Immuno-PET of Tissue Factor in Pancreatic Cancer

Hao Hong¹, Yin Zhang², Tapas R. Nayak¹, Jonathan W. Engle², Hing C. Wong³, Bai Liu³, Todd E. Barnhart², and Weibo Cai¹,²,⁴

¹Department of Radiology, University of Wisconsin, Madison, Wisconsin; ²Department of Medical Physics, University of Wisconsin, Madison, Wisconsin; ³Altor BioScience Corp., Miramar, Florida; and ⁴Carbone Cancer Center, University of Wisconsin, Madison, Wisconsin

Upregulation of tissue factor (TF) expression leads to increased patient morbidity and mortality in many solid tumor types. The goal of this study was to develop a PET tracer for imaging of TF expression in pancreatic cancer. Methods: ALT-836, a chimeric antihuman TF monoclonal antibody, was conjugated to 2-S-(4-isothiocyanatobenzyl)-1,4,7-triazacyclononane-1,4,7-triaceitic acid (p-SCN-Bn-NOTA) and labeled with ⁶⁴Cu. To compare the TF binding affinity of ALT-836 and NOTA-ALT-836, flow cytometry analysis was performed in 3 pancreatic cancer cell lines with different expression levels of TF (from low to high: PANC-1, ASPC-1, and BXPC-3). PET, biodistribution, blocking, and histology studies were performed on pancreatic tumor-bearing mice to evaluate the ability and specificity of ⁶⁴Cu-NOTA-ALT-836 to target TF in vivo. Results: There was no difference in TF binding affinity between ALT-836 and NOTA-ALT-836. ⁶⁴Cu-labeling was achieved with high yield and specific activity. Serial PET revealed that the uptake of ⁶⁴Cu-NOTA-ALT-836 in BXPC-3 tumors (high TF expression) was 5.7 ± 1.8, 10.4 ± 0.8, and 16.5 ± 2.6 percentage injected dose per gram at 4, 24, and 48 h after injection, respectively (n = 4), significantly higher than that in the PANC-1 and ASPC-1 tumors. Biodistribution data as measured by γ-counting were consistent with the PET findings. Blocking experiments and histology further confirmed the TF specificity of ⁶⁴Cu-NOTA-ALT-836. Conclusion: Herein we report the first successful PET imaging of TF expression. Persistent and TF-specific uptake of ⁶⁴Cu-NOTA-ALT-836 was observed in pancreatic cancer models.

Key Words: tissue factor (TF); pancreatic cancer; positron emission tomography (PET); monoclonal antibody; ⁶⁴Cu; immuno-PET

DOI: 10.2967/jnumed.112.105460

Tissue factor (TF), also known as platelet tissue factor, factor III, thrombokinase, or CD142, is a 47-kDa transmembrane glycoprotein receptor present in subendothelial tissue, platelets, and leukocytes (1). The principal function of TF is to localize the coagulation serine protease factor VII/VIIa to the cell surface, thereby initiating the coagulation cascade. In normal human tissues, TF is expressed only in extravascular cells that comprise the vascular adventitia and organ capsules, forming a hemostatic barrier surrounding the vasculature to prevent blood loss (2). Damages to the endothelial lining of blood vessels can lead to the initiation of TF expression, which ultimately results in activation of the coagulation system (3). The expression of TF is upregulated in several scenarios such as thrombosis, tumor growth, metastasis, and angiogenesis (4,5). Clinically, the pathologic consequences of unregulated activation of the coagulation system are a major cause of increased morbidity and mortality in cancer patients (6).

With about 44,030 estimated new cases and 37,660 estimated deaths in 2011 in the United States, pancreatic cancer is among the most incurable of malignant diseases, with a dismal 5-y survival rate of about 5% (7). Most pancreatic adenocarcinoma has been found to be associated with several hemostatic complications such as venous and arterial thrombosis, migratory thrombophlebitis, pulmonary embolism, disseminated intravascular coagulation, and simultaneously aggressive tumor angiogenesis and metastasis (8,9). It is now generally recognized that a strong correlation exists between the aberrant expression of TF and pancreatic cancer (10,11), as well as many other solid tumor types such as breast cancer (4), glioma (12), leukemia (13), and lung cancer (14). This correlation signifies the potential not only for inhibiting tumor angiogenesis/metastasis and the associated thrombosis through modulation of TF signaling but also for using TF as a prognostic marker (15).

ALT-836 (formerly known as Sunol-cH36) is a chimeric antihuman TF monoclonal antibody that binds to the factor X (FX)-binding site in TF with subnanomolar affinity, which can block the activation of FX and inhibit the coagulation cascade (16). ALT-836 did not exhibit a significant effect on TF-initiated clotting times in assays using plasma and lipitated TF from mice, indicating low binding affinity of ALT-836 to murine TF. Recently, ALT-836 has been evaluated in a phase I clinical trial in acute lung injury/acute respiratory distress syndrome (ALI/ARDS) (17). Based on the encouraging phase I results, a phase II clinical trial is currently ongoing. In addition, ALT-836 is in a phase I clinical trial for locally advanced or metastatic solid tumors.
PET has been widely used in clinical oncology for tumor staging and for monitoring therapeutic efficacy (18–21). Although anti-TF antibodies have been used for imaging of thrombus formation and atherosclerotic plaques in animal models (22,23), no PET study of TF has been reported to date. The goal of this proof-of-principle study was to investigate whether ALT-836 can be used for PET imaging of TF in preclinical pancreatic cancer models, using $^{64}$Cu as the radiolabel, which can open new avenues for future patient selection and stratification in TF-related clinical trials, as well as for monitoring therapeutic response. Various in vitro, in vivo, and ex vivo studies were performed to validate the PET findings by confirming the TF specificity of $^{64}$Cu-labeled ALT-836.

MATERIALS AND METHODS

Chemicals

ALT-836 was provided by Altor Bioscience Corp. Alexa Fluor 488 and Cy3-labeled secondary antibodies were purchased from Jackson ImmunoResearch Laboratories, Inc. S-2-(4-isothiocyanatobenzyl)-1,4,7-triazacyclononane-1,4,7-triacetic acid (p-SCN-Bn-NOTA) and Chelex 100 resin (50–100 mesh) were purchased from Macronova. Phosphate-buffered saline with 2% bovine serum albumin at room temperature, washed 3 times with cold phosphate-buffered saline with 2% bovine serum albumin at room temperature, washed 3 times with cold phosphate-buffered saline, and centrifuged at 1,000 rpm for 5 min. The cells were then

Cell Lines and Animal Models

BXPC-3, ASPC-1, and PANC-1 pancreatic cancer cell lines were purchased from the American Type Culture Collection and cultured according to the supplier’s instructions. Cells were used for in vitro and in vivo experiments when they reached approximately 75% confluence. All animal studies were conducted under a protocol approved by the University of Wisconsin Institutional Animal Care and Use Committee. Four- to 5-wk-old female athymic nude mice were purchased from Harlan, and tumors were established by subcutaneously injecting $5 \times 10^6$ cells—suspended in 100 $\mu$L of a 1:1 mixture of culture medium and Matrigel (BD Biosciences)—into the front flank of mice. Mice were used for in vivo experiments when the diameter of the tumors reached 5–8 mm (4–6 wk after inoculation).

Antibody Conjugation and Radiolabeling

NOTA-ALT-836 was prepared using a method similar to one previously described (24), with the reaction ratio of p-SCN-Bn-NOTA:ALT-836 being 25:1. $^{64}$Cu labeling and purification were also performed using a method similar to the one previously described (24), with 37 MBq of $^{64}$Cu per 25 $\mu$g of NOTA-ALT-836.

Flow Cytometry

The TF binding activities of ALT-836 and NOTA-ALT-836 to BXPC-3, ASPC-1, and PANC-1 cells were evaluated by flow cytometry. Briefly, cells were harvested and suspended in cold phosphate-buffered saline with 2% bovine serum albumin at a concentration of $5 \times 10^6$ cells/mL. The cells were incubated with ALT-836 or NOTA-ALT-836 (1 or 5 $\mu$g/mL) for 30 min at room temperature, washed 3 times with cold phosphate-buffered saline, and centrifuged at 1,000 rpm for 5 min. The cells were then

incubated with Alexa Fluor 488-labeled goat antihuman IgG for 30 min at room temperature. Afterward, the cells were washed and analyzed using a FACSCalibur 4-color analysis cytometer (Becton-Dickinson) with FlowJo analysis software (Tree Star, Inc.).

PET/CT and Biodistribution Studies

PET and CT scans, image reconstruction, and region-of-interest analysis of the PET data were performed using an Inveon microPET/microCT rodent model scanner (Siemens Medical Solutions USA, Inc.) as described earlier (24,25). Each tumor-bearing mouse was injected with $5–10$ MBq of $^{64}$Cu-NOTA-ALT-836 via the tail vein, and 5- to 10-min static PET scans were obtained at various time points after injection. Tracer uptake was presented as percentage injected dose per gram of tissue (%ID/g) (mean ± SD; 4 mice per group). Each mouse of another group of 4 BXPC-3 tumor-bearing mice was injected with 1 mg of unlabeled ALT-836 at 2 h before $^{64}$Cu-NOTA-ALT-836 administration to evaluate the TF specificity of the tracer in vivo (i.e., blocking experiment). Biodistribution studies were performed after the last PET scans at 48 h after injection to validate the PET data. The tumors, liver, spleen (i.e., tissues with significant uptake of $^{64}$Cu-NOTA-ALT-836), and muscle were also frozen and cryosectioned for histologic analysis.

Histology

Frozen tissue slices of 5-µm thickness were fixed with cold acetone for 10 min and dried in the air for 30 min. After rinsing with phosphate-buffered saline and blocking with 10% donkey serum for 30 min at room temperature, the slices were incubated with ALT-836 (2 $\mu$g/mL) for 1 h at $4^\circ$C and visualized using Alexa Fluor 488-labeled goat antihuman IgG. The tissue slices were also stained for endothelial marker CD31 as described previously (26,27). After washing with phosphate-buffered saline, the slices were incubated with rat antimouse CD31 antibody (2 $\mu$g/mL) for 1 h, followed by Cy3-labeled donkey antirat IgG for 30 min. All images were acquired with an Eclipse Ti microscope (Nikon).

Statistical Analysis

Quantitative data were expressed as mean ± SD. Means were compared using the Student $t$ test. $P$ values of less than 0.05 were considered statistically significant.

RESULTS

Characterization of NOTA-ALT-836

Based on flow cytometry analysis of both BXPC-3 and ASPC-1 cells (high and medium TF expression, respectively (15)), there were no observable differences between ALT-836 and NOTA-ALT-836 at 1 or 5 $\mu$g/mL (Fig. 1). For PANC-1 cells, which are TF-negative (15), neither ALT-836 nor NOTA-ALT-836 bound to the cells at these concentrations. Together, these results indicated that NOTA conjugation did not alter the antigen-binding affinity or specificity of ALT-836, with little nonspecific binding in cell culture. Furthermore, the binding level of ALT-836 and NOTA-ALT-836 to the 3 cell lines was in accordance with TF expression level in these cells (BXPC-3 > ASPC-1 > PANC-1).

Radiochemistry

$^{64}$Cu-labeling, including final purification using PD-10 columns, took $70 ± 10$ min ($n = 6$). The decay-corrected radiochemical yield was 79.6% ± 5.4%, based on 25 $\mu$g of
NOTA-ALT-836 per 37 MBq of $^{64}$Cu, with radiochemical purity greater than 95%. The ratio of $^{64}$Cu activity to antibody mass was 1.2 GBq/mg (4.86 Ci/μmol), assuming complete antibody recovery after size-exclusion chromatography.

PET

Time points of 4, 24, and 48 h after injection, based on the half-life of $^{64}$Cu (12.7 h), were chosen for serial PET scans after intravenous injection of $^{64}$Cu-NOTA-ALT-836 into tumor-bearing mice. The coronal slices that contained the tumors are shown in Figure 2A, and representative PET/CT images of a BXPC-3 tumor-bearing mouse at 24 h after injection are shown in Figure 2B. The quantitative data obtained from region-of-interest analysis are shown in Figure 3.

Liver uptake and blood-pool activity of $^{64}$Cu-NOTA-ALT-836 were prominent at early time points and gradually declined over time, similarly to other radiolabeled antibodies reported in the literature. In the PANC-1 tumor, which does not express a detectable level of TF, the uptake of $^{64}$Cu-NOTA-ALT-836 was low (1.7 ± 0.8, 2.8 ± 0.7, and 2.2 ± 0.6 %ID/g at 4, 24, and 48 h after injection, respectively; $n$ = 4; Figs. 2A and 3A). The liver uptake (18.2 ± 1.8, 16.7 ± 1.4, and 10.2 ± 0.9 %ID/g at 4, 24, and 48 h after injection, respectively) and blood radioactivity (10.6 ± 1.3, 6.8 ± 1.5, and 3.5 ± 0.7 %ID/g at 4, 24, and 48 h after injection, respectively; $n$ = 4; Figs. 2A and 3A) were similar to those in the other 2 tumor models. Uptake of $^{64}$Cu-NOTA-ALT-836 in the ASPC-1 tumors (medium TF expression) was higher than that in the PANC-1 tumors, at 2.1 ± 0.7, 4.6 ± 0.8, and 5.3 ± 0.9 %ID/g at 4, 24, and 48 h after injection, respectively ($n$ = 4; Figs. 2A and 3B). In BXPC-3 tumor-bearing mice, tumor uptake of $^{64}$Cu-NOTA-ALT-836 was clearly visible as early as 4 h after injection and gradually over time (5.7 ± 1.8, 10.4 ± 0.8, and 16.5 ± 2.6 %ID/g at 4, 24, and 48 h after injection, respectively; $n$ = 4; Figs. 2A and 3C). These values were significantly higher than those of the PANC-1 and ASPC-1 tumors at all time points ($P < 0.05$), indicating TF specificity of the tracer.

Administration of a blocking dose of ALT-836 at 2 h before $^{64}$Cu-NOTA-ALT-836 injection reduced the tumor uptake to 0.7 ± 0.4, 2.3 ± 0.9, and 2.8 ± 0.7 %ID/g at 4, 24, and 48 h after injection, respectively ($n$ = 4; Figs. 2A and 3D), which was significantly different from that of mice injected with $^{64}$Cu-NOTA-ALT-836 alone ($P < 0.05$ at all 3 time points). Slightly higher radioactivity in the blood was observed at early time points (12.4 ± 0.5 and 6.9 ± 2.1 %ID/g at 4 and 24 h after injection, respectively; $n$ = 4; Fig. 3D) in the blocking group, whereas liver uptake of $^{64}$Cu-NOTA-ALT-836 (15.6 ± 0.4, 12.8 ± 0.3, and 10.5 ± 0.6 %ID/g at 4, 24, and 48 h after injection, respectively; $n$ = 4; Fig. 3D) was comparable to that of mice injected with $^{64}$Cu-NOTA-ALT-836 alone. Overall, tracer uptake in all major organs was similar between the 2 groups, yet the BXPC-3 tumor uptake was significantly higher in the $^{64}$Cu-NOTA-ALT-836-dosed mice than in the blocking group, confirming the TF specificity of the tracer in vivo.

An additional group of 3 mice was injected with PANC-1 cells on the left flank and BXPC-3 cells on the right flank and subjected to serial $^{64}$Cu-NOTA-ALT-836 PET scans. Since BXPC-3 tumors grow faster than PANC-1 tumors, the tumor diameters at the time of the PET scans were about 7 mm for BXPC-3 and about 4 mm for PANC-1. Significant uptake of $^{64}$Cu-NOTA-ALT-836 in the BXPC-3 tumors could be observed at 24 h and 48 h after injection, whereas tracer uptake in the PANC-1 tumors was at the background level (Fig. 3E). Noninvasive detection of TF-positive tumors but not TF-negative tumors in the same mice further validated that $^{64}$Cu-NOTA-ALT-836 can specifically target TF in vivo.

Biodistribution Studies

All mice were euthanized after the terminal PET scans at 48 h after injection for biodistribution and histology studies to validate the in vivo PET data. Besides the tumors, the blood, liver, and spleen also had a significant accumulation of radioactivity at 48 h after injection, as expected for a radiolabeled antibody that typically has a long circulation half-life and hepatic clearance (Fig. 4A). Appreciable kidney uptake (−5 %ID/g) was also observed, likely due to a small fraction of $^{64}$Cu detaching from NOTA-ALT-836 and degrading of the tracer over time (lower-molecular-weight species can undergo renal clearance). $^{64}$Cu-NOTA-ALT-836 showed a similar biodistribution profile in BXPC-3, ASPC-1, and

**FIGURE 1.** Flow cytometry analysis of various concentrations of ALT-836 and NOTA-ALT-836 in 3 pancreatic cancer cell lines: PANC-1 (TF-negative), ASPC-1 (medium TF expression), and BXPC-3 (high TF expression).
PANC-1 tumor-bearing mice in all the major organs and tissues except the tumor. When compared with all major organs in mice, BXPC-3 tumor uptake of $^{64}$Cu-NOTA-ALT-836 was the highest and provided excellent tumor contrast, with a tumor-to-muscle ratio of $24.2 \pm 2.1$ at 48 h after injection ($n = 4$).

Preinjection of a blocking dose of ALT-836 led to a significantly decreased uptake in BXPC-3 tumor ($P < 0.01$; $n = 4$; Fig. 4B) but not in other normal tissues including the blood and liver, corroborating the PET findings. Overall, consistent quantitative data from ex vivo biodistribution studies and in vivo PET scans demonstrated that region-of-interest analysis of noninvasive PET scans accurately reflected tracer distribution in vivo, as well as TF specificity of $^{64}$Cu-NOTA-ALT-836.

**Histology**

Immunofluorescence TF/CD31 staining revealed that TF expression was prominent on the BXPC-3/ASPC-1 tumor cells but absent on the PANC-1 cells (Fig. 5). TF staining of mouse liver, spleen, and muscle gave a low signal, since not only do these tissues not express TF at a high level, but also the binding affinity of ALT-836 to murine TF is low. Therefore, uptake of $^{64}$Cu-NOTA-ALT-836 in mouse liver and spleen was largely unrelated to TF binding and more likely related to nonspecific capture by the reticuloendothelial system and hepatic clearance of the tracer.

**DISCUSSION**

TF is overexpressed in most cancer types (28,29). Mounting literature data have suggested a key role for TF in the development of cancer-associated thrombosis and tumor growth, angiogenesis, and metastasis. As a direct antagonist of TF, ALT-836 can be used to block TF signaling in cancer, thereby inhibiting cancer-associated venous thromboembolism and tumor development and metastasis. In preclinical studies including nonhuman primates, ALT-836 exhibited potent antithrombotic and antiinflammatory activities with a good safety profile (16). In patients with coronary artery disease and ALI/ARDS, single-dose administration of ALT-836 was found to be safe and exhibited anticoagulant and antiinflammatory effects (17). A phase II study using a multidose regimen of ALT-836 is being conducted in patients with ALI/ARDS.

With these encouraging results, a phase I, open-label, multicenter, dose-escalation study of ALT-836 in combination with gemcitabine for locally advanced or metastatic solid tumors is currently ongoing, with the primary outcome measures being maximum tolerated dose of ALT-836 in combination with gemcitabine, safety profile, number and severity of treatment-related adverse effects, clinical benefit, progression-free survival, and others (ClinicalTrials.gov identifier, NCT01325558). The development of a PET tracer based on ALT-836, which does not affect its binding affinity and specificity to TF, can play multiple roles in future clinical trials of ALT-836 such as patient selection, which can significantly improve the response rate in cancer patients. In addition, radiolabeled ALT-836 can be used to more effectively study the pharmacokinetics of this antibody, as well as to monitor the therapeutic effect of drugs that act through modulation of TF signaling.

Pancreatic cancer is the fourth leading cause of cancer death in the United States (30). With a 5-y survival rate of about 5% and a median survival of less than 6 mo, the diagnosis of pancreatic cancer carries one of the most dismal prognoses. Because of the lack of specific symptoms...
and limitations in diagnostic methods, pancreatic cancer is nearly undetectable during its formative stages. Many pancreatic cancer cells are naturally resistant to current chemotherapy and radiation therapy, and to date, most known pancreatic cancer antigens have generated a relatively weak immune response (30,31). Early detection methods are under development but do not yet exist for pancreatic cancer. The correlation between TF and pancreatic cancer development and metastasis makes TF a useful marker for diagnosis (10), and the development of a PET tracer for TF imaging is of critical importance.

To the best of our knowledge, imaging of TF in cancer has not been reported to date. Over the last decade, a myriad of preclinical and clinical data have suggested a promising future for immuno-PET in the management of cancer patients (32). In this study, we have successfully developed and characterized 64Cu-labeled ALT-836 for PET imaging of tumor TF expression in vivo. Our data showed that 64Cu-NOTA-ALT-836 specifically bound to TF-positive BXPC-3 tumors but not to TF-negative PANC-1 tumors, and the tumor uptake of 64Cu-NOTA-ALT-836 was closely associated with TF expression level in the tumor. Even within the same mouse, 64Cu-NOTA-ALT-836 can delineate and differentiate tumors with different TF expression (BXPC-3 vs. PANC-1, Fig. 3E). Such TF specificity in vivo makes 64Cu-NOTA-ALT-836 a PET tracer with broad potential applications in many clinical situations, not only in cancer but also in many other diseases where TF plays a key role (e.g., ALI/ARDS). To facilitate future clinical translation, further testing of radiolabeled ALT-836 in orthotopic tumor models will be needed since they have different tumor tissue microenvironments and are more clinically relevant than the subcutaneous models used in this proof-of-principle study.

Because the pancreas is near the liver, tracer uptake may be masked by appreciable signal in the liver. Usually, pancreatic cancer is diagnosed through the presence of metastases (33); therefore, we envision that 64Cu-NOTA-ALT-836 may be more useful in the metastatic setting than in initial detection of the primary tumor. Guiding surgery with molecularly targeted fluorescent agents has attracted enormous interest over the last decade (34). Future development of fluorescently labeled ALT-836, which emits in the near-infrared
range (700–900 nm (24,35)) and hence has good signal penetration with low autofluorescence background, may be useful for surgical guidance of tumor removal.

The advantages of antibody-based tracers are that they are antigen-specific and have high binding affinity and absolute tumor uptake, making them suitable for internal radiotherapy applications or targeted delivery of anticancer drugs. To provide more insight about the long-term behavior of ALT-836 in vivo, other longer-lived isotopes (e.g., $^{89}$Zr, which has a decay half-life of 3.3 d (25)) can be explored in future studies. The major limitations of antibody-based imaging are slow tumor accumulation and high background signal in the reticuloendothelial system, which may be overcome by peptide-, small-molecule–, or antibody fragment–based tracers (36,37). For pancreatic cancer, since the primary metastasis site is also liver, development of TF-targeting PET probes with low liver uptake will be mandatory for a successful clinical translation.

One of the key requirements for accurate PET imaging with $^{64}$Cu-labeled antibodies is that the tracer should be sufficiently stable during the imaging period, since the PET scanner detects the distribution of $^{64}$Cu rather than the antibody itself. It is now generally agreed that NOTA is one of the best chelators for $^{64}$Cu labeling. A recent elegant study comparing the effect of several bifunctional chelators on the biodistribution of a $^{64}$Cu-labeled antibody (38) concluded that thermodynamic stability of $^{64}$Cu–chelator complexes did not significantly influence tumor uptake of the tracer but dramatically affected distribution in normal tissues. Consistent with this report, our own studies also showed that $^{64}$Cu-NOTA– and $^{64}$Cu-DOTA (DOTA denotes 1,4,7,10-tetraazacyclododecane-1,4,7,10-tetraacetic acid)–based PET tracers behaved similarly in terms of tumor uptake yet the use of NOTA as the chelator resulted in significantly lower liver uptake (39,40).

**CONCLUSION**

Herein we have reported the development, characterization, and in vivo investigation of a $^{64}$Cu-labeled antibody for PET of TF expression in pancreatic cancer. Rapid, persistent, and TF-specific uptake of $^{64}$Cu-NOTA-ALT-836 in the BXPC-3 tumor was observed and was further validated by various in vitro, in vivo, and ex vivo experiments.
Furthermore, tumor uptake of ⁶⁴Cu-NOTA-ALT-836 in various pancreatic tumors was related to their TF expression level. On further optimization and development, such TF-targeted PET tracers can be translated into the clinic for improving the management of pancreatic cancer patients. Lastly, these tracers can also play multiple roles not only in many other solid tumors types but also in many other diseases for which TF signaling is upregulated.

DISCLOSURE STATEMENT

The costs of publication of this article were defrayed in part by the payment of page charges. Therefore, and solely to indicate this fact, this article is hereby marked “advertisement” in accordance with 18 USC section 1734.

ACKNOWLEDGMENTS

This work was supported, in part, by the University of Wisconsin Carbone Cancer Center, the Department of Defense (W81XWH-11-1-0644), the National Center for Advancing Translational Sciences (9U54TR000021), and the Elsa U. Pardee Foundation. No other potential conflict of interest relevant to this article was reported.

REFERENCES