

ORIGINAL ARTICLE

PRODUCTION OF GROWTH-RELATED ONCOGENE PROTEIN- α IN A HUMAN ORAL SQUAMOUS CELL CARCINOMA CELL LINE STIMULATED WITH TUMOR NECROSIS FACTOR- α : ROLE IN TUMOR ANGIOGENESIS AND TUMOR PROLIFERATION

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Abstract The CXC chemokine growth-related oncogene protein- α (GRO- α) has a wide variety of biological activities including as neutrophil trafficking or migration of vascular endothelial cells. In addition, studies have shown a crosstalk between tumor cells and vascular endothelial cells; GRO- α released by endothelial cells induces invasion of tumor cells toward endothelial cells, indicating an importance of GRO- α in a tumor environment. Oral squamous cells are reported to produce GRO- α in response to cytokines such as tumor necrosis factor- α (TNF- α). However, little is known about how GRO- α is involved in oral cancer. Here, we investigated the biological role of GRO- α for both tumor growth and angiogenesis in oral squamous cell carcinoma cells. We first evaluated the effect of TNF- α on GRO- α expression in three oral cancer cells from different origins. Among the cell lines we used, KOSC-2 cells expressed the highest amount of GRO- α mRNA in response to TNF- α . TNF- α -treated condition medium from KOSC-2 cells enhanced endothelial cell chemotaxis and the chemotactic activity was partially inhibited by the addition of neutralizing anti-GRO- α antibody. In addition, GRO- α exerted tumor cell migration of KOSC-2. From these results, we conclude that GRO- α may contribute to both angiogenesis and proliferation in oral cancer.

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Key words: growth-related oncogene protein- α (GRO- α); tumor necrosis factor- α (TNF- α); oral squamous cell carcinoma cells.

原 著

**培養ヒト口腔扁平上皮癌細胞における TNF- α 依存的な GRO- α の誘導：
GRO- α による血管新生作用と腫瘍増殖作用について**

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抄録 CXC ケモカインファミリーである GRO- α は、好中球走化性因子として知られている他、腫瘍増殖能や血管新生能を有することが明らかとなっている。これまでに口腔粘膜上皮の GRO- α 産生は報告されていたが、GRO- α の口腔癌における役割は不明である。そこで本研究では口腔癌における GRO- α を介した血管新生作用や腫瘍増殖効果について実験的に検討した。3 種類の口腔扁平上皮癌由来細胞に TNF- α 処理をしたところ、GRO- α の発現量は細胞間で大きく異なっており、TNF- α 依存的な GRO- α 産生は個々の腫瘍細胞の性質に依存することが示唆された。GRO- α を最も多く産生した KOSC-2(舌癌由来細胞株)の TNF- α 処理後の培養上清は、血管内皮細胞の走化性を亢進し、GRO- α 特異的な中和抗体の添加はその亢進を部分的に抑制した。さらに、ヒト組み換え型 GRO- α は KOSC-2 の増殖を促進した。これらの結果から、口腔癌において GRO- α は腫瘍の増悪因子である可能性が示唆された。

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キーワード: GRO- α ; 腫瘍壊死因子(TNF- α); 口腔扁平上皮癌。

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Introduction

Tumor necrosis factor- α (TNF- α) regulates a variety of biological functions related to inflammatory reactions, cell growth and apoptosis; and the most important source of TNF- α is macrophages¹. TNF- α affects carcinoma cells to induce the expressions of many cytokines^{2,3}. Constitutive activation of nuclear factor- κ B (NF- κ B) is observed in many types of cancer cells, strongly suggesting a critical role in cancer development and progression⁴. Among several carcinogens, TNF- α is thought to be the most potent activator of NF- κ B⁵. In the tumor mass, tumor-associated macrophage (TAM) should be major source of TNF- α ⁶.

Growth-related oncogene protein- α (GRO- α) / CXCL1 was first identified as a growth factor of melanoma^{7,8}. GRO- α belongs to the C-X-C chemokine family and has chemotactic activity for neutrophils⁹. Some types of the C-X-C chemokine family, which contain the sequence Glu-Leu-Arg (the ELR motif) in front of the C-X-C motif, have been shown to possess a potent angiogenic property¹⁰. Interleukin-8 (IL-8) / CXCL8, epithelial and neutrophil activating protein-78 (ENA-78) /CXCL5, and GRO- α are the members of this group¹⁰. A variety of chemokines including GRO- α are rapidly and markedly induced by TNF- α ¹¹. This indicates that GRO- α acts as the secondary mediator in response to TNF- α . TNF- α has also be reported to induce GRO- α normal oral keratinocytes; however, the role of GRO- α in oral squamous cells has not been proven by experimental analysis¹².

In healthy oral mucosa, IL-8 and monocyte chemotactic protein-1 (MCP-1) /CCL2 mRNA are constitutively expressed whereas mRNA expression of GRO- γ /CXCL3, a member of GRO family chemokine, is significant lower¹³. In contrast, high level of GRO- α expression

is shown to be correlated with both tumor angiogenesis and lymph node metastasis in oral cancer¹⁴. Furthermore, microarray analysis revealed that GRO- α is more markedly expressed in oral cancer cells than in normal oral epithelial cells¹⁵. These suggest the essential role of GRO- α in oral cancer cells.

Endothelial cells express NF- κ B-dependent GRO- α , mostly in response to TNF- α ¹⁶. CXC chemokines including GRO- α and IL-8 secreted by endothelial cells have been shown to induce tumor cell invasion¹⁷. On the other hand, the role of GRO- α , which is produced from oral squamous carcinoma, is incompletely understood.

We have been studying the effect of TNF- α on human oral squamous cell carcinoma, and here we report the expression of GRO- α is cell line-specific, even in response to TNF- α . We also studied the effect of GRO- α on tumor growth and endothelial cell chemotaxis.

Materials and Methods

Reagents

Cell culture medium Humedia EB-2 and its supplements were purchased from Kurabo (Osaka, Japan). Primer oligo(dT)₁₂₋₁₈ and M-Mulv reverse transcriptase were from GIBCO-BRL (Gaithersburg, MD, USA). Digoxigenen (DIG)-labeling and detection systems were obtained from Boehringer Mannheim (Mannheim, Germany) and a GRO- α enzyme-linked immunosorbent assay (ELISA) kit from R&D Systems (Minneapolis, MN, USA). An RNeasy total RNA isolation kit and *Taq* DNA polymerase were from Qiagen (Hilden, Germany). A Northern Max kit and a Lig'nScribe kit were from Ambion (Austin, TX, USA).

Cell culture

A cell line of human oral squamous cell carcinoma, KOSC-2, was a generous gift from

the National Institute of Health Science (Tokyo, Japan)¹⁸. The other human oral squamous cell carcinoma cell lines, HSC-3 and Ca9-22 were purchased from JCRB Cell Bank (Osaka, Japan). The cells were cultured using RPMI-1640 (KOSC-2) or DMEM (HSC-3 and Ca9-22) supplemented with 10% fetal bovine serum (FBS) and penicillin/streptomycin. The cells were subjected to the stimulation with TNF- α when they reached about 80% confluence.

Human umbilical vein endothelial cells (HUVECs) were purchased from KURABO Tokyo, Japan). The cells were cultured in Humedia EB-2 supplemented with 2% FBS, 10 ng/mL recombinant human (r(h)) epidermal growth factor, 5 ng/mL r(h) basic fibroblast growth factor, 1 μ g/mL hydrocortisone and 10 μ g/mL heparin. CD45+ cells were found in the cultures.

RNA extraction and quantitative reverse transcription-polymerase chain reaction (qRT-PCR)

Total RNA was extracted from the cells using an RNeasy total RNA isolation kit. Single-strand cDNA was synthesized from 1 μ g of total RNA using primer oligo(dT)₁₂₋₁₈ and M-Mulv reverse transcriptase. A CFX96 Real-Time PCR System (Bio-Rad) was used for quantitative analyses of GRO- α and 18S rRNA expression. The sequences of the primers were: GRO- α -F (5'-ATGGCCCCGCGTGCTCTCTCC-3'), GRO- α -R (5'-GTTGGATTTGTCACACTGTTTCAG-3'), 18S rRNA-F: 5'-ACTCAACACGGGAAACCTCA-3', and rRNA-R: 5'-AACCAGACAAATCGCTCCAC-3'. Amplifications were performed using iQ SYBR Green Supermix (Bio-Rad), according to the manufacturer's specifications. Cycling conditions were as follows: 50°C, 2 min; 95°C, 3 min; 40 cycles of 95°C (15s) + 58°C (30 s) + 72°C (30 s). A melting curve was generated by acquiring fluorescence measurements while slowly heating to 95°C at a rate of 0.1°C per second. Melting curves and quantitative

analysis of the data were performed using a CFX manager, as previously reported¹⁹.

ELISA for GRO- α

After the treatment with TNF- α , the KOSC-2 cells were washed twice with RPMI-1640 and incubated for 2 h in RPMI-1640 containing 0.5% human serum albumin (RPMI-HSA). The medium was collected and subjected to ELISA for GRO- α .

Endothelial cell chemotaxis

Endothelial cell chemotaxis was examined using a 24-well chemotaxis chamber as described previously²⁰. Briefly KOSC-2 cells were grown to confluence and stimulated for 4 h with 10 ng/mL TNF- α . Then the medium was replaced with Medium 199 containing 0.5% HSA (M199-HSA), and the cells were conditioned for 2 h. Aliquots (100 μ L) of the conditioned medium, M199-HSA containing 1 ng/mL r(h) GRO- α , 10 pg/mL vascular endothelial growth factor (VEGF), or control medium were placed in lower chambers and upper chambers filled with 100 μ L of HUVEC suspension (1x10⁵ cells/mL M199-HSA). When indicated, an anti-GRO- α neutralizing antibody was added to the medium. After incubating for 4 h at 37°C, the membrane from each chamber was fixed with methanol and stained with Giemsa solution. Transmigrated cells in random four low-power fields were counted under a microscope.

Wound assay

Confluent monolayers of KOSC-2 cells were wounded using a scalpel and a rubber policeman as described²¹. Then the cultures were washed with 20 mM phosphate-buffered saline, pH 7.4 (PBS), and further incubated in the conditioned medium of the cells stimulated for 4 h with 10 ng/mL TNF- α . The cells were washed with PBS, fixed with 10% formaldehyde, and photographed under a microscope. Control

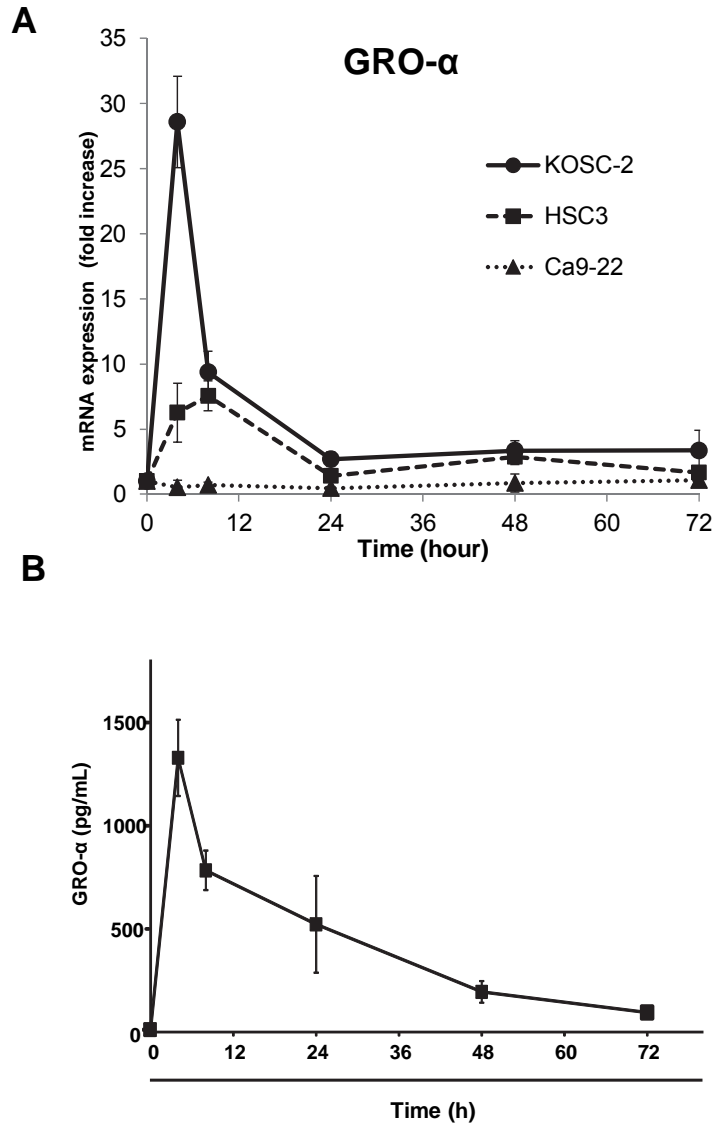


Fig. 1 Time course of the expression of GRO- α in KOSC-2 cells stimulated with TNF- α . (A) KOSC-2 (●), HSC-3 (■), and Ca9-22 (▲) cells were incubated with 1 ng/mL of TNF- α for 4-72 h. mRNA expression of GRO- α or 18s rRNA were analyzed by real-time RT-PCR. (B) KOSC-2 cells were incubated with 1 ng/ml of TNF- α for 4-72 h. The conditioned medium was collected and ELISA was performed. Means (\pm SD) of three experiments shown.

medium and the medium containing 1 ng/mL r(h)GRO- α were also tested in parallel.

Statistics

For chemotaxis assay (Fig.3.), data were analyzed using one-way analysis of variance (ANOVA) to compare the treatment effects. Tukey's post-hoc analyses were applied for multiple comparisons, with the statistical significance set at $P < 0.05$.

Results

Expression of GRO- α in oral squamous cell lines stimulated with TNF- α

We first asked whether most of the oral squamous cancer cells can induce GRO- α in response to TNF- α . In this study, we used three oral cancer cells from different donors to observe GRO- α expression in response to TNF- α . TNF- α (10 ng/mL) transiently expressed

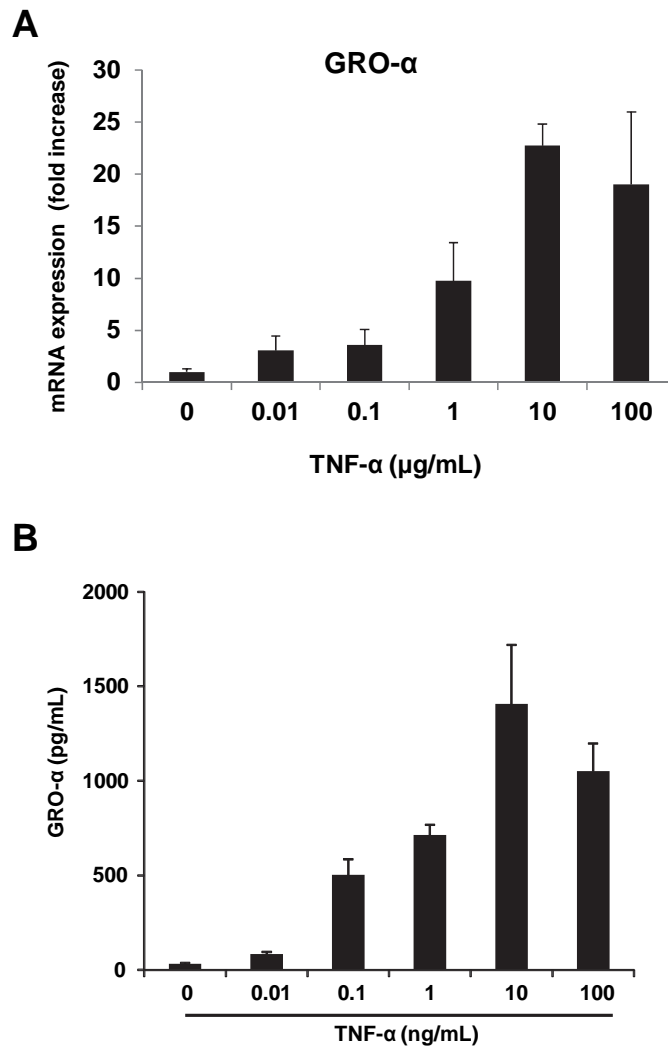


Fig. 2 Concentration-dependent induction of GRO- α by TNF- α . KOSC-2 cells were incubated with 0.01- 100ng/mL TNF- α for 4 h. (A) The expression of mRNA for GRO- α or GAPDH was analyzed by RT-PCR. (B) The conditioned medium of KOSC-2 cells was collected and subjected to ELISA for GRO- α . Means (\pm SD) of three experiments are shown.

GRO- α in KOSC-2 and HSC-3 cells. In both cells, GRO- α mRNA reached the maximal level 4 h after the stimulation with TNF- α (Fig 1A). The induced levels of mRNA levels of GRO- α in KOSC-2 were markedly higher than that in HSC-3 cells. In contrast, no such increase of GRO- α was observed in Ca9-22 (Fig 1A). These observations suggested that the induction of GRO- α in response to TNF- α varies depending on the cell type. The time course of GRO- α protein secretion corresponded with that of the mRNA expression (Fig. 1B).

TNF- α enhanced GRO- α mRNA expression of KOSC-2 cells in a concentration- dependent manner (Fig. 2A). The expression of GRO- α was observed from the treatment with 0.1 ng/mL TNF- α . TNF- α also stimulated the secretion of GRO- α protein and the maximal effect was observed at 10 ng/mL (Fig. 2B).

GRO- α has chemotactic activity for endothelial cells

The results of endothelial cell chemotaxis are summarized in Fig. 3. VEGF is known as

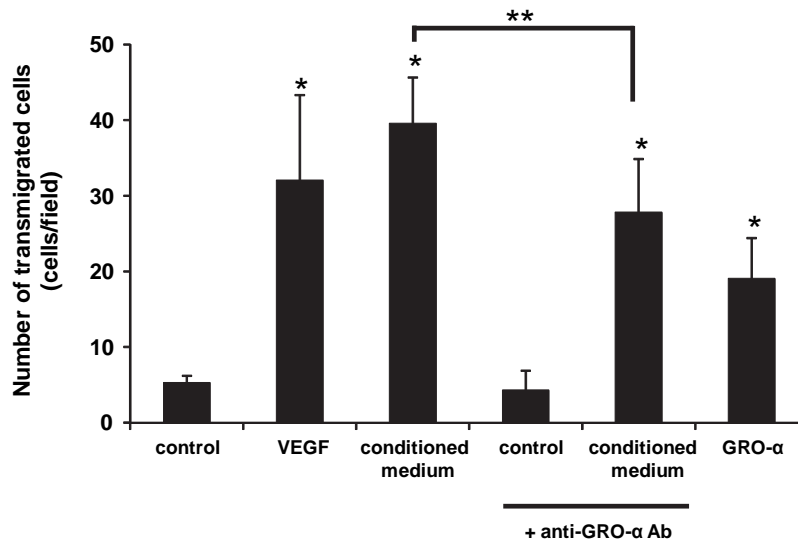


Fig. 3 Endothelial cell transmigration in response to GRO- α . Control medium (M199-HSA), VEGF (100 pg/mL, in M199-HSA), GRO- α (1 ng/mL, in M199-HSA) or conditioned medium (from KOSC-2 stimulated with TNF- α for 4 h) was placed in lower chambers, and upper chambers were filled with HUVEC suspension. When indicated, anti-neutralizing GRO- α antibody was added to the conditioned medium. After incubating for 4 h at 37°C, the membrane was fixed and stained with Giemsa solution. Transmigrated cells in random four fields were counted under a microscope. * P <0.05 statistically significant difference compared with the control, ** P <0.05 vs TNF- α -treated conditioned medium.

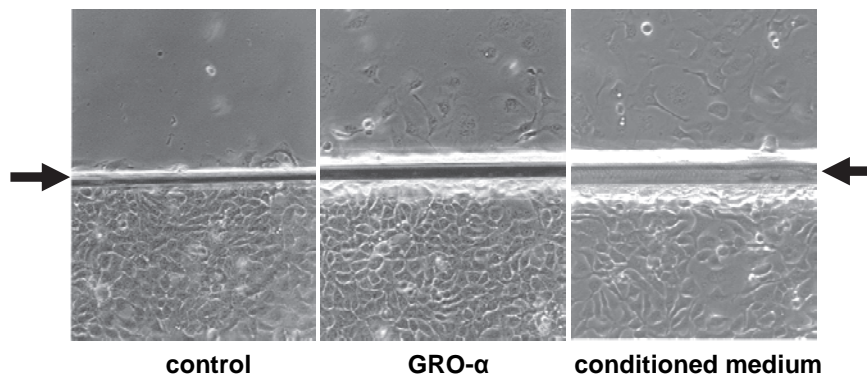


Fig. 4 KOSC-2 cells migration in the presence of GRO- α . Confluent monolayers of KOSC-2 cells were wounded as described in "Materials and Methods". The cells were incubated in the presence or absence of 1 ng/mL GRO- α , or conditioned medium (from KOSC-2 cells stimulated with TNF- α for 4 h) for 20 h, then fixed and photographed. The arrows point to the original edge of the wound. Data shown represent from two independent experiments.

a potent angiogenic factor, and thus we used VEGF as a positive control for this migration assay for endothelial cells. As we expected, only small amount of VEGF could induce chemotaxis in HUVECs. The conditioned medium from TNF- α -treated KOSC-2 cells significantly

enhanced the transmigration of endothelial cells, and r(h) GRO- α was also found to be active in this assay. To evaluate the possible role for GRO- α in the TNF- α -treated conditioned medium, we added anti-neutralizing antibody against GRO- α in the conditioned medium, and

found partial, but significant ($P < 0.05$) inhibition by GRO- α neutralization. These data suggest a positive role of GRO- α as a secondary mediator. In agreement with this result, r(h) GRO- α induces chemotaxis of HUVECs.

GRO- α promotes migration of KOSC-2 cells

The results of KOSC-2 cell migration in wound assay are shown in Fig. 4. Twenty hours after wound assay, control KOSC-2 cells grew into the wounded area. In contrast, the growth was faster in the cells incubated with the medium conditioned by TNF- α -treated cell. r(h) GRO- α also showed a migration promoting activity on KOSC-2 cells.

Discussion

TNF- α was first identified as a factor that induces necrosis of tumor cells; however, various functions of this cytokine have been demonstrated thereafter^{22, 23}. In some case, it serves as a "tumor growth factor"²⁴. TNF- α activates transcriptional factors such as AP-1 and NF- κ B, and subsequently induces the expression of various chemokines^{25, 26}.

In the present study, we initially found that TNF- α induces expression of GRO- α in KOSC-2 cells. GRO- α is known to be expressed in various types of cells including endothelial cells, bronchial epithelium, macrophages and polymorphonuclear neutrophils²⁷⁻³⁰. Previous report has shown the expression of GRO- α by TNF- α in oral keratinocytes¹². However, Ca9-22 derived from an oral squamous cell carcinoma did not express GRO- α in response to TNF- α . Moreover, super-induction of GRO- α was observed in TNF- α -treated KOSC-2 cells. These results suggested that the level of GRO- α is dependent on individual oral squamous cell carcinoma. GRO- α has a neutrophil chemotactic activity and plays an important role in inflammatory responses, but

the ubiquitous nature of its expression suggests that GRO- α is involved in biological events other than leukocyte chemotaxis^{9, 10}. In fact some members of C-X-C chemokines that contain ELR motif are demonstrated to act as an angiogenic factor, while the members that lack ELR- motif serve as an angiostatic factor¹⁰. In the present study, we found that the conditioned medium from TNF- α -treated KOSC-2 cells contained a substantial amount of GRO- α protein and enhanced endothelial cell transmigration. r(h) GRO- α was also found to enhance endothelial migration. Although the medium conditioned by the TNF- α -treated KOSC-2 cells contains many endothelial chemotactic factors, such as IL-8, ENA-78, or VEGF (data not shown), GRO- α may partly account for the activity in the conditioned medium.

GRO- α was originally found as a factor that promotes the growth of melanoma cells, and a subsequent report demonstrated the growth-enhancing effect on other malignant tumors^{7, 8, 31}. We demonstrated, in the wound assay, that GRO- α enhances the growth of KOSC-2 cells; and TNF- α may control the autocrine regulation mechanism of the growth of KOSC-2 cells.

In summary, TNF- α stimulates the secretion of GRO- α by KOSC-2 cells and may control the tumor spread through angiogenesis and growth of the tumor cells.

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