1	Cell proliferation effects of S-allyl-L-cysteine are associated with phosphorylation of
2	Janus kinase 2, insulin-like growth factor type-I receptor tyrosine kinase, and
3	extracellular signal-regulated kinase 2 in primary cultures of adult rat hepatocytes
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5	Hajime Moteki,* Masahiko Ogihara, and Mitsutoshi Kimura
6	
7	Department of Clinical Pharmacology, Faculty of Pharmaceutical Sciences, Josai
8	University; 1-1 Keyakidai, Sakado City, Saitama 350-0295, Japan
9	
10	*Correspondence: Department of Clinical Pharmacology, Faculty of Pharmaceutical
11	Sciences, Josai University; 1-1 Keyakidai, Sakado City, Saitama 350-0295, Japan.
12	Tel/Fax: +81-049-271-8078, E-mail: hmoteki@josai.ac.jp
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25 Abstract

26 The cell proliferation effect of S-allyl-L-cysteine (SAC) and its mechanisms were 27 examined in primary cultures of adult rat hepatocytes. In serum-free cultivation, SAC (10⁻ 28 ⁶ M)-stimulated hepatocytes showed significant proliferation compared to control at 5-h 29 culture; the effect was dependent on the culture time and the dose of SAC (EC₅₀ value 8.58×10^{-8} M). In addition, SAC-stimulated hepatocytes significantly increased mRNA 30 31 expression levels of *c-Myc* and *c-Fos* at 1 h and *cyclin B1* at 3.5 and 4 h, respectively. In 32 contrast, alliin and allicin, structural analogs of SAC, did not show these effects observed 33 with SAC. The SAC-induced hepatocyte proliferation effects were completely suppressed 34 by monoclonal antibodies against growth hormone receptor and insulin-like growth factor 35 type-I (IGF-I) receptor, respectively. Furthermore, the Janus kinase 2 (JAK2) inhibitor TG101209, phospholipase C (PLC) inhibitor U-73122, IGF-I receptor tyrosine kinase 36 37 (RTK) inhibitor AG538, PI3 kinase inhibitor LY294002, MEK inhibitor PD98059, and 38 mTOR inhibitor rapamycin completely suppressed the SAC-induced hepatocyte 39 proliferation. JAK2 (p125 kDa) phosphorylation in cultured hepatocytes peaked 5 min after SAC stimulation. SAC-induced IGF-I RTK (p95 kDa) and ERK2 (p42 kDa) 40 41 phosphorylation had slower rises than JAK2, peaking at 20 and 30 min, respectively. 42 These results indicate that SAC promoted cell proliferation by growth hormone 43 receptor/JAK2/PLC pathway activation followed by activation of the IGF-I 44 RTK/PI3K/ERK2/ mTOR pathway in primary cultures of adult rat hepatocytes.

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46 Keywords

47 S-Allyl-L-cysteine (SAC), proliferation, Janus kinase 2 (JAK2), Insulin-like growth

48 factor type I (IGF-I), Extracellular signal-regulated kinase 2 (ERK2)

49 **1. Introduction**

The liver has a self-regenerative function called liver regeneration, which causes rapid regeneration following liver damage or surgical resection (Fausto, 2000; Michalopoulos and DeFrances, 1997). The mitogen-activated protein (MAP) kinase pathway is a series of phosphorylation cascade reactions induced by the activation of receptor tyrosine kinase (RTK) during liver regeneration. This pathway ultimately activates a serine/threonine kinase called extracellular signal-regulated kinase (ERK) that promotes cell proliferation and differentiation (Fausto et al., 2006; Morrison, 2012).

57 S-Allyl-L-cysteine (SAC) is a water-soluble compound in which an allyl group is 58 bonded to the sulfur atom of cysteine. SAC is contained in a minuscule quantity in fresh 59 garlic, but its content increases up to about 36 times during aging of garlic for 3 months 60 (Colín-González et al., 2012). In the garlic aging process, SAC is biosynthesized from γ -61 glutamyl-S-allyl cysteine by a γ -glutamyl transferase. The synthesized SAC is oxidized 62 to alliin (S-allyl-L-cysteine sulfoxide), and it is hydrolyzed to allyl sulfenic acid 63 (Borlinghaus et al., 2014; Colín-González et al., 2012). Allicin (diallylthiosulfinate) is 64 produced by the condensation of this allyl sulfenic acid (Fig. 1). There is also a pathway 65 in which alliin and allicin are synthesized without going through SAC (Amagase et al., 2001). 66

Extracted substances of aged garlic and SAC have an antioxidant effect based on the removal of active oxygen or activation of nuclear factor-erythroid 2-related factor 2 (Nrf2) (Colín-González et al., 2015, 2012; Kalayarasan et al., 2008). Therefore, SAC is used as a supplement in Japan because it is effective for hypertension and hypercholesterolemia (Colín-González et al., 2012; Kim et al., 2006; Sohn et al., 2012). In addition, pharmacological effects of SAC other than these are becoming clear, and one

73 of them is a cell proliferation effect. For example, SAC has been demonstrated to promote 74 neuroblast division and differentiation by increasing the expression of serotonin 5-HT_{1A} 75 receptors in the dentate gyrus of mice (Nam et al., 2011). Furthermore, we have 76 previously reported that SAC, but not S-methyl-L-cysteine or cysteine, accelerated liver 77 regeneration by promoting DNA synthesis in remnant liver and restoring liver function in two-thirds partially hepatectomized rats (Kurihara et al., 2020). However, it is not clear 78 79 by what mechanism of action SAC exerts a cell proliferation-promoting effect on hepatic 80 parenchymal cells.

81 The aim of this study was to examine the detailed signal transduction pathways of cell 82 proliferation induced by SAC in primary cultures of adult rat hepatocytes. Thus, the cell 83 proliferation effect of SAC was compared to that of alliin or allicin by measuring the 84 number of hepatocyte nuclei, the progression of S-phase (DNA synthesis phase) in the 85 cell cycle, and the expression levels of genes, such as c-Myc, c-Fos, and cyclin B1 86 associated with liver regeneration. Moreover, the activity of growth-related signaling 87 elements such as RTK or MAPK involved in hepatocyte proliferation was measured using 88 various inhibitors of SAC-induced cell proliferation, and the mechanism of SAC on cell 89 proliferation in primary cultured hepatocytes was considered.

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91 **2. Materials and Methods**

92 2.1 Animals

Male Wistar rats weighing 180-200 g were obtained from Sankyo Labo Service Corp. (Tokyo, Japan). The pre-experimental acclimatization was set for 3 days, and during this period, rats had *ad libitum* access to food and water. The handling of all rats used in this study complied with the Guidelines for the Care and Use of Laboratory Animals of Josai 97 University (Nos. JU 20035 and JU 21035).

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99 **2.2 Isolation and culture of hepatocytes**

100 Rat hepatocytes were isolated and purified by two-step in situ collagenase perfusion 101 according to the method of Seglen (Seglen, 1975). In brief, under sodium pentobarbital 102 (45 mg/kg, i.p.) anesthesia, cannulation to the portal vein was performed. Then, the liver was refluxed in Ca²⁺-free Hanks as perfusion buffer for 10 min at 37 °C. Next, the solution 103 104 was changed to perfusion buffer containing 0.025% collagenase Type II (Worthington 105 Biochemical Corp., Freehold, NJ, USA.) and 5.0 mM CaCl₂ refluxed for 10-11 min. 106 Isolated hepatocytes were used at more than 96% cell viability by trypan blue exclusion. 107 Hepatocytes $(3.3 \times 10^4 \text{ cells/cm}^2)$ were plated and cultured for 3 h with Williams' medium 108 E (Sigma-Aldrich Co., St. Louis, MO, USA) containing newborn bovine serum (Thermo Fisher Scientific, Inc., Waltham, MA, USA.) and 10⁻¹⁰ M dexamethasone. After a 3-h 109 110 attachment period, the medium was exchanged to serum-free medium with various 111 reagents. The reagents added to hepatocytes were SAC, alliin, allicin (Tokyo Chemical 112 Industry Co., Tokyo, Japan), and growth-related signal-transducing factor inhibitors or 113 monoclonal antibodies. The inhibitors used were AG538 (Blum et al., 2003), LY294002 114 (Vlahos et al., 1994), PD98059 (Li et al., 2008), rapamycin (Dixon et al., 1999), 115 TG101209 (Pardanani et al., 2007), SH-4-54 (Haftchenary et al., 2013), and U-73122 116 (Thompson et al., 1991), which were obtained from Sigma-Aldrich Co. The monoclonal 117 antibodies used were against growth hormone receptor, insulin-like growth factor (IGF)-118 I receptor, epidermal growth factor (EGF) receptor, and hepatocyte growth factor (HGF) 119 receptor, which were obtained from Santa Cruz Biotechnology (Dallas, TX, USA).

121 **2.3** Counting the number of nuclei and hepatocytes

122 The cell proliferation effect of SAC was evaluated by measuring the number of 123 hepatocyte nuclei using a slight modification of the method of Nakamura (Nakamura et 124 al., 1983). Briefly, hepatocyte nuclei isolated in 0.1% Triton X-100 containing 0.1 M citric 125 acid were stained with 0.3% trypan blue, and the number of nuclei was measured with a 126 hemocytometer. In this study, the cell proliferation effect was evaluated by measuring the 127 number of nuclei rather than the number of cells, because the attachment between plates 128 or cell-cell contact is tight in hepatocytes, and a single cell cannot be isolated 129 quantitatively.

The numbers of total hepatocytes per 0.01 cm^2 (namely, $\times 10^2 \text{ cells/cm}^2$) were counted in a phase-contrast microscopic image. The values were measured from cells in microscopic images taken at any three locations, and the average was taken as one example.

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135 2.4 RNA extraction and quantification of *c-Myc*, *c-Fos*, and *cyclin B1* mRNA

136 The mRNA expression levels of *c-Myc*, *c-Fos*, and *cyclin B1* in cultured hepatocytes 137 were quantified using the TaqMan assay. The total RNA of the cultured hepatocytes was 138 extracted with Direct-zol RNA Miniprep (Zymo Research Co., Tustin, CA, USA), and 139 then a high-capacity cDNA reverse transcription kit (Applied Biosystems, Foster City, 140 CA, USA) was used for reverse transcription from the obtained RNA to cDNA. Next, 5 141 µg/µL of cDNA mixed with Fast Advanced Master Mix (Applied Biosystems) and 142 TaqMan probe (*c-Myc*, *c-Fos*, *cyclin B1*, or glyceraldehyde 3-phosphate dehydrogenase 143 [GAPDH]) were put in a StepOne Real-Time PCR System (Thermo Fisher Scientific, 144 Inc.), and each mRNA expression level was determined. GAPDH mRNA expression 145 levels were used as an endogenous control, and relative mRNA expression levels were 146 calculated by the $\Delta\Delta$ Ct method (Livak and Schmittgen, 2001).

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148 **2.5 Detection of the S-phase and the G₀/G₁-phase of the cell cycle**

149 DNA synthesis was determined by detecting the S-phase of the cell cycle. The S-phase 150 and the G_0/G_1 -phase in cultured hepatocytes were measured with a MuseTM cell analyzer, 151 using a slight modification to the manufacturer's instructions (Merck Millipore, 152 Darmstadt, Germany) (Kimura et al., 2014). Briefly, nuclei were isolated by lysing 153 hepatocytes as described in section 2.3 above. The isolated nuclei were mixed with 5% 154 propidium iodide (Fujifilm Wako Pure Chemical Corp., Tokyo, Japan) and incubated for 15 min. The Muse[™] cell analyzer is a cell analysis apparatus-based flow cytometer, and 155 156 the cell cycle, including the S-phase or G_0/G_1 -phase, was measured by this device.

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2.6 Measurement of phosphorylated p125 kDa Janus kinase (JAK) 2, p95 kDa receptor tyrosine kinase (RTK), and p44/42 kDa MAP kinase (ERK1/2)

160 Phosphorylated p125 kDa JAK2, p95 kDa RTK, and p44/42 kDa MAPK (ERK1/2) 161 were measured by Western blotting analysis (Kurihara et al., 2021b; Towbin et al., 1979). 162 Briefly, the cultured hepatocytes were lysed by Lysis buffer (Cell Signaling Technology, 163 Inc., Danvers, MA, USA) and centrifuged, and then the supernatant was mixed with a 164 Laemmli sample buffer (Bio-Rad Laboratories, Inc., Hercules, CA, USA). The protein 165 samples (30 µg/lane) were divided by polyacrylamide gel electrophoresis and transferred 166 to a PVDF transfer membrane (PerkinElmer, Inc., Waltham, MA, USA). Next, they were 167 immunoblotted with monoclonal antibodies against phosphorylated p125 kDa JAK2, p95 168 kDa RTK, or ERK1/2 (Cell Signaling Technology). The phosphorylated ERK1/2

169 antibody was diluted 2000-fold, and the phosphorylated p125 kDa JAK2 and p95 kDa 170 RTK antibodies were diluted 1000-fold. After incubation with HRP-labeled secondary 171 antibody (Cell Signaling Technology), the blot was detected by Western Lightning Plus-172 ECL (PerkinElmer, Inc.). Emission intensity was quantified by ChemiDoc Touch MP 173 (Bio-Rad Laboratories, Inc.). Phosphorylated p125 kDa JAK2, p95 RTK, or ERK1/2 174 activity ratios were calculated by dividing the respective expression levels (p125 kDa 175 JAK2, p95 kDa RTK, or ERK1/2 antibodies [Cell Signaling Technology]) by the total 176 activity. Each of these antibodies was diluted 1000-fold.

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178 **2.7 Statistical analysis**

Values are shown as means \pm standard error of the mean (SEM) of three separate experiments. For comparison with the control, Dunnett's test was performed, and the significance level was defined as 5%.

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183 **3. Results**

3.1 Time- and dose-dependent cell proliferation effects of SAC and its analogs on cultured hepatocytes

The cell proliferation effects of SAC and its analogs (alliin or allicin) were investigated in cultured hepatocytes. Hepatocytes attached to the plate and exchanged for serum-free medium were cultured with SAC, alliin, allicin, or a combination of SAC with a selective JAK2 inhibitor TG101209 or a selective signal transducer and activator of transcription 3/5 (STAT3/5) inhibitor SH-4-54. Fig. 2 shows a phase-contrast microscopic image of cultured hepatocytes for 5 h. It can be confirmed that the number of SAC-stimulated hepatocyte nuclei was greater than control (medium alone) (Fig. 2A, B). As shown in Fig.

2G, the number of total hepatocytes per field of view (per $0.01 \text{ cm}^2 \text{[} \times 10^2 \text{ cells/cm}^2\text{]}$) under 193 194 the microscope in the control treatment (medium only) was 309 ± 15 . In contrast, the number of total hepatocytes stimulated with 10^{-6} M SAC was 407 ± 19 , which was a 195 196 significant increase of approximately 1.32-fold compared to the control (Fig. 2G). The 197 correlation coefficient between the number of hepatocytes and nuclei stimulated by SAC 198 was 0.95 (Fig.2.H). On the other hand, no increase in the number of hepatocytes was 199 observed with alliin or allicin stimulation (Fig. 2C, D and G). In addition, the number of 200 total hepatocytes treated with SAC and TG101209 was significantly reduced to control 201 levels, but no reduction in the number of hepatocytes was observed in the combination of 202 SAC with SH-4-54 (Fig. 2E-G).

203 Next, the number of hepatocyte nuclei was measured to assess the time- and dose-204 dependent proliferative effects of SAC or its analogs (Fig. 3A, B). As shown in Fig. 3A, 205 SAC-stimulated hepatocyte nuclei were observed to proliferate depending on the culture time. In 10⁻⁶ M SAC-treatment, a significantly higher number of nuclei on cultured 206 207 hepatocytes was observed at about 4 h, peaked at approximately 5 h, and then continued 208 for up to 21 h. In contrast, allicin and alliin did not show a significant increase in 209 hepatocyte nuclei (Fig. 3A). Next, fixing the culture time at 5 h in serum-free culture, effects of SAC or its derivatives at concentrations of 10⁻¹⁰ to 10⁻⁵ M on the number of 210 hepatocyte nuclei were examined. SAC-stimulated hepatocyte nuclei were increased 211 dose-dependently and plateaued at 10^{-6} M, with the 50% effective concentration (EC₅₀) 212 value of 8.58×10^{-8} M. In contrast, no hepatocyte proliferation effect of alliin or allicin 213 was observed in the range of 10^{-10} to 10^{-5} M (Fig. 3B). 214

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216 **3.2 Effects of SAC and its analogs on** *c-Myc*, *c-Fos*, and *cyclin B1* mRNA expression

217 levels in cultured hepatocytes

218 It is known that oncogenes such as *c-Myc* and *c-Fos* are involved in cultured hepatocyte 219 proliferation and liver regeneration (Fausto, 2000). The StepOne Real-Time PCR System 220 was used to investigate whether *c-Myc* and *c-Fos* mRNAs were expressed in hepatocytes 221 stimulated with SAC and its analogs. Fig. 4A shows the time course associated with c-222 Myc mRNA expression levels in hepatocytes stimulated with SAC and its analogs. The 223 expression level of *c-Myc* mRNA was significantly increased in SAC-stimulated hepatocytes. The peak was observed 1 h after addition of SAC (10⁻⁶ M), and the 224 225 expression level increased approximately 1.62-fold compared with time 0. On the other hand, stimulation with alliin (10⁻⁶ M) or allicin (10⁻⁶ M) did not significantly increase c-226 Myc mRNA expression levels. Fig. 4B shows c-Fos mRNA expression levels in SAC- or 227 its analog-stimulated hepatocytes. The c-Fos mRNA expression levels showed a time 228 229 course similar to those of *c-Myc*, and the increase was approximately 1.85-fold in SAC-230 stimulated hepatocytes 1 h after stimulation (Fig. 4B).

In addition, mRNA expression levels of *cyclin B1* were measured as a G_2/M phase marker in SAC or analog-stimulated hepatocytes. As shown in Fig. 4C, *cyclin B1* mRNA expression levels in SAC (10⁻⁶ M)-stimulated hepatocytes began to increase at 3 h and peaked at 3.5 to 4 h. At the peak times, the proportion of SAC-induced *cyclin B1* was significantly increased by approximately 1.72-fold compared to controls (Fig.4C). Alliin (10⁻⁶ M) or allicin (10⁻⁶ M) did not show a significant increase in *cyclin B1* mRNA expression in hepatocytes (Fig.4C).

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3.3 Time course effects of SAC and its analogs on S- and G₀/G₁-phase progression in
cultured hepatocytes

To investigate whether SAC promotes DNA synthesis in cultured hepatocytes, the Sphase in the cell cycle of hepatocytes stimulated with SAC and its analogs was measured. As shown in Fig. 5A, the proportion of S-phase of SAC (10^{-6} M)-stimulated hepatocytes was significantly higher over 3 to 5 h compared to controls. In contrast, alliin (10^{-6} M)or allicin (10^{-6} M)-stimulated hepatocytes did not show an effect on the proportion of Sphase (Fig. 5A). Furthermore, the proportion of hepatocyte nuclei for the G₀/G₁-phase further decreased significantly in SAC (10^{-6} M) than in control at 3 and 4 h (Fig. 5B).

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3.4 Effects of monoclonal antibodies against growth factor receptors on SAC induced hepatocyte proliferation

251 To investigate whether SAC showed a cell proliferation effect via growth factor 252 receptors expressed in cultured hepatocytes, the number of nuclei and the progression of 253 the S-phase of SAC-stimulated hepatocytes in the presence of monoclonal antibodies to 254 growth factor receptors associated with cell proliferation were measured. As shown in Fig. 6A, B, SAC-stimulated progression of the S-phase and increase in hepatocellular 255 256 nuclei were completely suppressed by monoclonal antibodies to GH receptor and IGF-I 257 receptor, respectively. In contrast, monoclonal antibodies against HGF and EGF receptors 258 did not suppress the SAC-induced hepatocyte proliferation (Fig. 6A, B).

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3.5 Effects of SAC on hepatocyte proliferation in the presence of specific signal transduction factor inhibitors

Next, to investigate intracellular signaling pathways associated with SAC-induced cell proliferation and DNA synthesis, the number of nuclei and progression of S-phase of SAC-stimulated hepatocytes in the presence of specific signaling factor inhibitors associated with cell proliferation were measured. Fig. 7A, B show that the hepatocyte
proliferation effects of SAC were completely inhibited to control level by TG101209, a
selective PLC inhibitor (U-73122), a selective IGF-I RTK inhibitor (AG538), a selective
phosphoinositide 3-kinase inhibitor (LY294002), a selective MAPK/ERK kinase (MEK)
inhibitor (PD98059), and a selective mammalian target of rapamycin (mTOR) inhibitor
(rapamycin). On the other hand, SH-4-54 did not inhibit the SAC-induced hepatocyte
proliferation (Fig. 7A, B).

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3.6 Time courses of the effects of SAC on p125 kDa JAK2, p95 kDa IGF-I RTK, and ERK1/2 phosphorylation in cultured hepatocytes

275 JAK2, IGF-I RTK, and ERK1/2 phosphorylation patterns of the SAC-stimulated 276 hepatocytes were measured by Western blot analysis. Fig. 8A shows that phosphorylated 277 p125 kDa JAK2 increased sharply after the addition of SAC and peaked at 5 min; p125 278 kDa JAK2 phosphorylation activity by SAC at the peak was approximately 2.3-fold the 279 activity ratio compared to the control. Then, the phosphorylation activities began to 280 decrease to control levels after 30 and 60 min of culture (Fig. 8A). Phosphorylation by 281 SAC of IGF-I RTK and ERK2 was observed later than that of JAK2, with peaks 20 and 282 30 min after SAC addition, respectively. However, no increase in ERK1 phosphorylation 283 was observed with SAC (Fig. 8B, C).

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3.7 Effects of specific signal transduction factor inhibitors on phosphorylation of

286 JAK2, p95 kDa IGF-I RTK, and ERK2 induced by SAC

287 Next, the effects of specific inhibitors of growth-related signal transducers on JAK2,

288 IGF-I RTK, and ERK2 phosphorylated by SAC were investigated. Fig. 9 shows that SAC-

induced JAK2 phosphorylation was suppressed only by TG101209 (Fig. 9A). SACinduced IGF-I RTK phosphorylation was suppressed not only by TG101209, but also by
U-73122 and AG538 (Fig. 9B). SAC-induced ERK2 phosphorylation was suppressed by
LY294002 and PD98059, in addition to the above inhibitors (Fig. 9C). SH-4-54 or
rapamycin did not affect SAC-induced phosphorylation of both IGF-I RTK and ERK2
(Fig. 9A-C).

295

296 **4. Discussion**

In our previous studies, we demonstrated that SAC accelerated liver regeneration and promoted cell proliferation *in vivo* in two-thirds partially hepatectomized rats (Kurihara et al., 2020). The aim of this study was to examine the mechanism of the proliferative effect of SAC compared to its analogues in primary cultures of adult rat hepatocytes *in vitro*.

Fig. 2 showed that SAC, but not alliin or allicin, significantly increased the number of hepatocytes at 5 h of culture. These results showed that there was a close correlation between the increase in the number of cells and the number of nuclei shown in Figs. 2H, 3, and 7. Furthermore, the effects of SAC on hepatocyte numbers were completely suppressed by TG101209, but not SH-4-54. The results suggest that the effects of SAC are mediated by JAK2, but not STATs.

c-Myc and c-Fos are proto-oncogenes expressed during early liver regeneration and are involved in the transition from the G₀ phase to the G₁-phase in the cell cycle of regenerated hepatocytes (Corral et al., 1988; Morello et al., 1990). Accordingly, since SAC significantly increased mRNA expression levels of c-Myc and c-Fos, SAC can advance the G₀ phase cells to G₁-phase (Fig. 4A and B). In addition, since SAC also 313 increased mRNA expression levels of *cyclin B1*, SAC promoted progression to S- and

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G₂/M-phases in cultured hepatocytes (Fig. 4C, 5).

315 Rapid cell proliferation effects on hepatocytes have been demonstrated not only with 316 SAC, but also with growth factors such as HGF and IGF-I (Kimura and Ogihara, 1998, 1997). It has been shown that the proliferative effects of growth factors in cultured 317 318 hepatocytes are affected by cell density and dexamethasone. Nakamura et al. reported 319 that incorporation of $[^{3}H]$ -thymidine into the nucleus (as a DNA synthesis index) in hepatocytes treated with epidermal growth factor and insulin was faster at low cell density 320 $(2.5 \times 10^4 \text{ cells/cm}^2)$ than at high cell density $(1.0 \times 10^5 \text{ cells/cm}^2)$ (Nakamura et al., 1983). 321 322 Furthermore, we demonstrated that the hepatocellular proliferation effect of HGF was suppressed depending on the dose of dexamethasone $(10^{-10}-10^{-7} \text{ M})$ (Kimura et al., 2011). 323 The rapid cell proliferation effect of SAC may be affected by the cell density $(3.3 \times 10^4$ 324 cells/cm²) and dexamethasone (10⁻¹⁰ M) concentration in the culture conditions of this 325 326 study.

327 Alliin has a structure in which oxygen is added to the sulfur part of SAC, and its 328 structure is very similar to SAC (Fig. 1). Not only SAC, but also alliin and its metabolite, 329 allicin, have antioxidant effects based on the removal of reactive oxygen or activation of 330 the Nrf2-Keap pathway (Chung, 2006; Li et al., 2012b, 2012a). In other words, the results 331 in Figs. 2 to 5 imply that the proliferative effect of SAC on hepatocytes does not go 332 through the Nrf2-Keap pathway in cultured hepatocytes. In addition, we have 333 demonstrated that antioxidants such as isoascorbic acid and vitamin E do not cause 334 hepatocyte proliferation (Moteki et al., 2012). These results suggest that the hepatocyte 335 proliferation effect of SAC proceeds by a mechanism different from the antioxidant effect. To analyze the target of SAC and the signal transduction pathways, effects of some 336

337 monoclonal antibodies against growth factors on SAC-induced hepatocyte proliferation 338 were examined. Fig. 6 shows that SAC-induced progression of the S-phase and the 339 number of nuclei in hepatocytes were completely suppressed by monoclonal antibodies 340 to the GH receptor and IGF-I receptor. In the past, we demonstrated that GH secretes IGF-341 I by stimulation of the GH receptor/JAK2/PLC pathway, and that the secreted IGF-I 342 promotes hepatocyte DNA synthesis and cell proliferation via the IGF-I 343 RTK/PI3K/ERK2/mTOR pathway in primary cultures of adult rat hepatocytes (Kurihara 344 et al., 2021b, 2021a). Based on these results, we hypothesized that SAC promotes cell 345 proliferation via the GH signaling pathway. In support of this notion, the hepatocyte 346 proliferation effect of SAC was completely suppressed by inhibitors related to the 347 JAK2/PLC and IGF-I signaling pathways (Fig. 7). In addition, as shown in Fig. 8, Western 348 blot analysis confirmed that SAC phosphorylated p125 kDa JAK2, p95 kDa RTK (i.e., 349 IGF-I RTK), and ERK2 in cultured hepatocytes. The ERK2, but not ERK1, 350 phosphorylation-enhancing effect of SAC was also observed in the remaining liver in 351 SAC-treated partially hepatectomized rats (Kurihara et al., 2020). Fig. 8 shows that the 352 peak of SAC-induced JAK2 phosphorylation was faster than that of SAC-induced RTK 353 and ERK2 phosphorylation. Furthermore, Fig. 9 shows that SAC-induced RTK phosphorylation was suppressed by TG101209 and U-73122. These results indicate the 354 355 presence of IGF-I RTK downstream of JAK2/PLC in the SAC-induced hepatocyte 356 proliferation signaling pathway. It has been reported that JAK1, which is one of the JAK 357 family, interacts with PLC (Chang et al., 2004; Clark et al., 2014). However, TG101209, 358 which inhibited SAC-induced PLC, has high specificity for JAK2 and low affinity for 359 JAK1 and JAK3 (Pardanani et al., 2007). Therefore, the interaction of JAK with PLC 360 may be mainly mediated by JAK2. On the other hand, LY294002 and PD98059

suppressed SAC-induced ERK2 phosphorylation, but not SAC-induced p95 IGF-I RTK phosphorylation. In contrast, rapamycin did not suppress both SAC-induced RTK and ERK2 (Fig. 9). These results indicate that PI3K and MEK are upstream elements of ERK2, and mTOR is a downstream element of ERK2. Taken together, these results support the notion that SAC activates the IGF-I RTK/PI3K/MEK/ERK2/mTOR pathway expressed in cultured hepatocytes. In addition, we have previously demonstrated that SAC stimulates autocrine secretion of IGF-I (Moteki et al., 2022).

In conclusion, the results of the present study showed that SAC rapidly stimulated proliferation in primary cultures of adult rat hepatocytes, and that the hepatocyte proliferation effect of SAC was based on activation of the GH receptor/JAK2/PLC pathway followed by activation of the IGF-I RTK/PI3K/ERK2/mTOR pathway.

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373 CRediT authorship contribution statement

Hajime Moteki participated in conducting all the experiments in this study and writing a
draft of this manuscript. Mitsutoshi Kimura participated in supervising the planning and

376 execution of the study. Masahiko Ogihara edited the manuscript and managed the project

377 for this research. The final manuscript was approved by all authors.

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379 **Declarations of interest:** none

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1 Figures







16	Fig. 2 Phase-contrast micrographs of hepatocytes cultured with SAC and its analogs for
17	5 h. As described in the Methods section, after a 3-h attachment period, hepatocytes (3.3
18	\times 10 ⁴ cells/cm ²) were cultured in serum-free medium containing SAC, alliin, or allicin
19	with growth-related signal-transducing factor inhibitors for 5 h. Photos are (A), control
20	(medium alone); (B), SAC (10 ⁻⁶ M); (C), alliin (10 ⁻⁶ M); (D), allicin (10 ⁻⁶ M); (E), SAC
21	(10 ⁻⁶ M) + TG101209 (10 ⁻⁶ M); (F), SAC (10 ⁻⁶ M) + SH4-54 (10 ⁻⁶ M); Scale bar, 200 μm.
22	(G), Number of hepatocytes per field of view (per 0.01 $\text{cm}^2[\times 10^2 \text{ cells/ cm}^2]$) under the
23	microscope. Values are shown as means \pm S.E.M. (n=3). ** (P<0.01) shows comparisons
24	with respective control. ## (P<0.01) shows comparisons with respective SAC treatment.
25	(H), Correlation between the number of hepatocytes and the number of nuclei. Open
26	circles (\circ), control (medium only, n=3); closed circles (\bullet), SAC (10 ⁻⁶ M) (n=3); closed
27	diamonds (\blacklozenge), SAC (10 ⁻⁶ M) + TG101209 (10 ⁻⁶ M) (n=3); closed squares (\blacksquare), SAC (10 ⁻⁶ M) (n=3); close
28	⁶ M) + SH4-54 (10 ⁻⁶ M) (n=3).
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Fig. 3 Time- and dose-dependent cell proliferation effects of SAC and its analogs on 38 cultured hepatocytes. As described in the Methods section, hepatocytes (3.3×10^4) 39 40 cells/cm²) attached to the plate were exchanged for serum-free medium and then SAC, alliin, or allicin was added. (A) Time-dependent cell proliferation effects of SAC (• 10⁻⁶ 41 M), alliin ($\triangle 10^{-6}$ M), or allicin ($\mathbf{\nabla} 10^{-6}$ M) in cultured hepatocytes. The arrow shows the 42 addition of the drugs. (B) Dose-dependent cell proliferative effects of SAC (• 10⁻¹⁰-10⁻⁶ 43 M), alliin ($\triangle 10^{-10}$ -10⁻⁶ M), or allicin ($\mathbf{\nabla} 10^{-10}$ -10⁻⁶ M) on cell proliferation in cultured 44 hepatocytes for 5 h of drug stimulation. Values are shown as means \pm S.E.M. (n=3). 45 *(P<0.05) and ** (P<0.01) show comparisons with respective control. 46



48 Fig. 4 Effects of SAC and its analogs on *c-Myc*, *c-Fos*, and *cyclin B1* mRNA expression 49 levels in cultured hepatocytes. The mRNA expression levels of c-Myc, c-Fos, and cyclin B1 in hepatocytes were quantified using the TaqMan assay, as described in the Methods 50 section. (A) *c-Myc*; (B) *c-Fos*; (C) *cyclin B1*. Control (\circ medium alone), SAC (\bullet 10⁻⁶ M), 51 alliin ($\triangle 10^{-6}$ M), or allicin ($\mathbf{\nabla} 10^{-6}$ M) were added at the arrow point. The value for 52 each sample was normalized to the GAPDH mRNA expression levels in each sample. 53 Values are shown as means \pm S.E.M. (n=3). *(P<0.05) and ** (P<0.01) show comparison 54 with respective control. 55



Fig. 5 Time course of the effects of SAC and its analogs on S- and G_0/G_1 -phase progression in cultured hepatocytes. S- and G_0/G_1 -phases in SAC- (• 10⁻⁶ M), alliin- (Δ 10⁻⁶ M), or allicin- ($\mathbf{\nabla}$ 10⁻⁶ M) stimulated, as described in the Methods section. (A) Time course of the percentage in the S-phase. (B) Time course of the percentage in the G_0/G_1 phase. The arrow shows the addition of the drugs. Values are shown as means ± S.E.M. (n=3). *(P<0.05) and ** (P<0.01) show comparisons with respective control.

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Fig. 6 Effects of monoclonal antibodies to growth factor receptors on SAC-induced 67 hepatocyte proliferation. As described in the Methods section, hepatocytes (3.3×10^4) 68 69 cells/cm²) attached to the plate were exchanged for serum-free medium. Then, 70 hepatocytes were cultured in the presence of SAC with or without growth-related signal-71 transducing inhibitors. S-phase progression (DNA synthesis [A]) and number of nuclei 72 (cell proliferation [B]) were measured 5 h after SAC (10⁻⁶ M) was added. Monoclonal 73 antibodies were to growth hormone receptor (anti-GHR mAb), insulin-like growth factor 74 type-I receptor (anti-IGF-I R mAb), epidermal growth factor receptor (anti-EGFR mAb),

75	and hepatocyte growth factor receptor (anti-HGFR mAb). IgG (as a negative control).
76	The antibody concentrations are 100 ng/mL each. Values are shown as means \pm S.E.M.
77	(n=3). ** (P<0.01) shows comparisons with respective SAC treatment.
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Fig. 7 Effects of SAC on hepatocyte proliferation in the presence of specific signal transduction factor inhibitors. After cell attachment and exchanging for serum-free medium as described in the Methods section, hepatocytes were cultured in the presence of SAC with or without growth-related signal-transducing factor inhibitors. S-phase progression (DNA synthesis [A]) and number of nuclei (cell proliferation [B]) were measured 5 h after SAC (10⁻⁶ M) was added. Inhibitors used were TG101209 (10⁻⁶ M), SH-4-54 (10⁻⁶ M), U-73122 (10⁻⁶ M), AG538 (10⁻⁷ M), LY294002 (10⁻⁷ M), PD98059

106	(10 ⁻⁶ M), and rapamycin (10 ng/mL). Values are shown as means \pm S.E.M. (n=3).**
107	(P<0.01) shows comparisons with respective SAC treatment.
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131	Fig. 8 Time course of the effects of SAC on p125 JAK2, p95 kDa IGF-I RTK, and ERK1/2
132	phosphorylation in cultured hepatocytes. Hepatocytes $(3.3 \times 10^4 \text{ cells/cm}^2)$ attached to
133	the plate were replaced with serum-free medium as described in the Methods section.
134	Phosphorylation of p125 JAK2, p95 kDa RTK, and ERK1/2 was determined by Western
135	blot analysis, as described in the Methods. (A) Percentages of phosphorylated p125 JAK2
136	(P-p125 kDa JAK2) / total p125 JAK (p125 kDa JAK2); (B) Percentages of
137	phosphorylated p95 RTK (P-p95 kDa RTK) / total p95 RTK (p95 kDa RTK); (C)
138	Percentages of phosphorylated ERK (P-ERK) 2 / total ERK (ERK) 2. Western blot images
139	are shown at the top of each graph. The arrow shows the addition of SAC (10^{-6} M). Values
140	are shown as means \pm S.E.M. (n=3). *(P<0.05) and ** (P<0.01) show comparisons with
141	respective control.
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156	Fig. 9 Effects of specific signal transduction factor inhibitors on p125 JAK2, p95 kDa
157	IGF-I RTK, and ERK2 phosphorylation induced by SAC. After cell attachment and
158	exchanging for serum-free medium as described in the Methods, hepatocytes were
159	cultured in the presence of SAC with or without growth-related signal-transducing
160	inhibitors. Phosphorylation of p125 JAK2, p95 kDa RTK, and ERK1/2 was determined
161	by Western blot analysis, as described in the Methods. (A) Phosphorylation of p125 kDa
162	JAK2 5 min after SAC (10 ⁻⁶ M) stimulation. (B) Phosphorylation of p95 kDa RTK 20
163	min after SAC (10 ⁻⁶ M) stimulation. (C) Phosphorylation of ERK2 30 min after SAC (10 ⁻
164	⁶ M) stimulation. Western blot images are shown at the top of each graph. Inhibitors used
165	were TG101209 (10 ⁻⁶ M), SH-4-54 (10 ⁻⁶ M), U-73122 (10 ⁻⁶ M), AG538 (10 ⁻⁷ M),
166	LY294002 (10 ⁻⁷ M), PD98059 (10 ⁻⁶ M), and rapamycin. Values are shown as means \pm
167	S.E.M. (n=3).** (P<0.01) shows comparisons with respective SAC treatment.
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CRediT Author Statement

Title of Manuscript:

Cell proliferation effects of *S*-allyl-L-cysteine are associated with phosphorylation of Janus kinase 2, insulin-like growth factor type-I receptor tyrosine kinase, and extracellular signal-regulated kinase 2 in primary cultures of adult rat hepatocytes.

Hajime Moteki: participated in conducting all the experiments in this study and writing a draft of this manuscript. **Mitsutoshi Kimura**: participated to supervise the planning and execution of the study. **Masahiko Ogihara:** edited the manuscript and managed the project for this research. The final manuscript was approved by all authors.

Declaration of interests

⊠ The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

□ The authors declare the following financial interests/personal relationships which may be considered as potential competing interests: