

Figure 3. Stabilization of complexes by use of cholesterol-conjugated siRNA. (A) The portion of siRNA dissociated from TEPA-PCL by PEGylation was separated by gel electrophoresis and stained with ethidium bromide. (B) The interaction between TEPA-PCL and siRNA was improved by modifying siRNA with cholesterol at the 3'-prime of the sense strand. The portion of siRNA or siRNA-C dissociated from TEPA-PCL by PEGylation was visualized by a similar procedure. (C, D) RNAi efficiency of siRNA-C transfected with TEPA-PCL was determined. A549-luc-C8 (C) or B16-F10-luc2 (D) cells (5×10^5 cells/well) were precultured overnight. TEPA-PCL complexed with siRNA (siLuc, siLuc-C; 10 pmol, N/P = 18 equiv (C), siCont-C, or siLuc2-C, 20 pmol, N/P = 18 equiv (D)) was added to these cells and removed after a 24 h incubation. These cells were subsequently cultured for additional 24 h and used for the luciferase assay. B16-F10-luc2 cells were also transfected with siLuc2 mixed with LFA2K according to the manufacturer's instructions. The control is that of cells not transfected with siRNA. Data are presented as the percentage of control luminescence intensity.

clearance from the bloodstream. The biodistribution of TEPA-PCL modified with 10% PEG2000 was almost similar to that of TEPA-PCL without PEG. On the other hand, TEPA-PCL modified with 20% PEG2000 avoided RES uptake and showed long-term circulation in the bloodstream. The length of PEG on TEPA-PCL also affected the avoidance of PEGylated TEPA-PCL from the RES uptake. TEPA-PCL modified with PEG6000 avoided RES uptake and showed long-term circulation in the bloodstream even when the percent molar ratio of DSPE-PEG to total lipids was 10%. The long-term circulation property of

TEPA-PCL modified with PEG6000 was superior to that with PEG2000. On the basis of these data, TEPA-PCL modified with 10% PEG6000 was adopted for subsequent experiments.

Stabilization of Complexes by Use of Cholesterol-Conjugated siRNA. Complex formation of PEGylated TEPA-PCL/siRNA was examined by electrophoresis in 1.5% acrylamide gels. As a result, free siRNA dissociated from TEPA-PCL was detected after PEGylation (Figure 3A). In contrast, siRNA was stably associated to TEPA-PCL by modifying the siRNA with cholesterol at the 3'-prime of the sense strand (Figure 3B). In the case of TEPA-PCL/siRNA-C complexes, free siRNA-C was not detected when they were formed at an N/P ratio of 18–30. In contrast, free siRNA dissociated from TEPA-PCL/siRNA complexes at all N/P ratios tested. The influence of cholesterol-conjugation of siRNA on knockdown efficiency was examined by using A549-luc-C8 cells and siLuc-C. TEPA-PCL complexed with siLuc or siLuc-C was used to transfect A549-luc-C8 cells. As a result, cholesterol-conjugation at the 3'-prime end of the sense strand of the siRNA did not affect the knockdown efficiency (Figure 3C). The knockdown efficiency of TEPA-PCL was also examined in another experimental model using B16-F10-luc2 cells and siLuc2-C. The knockdown efficiency of TEPA-PCL/siLuc2-C was higher than that obtained with LFA2K (Figure 3D).

NIRF Imaging of siRNA in Tumor-Bearing Mice. The biodistribution of AF750-siCont-C after intravenous injection in Colon26 NL-17 carcinoma-bearing mice was imaged *in vivo* by use of IVIS (Figure 4A). *In vivo* imaging data showed that fluorescent signals of AF750 were observed in the whole body of all mice at 10 min and 3 h after the injection. Accumulation of AF750 in the bladder was also observed in all groups up to 24 h after the injection. The total fluorescence of AF750 injected with TEPA-PCL was stronger than that without TEPA-PCL at 24 h after the injection. Importantly, AF750-siCont-C formulated in APRPG-TEPA-PCL accumulated in the tumor 24 h after the injection. *Ex vivo* images of each organ were obtained 96 h after the injection (Figure 4B). The total fluorescence of AF750 in the liver was observed with PEG-TEPA-PCL. In addition, strong fluorescent signals of AF750 in the tumor were observed only in the case of APRPG-TEPA-PCL.

DISCUSSION

Our previous results showed that siRNA formulated in cetyl-PEI-PCL showed efficient gene silencing in EGFP/HT-1080 cells and human umbilical vein endothelial cells.¹⁷ However, cetyl-PEI-PCL modified with PEG6000 accumulated in the RES immediately after intravenous injection into mice. We considered that PEG6000 modification could not negate the undesirable properties of cetyl-PEI-PCL because of the molecular size of PEI [1800]. On the basis of these results, DCP-TEPA was newly synthesized and liposomalized for systemic siRNA delivery. Since TEPA is a relatively short polycation, we expected it to be stably presented on the liposomal membrane and not to interfere with the beneficial effects of PEGylation. In fact, PEG6000 endowed TEPA-PCL with the property of long circulation. On the other hand, we previously reported gene delivery with liposomal polyamine (spermidine, spermine, and PEI[1800])-dialkyl (dilauryl (DLP), dimyristyl (DMP), and DCP) phosphate conjugates.¹⁴ Our previous results demonstrated that not only the polyamine but also the dialkyl parts affect their transfection efficiencies and that DCP is favorable for polyamine-mediated

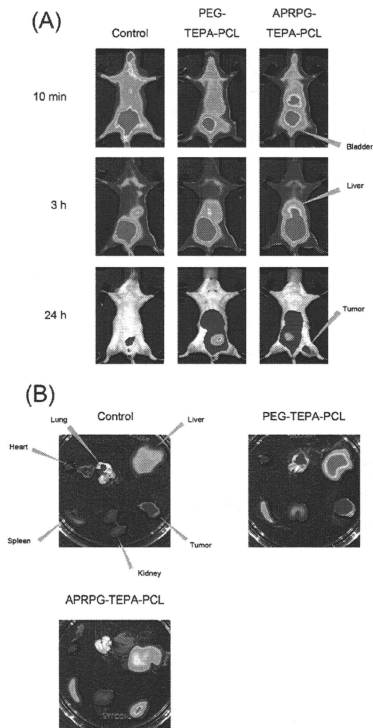


Figure 4. NIRF-imaging of APRPG-TEPA-PCL/AF750-siCont-C in tumor-bearing mice. (A, B) Colon26 NL-17 carcinoma-bearing BALB/c mice ($n = 3$) were intravenously injected with AF750-siCont-C ($15 \mu\text{g}$), PEG-TEPA-PCL/AF750-siCont-C or APRPG-TEPA-PCL/AF750-siCont-C ($15 \mu\text{g}$, $N/P = 18$ equiv) via the tail vein under anesthesia. The biodistribution of AF750 in these mice was assessed with a Xenogen IVIS Lumina System coupled to *Living Image* software for data acquisition. (A) The fluorescence of AF750 in these mice was acquired *in vivo* 10 min, 3 h, and 24 h after the injection. (B) The fluorescence of AF750 in each organ (heart, lungs, liver, spleen, kidneys, and tumor) was acquired *ex vivo* at 96 h after the injection.

gene delivery. For these reasons, we designed DCP-TEPA for systemic siRNA delivery.

In the present study, we first determined the formulation of TEPA-PCL/siRNA complexes by performing knockdown experiments. Interestingly, the amount of DCP-TEPA in PCL clearly affected the knockdown efficiency, but did not show dose dependency. This phenomenon might be related to the stability of TEPA-PCL, since the hydrophilic portion of DCP-TEPA is relatively large compared with conventional liposomal components. Our data suggest that the amount of DCP-TEPA in TEPA-

PCL and the N/P ratio of TEPA-PCL/siRNA are important for gene silencing. On the other hand, PEGylation of TEPA-PCL/siRNA complexes caused dissociation of a part of siRNA from the TEPA-PCL, even though the formation of TEPA-PCL/siRNA complexes is based on electrostatic interactions. However, we could prevent this undesirable phenomenon by adding hydrophobic cholesterol to complexes by using siRNA-C. It is still not presently clear how DSPE-PEG removed siRNA from the TEPA-PCL. Modification of siRNA with cholesterol at the 3'-prime of the sense strand did not affect the knockdown efficiency, as previously reported.¹⁰ Thus, modification of siRNA with certain compounds to improve the properties of nanoparticles could be a useful strategy and applicable to other situations. In addition, the transfection efficiency of TEPA-PCL/siRNA-C was higher than that of LFA2K. TEPA-PCL/siRNA-C would be internalized into cells by endocytosis and efficiently escape from endosomes possibly through the buffering effects, which have been observed in the case of cetyl-PEI[1800]-based PCL.¹⁵

The biodistribution study demonstrates that PEG6000 is suitable for PEGylation of TEPA-PCL. In general, the length of PEG2000 is adequate to endow neutral liposomes with long circulation. In fact, Doxil, which is a liposomal doxorubicin modified with MPEG2000, has such a property and is used worldwide as a cancer drug. In contrast, modification of TEPA-PCL with 10% PEG2000 did not alter the biodistribution of it after intravenous administration. This phenomenon might have been due to TEPA exposure on the liposomal surface despite PEGylation. Our data suggest that relatively long PEG chains would be needed to improve the biodistribution of positively charged liposomes in some cases. In addition, we modified PEGylated TEPA-PCL with APRPG for efficient delivery of siRNA into target cells since it is assumed that PEGylation of TEPA-PCL attenuates cellular uptake and gene silencing effects of siRNA formulated in TEPA-PCL. A steric barrier formed on the surface of liposomes by PEGylation is known to inhibit binding of liposomes to cells. In fact, our previous *in vitro* study using human umbilical vein endothelial cells revealed that PEGylation of cetyl-PEI-PCL attenuates cellular uptake of siRNA formulated in cetyl-PEI-PCL, whereas modification of PEGylated cetyl-PEI-PCL with APRPG improves this attenuation.²⁵

In the NIRF imaging, we obtained not only *in vivo* but also *ex vivo* data, since the fluorescence intensities of *in vivo* NIRF imaging are strongly affected by tissue depth in animals.¹⁸ As a result, both our *in vivo* and *ex vivo* data showed that AF750-siCont-C formulated in APRPG-TEPA-PCL accumulated in the tumor after intravenous injection into Colon26 NL-17 carcinoma-bearing mice. *In vivo* data also showed that the total fluorescent signals of AF750 observed in the whole body were almost similar between PEG-TEPA-PCL and APRPG-TEPA-PCL, suggesting that the presence of APRPG did not diminish the long-circulating property achieved by PEGylation. These data indicate that the accumulation of AF750-siCont-C formulated in APRPG-TEPA-PCL in the tumor would be achieved by both active targeting to angiogenic vessels and passive targeting by the EPR effect. These results are consistent with our previous results showing that modification of liposomes with APRPG-PEG conjugate should be useful for both drug and gene delivery to angiogenic vessels in tumors.^{22,26–28}

In conclusion, we developed TEPA-PCL for efficient delivery of siRNA into tumor cells and angiogenic endothelial cells. Furthermore, we also developed APRPG-TEPA-PCL for targeting angiogenic vessels and achieved systemic delivery of siRNA to the tumors in *in vivo* study.

■ ASSOCIATED CONTENT

Supporting Information. The ζ -potential of TEPA-PCL under various pH conditions (pH = approximately 4.1–7.9) was determined as an indicator of the buffering effects of TEPA in PCL and compared with that of 1,2-dioleoyl-3-trimethylammonium-propane (DOTAP) liposomes. The ζ -potential of TEPA-PCL under low pH conditions was higher than that under the neutral condition, whereas that of DOTAP liposomes did not change at any pH tested. This material is available free of charge via the Internet at <http://pubs.acs.org>.

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Suppression of Choroidal Neovascularization by Intravitreal Injection of Liposomal SU5416

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Objective: To clarify whether use of angiogenic vessel-homing peptide, Ala-Pro-Arg-Pro-Gly (APRPG)-modified liposomes encapsulating 3-[(2,4-dimethylpyrrol-5-yl) methylidene]-indolin-2-one (SU5416), an angiogenesis inhibitor, can inhibit the development of experimental choroidal neovascularization (CNV) in rats.

Methods: Liposomes were prepared using the thin-film hydration method. To set up the rat experimental CNV model, intense fundus laser photocoagulation at 6 spots per eye was performed on pigmented rats. After photocoagulation, the rats were divided into 4 groups (6 rats in each group): an APRPG-liposomal SU5416 treatment group and control groups treated with a balanced salt solution, APRPG liposomes, or soluble SU5416. Each rat received a single intravitreal injection immediately after the injury. One week or 2 weeks after laser injury, the extent of CNV was evaluated by perfu-

sion with high-molecular-weight fluorescein isothiocyanate-dextran.

Results: Two weeks after injection, the CNV area was significantly ($P < .05$) smaller in the APRPG-liposomal SU5416-treated group compared with the CNV area in the balanced salt solution- and APRPG liposome-treated groups.

Conclusion: Liposomes modified by APRPG and encapsulating SU5416 constitute a potential drug formulation for CNV treatment that would require only a single intravitreal injection.

Clinical Relevance: This liposomal delivery may enable the sustained release of small molecules and be a new treatment modality for CNV.

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CHOROIDAL NEOVASCULARIZATION (CNV) is a key pathologic change, causing severe vision loss as part of the pathogenesis of several diseases. The most prevalent disease of this type is exudative age-related macular degeneration (AMD), resulting in irreversible vision loss, which primarily affects people in developed countries.^{1,2} In relation to pathomorphologic disorders, various angiogenic growth factors, such as vascular endothelial growth factor (VEGF), have been reported³⁻⁵ to be expressed in CNV tissue and in retinal pigment epithelium (RPE) cells. Vascular endothelial growth factor is believed to be one of the major regulators in a variety of physiologic and pathologic angiogenesis processes, including CNV.⁶⁻⁹ In addition, the molecular expression of VEGF and its receptors (VEGFRs) is upregulated in rat experimental CNV models.^{3,10}

Two types of endothelium-specific tyrosine kinase receptors, Flt-1 (VEGFR1) and KDR/Flk-1 (VEGFR2), are known to

mediate the function of VEGF, as receptor molecules, in physiologic and pathologic neovascularization.^{11,13} In addition, VEGFR2 is thought to mediate VEGF-induced signals such as vascular hyperpermeability and most angiogenic properties, including mitogenicity/chemotactic activity.¹⁴

To date, 2 antiangiogenic therapies have been approved for the treatment of exudative AMD: pegaptanib sodium, an oligonucleotide (Macugen; Eyetech Pharmaceuticals/Pfizer, New York, New York), and ranibizumab, an anti-VEGF antibody Fab fragment (Lucentis; Genentech/Novartis, Basel, Switzerland). Another VEGF blocker in ongoing clinical studies is VEGF trap, a fusion protein of domain 2 from VEGFR1 and domain 3 from VEGFR2. This recombinant fusion protein has a high affinity for VEGF and interacts with all VEGF isoforms as well as placental growth factors. All clinical trials to date have demonstrated satisfactory results with intravitreal delivery of these agents in patients with AMD. However,

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there are serious clinical problems with these approaches. The desired therapeutic outcome often requires frequent injections. Repeated ocular injections are generally accompanied by risk of endophthalmitis, retinal detachment, and vitreous hemorrhage. Therefore, drug formulations, and especially drug delivery systems (DDSs), need to be improved to reduce the frequency of intravitreal injection. Vascular endothelial growth factor signal transduction inhibitors have been also used as antiangiogenic factors in chemotherapy. For example, sunitinib (Sutent, SU1248, Pfizer), a small molecule inhibitor of receptor tyrosine kinases of VEGFR and platelet-derived growth factor receptor, has been approved for the treatment of cancer.¹⁵ The antiangiogenic compound SU5416 (3-[(2,4-dimethylpyrrol-5-yl) methylidene]-indolin-2-one) is a potent and selective inhibitor of VEGFR2.¹⁶ It has been shown¹⁷ to suppress VEGF-mediated angiogenesis *in vitro* and *in vivo* through the inhibition of autophosphorylation of VEGFR2 by blocking the adenosine monophosphate-binding site within the kinase domain. In contrast to the successful results of phase 1 and 2 trials for the treatment of advanced cancers, SU5416 did not show any significant clinical benefit, and some patients experienced severe toxic reactions to the solvent used for dissolving SU5416, polythoxylylated castor oil.¹⁸

It has been demonstrated¹⁹ that the liposomal formulation of SU5416 does not require polythoxylylated castor oil for administration. Furthermore, a liposomal DDS using the angiogenic vessel-homing peptide Ala-Pro-Arg-Pro-Gly (APRPG) for active targeting of the drug carrier to the angiogenic site was useful in the treatment of tumor-bearing mice.²⁰⁻²² Thus, this chemical compound is expected to be a new drug for treatment of AMD if an appropriate and safe DDS is developed.

Liposomes are small lipid vesicles and one of the most advanced drug nanocarriers in DDS studies.²³ The use of liposomal vehicles to prolong clearance and limit peak concentration after intravitreal injection might avert the risks of frequent intravitreal injections.²⁴ The aim of the present study was to clarify whether SU5416 can be applied in CNV treatment using a liposomal DDS.

METHODS

ANIMALS

Sexually mature male Brown Norway rats (weight, 180-200 g; Japan SLC, Inc, Shizuoka, Japan) were maintained under 12-hour light and 12-hour dark conditions and fed a standard rat chow and water *ad libitum*. All experiments adhered to the Association for Research in Vision and Ophthalmology Statement for the Use of Animals in Ophthalmic and Vision Research²⁵ and Juntendo University's guidelines for care and use of laboratory animals.

PREPARATION OF LIPOSOMAL SU5416

Liposomes were prepared as previously described.²¹ In brief, DPPC (dipalmitoylphosphatidylcholine), POPC (palmitoyloleoylphosphatidylcholine), cholesterol, and SU5416 solutions in chloroform were mixed (10:10:5:1 molar ratio) and dried under reduced pressure to make a thin lipid film. The APRPG peptide

conjugated DSPE-PEG (distearoylphosphatidylethanolamine-polyethylene glycol) solution was added to the initial lipid solutions in the proportion of 10 mole percent (mol%) to phosphatidylcholine to modify the liposomes. For histochemical analysis, DiI-C₁₈ (Molecular Probes, Inc, Eugene, Oregon) was added to the initial lipid solutions in the proportion of 10 mol% to phosphatidylcholine. The thin lipid films were hydrated with 20mM HEPES (2-[4-(2-hydroxyethyl)-1-piperazinyl]ethanesulfonic acid)-buffered saline (pH 7.4), and the liposome solutions underwent 3 freeze-thaw cycles using liquid nitrogen. The size of the liposomes was then adjusted by extruding through a 100-nm pore polycarbonate filter. The particle size and zeta potential of liposomal SU5416 were measured (Zetasizer; Malvern Instruments, Malvern, Worcestershire, England), and these values were approximately 130 nm and -2 mV, respectively.

INDUCTION OF CNV

Laser-induced CNV is widely used as an animal model for exudative AMD.²⁶ Using this model, we observed new vessels from the choroid that invaded the subretinal space after photocoagulation. Laser photocoagulation was performed in 6 spots per eye around the optic disc (wavelength, 532 nm; power, 90 mW; spot diameter, 100 μ m) using a slit delivery system (NOVUS Spectra; Lumens, Tokyo, Japan), as described.^{27,28} After photocoagulation, the rats were divided into 4 experimental groups (6 rats in each group) and were given APRPG-liposomal SU5416, balanced salt solution (BSS), APRPG liposomes, or soluble SU5416 for administration of the drug. The SU5416 concentration used was 0.625 mg/mL. Immediately after photocoagulation, each rat received a single 10- μ L intravitreal injection using a 30-gauge needle.

VISUALIZING AND QUANTIFYING CNV

One week or 2 weeks after injury, blood flow was visualized by vascular perfusion with high-molecular-weight (2×10^5 -MW [2-MMW]) fluorescein isothiocyanate (FITC)-dextran (Sigma-Aldrich Corp, St Louis, Missouri) using the method described by D'Amato et al.²⁹ Briefly, animals were anesthetized with intraperitoneal injection of sodium pentobarbital, 30 mg (Nembutal; Dainippon Sumitomo Pharma Co, Ltd, Osaka, Japan) per kilogram of body weight. The left ventricle was then perfused with approximately 50 mL of lactated Ringer solution, followed with 20 mL of lactated Ringer solution containing 10% (wt/vol) gelatin and 5-mg/mL FITC-dextran. After perfusion, treated eyes were enucleated and fixed with 4% paraformaldehyde solution for 20 minutes. Retinal pigment epithelium-choroid-sclera flat mounts were obtained by hemisectioning the eye and peeling the neural retina away from the underlying RPE. Radial cuts allowed the eyecup to be laid flat onto a microscope slide with the RPE side facing up. Flat mounts were fixed by mounting medium (Vectashield; Vector Laboratories, Inc, Burlingame, California), and the specimens were examined with a microscope (Biorovo BZ-9000; Keyence, Osaka, Japan) equipped with an FITC-detecting filter unit. The CNV areas were photographed with a charge-coupled device camera and captured with an image analysis system (BZ-II; Keyence). The FITC-dextran-perfused vessels representing neovascularization were quantified by an experienced clinician (M.H.) delineating the perimeter of the fibrovascular membranes.

ACCUMULATION OF FLUORESCEN LIPOSOMAL SU5416 IN CNV

To investigate the accumulation of APRPG-liposomal SU5416 in CNV, double-labeling experiments including DiI-C₁₈ (red fluo-

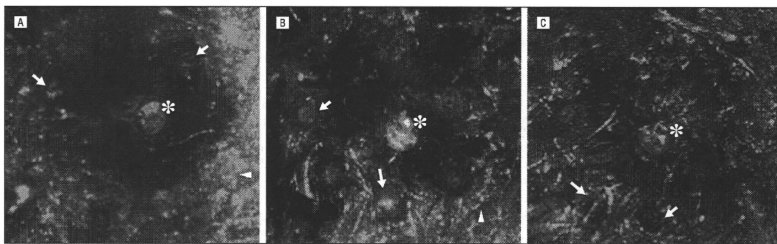


Figure 1. Liposomal SU5416 (3-[(2,4-dimethylpyrrol-5-yl) methylidene]-indolin-2-one) (red) and choroidal vascularization (green) double labeling. Localization of (Ala-Pro-Arg-Pro-Gly [APRPG]-free) liposomal SU5416 (A) and APRPG-liposomal SU5416 (B) 4 days after intravitreal injection, as well as APRPG-liposomal SU5416 at 2 weeks after injection (C). Asterisks indicate optic nerve; arrows, choroidal neovascularization region; arrowheads, normal choroidal vessels.

rescence) were performed. Just after laser injury, (APRPG-free) liposomal SU5416 and APRPG-liposomal SU5416 were intravitreally injected in rats. Four days or 2 weeks after injection, the accumulation of liposomal SU5416 in choroidal flat-mount was examined using an epifluorescence microscope.

STATISTICAL ANALYSIS

Experimental results of the quantitative analysis are expressed as the mean (SE). Data were analyzed by analysis of variance with the Tukey test. A probability of $P < .05$ was considered statistically significant.

RESULTS

To evaluate the effect of vascular-targeted liposomal SU5416 on CNV, the morphologic effect of the compound was examined by angiography using 2-MMW FITC-dextran and by epifluorescence microscopy. The rat experimental model was chosen for this study to determine the pharmacologic effect of the liposomal chemical compound on CNV, since this animal model of laser-induced CNV has been widely used^{20,31} in the evaluation of experimental AMD treatments. The rat model of laser-induced CNV has a high reproducibility.

As an initial experiment, we examined the accumulation of liposomal SU5416 after intravitreal injection by using DiI_{C18} (red fluorescence) to clarify whether the efficacy of liposomal SU5416 might be enhanced by the angiogenic-homing peptide (APRPG) or only by sustained release from liposomes. Four days after laser injury and intravitreal injection, significant amounts of APRPG-liposomal SU5416 recruited to the developing CNV region were compared with the control (APRPG-free liposomal SU5416) (Figure 1A and B). The APRPG-liposomal SU5416 remained in the CNV region 2 weeks after intravitreal injection (Figure 1C).

The typical CNV observed by fluorescent microscopy in the present study is shown in Figure 2. Two weeks after the injury, CNV was observed as relative hyperfluorescence areas overlying the laser lesions in the RPE-choroid-sclera flat mounts. Although CNV in the group treated with soluble SU5416 appeared to be somewhat inhibited (Figure 2C), the inhibitory effects on the induced CNV were enhanced when the animals

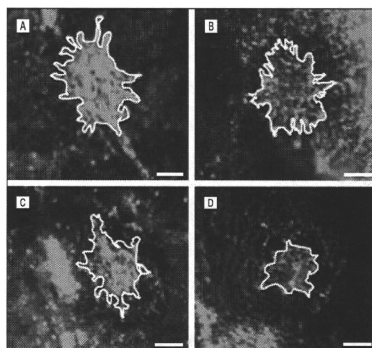


Figure 2. Morphologic changes of laser-induced choroidal neovascularization (CNV) after intravitreal injection. Typical fluorescein angiogram of retinal pigment epithelium 2 weeks after injury treated with balanced salt solution (A), Ala-Pro-Arg-Pro-Gly (APRPG) liposomes (B), soluble SU5416 (3-[(2,4-dimethylpyrrol-5-yl) methylidene]-indolin-2-one) (C), or APRPG peptide-modified liposomal SU5416 (D). Regions of CNV are outlined by a solid line. Bars represent 100 μm .

were treated with APRPG peptide-modified liposomal SU5416 (Figure 2D) compared with the controls (Figure 2A and B).

Attempts were then made to quantify the analysis by measuring the CNV areas in images of each experimental group. One week after laser injury and intravitreal injection, neovascular regions in RPE showed that the area of CNV in the soluble SU5416-treated group (19 598 [1128] μm^2) was significantly smaller than that in each of the other 2 groups (BSS, 25 092 [1584] μm^2 ; APRPG liposomes, 24 987 [1641] μm^2). No significant difference was observed between the group treated with APRPG-liposomal SU5416 (20 177 [1339] μm^2) and each of the other control groups (BSS and APRPG liposomes) (Figure 3A).

Data obtained from traced neovascular regions in RPE 2 weeks after injury showed that the area of CNV in the APRPG-liposomal SU5416-treated group (28 431 [2237]

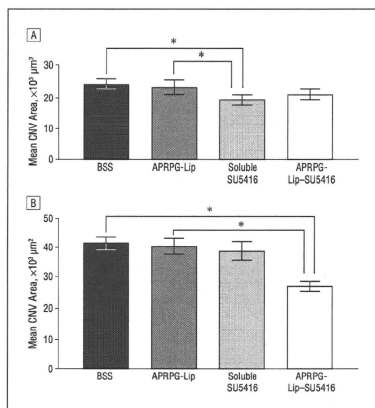


Figure 3. Quantitative analysis of the choroidal neovascularization (CNV) area in each experimental group 1 week (A) and 2 weeks (B) after intravitreal injection. The results are expressed as simple mean (SE) of CNV area ($n=6$). *Statistical significance between indicated groups at $P<.05$. APRPG-Lip indicates Ala-Pro-Arg-Pro-Gly-modified liposomes; BSS, balanced salt solution; and SU5416, 3-[(2,4-dimethylpyrrol-5-yl) methylidene]-indolin-2-one.

μm^2) was significantly smaller than that in each of the other 2 groups (BSS and APRPG liposomes). However, no significant difference was observed between the group treated with soluble SU5416 (38 160 [3500] μm^2) and that treated with APRPG liposomes (40 788 [2845] μm^2) or BSS (421 437 [2468] μm^2) (Figure 3B).

COMMENT

The data from this study clearly demonstrate that a single intravitreal injection of the vascular-targeted liposomal SU5416 significantly reduced laser-induced CNV in the experimental rat model. This implies that the liposomal compound may be useful in the treatment of CNV in clinical practice via its inhibitory effect on VEGF. Indeed, intravitreal injections using liposomal chemicals such as antibiotics (clindamycin,^{32,33} gentamicin,³⁴ and amikacin³⁵), a polyene antifungal (amphotericin B^{36,37}), an immunosuppressant (cyclosporine³⁸), and antimetabolites (flourouracil, fluorouridine^{39,40}) have been reported to be effective.

SU5416 inhibits the phosphorylation of VEGFR2 in a dose-dependent manner *in vitro*.¹⁸ It is assumed that SU5416 inhibits neovascularization by blocking VEGFR2 downstream signal transduction *in vivo*. Previous animal model studies on SU5416 demonstrated its tumor-suppressive effect. For example, a 73% decrease in tumor volume has been demonstrated¹⁸ by SU5416 administration in a null/null mouse model after rat C6 glioma cells were transplanted into the colon serosa. The inhibition of VEGFR2 phosphorylation by in-

traperitoneal injection of SU5416 once daily for 2 weeks has been shown in the rat model, and SU5416 also showed a tumor-suppressive effect in a rat experimental CNV model.⁴¹ In addition, apoptosis has been induced in murine CNV endothelial cells after administration of SU5416 by intraperitoneal injection every other day for 14 days.²⁸

Treatment using soluble SU5416 inhibited CNV development 1 week (but not 2 weeks) after injection compared with APRPG-liposomal SU5416 (Figure 3). Because APRPG-liposomal SU5416 was observed at the CNV region 2 weeks after intravitreal injection (Figure 1C), these results suggest that APRPG-liposomal SU5416 has a long-standing effect for CNV reduction.

Based on the inhibitory effects of SU5416 on CNV *in vivo*, the present study evaluated the validity of a novel DDS—a single intravitreal injection of an angiogenic vessel-homing peptide, APRPG—for the active targeting of drug carriers to angiogenic sites. Healthy choroidal vessels in the APRPG-liposomal SU5416-treated group were not noted (Figure 2). However, Shimotake et al¹² recently reported that VEGF inhibition could affect normal vessels as well as angiogenic vessels; further studies concerning the effect of liposomal SU5416 on healthy tissues are necessary to clarify the issue.

We believe that the liposomal DDS provided significant basic data for clinical use for the following reasons: (1) the efficiency of the liposomal method used in this study was satisfactory for use in an *in vivo* study, (2) it is believed that APRPG-liposomal SU5416 may be stable in the vitreous humor, and (3) APRPG has been reported²¹ in studies of tumor therapies to be effective in active targeting of drug carriers to angiogenic sites in the vitreous body.

Although further investigation of the precise mechanism and direct molecular basis of the effect of vitreous injection of APRPG-liposomal SU5416 on CNV inhibition is necessary, the findings in this study suggest that APRPG-liposomal SU5416 is a strong candidate for development as an agent for the treatment of conditions such as exudative AMD by a single vitreous injection. The liposomal delivery of small compounds may be a new treatment modality for CNV.

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PET imaging of brain cancer with positron emitter-labeled liposomes

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ABSTRACT

Since nanocarriers such as liposomes are known to accumulate in tumors of tumor-bearing animals, and those that have entrapped a positron emitter can be used to image a tumor by PET, we applied ¹⁸F-labeled 100-nm-sized liposomes for the imaging of brain tumors. Polyethylene glycol (PEG)-modified liposomes, which are known to accumulate in tumors by passive targeting and those modified with Ala-Pro-Arg-Pro-Gly, which are known to home into angiogenic sites were used. Those liposomes labeled with Dil fluorescence accumulated in a glioma implanted in a rat brain 1 h after the injection, although they did not accumulate in the normal brain tissues due to the protection afforded by the blood–brain barrier. Preformed liposomes were easily labeled with 1-[¹⁸F]fluoro-3,6-dioxatetracosane, and enabled the imaging of gliomas by PET with higher contrast than that obtained with [¹⁸F]deoxyfluoroglucose. In addition, the smallest tumor among those tested, having a diameter of 1 mm was successfully imaged by the liposomal ¹⁸F. Therefore, nanocarrier-based imaging of brain tumors is promising for the diagnosis of brain cancer and possible drug delivery-based therapy.

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1. Introduction

Brain cancers such as glioblastomas are known to be aggressive and invasive (Wen and Kesari, 2008). Therefore, diagnosis of brain cancer at the early stages is quite important. Positron emission tomography (PET) is one of the strong tools for diagnosis, therapeutic evaluation, and prognostic evaluation of cancer. [2-¹⁸F]-2-deoxyfluoro-D-glucose (FDG) is the most widely used positron emitter for cancer diagnosis (Scott et al., 2008; Nakamoto et al., 2009). However, the utility of FDG-PET imaging for detection of brain cancer is controversial due to the high demand for glucose in the brain (Takeda et al., 2003; Chen, 2007). To reduce this bothersome background various compounds such as [¹¹C]choline (Tian et al., 2004; Kato et al., 2008), [¹¹C]acetate (Yamamoto et al., 2008), and amino acid analogues such as [¹¹C]methionine (Hatakeyama et al., 2008; Jager et al., 2001), L-[methyl-¹¹C]methionine (Nojiri et al., 2009; Ullrich et al., 2009), O-[¹¹C]methyl-L-tyrosine (Ishiwata et al., 2004), O-[¹⁸F]fluoromethyl-L-tyrosine (Ishiwata et al., 2004),

O-[¹⁸F]fluoroethyl-L-tyrosine (Heiss et al., 1999; Langen et al., 2006; Floeth et al., 2007; Mehrkens et al., 2008; Pauleit et al., 2009), and O-[¹⁸F]fluoropropyl-L-tyrosine (Tang et al., 2003) have been synthesized and evaluated as PET imaging agents for the diagnosis and detection of recurrence of brain tumors. Among them, the amino acid analogues show relatively low accumulation in normal peripheral tissue (low tissue-to-blood ratio) and rapid blood clearance and have been used for detecting brain tumors and other tumors as well. We also demonstrated that the D-amino acid analogue O-[¹⁸F]fluoromethyl-D-tyrosine is useful for tumor imaging by PET (Urakami et al., 2009).

Nanomedicines such as liposomal drugs are known to accumulate in tumors due to the enhanced permeability of tumor blood vessels and the retention effect (Maeda et al., 2000, 2009). This drug delivery system (DDS) strategy is based on the nature of tumors: tumor cells demand oxygen and nutrition and cause angiogenesis for obtaining them, and angiogenic vessels are leaky enough to be permeated by nano-sized materials. Liposomes are known as one of the most effective drug carriers for cancer therapy. In liposomal DDS technologies, polyethylene glycol (PEG)-modified liposomes are useful drug carriers for cancer therapy; for they have a characteristically long circulation time in the bloodstream due to avoidance of being trapped by the reticuloendothelial system (RES) such as in the liver and spleen (Sakakibara et al., 1996;

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Gabizon, 2001). PEG-modified liposomes tend to accumulate in tumor tissues through passive targeting, and PEG-modified liposomes containing doxorubicin have been used in clinical cancer therapy.

We previously demonstrated that PEG-liposomes encapsulating [^{18}F]FDG accumulate in solid tumors and can be effectively imaged by PET (Oku et al., 1996). Although the encapsulation efficiency of [^{18}F]FDG is not so high, the result indicated the usefulness of liposomes or nanocarriers for cancer imaging. Moreover, actively targeting nanocarriers specifically associating with tumor cells or angiogenic vessels are an attractive approach. For example, we previously demonstrated that liposomes modified with a peptide specifically recognized by membrane type-1 matrix metalloproteinase (MT1-MMP), which is expressed on the surface of angiogenic endothelial cells, accumulate in tumors (Kondo et al., 2004). In that study, we encapsulated [^{18}F]FDG in the liposomes and determined the distribution of the liposomes by PET. We also performed *in vivo* biopanning of a phage-displayed peptide library using an angiogenesis mouse model to obtain specific probes for angiogenic endothelial cells, and identified the Ala-Pro-Arg-Pro-Gly (APRPG) motif as a novel peptide homing to angiogenic vessels (Oku et al., 2002; Asai et al., 2002). Liposomes modified with APRPG and labeled with [^3H]cholesterol hexadecylether actually accumulate in tumors of colon 26 NL-17 carcinoma-implanted mice (Maeda et al., 2004). Moreover, the accumulation of APRPG-modified liposomes in such tumors was also confirmed by PET analysis using [^{18}F]FDG-encapsulated liposomes (Maeda et al., 2006).

In the present study, we applied PEG-modified and APRPG-PEG-modified liposomes entrapping a positron-emitter to brain tumor imaging by PET. At first, we analyzed distribution of fluorescence-labeled liposomes in glioma-bearing brain of rats by use of fluorescence microscopy and an *in vivo* fluorescence-based imaging system. Then PET imaging of brain cancers was achieved by use of 1- ^{18}F]fluoro-3,6-dioxatetracosane ([^{18}F]SteP2)-labeled liposomes. Our results indicate that even a 1-mm diameter brain tumor could be imaged by PET, thus demonstrating the usefulness of these liposomes for PET imaging of brain cancer.

2. Materials and methods

2.1. Preparation of liposomes

All lipids were the products of Nippon Fine Chemical, Co. Ltd. (Takasago, Hyogo, Japan). Distearoylphosphatidylcholine (DSPC) and cholesterol along with DSPE-PEG or DSPE-PEG-APRPG (10:5:1 as a molar ratio) were dissolved in chloroform/*t*-butanol to formulate PEG-modified liposomes (PEG-liposomes) or APRPG-PEG-modified liposomes (APRPG-liposomes), respectively. For fluorescence labeling of liposomes, 1,1'-dioctadecyl-3,3,3',3'-tetramethyl-indocarbocyanine perchlorate (DIL, Molecular Probes Inc., Eugene, OR, USA) was added to the initial chloroform/*t*-butanol solution at a concentration of 1 mol% of DSPC. After the lipids had been dried under reduced pressure and stored *in vacuo* for at least 1 h, liposomes were prepared by hydration of the thin lipid film with 0.3 M glucose solution and frozen and thawed for 3 cycles by using liquid nitrogen. Then, the liposomes were sized by extruding them three through a polycarbonate membrane filter with 100-nm pores (Nuclepore track-Etch Membrane, Lipex). Particle size and ζ -potential of PEG-liposomes and APRPG-liposomes were measured by use of a Zetasizer Nano ZS (MALVERN, Worcester-shire UK, USA) and found to be 113 nm and -2.0 mV, respectively, for PEG-liposomes and 107 nm and -3.3 mV, respectively, for the APRPG-liposomes.

2.2. Preparation of brain cancer-bearing model rats

The glioma-bearing rat model was prepared by a modification of the procedure described previously (Takeda et al., 2003). Nine-week-old Fischer 344 male rats (Japan SLC, Hamamatsu, Japan) were cared for according to the Animal Facility Guidelines of the University of Shizuoka. For the implantation of tumor cells, the rats were anesthetized with chloral hydrate (40 mg/kg) in saline and individually placed in a stereotaxic apparatus. C6 glioma cells were maintained in Dulbecco's modified Eagle medium (DMEM, Wako Fine Chemical Co. Ltd., Osaka, Japan) supplemented with 10% heat-inactivated fetal bovine serum (Japan Bioserum Co. Ltd., Japan), penicillin G (100 U/mL) and streptomycin (100 $\mu\text{g}/\text{mL}$) at 37 °C in 5% CO₂ atmosphere. After harvesting of these cells, the cells were suspended in DMEM containing 1% gelatin (2×10^7 cells/mL). Ten microliters of the cell suspension or vehicle (DMEM containing 1% gelatin) was injected at a rate of 0.7 $\mu\text{L}/\text{min}$ into the left midbrain of each rat (4.3 mm posterior to bregma, 3.9 mm lateral to the midline suture, and 7.0 mm in depth) with an infusion pump (11 Plus, Harvard Apparatus, USA).

2.3. Intratumoral localization of Dil-labeled liposomes

C6 glioma cells (2.0×10^5 cells/rat) were inoculated as described above. Dil-labeled liposomes were administered via a tail vein of the rat at 11 days after tumor implantation. One hour or 24 h after the injection of liposomes (0.5 mL/rat, 5 μmol as DSPC), the rats were anesthetized with chloral hydrate, perfused with ca. 200 mL saline and with ca. 100 mL 20% formaldehyde for the fixation. The brain was removed and sliced at a 2-mm thickness with a cryostat microtome (HM 505E, Microm, Walldorf, Germany). The slices were then incubated twice in PBS for 4 h each time at 4 °C, and then overnight in 30% sucrose solution at 4 °C. After removal of water, the slices were embedded in the optimal cutting temperature compound (Sakura Finetech, Co. Ltd., Tokyo, Japan) and frozen at -80 °C. Brain sections (10 μm) were prepared with a cryostat microtome, and mounted on MAS-coated slides (Matsunami Class Ind., Ltd., Japan) for observation with a fluorescence microscope (IMT-2, Olympus, Japan). Then, the slices were stained with hematoxylin and eosin (HE) and observed under the microscope.

2.4. Ex vivo imaging of the distribution of Dil-labeled liposomes

Dil-labeled liposomes (0.5 mL/rat, 5 μmol as DSPC) were administered via a tail vein of a rat at 9 days after the implantation of C6 glioma cells. One hour after the injection, the rats were sacrificed; and brain slices (2-mm thickness) were then prepared with a cryostat microtome without perfusion and fixation. Fluorescent images of those slices were obtained with a fluorescence imaging system (IVIS Lumina, Xenogen).

2.5. Preparation of positron emitter-labeled liposomes

Preparation of 1- ^{18}F]fluoro-3,6-dioxatetracosane (SteP2) and liposome labeling by the SphIT method were performed as described previously (Urakami et al., 2007). About 100 MBq of [^{18}F]SteP2 in ethanol solution was transferred to a glass test tube, and the solvent was removed completely at 90 °C with a helium gas flow. Liposomal solution (1 mL, 10 μmol as DSPC) was added to the vial having a thin film of ^{18}F -radiolabeled compound and incubated at 65 °C for 15 min with 5-s mixing by a vortex stirrer every 3 min. After the incubation, the liposomal solution was washed with PBS by centrifugation at 100,000 $\times g$ for 15 min (Beckman, Fullerton, CA, USA), and the pellet was resuspended and diluted in PBS to make a 22 MBq/mL solution. Radioactivity was measured with a curie meter (IGC-3, Aloka, Japan).

Rats anesthetized with chloral hydrate at 11 days after the implantation of C6 glioma cells were placed on an animal CT (Clairvivo CT, Shimadzu, Japan) to obtain CT images. Then, [^{18}F]SteP2-labeled liposomes (10 MBq/rat) were administered via a tail vein. PET scans were started immediately after the injection and continued for 60 min by use of an animal PET apparatus (Clairvivo PET, Shimadzu, Japan).

FDG-PET was similarly performed. [^{18}F]FDG (10 MBq/rat) was injected into the rats after having obtained the CT images.

2.6. PET analysis and autoradiography

Before PET analysis, rats anesthetized with chloral hydrate at 11 days after the implantation of C6 glioma cells were placed on the animal CT to obtain CT images. Then, [^{18}F]SteP2-labeled liposomes (10 MBq/rat) were administered via a tail vein. Then the PET scan was started immediately after the injection and continued for 60 min. FDG-PET was similarly performed. [^{18}F]FDG (10 MBq/rat) was injected into the rats after having obtained the CT images.

After the PET scan, the rats were sacrificed; and the brain was then excised. Thereafter, 2-mm slices were prepared and set on an imaging plate. Autoradiograms were obtained by using a bioimaging analyzer system (BAS2000, Fuji Film, Japan).

2.7. Biodistribution of liposomes

Glioma-bearing rats were injected with [^{18}F]SteP2-labeled PEG-liposomes (10 MBq/rat) via a tail vein as described above. At 1 h after the injection, the rats were sacrificed, and blood and organs (heart, lung, liver, spleen kidney normal part of brain, brain tumor) were removed. The radioactivity of blood and each organ was measured by using an auto gamma counter (1480 Wizard 3, Perkin Elmer, USA).

2.8. Statistical analysis

Differences between groups were evaluated by analysis of variance (ANOVA) with the Tukey *post hoc* test.

3. Results

3.1. Intratumoral distribution of fluorescence-labeled PEG- and APRPG-liposomes

At first, we confirmed that the angiogenic vessels formed by experimental glioma in the rat brain were leaky enough to allow permeation by nanomedicines such as liposomes. As shown in Fig. 1B and E, the Dil fluorescence of Dil-labeled PEG-liposomes and APRPG-liposomes, respectively, was observed in the glioma at 1 h after intravenous injection of the liposomes. Since Dil fluorescence was not observed in the normal brain tissues near the tumor, we concluded that the liposomes had accumulated in the tumor through angiogenic vessels. The fluorescence intensity was increased at 24 h after the injection (Fig. 1C and F). Interestingly, the Dil fluorescence of the PEG-liposomes accumulated widely in the tumor, although that of APRPG-liposomes accumulated intensely in rather specific areas, suggesting that these liposomes had accumulated in the angiogenic vessels. These patterns of liposomal fluorescence are consistent with our previous data obtained by use of tumor-bearing mice that had been implanted subcutaneously with C26 NL-17 colon carcinoma (Maeda et al., 2006).

To understand the structure of the brain vasculature, we examined normal and glioma-implanted brain specimens after HE-staining. As shown in Fig. 2, capillaries in the normal brain tissue were covered with astrocytes (Fig. 2A); whereas capillaries in the brain tumor tissues were surrounded directly by glioma cells (Fig. 2B). These results support the idea that the blood–brain barrier (BBB) is immature in angiogenic vessels of brain tumors. The possible architecture based on our observation and the literature (Standring, 2005) is shown in Fig. 2C and D.

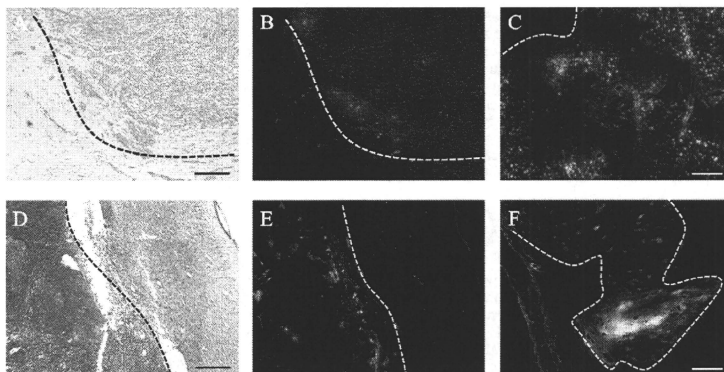


Fig. 1. Intratumoral distribution of fluorescence-labeled liposomes in brain tumor after intravenous injection. Rats were intravenously injected with Dil-labeled PEG-liposomes (A–C) or APRPG-liposomes (D–F) at day 11 after implantation of C6 glioma cells into the left midbrain. At 1 h (A, B, D, and E) or 24 h (C and F) after the injection, the brain tumors were dissected; and 10- μm frozen sections were prepared as described in Section 2. Then, liposomal distribution in the brain was observed under a fluorescence microscope (B, C, E, and F). Red fluorescence shows the liposomal localization. A and D show the images stained with hematoxylin and eosin, corresponding to the fluorescent images of “B” and “E”, respectively. The dotted lines show the borders between normal and tumor tissue. Deep purple regions in “A” and “D” indicate the tumor tissues. The scale bar represents 100 μm .

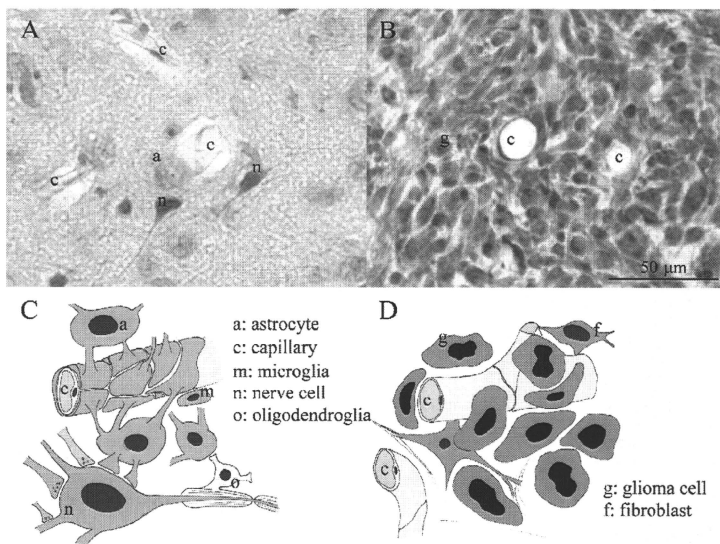


Fig. 2. Structure around blood capillaries in normal brain and brain tumor tissues. A rat brain tumor model was prepared by the implantation of C6 glioma cells into the left midbrain. The brain was dissected, and 10- μ m sections of normal right brain (A) and tumor-implanted left brain (B) were stained with hematoxylin and eosin. Schematic interpretation of the histology around blood capillaries in normal brain (C) and brain tumor tissues (D) are shown.

3.2. *Ex vivo* study on the accumulation of fluorescence-labeled liposomes in brain tumor

Although microscopic study provided data on the intratumoral distribution of the liposomes, the liposomal distribution in the whole brain could not be understood. Therefore, we next examined the distribution of Dil-labeled liposomes in the glioma-bearing brain by use of whole-brain slices. As a result, both PEG-liposomes and APRPG-liposomes accumulated in the brain tumor 1 h after the injection via a tail vein (Fig. 3B and D, respectively). No specific accumulation could be observed in the sham-operated brain after the injection of the Dil-labeled PEG-liposomes (Fig. 3E and F). The fluorescence intensity of the tumor region was 4.0-fold and 3.1-fold higher than that of the contralateral regions after the injection of Dil-labeled PEG-liposomes and Dil-labeled APRPG-liposomes, respectively.

3.3. Brain tumor imaging with positron-emitter-labeled liposomes by PET

Finally, PEG-liposomes and APRPG-liposomes were labeled with the positron emitter [18 F]SteP2, by the SophT method, and injected into the glioma-bearing mice. As shown in Fig. 4, the positron emitter accumulated in the tumor region and imaged the tumor after injection in PEG-liposomes (Fig. 4, top panel) or APRPG-liposomes (Fig. 4, middle panel). Interestingly, the other regions of the brain showed a low background. On the contrary, [18 F]FDG imaged the whole brain, although the accumulation was higher in the tumor region (Fig. 4, bottom panel). BAS images (autoradiograms) shown in right panel confirmed the region of tumor. The tumor sizes of preparations varied to some extent in the present experiment, and

the smallest tumor was revealed to be only about 1 mm in diameter. Interestingly, this small tumor was imaged by PET but was hardly detectable by CT when [18 F]SteP2-labeled APRPG-liposomes had been injected (Fig. 5).

The biodistribution of 18 F at 1 h after the injection of [18 F]SteP2-labeled PEG-liposomes or [18 F]FDG is shown in Fig. 6. The liposome-bearing label was maintained in the bloodstream and highly accumulated in the spleen. This result is consistent with our previous study (Maeda et al., 2004). In the brain, 18 F in the liposomes or as FDG significantly accumulated more in the tumor than in the normal tissues. In contrast to the distribution of 18 F after injection of liposomes bearing it, 18 F as FDG was cleared from the bloodstream quite fast and accumulated in the heart and normal region of brain.

4. Discussion

Since the usefulness of FDG-PET for the detection of brain tumors is controversial (Takeda et al., 2003; Chen, 2007), other PET imaging agents have been developed. In the present study, we aimed at imaging brain tumors not by the synthesis of molecules specifically taken up by tumor cells, but by use of DDS technology. Nanocarriers such as liposomes are well known to accumulate in solid tumors, especially in hypervascular tumors, due to the EPR effect (Maeda et al., 2000, 2009). In fact, liposomal radionuclides show effective tumor imaging (Oku et al., 1996; Geng et al., 2004; Elbayoumi and Torchilin, 2006). Moreover, liposome-based radionuclide delivery is not only useful for cancer imaging but also for cancer therapy when the carriers entrapping radiopharmaceuticals or other anticancer agents are used (Kostarelos and

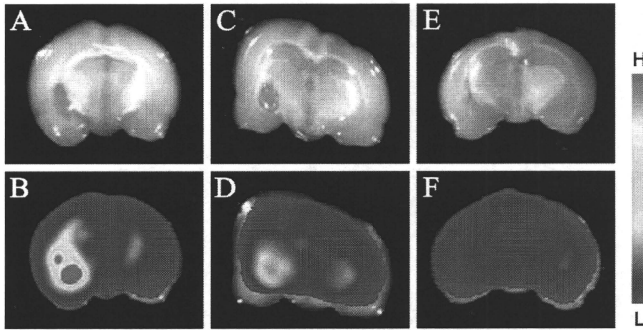


Fig. 3. Ex vivo imaging of liposomal distribution in rat brain bearing glioma. Rats were intravenously injected with Dil-labeled PEG-liposomes (A, B, E, and F) or APRPG-liposomes (C and D) at day 9 after implantation of C6 glioma cells (A–D) or medium alone (E and F) into the left midbrain. At 1 h after the injection, the brains were sliced into 2-mm sections, and the distribution of liposomes labeled with Dil was scanned with a fluorescence imaging system, IVIS (B, D, and F). Photos are also shown (A, C, and E). Two separate experiments gave similar results.

Emfietzoglou, 2000; Syme et al., 2003; Hamoudeh et al., 2008). Another advantage of carrier-based imaging is the active targeting by modifying carriers with some specific probes. We previously modified liposomes with angiogenic vessel-specific peptides and

observed the efficient delivery of targeting liposomes to cancer by PET (Kondo et al., 2004; Maeda et al., 2006). Therefore, cancer imaging with DDS technology has various advantages including cancer treatment.

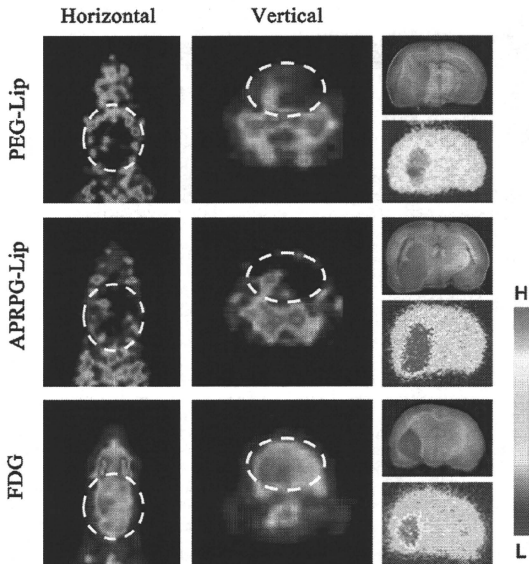


Fig. 4. PET imaging of brain tumor with [^{18}F]SteP2-liposomes and [^{18}F]-FDG. Rats were intravenously injected with 10 MBq of [^{18}F]SteP2-labeled PEG-liposomes (top panel), [^{18}F]SteP2-labeled APRPG-liposomes (middle panel) or [^{18}F]FDG (bottom panel) at day 11 after implantation of C6 glioma cells into the left midbrain. The biodistribution pattern of samples was determined by taking 1 frame/min for 1 h with the Clairvivo PET, and averaged data from 40 to 60 min are shown. Horizontal (left panel) and vertical (center panel) images are shown. After PET imaging, the brains were sliced into 2-mm sections, and the autoradiograms (right lower panels) were obtained with a bioimaging analyzer BAS 2000, and the pictures were taken (right upper panels). The ovals formed by the white dotted line show the estimated brain position.

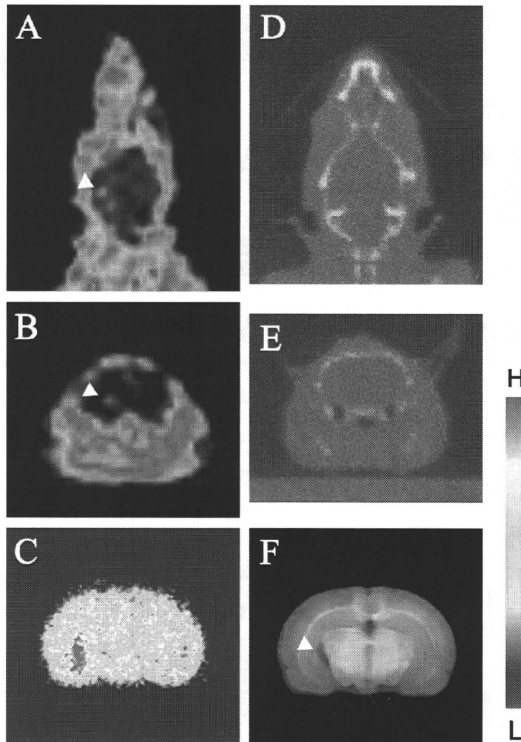


Fig. 5. PET imaging of small brain tumor with [^{18}F]SteP2-liposomes. PET imaging was performed as described in the legend of Fig. 4. Even the smallest tumor, 1-mm diameter, seen among the prepared model rats was imaged by PET. PET images (A and B) and CT images (D and E) of a rat brain are shown. Horizontal (A and D) and vertical (B and E) images are shown. (C) BAS image of a rat brain slice. (F) Photo of the brain slice. Arrowheads indicate the tumor.

It is well known that glioblastoma, the most prevalent brain cancer, is a hypervascular tumor due to angiogenesis (Verhoeff et al., 2009; Brastianos and Batchelor, 2009); and thus nanocarriers such as liposomes would be expected to accumulate in the brain cancer, since the permeability of tumor angiogenic vessels was higher than that of normal blood vessels. Moreover, brain blood vessels especially protect from the penetration of some molecules due to the BBB. Therefore, liposomes or other nanocarriers would be specifically accumulated into brain tumor region after circulating bloodstream, and be useful for brain cancer imaging. For labeling nanocarriers quite conveniently with PET probes, we previously synthesized a novel ^{18}F -labeled amphiphilic compound known as 1- ^{18}F fluoro-3,6-dioxatetracosane (Urakami et al., 2007). There are several methods for labeling liposomes, although radionuclides need to be incorporated during formulation of the liposomes in most of those methods. For example, to label liposomes with [^{18}F]FDG, [^{18}F]FDG was incorporated into liposomal aqueous phase during hydration step of liposomal lipids. In addition, the encapsulation efficiency of [^{18}F]FDG into the liposomes was very low. The

advantage of the "solid-phase transition" (SopHT) method is that it is applicable to preformed liposomes with quite high labeling efficiency. Moreover, [^{18}F]SteP2 in DSPC-based liposomes is stable in the presence of serum (Urakami et al., 2007). This universal method of liposomal modification can be used for various kinds of liposomes and lipid nanoparticles and bring the enhanced detection of a target tissue.

Aim of the present study is to demonstrate the application of positron-labeled liposomes for brain tumor imaging by PET. We firstly examined the intratumoral distribution of fluorescence-labeled PEG- and APRPG-liposomes by fluorescence microscopy, and observed the accumulation of both DiI-labeled PEG-liposomes and APRPG-liposomes in the brain tumor region at 1 h after intravenous injection of the liposomes, although the intratumoral distribution of each liposome was different. We previously demonstrated that PEG-liposomes were accumulated around neovessels and fluorescence of DiI-labeled-APRPG-liposomes was colocalized with CD31 stain in colon 26 NL-17 solid tumor (Maeda et al., 2006). Therefore, the differential accumulation

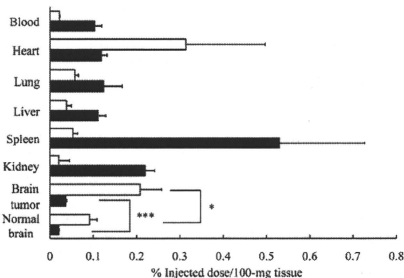


Fig. 6. Biodistribution of ¹⁸F after injection of [¹⁸F]SteP2-labeled PEG-liposomes or [¹⁸F]-FDG. Injection of [¹⁸F]SteP2-labeled liposomes and [¹⁸F]-FDG was performed as described in the legend of Fig. 4. At 1 h after the injection, blood was collected, and the selected organs were then excised for the determination of the radioactivity. Data are presented as the mean distribution of ¹⁸F after injection of [¹⁸F]SteP2-labeled PEG-liposomes (closed bars, n=3) or [¹⁸F]FDG (open bars, n=4). Significant differences between normal and tumor tissues are shown: *p<0.05, ***p<0.001.

of PEG-liposomes and APRPG-liposomes observed in the tumor region, might be caused by the APRPG-modification of the latter liposomes.

In the present study HE-staining of normal brain tissue indicated that the capillaries were wrapped with astrocytes, whose arrangement may strengthen the BBB function (Standing, 2005). In contrast, astrocytes and nerve cells were not observed in the brain tumor tissues, and the capillaries directly faced the surrounding glioma cells: This might support the extravasation of PEG-liposomes into the tumor tissues. On the other hand, APRPG-liposomes might specifically accumulate in angiogenic vessels in brain tumor tissue. *Ex vivo* imaging of Dil with the *in vivo* fluorescence imaging system indicated the accumulation of both PEG-liposomes and APRPG-liposomes in the brain tumor was 3–4 fold higher than that in the surrounding normal brain tissue.

Our PET study indicated the accumulation of ¹⁸F-labeled liposomes and imaging of glioma in the tumor-bearing rat brain. The tumor images obtained with liposomes were clearer than those obtained with FDG-PET. In the case of [¹⁸F]FDG imaging, the background level represented by imaging of normal brain region was high, since persistent demand for glucose to normal brain cells (Fig. 5). In contrast, brain tumor was specifically imaged by using [¹⁸F]-labeled PEG- or APRPG-liposomes with quite low background. Additionally, those liposomes also can be used for the brain cancer therapy.

The efficiency for tumor imaging by both [¹⁸F]SteP2-labeled PEG-liposomes and APRPG-liposomes was similar, although the intratumoral distribution was different between PEG-liposomes and APRPG-liposomes. These data are consistent with our previous results showing that the accumulation of PEG-liposomes and APRPG-liposomes in solid tumors was not much different (Maeda et al., 2004). Therefore, we conclude that [¹⁸F]-labeled liposomes having either passive or active targeting characteristics are useful for brain tumor imaging: However, active targeting liposomes to tumor angiogenic endothelium would be more potent when these are used for both imaging and therapy. In the present study, it should be also noted that a tumor with a diameter of only 1 mm could be successfully imaged by using [¹⁸F]SteP2-labeled liposomes.

5. Conclusions

In the present study, brain tumor imaging by a DDS nanocarrier, i.e., liposomes, was explored. Liposomes were ¹⁸F-labeled by the SophT method and injected intravenously into rats bearing a glioma-type brain tumor. Liposomes failed to accumulate in the normal surrounding brain tissue due to BBB protection and a brain tumor was specifically imaged with the liposomes via the different structure of brain tumor vessels. Since radiopharmaceutics or anticancer agents can be incorporated into DDS nanocarriers, these nanocarriers including liposomes should be useful for diagnosis and therapy of brain cancer.

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Cancer antineovascular therapy with liposome drug delivery systems targeted to BiP/GRP78

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Angiogenesis is crucial for tumor growth and hematogenous metastasis. Specifically expressed and functional protein molecules in angiogenic endothelial cells, especially on the plasma membrane, may be molecular targets for antiangiogenic drugs and drug delivery systems (DDS) in cancer therapy. To discover such target molecules, we performed subcellular proteome analysis of human umbilical vein endothelial cells (HUVECs) treated with or without vascular endothelial growth factor (VEGF) using 2-dimensional difference in-gel electrophoresis (2D-DIGE) and matrix-assisted laser desorption/ionization tandem time-of-flight mass spectrometry (MALDI-TOF/TOF-MS). Among the identified proteins, BiP/GRP78, a molecular chaperone, was highly expressed in the membrane/organelle fraction of HUVECs after VEGF treatment. The involvement of BiP in VEGF-induced angiogenesis was examined by RNA interference. BiP knockdown significantly suppressed VEGF-induced endothelial cell proliferation and VEGF-induced phosphorylation of extracellular-regulated kinase 1/2, phospholipase C- γ , and VEGF receptor-2 in HUVECs. Cell surface biotinylation analysis revealed that the cell surface expression of BiP was elevated in VEGF-activated HUVECs. Aiming to apply BiP to a target molecule in liposomal DDS, we developed liposomes modified with the WIFPWIQL peptide, which has been shown to bind to BiP, and investigated its potential for cancer therapy. The WIFPWIQL-modified liposomes (WIFPWIQL liposomes) were significantly taken up by VEGF-activated HUVECs as compared to peptide-unmodified liposomes. WIFPWIQL liposomes appeared to accumulate in tumor endothelial cells *in vivo*. WIFPWIQL liposomes containing doxorubicin significantly suppressed tumor growth and prolonged the survival of colon26 NL-17 carcinoma cell-bearing mice. In summary, BiP may regulate VEGF-induced endothelial cell proliferation through VEGFR-2-mediated signaling and be an effective target molecule for cancer antineovascular therapy.

Angiogenesis, the process of sprouting new blood vessels from preexisting vessels, is critical for tumor growth and blood-borne metastasis.¹ The overall process of angiogenesis is regulated by the balance between proangiogenic and anti-angiogenic factors secreted from several cell types in a tumor

microenvironment, such as tumor cells, fibroblasts and macrophages.^{2,3} Endothelial cells in tumors differentially express various protein molecules compared to those in normal tissues.⁴ In intratumoral regions, under hypoxic conditions, tumor cells produce vascular endothelial growth factor

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Abbreviations: 2D-DIGE: 2-dimensional difference in-gel electrophoresis; BiP: immunoglobulin heavy-chain binding protein; DDS: drug delivery systems; DSPC: distearylphosphatidylcholine; DSPG: distearylphosphatidylglycerol; DSPE-PEG: distearylphosphatidylethanolamine conjugated polyethyleneglycol 2000; DOX: doxorubicin; EBM-2: endothelial basal medium-2; EGM-2: endothelial growth medium-2; ER: endoplasmic reticulum; ERK: extracellular-regulated kinase; FBS: fetal bovine serum; GFP: green fluorescent protein; GRP78: glucose-regulated protein 78 kDa; HUVECs: human umbilical vein endothelial cells; MALDI-TOF/TOF-MS: matrix-assisted laser desorption/ionization tandem time of flight mass spectrometry; MAPK: mitogen-activated protein kinase; PEG-Lip-DOX: PEG-liposomal DOX; PLC γ : phospholipase C- γ ; RNAi: RNA interference; siRNA: small interfering RNA; VEGF: vascular endothelial growth factor; WIFPWIQL liposomes: WIFPWIQL peptide-modified liposomes; WIFPWIQL-Lip-DOX: WIFPWIQL-liposomal DOX
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(VEGF), a principal proangiogenic factor, through hypoxia-inducible factor 1-dependent transcription.⁵ VEGF and its receptor (VEGFR) have been well studied in basic and clinical cancer research, and inhibitors of VEGF signal transduction have been developed as anticancer drugs.⁶ The activation of VEGFRs initiates the proliferation, migration and invasion of endothelial cells. VEGF induces the phosphorylation of VEGFR tyrosine kinases, and thereby activates mitogen-activated protein kinase (MAPK) mediated by phospholipase C- γ (PLC γ) or the AKT/PKB signal cascade mediated by phosphatidylinositol-3 kinase (PI3K).⁷

Comparative proteome analysis is a powerful tool to find differentially expressed proteins,^{8,9} and it is rendered more powerful by combining it with subcellular fractionation, whereby it can identify low-abundance proteins in cellular organelles by subtracting out common protein contaminants.^{10,11} The proteins on the plasma membrane play a role in fundamental cellular processes, including ion and solute transports, signal transduction or cell adhesion. Since alterations in the expression of membrane proteins are often related to diseases, such proteins may become molecular targets for diagnostic and therapeutic approaches.¹²⁻¹⁴ Comprehensive profiling of these proteins may also provide a better understanding of how a cell responds to a variety of intracellular and extracellular signals. Therefore, expression profiling of membrane proteins in specific cell types under defined conditions has become a central point of many molecular biology investigations.

Inhibition of angiogenesis has curative benefits such as inhibition of primary tumor growth as well as suppression of blood-borne metastasis.¹⁵ Therapeutic vascular targeting has been categorized into antiangiogenic approaches, which aim to prevent the neovascularization processes in tumors, whereas vascular-disrupting approaches—using vascular disrupting agents—aim to damage the established tumor vasculature.¹⁶ Antineovascular therapy is a strategy targeted against angiogenic neovasculature by using liposome drug delivery systems (DDS).¹⁷⁻¹⁹ We and other groups have previously reported that tumor vasculature-targeted liposomes can deliver encapsulated agents to angiogenic endothelial cells and have demonstrated their therapeutic advantages.^{20,21} Cell surface proteins specifically expressed in angiogenic endothelial cells are possible molecular targets for DDS because these cells are in direct contact with the blood stream, making it easy for drug carriers to recognize them after modifying these carriers with appropriate targeting tools such as antibodies, peptides or carbohydrates. However, it appears that there are few target molecules for tumor vasculature-targeted drug delivery, and the identification of novel target molecules would contribute to the development of DDS in cancer therapeutics.

Here, we describe subcellular proteome analysis of human umbilical vein endothelial cells (HUVECs) treated with or without VEGF to discover the target molecules expressed on the cellular membranes of angiogenic endothelial cells using 2-dimensional difference in-gel electrophoresis (2D-DIGE)

and matrix-assisted laser desorption/ionization tandem time-of-flight mass spectrometry (MALDI-TOF/TOF-MS); and then describe the functional analysis of BiP (immunoglobulin heavy chain-binding protein), which is also called glucose-regulated protein 78 kDa (GRP78) and is an identified protein in the proteomic analysis. Moreover, we developed BiP-targeted liposomes and evaluated the utility of BiP-targeted drug delivery in cancer antineovascular therapy.

Material and Methods

Cell culture

HUVECs (Cambrex Corporation, Walkersville, MD) were maintained in endothelial growth medium-2 (EGM-2, Cambrex Corporation) at 37°C under 5% CO₂ in a humidified chamber, as described previously.¹⁷ HUVECs were used at passage 7 or less in all the experiments. Mouse colon26 NL-17 carcinoma cells (C26) and human prostate carcinoma DU145 cells were cultured in an appropriate medium supplemented with streptomycin (100 μ g/mL), penicillin (100 units/mL) and 10% heat-inactivated fetal bovine serum (FBS) at 37°C in 5% CO₂.

Cellular protein fractionation

HUVECs were grown to subconfluence (80–90% confluence) in T-150 flasks, and the culture medium was replaced with 0.5% FBS-containing endothelial basal medium-2 (EBM-2, Cambrex Corporation). After serum starvation by overnight incubation, recombinant human VEGF₁₆₅ (rhVEGF, 20 ng/mL as final concentration; BD Biosciences, San Diego, CA) was added to the starved cells, which were then further incubated for 24 hr at 37°C. The cells were washed with ice-cold PBS (pH 7.4), and the proteins in the membrane/organelle were separately extracted using a Subcellular Proteome Extraction Kit (Calbiochem, San Diego, CA) according to the manufacturer's recommended protocol. The proteins in each fraction were precipitated using a Protein Precipitant Kit (Calbiochem) to concentrate the proteins and to remove interfering substances according to the manufacturer's recommended protocol. The precipitated proteins were then resolubilized in lysis buffer for 2D-DIGE (7 M urea, 2 M thiourea, 4% CHAPS, 30 mM Tris-HCl; pH 8.5), and the protein concentration was determined by the Bradford protein assay method (Bio-Rad Laboratories, Tokyo, Japan).

2D-DIGE

2D-DIGE was performed as previously described,²² and reagents used for 2D-DIGE were purchased from GE Healthcare UK (Buckinghamshire, UK). An internal standard was prepared by pooling an equal amount of each protein from VEGF-stimulated and nonstimulated HUVECs. Then, 50 μ g of protein from the internal standard, nontreated sample and VEGF-stimulated sample was fluorescence-labeled with 400 pmol of Cy2, Cy3 and Cy5, respectively, on ice for 30 min in the dark. The labeling reaction was quenched with 10 mM L-lysine. The fluorescently labeled samples were mixed and

adjusted to a final volume of 450 μ L with rehydration buffer (7 M urea, 2 M thiourea, 2% CHAPS, 1% DTT and 0.002% bromophenol blue). The samples were applied to isoelectric focusing polyacrylamide gel electrophoresis (IEF) using IPG strips (pH 3–10, nonlinear, 24 cm) and IPGphor II. The IPG strips were rehydrated with CyDye-labeled samples at 20°C for 12 hr, and IEF was performed in IPGphor at 20°C with a total of 80 kVh. The IPG strips were equilibrated for 15 min by shaking gently in an equilibration buffer (6 M urea, 30% glycerol, 2% SDS, 1.5 M Tris-HCl; pH 8.8) containing DTT (10 mg/mL) and then for 15 min in an equilibration buffer containing iodoacetamide (25 mg/mL) instead of DTT. The equilibrated IPG strips were transferred onto 10% polyacrylamide gels and 2D electrophoresis was performed using the Ettan Dalt II system. The 2D-gel images were scanned at appropriate wavelengths for Cy2, Cy3 and Cy5 dyes with a Typhoon 9400™ scanner; and the spot volumes were analyzed quantitatively using the biological variation analysis (BVA) mode of DyCyder software (version 5.0). Spots of interest were defined from average ratios of VEGF-stimulated over nonstimulated, which were above 1.5 or below -1.5-fold (0.67-fold) with significant difference (4 separate gels, Student's *t*-test, $p < 0.05$).²³

Mass spectrometry and protein identification

For mass spectrometry analysis, 500 μ g of the pooled proteins was applied to 2D electrophoresis and then stained with Deep Purple, according to the manufacturer's instructions. The gel was imaged with a Typhoon 9400 scanner. The matched spots of interest were picked with Ettan Spot Picker. In-gel digestion was performed as follows. The gel pieces picked were destained with 50 mM NH_4HCO_3 in 50% acetonitrile (ACN) and dehydrated with 100% ACN. The dried gel pieces were added to trypsin solution (25 ng/mL, Promega, Madison, WI) and incubated overnight at 37°C. The digested peptides were extracted with 1% trifluoroacetic acid (TFA) in 80% ACN and concentrated with a vacuum centrifuge. The peptide solutions were desalted and concentrated with Zip-Tip C18 μ (Millipore, Bedford, MA). Then, the peptide solutions were mixed with α -cyano-4-hydroxycinnamic acid (Sigma, Tokyo, Japan) and applied onto a target plate (Bruker Daltonics, Bremen, Germany). The MS and MS/MS spectra were obtained using an Ultraflex mass spectrometer (Bruker Daltonics) in the reflector mode. The acquired spectra were analyzed using FlexAnalysis software (version 2.2, Bruker Daltonics) in the default mode. The protein species were identified by peptide mass fingerprinting and/or combined search using Biotool software (version 2.2, Bruker Daltonics) of the Mascot search engine (version 2.0, Matrix Science, London, UK) against the NCBI database.

Western blot analysis

Western blotting was performed as described previously²⁴ with some modifications. The cells were washed with ice-cold PBS and lysed with 10 mM Tris-HCl (pH 7.4) containing

0.1% SDS, 2 mM phenylmethylsulfonyl fluoride (PMSF), 50 μ g/mL aprotinin, 200 μ M leupeptin and 100 μ M pepstatin A. To examine the phosphorylation of VEGFR-2, PLC γ and ERK1/2, we stimulated starved HUVECs with rhVEGF for 5 or 10 min. The protein concentration was determined by using the BCA protein assay kit (Pierce, Tokyo, Japan). The protein samples were mixed with SDS-PAGE sample buffer (2% SDS, 10% glycerol, 6% 2-mercaptoethanol, 50 mM Tris-HCl; pH 6.8), and an equal amount of proteins in each sample was subjected to SDS-PAGE. The separated proteins were transferred to a PVDF membrane (Millipore) and blocked with 5% skim-milk in TBST (0.9% NaCl, 0.1% Tween20, 20 mM Tris-HCl; pH 7.4). The primary antibodies used were anti-GRP78 (BiP) antibody (Santa Cruz Biotechnology, San Diego, CA), antiactin antibody (Sigma), anti-GAPDH antibody (Abcam, Cambridge, UK), antiphospho-ERK1/2 (Thr183/Tyr185), anti-ERK1/2 (Promega), antiphospho-PLC γ 1 (Tyr783), anti-PLC γ 1, antiphospho-VEGFR-2 (Tyr1175) and anti-VEGFR-2 (Cell Signaling Technology, Beverly, MA). Horseradish peroxidase (HRP)-conjugated antibodies were used as secondary antibodies. The PVDF membrane was developed with ECL reagent (GE Healthcare). The band density was measured with ImageJ software.

Cell-surface biotinylation

Extraction of cell-surface proteins was performed with a Cell Surface Biotinylation Kit (Pierce) according to the manufacturer's instructions. In brief, HUVECs were washed with ice-cold PBS, treated with NHS-SS-Biotin, and further incubated at 4°C for 30 min to conjugate the cell-surface proteins with NHS-SS-Biotin. After quenching the protein-NHS reaction, the cells were scraped and centrifuged at 500g for 3 min at 4°C. The cell pellets were then solubilized by sonication in the manufacturer's supplied lysis buffer and centrifuged at 10,000g for 3 min at 4°C. The protein samples in the supernatant were applied to a NeutrAvidin™ column and the biotin-conjugated cell-surface proteins were eluted with 50 mM DTT-containing buffer (0.1% SDS, 62.5 mM Tris-HCl; pH 6.8).

RNA interference

RNA interference (RNAi) was performed with DharmaFECT™ reagent (GE Healthcare) according to the manufacturer's instructions. HUVECs were seeded at 2.0×10^5 cells (80–90% confluence) in an antibiotic-free EGM-2 medium and incubated overnight. The cells were transfected with 100 nM of small interfering RNA (siRNA, Hokkaido System Sciences, Hokkaido, Japan) using DharmaFECT™ reagent. The sequences of siRNA for BiP (BiP siRNA) were 5'-GGU UAC CCA UGC AGU UGU UTT-3' (sense) and 5'-AAC AAC UGC AUG GGU AAC CTT-3' (antisense), as previously described.²⁵ The sequences of siRNA for green fluorescent protein (GFP siRNA used as non-silencing control siRNA) have been described previously.²⁶ After incubation with these siRNAs for 48 hr at 37°C, the effects of each siRNA on

protein expression were determined by Western blot analysis and real-time PCR.

RNA isolation and real-time PCR

Total RNA was isolated using ISOGEN (Nippon Gene, Tokyo, Japan) and the resulting RNA was reverse transcribed. TaqMan real-time PCR assay was performed using TaqMan gene expression assays on a StepOnePlus (Applied Biosystems, Tokyo, Japan). The relative real-time PCR quantification was based on a comparative quantification method. The endogenous reference gene was GAPDH. The specificity of the PCR reactions was confirmed by a single band of the predicted size after agarose gel electrophoresis (data not shown).

Cell proliferation assay

HUVECs were transfected with each siRNA as described above and incubated for 24 hr at 37°C. The transfected cells were harvested and reseeded at 2.0×10^4 cells per well on a 24-well plate in 0.5% FBS-containing EBM-2 medium with or without rhVEGF (20 ng/mL). The cells were further incubated for 24 or 48 hr, and the cell viability was determined at each time with TetraColor One™ (Seikagaku, Tokyo, Japan). The growth ratio was calculated by the absorbance (450 nm) at 24 hr/3 hr and 48 hr/3 hr. The absorbance at 3 hr was considered to reflect the cell number immediately after seeding the cells without cell proliferation.

Preparation of liposomes

Distearyl phosphatidylcholine (DSPC), distearyl phosphatidylglycerol (DSPG), cholesterol and distearyl phosphatidylethanolamine-conjugated polyethyleneglycol 2000 (DSPE-PEG) or the DSPE-PEG conjugate of GWIFWQIQL-peptides, which are shown to bind to BiP/GRP78,²⁷ were dissolved in tert-butylalcohol (10:10:10:1 as a molar ratio). The lipid solutions were lyophilized, and the lyophilizates were hydrated with 0.3 M sucrose solution (pH 7.4). The liposome solutions were frozen and thawed for 3 cycles with liquid nitrogen. Then, the liposome size was adjusted by extruding through polycarbonate filters of 100-nm pore size (Nuclepore, Cambridge, MA). For the preparation of doxorubicin (DOX)-containing liposomes, DOX solutions were added to the initial lipid solutions in the proportion of 30 mol % to the composed lipids (DSPC + DSPG). Unencapsulated DOX was removed by ultracentrifugation at 604,000g for 15 min (HITACHI, Tokyo, Japan), and the amount of DOX in the liposome was determined by measuring the absorbance at 484 nm. The liposome size was measured using ZETASIZER (Malvern Instruments, Malvern, UK).

Cellular uptake of liposomes

The liposomes were radiolabeled by adding [³H]cholesteryl hexadecyl ether solution (370 kBq/mL, GE Healthcare) to the initial lipid solution. HUVECs were seeded (2.0×10^5 cells) on a 35-mm dish and stimulated with rhVEGF as described above. C26 or DU145 cells (both 1.0×10^5 cells) were also

seeded on a 35-mm dish and incubated overnight. The radiolabeled liposomes were added to the cells and further incubated for 2 hr. The cells were washed with PBS and solubilized with 10 mM Tris-HCl buffer containing 0.1% SDS (pH 7.4). The cell lysates were recovered and transferred to HionicFluor (Perkin Elmer, Foster City, CA). Radioactivity was determined with a liquid scintillation counter (LSC-3500; Aloka, Tokyo, Japan).

Intratumoral distribution of WIFWQIQL liposomes

The liposomes were fluorescently labeled by adding the Dil C18 (Molecular Probes, Eugene, OR) solution to the initial lipid solution. C26 cells were subcutaneously implanted into the posterior flank of 5-week-old BALB/c male mice. The Dil C18-labeled liposomes were intravenously injected into the mice 10 days after tumor implantation. At 3 or 6 hr after injection of the liposomes, the mice were sacrificed using diethyl ether anesthesia, and the tumors were dissected. Immunostaining was performed as described previously. The tumor tissues were embedded and frozen with dry ice/ethanol. Tumor sections (10 μm) were prepared with cryostatic microtome (HM 505E; Microm, Walldorf, Germany) and air-dried for at least 1 hr. The sections were incubated in 1% bovine serum albumin containing PBS for 10 min at room temperature for protein blocking, with biotinylated anti-mouse CD31 rat monoclonal antibody (BD Pharmingen, Franklin Lakes, NJ) for 18 hr at 4°C, and then with streptavidin-Alexa Fluor 488 conjugates (Molecular Probes) for 30 min at room temperature. Finally, the sections were mounted with Perma Fluor Aqueous Mounting Medium (Thermo Shandon, Pittsburgh, PA) and fluorescently observed with a microscopic LSM system (Carl Zeiss, Germany).

Dorsal air sac model

Dorsal air sac model mice were prepared as follows: C26 cells (1.0×10^7 cells/150 μL) were loaded into a chamber ring (Millipore) covered with filters (0.45-nm pore size, Millipore). The chamber ring was then implanted subcutaneously into the dorsal skin of the mice. At days 2 and 3, 0.3 M sucrose solution (control), PEG-liposomal doxorubicin (PEG-Lip-DOX) or WIFWQIQL-liposomal doxorubicin (WIFWQIQL-Lip-DOX) was intravenously administered (5 mg/kg/day as dose of DOX). At day 4, the mice were sacrificed using diethyl ether and the dorsal skin oscillating the chamber ring was observed.

Therapeutic experiment

PEG-Lip-DOX, WIFWQIQL-Lip-DOX (5 mg/kg/day as dose of DOX) or 0.3 M sucrose solution (control) was intravenously administered to the C26-bearing mice at days 6, 9, 12 and 15 after tumor implantation. The tumor size and body weight of each mouse were monitored. The animals used in the experiments were cared for according to the guidelines for the care and use of laboratory animals of the University of Shizuoka.