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Research Article

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# Liver acid sphingomyelinase inhibits growth of metastatic colon cancer

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**Acid sphingomyelinase (ASM) regulates the homeostasis of sphingolipids, including ceramides and sphingosine-1-phosphate (S1P). These sphingolipids regulate carcinogenesis and proliferation, survival, and apoptosis of cancer cells. However, the role of ASM in host defense against liver metastasis remains unclear. In this study, the involvement of ASM in liver metastasis of colon cancer was examined using *Asm*<sup>-/-</sup> and *Asm*<sup>+/+</sup> mice that were inoculated with SL4 colon cancer cells to produce metastatic liver tumors. *Asm*<sup>-/-</sup> mice demonstrated enhanced tumor growth and reduced macrophage accumulation in the tumor, accompanied by decreased numbers of hepatic myofibroblasts (hMFs), which express tissue inhibitor of metalloproteinase 1 (TIMP1), around the tumor margin. Tumor growth was increased by macrophage depletion or by *Timp1* deficiency, but was decreased by hepatocyte-specific ASM overexpression, which was associated with increased S1P production. S1P stimulated macrophage migration and TIMP1 expression in hMFs in vitro. These findings indicate that ASM in the liver inhibits tumor growth through cytotoxic macrophage accumulation and TIMP1 production by hMFs in response to S1P. Targeting ASM may represent a new therapeutic strategy for treating liver metastasis of colon cancer.**

## Introduction

Colon cancer, one of the most common malignancies, frequently metastasizes to the liver. Acid sphingomyelinase (ASM) is involved in various physiological cellular functions and diseases, including cancer (1), and hydrolyzes sphingomyelin into ceramide and phosphorylcholine. Ceramide, a bioactive mediator of numerous cellular functions, such as apoptosis and cell cycle regulation (2, 3), is in turn hydrolyzed by ceramidase into sphingosine, which is subsequently phosphorylated into sphingosine-1-phosphate (S1P) by sphingosine kinase (SphK). Although these sphingolipids are involved in colon carcinogenesis and colon cancer cell survival (4, 5), the roles of ASM and S1P in host antitumor defenses (i.e., inhibiting the progression of colon cancer metastasis to the liver) remain unclear.

Tumors contain stromal cells, such as immune cells and fibroblasts (6). Infiltrated tumor-associated macrophages (TAMs) have recently been reported to function as promoters of tumor progression (7–9), with several clinical studies demonstrating an association between the presence of TAMs and poor prognosis in various cancers (10). In contrast, opposing data have shown that the presence of TAMs is correlated with improved survival and that these cells have protective potential in colon cancer (11, 12). Although tumor growth-promoting TAMs have been observed to resemble regulatory M2 macrophages (13, 14), the density of classically activated M1 macrophages is positively correlated with the survival time of patients with non-small-cell lung cancer (15). These findings indicate that macrophages have contrasting roles in cancer, depending on their phenotype (13).

S1P is a ceramide-derived metabolite that is involved in various cellular functions and increases migration of macrophages (16, 17), resulting in macrophage recruitment. S1P released from apoptotic tumor cells induces a switch from the M1 to the M2

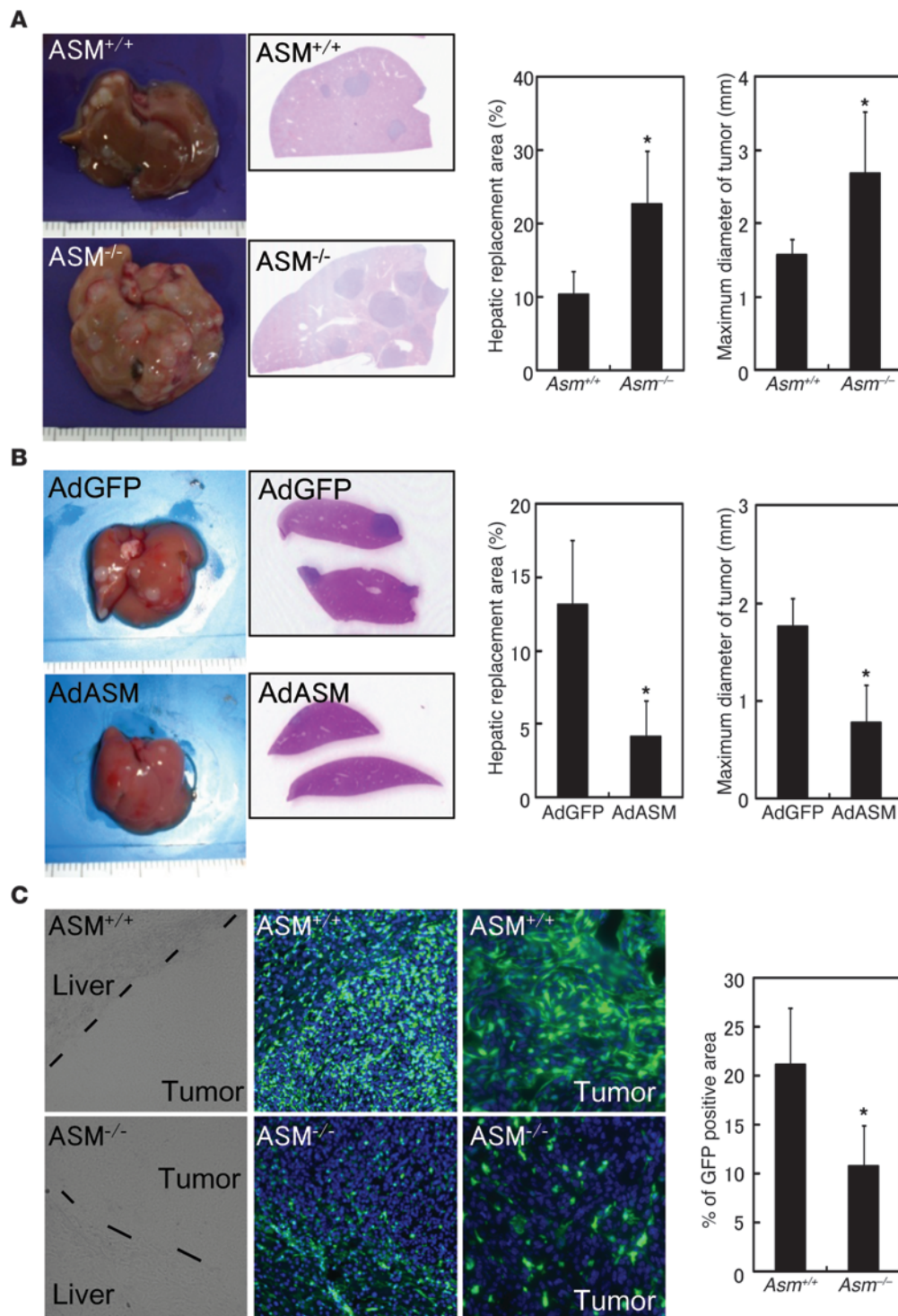
macrophage subtype (18). In contrast, the S1P analog FTY720 does not alter the ratio of M1 and M2 subtypes in mouse peritoneal macrophages (19). Because of these contrasting findings, the effect of S1P on macrophages remains controversial and poorly understood. Whereas MMP cleavage of the ECM, a primary barrier against tumor invasion produced by hepatic myofibroblasts (hMFs) in the liver, promotes cancer cell migration, invasion, and metastasis, MMP inhibitors reduce colon cancer metastasis to the liver (20). Specifically, overexpression of tissue inhibitors of metalloproteinases (TIMPs), endogenous inhibitors of MMPs, inhibits liver metastasis of colon cancer in animal models (21), whereas decreased TIMP1 expression in the liver results in progression of SV40T antigen-induced hepatocellular carcinoma (22) and increased metastatic colonization of T cell lymphoma (23). However, it has been reported that elevated TIMP levels are associated with cancer progression (24). Thus, although S1P stimulates hMF accumulation and TIMP1 induction in the liver (25, 26), the precise roles of S1P in tumor growth remain unclear. To attempt to clarify these precise roles, we here investigated the involvement of ASM in the progression of liver metastasis of colon cancer.

## Results

*Role of ASM deficiency and overexpression in the progression of metastatic liver tumors of colon cancer.* To determine the role of host ASM on metastatic liver tumor growth caused by colon cancer, we created metastatic liver tumors by intrasplenic injection with SL4 cells, a mouse colon cancer cell line, and examined expression of ASM and ceramide by immunostaining. ASM and ceramide expression increased in the liver cells around the tumors (Supplemental Figure 1A; supplemental material available online with this article; doi:10.1172/JCI65188DS1), which suggests that tumor cells stimulate ASM expression and promote its activity in liver cells. To examine the effect of ASM deficiency on metastatic liver tumors, the extent of tumor growth in *Asm*<sup>+/+</sup> and *Asm*<sup>-/-</sup> mice

**Conflict of interest:** The authors have declared that no conflict of interest exists.

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**Figure 1**

ASM deficiency increased, and ASM overexpression decreased, metastatic tumor growth in the liver. (A and B) *Asm*<sup>+/+</sup> and *Asm*<sup>-/-</sup> mice were intrasplenically injected with  $2 \times 10^4$  SL4 cells (A). Wild-type mice were infected with AdGFP or AdASM 24 hours before inoculation (B). Mice were sacrificed 14 days after inoculation, and livers were excised and photographed. Liver sections were stained with H&E (loupe magnification), and intrahepatic tumor load was presented as hepatic replacement area and maximum diameter, based on measurement of 3 nonsequential sections. (C) *GFP*<sup>+</sup>*Asm*<sup>+/+</sup> and *GFP*<sup>+</sup>*Asm*<sup>-/-</sup> mice were intrasplenically injected with SL4 cells. At 14 days after inoculation, GFP expression in the liver was assessed by fluorescent microscopy, measurement of the GFP<sup>+</sup> area was performed, and nuclei were stained with DAPI. Shown are bright field images at the border of the metastatic tumor and liver (left), GFP/DAPI merge in the same bright field images (middle), and higher-magnification views of GFP/DAPI merge in the tumor area (right). Original magnification,  $\times 200$  (left and middle);  $\times 400$  (right). Results are mean  $\pm$  SD of data obtained from at least 5 independent experiments. \**P* < 0.05, 2-tailed Student's *t* test.

was compared. No tumors were evident in livers without SL4 cell inoculation (Supplemental Figure 1B). After inoculation with SL4 cells, the hepatic replacement area, maximum tumor diameter, and liver weight were increased in *Asm*<sup>-/-</sup> compared with *Asm*<sup>+/+</sup> mice (Figure 1A and Supplemental Figure 1C). In contrast, a similar number of the initially transplanted cells was observed in the livers of *Asm*<sup>+/+</sup> and *Asm*<sup>-/-</sup> mice 6 hours after inoculation (Supplemental Figure 2A). These results indicate that colon cancer

cells grow more rapidly in the *Asm*<sup>-/-</sup> liver. Conversely, administration of adenovirus-expressing ASM (AdASM) to the mice prior to cancer cell inoculation inhibited tumor growth compared with adenovirus-expressing GFP (AdGFP) administration (Figure 1B). AdASM-derived GFP was expressed in the hepatocytes, but not in the tumor cells (Supplemental Figure 2B). In support of the results of our previous study (27), AdASM increased ASM activity in the liver (Supplemental Figure 2C). In addition, ASM-chimeric mice,



in which ASM was deficient only in bone marrow-derived cells, demonstrated tumor growth that was comparable to mice containing *Asm*<sup>+/+</sup> bone marrow cells (Supplemental Figure 3A). These results indicate that ASM in hepatocytes, but not in bone marrow-derived cells, is involved in the inhibition of tumor growth.

To explore the mechanisms of enhanced tumor growth in *Asm*<sup>-/-</sup> mice, we generated *GFP*<sup>+</sup>*Asm*<sup>-/-</sup> mice and examined the host-tumor interaction by inoculating *GFP*<sup>+</sup>*Asm*<sup>+/+</sup> and *GFP*<sup>+</sup>*Asm*<sup>-/-</sup> mice with SL4 cells. After inoculation, spindle-shaped host-derived *GFP*<sup>+</sup> cells, most of which expressed F4/80 (Figure 2A), were observed in the metastatic liver tumors, while fewer host-derived cell infiltrates were observed in the *Asm*<sup>-/-</sup> livers (Figure 1C). Spindle-shaped bone marrow-derived cells were also observed in the tumors of GFP-chimeric mice (expressing GFP only in bone marrow-derived cells; Supplemental Figure 3B), which suggests that these cells were F4/80<sup>+</sup> macrophages originating from the bone marrow. F4/80<sup>+</sup> cells were observed in the tumors; compared with the control livers, a smaller number of F4/80<sup>+</sup> cells was observed in the *Asm*<sup>-/-</sup> livers, while a greater number of these cells was present in AdASM-infected livers (Figure 2, B and C). The induction of classically activated M1 macrophage markers (*Cd11c*, *Il12p40*, and *Ifng*) by SL4 cell inoculation was more remarkable than that of *Cd163*, mannose receptor, and *Il10*, markers for regulatory M2 macrophages (Supplemental Table 1). Thus, these findings suggested that the accumulated macrophages were M1 dominant. In *Asm*<sup>+/+</sup> mice transplanted with *GFP*<sup>+</sup>*Asm*<sup>-/-</sup> bone marrow, the number of bone marrow-derived *GFP*<sup>+</sup>*Asm*<sup>-/-</sup> cells in the tumor was comparable to that of *GFP*<sup>+</sup>*Asm*<sup>+/+</sup> cells in the control *Asm*<sup>+/+</sup> mice transplanted with *GFP*<sup>+</sup>*Asm*<sup>+/+</sup> bone marrow (Supplemental Figure 3C). Therefore, it is likely that ASM in hepatocytes, but not in bone marrow-derived cells, contributes to the accumulation of macrophages in tumors.

To examine whether macrophage accumulation in tumors is involved in the inhibition of tumor growth, liposome-encapsulated alendronate was administered to the SL4 cell-inoculated mice to decrease the number of F4/80<sup>+</sup> cells in the tumors (Supplemental Figure 3D). Alendronate administration increased the severity of tumor growth by inducing macrophage depletion (Figure 2D), which is suggestive of macrophage involvement in tumor growth suppression. These data indicate that hepatocytic ASM leads to accumulation of antitumor macrophages in the metastatic liver tumors of colon cancer, thereby suppressing tumor growth.

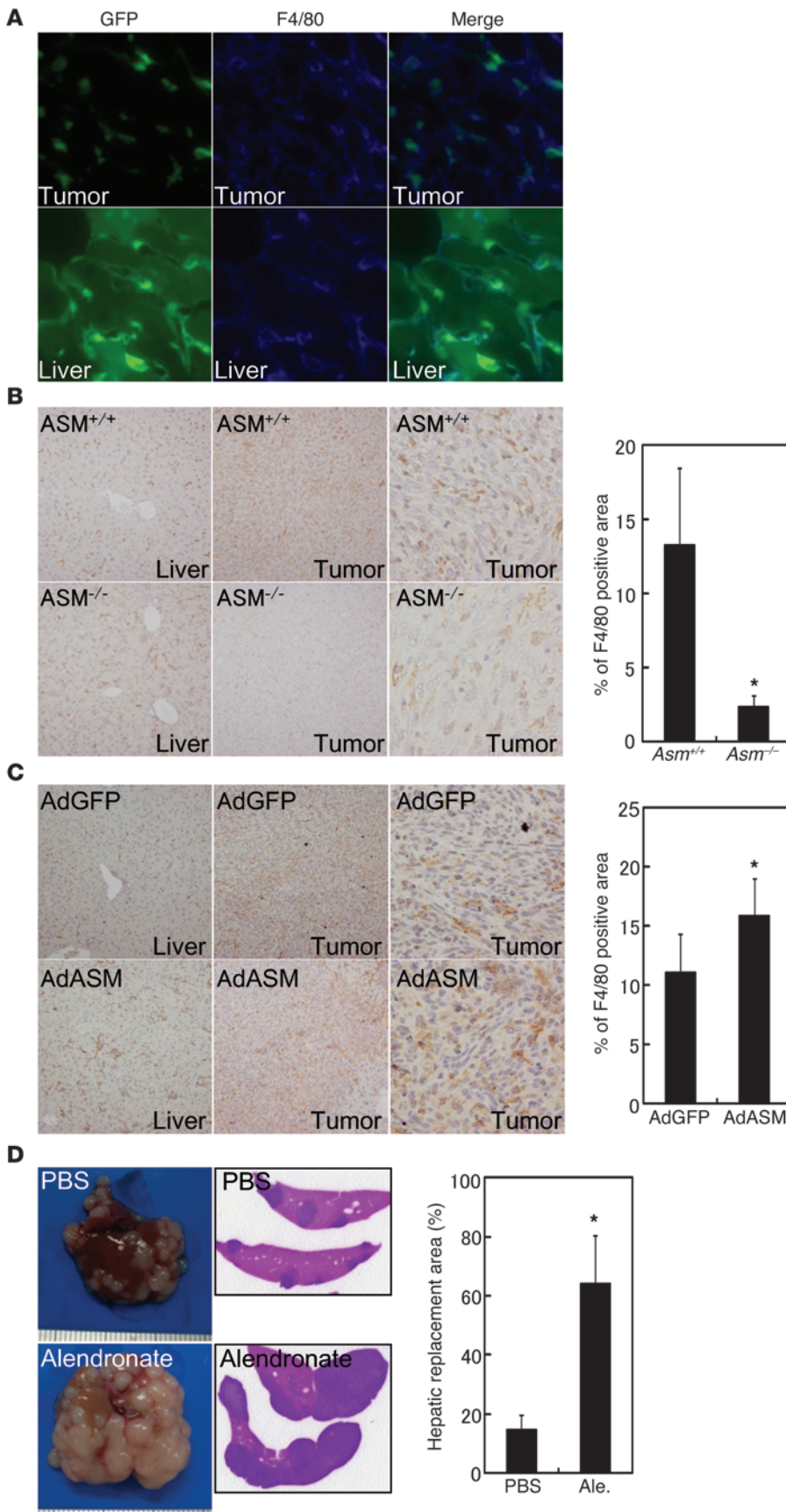
*Macrophage-induced hMFs accumulate around invasive margins of metastatic liver tumors.* Macrophages have been shown to exhibit antitumor potential as tumoricidal cells and major antigen-presenting cells (28–31). Liver macrophages also modulate the host immune response to cancer cells by releasing cytotoxic products and immune-stimulating factors, including IFN- $\gamma$ . In accordance with these findings, CD3<sup>+</sup> lymphocytes infiltrated into the tumors, and fewer CD3<sup>+</sup> cells were observed in *Asm*<sup>-/-</sup> and alendronate-treated mice (Supplemental Figure 4, A and B). Moreover, a NKT cell activator,  $\alpha$ -galactosylceramide, suppressed tumor growth, even in *Asm*<sup>-/-</sup> mice (Supplemental Figure 4C), which suggests that the antitumor immunity of NKT cells was not impaired in *Asm*<sup>-/-</sup> mice. Thus, NKT cells contributed to the antitumor defense mechanism.

Activated macrophages release various types of inflammatory cytokines and growth factors. Among these, TNF- $\alpha$  is thought to induce tumor necrosis. However, in the model used in this study, mRNA expression of *Tnfa* was comparable in SL4 cell-inoculated *Asm*<sup>-/-</sup> and *Asm*<sup>+/+</sup> livers (Supplemental Table 2), and tumor growth was not increased in *Tnfa*<sup>-/-</sup> mice (Supplemental Figure 4D), which

suggests that TNF- $\alpha$  is not involved in tumor development in *Asm*<sup>-/-</sup> mice. The reduced induction of *Il1b*, *Cxcl1*, and *Tgfb* mRNA expression by SL4 cell inoculation in *Asm*<sup>-/-</sup> compared with *Asm*<sup>+/+</sup> livers (Supplemental Table 2) can be explained by the reduction in the number of macrophages in the *Asm*<sup>-/-</sup> liver, as these factors are released from liver macrophages. The mRNA expression levels of *Des* (encoding desmin, a marker of hMFs), *Acta2* (encoding  $\alpha$ -SMA, a marker of activated hMFs), and *Col1a1* (encoding collagen  $\alpha$ 1[I], a product of activated hMFs) were also increased after SL4 cell inoculation in *Asm*<sup>+/+</sup> livers (Supplemental Table 2), which suggests that the metastatic tumors stimulated hMF proliferation and activation. In the SL4 cell-inoculated mice, the number of desmin<sup>+</sup> and  $\alpha$ -SMA<sup>+</sup> cells were increased around the invasive margins of the tumors in the liver (Figure 3). Desmin<sup>+</sup> cells were observed to express GFP in the *GFP*<sup>+</sup> mice, but not in the GFP-chimeric mice (data not shown), which suggests that the hMFs were derived from host, not from bone marrow. The increase of *Des*, *Acta2*, and *Col1a1* mRNA after SL4 cell inoculation was lower in *Asm*<sup>-/-</sup> than in *Asm*<sup>+/+</sup> livers (Supplemental Table 2). Moreover, whereas the number of desmin<sup>+</sup> and  $\alpha$ -SMA<sup>+</sup> cells was decreased by *Asm* deficiency (Figure 3A) and by macrophage depletion (Figure 3C), it was increased by ASM overexpression (Figure 3B). These findings suggest that increased macrophage accumulation in the metastatic liver tumor by ASM stimulates hMF accumulation and activation. Because the number of desmin<sup>+</sup> and  $\alpha$ -SMA<sup>+</sup> cells was inversely correlated with tumor growth, we concluded that hMFs may contribute to tumor suppression.

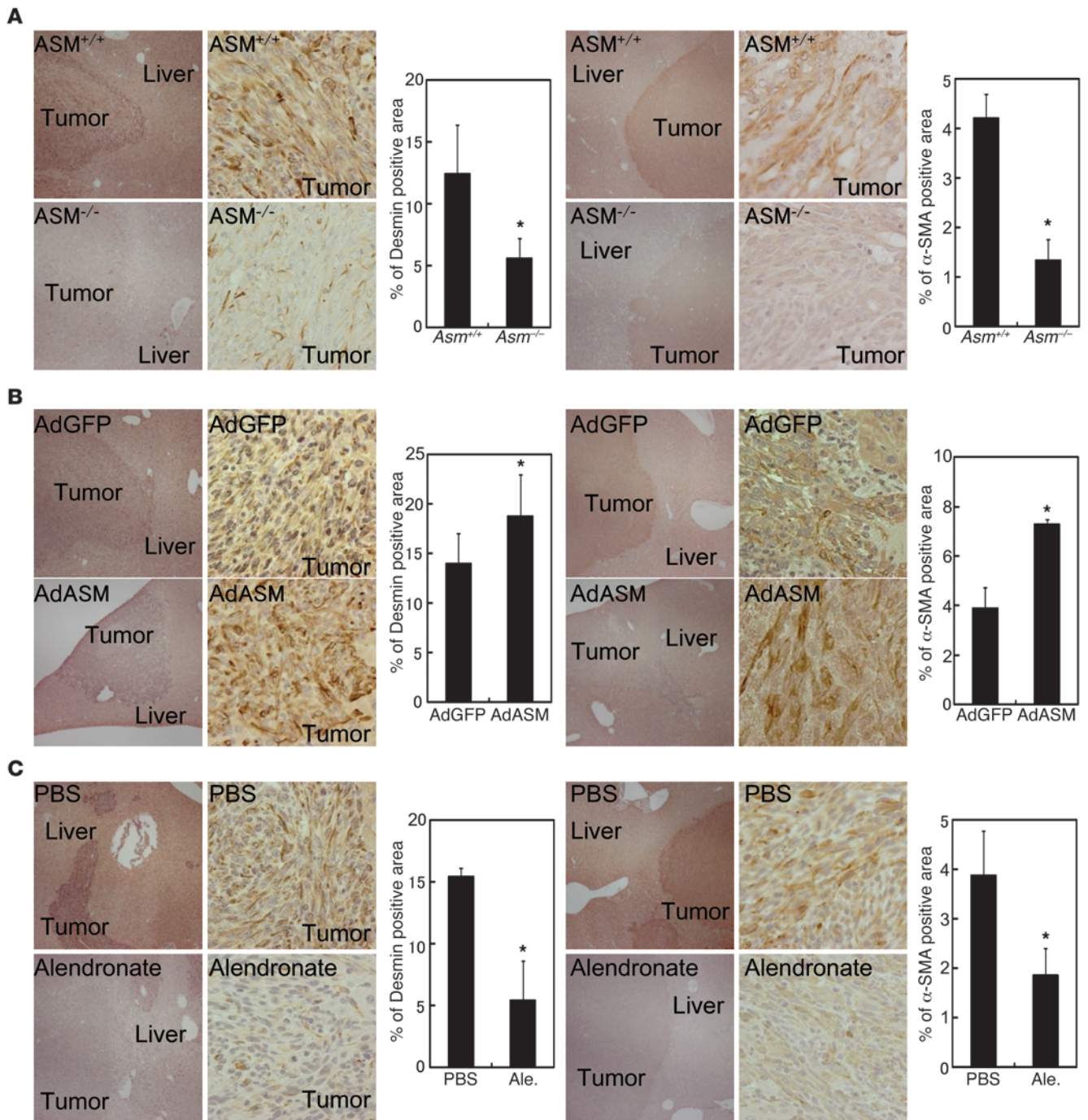
*Role of TIMP1 produced by hMFs in growth suppression of metastatic liver tumors.* Although it has been reported that hMFs contribute to cancer progression, the model used in this study indicated that increased hMF accumulation is negatively correlated with tumor growth. This finding led us to hypothesize that hMFs contain antitumor factors that are increased in number and magnitude by ASM overexpression and decreased by ASM deficiency. To test this hypothesis, we focused on examining TIMP1, which exhibits antitumor potential (21) and is expressed by hMFs (32). In the metastatic tumors, most of the desmin<sup>+</sup> cells showed double staining for TIMP1 (Figure 4A), which suggests that hMFs are TIMP1<sup>+</sup>. *Timp1* mRNA expression increased in *Asm*<sup>+/+</sup> livers by SL4 cell inoculation, an increase that was attenuated in *Asm*<sup>-/-</sup> livers (Figure 4B). Nevertheless, the expression levels of *Mmp2*, *Mmp9*, and *Mmp13* were comparable in *Asm*<sup>+/+</sup> and *Asm*<sup>-/-</sup> livers (Supplemental Table 2). The increased number of TIMP1<sup>+</sup> cells observed around the invasive margin of the tumors was attenuated by *Asm* deficiency as well as by macrophage depletion (Figure 4, C and E), whereas ASM overexpression further increased TIMP1<sup>+</sup> cell number (Figure 4D), similar to our findings with respect to hMF induction. These results, in addition to the increased tumor growth observed in *Timp1*<sup>-/-</sup> mice (Figure 4F), suggest that the increased expression of TIMP1 in hMFs by metastatic tumors is involved in the suppression of tumor growth.

*ASM suppression of tumor growth via S1P production.* In accordance with previous reports that ASM hydrolyzes sphingomyelin into ceramide, which is then hydrolyzed by ceramidase and phosphorylated by SphK to form S1P, we observed the level of S1P to increase in AdASM-infected livers (Figure 5A). Although exogenous administration of S1P did not affect *Tnfa*, *Il1b*, *Cxcl1*, *Tgfb*, *Il12p40*, or *Il10* mRNA expression in isolated peritoneal CD11b<sup>+</sup> macrophages (Supplemental Table 3), these cells showed increased migration toward S1P (Figure 5B), as previously reported regarding RAW 264.7 macrophages (33). These findings suggest that macrophage accumulation in tumors might be caused by S1P-induced macrophage recruitment. Although it



**Figure 2**

Macrophage depletion increased metastatic tumor growth in the liver. (A) GFP<sup>+</sup> mice were intrasplenically injected with  $2 \times 10^4$  SL4 cells. GFP fluorescence was visualized with fluorescent microscopy, and macrophages were detected in metastatic tumor and liver by immunofluorescent staining for F4/80 (blue). Original magnification,  $\times 400$ . (B and C) ASM<sup>+/+</sup> and ASM<sup>-/-</sup> mice (B) and AdGFP- and AdASM-infected wild-type mice (C) were intrasplenically injected with  $2 \times 10^4$  SL4 cells. Expression of F4/80 in the liver and metastatic tumor was examined by immunohistochemistry with an anti-F4/80 antibody to assess the number of macrophages. Original magnification,  $\times 100$  (left and middle);  $\times 400$  (right). (D) Wild-type mice were injected with SL4 cells and treated with PBS or alendronate before sacrifice 14 days after inoculation. Images of livers after excision and liver sections stained with H&E (loupe magnification) are shown. Intrahepatic tumor load is presented as hepatic replacement area, based on measurement of 3 nonsequential sections. Results are mean  $\pm$  SD of data collected from at least 5 independent experiments. \* $P < 0.05$ , 2-tailed Student's *t* test.

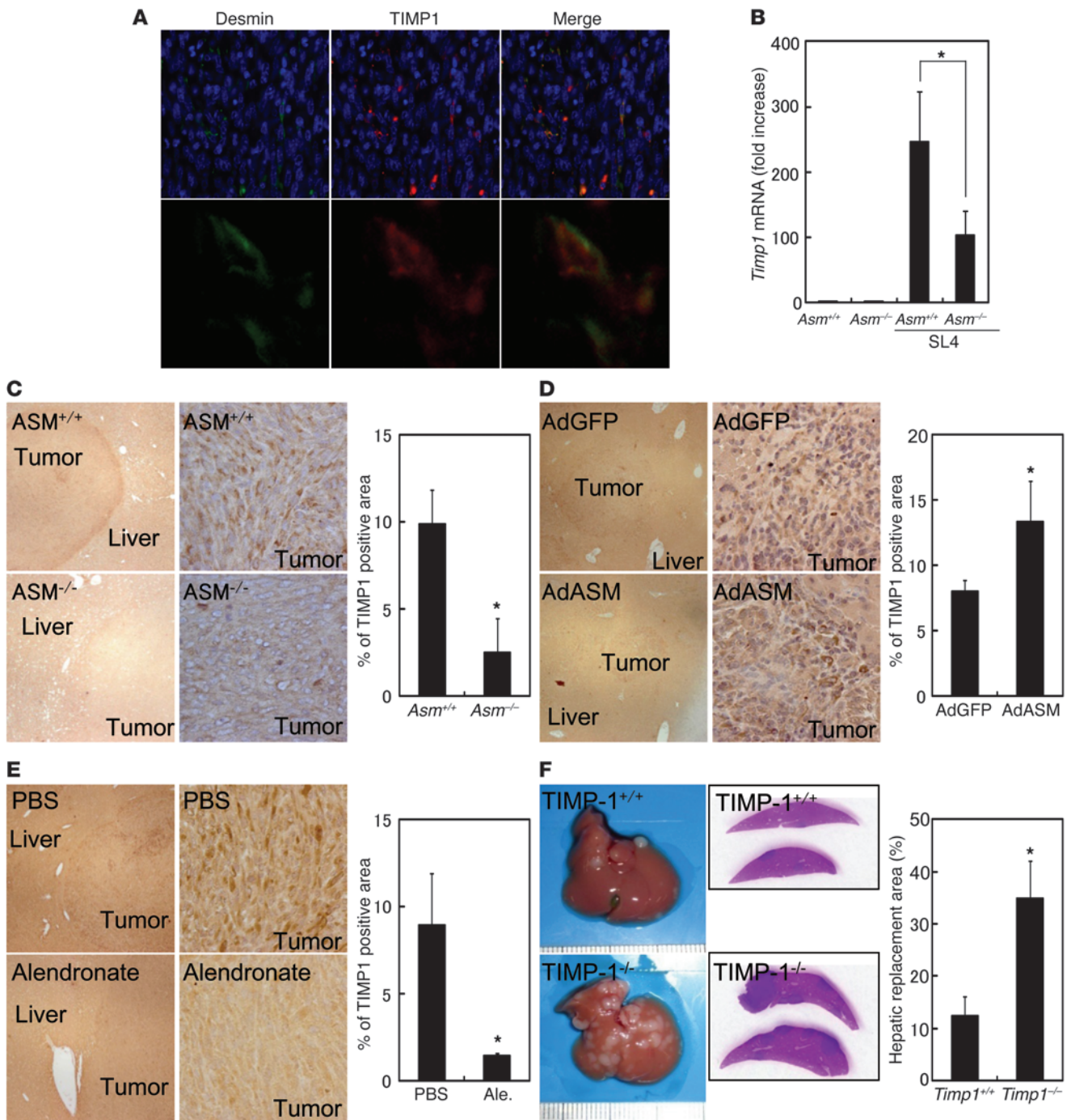


**Figure 3**

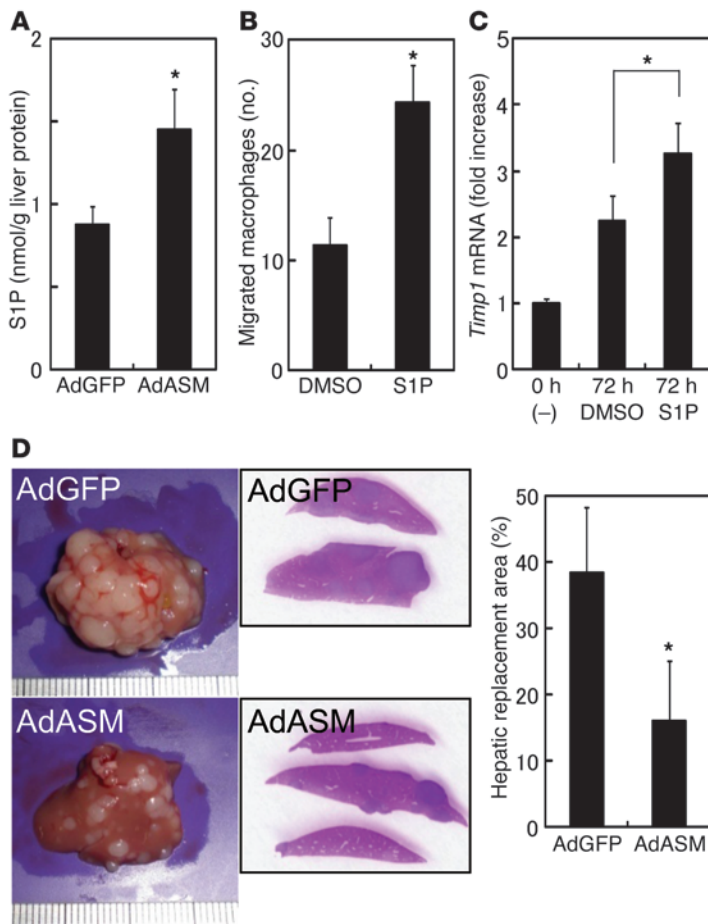
ASM deficiency or macrophage depletion decreased hMFs, and ASM overexpression increased hMFs, around the tumor invasive margin. *Asm*<sup>+/+</sup> and *Asm*<sup>-/-</sup> mice (A), AdGFP- or AdASM-infected wild-type mice (B), and wild-type mice treated with PBS or alendronate (C) were intrasplenically injected with 2 × 10<sup>4</sup> SL4 cells and sacrificed 14 days after inoculation. Expression of desmin (left) and α-SMA (right) around the tumor invasive margin was examined by immunohistochemistry, and measurement of immunostain-positive area was performed. Original magnification, ×40 (left); ×400 (right). Results are mean ± SD of data collected from at least 5 independent experiments. \**P* < 0.05, 2-tailed Student's *t* test.

has been reported that S1P induces a switch from the M1 to the M2 macrophage subtype (18), in the present study, S1P did not alter the mRNA expression levels of M1 and M2 markers, which suggests that S1P does not affect the macrophage phenotype. Instead, our observations that 1 μM S1P treatment for 72 hours increased *Timp1*

mRNA expression in isolated hMFs (Figure 5C) and that adenoviral overexpression of SphK1 (AdSphK) in hepatocytes prior to cancer cell inoculation inhibited tumor growth (Supplemental Figure 5A) suggest that ASM suppresses tumor growth via S1P production. The fold increase of *Timp1* mRNA induced by S1P did not seem to

**Figure 4**

TIMP1 deficiency increased metastatic tumor growth in the liver. (A) Wild-type mice were intrasplenically injected with  $2 \times 10^4$  SL4 cells and sacrificed 14 days after inoculation. Expression of desmin (green) and TIMP1 (red) around the invasive margin of the tumors was examined by immunofluorescent staining. Nuclei were stained with DAPI (blue, top). Original magnification,  $\times 400$  (top);  $\times 1,600$  (bottom). (B) *Asm*<sup>+/+</sup> and *Asm*<sup>-/-</sup> mice were intrasplenically injected with  $5 \times 10^5$  SL4 cells and sacrificed 7 days after inoculation. *Timp1* mRNA expression in the liver was determined by quantitative real-time RT-PCR. (C–E) *Asm*<sup>+/+</sup> and *Asm*<sup>-/-</sup> mice (C), AdGFP- or AdASM-infected wild-type mice (D), and wild-type mice treated with PBS or alendronate (E) were intrasplenically injected with  $2 \times 10^4$  SL4 cells and sacrificed 14 days after inoculation. TIMP1 expression around the tumor invasive margin was examined by immunohistochemistry, and measurement of TIMP1<sup>+</sup> area was performed. Original magnification,  $\times 40$  (left);  $\times 400$  (right). (F) *Timp1*<sup>+/+</sup> and *Timp1*<sup>-/-</sup> mice were intrasplenically injected with  $2 \times 10^4$  SL4 cells and sacrificed 14 days after inoculation. Images of livers after excision and liver sections stained with H&E (loupe magnification) are shown. Intrahepatic tumor load is presented as hepatic replacement area, based on measurement of 3 nonsequential sections. Results are mean  $\pm$  SD of data collected from at least 5 independent experiments. \* $P < 0.05$ , 2-tailed Student's *t* test.



**Figure 5**

S1P promoted macrophage migration and *Timp1* mRNA expression in hMFs. (A) Wild-type mice were infected with AdGFP or AdASM, and S1P was examined by MS analysis 3 days after infection. (B) After addition of CD11b<sup>+</sup> peritoneal macrophages and 1 μM S1P to the upper and lower chambers, respectively, of transwell plates, cells were incubated for 4 hours. The number of macrophages that migrated to the underside of the chamber was determined. (C) Primary rat hMFs were incubated with or without 1 μM S1P on plastic dishes for 72 hours before determination of *Timp1* mRNA levels by quantitative real-time RT-PCR. (D) Wild-type mice were intrasplenically injected with 2 × 10<sup>4</sup> SL4 cells, infected with AdGFP or AdASM 5 days after inoculation, and sacrificed 14 days after infection (i.e., 19 days after inoculation). Images of livers after excision and liver sections stained with H&E (loupe magnification) are shown. Intrahepatic tumor load is presented as hepatic replacement area, based on measurement of 3 nonsequential sections. Results are mean ± SD of data collected from 4 (A) or at least 5 (B–D) independent experiments. \**P* < 0.05, 2-tailed Student's *t* test.

be large (Figure 5C). However, a 48-hour treatment with TGF-β has previously been reported to induce an approximately 2-fold increase of *TIMP1* mRNA in cultured human hMFs (34). This suggests that the induction of *Timp1* mRNA by S1P was not a small change, and it is possible that the induction might be sufficient. 5 μM S1P induced *TIMP1* expression to a similar degree as that observed using 1 μM S1P (data not shown), which suggests that 1 μM S1P fully induces activation of S1P receptors.

As described above, ASM exerted an antitumor effect through macrophage accumulation and *TIMP1* production from hMFs via S1P. To evaluate AdASM administration as a form of treatment against metastatic liver tumors of colon cancer, AdASM was administered to mice 5 days after SL4 cell inoculation. AdASM suppressed tumor growth, even when it was administered after cell inoculation (Figure 5D). However, AdSphK administration did not suppress tumor growth (Supplemental Figure 5B), although the accumulation of F4/80<sup>+</sup> and desmin<sup>+</sup> cells was similar to that in AdASM-treated mice (Supplemental Figure 6A). In addition, accumulation of F4/80<sup>+</sup> and desmin<sup>+</sup> cells was observed to be equal, regardless of whether mice were infected with AdSphK before or after SL4 cell inoculation (data not shown). Adenovirus-mediated GFP expression was observed in the tumors when the virus was administered after cell inoculation (Supplemental Figure 6B), which suggests that the adenovirus infects not only hepatocytes, but also inoculated SL4 cells. The observation that AdSphK increased proliferation of cultured SL4 cells (Supplemental

Figure 6C) indicated that the antitumor effects of overexpressed SphK1 in the liver may be counteracted by the tumor-promoting effects of SphK1 on SL4 cell proliferation.

## Discussion

In the present study, we investigated the contribution of ASM to the suppression of tumor growth in liver metastasis of colon cancer cells. Our results indicated that liver ASM inhibits tumor growth through S1P formation and subsequent macrophage accumulation and *TIMP1* production from hMFs. These results suggest novel therapeutic possibilities for treating metastatic liver tumors of colon cancer.

ASM in cancer cells is involved in cell death and plays an important role in the host response to a variety of anticancer treatments, including those for colon cancer (1). In the present study, metastatic tumors of colon cancer cells stimulated ASM expression and ceramide in liver cells. Moreover, host ASM deficiency increased, while overexpression of ASM in the liver decreased, the growth of liver tumors after inoculation with colon cancer cells, which suggests that ASM in host cells contributes to antitumor defense. Among the roles that ASM plays in liver cells, ASM in hepatocytes stimulates glucose uptake, resulting in improvement of glucose tolerance in mice through S1P production (27). ASM has pleiotropic signaling functions related to posttranslational processing of ASM, and secretory and lysosomal ASM (35) may have distinct functions. Although it is unclear which type of ASM was





involved in S1P production in our study, hepatocytes represented one source of S1P, as the S1P elevation by AdASM was inhibited in both SphK1-deficient livers and SphK1-deficient primary cultured rat hepatocytes (data not shown). Previous studies have shown that the generated S1P can be secreted and act as a ligand for S1P receptors (S1PRs; refs. 27, 36, 37), while the S1P/S1PR1 axis controls the trafficking and migration of immune cells, including macrophages (16, 17, 36). In accordance with these prior reports, S1P increased the migration of peritoneal macrophages *in vitro*, with less and more macrophage accumulation, respectively, in the tumors of *Asm*<sup>-/-</sup> and AdASM-infected mice compared with controls. These findings indicate that ASM in hepatocytes leads to macrophage accumulation in tumors via S1P production and that this process exerts an antitumor effect, in contrast to macrophage depletion. Although it has been reported that TAMs, which resemble regulatory M2 macrophages (13, 14), function as promoters of tumor progression (7–9), the macrophages in the tumors inoculated with SL4 colon cancer cells in this study were M1 dominant. It has been proposed that a gradual shift of TAM polarization from the M1 to the M2 subtype occurs during different stages of tumor progression, due to dynamic changes in the tumor microenvironment (38), and is paralleled by the gradual inhibition of NF- $\kappa$ B (39). In this study, we observed immunofluorescent staining of NF- $\kappa$ B p65 in the nuclei of F4/80<sup>+</sup> cells in the tumors (data not shown), which suggests that NF- $\kappa$ B is activated in these macrophages. This finding may indicate that in metastatic liver tumors characterized by rapid growth, such as by increased numbers of SL4 cells (40), the accumulated TAMs may retain an M1 phenotype that displays a cytotoxic proinflammatory phenotype. Thus, the number of macrophages present in the tumor is inversely correlated with tumor growth.

Myofibroblasts represent another important component of the tumor stroma; these cells are driven by migration and proliferation-stimulating factors released from cancer cells. Whereas some studies have reported a supportive role for myofibroblasts in cancer cell development (41) and a direct correlation between their presence and poor prognosis (42), others have reported a protective role of myofibroblasts against cancer cells (43, 44), with p53 in stromal fibroblasts demonstrated to inhibit tumor growth and malignant progression (45). In the current study using SL4 cell inoculation, hMF accumulation was found to be dependent on macrophage accumulation and inversely correlated with tumor growth. Specifically, hMF accumulation was increased only around the invasive margins of the tumors, and no hMF infiltration or collagen deposition was observed in the central part of tumors (data not shown). These findings indicate that the supportive effects of hMFs in tumor growth may be limited. However, the observation of enhanced tumor growth in *Timp1*<sup>-/-</sup> mice indicates that TIMP1 produced from hMFs, stimulated by S1P, might inhibit tumor growth through inhibition of ECM degradation. Although previous studies reported a protumor role for TIMP1, TIMP1 staining was observed only around the tumor invasive margin in the current study, which indicates that the supportive effects of TIMP1 in tumor growth may also be limited.

Whereas ceramide acts as a signaling molecule in the induction of cell responses, such as apoptosis and growth arrest (3, 46, 47), S1P acts as a potent mitogen for a variety of cell types (48), with SphK1/S1P contributing to colon carcinogenesis (49). Ceramide levels in colon carcinoma tissue are significantly lower than those in normal tissue (50), reflecting the fact that the dynamic balance

between intracellular levels of ceramide and S1P (the “ceramide/S1P rheostat”) determines cell growth and cell death in colon cancer. In the present study, adenoviral overexpression of SphK1 in the liver inhibited tumor growth when colon cancer cells were inoculated after infection. In contrast to AdASM, AdSphK did not exert an inhibitory effect when administered after cell inoculation, which suggests that its inhibitory effect on existing tumors is less than that of AdASM. The observation that AdSphK increased the proliferation of cultured SL4 cells suggests that it decreased the ceramide/S1P ratio in the cancer cells, which may promote their proliferation. In contrast, AdASM might increase the ceramide/S1P ratio, which may inhibit SL4 cell proliferation. We therefore conclude that AdASM, but not AdSphK, may be a useful candidate for gene therapy against metastatic liver tumors of colon cancer.

In *Asm*<sup>-/-</sup> mice, B16C2M melanoma cells formed small metastatic lesions in the liver (4 of 9, 44.4%) whereas none of the *Asm*<sup>+/+</sup> mice developed metastases (0 of 11, 0%; Supplemental Figure 7), which suggests that the antitumor effect of ASM is not specific for colon cancer cells. However, use of the *Asm* knockout may have secondary effects due to the severe lipid storage defect in these mice, particularly in macrophages. *Asm*<sup>-/-</sup> mice are known to have severe immune defects (51, 52) and display severe neurocutaneous disease by 8 weeks of age. Thus, *Asm*<sup>-/-</sup> mice may have various abnormalities in the liver as a normal pathology. In our study, the livers of *Asm*<sup>-/-</sup> mice without injection of SL4 cells did not show the nodular appearance. In addition, liver weight of *Asm*<sup>-/-</sup> mice without injection of SL4 cells was comparable to that of *Asm*<sup>+/+</sup> mice. Thus, it seems unlikely that the changes are due to normal *Asm*<sup>-/-</sup> mouse pathology. In addition, besides ASM, ceramide synthesized by the *de novo* pathway can be involved in TIMP1 regulation. These data do not provide direct evidence that S1P production in hepatocytes recruits macrophages, and experiments using mice with conditional knockout of SphK are also required to confirm the involvement of SphK and S1P in macrophage recruitment and TIMP1 regulation. Moreover, the mechanism by which tumor cells stimulate ASM expression in hepatocytes remains unclear, and there is no direct evidence of a positive correlation between ASM activity in hepatocytes and TIMP1 expression in hMFs; further studies are needed to resolve these uncertainties. In conclusion, we found here that ASM in hepatocytes inhibited tumor growth via S1P formation and subsequent cytotoxic macrophage accumulation. This S1P formation in turn stimulated TIMP1 production by hMFs, leading to tumor suppression. Thus, targeting ASM may represent a new therapeutic strategy for treating liver metastasis of colon cancer.

## Methods

**Animals.** *Asm*<sup>-/-</sup> mice (C57BL/6 background; ref. 47) were bred for use in this study. *GFP*<sup>+</sup> mice (C57BL/6 background), which express EGFP ubiquitously under the CAG promoter (chicken  $\beta$ -actin promoter, rabbit  $\beta$ -globin poly A, CMV-IE enhancer), were obtained from Riken Bio Resource Center. *Timp1*<sup>-/-</sup> mice (C57BL/6 background) were obtained from Jackson Laboratory. Male Wistar rats and female C57BL/6J wild-type mice were obtained from Japan SLC. *GFP*<sup>+</sup>*Asm*<sup>-/-</sup> mice were generated for this study.

**Cell culture.** A red fluorescent protein-expressing mouse colon adenocarcinoma cell line (SL4; Anti-Cancer Japan, Osaka, Japan) was maintained as a monolayer culture in RPMI-1640 (Invitrogen) containing 10% FBS supplemented with penicillin and streptomycin (Invitrogen). Cells were harvested using trypsin and EDTA, washed with PBS, and then resuspended in serum-free RPMI-1640 ( $2 \times 10^5$  or  $5 \times 10^6$  cells/ml).



**Liver metastasis model.** Female wild-type (C57BL/6J) mice aged 8–10 weeks were divided into 2 groups. After a small incision was made under anesthesia to expose the spleen, 0.1 ml of a viable cell suspension containing  $5 \times 10^5$  cells/mouse was injected into the spleen of group 1, and the same suspension containing  $2 \times 10^4$  cells/mouse was injected into group 2. Mice were sacrificed on day 7 (group 1) and day 14 (group 2) after cell inoculation. The livers of group 1 were immediately removed, washed in ice-cold PBS, and weighed, after which a part of the dissected liver tissue was frozen in liquid nitrogen. For assessment of liver metastatic tumors in group 2, the intrahepatic tumor was scored as the hepatic replacement area and maximum diameter of tumor, based on examination of 3 nonsequential H&E-stained sections using ImageJ software (NIH). For detection of GFP fluorescence, livers were first perfused in situ with 0.5 mM EGTA containing salt solution to remove peripheral blood cells. The livers were then perfused with 4% paraformaldehyde, fixed with 4% paraformaldehyde for 24 hours, soaked in 15% sucrose in PBS for 12 hours, soaked in 30% sucrose for 24 hours at 4 °C under constant agitation, and then embedded in OCT compound to cut sections on a cryostat. The GFP<sup>+</sup> area was determined using ImageJ software and shown as a percentage of the total section area.

**Histological analysis.** F4/80, desmin,  $\alpha$ -SMA, and TIMP1 were stained with anti-F4/80 (Santa Cruz Biotechnology), anti-desmin (Lab Vision), anti- $\alpha$ -SMA (clone 1A4; Sigma-Aldrich), and anti-TIMP1 (R&D Systems) antibodies, respectively, using Vectastain Elite ABC Kit or M.O.M Immunodetection Kit (Vector Laboratories). Diaminobenzidine tetrahydrochloride was used as peroxidase substrate, and sections were counterstained with hematoxylin. The immunostain-positive area was determined using ImageJ software and shown as a percentage of the total section area. In some experiments, fluorescent dye-labeled secondary antibodies (Alexa Fluor 350 anti-rat for F4/80, Alexa Fluor 488 anti-rabbit for desmin, and Alexa Fluor 594 anti-goat for TIMP1) were used for detection of primary antibodies.

**Depletion of macrophages.** Liposome-encapsulated alendronate (Sigma-Aldrich) was generated as previously reported (53). The alendronate was injected in the mice 4 times: 1 day before cell inoculation and 4, 7, and 10 days after. Liposome-encapsulated PBS was injected into the mice as a control.

**Recombinant adenoviruses.** Mice were infected with 2 adenoviruses ( $5 \times 10^8$  pfu/mouse) by intravenous injection 1 day before or 5 days after cell inoculation. Of the 2 adenovirus-5 variants, AdGFP expresses GFP, whereas AdASM expresses both ASM and GFP, because it contains a GFP sequence (27).

**Mass spectrometric (MS) analysis of S1P.** Electrospray ionization MS/MS analysis was performed using a Thermo-Finnigan TSQ 7000 Triple Quadrupole Mass Spectrometer (GenTEch Scientific Inc.) in multiple-reaction monitoring positive ionization mode, as previously reported (54). S1P level was normalized by total protein per sample.

**Macrophage migration assay.** For isolation of peritoneal macrophages, wild-type mice were injected intraperitoneally with cold normal saline. Peritoneal macrophages were purified using MACS Cell Separation Columns with CD11b microbeads (Miltenyi Biotec) from the lavage fluid. The cell migra-

tion assay was performed using 24-well transwell plates (5- $\mu$ m pore size; Costar). The peritoneal macrophages ( $1 \times 10^5$  cells) in DMEM containing 1% BSA were loaded into the upper chambers, while the lower chambers were filled with medium with or without 1  $\mu$ M S1P (Matreya LLC). Cells were incubated for 4 hours, then the migrated macrophages on the underside of the chambers were fixed and stained with Giemsa solution (Wako). Migration was expressed as the number of cells per field.

**Quantitative real-time RT-PCR.** Rat primary liver hMFs were isolated as previously described (25). The cells were cultured in DMEM with 10% FBS on plastic dishes for 4 hours and administered S1P or vehicle (1  $\mu$ M), and RNA was extracted 0 or 72 hours after S1P stimulation. The RNeasy and DNase Kits (Qiagen) was used for RNA extraction from liver tissue and cultured cells, and the High-Capacity cDNA Reverse Transcription Kit (Applied Biosystems) was used for reverse transcription. Quantitative real-time RT-PCR was performed using the SYBR Premix Ex Taq (Takara) for *Timp1* (forward, TGGGGAACCCATGAATTTAG; reverse, TCTGGCATCCTCTTGTTGC) and probe and primer sets (Applied Biosystems) for 18S with LightCycler 480 (Roche Applied Science). The changes were normalized based on 18S rRNA values.

**Statistics.** Data are expressed as mean  $\pm$  SD of data collected from 5 independent experiments. Data between groups were analyzed by 2-tailed Student's *t* test. A *P* value less than 0.05 was considered significant.

**Study approval.** All experiments were conducted in accordance with the institutional guidelines of Gifu University and were approved by the animal research committee and the committee on living modified organisms of Gifu University.

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