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血管新生とリンパ管新生の同時制御による
制癌法の確立に関する研究

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厚生労働科学研究費補助金（第3次対がん総合戦略研究事業）
（総合）研究報告書

血管新生とリンパ管新生の同時制御による制癌法の確立に関する研究

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研究要旨

研究代表者が発見した新規血管新生抑制因子 Vasohibin-1 (VASH1) と、そのホモログの Vasohibin-2 (VASH2) の血管新生調節における役割について解析し、VASH1 は内皮細胞に発現して血管新生を終息させるのに対し、VASH2 は間質に浸潤する単核球や癌細胞に発現して血管新生を促進することを明らかにした。VASH1 は、広いスペクトルで血管新生とリンパ管新生を同時に阻害し、腫瘍の発育とリンパ節転移を抑制する活性を有しているが内皮細胞は障害せず、腫瘍血管を成熟化させることで抗癌剤との併用効果が示された。特に、VASH1 は内皮細胞を障害しないばかりか、内皮細胞のストレス耐性を増し、このことは他の血管新生阻害剤と際違った相違点である。大腸菌を用いて、活性を保持した組換え VASH1 蛋白の調整法を確立した。組換え VASH1 蛋白の全身投与による治療は、蛋白のデリバリーの点で未だ改善の余地があるが、腹腔内など局所療法の有用性が示された。一方、がん組織に発現する VASH2 は腫瘍血管新生を促進して腫瘍発育に寄与していることから、がん治療の新たな分子標的になることが示された。

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A. 研究目的

癌の治療標的として血管新生とリンパ管新生が注目されている。理想としては血管新生とリンパ管新生の双方を同時に制御する治療法が望ましいが、そのような治療法は未だ確立していない。主任研究者は、血管新生刺激に反応して血管内皮細胞が産生して血管新生を抑制する新規血管新生抑制因子 Vasohibin-1 とそのホモログの vasohibin-2 を発見した。本研究は、以上のオリジナルな研究成績を背景として、血管新生とリンパ管新生の同時制御による制癌法の確立を目指している。本研究を推進することにより、癌の血管新生とリンパ管新生の双方に対し安全で効果的な制御法を確立し、本邦発の新しい治療法として国民のがん医療に大きく貢献することを目的としている。

B. 研究方法

(1) マウスの皮下血管新生モデルを用いて VASH1 と VASH2 の時間・空間的発現様式を明らかにする。作製した VASH1 と VASH2 の遺伝子改変マウスについて、血管新生を惹起させてその過程を野生型マウスと比較する。ヒト

VASH1 または VASH2 遺伝子搭載非増殖型アデノウイルスを尾静注して肝臓で発現させ、血管新生に与える効果を観察する。

(2) マウス角膜法を用いて、VEGF-A、VEGF-C、FGF-2、PDGF-BB など種々の増殖因子によって惹起される血管新生、リンパ管新生に対する VASH1 の効果を検討する。

(3) 高転移性ヒト非小細胞肺癌細胞を SCID マウスに皮下移植、ヒト VASH1 遺伝子搭載非増殖型アデノウイルスを尾静注して肝臓で発現させ、移植 4 週間後に屠殺して癌の発育、血管新生、リンパ管新生、リンパ節転移に対する効果を観察する。この治療後の気管粘膜の血管とリンパ管の形態をコントロールの LacZ 遺伝子搭載非増殖型アデノウイルス注射群と比較する。

(4) VASH1 ノックアウトマウスに癌細胞を皮下移植して、腫瘍の発育と血管新生の程度を野生型マウスと比較することで内因性 VASH1 の意義を検討する。野生型マウスに癌細胞を皮下移植し、ヒト VASH1 遺伝子搭載非増殖型アデノウイルスを尾静注して肝臓で発現させることで、外因性 VASH1 を腫瘍に作用させたときの効果を検討する。

(5) 培養内皮細胞の内因性 VASH1 の発現を siRNA でノックダウンするか、あるいは VASH1 遺伝子導入して高発現させ、血清飢餓

あるいはH₂O₂処理して、VASH1の内皮細胞のviabilityに与える影響を検討する。

(6)大腸菌用コドンに置き換えることで大腸菌でのレコンビナント VASH1 蛋白の産生を行い、蛋白の調整法について、精製方法やリフォールディングなどの条件を検討する。その抗血管新生活性を角膜アッセイで確認する。さらに、レコンビナント VASH1 蛋白をポリエチレングリコール(PEG)化修飾して体内動態を検討する。

(7) 精製したレコンビナントVASH1の抗腫瘍活性を、マウスモデルを用いて検証する。

(8) Vasohibinファミリーのもう1つの分子であるVASH2のがん組織における発現と局在を検証する。また、がん細胞におけるVASH2の発現を変動させ、腫瘍発育や腫瘍血管新生における意義を動物モデルを用いて検証する。

(倫理面への配慮)

全ての動物実験は所属施設での審査を受けた後に行う。

C. 研究結果

平成19年度

(1) マウス皮下の血管新生モデルを用いて VASH1, VASH2 の時間的・空間的発現を詳細に検討したところ、野生型マウスでは、VASH1 は、血管新生先端の発芽部位でなく、後方の血管新生終止部位の血管内皮に発現するのに対し、VASH2 は主に発芽部位に浸潤する骨髄由来の単核球に発現していた。これに対し、VASH1 遺伝子改変マウスでは血管新生が終息しにくかったのに対して、VASH2 遺伝子改変マウスでは発芽部位での血管新生が障害されていた。アデノウイルスベクターを用いて VASH1 や VASH2 を補充すると、それらの異常は正常化した。

(2) VASH1 は種々の増殖因子によって惹起される血管新生、リンパ管新生を広い作用スペクトルで抑制した。

(3) 高転移性ヒト非小細胞肺癌細胞の移植実験では、LacZ 遺伝子搭載非増殖型アデノウイルス注射群と比較して、ヒト VASH1 遺伝子搭載非増殖型アデノウイルス注射群では、腫瘍の発育、腫瘍内血管密度、腫瘍周囲リンパ管密度、所属リンパ節の転移はいずれも有意に抑制された。また、気管粘膜の血管とリンパ管の形態はヒト VASH1 遺伝子搭載非増殖型アデノウイルス注射群で変化を認めなかった。

(4) 培養内皮細胞に VASH1 を発現させると、血管内皮細胞の増殖は抑制され、血清飢

餓や H₂O₂ など各種の細胞死を惹起する処理に曝されても血管内皮細胞の細胞死は顕著に抑制された。

平成20年度

(1) VASH1 ノックアウトマウスでは、移植腫瘍は腫瘍血管新生に富んでおり、内因性 VASH1 は腫瘍血管新生制御に機能していることが確認された。

腫瘍を移植した野生型マウスにヒト VASH1 遺伝子搭載非増殖型アデノウイルスを尾静注すると、外因性 VASH1 は腫瘍血管を成熟化させた。また、その結果として血流が改善し、抗癌剤との併用効果が観察された。

(2) 一般に血管新生抑制因子は内皮細胞のアポトーシスを促進するが、VASH1 は内皮細胞のストレス耐性を増し、内皮細胞の生存を助長することが確認された。その機序として VASH1 は SOD2 の発現を介して ROS の生成量を制御すると共に、長寿遺伝子 SIRT1 の発現を増すことが判明した。

(3) 治療応用に資するためレコンビナント VASH1 蛋白の大量調整法の検討を進め、大腸菌を用いて活性を有するレコンビナント蛋白の調整を可能とし、本蛋白が、動物実験においても活性を発揮することを確認した。さらに、レコンビナント VASH1 蛋白の PEG 化修飾による体内での安定性の改良を行った。

平成21年度

(1) 内皮細胞を障害せずに、広いスペクトルで血管新生とリンパ管新生を抑制する VASH1 について、大腸菌由来組換え蛋白のリフォールディングによる分散画分(活性画分)調製法の改良を進め、活性を有する組換え蛋白の調整法を確立した。

(2) 動物モデルにおいて、このレコンビナント VASH1 蛋白を腫瘍局所に注射すると抗腫瘍効果は得られるが、血中での安定化を目的に PEG 化修飾した蛋白の全身投与では、PEG 化によって活性化が低下するため十分な抗腫瘍効果は得られなかった。一方、腹膜播種すると腹水を形成して急速に個体を死に至らしめるヒト卵巣癌由来 SKOV-3 細胞に VASH1 遺伝子を安定導入すると、mock コントロールではヌードマウスの腹腔内移植で 4 週間以内に全て死亡するのに対し、VASH1 遺伝子導入群では腹水も生じず、全て 10 週間以上生存するという顕著な効果を観察した。

(3) Vasohibin ファミリーメンバーの VASH2 は、血管新生局所に浸潤する骨髄由来単核球に発現し、VASH1 と拮抗して血管新生を促進することが判明しているが、癌細胞の一部にも発現することが判明した。そこで

VASH2 発現陰性の腫瘍細胞に VASH2 を導入してマウスに移植すると、腫瘍血管新生と腫瘍発育が増強することが判明した。

D. 考察

VASH 1 は、発芽部位より後方の血管新生終止部位の血管内皮に発現して、血管新生を止める働きがあるのに対して、VASH2 は主に発芽部位に局在し、VASH1 と拮抗して血管新生を促進すると考えられた。

VASH1 は、血管新生のみならずリンパ管新生をも広い作用スペクトルで抑制するが、正常内皮を障害せず、癌のリンパ節転移制御に対しても有用と考えられた。

VASH1 は、広いスペクトルで血管新生とリンパ管新生を阻害するが、内皮細胞を障害せず、逆にストレス耐性を増ことは他の血管新生阻害剤と際違った相違点である。腫瘍血管を成熟化させるため、抗癌剤との併用が効果的と考えられる。

レコンビナント VASH1 蛋白の調整法は確立できたが、血管をルートとして全身投与する治療法への応用は、現時点では未だ困難である。しかし、腹腔内投与など、局所療法への応用は、可能性の高い手法と考えられた。VASH2 はがん細胞に発現し、腫瘍血管新生を促進して腫瘍発育に寄与している。従って、VASH1 を外因性に投与すると同時に VASH2 の作用を阻害することが、がん治療における望ましい治療法となると期待される。

E. 結論

VASH1 は、血管新生のみならずリンパ管新生に対しても広いスペクトルで抑制作用を有しており、血管新生とリンパ管新生の同時制御による制癌法への応用が期待される。

内皮細胞が産生する血管新生・リンパ管新生抑制因子 VASH1 の特異な性質を明らかにし、さらに治療応用の一環としてのレコンビナント蛋白の大量調整と体内安定化に目処がたった。

レコンビナント VASH1 蛋白の大量調整法を確立すると共に、がんにおける VASH2 の意義が明らかにし、VASH2 が、がん治療の新たな分子標的となえう可能性を示した。がん治療においては、外因性のレコンビナント VASH1 蛋白を投与に VASH2 の作用阻害を併用することの有用性が示唆された。

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G. 知的所有権の取得状況

1. 特許取得

特許出願「バソヒビンからなるリンパ管新生抑制剤」

基礎出願 2006-324773

WO PCT/JP2007/072838

特許出願「バソヒビン1を認識するモノクローナル抗体および該抗体を用いるバソヒビンの免疫測定方法」

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佐藤靖史 血管新生促進因子 特許出願

PCT/JP2009/051359

2. 実用新案登録

なし

3. その他

なし

研究成果の刊行に関する一覧表レイアウト

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Puromycin Insensitive Leucyl-Specific Aminopeptidase (PILSAP) Affects RhoA Activation in Endothelial Cells

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Puromycin insensitive leucyl-specific aminopeptidase (PILSAP) expressed in endothelial cells (ECs) plays an important role in angiogenesis due to its involvement in migration, proliferation and network formation. Here we examined the biological function of PILSAP with respect to EC morphogenesis and the related intracellular signaling for this process. When mouse endothelial MSS3I cells were cultured, a dominant negative PILSAP mutant converted cell shape to disk-like morphology, blocked stress fiber formation, and augmented membrane ruffling in random directions. These phenotypic changes led us to test whether PILSAP affected activities of Rho family small G-proteins. Abrogation of PILSAP enzymatic activity or its expression attenuated RhoA but not Rac1 activation during cell adhesion. This attenuation of RhoA activation was also evident when G-protein coupled receptors such as proteinase-activated receptor or lysophosphatidic acid receptor were activated in ECs. These results indicate that PILSAP affects RhoA activation and that influences the proper function of ECs.

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Angiogenesis is the formation of new blood vessels through endothelial cell (EC) proliferation and migration in combination with tubular morphogenesis. Angiogenesis is indispensable for various physiological and pathological processes, such as embryonic development, wound healing, diabetic retinopathy, and solid tumor growth. A number of molecules regulate angiogenesis both positively and negatively. However, the molecular mechanism of angiogenesis is not yet completely understood.

We searched for novel molecules involved in angiogenesis regulation, and isolated puromycin insensitive leucyl-specific aminopeptidase (PILSAP) with the use of subtraction strategy whose expression was augmented during the *in vitro* differentiation of murine embryonic stem (ES) cells to ECs (Miyashita et al., 2002). The expression of PILSAP in ECs is evident at the site of angiogenesis *in vivo*, and is regulated, at least in part, by a transcription factor polyomavirus enhancer-binding protein 2 (PEBP2) (Miyashita et al., 2002; Niizeki et al., 2004). Interestingly, endoplasmic reticulum aminopeptidase associated with antigen processing (ERAAP) was found to be identical to PILSAP, which was involved in the cleavage of various peptides for antigen presentation by MHC class I molecules (Serwold et al., 2002). Nevertheless, our analyses have revealed that PILSAP plays an important role in angiogenesis by its involvement in migration, proliferation and network formation (Akada et al., 2002; Miyashita et al., 2002). Aminopeptidases catalyze the sequential removal of amino acids from unblocked N-termini of peptides and proteins and play important roles in various biological processes, such as maturation, activation, modulation, degradation of bioactive peptides (Taylor, 1993). PILSAP belongs to the M1 aminopeptidase family, which contains an HEXXH(18X)E motif and a central Zn²⁺ ion essential for enzymatic activity. We examined the mechanism by which PILSAP regulates vascular endothelial growth factor (VEGF)-stimulated proliferation of ECs. As PILSAP is an aminopeptidase, PILSAP is expected to modulate cell function by catalyzing its physiological substrates. Our analysis have revealed that PILSAP binds to

phosphatidylinositol-dependent kinase I (PDK1) and removes 9 amino acids from the PDK1 N-terminus, which subsequently allows S6 kinase (S6K) to associate with PDK1 and PILSAP upon VEGF stimulation (Yamazaki et al., 2004).

Cell migration is a mechanically integrated molecular process that involves dynamic, coordinated changes in cell adhesions and cytoskeletal reorganization. The migration process includes protrusion of the leading edge, formation of new adhesions at the front, cell contraction, and the release of adhesions at the rear (Lauffenburger and Horwitz, 1996; Sheetz et al., 1998; Li et al., 2005). Reorganization of actin cytoskeleton generates locomotive force, and this process is regulated by Rho family small GTPases such as RhoA, Rac, and Cdc42. Rho family small GTPases act as molecular switches by cycling between GTP- and GDP-bound states and transmit extracellular chemotactic signals to downstream effectors. Activated Rac and Cdc42 induce reorganization of actin cytoskeleton at the leading edge. This reorganization of actin

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cytoskeleton induces the formation of membrane protrusion such as membrane ruffling (Small et al., 2002). In contrast, RhoA regulates the assembly of contractile acto-myosin filaments. This RhoA-mediated acto-myosin contractile force promotes locomotion of the cell body and the trailing edge (Nobes and Hall, 1999). These organized activities of Rho family small GTPases make the proper cell polarity. We previously showed that PILSAP did not affect integrin expression, but was involved in adhesion to extracellular matrix proteins (Akada et al., 2002). Here, we extended our analysis to elucidate the molecular mechanism as to how PILSAP regulates adhesion of ECs. Our present analysis revealed that PILSAP takes part in the activation of RhoA in ECs.

Materials and Methods

Materials

The following materials were used: growth factor-reduced Matrigel (Collaborative Research, Bedford, MA); α -minimum essential medium (α MEM), Opti-MEM, Lipofectamine, Superscript II reverse transcriptase, oligo(dT)12–18 primer, and oligofectamine (Gibco BRL, Rockville, MD); anti-RhoA Ab and anti-Rac1 Ab (Santa Cruz Biotechnology, Santa Cruz, CA); Isogen (Nippon Gene, Toyama, Japan); PAR1 agonist peptide (PAR-1 AP) (TFLLR-NH₂, Tocris Cookson, Bristol, UK); Y27632 (EMD Bioscience, San Diego, CA); lysophosphatidic acid (LPA; Biomol Laboratories, Plymouth Meeting, PA); nitrocellulose membranes (Amersham Biosciences, Buckinghamshire, UK); FuGENE 6 Reagent (Roche Diagnostics, Mannheim, Germany); C3 exoenzyme (Calbiochem, La Jolla, CA). Other chemicals were purchased from Sigma (St. Louis, MO).

Cell culture

Mouse endothelial MSS31 cells isolated from mouse spleen microfossils (Yanai et al., 1991) were routinely cultured in α MEM containing 5% fetal bovine serum (5% FBS/ α MEM; Oda et al., 1999). MSS31 cells were stably transfected with pcDNA4A (Invitrogen, Carlsbad, CA) empty vector (Mock), Wt-PILSAP or Mut-PILSAP (Yamazaki et al., 2004). Established transfectants in bulk were maintained in 5% FBS/ α MEM containing 100 μ g/ml of zeocin (Invitrogen). Mut-PILSAP acts as a dominant negative molecule in which glutamate residue 343 is substituted with an alanine (E343A) in the aminopeptidase motif of PILSAP.

Immunofluorescence staining

For the inhibition of RhoA, cells were treated with C3 exoenzyme according to the method described by Minambres et al. (2006). Briefly, C3 exoenzyme (2.5 μ g) was precomplexed with FuGENE 6 Reagent (5 μ l) in 100 μ l of opti-MEM, and C3 exoenzyme complex was added to the cultures. For the inhibition of Rho kinase, cells were incubated in 5% FBS/ α MEM on type-I collagen coated dishes for 8 or 10 h with or without Y27632. In addition, cells were cultured in 1% FBS/ α MEM prior to inoculation onto type-I collagen coated dishes for 10 h and then stimulated with or without PAR-1 AP (40 μ M) or LPA (10 μ M). Thereafter, cells were fixed with 3.8% formaldehyde for 10 min at room temperature and permeabilized with 0.1% Triton X-100 in phosphate buffered saline (PBS). Non-specific binding sites were blocked with 1% bovine serum albumin (BSA) in PBS. Filamentous actin (F-actin) was detected by rhodamine phalloidine (Molecular Probes, Eugene, OR) and focal adhesion complexes were detected by indirect immunofluorescence by using anti-paxillin Ab (Transduction Laboratories, Lexington, KY) and FITC-labeled secondary Ab (Jackson ImmunoResearch, West Grove, PA). Then, cells were observed by confocal microscopy (LSM410, Carl Zeiss Jena GmbH, Jena, Germany).

Cell morphology during movement

Transfected cells were plated onto type-I collagen coated 35 mm dishes for 10 h in 5% FBS/ α MEM at a sparse density, so that cells did not affect the movement of each other. Cells were cultured at 37°C in 5% CO₂. Next, cells were photographed by phase-contrast time lapse microscopy in random high-power (200 \times) fields after additional incubation of 2–6 min.

RhoA and Rac1 activities

Pull-down assay kits (Rho activation assay kit and Cdc42 activation kit, Upstate Biotechnology, Lake Placid, NY) were used to measure RhoA, Rac1 and Cdc42 activities in stable transfectants, leucinethiol (LT), a specific inhibitor of leucine aminopeptidase, treated parental cells or siRNA transfected parental cells. The Cdc42 activation kit includes p21 activated kinase 1 (PAK-1) binding domain agarose. PAK-1 binds both Rac1 and Cdc42 thus; it can measure both Rac1 and Cdc42 activity. Cells were cultured in 1% FBS/ α MEM for 24 h and replated in 1% FBS/ α MEM on type-I collagen coated dishes for 10 h. In LT treatment experiments, parental cells were plated onto type-I collagen coated dishes for 10 h in 1% FBS/ α MEM with or without LT. In some experiments, cells were treated with or without 1% FBS/ α MEM containing PAR-1 AP (40 μ M) or LPA (10 μ M) for 5 min. Cells were extracted with lysis buffer A (25 mM HEPES pH 7.5, 150 mM NaCl, 1% Igepal CA-630, 10 mM MgCl₂, 1 mM EDTA, and 10% glycerol). The pull-down of activated RhoA, Rac1, or Cdc42 was performed according to the manufacturer's protocol. A protein assay was performed to equalize the protein amount of each treatment group.

Western blot analysis

The proteins extracted by lysis buffer A or the samples obtained by pull-down assay were separated by SDS-polyacrylamide gel electrophoresis on a 10% gel and then transferred to nitrocellulose membranes (Iwasaka et al., 1996). The membranes were blocked for 1 h at room temperature with Tris-HCl-buffered saline (TBS), pH 7.4, containing 5% skim milk, and then incubated for 1 h at room temperature in TBS containing 0.05% Tween-20 (T-TBS), 1% BSA, and anti-RhoA Ab (1:1,000) or anti-Rac1 Ab (1:1,000). The filters were washed three times with T-TBS and incubated for 1 h with horseradish peroxidase-conjugated protein G (Bio-Rad, Hercules, CA) diluted 1:3,000 in T-TBS. After the filters were washed with T-TBS three times, signal was detected by an enhanced chemiluminescence method with the ECL Western blotting detection kit (Amersham Bioscience). The results were visualized with LAS-1000 (Fuji Film, Tokyo, Japan).

siRNA transfection

RNA interference of the ERAAP gene (identical to PILSAP) was described by Serwold et al. (2002). We generated another siRNA to strengthen the siRNA-mediated knock down of PILSAP. The coding strands of the two pair of siRNA oligonucleotide directed to the 5' end of mouse PILSAP messenger RNA were 5'-AGCUAGUAAUGGAGACUCATT-3', 5'-UGAGUCUCCAUUACUAGCUTT-3' and 5'-CCUCAGCACUCGACUUUCTT-3', 5'-GAAAGUCAGAGUGCUGAGGTT-3' and scramble RNAs, the negative control of siRNA, were 5'-AAGAUUCGACGAGCUAUAGTT-3', 5'-CUAUAGCUCGUCGAAUCUUTT-3' and 5'-UUAGCCCGUCUACGAAUUUTT-3', 5'-AAAUUCGUAGACGGGCUAATT-3'. RNA duplexes were denatured in annealing buffer (100 mM NaCl, 50 mM Tris-HCl pH 7.5) at 90°C for 1 min and subsequently annealed at 60°C for 1 h. MSS31 cells were cultured in 5% FBS/ α MEM for 24 h, and then oligonucleotides were transfected into cells using oligofectamine according to the manufacturer's instructions. Twenty-four hours

after transfection, MSS31 cells were cultured in 5% FBS/ α MEM for 24 h. The gene silencing effect was confirmed by a quantitative real time RT-PCR.

Quantitative real time RT-PCR

Quantitative real time RT-PCR was performed using a Light Cycler System (Roche Diagnostics) as described previously (Shibuya et al., 2006). Total RNA isolated from parental MSS 31 cells was extracted by ISOGEN according to the manufacturer's instructions. RNA was reverse transcribed with AMV reverse transcriptase (Roche Diagnostics) and oligo(dT)12–18 primer according to the manufacturer's instructions. PCR conditions consisted of an initial denaturation step at 95°C for 10 min, followed by 40 cycles of 15 sec at 95°C, 5 sec at 60°C and 15 sec at 72°C. The primer pairs used were: PILSAP 5'-GATGATGGATGGGCTTCTCT-3' (forward primer) and 5'-GGCTTTTCTCAGTACTAGAC-3' (reverse primer); mouse β -actin 5'-TCGTGCGTGACATCAAAGAG-3' (forward primer) and 5'-TGGACAGTGAGGCCAGGATG-3' (reverse primer). Each mRNA level was measured as a fluorescent signal corrected according to the signal for β -actin.

Network formation

The transfected cells were harvested with 0.25% trypsin and 1 mM EDTA, resuspended in 5% FBS/ α MEM with or without Y27632 (10 μ M) in a final volume of 1 ml, replated (2×10^5 cells per dish) onto 35 mm dishes coated with Matrigel (700 μ l per dish), and incubated at 37°C for 12 h. Cells were observed by phase-contrast microscopy. The length of network structures was quantified with Soft Imaging System Analysis.

Calculations and statistical analysis

The statistical significance of differences in the data was evaluated by the use of unpaired analysis of variance. *P* values were calculated

by the unpaired Student *t*-test. *P* < 0.05 was accepted as statistically significant.

Results

PILSAP is involved in F-actin formation during cell adhesion

The consensus HEXXH(18X)E motif is defined as a unique signature for zinc metalloproteinase and glutamate residues are essential for catalytic activity (Hooper, 1994). We previously established a plasmid in which glutamate residue 343 is substituted with an alanine (E343A, namely, HAXXH(18X)E) in the aminopeptidase motif of PILSAP, and this mutant PILSAP acts as a dominant negative molecule (Yamazaki et al., 2004). To investigate the biological function of PILSAP in ECs, we transfected MSS31 cells with empty vector (Mock), wild-type PILSAP (Wt-PILSAP) expressing vector or mutant PILSAP (Mut-PILSAP) expressing vector, and established the respective stable transfectants. Cells were routinely cultured on type-I collagen coated dishes. The most significant change that we found was cell morphology. As shown in Figure 1A, Mut-PILSAP transfectants showed disk-like morphology and lost cell polarity in sparse to subconfluent conditions. Because of this phenotypic change, we examined the actin cytoskeleton organization. Stress fibers were scarcely formed in Mut-PILSAP transfectants (Fig. 1B). In contrast, formation of membrane ruffling was increased but in random directions in Mut-PILSAP transfectants (Fig. 1C).

ECs use at least a type I collagen receptor integrin $\alpha 2\beta 1$, a fibronectin receptor $\alpha 5\beta 1$, and vitronectin receptors $\alpha v\beta 3$ and $\alpha v\beta 5$ for angiogenesis (Brooks et al., 1994; Collo and Pepper, 1999). Expression of these integrins was unaltered, and identical phenotypic changes were observed when transfectants were plated on fibronectin or vitronectin coated dishes (data not shown).

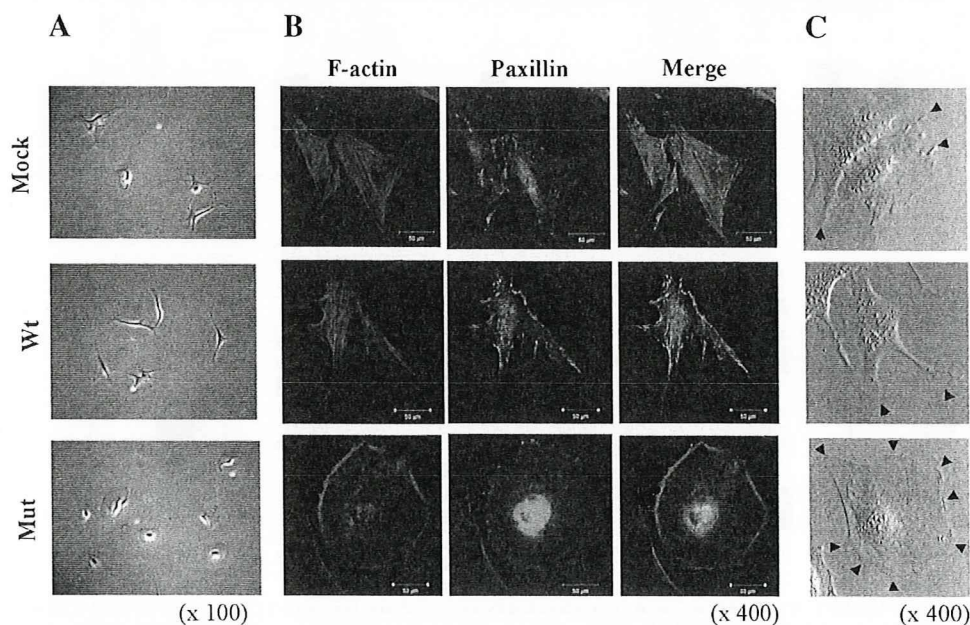


Fig. 1. Involvement of PILSAP in cell shape change, actin organization and membrane ruffling. **A:** Transfectants were plated onto type-I collagen coated dishes and incubated for 8 h in 5% FBS/ α MEM. Cell shape was observed by a phase-contrast microscopy (upper part, 40 \times , lower part, 100 \times). **B:** Transfectants were plated onto type-I collagen coated dishes, and incubated for 8 h in 5% FBS/ α MEM, and then stained with rhodamine-phalloidin (red) and anti-paxillin Ab (green). A scale bar indicates 50 μ m. **C:** Cell morphology during movement was observed by time lapse microscopy. Typical pictures are shown here. Arrow heads indicate membrane ruffling.

PILSAP is involved in RhoA activation for EC morphogenesis

As Rho family small GTPases play a pivotal role in actin cytoskeleton reorganization, we examined the activities of RhoA, Rac1 and Cdc42 in transfectants upon cell adhesion. We observed that RhoA activity was attenuated, while Rac1 activity was augmented in Mut-PILSAP transfectants (Fig. 2A). Cdc42 activity was hardly detected (data not shown). Involvement of PILSAP in RhoA activation was confirmed by two additional treatments. We previously showed that PILSAP was highly sensitive to LT, but was insensitive to puromycin (Miyashita et al., 2002). LT inhibited spreading of MSS31 cells upon extracellular matrix such as type I collagen, fibronectin and vitronectin (Akada et al., 2002). Here we examined whether LT influenced RhoA activity. As shown in Figure 2B, LT inhibited RhoA activation in parental MSS31 cells when added to the medium. Moreover PILSAP siRNAs, which knocked down to 16% of the control level of PILSAP mRNA (Fig. 2D), decreased RhoA activity of parental MSS31 cells (Fig. 2C). RhoA regulates cell contractility through its downstream Rho kinase (Alblas et al., 2001). To further confirm the involvement of RhoA activity in EC morphogenesis, we employed specific RhoA inhibitor, C3 exoenzyme, or a Rho kinase inhibitor, Y27632. When parental MSS31 cells were treated with C3 exoenzyme or Y27632, cells exhibited cell-shape changes identical to that in Mut-PILSAP transfectants (Fig. 3A,B). ECs form network-like structures when plated on Matrigel. This network formation was aborted in Mut-PILSAP

transfectants, or by the treatment of Wt-PILSAP transfectants with Y27632 (Fig. 4). These results indicate that PILSAP is involved in RhoA activation for proper morphogenesis and organization of ECs.

PILSAP is involved in RhoA activation via G-protein coupled receptors

During the maintenance of transfectants in culture, we noticed that Mut-PILSAP transfectants showed delayed cell shrinkage upon trypsin/EDTA treatment for cell harvest. We reasoned this delay of cell shrinkage was due to impaired RhoA activation. Indeed, when transfectants were pretreated with Y27632, cell shrinkage upon trypsin/EDTA treatment was inhibited in Mock or Wt-PILSAP transfectants (data not shown). Trypsin activates proteinase-activated receptors (PARs). PARs belong to the G-protein coupled receptors (GPCRs), and ECs express PAR1 and PAR2 (Brass and Molino, 1997). Western blot analysis revealed that protein levels of PAR1 and PAR2 were identical between the three transfectants (data not shown). It was previously reported that PAR1 or PAR2 agonist induced RhoA activation in HUVEC (Vouret-Craviari et al., 2003). Here we used the PAR1 AP to induce RhoA activity in our system. There was no significant difference in the basal level of RhoA activity between transfectants. However when transfectants were treated with the PAR1 AP, RhoA activation was almost completely abolished in Mut-PILSAP transfectants (Fig. 5A). Moreover, with respect to actin reorganization, PAR1 AP

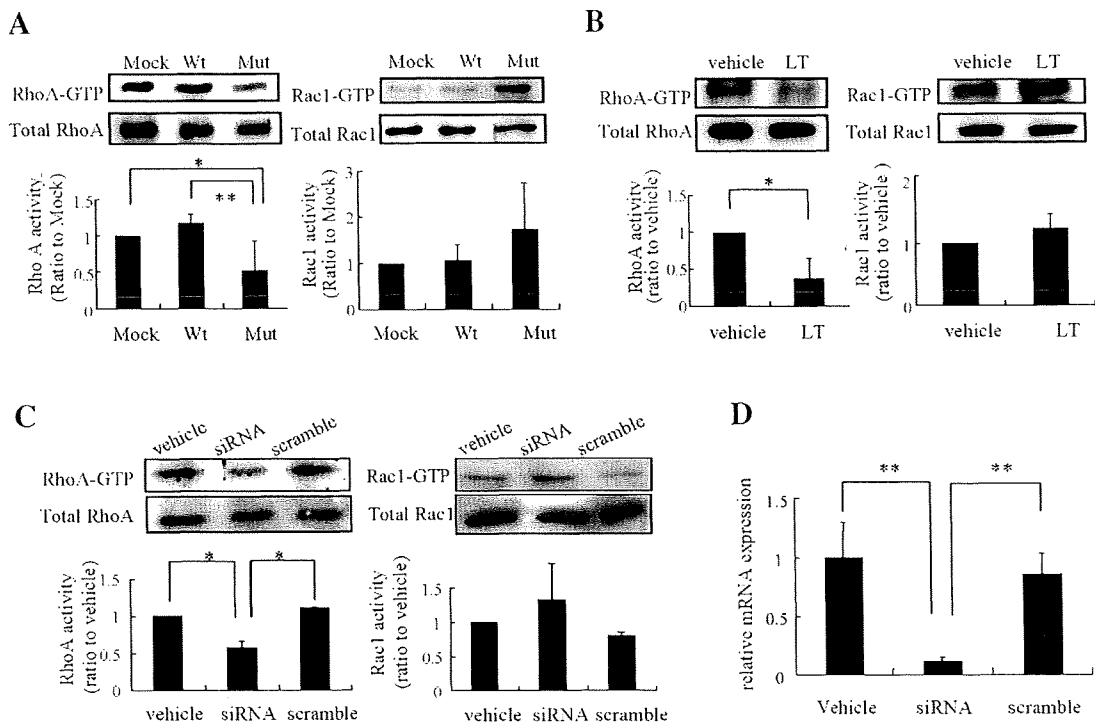


Fig. 2. Involvement of PILSAP in RhoA activation. **A:** Transfectants were incubated on type-I collagen for 10 h in 1% FBS/ α MEM. Then, RhoA and Rac1 activities were determined. **B:** Parental MSS31 cells were incubated on type-I collagen for 10 h in 1% FBS/ α MEM with or without LT (10 μ M/L). Next, RhoA and Rac1 activities were determined. GTP-RhoA or GTP-Rac1 was quantified by density and normalized to that of total RhoA or total Rac1. The values are expressed as the mean \pm SD from three independent experiments; * P < 0.05, ** P < 0.01. **C:** Parental MSS31 cells were transfected with PILSAP siRNA and incubated in 1% FBS/ α MEM for 24 h. Then, transfectants were plated onto type-I collagen coated dishes and incubated for 10 h. Next, RhoA and Rac1 activities were determined. The values are expressed as the mean \pm SD from two independent experiments; * P < 0.05. **D:** The gene silencing effect of PILSAP siRNA was confirmed by quantitative real time RT-PCR. The values are expressed as the mean \pm SD from three samples; ** P < 0.01.

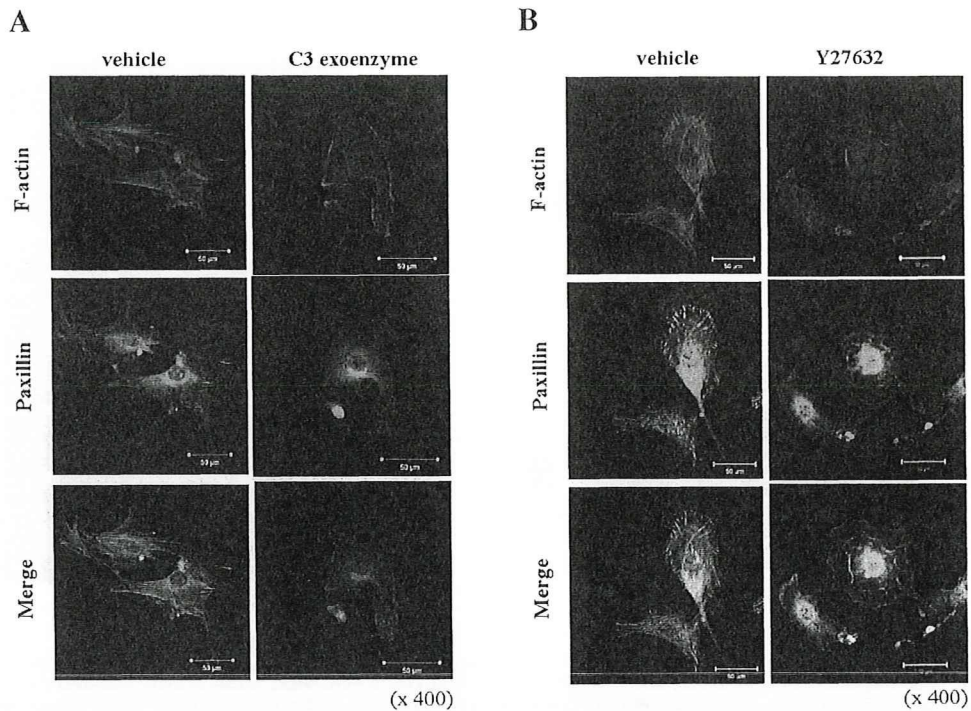


Fig. 3. Effects of RhoA and Rho kinase inhibitors on cell morphology. **A:** Parental MSS31 cells were plated onto type-I collagen coated dishes and incubated for 1 h in 5% FBS/ α MEM. Next, cells were treated with or without C3 exoenzyme precomplexed with FuGENE 6 Reagent for additional 7 h. Cells were then stained with rhodamine-phalloidin (red) and anti-paxillin Ab (green). **B:** Parental MSS31 cells were plated onto type-I collagen coated dishes and incubated for 8 h in 5% FBS/ α MEM. Next, cells were treated with or without Y27632 (10 μ mol/L) for 30 min. Cells were then stained with rhodamine-phalloidin (red) and anti-paxillin Ab (green).

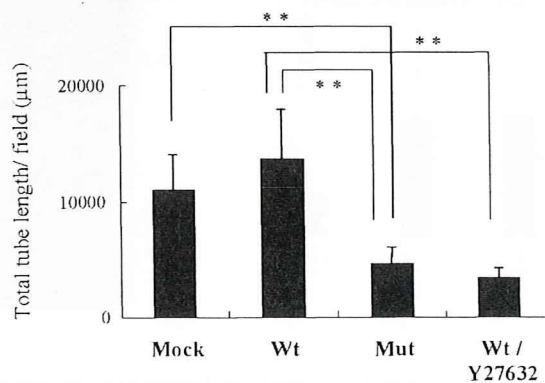
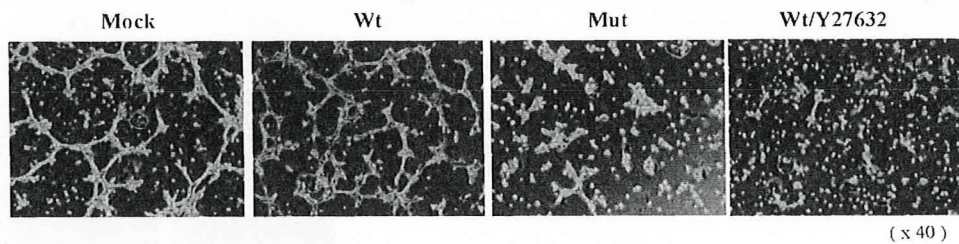


Fig. 4. Role of Rho kinase in alteration of network formation mediated by PILSAP. Transfectants were plated onto Matrigel and incubated in 5% FBS/ α MEM for 12 h. In some experiments, cells were incubated in the presence of Y27632 (10 μ mol/L). The total length of network structures per field ($\times 40$ magnification) was quantified with Soft Imaging System Analysis. The values are expressed as the mean \pm SD of four fields; ** $P < 0.01$.

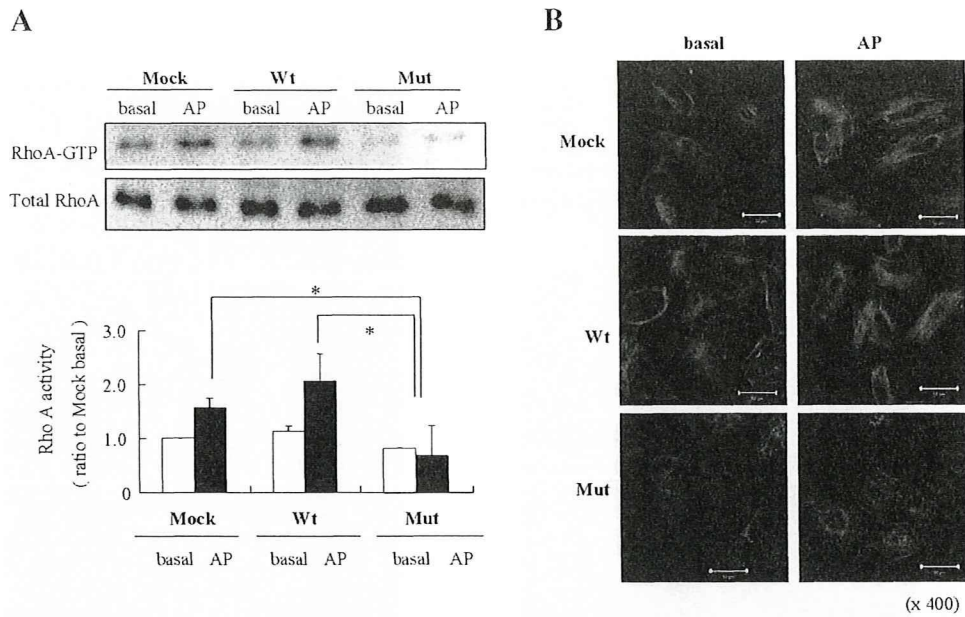


Fig. 5. Involvement of PILSAP in RhoA activation upon stimulation through PARs. **A:** Transfectants were incubated on type-I collagen for 10 h in 1% FBS/ α MEM, and then treated with or without PAR-1 AP (AP) (40 μ mol/L) for 5 min. Next, RhoA activity was determined. GTP-RhoA was quantified by density and normalized to that of total RhoA. The values are expressed as the mean \pm SD from four independent experiments; * $P < 0.05$. **B:** Transfectants were incubated on type-I collagen for 10 h in 1% FBS/ α MEM, and then treated with or without PAR-1 AP (AP) (40 μ mol/L) for 5 min. Subsequently, they were fixed and stained with rhodamine-phalloidine. A scale bar indicates 50 μ m.

induced stress fiber formation in Mock or Wt-PILSAP transfectants but not in Mut-PILSAP transfectants (Fig. 5B). LPA binds to the LPA receptor and activates RhoA for F-actin formation in ECs (Panetti, 2002). When transfectants were treated with LPA, RhoA activation was significantly lower in

Mut-PILSAP transfectants (Fig. 6A). Stress fiber formation was not induced by LPA in Mut-PILSAP transfectants (Fig. 6B). Moreover, LT or PILSAP siRNA inhibited LPA-stimulated activation of RhoA in parental MSS31 cells (Fig. 7A,B). These results indicate that the involvement of PILSAP in RhoA

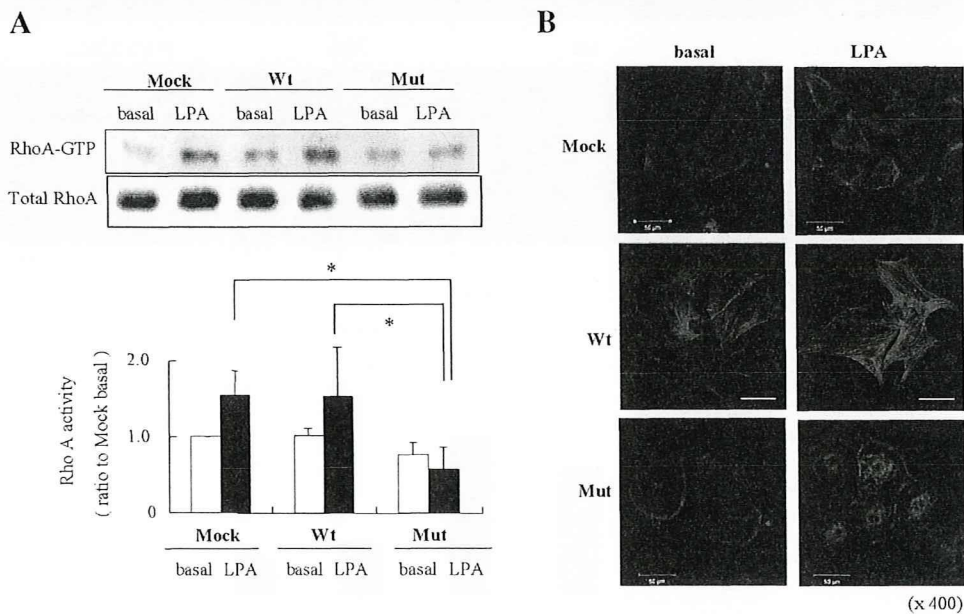


Fig. 6. Involvement of PILSAP in RhoA activation upon LPA stimulation. **A:** Transfectants were incubated on type-I collagen for 10 h in 1% FBS/ α MEM, and then treated with or without LPA (10 μ mol/L) for 5 min. Next, RhoA activity was determined. GTP-RhoA was quantified by density and normalized to that of total RhoA. The values are expressed as the mean \pm SD from four independent experiments; * $P < 0.05$. **B:** Transfectants were incubated on type-I collagen for 10 h in 1% FBS/ α MEM, and then treated with or without LPA (10 μ mol/L) for 5 min. Subsequently, they were fixed and stained with rhodamine-phalloidine. A scale bar indicates 50 μ m.