A Physiologic Imaging Pilot Study of Breast Cancer Treated with AZD2171

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Abstract  Background: This pilot study combined physiologic imaging, microcomputed tomography, and histologic tumor evaluation with a xenograft model of breast cancer to identify surrogates likely to correlate with response to AZD2171, an inhibitor of the vascular endothelial growth factor (VEGF) receptor tyrosine kinases.

Experimental Design: MCF-7 cells transfected with vector (MCF-7neo) or VEGF (MCFVEGF) were implanted in the right and left mammary fat pads of 75 athymic mice. Treatment with AZD2171 (5 mg/kg/d) or vehicle control was initiated once tumors were established. Positron emission tomography with [11C]carbon monoxide to measure blood volume, [18F]fluoromethane to measure perfusion, and [18F]fluorodeoxyglucose to measure glucose utilization was done at baseline, and after 24 hours, 72 hours, and 4 weeks of treatment. After imaging, tumors were analyzed for microvessel density, proliferation, and VEGF expression.

Results: AZD2171 induced significant inhibition of tumor growth in established MCF-7neo xenografts and regression of established MCF-7VEGF xenografts. An acute decrease in blood flow was detected in MCF-7VEGF tumors at 24 hours (P = 0.05). Tumor blood volume was increased in the MCF-7VEGF tumors but correlated with tumor size; blood volume did not change with AZD2171 therapy. Glucose utilization correlated with tumor size and did not change with acute or chronic AZD2171 therapy. Unlike blood flow and blood volume, glucose utilization was similar in MCF-7neo and MCF-7VEGF tumors. Microvessel density and proliferation acutely decreased in MCF-7VEGF tumors but returned to baseline during chronic therapy.

Conclusions: [18F]Fluoromethane imaging may be a useful surrogate for biological activity of AZD2171 with changes identified within 24 hours of starting therapy.

Over the last two decades, substantial laboratory and indirect clinical evidence has accumulated to support the central role of angiogenesis in breast cancer progression (1, 2). Multiple angiogenic factors are commonly expressed by invasive human breast cancers with the 121-amino-acid isoform of vascular endothelial growth factor (VEGF) predominating (3). VEGF is a highly conserved, homodimeric, secreted, heparin-binding endothelial growth factor (VEGF) predominating (3). VEGF is breast cancers with the 121-amino-acid isoform of vascular angiogenic factors are commonly expressed by invasive human clinical evidence has accumulated to support the central role

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Over the last two decades, substantial laboratory and indirect clinical evidence has accumulated to support the central role of angiogenesis in breast cancer progression (1, 2). Multiple angiogenic factors are commonly expressed by invasive human breast cancers with the 121-amino-acid isoform of vascular endothelial growth factor (VEGF) predominating (3). VEGF is a highly conserved, homodimeric, secreted, heparin-binding glycoprotein whose dominant isoform has a molecular weight of ~45,000 (4). The biological effects of VEGF are mediated through binding to one of three endothelial surface receptors VEGF receptor-1 (VEGFR-1, Flt-1), VEGFR-2 (Flk-1/KDR), and VEGFR-3 (Flt-4); binding to the coreceptor neuropilin enhances signaling (5, 6). The VEGFRs share considerable overlap in ligand binding, downstream effector interaction, and biological function; however, predominant actions have been identified for each. VEGFR-1 promotes differentiation and vascular maintenance (7); VEGFR-2 induces endothelial cell mitogenesis, survival, and migration and increases vascular permeability (8); VEGFR-3 stimulates lymphangiogenesis (9, 10).

AZD2171, a novel indole-ether quinazoline, is a highly potent (IC50 < 1 nmol/L) inhibitor of the VEGF-2 tyrosine kinase. AZD2171 inhibits VEGF-stimulated endothelial cell proliferation and in vitro vessel growth and sprouting. In vivo AZD2171 inhibits experimental angiogenesis, endochondral ossification, and growth of established human tumor xenografts in a dose-dependent fashion (11). Early clinical trials with AZD2171 in patients with advanced cancer are ongoing (12, 13).

Correlative laboratory studies assessing biologically meaningful intermediate end points are a necessity for the successful development of antiangiogenic therapy. Although, as yet, no clear standard has emerged, the search for reliable surrogates of antiangiogenic activity has focused on two main areas: soluble factors and imaging the tumor vasculature. This pilot study combines functional and anatomic imaging using small animal
positional emission tomography (PET) and microcomputed tomography (micro-CT) techniques in a xenograft model of breast cancer to identify PET tracers likely to correlate with response to AZD2171. Tumor growth and imaging were compared with histologic changes in tumor vascularity, proliferation, and VEGF expression.

Materials and Methods

Orthotopic xenograft model
Six- to eight-week-old female athymic (nu/nu) mice (n = 75) were purchased from Harlan Sprague-Dawley (Indianapolis, IN) and acclimated for 7 to 10 days. As MCF-7 tumor cells require supplemental estrogen for growth in athymic mice, a controlled release estradiol pellet (0.72 mg, 60-day formulation) was injected s.c. via a sterile 14-gauge trocar at least 24 hours before tumor implantation. Under direct visualization, the right and left mammary fat pads were injected with $5 \times 10^6$ MCF-7 cells (non-VEGF expressing) or $1 \times 10^6$ MCF-7 cells (VEGF expressing) cells in 0.1 mL of Hanks’ buffered saline (14, 15). This dual implantation technique allowed comparison of physiologic imaging, tumor growth, and response to AZD2171 in tumors with high and low levels of VEGF production within the same animal. In preliminary studies, this dual implantation technique did not alter tumor growth or vascularity compared with tumors implanted singly (data not shown).

Once tumors were established (~5-8 mm diameter), mice were divided into five groups; baseline imaging (n = 15). Imaging 24 hours after a single dose of AZD2171 (n = 15), imaging 72 hours after initiation of AZD2171 treatment (n = 15), and imaging after 4 weeks of treatment with either AZD2171 (n = 15) or vehicle control (n = 15). AZD2171 (5 mg/kg/d) or vehicle (100 L) was administered by oral gavage once daily, starting on day 2 after implantation. A single observer (K.D. Miller) measured tumor diameter in two dimensions with vernier calipers twice weekly throughout treatment; tumor volume was calculated as $3.14 \times 6 \times$ largest diameter $\times$ smallest diameter squared. All animal experiments were done under a protocol approved by the Indiana University Institutional Animal Care and Use Committee.

Positron emission tomography imaging
PET imaging was done to measure tumor blood volume with [11C]carbon monoxide ([11C]CO; refs. 16, 17), perfusion with [18F]fluoromethane ([18F]FCH3; refs. 18, 19), and glucose utilization with [18F]fluorodeoxyglucose ([18F]FDG). Studies were done using the IndyPET-II scanner (20) using 120 CTI HR detector modules arranged in four detector banks. The detector banks are arranged to produce a transaxial field of view of 23 cm and an axial field of view of 15 cm. Within an axial plane, the average full-width-at-half-maximum resolution is 2.5 mm at the center of the field of view with minimal resolution degradation over the central 10 cm of the field of view (20). The NEMA-2001 sensitivity is 9,030 cps/MBq at the center of the scanner and 4,250 cps/MBq at a distance of 10 cm from the center of the field of view. All data are acquired with a 350 to 650 keV energy window. The IndyPET-II scanner collects all data in list mode (the time and location of individual events is recorded). Following data acquisition, events are combined to form event versus position histograms (sinogram) used for image reconstruction. List mode data acquisition enables the retrospective formation of sinogram data over discrete time intervals for tracer kinetic analysis.

Radioisotopes. [14N(p,α)[13C]CO was produced via the 14N(p,α)[13C] nuclear reaction starting from oxygen-18–enriched water, was azeotroped to dryness with acetonitrile in the presence of 10 mg Kryptofix 2.2.2, 2 mg K2CO3, and $-20$ mg Ag2O. Upon addition of 0.1 mL CH3I and 0.5 mL acetonitrile, [13C]CO was liberated and swept from the reaction vial with nitrogen gas. Flow was diverted through a water-ice bath and a C18 Sep-Pak cartridge to reduce the amount of carrier CH3I and acetonitrile collected in the gas bag. Typical production yielded $>300$ mCi of [18F]FCH3 in 100 mL of nitrogen carrier gas in the gas bag at end of bombardment. Ten-milliliter aliquots containing ~30 mCi of product were delivered for unit-dose preparation and administration.

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Gas tracer administration. An in-house gas administration system was developed for the [11C]CO and [18F]FCH3 studies (Fig. 1A). The mouse is positioned on the scanning bed with a mask surrounding the entire head. The mask, made from a 60 mL syringe, is cylindrical, ~25 mm in diameter, and 50 mm in length with a total internal volume of ~25 mL. A latex membrane with a central hole, through which the head of the mouse is inserted, covers the open end of the mask. This membrane provides a seal around the nose of the mouse while still being loose enough that it does not constrict blood flow or respiration. The mask has two female leur fittings. The first leur fitting is at the end of the syringe and the second is mounted in the side of the syringe. The side fitting is attached to a stopcock and a small air pump. The gas tracer
is prepared by loading ~37 MBq (~1 mCi) of tracer in 1 to 2 mL into a 3 mL syringe. This syringe is coupled to the end of the mask with a stopcock. After starting image acquisition, the stopcock to the air pump is closed and the gas tracer is introduced into the mask. The mouse is allowed to rebreathe the tracer for 2 minutes. After this uptake period, the stopcocks are opened to allow the air pump to circulate ~100 mL of room air through the mask and out the end, flushing the remaining tracer out of the mask.

\[^{[18]F}\]Fluorodeoxyglucose administration. \[^{[18]F}\]FDG (1 mCi) was administered i.v. to the mouse via tail vein.

**Imaging protocol.** Before each imaging session, anesthesia was administered to the animals in a laminar flow hood. Once anesthetized with ketamine, animals were placed in a small isolation chamber for the imaging procedures. The isolation chamber is integrated with the gas administration system previously described to isolate each animal from the external environment. Each animal imaging session began with a high resolution X-ray CT scan using the EVS MS-9 scanner (Enhanced Vision Systems Corp., London, Ontario, Canada) in the 100 μm voxel resolution mode. Images were reconstructed using a Feldcamp cone beam reconstruction algorithm. The CT image data serves as an anatomic reference to aid interpretation of the PET image data and is used to generate attenuation correction information to reconstruct quantitative images of radionuclide concentrations. PET data is acquired in list mode for 30 minutes and sorted into 17 frames, 4, 60-second frames, and 4, 300-second frames after the acquisition is complete. Frames are reconstructed with an 80 mm transaxial field of view, 0.625 mm pixels, and 47 slices spaced 3.15 mm apart. Reconstruction is done with filtered back-projection using a Hanning filter with a cutoff frequency of 4.2/ cm.

**Image registration and fusion.** We have developed a method for registering small animal scans acquired with the IndyPET-II PET scanner and the EVS RS-9 micro-CT scanner. After sedation, the subject is strapped to a scanning bed (previously described), which is mounted on the EVS micro-CT imaging stage for CT image acquisition. Following the CT scan, the bed is moved from the EVS micro-CT stage to the IndyPET II imaging stage. Both stages have identical mounts with precision pins so that the bed position is reproducible. A six-variable rigid body registration transform of the PET image to the CT image space is calculated using PET and CT images of a registration phantom consisting of parallel and perpendicular tubes filled with a mixture of \[^{[18]F}\]FDG and iodinated contrast agent. A CT image of the registration phantom is shown in Fig. 1B. Transformation variables were calculated using an algorithm that finds the ends of the tubes and calculates a geometric transform using the locations of the end points in the two image pairs. This method also provides a measure of fiducial registration error. Typically, the overall fiducial registration error is <0.4 mm. The phantom-derived registration transformation variables were applied to the animal CT and PET data sets to register the anatomic and functional image data sets for further evaluation.

**Positron emission tomography data analysis**

PET and CT images are registered and regions of interest are drawn on the tumors, thigh muscle, heart blood pool, and the left and right lungs using the CT image to identify the tumor and tissue regions. For the \[^{[18]F}\]FCH\(_3\) studies, time-activity curves are calculated by applying the regions of interest to each frame of the PET image.

**Blood volume estimates.** The \[^{[11]C}\]CO tracer tightly binds to RBC following administration and provides an image of RBC distribution in tissue. Fractional blood volume estimates are obtained by taking the ratio of \[^{[11]C}\]CO concentration in a tissue region of interest \(C_r(t)\) to the \[^{[11]C}\]CO concentration in a blood volume region of interest \(\text{left ventricular chamber of the heart; } C_c(t)\). To reduce statistical uncertainty, the blood volume fraction (BVF) estimate image data is integrated over a period of 10 minutes following equilibration of the \[^{[11]C}\]CO throughout the blood pool. A period of 5 minutes following tracer administration is used to assure equilibration of \[^{[11]C}\]CO within the vascular space before integration of the tissue and blood pool signal. BVF estimates were calculated using Eq. A.

\[
\text{BVF} = \frac{\int_0^5 C_r(t)dt}{\int_0^5 C_c(t)dt} = \frac{15}{5} C_r(t)dt
\]

Estimates of tumor blood volume were calculated by multiplying BVF by the physical tumor volume \(V_T\) obtained by placing regions of interest on the X-ray CT images.

\[
\text{BF} = \text{BVF} \cdot V_T
\]

**Perfusion estimates.** A one-compartment tracer kinetic model is fit to the tissue-activity curves using the lung curves as the input function. This model fits Eq. C to the measured tissue curve \(C_r(t)\). This model is the Kety-Schmidt single-compartment model for diffusible tracers (23, 24) with an additional term added to account for tracer in the vascular compartment.

\[
C_r(t) = (1 - \text{BVF}) \int_{t_1}^{t_2} K_i C_c(s) e^{-K_i V_c g'/t} ds + \text{BVF} \int_{t_1}^{t_2} C_c(t)dt
\]

The variables of the fit are \(K_v, V_v, \text{ and } \text{BVF}\), which are related to the traditional blood flow (perfusion) variable, distribution volume, and BFV. The applicability of this model follows from our assumption that the tracer concentration in the lungs is in equilibrium with the tracer concentration in the blood. The unknown ratio of blood to lung activity concentration is folded into the blood volume variable BVF.

Total blood flow (TBF) through the tumor was calculated by using Eq. D.

\[
\text{TBF} = K_i V_T
\]

**Fluorodeoxyglucose utilization estimates.** FDG uptake estimates are generated by placing regions of interest on PET images acquired over the time period from 45 to 60 minutes after tracer administration. This imaging time window was selected to enable sufficient clearance of nonmetabolized FDG from the tissue to provide an image that is dominated by tracer that has been phosphorylated by hexokinase and trapped in the tissue. Indices of tumor FDG uptake were generated by calculating the total FDG uptake in the tumor normalized by the mass of the animal and the injected dose of tracer. Eq. E was used to generate FDG uptake indices (UIFDG). \(\text{UIFDG}\) is the fraction of the injected dose in the tumor (\(\text{FDG}_T\)) normalized by the mass of the animal.

\[
\text{UIFDG} = \frac{V_T \int_{t_1}^{t_2} C_r(t)dt}{(20(60-45)/M/D)} = \frac{\text{FDG}_T M}{20} = \frac{V_T S IUV}{20}
\]

In Eq. E, \(V_T\) is the tumor volume derived from the CT images, \(C_r(t)\) is the PET measured tracer uptake in the tumor, \(M\) is the mass of the animal, and \(D\) is the injected dose of FDG. The ratio of \(M/20\) in this equation is to normalize the fraction of the injected dose to a standard 20 g mouse. The \(\text{UIFDG}\) therefore, represents the fraction of the injected dose that accumulates in the tumor for a standard 20 g mouse. In Eq. E, S IUV is the standardized uptake value normally used in clinical PET studies.

**Histology**

Primary tumors were resected after imaging and fixed in 10% buffered neutral formalin for subsequent histologic analysis of microvessel density (CD31), proliferation (Ki-67), and VEGF expression. Tumors after 4 weeks of treatment with AZD2171 or control frequently had necrotic centers; only viable tumor was processed for analysis.
Microvessel density

Specimens were dewaxed in xylene and rehydrated through descending grades of alcohol. Antigen retrieval was done in DAKO citrate buffer (pH 6; DAKO, Carpinteria, CA) for 20 minutes in a 95°C water bath. Endogenous peroxidase activity was blocked using 3% hydrogen peroxide. Endogenous biotin and avidin were blocked using DAKO avidin/biotin each for 10 minutes; nonspecific binding was minimized using a protein block for 20 minutes. Slides were then incubated for 30 minutes with DAKO prediluted mouse monoclonal CD31 then followed by sequential application of DAKO-LSAB2 biotinylated link antibody and DAKO-LSAB2 streptavidin labeled with alkaline phosphatase for 15 minutes each. 3,3′-Diaminobenzidine tetrahydrochloride was used as the chromogen for visualization; sections were counterstained with hematoxylin QS, dehydrated through ascending grades of alcohol, cleared in xylene, and mounted. The Ki-67 labeling index was determined by counting the number of positive nuclei in 1,000 tumor cells in at least five representative high-power fields (0.7386 mm² per 200× field). Any brown staining endothelial cell or cell cluster clearly separated from adjacent microvessels was counted. A vessel lumen or the presence of red cells was not required; microvessels with a diameter >50 μm were excluded. Both the maximum and average of all three fields were reported.

Proliferation

Sections were dewaxed in xylene and rehydrated through descending grades of alcohol. Antigen retrieval was done in DAKO Target Retrieval for 20 minutes in a 95°C water bath. Endogenous biotin and avidin were blocked using DAKO avidin/biotin each for 10 minutes. Endogenous peroxidase activity was blocked using 3% hydrogen peroxide. The slides were incubated for 30 minutes with DAKO mouse monoclonal Ki-67 (Mib-1) at a dilution of 1:60, followed by sequential application of DAKO-LSAB2 biotinylated link antibody and DAKO-LSAB2 streptavidin labeled with alkaline phosphatase for 15 minutes each. 3,3′-Diaminobenzidine tetrahydrochloride was used as the chromogen for visualization; sections were counterstained with hematoxylin QS, dehydrated through ascending grades of alcohol, cleared in xylene, and mounted. The Ki-67 labeling index was determined by counting the number of positive nuclei in 1,000 tumor cells in at least five representative high-power fields (>400) with selection of fields that contained viable tumor cells (28).

Vascular endothelial growth factor expression

Sections were dewaxed in xylene and rehydrated through descending grades of alcohol; slides were rinsed between grades with TBS. Antigen retrieval was done in DAKO Target Retrieval for 20 minutes in a 95°C water bath. Endogenous biotin and avidin were blocked using DAKO avidin/biotin each for 10 minutes. Endogenous peroxidase activity was blocked using 3% hydrogen peroxide. Nonspecific binding of antibody was minimized using a protein block for 20 minutes; excess protein block was tapped off the slides without rinsing. Slides were incubated for 1 hour with Biogenix (Bogota, Columbia) prediluted rabbit polyclonal VEGF diluting it further in a 1:2 ratio, followed by sequential application of DAKO-LSAB2 biotinylated link antibody and DAKO-LSAB2 streptavidin labeled with alkaline phosphatase for 15 minutes each. 3,3′-Diaminobenzidine tetrahydrochloride was used as the chromogen; sections were counterstained with hematoxylin QS. Slides were dehydrated through ascending grades of alcohol followed by clearing in xylene and mounted. VEGF expression was reported using the following scoring system: 0, negative; 1, weak staining in <10% of tumor cells; 2, moderate or high intensity staining in >10% of tumor cells. At least five representative high-power fields (>400) containing viable tumor cells were evaluated.

Statistical analyses

All statistical analyses were done using Student’s t test. P < 0.05 denoted the presence of a statistically significant difference. In previous studies, we found no acute alteration in any of the PET or histologic variables with vehicle control. To limit the number of animals required, acute time points (24 and 72 hours) were compared with baseline studies; tumor size was comparable in these groups. In vivo tumor growth analyses were done using GraphPad Prism version 3.02 for Windows (GraphPad Software, San Diego, CA; www.graphpad.com).

Results

Inhibition of in vivo tumor growth. As expected, MCF-7VEGF xenografts grew more rapidly and formed larger tumors than MCF-7neo xenografts in the absence of active treatment. Chronic once-daily oral administration of AZD2171 (5 mg/kg) induced a statistically significant (P = 0.027) regression of established MCF-7VEGF xenografts (Fig. 2). In contrast, AZD2171 treatment did not induce regression, but inhibited the growth of MCF-7neo tumors, although this did not reach statistical significance using a Kaplan-Meier–based analysis.

Tumor histology. MCF-7VEGF tumors were more vascular (P = 0.0001) and had a higher proliferative fraction (P = 0.002) than MCF-7neo tumors at baseline. An acute decrease in tumor microvessel density was apparent 24 hours after treatment with AZD2171 in MCF-7VEGF tumors (P = 0.057) with an even more pronounced decrease at 72 hours (P = 0.0003). Despite continued control of tumor growth, MCF-7VEGF tumors exposed to chronic daily AZD2171 therapy for 4 weeks had a microvessel density similar to MCF-7VEGF tumors treated with vehicle alone. Similarly, proliferation as measured by Ki-67 index acutely decreased in MCF-7VEGF tumors (P = 0.036 at 24 hours and P = 0.0016 at 72 hours) but returned to baseline during chronic therapy. In contrast, no significant difference in tumor microvessel density or proliferation was seen in MCF-7neo tumors after either acute or chronic exposure to AZD2171.
AZD2171 (Fig. 3). As anticipated, a higher VEGF expression was detected in the MCF-7VEGF tumor compared with the MCF-7neo tumors. No acute change in VEGF expression was detected with AZD2171 therapy; however, chronic therapy with AZD2171 increased VEGF expression in both MCF-7 neo and MCF-7 VEGF tumors (P = 0.02 and P = 0.01, respectively; Fig. 3).

Physiologic imaging. Perfusion (tissue blood flow) is increased >20-fold and is more variable in MCF-7VEGF compared with MCF-7neo tumors (Fig. 4A). An acute decrease in perfusion is apparent in MCF-7 VEGF tumors 24 hours after starting AZD2171 therapy (P = 0.05). Perfusion is more dramatically decreased after 72 hours of AZD2171 treatment (P = 0.03). Perfusion in MCF-7neo tumors is similar to that of normal muscle and is not altered by AZD2171. [18F]FCH$_3$ volume of distribution is not altered by treatment with AZD2171 (Fig. 4B). Total tumor blood flow does not correspond with tumor volume, confirming that the treatment-induced perfusion changes are not due to differences in tumor size (Fig. 4C).

Although we detected a statistically significant decrease in blood volume in MCF-7VEGF tumors 24 hours after AZD2171 therapy by [13C]CO (data not shown), no difference was seen at 72 hours or after 4 weeks of therapy. Importantly, we observed a consistent relationship between total tumor blood volume and tumor volume independent of treatment in MCF-7VEGF tumors (Fig. 5A). Tumor-associated blood volume was increased 3-fold in MCF-7VEGF compared with MCF-7neo tumors (data not shown); no consistent change with treatment was identified in either xenograft model (Fig. 5B).

Glucose utilization, as estimated by the [18F]FDG uptake index, was associated with tumor size and did not change with acute or chronic AZD2171 therapy in MCF-7VEGF tumors (data not shown). Unlike blood flow and blood volume, glucose utilization was similar in MCF-7neo and MCF-7VEGF tumors (Fig. 6).

Discussion

We developed a breast cancer xenograft model system allowing direct comparison of tumor response, histology, and physiologic imaging based on VEGF expression. Dual implantation of controls and VEGF-transfected MCF-7 cells allows each animal to serve as its own control, decreasing both the variability and number of animals required. As expected, MCF-7VEGF tumors were more vascular than MCF-7neo tumors (14). The increased microvessel density seen with MCF-7VEGF tumors was associated with an increase in proliferation as measured by Ki-67. The increase in tumor growth, microvessel density, proliferation, and perfusion confirm that VEGF secreted from the MCF-7VEGF xenograft had little effect on the MCF-7neo xenograft in the contralateral fat pad. Using this model, we have shown that the
Treatment with AZD2171 induces an acute decrease in tumor perfusion in MCF-7VEGF tumors that can be detected within 24 hours; a further decrease was detected at 72 hours. Histologic analysis of viable tissue from treated tumors found decreases in tumor microvessel density and proliferation in a time course that paralleled the decrease in perfusion. Although tumor blood volume was increased in the MCF-7VEGF tumors, blood volume was not altered by AZD2171 therapy. The tumor blood volume measured by PET represents blood in both large and small vessels within the tumor. The inability to detect blood volume changes using PET in the setting of observed changes in microvessel density by histology suggests that small vessels, easily quantified by histology, are most affected by AZD2171 but carry only a small fraction of the total tumor blood volume. Compensatory mechanisms, as yet not fully defined, restore perfusion during 4 weeks of continuous daily therapy in our model. We detected an increase in VEGF expression in both MCF-7neo tumors and MCF-7VEGF tumors exposed to chronic AZD2171 therapy. Inhibition of VEGF signaling by AZD2171 could induce tumor hypoxia, with a resultant increase in VEGF expression. Other potential compensatory mechanisms, not investigated in this study, include induction of other proangiogenic peptides, mutation of the VEGFR tyrosine kinase, or co-option of existing host vessels. As our study did not include pharmacokinetic assessments, we cannot exclude increased clearance or altered distribution of AZD2171 during chronic therapy.

Physiologic imaging using the IndyPET-II system was able to distinguish VEGF and non-VEGF-expressing tumors within a single animal. Interestingly, blood volume and perfusion of the non-VEGF-expressing MCF-7neo tumors is similar to that of normal muscle and is not altered by AZD2171. B, [18F]FCH3, volume of distribution is not altered by treatment with AZD2171. C, total tumor blood flow does not correlate with tumor volume in MCF-7VEGF tumors (or in MCF-7neo tumors; data not shown). Columns (A and B), mean; bars, SE.

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Effect of the small-molecule VEGFR-2 tyrosine kinase inhibitor AZD2171 on tumor growth is at least partially dependent on VEGF expression. MCF-7neo tumors remained growth inhibited, whereas VEGF-expressing MCF-7VEGF significantly regressed. Physiologic imaging using the IndyPET-II system was able to distinguish VEGF and non-VEGF-expressing tumors within a single animal. Interestingly, blood volume and perfusion of the non-VEGF-expressing MCF-7neo tumors was similar to normal skeletal muscle in this model. Although PET imaging of tumor perfusion and/or blood volume has been incorporated into early clinical trials (18, 29, 30), few studies have compared perfusion in tumor and normal tissue. Wells et al. (31) found that [15O]CO2 PET measurements of regional blood flow are reproducible in patients with predominantly intra-abdominal malignancies but did not compare perfusion to normal tissues. In one study of eight patients with simultaneous parallel measurements, renal tumor perfusion was significantly lower than perfusion of the normal kidney (32).
therapy although this has not been observed previously in other studies.

The increase in microvessel density we observed during chronic therapy differed from the results reported by Wedge et al. (11) using a s.c. Calu-6 lung cancer xenograft model. Treatment with AZD2171 produced a 47% reduction in vessel number and a 40% reduction in CD31 area after 52 hours; vessel number and CD31 area continued to decrease throughout the 21-day treatment period (≤70% at day 21). Imaging to assess perfusion or blood volume was not done. Critical differences in methodology may account for the difference in results. First, the complex interaction between tumor cells and host microenvironment has been recognized for over a century (33–35). More recently, the effect of the local tumor microenvironment on angiogenesis and response to antiangiogenic therapy has become increasingly clear (36–38). We would expect the tumor microenvironment to affect compensatory mechanisms and the development of resistance as well. Second, Wedge et al. analyzed the entire tumor, normalizing the number of vessels or CD31 area to tumor size. In contrast, we scanned the periphery of the tumor for the area of greatest vascularity. As such, small areas of intense vascularity would be missed.

Importantly, although perfusion, microvessel density, and proliferation returned to baseline levels during chronic therapy in our study, tumor growth control was maintained. One possible explanation for these seemingly disparate results might be that acute treatment prunes the immature vessels at the tumor periphery leaving the mature vessels in the center of the tumor (39). These mature, normal vessels would support persistence of a residual smaller, but well-vascularized, tumor with proliferation matched by apoptosis (40, 41). Unfortunately, we were unable to assess apoptosis to investigate this hypothesis. Alternatively, overt resistance and tumor growth may have ensued quickly with further therapy, although resistance to VEGFR tyrosine kinase inhibitors (including AZD2171) has not been documented previously in tumor xenograft models (11, 42, 43).

In our study, total [18F]FDG uptake correlated only with tumor size; no association was found with VEGF expression, [18F]FCH₃, or [11C]CO uptake or AZD2171 treatment. Tumor glucose utilization is tightly controlled. Consequently, FDG uptake is complex and may be affected by many factors besides perfusion and blood volume (44, 45). Two small studies found a correlation with [18F]FDG uptake and tumor blood flow as assessed by [13O]H₂O in primary breast cancers (46–48). Nonetheless, our results suggest that [18F]FDG imaging is not a reliable surrogate for changes in tumor vasculature and activity of antiangiogenic agents.

Using this model, we previously showed a dose-dependent decrease in tumor blood volume and perfusion after 4 weeks of treatment with ZD6474, an inhibitor of the VEGFR-2 and epidermal growth factor tyrosine kinases; the acute effects of ZD6474 therapy were not imaged (49). In contrast, our results differ from studies using another VEGFR tyrosine kinase inhibitor in this model. Whereas AZD2171 had little effect on tumor blood volume, SU11248 treatment induced an acute decrease in tumor blood volume that was partially attenuated during chronic therapy. Perfusion increased acutely but decreased throughout chronic SU11248 therapy (50). Although both agents inhibit the VEGFR tyrosine kinases, the spectrum of other kinases they inhibit differs. These results show the need for careful consideration of the biological variable measured when using these imaging techniques as surrogates of antiangiogenic activity. Although the image surrogates used in these studies (blood volume, perfusion, and glucose utilization) are all potentially linked to AZD2171 and SU11248 through modulation of the VEGFR tyrosine kinases, none directly measure kinase inhibition. Consequently, each imaging technique needs to be developed and tested specifically for each agent rather than as a general surrogate for antiangiogenic activity.

The ultimate test of any surrogate marker or imaging technique will come in the clinic. The lack of gold standard and variability of physiologic imaging with different antiangiogenic agents in the same model argues for increased preclinical study. Xenograft models like the one we used here can be used to screen potential imaging techniques, allowing selection of the method most predictive of response to a specific agent for incorporation into clinical trials.

Acknowledgments

We thank Kerry Sanders, Winston Baity, Tanya Martinez, Kristan Boling, and Terry McBride for technical assistance, and Julianne Jürgensmeier and Stephen Wedge for helpful discussions during the design and analysis of this study.

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Fig. 6. Tumor glucose utilization as measured by [18F]FDG. Glucose utilization was similar in MCF-7neo and MCF-7VEGF tumors and does not change with acute or chronic AZD2171 therapy. The percentage of the administered FDG that accumulated in the tumor normalized to a reference animal weight of 20 g ([NPID/20]: percentage of FDG uptake in tumor expected if individual animals weighed 20 g) is plotted versus tumor volume. The close correspondence of each curve serves as an indicator that the FDG uptake per unit volume or cell within the tumor (slope of curves) is unchanged between groups and following AZD2171 therapy.
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