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Radioiodinated VEGF to image tumor angiogenesis in a LS180 tumor xenograft model

Mitsuyoshi Yoshimoto^{a,*}, Seigo Kinuya^b, Atsuhiro Kawashima^c, Ryuichi Nishii^{d,e}, Kunihiko Yokoyama^b, Keiichi Kawai^{a,e}

^aDivision of Health Sciences, Graduate School of Medical Science, Kanazawa University, Ishikawa 920-0942, Japan

^bDepartment of Biotracer Medicine, Graduate School of Medical Science, Kanazawa University, Ishikawa 920-8640, Japan

^cKanazawa Medical Center, Ishikawa 920-8650, Japan

^dDepartment of Radiology, Fujimoto Hayasuzu Hospital, Miyazaki 885-0055, Japan

^eBiomedical Imaging Research Center, University of Fukui, Fukui 910-1193, Japan

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Abstract

Introduction: Angiogenesis is essential for tumor growth or metastasis. A method involving noninvasive detection of angiogenic activity in vivo would provide diagnostic information regarding antiangiogenic therapy targeting vascular endothelial cells as well as important insight into the role of vascular endothelial growth factor (VEGF) and its receptor (flt-1 and KDR) system in tumor biology. We evaluated radioiodinated VEGF₁₂₁, which displays high binding affinity for KDR, and VEGF₁₆₅, which possesses high binding affinity for flt-1 and low affinity for KDR, as angiogenesis imaging agents using the LS180 tumor xenograft model.

Methods: $VEGF_{121}$ and $VEGF_{165}$ were labeled with ¹²⁵I by the chloramine-T method. Biodistribution was observed in an LS180 human colon cancer xenograft model. Additionally, autoradiographic imaging and immunohistochemical staining of tumors were performed with ¹²⁵I-VEGF₁₂₁.

Results: ¹²⁵I-VEGF₁₂₁ and ¹²⁵I-VEGF₁₆₅ exhibited strong, continuous uptake by tumors and the uterus, an organ characterized by angiogenesis. ¹²⁵I-VEGF₁₂₁ uptake in tumors was twofold higher than that of ¹²⁵I-VEGF₁₆₅ (9.12±98 and 4.79±1.08 %ID/g at 2 h, respectively). ¹²⁵I-VEGF₁₂₁ displayed higher tumor to nontumor (T/N) ratios in most normal organs in comparison with ¹²⁵I-VEGF₁₆₅. ¹²⁵I-VEGF₁₂₁ accumulation in tumors decreased with increasing tumor volume. Autoradiographic and immunohistochemical analyses confirmed that the difference in ¹²⁵I-VEGF₁₂₁ tumor accumulation correlated with degree of tumor vascularity.

Conclusion: Radioiodinated VEGF₁₂₁ is a promising tracer for noninvasive delineation of angiogenesis in vivo.

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Keywords: Tumor imaging; Angiogenesis; VEGF; VEGF receptors

1. Introduction

Tumors require angiogenesis in order to grow beyond the size of 1–2 mm in diameter, a condition in which nutrients and oxygen are not adequately supplied via passive diffusion through existing blood vessels [1]. Furthermore, the metastatic property of tumors strongly correlates with tumor angiogenesis [2,3]. In particular, the balance between various activators and inhibitors closely associated with overexpression of vascular endothelial growth factor (VEGF) and its receptors serves to regulate angiogenesis.

It is well known that VEGF plays a pivotal role in angiogenesis — proliferation and migration of endothelial cells. Five VEGF isoforms, produced by alternative splicing of responsible genes, have been identified. VEGF₁₂₁ and VEGF₁₆₅ are the predominant isoforms [4]. Endothelial cells express two VEGF receptors (VEGFRs): flt-1 (K_d =26±6 pM for VEGF₁₆₅) and KDR (K_d =80±13 pM for VEGF₁₂₁, K_d =2070±520 pM for VEGF₁₆₅) [5]. These receptors are up-regulated in endothelial cells in tumor tissue [6,7].

Numerous compounds possessing an antiangiogenic profile have been investigated [8–10]. Conventional antitumor agents act directly on tumor cells. In contrast, most antiangiogenic agents primarily target endothelial cells,

^{*} Corresponding author. Tel.: +81 76 265 2539; fax: +81 76 234 4366. *E-mail address:* myoshi@mhs.mp.kanazawa-u.ac.jp (M. Yoshimoto).

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which enables their application to various types of tumors. Most antiangiogenic agents target the VEGF–VEGFR system, which inhibits proliferation of endothelial cells [11–14].

Despite its invasive nature, conventional angiography has been the standard imaging tool for evaluation of tumor vascularity. Less invasive approaches such as computed tomographic angiography and, more recently, three-dimensional magnetic resonance angiography can effectively demonstrate the vasculature. These modalities, however, merely delineate preexisting vasculature; they do not disclose the dynamic changes in blood vessels (endothelial cells) attributable to antiangiogenic therapy. Positron emission tomography and single photon emission computed tomography may enable lesional characterization in the management of cancer patients. Positron tracers such as 2-18F-fluoro-2-deoxyglucose (18F-FDG) and 3'-deoxy-3' -18 F-fluorothymidine (18 F-FLT) would afford data corresponding to tumor therapeutic responses. The major action of antiangiogenic therapy is the induction of dormancy of tumors, which may alter glycolysis or DNA synthesis in tumors. However, alternations in these cellular functions in antiangiogenic therapy occur secondarily, following the primary effects of antiangiogenic agents on endothelial cells. So-called perfusion agents, e.g., ²⁰¹Tl, 99mTc-sestamibi and 99mTc-tetrofosmin, would also be inadequate, as factors other than tissue perfusion substantially affect the distribution of these tracers. For instance, ^{99m}Tc-sestamibi accumulation in brain tumors is independent of tumor vascularization [15]. Neo-angiogenesis in tumors is strongly correlated with patient prognosis; consequently, a method enabling noninvasive estimation of angiogenesis in vivo would surely provide beneficial clinical information. Furthermore, such information may permit selection of those patients most likely to benefit from antiangiogenic therapy as well as facilitate monitoring of its efficacy.

In order to develop noninvasive, specific techniques for estimation of tumor vascularity, this investigation utilized VEGF₁₂₁ and VEGF₁₆₅ radiolabeled with ¹²⁵I. The specificity and binding ability of VEGF₁₂₁ and VEGF₁₆₅ to receptors have been well established with ¹²⁵I-VEGF₁₂₁ and ¹²⁵I-VEGF₁₆₅ in biochemical field. These compounds were evaluated as candidates for angiogenesis imaging using a human cancer xenograft model.

2. Materials and methods

2.1. Radioiodination of VEGF

Radioiodination of human recombinant VEGF₁₂₁ (R&D Systems, Minneapolis, MN) and VEGF₁₆₅ (PeproTech EC, London, UK) was performed via the chloramine-T method described previously [5,16]. Briefly, 1.0 μ g of VEGF₁₂₁ or VEGF₁₆₅ was labeled with approximately 18.5 MBq of Na¹²⁵I (Amersham Pharmacia, Buckinghamshire, UK) and

25 µg of chloramine-T in 20 mM phosphate buffer (pH 7.2) in a final volume of 110 µl. Iodination was allowed to proceed at room temperature for 1 min. The reaction was terminated upon the addition of 5 µl of sodium metabisulfite (5 mg/ml). ¹²⁵I-VEGF₁₂₁ and ¹²⁵I-VEGF₁₆₅ were purified on a gel filtration (Sephadex G-25) column (Amersham Biosciences, Uppsala, Sweden) and on a Heparin-Sepharose column (Amersham Biosciences), respectively. ¹²⁵I-VEGF₁₂₁ and ¹²⁵I-VEGF₁₆₅ displayed specific activities of 13.0–18.5 and 7.2–8.0 GBq/mg, respectively.

2.2. Cell culture

LS180 human colorectal adenocarcinoma cell line was purchased from Dai-Nippon Seiyaku Co. Ltd. (Japan) and grown in Eagle's minimal essential medium (Sigma, St. Louis, MO) containing 2 mM L-glutamine, nonessential amino acids and 10% FBS. Cells were cultured in a 5% CO_2 -humidified atmosphere at 37°C.

2.3. Serum stability

Mouse serum (50 µl) was mixed with 18.5 kBq of 125 I-VEGF₁₂₁ and 125 I-VEGF₁₆₅, respectively, and was incubated at 37°C. After incubation for 0, 1, 2 and 4 h, trichloroacetic acid (TCA) was added to the samples. Precipitates were filtrated on glass filters, and the filters were washed with 5% TCA. Radioactivity in precipitates at 0 h was used as control. Percent precipitability of control was calculated.

2.4. Biodistribution of ¹²⁵I-VEGF

Animal studies were performed in compliance with guidelines for the care and use of laboratory animals of Kanazawa University. Biodistribution studies were conducted in nude mice bearing LS180 tumor. Balb/c nu/nu female mice, 5-6 weeks of age (Japan SLC, Inc., Shizuoka, Japan), were xenografted subcutaneously in the left thigh with 5×10^6 LS180 cells. LS180 displays considerable VEGF production and no VEGF receptors; moreover, this growth is suppressed by an anti-VEGF antibody in vivo [17]. Biodistribution was examined at 7, 10 and 14 days after implantation in order to include various tumor sizes. Mice were injected via the tail vein with 55.5 kBq/100 μ l (saline containing 1% bovine serum albumin) ¹²⁵I-VEGF₁₂₁ or ¹²⁵I-VEGF₁₆₅. Mice were sacrificed at 0.5, 1, 2, 4 and 24 h postinjection, and tissue samples were excised. Tissue samples were weighed and radioactivity was measured with a γ-counter (ARC-1000M, Aloka, Tokyo, Japan). Uptake in

Table 1 Serum stability of ¹²⁵I-VEGF₁₂₁ and ¹²⁵I-VEGF₁₆₅ evaluated by TCA precipitability measurements

	Incubation time			
	1 h	2 h	4 h	
¹²⁵ I-VEGF ₁₂₁	93.8±2.0	92.5±1.2	94.9±2.6	
¹²⁵ I-VEGF ₁₆₅	97.3 ± 2.2	96.8 ± 0.9	97.5±0.7	

The stability was expressed as the percent precipitant of control (n=2-3).

Table 2	
Biodistribution of ¹²⁵ I-VEGF ₁₂₁	in LS180 tumor-bearing mice $(n=4)$

Organ	Time after injection					
	0.5 h	1 h	2 h	4 h	24 h	
Tumor	5.94 ± 0.41	8.43 ± 0.58	9.12 ± 0.98	9.02 ± 1.59	2.55 ± 0.28	
Blood	40.32 ± 2.45	38.69 ± 4.34	32.03 ± 0.59	24.90 ± 2.99	7.19 ± 0.57	
Brain	0.88 ± 0.13	0.89 ± 0.17	0.70 ± 0.08	0.44 ± 0.03	$0.16 {\pm} 0.02$	
Liver	7.16 ± 0.62	6.86 ± 1.15	5.49 ± 0.44	5.21 ± 0.57	1.44 ± 0.11	
Uterus	6.62 ± 2.39	11.24 ± 3.70	13.76 ± 2.19	15.58 ± 4.13	3.72 ± 0.88	
Heart	9.36 ± 1.59	9.43 ± 1.04	8.39 ± 0.41	6.48 ± 1.05	2.09 ± 0.18	
Lung	18.88 ± 3.11	18.54 ± 2.26	13.31 ± 2.05	14.11 ± 1.62	$3.56 {\pm} 0.37$	
Stomach	4.49 ± 0.39	7.32 ± 0.94	8.81 ± 1.87	10.85 ± 2.18	4.23 ± 1.22	
Kidney	14.91 ± 0.60	14.16 ± 0.78	11.46 ± 0.78	9.21±1.04	3.00 ± 0.14	
Pancreas	1.97 ± 0.20	3.27 ± 0.53	2.30 ± 0.73	2.91 ± 0.15	1.17 ± 0.21	
Spleen	6.77 ± 0.89	6.46 ± 0.34	5.90 ± 1.36	4.53 ± 0.79	1.67 ± 0.11	
Muscle	1.27 ± 0.33	1.33 ± 0.06	2.48 ± 0.86	1.74 ± 0.44	0.90 ± 0.10	
Bone	3.32 ± 0.66	4.19±1.22	4.90 ± 1.05	2.76 ± 0.18	1.12 ± 0.28	
Intestine	3.40 ± 0.69	3.93 ± 0.74	3.84 ± 0.36	3.10 ± 0.19	1.08 ± 0.13	

Radioactivity in organs was expressed as %ID/g tissue (mean±S.D.). Tumor volume used in this study was less than 200 mm³.

organs was expressed as percent injected dose per gram (%ID/g) of tissue.

2.5. Autoradiography of ¹²⁵I-VEGF₁₂₁ in tumor

¹²⁵I-VEGF₁₂₁ displayed higher tumor to nontumor (T/N) ratio in comparison to ¹²⁵I-VEGF₁₆₅; as a result, intratumoral distribution of ¹²⁵I-VEGF₁₂₁ was analyzed autoradiographically (ARG). Mice were injected via the tail vein with 740–999 kBq/100 µl. Tumors were excised from mice at 4 h postinjection. Specimens were embedded in OCT compound (Sakura Finetechnical Co. Ltd., Tokyo, Japan) and frozen in dry ice–acetone. Frozen sections were cut to 20-µm thickness with a cryostat and mounted on glass slides. Glass slides were placed on an imaging plate (BAS-SR 2025, Fuji Photo Film Co. Ltd., Tokyo, Japan); subsequently, the exposed plate was scanned with a bio-imaging analyzer (BAS-5000, Fuji Photo Film) to detect radioactivity.

Table 5	
Biodistribution of ¹²⁵ I-VEGF ₁₆₅ in LS180	tumor-bearing mice $(n=3-5)$

Table 2

2.6. Immunohistochemistry by CD34 in tumors

Frozen sections adjacent to the ARG specimens were sliced to 5-µm thickness and mounted on silanized slides. The sections were air-dried and fixed in cold acetone for 5 min. Subsequently, sections were rinsed in PBS to remove the OCT compound. Endogenous peroxidase activity was blocked with 0.3% H₂O₂, and nonspecific binding was blocked with 10% normal rabbit serum. After washing in PBS, rat antimouse CD34 monoclonal antibody (BD pharmingen, La Jolla, CA), diluted 1:20, was applied to the sections and incubated at room temperature (RT) for 30 min. Next, after a PBS wash, biotinylated rabbit antirat IgG antibody (Vector Laboratory, Burlingame, CA), diluted 1:200, was applied to the sections and incubated at RT for 30 min. After washing again in PBS, streptavidine (Dako-Cytomation, Glostrup, Denmark), diluted 1:300, was applied to the sections, incubated at RT for 30 min, and washed again in PBS. The sections were then developed with diamino-

Organ	Time after injection				
	0.5 h	1 h	2 h	4 h	24 h
Tumor	2.06 ± 0.40	3.09 ± 0.72	4.79 ± 1.08	1.84 ± 0.08	$0.17 {\pm} 0.03$
Blood	1.88 ± 0.25	3.33 ± 0.46	5.15 ± 0.55	2.48 ± 0.45	0.61 ± 0.52
Brain	3.17 ± 0.22	2.53 ± 0.14	1.88 ± 0.51	0.76 ± 0.06	0.31 ± 0.01
Liver	16.15 ± 2.05	7.71 ± 0.23	5.72 ± 0.95	4.26 ± 0.05	2.58 ± 0.31
Uterus	6.24 ± 0.47	4.93 ± 1.02	5.24 ± 0.95	2.56 ± 0.40	0.63 ± 0.30
Heart	14.36 ± 0.44	9.93 ± 1.18	6.59 ± 0.95	2.96 ± 0.48	1.31 ± 0.11
Lung	31.30 ± 7.56	20.32 ± 2.95	15.71 ± 1.53	7.84 ± 1.68	4.12 ± 0.37
Stomach	12.65 ± 1.34	17.96 ± 6.90	32.79 ± 10.06	23.93 ± 8.71	1.11 ± 0.19
Kidney	25.19 ± 1.37	18.64 ± 1.35	12.45 ± 1.71	8.36 ± 0.46	6.67 ± 1.38
Pancreas	10.07 ± 1.30	9.43 ± 1.16	6.82 ± 0.68	3.68 ± 0.56	1.00 ± 0.13
Spleen	17.80 ± 2.39	9.00 ± 1.62	6.78 ± 1.75	4.14 ± 0.46	2.53 ± 0.54
Muscle	4.68 ± 2.78	3.71 ± 1.31	3.91 ± 0.76	1.66 ± 0.08	0.50 ± 0.04
Bone	2.55 ± 0.10	2.57 ± 0.35	2.28 ± 0.17	1.36 ± 0.29	0.25 ± 0.05
Intestine	4.13 ± 0.16	3.87 ± 0.74	4.15 ± 0.36	2.53 ± 0.51	0.43 ± 0.05

Radioactivity in organs was expressed as %ID/g tissue (mean±S.D.).

Tumor volume used in this study was less than 220 mm³.

benzidine (DakoCytomation) and counterstained with hematoxylin (Merck, Darmstadt, Germany). Additional slices were also examined using hematoxylin–eosin staining.

3. Results

3.1. Serum stability

Stability study showed no decrease in precipitability of $^{125}\text{I-VEGF}_{121}$ and $^{125}\text{I-VEGF}_{165}$ in mouse serum (Table 1). In serum, no obvious deiodination of $^{125}\text{I-VEGF}_{121}$ and $^{125}\text{I-VEGF}_{165}$ was found.

3.2. Biodistribution and tumor accumulation of ¹²⁵I-VEGF

Tables 2 and 3 summarize the results of biodistribution studies of $^{125}\text{I-VEGF}_{121}$ and $^{125}\text{I-VEGF}_{165}$ in LS180-bearing mice, respectively. Accumulation of $^{125}\text{I-VEGF}_{121}$ in tumors gradually increased with time, reaching a peak (9.12 %ID/g) at 2 h postinjection; subsequently, $^{125}\text{I-VEGF}_{121}$ accumulation remained unchanged through 4 h postinjection. The uterus demonstrated a $^{125}\text{I-VEGF}_{121}$ accumulation pattern similar to that of tumors.

¹²⁵I-VEGF₁₆₅ exhibited higher accumulation in normal tissues, with the exception of blood, than ¹²⁵I-VEGF₁₂₁;



Fig. 1. Tumor to nontumor ratios in LS180 tumor-bearing mice at 0.5, 1, 2, 4 and 24 h after injection of ¹²⁵I-VEGF₁₂₁ (A) and ¹²⁵I-VEGF₁₆₅ (B). Columns, mean; bars, SD (n=3-5).



Fig. 2. Relationship between ¹²⁵I-VEGF₁₂₁ accumulation and tumor volume. Data correspond to ¹²⁵I-VEGF₁₂₁ accumulation at 4 h postinjection. LS180-xenografted mice were examined at 7, 10 and 17 days after implantation. Tumor volume was calculated as $\pi/6 \times (r_1 \times r_2 \times r_3)$; (r_1 : length; r_2 : width; r_3 : depth).

however, ¹²⁵I-VEGF₁₆₅ was cleared more rapidly than ¹²⁵I-VEGF₁₂₁, particularly in liver, heart, lung, kidney and spleen. ¹²⁵I-VEGF₁₂₁ accumulation in tumors was two- and fivefold greater than that of ¹²⁵I-VEGF₁₆₅ at 2 and 4 h postinjection, respectively.



Fig. 3. Autoradiography image of intratumoral distribution at 4 h postinjection of 125 I-VEGF₁₂₁. Images were obtained from LS180 tumor tissues displaying volumes of (A) 101.9 mm³ and (B) 361.9 mm₃. Tissue slice photos (C) and (D) correspond to ARG images (A) and (B), respectively (T: tumor; M: muscle).



Fig. 4. Vascular staining with an anti-CD34 mAb in LS180 tumor. Images (A) and (B) are representative areas of Fig. 3A and B, respectively. Bar indicates 100 im in (A).

Fig. 1A and B displays T/N ratios of ¹²⁵I-VEGF₁₂₁ and ¹²⁵I-VEGF₁₆₅, respectively. Tumor-to-nontumor ratios of ¹²⁵I-VEGF₁₂₁ were markedly higher than those of ¹²⁵I-VEGF₁₆₅, with the exception of blood. Tumor-to-blood ratio of ¹²⁵I-VEGF₁₂₁ increased to 4 h postinjection, although the ratio was strikingly low. In contrast, the tumor-to-blood ratio of ¹²⁵I-VEGF₁₆₅ was maximal at 30 min; subsequently, it decreased with time.

3.3. Relationship between tumor volume and ¹²⁵I-VEGF121 accumulation in tumors

Fig. 2 illustrates the relationship between tumor volume (mm^3) and accumulation in tumors. ¹²⁵I-VEGF₁₂₁ exhibited a tendency toward diminishing accumulation in tumor with increasing tumor volume.

3.4. Autoradiography and histopathological findings

Fig. 3 displays intratumoral distribution of ¹²⁵I-VEGF₁₂₁. ¹²⁵I-VEGF₁₂₁ demonstrated overall high localization in small tumors (101.9 mm³) (Fig. 3A). As shown in Fig. 3B, high localization of ¹²⁵I-VEGF₁₂₁ was evident in portions of the muscle area in large tumors (361.9 mm³). Fibrosis was induced at regions of high ¹²⁵I-VEGF₁₂₁ localization in tumoral area. Homogeneous accumulation of ¹²⁵I-VEGF₁₂₁ was observed in the tumor area, with the exception of areas of fibrosis.

Fig. 4A and B indicates microvessel staining of tumor tissues by CD34 in Fig. 3A and B, respectively. As shown in Fig. 4A, abundant vasculature was observed in small tumors. On the other hand, poor vasculature was apparent in large tumors (Fig. 4B).

4. Discussion

Neo-angiogenesis is a common characteristic of all proliferating and metastatic tumors. Development of novel drugs possessing anti-angiogenic profile is a major area of interest in clinical oncology. In vivo detection of angiogenesis clearly affords valuable information in terms of monitoring the efficacy of anti-angiogenic therapy. The VEGF-VEGFR system would be the most suitable imaging target for this purpose as this system plays a key role in angiogenesis in tumors. Positron-labeled monoclonal antibody (VG76e) that binds human VEGF has been developed in order to measure VEGF expression in tumor tissues [18]. However, substances for imaging VEGFR density would be required to monitor therapeutic responses to anti-angiogenic agents given that most agents act as inhibitors of VEGFR. Our findings provide a rationale for VEGFR imaging in tumors utilizing radiolabeled VEGF.

¹²⁵I-VEGF₁₂₁ and ¹²⁵I-VEGF₁₆₅ displayed similarities and differences in terms of in vivo behavior in tumorxenografted mice. High initial uptake was observed in the lung and heart, probably due to abundant vascularity in these organs. Based on the molecular sizes, ¹²⁵I-VEGF₁₂₁ and ¹²⁵I-VEGF₁₆₅ were filtered by the kidney, resulting in early large accumulation in that tissue. In addition, radioiodinated metabolites may also contribute to the accumulation in kidney. In contrast, prominent 125 I-VEGF₁₆₅ accumulation in the stomach in comparison to 125 I-VEGF₁₂₁ suggests that the former was more prone to deiodination than the latter. Furthermore, differences in tissue distribution between these two compounds were observed in various other organs, including the brain and spleen, which was likely attributable to the difference corresponding to binding sites recognized by each tracer: VEGF₁₂₁ binds only to KDR with high affinity [5,19], whereas VEGF₁₆₅ binds to flt-1 with higher affinity but possesses lower affinity for KDR [5,20]. KDR mRNA is primarily expressed in the kidney, heart, spleen and lung [21], whereas flt-1 mRNA is expressed in the brain, heart, liver, kidney and lung [22]. In kidney, heart and lung, which express both VEGFRs, ¹²⁵I-VEGF₁₆₅ uptake was greater than that of ¹²⁵I-VEGF₁₂₁. Although the precise reasons regarding the different accumulation in these organs remain unclear, previous reports have suggested a possible mechanism for this

phenomenon: the difference in the expression between flt-1 and KDR on endothelial cells in adult tissues. That is, endothelial cells in adult tissues express flt-1 mRNA in a manner similar to those in embryonic tissues [23], whereas KDR is down-regulated on endothelial cells in adult tissues relative to corresponding cells in fetal tissues [24]. These findings indicate that VEGF₁₂₁ would be the more appropriate targeting molecule of VEGFRs in terms of low background activities.

The superiority of ¹²⁵I-VEGF₁₂₁ was also suggested by its greater tumor accumulation as compared to ¹²⁵I-VEGF₁₆₅. We cannot exclude the possibility that the radioactivity in blood contributes to the radioactivity in tumor especially at early time points. However, the fact that tumor-to-blood ratios gradually increased through 4 h affords evidence of specific binding to KDR. Similarity of uptake profile of ¹²⁵I-VEGF₁₂₁ in tumor to that of the uterus where angiogenesis occurs physiologically also indicated its specific uptake by the receptors. Therefore, the difference in expression amount and/or binding affinity between flt-1 and KDR is an essential factor with respect to determination of targeting level in tumors between ¹²⁵I-VEGF₁₂₁ and ¹²⁵I-VEGF₁₆₅. In addition, the up-regulation of VEGFRs may affect the binding affinity of ¹²⁵I-VEGF₁₂₁ to flt-1 [25], resulting in high accumulation of ¹²⁵I-VEGF₁₂₁ in tumors.

In the clinical setting, ¹²³I-VEGF₁₆₅ scintigraphy has successfully delineated gastrointestinal tumors and their metastases in lung and liver despite the high background activity in these organs [26]. Our present results suggest that ¹²³I-VEGF₁₂₁ would lead to improved detection ability in terms of abdominal lesions relative to ¹²³I-VEGF₁₆₅ consequent to higher T/N ratios expected with ¹²⁵I-VEGF₁₂₁.

¹²⁵I-VEGF₁₂₁ uptake is greater in tumors characterized by abundant vascularity than in tumors displaying poor vascularity. Greater expression of KDR in metastatic tumors with larger vessel counts in comparison to nonmetastatic tumors with smaller vessel counts has been documented [27]. Although we could not accurately measure microvessel density due to the well-developed nature of the sinusoidal vascular network in this experimental tumor model, this previous report supports our findings.

Microvessel density is correlated with invasion and metastasis, which suggests that this parameter may serve as a powerful prognostic marker [2,3]. In addition, antiangiogenic therapy leads to diminished microvessel density [11,28]. Therefore, radioiodinated VEGF may be useful for tumor diagnosis and monitoring of anti-angiogenic therapy.

Angiostatin and endostatin are endogenous angiogenesis inhibitors and are specific inhibitors of endothelial cell proliferation. The potential of ^{123/125}I-angiostatin and ^{99m}Tc-EC-endostatin as markers for evaluating angiogenic activity has been demonstrated [29,30]. tumor-to-nontumor ratios for liver and muscle, which are comparable, are even higher for ¹²⁵I-VEGF₁₂₁ in comparison with these compounds: the ratio of tumor to liver and muscle is 1.74 and 5.49 for ¹²⁵I- VEGF₁₂₁, 0.69 and 3.54 for ^{123/125}I-angiostatin and 0.16 and 6.1 for ^{99m}Tc-EC-endostatin, respectively. Moreover, ¹²⁵I-VEGF₁₂₁ produced a higher tumor-to-intestine ratio of 2.92 relative to that of ^{123/125}I-angiostatin and ^{99m}Tc-ECendostatin. These findings are suggestive of the feasibility of radioiodinated VEGF₁₂₁ as a targeting agent.

A possible critical problem associated with radioiodinated VEGF₁₂₁ involves prolonged elevated radioactivity in the circulation. A plausible explanation for this phenomenon corresponds to the binding of VEGF₁₂₁ to endothelial progenitor cells and/or KDR-positive cells in the circulation. Endothelial progenitor cells that express KDR originate in the bone marrow and circulate to tissues where these cells contribute to angiogenesis [31,32]. This obstacle may be overcome by co-injection of unlabeled VEGF₁₂₁, which would suppress binding of radioiodinated VEGF₁₂₁ to KDR, leading to increased availability of free radioiodinated VEGF₁₂₁; thus, accumulation of radioiodinated VEGF₁₂₁ in tumors would be enhanced in a manner similar to that observed with ¹²⁵I-insuline-like growth factor (IGF)-I binding to IGF-binding proteins [33].

5. Conclusion

Diagnosis of angiogenic activity in tumors provides beneficial information concerning the efficacy of antiangiogenic therapy and risk of distant metastasis. In this report, we confirmed that radioiodinated VEGF₁₂₁ is a promising tracer for this purpose. In terms of VEGFRs, KDR is a superior target in comparison to flt-1.

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