) [\(Figure 4b](#page-6-0)), and enhanced the TEER of an MS1 cell monolayer without hampering the viability of these cells [\(Figures 4c](#page-6-0) and d). These results indicated that MS-275 enhances vascular and endothelial permeability.

MS-275 enhanced cell migration in an experimental metastasis model

To investigate whether MS-275-induced vascular hyperpermeability regulates cell extravasation from the vasculature, experimental

Figure 2. Knockdown of HDAC3 downregulates Robo4 expression in endothelial cells. (A) Effect of knockdown of HDACs on Robo4 expression. HUVECs were transfected with the siRNAs against HDAC1 (A), HDAC2 (B), and HDAC3 (C), and the expression levels of HDACs and Robo4 were measured by quantitative PCR (n = 4). Data are expressed as the mean \pm s.e.m. Statistical significance was determined by the unpaired *t*-test (***p* < .01; N.S., not significant).

Figure 3. (a) Schematic illustration of the Robo4 promoter and binding sites of SP1 and GABP in the proximal region.^{[7](#page-9-4),8} (b) Effect of MS-275 on Robo4 promoter activity. A reporter plasmid containing the 3 kb Robo4 promoter was transfected into HVUECs treated with or without 20 μ M MS-275. Luciferase activity was measured ($n = 4$). (c, d) Expression of SP1 and GABP subunits in the HUVECs treated with MS-275 (c) or siRNA against HDAC3 (d) was measured by quantitative PCR. Data are expressed as the mean \pm s.e.m. Statistical significance was determined by the unpaired *t*-test ($p < .05$, $*p < .01$; N.S., not significant).

metastasis models were established by intravenously injecting B16-F10 melanoma cells into mice pretreated with or without MS-275. MS-275 treatment increased the number of B16-F10 colonies in the lungs [\(Figure 5a\)](#page-7-0). Consistent with this *in vivo* result, treatment of the MS1 cell monolayer with MS-275 enhanced the transendothelial migration of B16-F10 cells without affecting the viability of MS1 cells [\(Figure 5b](#page-7-0)). Taken together, these results indicated that MS-275 enhances cell extravasation from the vasculature.

Downregulation of Robo4 enhanced cell extravasation

To investigate whether the downregulation of Robo4 contributes to cell extravasation, the experimental metastasis model was subjected to Robo4-deficient mice. The injection of B16-F10 colonies into Robo4−/− mice resulted in an increased number of colonies in the lungs compared with those in the Robo $4^{+/+}$ and Robo $4^{-/+}$ mice [\(Figure 6a\)](#page-8-0). In the same model, the use of Robo $4^{-/-}$ mice resulted in increased leakage of

Figure 4. MS-275 enhances endothelial and vascular permeability. A) Vascular leakage in the lungs of mice treated with or without MS-275. Mice were administered MS-275 (30 mg/kg/day) or dimethyl sulfoxide (DMSO; Vehicle) by oral gavage for four consecutive days and intravenously injected with Evans blue dye. The extravasated dye was quantified $(n = 3)$. (B) Robo4 expression in MS1 cells treated with or without MS-275 ($n = 4$). (C) Effect of MS-275 on endothelial permeability. MS1 cells were treated with MS-275 or DMSO and transendothelial electrical resistance (TEER) was measured (n = 3). (D) Effect of MS-275 on the viability of endothelial cells. WST-8 assays were performed using MS1 cells with or without MS-275 treatment ($n = 6$). Data are expressed as the mean \pm s.e.m. Statistical significance was determined by the unpaired *t*-test or Bonferroni correction (**p* < .05, ***p* < .01).

Evans blue dye in the lungs ([Figure 6b](#page-8-0)). Consistent with this, the siRNA-mediated knockdown of Robo4 in MS1 cells enhanced the transendothelial migration of B16-F10 cells without impacting the viability of MS1 cells [\(Figures 6c](#page-8-0) and d). Taken together, these results indicated that the downregulation of Robo4 enhances endothelial and vascular permeability and cell extravasation.

Discussion

Robo4 stabilizes the VE-cadherin-mediated adherens junction and suppresses vascular permeability. It improves the survival of mice under severe infectious and systemic inflammation.^{[3,](#page-9-2)[6](#page-9-3)} Therefore, an understanding of the mechanism underlying the regulation of Robo4 expression and devising a strategy for controlling its expression is expected to be useful for the development

Figure 5. MS-275 promotes the transmigration of cells through the vasculature and endothelial monolayer. A) Effect of MS-275 on the extravasation of B16-F10 cells in the experimental metastasis model. A schedule for the treatment of MS-275 and injection of B16-F10 cells and for colony counting is indicated. Pictures are representative images of the lungs obtained from mice treated with the vehicle and MS-275. The graph indicates the number of colonies in the lungs from mice treated with the vehicle or MS-275 ($n = 5$). (B) Effect of MS-275 on the transmigration of B16-F10 cells through the MS1 cell monolayer. B16-F10 cells that transmigrated through the MS1 monolayer treated with or without MS-275 were quantified ($n = 4$). Data are expressed as the mean \pm s.e.m. Statistical significance was determined by the unpaired *t*-test (***p* < .01).

of future therapy. In the present study, we investigated the effects of HDAC inhibitors on Robo4 expression. We demonstrated that MS-275 suppresses Robo4 expression and induces endothelial hyperpermeability and cell extravasation. Mechanistically, MS-275-mediated inactivation of HDAC3 suppressed Robo4 expression by downregulating the expression of GABP. Thus, we successfully demonstrated a novel mechanism for the regulation of Robo4 expression by HDAC and its inhibitor and proposed a strategy for controlling vascular permeability using small molecules.

MS-275 inhibits the activity of HDAC1, HDAC2, and HDAC3, which are ubiquitously expressed in tissues.^{[15](#page-10-3)} HDAC1 and HDAC2 have almost identical protein structures that contain tandem phosphorylation sites and catalytic domains for deacetylation activity, and are localized only in the nucleus.¹⁶ In contrast, HDAC3 contains only one phosphorylation site and shuttles between the nucleus and the cytoplasm using the nuclear export signal.¹⁷ HDAC3 is exclusively recruited to the SMRT/NCoR complex, whereas HDAC1 and 2 are found in multiple corepressor complexes.^{[16](#page-10-4)} Thus, HDAC3 possesses a unique structure and function compared with HDAC1 and HDAC2. Consistent with this fact, siRNA-mediated knockdown of HDAC3, but not of HDAC1 and 2, was observed to suppress Robo4 expression by

downregulating the expression of GABP, indicating a unique function of HDAC3 in endothelial cells. However, it remains to be answered as to how MS-275 downregulates GABP by inhibiting the HDAC3 activity. In a previous study, it was suggested that GABPα expression is regulated by a protein complex containing GABP and $SP1¹⁸$ $SP1¹⁸$ $SP1¹⁸$ In addition, physical and functional interactions between SP1 and HDAC3 in gene regulation have been reported.^{[19,](#page-10-7)20} These findings indicated the possibility that GABPα expression is regulated by the SP1–HDAC3 complex and that MS-275 downregulates Robo4 expression by inhibiting the functioning of this complex. However, there are relatively few studies demonstrating the mechanisms for the regulation of GABP. The MS-275- and HDAC3-mediated gene regulation of GABP needs to be investigated in the future.

MS-275 has been shown to regulate the expression of both pro- and anti-inflammatory genes. 21 21 21 It activates NF-κB by augmenting its acetylation and nuclear translocation.^{[22](#page-10-10)} MS-275 also exerts an anti-tumor effect by targeting tumor and immune cells. In the present study, we identified endothelial cells as the novel target cell type for MS-275 and unraveled a unique effect of MS-275 on vascular permeability. We used an experimental metastasis model to readily evaluate the effect of MS-275 treatment on cell extravasation. However, this does not elucidate the effect of

Figure 6. Downregulation of Robo4 promotes vascular permeability and cell transmigration through the vasculature and endothelial monolayer. **A** Quantification of the number of colonies in the lungs of wild-type (Robo4^{+/+}) and Robo4-deficient (Robo4^{+/−} and Robo4^{-/-}) mice intravenously injected with B16-F10 cells (n = 5). Pictures are representative images of lungs from the wild-type and Robo4-deficient mice. (B) Vascular leakage in the lungs of wild-type or Robo4 knockout mice injected with B16-F10 cells ($n = 5$). (C) Effect of the knockdown of Robo4 on the transendothelial migration of tumor cells. Migration of B16-F10 cells through MS1 cells transfected with siRNA specific for Robo4 was quantified $(n = 6)$. (D) Effect of the knockdown of Robo4 on endothelial cell viability. WST-8 assays were performed using MS1 cells transfected with siRNA ($n = 6$). Data are expressed as the mean \pm s.e.m. Statistical significance was determined by Dunnett's test or the unpaired *t*-test (**p* < .05, ***p* < .01; N.S., not significant).

MS-275 on autonomous tumor metastasis because this model does not completely reproduce all the processes involved in metastasis, including the migration of tumor cells from the primal tumor into the blood. In fact, the antimetastasis effect of MS-275 was demonstrated in previous studies on the spontaneous metastasis model (subcutaneous injection model).^{[23](#page-10-11),24} In addition, we have previously shown that downregulation of Robo4 increases the transmigration of monocytic cells through the endothelial cell monolayer, 3 indicating that MS-275 also promotes the extravasation of immune cells, which may contribute to the anti-tumor effect of MS-

275. Furthermore, a recent study showed that Robo4 regulates the expression of α-SMA, Snail, and VE-cadherin, implying that Robo4 modulates endothelial–mesenchymal transition, which is associated with tumor metastasis.^{[25](#page-10-13),[26](#page-10-14)} Further investigations on the effect of MS-275 on tumor metastasis are warranted.

In summary, we have successfully demonstrated a novel mechanism for the regulation of Robo4 expression and vascular permeability via HDAC and its inhibitors. The results presented herein would be useful for developing therapies against severe infectious and systemic inflammatory diseases in the future by targeting vascular permeability.

Notes on contributors

T. Kashio, K. Shirakura, M. Kinoshita, N. Hino, S. Koyama, R., Suzuki, Y. Yoshioka, T. Aoshi, T. Doi, and Y. Okada designed and conceived the study. T. Kashio, M. Kinoshita, M. Morita, R. Ishiba, K. Muraoka, T. Kanbara, M. Tanaka, and R. Funatsu performed the experiments. T. Kashio, K. Shirakura, M. Kinoshita, N. Hino, S. Koyama, R. Suzuki, Y. Yoshioka, T. Aoshi, T. Doi, and Y. Okada interpreted the data. T. Kashio, K. Shirakura, and Y. Okada wrote the manuscript.

Disclosure statements

Y. Yoshioka is employed by the Research Foundation for Microbial Diseases of Osaka University. The remaining authors declare no conflicts of interest.

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