



SPECT/CT Imaging, Biodistribution and Radiation Dosimetry of a ^{177}Lu -DOTA-Integrin $\alpha\nu\beta 6$ Cystine Knot Peptide in a Pancreatic Cancer Xenograft Model

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Introduction: Pancreatic ductal adenocarcinoma (PDAC) is one of the most aggressive malignant neoplasms, as many cases go undetected until they reach an advanced stage. Integrin $\alpha\nu\beta 6$ is a cell surface receptor overexpressed in PDAC. Consequently, it may serve as a target for the development of probes for imaging diagnosis and radioligand therapy. Engineered cystine knottin peptides specific for integrin $\alpha\nu\beta 6$ have recently been developed showing high affinity and stability. This study aimed to evaluate an integrin $\alpha\nu\beta 6$ -specific knottin molecular probe containing the therapeutic radionuclide ^{177}Lu for targeting of PDAC.

Methods: The expression of integrin $\alpha\nu\beta 6$ in PDAC cell lines BxPC-3 and Capan-2 was analyzed using RT-qPCR and immunofluorescence. In vitro competition and saturation radioligand binding assays were performed to calculate the binding affinity of the DOTA-coupled tracer loaded with and without lutetium to BxPC-3 and Capan-2 cell lines as well as the maximum number of binding sites in these cell lines. To evaluate tracer accumulation in the tumor and organs, SPECT/CT, biodistribution and dosimetry projections were carried out using a Capan-2 xenograft tumor mouse model.

Results: RT-qPCR and immunofluorescence results showed high expression of integrin $\alpha\nu\beta 6$ in BxPC-3 and Capan-2 cells. A competition binding assay revealed high affinity of the tracer with IC_{50} values of 1.69 nM and 9.46 nM for BxPC-3 and Capan-2, respectively. SPECT/CT and biodistribution analysis of the conjugate ^{177}Lu -DOTA-integrin $\alpha\nu\beta 6$ knottin demonstrated accumulation in Capan-2 xenograft tumors ($3.13 \pm 0.63\% \text{IA/g}$ at day 1 post injection) with kidney uptake at $19.2 \pm 2.5\% \text{IA/g}$, declining much more rapidly than in tumors.

Conclusion: ¹⁷⁷Lu-DOTA-integrin $\alpha\beta 6$ knottin was found to be a high-affinity tracer for PDAC tumors with considerable tumor accumulation and moderate, rapidly declining kidney uptake. These promising results warrant a preclinical treatment study to establish therapeutic efficacy.

Keywords: tumor targeting, integrin $\alpha\beta 6$, radioligand therapy, knottin peptide, pancreatic ductal adenocarcinoma, dosimetry

INTRODUCTION

Pancreatic ductal adenocarcinoma (PDAC) is one of the most aggressive malignant neoplasms and accounts for 80-90% of all pancreatic cancer cases. Although the incidence of PDAC is low, in cancer-related deaths, it ranks seventh globally (1). Owing to its poor prognosis, the 5-year survival rate is 9% with merely 24% of patients surviving for a year (2). This is mainly because PDAC patients rarely exhibit symptoms before an advanced stage of the disease has been reached, and due to the lack of appropriate diagnostic and therapeutic options. Surgical resection of the tumor, part of the pancreas, and other nearby digestive tract organs remains to be the only curative treatment for early-stage PDAC patients. Gemcitabine has been used for several years as a baseline chemotherapeutic treatment. Lately, combination therapy of gemcitabine with folirinox and nab-paclitaxel demonstrated improved results in comparison to the use of gemcitabine alone (3, 4). However, high resistance of PDAC to chemotherapy dilutes its efficacy (5, 6).

Integrins are heterodimeric transmembrane cell surface proteins that mediate cell-to-cell and cell-to-extracellular matrix (ECM) adhesion (7, 8). Many integrins, including integrin $\alpha\beta 6$, were reported to be upregulated in various cancers such as breast cancer, gastric cancer, colorectal cancer, lung cancer, ovarian cancer and PDAC (9–14). In well-differentiated PDACs, integrin $\alpha\beta 6$ overexpression was identified in 100% of the samples (13, 14). Moreover, integrin $\alpha\beta 6$ overexpression has been recognized as a prognostic marker for reduced survival in non-small cell lung cancer (15), gastric carcinoma (16), colorectal cancer (17) cervical squamous cell carcinoma (18) and PDAC (19). Remarkably, integrin $\alpha\beta 6$ expression was found to be higher in PDAC than in chronic pancreatitis (20). These findings support the utilization of integrin $\alpha\beta 6$ as a target for the development of new diagnostic and therapeutic tools.

Cystine knot peptides (knottins) represent small peptides of approximately 4 kDa with three threaded disulfide bonds that form a rigid topological knot constraining the peptide's

conformational flexibility. Such a structural motif is known as a cystine knot (21, 22). One of the advantages of knottins is the high variability of backbone residues that may be used to modulate tumor and kidney uptake (23). Knottins are well suited for *in vivo* tumor-targeting applications as their disulfide-bonded core confers outstanding thermal stability and resistance against proteolytic degradation; moreover, they have been shown to be nonimmunogenic. Furthermore, knottins may be chemically modified to tailor their *in vivo* pharmacokinetic properties for a variety of clinical applications. Previously, we have developed the optimized knottin R₀I-MG that shows low single-digit nanomolar binding affinity for integrin $\alpha\beta 6$ (23, 24). Recently, the first clinical study with R₀I-MG demonstrated clinical potential for targeting PDAC in humans (24).

In the current imaging, biodistribution and dosimetry study, a lutetium-177 DOTA conjugate of this knottin was evaluated as a candidate for therapeutic purposes in a PDAC xenograft model. Binding affinity of the ¹⁷⁷Lu tracer was found to be in the low single-digit nanomolar range. SPECT/CT imaging and biodistribution of the tracer ¹⁷⁷Lu-DOTA-integrin $\alpha\beta 6$ knottin revealed substantial accumulation of the tracer in the tumor as well as faster renal clearance. Our studies demonstrate the potential of the ¹⁷⁷Lu-DOTA integrin $\alpha\beta 6$ knottin as a therapeutics candidate in PDAC.

MATERIAL AND METHODS

Cell Culture

The human pancreatic adenocarcinoma cell lines BxPC-3 and Capan-2 were obtained from ATCC/LGC Standards (Wesel, Germany). They were cultured in RPMI 1640 and McCoy's 5A (Modified) Medium (both Biochrom, Berlin, Germany), respectively, and supplemented with 10% fetal calf serum (Biochrom, Berlin, Germany) for cultivation in a humidified atmosphere at 37°C with 5% CO₂.

Reverse Transcription Quantitative Real-Time PCR (RT-qPCR)

Total RNA was isolated from cell lines or kidney and liver of mouse using the RNeasy Mini Kit (Qiagen, Hilden, Germany) according to the manufacturer's protocol. RNA was treated with DNase I (Sigma Aldrich, Munich, Germany) before reverse transcription with the High Capacity cDNA Reverse Transcription Kit (Applied Biosystems, Waltham, MA, USA). Quantitative real-time PCR was performed with Blue S'Green qPCR 2X mix (Biozym Scientific GmbH, Oldendorf, Germany),

Abbreviations: % IA/g, percent injected activity per gram tissue; ¹⁷⁷Lu, lutetium-177; BERIC, Berlin Experimental Radionuclide Imaging Center; BSA, bovine serum albumin; CPM, counts per minute; DMSO, dimethyl sulfoxide; DOTA, 1,4,7,10-tetraazacyclododecane-1,4,7,10-tetraacetic acid; ECM, extracellular matrix; HPLC, high-performance liquid chromatography; IC₅₀, half maximal inhibitory concentration; n.d., not determined; NMRI, Naval Medical Research Institute; OATP, organic anion transporter protein; p.i., post injection; PBS, phosphate-buffered saline; PDAC, pancreatic ductal adenocarcinoma; PFA, paraformaldehyde; PET, positron emission tomography; RPMI1640, Roswell Park Memorial Institute medium 1640; SD, standard deviation; SEM, standard error of mean; TFA, trifluoroacetic acid.

0.5 μM primer and 20 ng cDNA in a total reaction volume of 10 μL on a Bio-Rad CFX96 Real-Time-System. PCR settings were 95°C for 30 sec, followed by 45 cycles of 95°C for 5 sec and 61°C for 30 sec. All primers were designed using NCBI Primer-BLAST software. They were manufactured by Tib-MolBiol (Berlin, Germany) and their sequences are indicated below. Plotted values were normalized on the geometric mean of UBC and GAPDH (for cells and xenografts) or GAPDH only (for mouse kidney and liver) using the $\Delta\Delta\text{C}_t$ method.

hITGB6-Fwd: 5'-ACTGCCTGCTTATTGGACCTC-3'

hITGB6-Rev: 5'-ATCACACCTTTCGCCAACTC-3'

hUBC-Fwd: 5'-ATTTGGGTCGCAGTTCTTG-3'

hUBC-Rev: 5'-TGCCTTGACATTCTCGATGGT-3'

hGAPDH-Fwd: 5'-TGCACCACCAACTGCTTAGC-3'

hGAPDH-Rev: 5'-GGCATGGACTGTGGTCATGAG-3'

mITGB6-Fwd: 5'-CTCACgggTACAgTAACgCAT-3'

mITGB6-Rev: 5'-AATgAgCTCTCAggCaggCT-3'

Immunofluorescence

For immunofluorescent staining of tumor tissue, cells grown on glass coverslips were fixed with 1:1 methanol/acetone for two minutes and air-dried. After washing with PBS (Biochrom, Berlin, Germany) and blocking with 5% goat serum in PBS for 30 minutes, coverslips were incubated with a rabbit IgG against human integrin $\beta 6$ (#HPA023626, Atlas Antibodies, Bromma, Sweden) diluted in 0.1% BSA in PBS in a wet chamber for one hour at room temperature. After washing, coverslips were incubated with the secondary antibody goat-anti-rabbit-Cy3 (Jackson ImmunoResearch, West Grove, USA; 2.5 $\mu\text{g}/\text{mL}$ diluted in 0.1% BSA in PBS) for 30 minutes. After washing with PBS, nuclei were stained with 1 $\mu\text{g}/\text{mL}$ TOTO-3 (Invitrogen) in PBS for 5 minutes. Finally, the cells were fixed with 96% ethanol for two minutes, embedded with Immu-Mount (Thermo Fisher Scientific, Waltham, USA) and analyzed using a confocal laser-scanning microscope (LSM510, Carl Zeiss, Jena, Germany).

Radioiodination of the DOTA Integrin $\alpha\beta 6$ Knottin

The previously described cystine knot peptide specific for integrin $\alpha\beta 6$ R₀1-MG (DOTA-GCILNGRTDLGTLFR-CRRSDCPGACICRGNGYCG, DOTA coupled to the aminoterminal glycine *via* N-hydroxysuccinimide) was utilized in this study. The DOTA-integrin $\alpha\beta 6$ knottin was labelled with ¹²⁵I using the chloramine-T method. Chloramine-T and sodium metabisulfite solutions were freshly prepared in water. For labelling, 10 μL of DOTA-integrin $\alpha\beta 6$ knottin (stock 1 mM) with 15 μL of sodium phosphate buffer (0.5 M, pH 7.6) was mixed with 37 MBq carrier-free [¹²⁵I]NaI (NEZ033L010MC, Perkin Elmer, Waltham, US). 4 μL chloramine T (1 mg/ml) were added to start the reaction and after 30 seconds, 4 μL sodium metabisulfite (2 mg/ml) were added to stop the iodination. Labelled radioactive peptide was separated from unlabeled peptide by HPLC purification (Agilent ZORBAX 300 Extend-

C18, 46 x 150 mm, 5 μm column) using a gradient from 20-50% acetonitrile (+0.1% TFA) against water (+0.1% TFA) for 20 min. To determine the retention time of the radioactive peptide, 1-2 μL of the reaction mixture was analyzed before the purification run. The fraction containing the radiolabeled peptide peak (purity after purification according to HPLC: 97%) was then collected, diluted with binding buffer (50 mM HEPES pH 7.4, 5 mM MgCl_2 , 1 mM CaCl_2 , 0.5% BSA, cOmplete™ protease inhibitors [Roche, Mannheim, Germany]) to prevent radioautolysis, aliquoted and stored at -80°C.

Non-Radioactive Metalation With ^{nat}Lu

Non-radioactive complexes of DOTA-integrin $\alpha\beta 6$ knottin with natural lutetium were generated by incubation of 0.9 nmol (final concentration of 30 μM) peptide conjugate dissolved in 5 μL buffer (sodium acetate/acetic acid buffer, 0.5 M, pH 5.4) with 27 nmol (a 30-fold molar excess) of the Lu³⁺ ion (^{nat}Lu-DOTA-knottin) or no metal ion (control DOTA-knottin). The reaction volume was made up to 30 μL with water. The reaction was carried out for 10 minutes at 80°C. HPLC showed >98% purity of the ^{nat}Lu-labeled peptide.

Saturation Binding Assay

To determine the dissociation constant (K_d) and the maximum number of binding sites (B_{max}) of the radioligand, saturation binding assay was performed on BxPC-3 and Capan-2 cell lines. For this, approximately 50,000 cells per well were seeded in a 96 well plate and incubated overnight at 37°C. Cells were then incubated with binding buffer and varying concentration (0, 0.01, 0.05, 0.1, 0.2, 0.5, 1, 2, 5, 10, 15, 20 nM) of ¹²⁵I-DOTA-integrin $\alpha\beta 6$ knottin either with (non-specific binding) or without 1 μM of DOTA-integrin $\alpha\beta 6$ knottin (total binding). Non-specific binding was subtracted from total binding to obtain specific binding. All three datasets (non-specific binding, specific binding and total binding) were plotted with GraphPad Prism and fitted using nonlinear regression (one site - total and non-specific, one site - specific binding). The software provides B_{max} in the same value as the respective y-axis, in this case cpm. The following calculations were performed to obtain B_{max} in receptor sites per cell. In the first step, the specific activity of ¹²⁵I (2175 Ci/mmol) was transformed into dpm/mmol by multiplication with 2.22×10^{12} . This was multiplied with the counter efficiency of 50% to get cpm/mmol and subsequently converted to cpm/fmol:

$$\text{cpm/fmol} = [2175 \text{ Ci/mmol} * 2.22 * 10^{12} * 0.65] / 10^{12}$$

In a second step, the B_{max} value, calculated by the software in cpm, was divided by cpm/fmol to derive the amount of substance in fmol. This was multiplied with the Avogadro constant (6.02×10^8 /fmol) to obtain the number of molecules before division by seeded cells (50,000 in this case):

$$\text{sites/cell} = [B_{\text{max}}(\text{cpm}) * 6.02 * 10^8 \text{ fmol}^{-1}] / \text{cpm/fmol} * 50,000$$

Competition Binding Assay

For competitive radioligand binding assay, approximately 50,000 cells (BxPC-3 or Capan-2) per well were seeded in a 96 well plate

incubated overnight at 37°C. Next day, cells were incubated in binding buffer containing 100,000 cpm of ¹²⁵I-DOTA-integrin α v β 6 knottin and increasing concentrations (0, 0.00001, 0.00003, 0.0001, 0.0003, 0.001, 0.003, 0.01, 0.03, 0.1, 0.3 and 1 μ M) of unlabeled peptide (DOTA-knottin). After 2 hours of incubation, cells were washed for 2-3 times with ice-cold washing buffer (50 mM Tris-HCl, pH 7.4, 125 mM NaCl, 0.05% BSA) and lysed with 1 N NaOH (80 μ l/well). The lysed cells were transferred to vials and measured in a gamma counter (Wallac 1470 Wizard, Perkin Elmer, Waltham, MA, USA). Obtained cpm values were plotted with GraphPad Prism and the data fitted using nonlinear regression (one site - fit Ki, one site - fit logIC₅₀).

Cell Irradiation

Cells were irradiated using a GSR D1 gamma irradiator (Gamma-Service Medical GmbH, Leipzig, Germany) with a cesium-137 source. The cells with the respective culture medium were placed inside the radiation chamber and required dose was achieved by adjusting the tray level and duration of the exposure. After irradiation, the cell lines were incubated at 37°C without medium change for the appropriate duration.

Cell Count (DAPI Staining)

Cells were seeded at a density of 5,000 cells per well in a 96 well plate and incubated overnight at 37°C. After irradiating with 0, 0.2, 2, 4, 6, 8, 10, 15 or 30 Gy, cells were incubated at 37°C for 96 hours. Cells were then washed with PBS and fixed with 4% PFA for 10 min at room temperature. Thereafter, cells were stained with DAPI (1:5,000 in PBS/0.1% Tx-100) and incubated at room temperature for 10 min. Before image acquisition, wells were filled with 80 μ L of PBS and images were acquired using an automated microscope (IN Cell Analyzer 1000; GE, Reading, UK) with a 4x objective. Image stacks were analyzed and nuclei were counted by the IN Cell software.

Clonogenic Assay

Cells were seeded at a density of 5,000 cells per well in a 12 well plate and incubated overnight at 37°C. After irradiation with 0, 2, 4, 6 or 8 Gy, cells were incubated at 37°C without medium change for 7 days. Cells were then washed with PBS and fixed with 70% ethanol for 10 min, stained with 0.2% crystal violet solution for another 10 min and carefully rinsed with tap water. Plates were dried overnight and digitized using an Odyssey infrared scanner (700 nm channel, intensity 3, 84 μ m resolution and medium quality). For quantification, images were analyzed using the Colony Area plugin for ImageJ.

Xenografts

For *in vivo* experiments, at least 8-week-old female athymic NMRI-Foxn1^{nu}/Foxn1^{nu} mice (Janvier Labs, Saint-Berthevin, France) were used. Animal care followed institutional guidelines and all experiments were approved by local animal research authorities. For the generation of tumor xenografts, 5 x 10⁶ cells of Capan-2 cells were inoculated subcutaneously into the left and right shoulder (1:1 phosphate-buffered saline [PBS]/

Matrigel Basement Membrane Matrix High Concentration, Corning, Corning, USA). Tumors were allowed to grow for two to four weeks (tumor volume > 100 mm³) after cell inoculation.

Radiochemical Labeling With Lutetium-177 (¹⁷⁷Lu)

Radiolabeling of the DOTA-integrin α v β 6 knottin was carried out manually using [¹⁷⁷Lu]LuCl₃ and a reagent kit from ITG Isotope Technologies Garching GmbH (Garching, Germany). A total of 40 μ L (stock solution of peptide at 1 μ g/ μ L in 10% DMSO in water) was added to 500 μ L of ascorbate buffer solution pH 5.0, prepared from the kit (composition not disclosed by vendor) and the resulting mixture was then added to 35 μ L of [¹⁷⁷Lu]LuCl₃ (2 GBq in aqueous 0.04 M HCl) and the reaction mixture was heated at 80 °C for 35 minutes followed by cooling for 10 minutes at room temperature. The product was then diluted with 0.5 mL of saline and the pH was adjusted to 7 using a 1 M aqueous solution of NaHCO₃ prior to injection in mice. A reverse-phase HPLC system (Knauer GmbH, Germany) with a column (Eurospher II, C18, 250 x 4 mm) was used to determine the radiochemical purity of the radiotracer. The HPLC system was equipped with an Azura P.6.1L pump coupled with ultraviolet (Azura UVD 2.1L) and radiometric (γ -Raytest-Isotopenmessgeraete GmbH, Germany) detectors. The gradient elution system used mobile phase A (100% acetonitrile) and mobile phase B (deionized H₂O containing 0.1% trifluoroacetic acid) and a flow rate of 1.0 mL/min. Starting with 0% A and 100% B, the gradient was increased to 100% A over 25 min and finally returned to initial gradient conditions within 5 min. The retention time of the tracer was found to be 10.37 min and the peak of free Lu-177 was observed at 2.64 min. Radiochemical purity determined by HPLC was found to be higher than 97% (**Supplementary Figure S1**) in all cases and the product had a molar activity of approximately 44 GBq/ μ mol.

SPECT/CT Imaging

SPECT and CT imaging were performed using the nanoSPECT/CTplus scanner (Mediso, Hungary/Bioscan, France). Mice were anesthetized using 1-2% isoflurane with oxygen at a flow rate of approximately 0.5 l/min. After a low-dose CT scan for positioning the tumors and kidneys in the scan range the SPECT acquisition was started directly before intravenous injection of approximately 50 MBq of ¹⁷⁷Lu-DOTA-integrin α v β 6 knottin (0.1-0.15 ml).

Nine consecutive multi-pinhole SPECT images of 10 min duration each (5 angular steps a 60 sec, 2 bed positions) were acquired. Additional individual scans of 35-60 min duration (5 angular steps a 60-180 sec, 2 bed positions) were performed up to 8 days to assess biodistribution kinetics.

Biodistribution Studies

Tumor-bearing mice (n=4-5 per time point) were injected with approximately 1 MBq (22 pmol) of ¹⁷⁷Lu-DOTA-integrin α v β 6 knottin to the tail vein *via* a catheter. Mice (n=4-5 per time point) were sacrificed and dissected one, two,

three and eight days post injection. Tumor, blood, stomach, pancreas, small intestine, colon, liver, spleen, kidney, heart, lung, muscle and femur samples were weighed and uptake of radioactivity was measured by a gamma counter.

Dosimetry of Tissue and Tumor

Assuming that the organ-to-whole-body activity concentration ratio in mice would equal that in humans, the injected activity concentration (%IA/g) acquired from the mouse biodistribution study was transformed to human whole-organ percentage of injected activity per gram of tissue (%IA/g)_{human}. The mouse uptake data were extrapolated to humans by relative scaling of mass and time using the following two equations,

$$(\% IA/organ)_{human} = [(\% IA/g)_{mouse} \times M_{mouse}(kg) \times [m(g)/M(kg)]_{human}] \quad (1)$$

$$t_{human} = t_{animal} [M_{human}/M_{animal}]^{0.25} \quad (2)$$

where %IA/g is percentage of injected activity per gram of tissue, %IA/organ is percentage of injected activity per organ; m is organ mass of mouse or human, M is body mass of mouse or human; t is time. The absorbed doses per activity were calculated by using the extrapolated human source organ residence times as input in the OLINDA/EXM 1.1 software (Organ Level Internal Dose Assessment Code, Vanderbilt University, Nashville, USA), with the reference adult male (25).

Statistical Analysis and Data Availability

All statistical analyses were performed using GraphPad Prism 5.04. IC₅₀ values were determined by nonlinear sigmoidal curve fitting with variable slope setting. All presented data are based on independent experiments. Numerical data for this study have been deposited in an open data repository for public access: <http://doi.org/10.5281/zenodo.4362503>.

RESULTS

Target Expression and Tracer Affinity

As integrin $\beta 6$ forms heterodimers only with integrin αv , detection of the $\beta 6$ subunit mRNA or protein will be informative about the dimer, too. To identify the optimal animal model for imaging, integrin $\beta 6$ mRNA was measured by RT-qPCR in human PDAC cell lines, corresponding xenografts, A549 lung adenocarcinoma and HT29 colorectal adenocarcinoma cells as well as mouse kidney and liver. Capan-2 and BxPC-3 cells and xenografts showed the highest abundance of integrin $\beta 6$ mRNA (**Figure 1A**). Mouse kidney and liver showed around 1,000-fold less integrin $\beta 6$ mRNA, indicating a significantly lower expression of the corresponding mouse receptor in these organs. Immunofluorescence staining detected integrin $\beta 6$ on the plasma membrane of Capan-2 cells (**Figure 1B**).

To determine the binding affinity of the DOTA-integrin $\alpha\beta 6$ knottin to its target, saturation and competition

binding assays on BxPC-3 and Capan-2 cells were performed (**Figures 1C, D** and **Table 1**). Saturation binding assays were performed to determine dissociation constant (K_d) and maximum number of binding sites per cell (B_{max}). K_d values for BxPC-3 and Capan-2 cells were found to be 0.30 nM and 0.75 nM, respectively. Correspondingly, the B_{max} values for BxPC-3 and Capan-2 were shown to be approximately 11,800 and 11,500 binding sites/cell. ¹²⁵I-labeled DOTA-integrin $\alpha\beta 6$ knottin showed binding to both cell lines, BxPC-3 and Capan-2, displaced by the unlabeled peptide in a concentration-dependent manner. The inhibitory constant (K_i) values of DOTA-integrin $\alpha\beta 6$ knottin for BxPC-3 and Capan-2 were calculated to be 1.69 nM and 9.46 nM respectively. To determine the effect of lutetium chelation on the probe's affinity for integrin $\alpha\beta 6$, binding of ^{nat}Lu-DOTA-knottin and control-DOTA-knottin (without ^{nat}Lu) on BxPC-3 and Capan-2 was examined. The complexation of the ^{nat}Lu ion did not compromise the high affinity of the tracer (**Table 1**).

Radiosensitivity of BxPC-3 and Capan-2 Cells

To evaluate the suitability of BxPC-3 and Capan-2 cells as model cell lines in terms of their radiosensitivity, two approaches were taken. Both cell lines were irradiated with different doses (0, 0.05, 0.2, 2, 4, 6, 8, 10, 15 and 30 Gy) of gamma radiation from a ¹³⁷Cs source. After 96 hours of incubation, nuclei were stained and counted. The IC₅₀ values for the resulting growth inhibition/cell death in BxPC-3 and Capan-2 cells were found to be 4.3 Gy and 5.5 Gy, respectively (**Figures 2A, B**). To evaluate radiation effects on colony formation, both cell lines were treated with different doses (0, 2, 4, 6 and 8 Gy) of radiation. After 7-8 days of incubation, colonies were fixed, stained and counted (**Figures 2C, D**). BxPC-3 cells (IC₅₀ 1.3 Gy) appeared to be slightly more radiation-sensitive compared to Capan-2 (IC₅₀ 2.2 Gy) in this assay.

SPECT Imaging in a Capan-2 Xenograft Mouse Model of Pancreatic Cancer

While the integrin $\alpha\beta 6$ knottin and its DOTA conjugate previously had been used for tumor imaging studies employing ¹⁸F, ⁶⁴Cu or ^{99m}Tc, no data regarding the biodistribution of the ¹⁷⁷Lu-DOTA-integrin $\alpha\beta 6$ knottin were available. To obtain such data, ¹⁷⁷Lu-DOTA-integrin $\alpha\beta 6$ knottin was synthesized, yielding a product with 98% radiochemical purity and a molar activity of approximately 44 GBq/ μ mol. The tracer was injected intravenously in mice bearing Capan-2 xenografts on both shoulders. SPECT/CT images were taken at different times post injection. **Figure 3A** shows SPECT/CT images taken at 22 hours p.i. The maximum intensity projection (MIP) reveals an accumulation of ¹⁷⁷Lu-DOTA-integrin $\alpha\beta 6$ knottin in both the tumors and the kidneys. Uptake by other organs was low or moderate. Short-term and long-term tracer kinetics for kidney and tumor were quantified from SPECT images, up to 3 and 187 hours post-injection, respectively. Indeed, kidneys showed a higher initial accumulation of ¹⁷⁷Lu-DOTA-integrin $\alpha\beta 6$

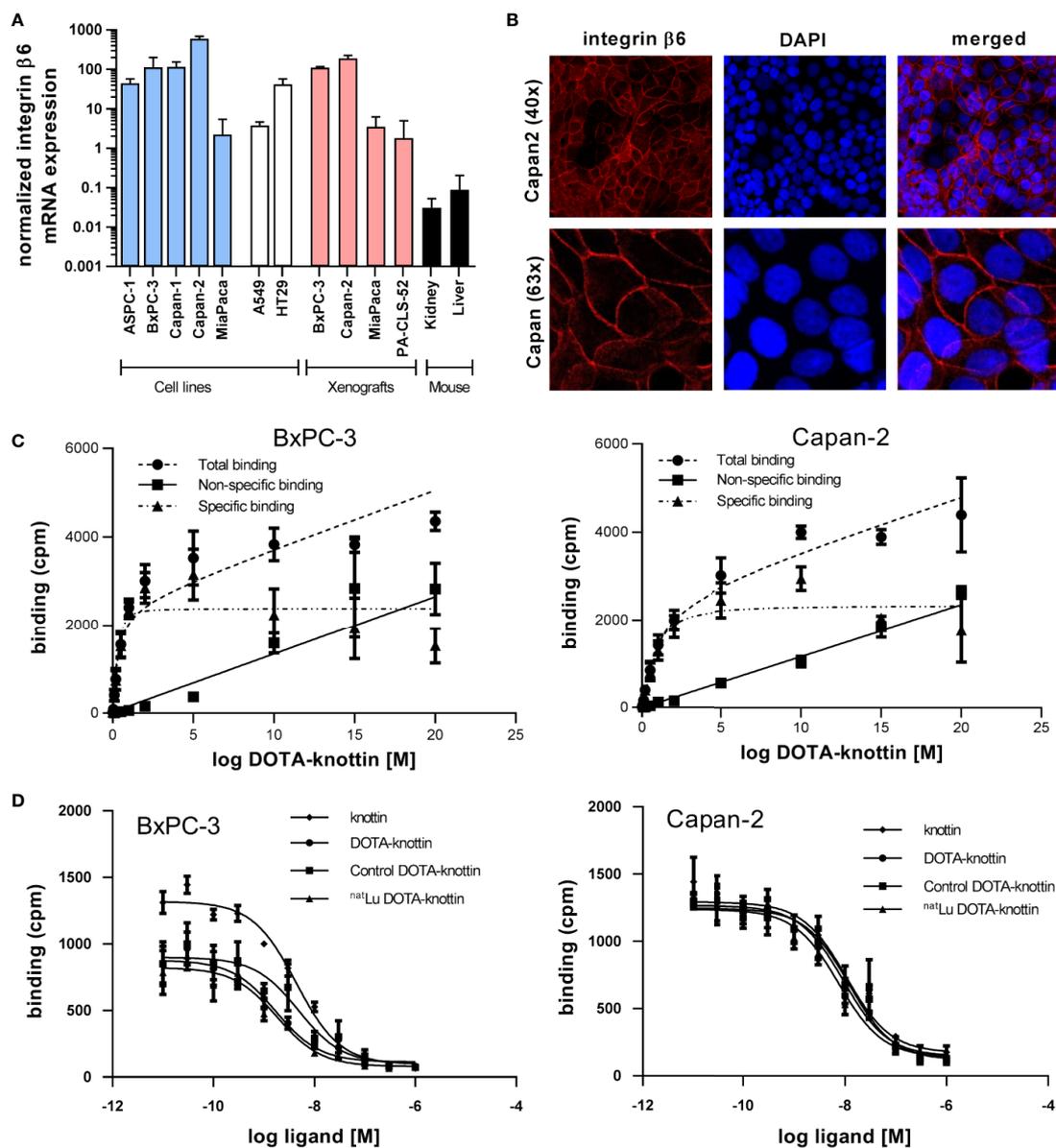


FIGURE 1 | Integrin $\beta 6$ mRNA, protein localization, binding sites and tracer affinity. **(A)** Integrin $\beta 6$ mRNA was measured by RT-qPCR in human cell lines of different origin, xenografts and mouse kidney and liver. Graphs represent normalized $\beta 6$ mRNA expression. **(B)** Fluorescence images show staining of integrin $\beta 6$ (antibody, red) and nucleus (TOTO-3, blue) in Capan-2 cells. **(C)** Dissociation constant (K_d) and number of binding sites per cell were determined in BxPC-3 and Capan-2 cells using a saturation binding assay with ¹²⁵I-DOTA-integrin $\alpha\beta 6$ knottin. **(D)** A competition binding assay using ¹²⁵I-DOTA-integrin $\alpha\beta 6$ knottin was performed to derive the inhibitory constant (K_i) of DOTA-knottin in both cell lines. The binding curves show tracer displacement by increasing concentrations of unlabeled knottin, DOTA-knottin (DOTA-integrin $\alpha\beta 6$ knottin), control-DOTA-knottin and ^{nat}Lu-DOTA-knottin. For the latter two, DOTA-integrin $\alpha\beta 6$ knottin was either complexed with natural lutetium (^{nat}Lu-DOTA-knottin) or treated the same but without the metal ion (control-DOTA-knottin). Data represent mean \pm S.E.M. ($3 \leq n \leq 4$).

TABLE 1 | Saturation and competition binding assay results using ¹²⁵I-DOTA-integrin $\alpha\beta 6$ knottin for BxPC-3 and Capan-2 cells.

	Kd (nM)	Bmax (Sites/cell)	Ki (nM)			
			knottin	DOTA knottin	control DOTA knottin	^{nat} Lu DOTA knottin
BxPC-3	0.30 \pm 0.08	11,874 \pm 897	4.36 \pm 0.16	1.69 \pm 1.3	4.85 \pm 1.94	1.82 \pm 1.39
Capan-2	0.75 \pm 0.19	11,545 \pm 887	10.79 \pm 4.03	9.46 \pm 5.21	12.06 \pm 2.87	7.54 \pm 1.27

Data are presented as mean \pm S.E.M. ($3 \leq n \leq 4$).

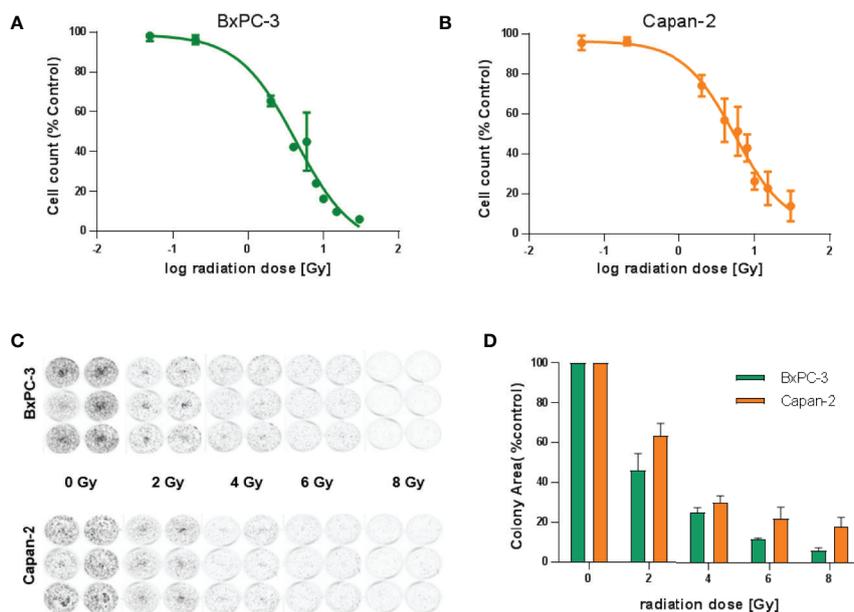


FIGURE 2 | Radiosensitivity of BxPC-3 and Capan-2 cells. **(A)** BxPC-3 and Capan-2 **(B)** cells were exposed to different doses of radiation (0, 0.05, 0.2, 2, 4, 6, 8, 10, 15 and 30 Gy). 96 hours later, nuclei were stained and counted. **(C, D)** BxPC-3 and Capan-2 cells were irradiated with 0, 2, 4, 6 or 8 Gy. Cells were incubated for 1 week. Colonies were stained and colony area was measured. Data represent mean \pm S.E.M. ($n = 5$).

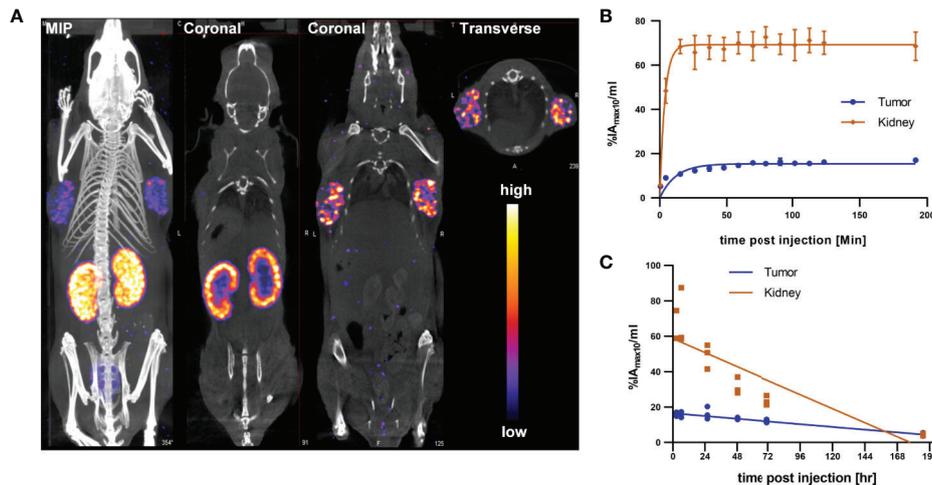


FIGURE 3 | SPECT/CT imaging with ^{177}Lu -DOTA-integrin $\alpha\beta 6$ knottin. **(A)** SPECT images show maximum intensity projection (MIP), coronal and transverse projections fused with CT images of ^{177}Lu -DOTA-integrin $\alpha\beta 6$ knottin in a Capan-2 xenograft model at 22 hours post injection of 62 MBq tracer. Nude mice were carrying xenografts on left and right shoulder. **(B)** Early SPECT kinetics data show the uptake of ^{177}Lu -DOTA-integrin $\alpha\beta 6$ knottin in tumor and kidney. **(C)** SPECT-based time-activity curve (2-187 hours p.i.) shows faster clearance of ^{177}Lu -DOTA-integrin $\alpha\beta 6$ knottin from kidney than from tumor.

knottin (**Figure 3B**), yet clearance was also faster than from the tumor (**Figure 3C**).

Biodistribution Analysis of Organ Uptake

Ex vivo biodistribution analysis of mice bearing Capan-2 xenografts demonstrated a tumor uptake of the ^{177}Lu -DOTA-

integrin $\alpha\beta 6$ knottin of 3.1 ± 0.6 , 2.5 ± 0.4 , 3.5 ± 0.9 and $1.2 \pm 0.2\%$ IA/g (mean \pm S.E.M.) on day 1, 2, 3 and 8, respectively (**Figure 4A**). Nevertheless, tracer uptake by the kidney of 19.2 ± 2.5 , 12.5 ± 0.6 , 14.7 ± 4.5 and $2.3 \pm 0.4\%$ IA/g was detected on days 1, 2, 3 and 8 p.i., respectively. On the other hand, low or moderate activity was discovered in organs like stomach, colon

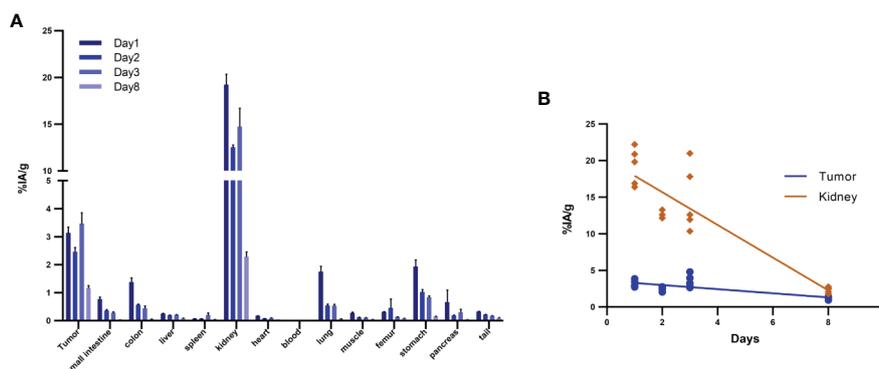


FIGURE 4 | Biodistribution of ¹⁷⁷Lu-DOTA integrin $\alpha\beta 6$ knottin in Capan-2 xenograft-bearing mouse. **(A)** Quantitative analysis of organ activity at day 1, 2, 3 and 8 p.i. **(B)** Time-activity curve of ¹⁷⁷Lu-DOTA integrin $\alpha\beta 6$ knottin biodistribution data for tumor and kidney from days 1, 2, 3 and 8 p.i. Data represent mean \pm S.E.M. ($4 \leq n \leq 5$).

and lung at later time points (Table 2). The time-activity curve from *ex vivo* biodistribution data (Figure 4B) confirms the analysis of SPECT kinetics (Figure 3C) with activity from kidneys being washed out more rapidly than from tumors. Based on *ex vivo* biodistribution inputs, tumor-to-organ ratios were calculated (Table 2 and Figure 5). Except for kidney, these ratios were favorable in pancreas, liver, blood, lung and muscle. Ex-vivo analysis of H&E-stained xenografts did not reveal significant differences in tissue structure between samples obtained on day 1 or day 8 after injection of the tracer (Supplementary Figure S2).

Dosimetric Calculation

Dosimetric calculations for the human male adult for ¹⁷⁷Lu were generated (Table 3). The expected absorbed doses per injected

activity in humans were calculated using the mouse biodistribution data. The two interspecies scaling (mass and time) model (25, 26) was implemented for extrapolating animal to human data. The calculated expected effective dose was 0.04 mSv/MBq. Additionally, the calculated absorbed doses for the common organs are indicated. As expected from the mouse biodistribution data, kidney showed the highest absorbed dose of 0.02 mSv/MBq followed by lung and stomach of 0.01 and 0.005 mSv/MBq respectively (Table 3).

DISCUSSION

Due to its high and selective tumor overexpression, integrin $\alpha\beta 6$ is emerging as a target in cancer for nuclear imaging.

TABLE 2 | Biodistribution (%IA/g, mean \pm SD) and tumor-to-organ ratios of ¹⁷⁷Lu-DOTA-integrin $\alpha\beta 6$ knottin in nude mice bearing Capan-2 xenografts.

	24 h		48 h		72 h		192 h	
tumor	3.13	± 0.63	2.45	± 0.39	3.46	± 0.85	1.16	± 0.21
small intestine	0.76	± 0.17	0.35	± 0.08	0.27	± 0.10	0.02	± 0.00
colon	1.37	± 0.36	0.56	± 0.06	0.43	± 0.21	0.04	± 0.01
liver	0.24	± 0.03	0.18	± 0.04	0.20	± 0.04	0.08	± 0.02
spleen	0.06	± 0.01	0.07	± 0.01	0.18	± 0.21	0.04	± 0.01
kidney	19.21	± 2.53	12.50	± 0.55	14.71	± 4.48	2.28	± 0.40
heart	0.16	± 0.03	0.07	± 0.01	0.08	± 0.04	0.04	± 0.01
blood	0.01	± 0.00	0.00	± 0.00	0.01	± 0.00	0.00	± 0.00
lung	1.74	± 0.46	0.53	± 0.13	0.52	± 0.13	0.06	± 0.01
muscle	0.26	± 0.09	0.11	± 0.02	0.10	± 0.02	0.04	± 0.01
femur	0.30	± 0.04	0.45	± 0.65	0.13	± 0.03	0.07	± 0.02
stomach	1.92	± 0.55	1.01	± 0.19	0.83	± 0.13	0.14	± 0.03
pancreas	0.65	± 0.98	0.17	± 0.05	0.29	± 0.26	0.03	± 0.01
tail	0.31	± 0.05	0.20	± 0.05	0.16	± 0.03	0.08	± 0.05
tumor-to-blood	497.70	± 79.30	851.38	± 266.49	697.02	± 248.45	2123.36	± 553.90
tumor-to-kidney	0.17	± 0.05	0.19	± 0.04	0.24	± 0.02	0.51	± 0.07
tumor-to-muscle	13.13	± 5.06	23.15	± 4.30	35.80	± 3.35	32.86	± 13.74
tumor-to-liver	12.99	± 1.72	13.58	± 2.66	18.35	± 6.29	14.64	± 1.79
tumor-to-pancreas	13.07	± 7.99	15.56	± 7.69	26.96	± 22.55	38.69	± 10.34

Data are presented as mean \pm SD %IA/g of tissue ($4 \leq n \leq 5$).

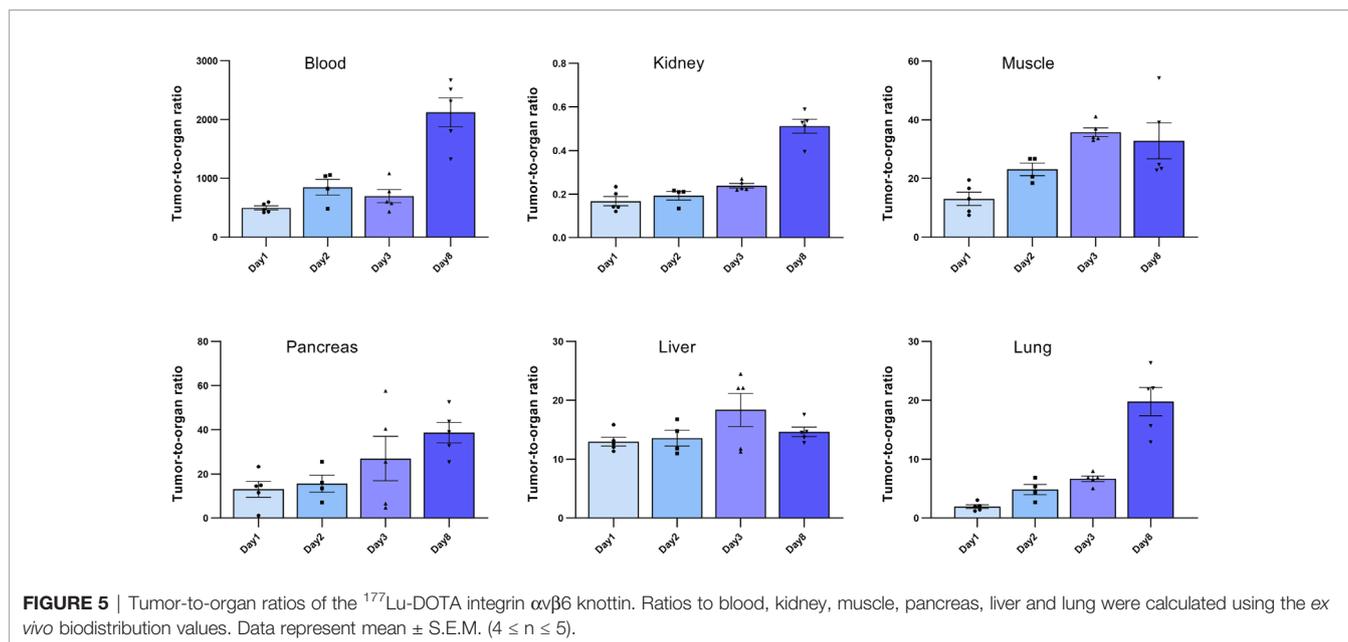


TABLE 3 | Expected absorbed doses of ¹⁷⁷Lu-DOTA integrin $\alpha\beta6$ knottin in humans.

Target Organ	mSv/MBq	rem/mCi
Adrenals	6.78E-06	2.51E-05
Brain	3.42E-08	1.26E-07
Breasts	2.36E-05	8.71E-05
Gallbladder Wall	0.00E+00	0.00E+00
LLI Wall	7.21E-05	2.67E-04
Small Intestine	1.82E-04	6.74E-04
Stomach Wall	5.15E-03	1.90E-02
ULI Wall	4.30E-06	1.59E-05
Heart Wall	0.00E+00	0.00E+00
Kidneys	2.06E-02	7.62E-02
Liver	9.34E-04	3.46E-03
Lungs	1.58E-02	5.85E-02
Muscle	1.40E-06	5.17E-06
Ovaries	1.76E-04	6.51E-04
Pancreas	5.59E-06	2.07E-05
Red Marrow	1.00E-04	3.71E-04
Osteogenic Cells	8.15E-06	3.01E-05
Skin	2.22E-06	8.20E-06
Spleen	2.35E-05	8.69E-05
Testes	0.00E+00	0.00E+00
Thymus	1.37E-06	5.08E-06
Thyroid	7.36E-06	2.72E-05
Urinary Bl Wall	1.09E-05	4.02E-05
Uterus	1.99E-06	7.36E-06
Effective dose	4.31E-02	1.60E-01

Several tumor-targeting strategies based on integrin $\alpha\beta6$ have been developed for either diagnostic or therapeutic purposes (15, 27–31). To that end, ligands had been conjugated either with a radionuclide (¹⁸F, ⁶⁴Cu, ¹¹¹In or ¹⁷⁷Lu) or an anti-cancer drug (e.g. tesirine) (31). The majority of approaches involved the use of linear peptides derived from foot-and-mouth disease virus, which show a high affinity to the receptor. However, due to

reduced stability and specificity for the receptor, *in vivo* results were ambiguous. In a recent study, a cyclic radiotracer specific for integrin $\alpha\beta6$ (⁶⁸Ga-cycratide) was used in a pancreatic mouse model for PET imaging (32). The DOTA-integrin $\alpha\beta6$ knottin applied here has recently been used in a first-in-human clinical study and has demonstrated a high potential for PDAC targeting (24). In this study, the same DOTA-conjugated engineered cystine knot peptide (knottin) specific for integrin $\alpha\beta6$ was chosen. The knottin peptide's stability and affinity to the receptor were not compromised *in vivo* and it had previously exhibited favorable tumor uptake (21, 23). Previously, ⁶⁴Cu, ¹⁸F and ^{99m}Tc have been investigated in conjugation with this knottin. However, no data were available on the pharmacology of the tracer coupled to a therapeutic radionuclide such as ¹⁷⁷Lu. This study was designed to examine the *in vitro* pharmacology, biodistribution and dosimetry of a ¹⁷⁷Lu-labeled DOTA-knottin in a xenograft mouse model, to pave the way for a potential clinical translation of this compound. Likewise, fluorescence-labeled knottins specific for integrin $\alpha\beta6$ had shown promising preclinical results, which could be further translated for the early detection of PDAC in patients (33, 34). Such an optical imaging agent could also play a role in fluorescence-guided surgery for PDAC patients. Indeed, the specificity of the agent would assist in discerning PDAC from pancreatitis and normal pancreatic tissue.

In the current study, integrin $\alpha\beta6$ mRNA expression in BxPC-3 and Capan-2 was found to be highest among all tested PDAC and other cell lines. Additionally, sustained expression of integrin $\alpha\beta6$ in mouse xenografts of these cell lines was confirmed. More importantly, mRNA expression of integrin $\alpha\beta6$ in mouse liver and kidney was found to be very low compared to BxPC-3 and Capan-2 cell lines. This, however, does not rule out expression in specific anatomical substructures of these two organs of excretion. Immunofluorescence

experiments established the expression of integrin $\alpha\beta 6$ on the surface of Capan-2 cells. This is of relevance as RT-qPCR data will only be informative about mRNA levels, not protein. In addition, many cell membrane receptors occur in equilibrium of distribution between plasma membrane and intracellular compartments, e.g. the trans-Golgi network or endosomes. Proof of a high degree of surface expression may therefore be a meaningful predictor of *in vitro* and *in vivo* tracer binding.

Along with mRNA and protein expression data, the presence of a substantial number of functional receptors on the cell surface is an essential benchmark for nuclear imaging. In agreement with previous findings, the affinity of the DOTA-integrin $\alpha\beta 6$ knottin for BxPC-3 and Capan-2 cells was found to be in the low nanomolar range (1.69 and 9.46 nM) as revealed by radioligand binding assay. It is interesting to note that the affinity of the knottin did not change upon either the addition of a DOTA chelator moiety or incorporation of natural non-radioactive lutetium (^{nat}Lu) into the chelator.

Compared to previous biodistribution studies (¹⁸F, ^{99m}Tc) using the same knottin (35, 36), a higher uptake of the ¹⁷⁷Lu tracer in tumors was observed here. Some organs like lung, stomach, colon and kidney showed tracer uptake at early time points yet this was washed away at later time points. As integrin $\alpha\beta 6$ expression in mouse kidney had been found several orders of magnitude lower than in xenograft tumors (Fig. 1A), this uptake in the renal medulla may be attributed to unspecific uptake by transporters, e.g. OATPs. A high amount of tracer was accumulated in the kidney at early time points (day 1: $19.2 \pm 2.5\%$ IA/g); there was a significant reduction at later time points (day 8: $2.3 \pm 0.40\%$ IA/g). However, in the tumor the activity was cleared at a much lower rate than in the kidney. Higher tracer activity in the kidney resulted in a lower tumor-to-kidney ratio at day-1, however, Capan-2 tumors were clearly recognized. Similarly, a higher tumor-to-organ ratio at all-time points gives an advantage for contrast and favorable tumor imaging. These findings again underline that the tracer binds in the tumor specifically and with greater affinity. However, tumor activity is still limiting the full impact of this targeting approach. In addition to characterizing the principal tracer pharmacology with the therapeutic radionuclide ¹⁷⁷Lu, this study was also designed to see whether tumor uptake would be higher with this nuclide than with the previously tested ones. As ¹⁷⁷Lu has a comparatively long half-life and potentially allows for a smaller peptide/radiometal ratio, the resulting higher molar activity was hoped to lead to higher tumor accumulation. Indeed, a molar activity of 44 GB/ μ mol was achieved, which was 2.4-fold higher than the one reported for the ⁶⁴Cu conjugate before (19). Unfortunately, tumor uptake was not found to be higher than in this study. A further improved initial tracer uptake in the tumors should be the goal for the continued path towards translation into the clinic. With affinities already in a very favorable range, overall design of the conjugate could be modulated, e.g. by the introduction of a different spacer. In addition, increasing the molar activity of the tracer formulation may result in increased tumor uptake.

For dosimetry, biodistribution data were extrapolated to human adult male using OLINDA/Exm software for the therapeutic radionuclide ¹⁷⁷Lu. This dosimetric projection provided a preliminary estimate for assessing the therapy-associated risk of radiation damage. The absorbed doses in the kidney propose it as the dose-limiting organ. As this method involves two scaling methods to extrapolate human data, the resulting prediction is of preliminary nature and needs further confirmation, e.g. by collecting corresponding data from human subjects.

In the framework of this study, only a single dose of tracer was injected into the mice, and the animals were monitored for no longer than eight days. Consequently, no significant impact was observed in H&E-stained tissue sections of day 1 and day 8 post injections. A therapy study, potentially involving multiple/repeated administrations of tracer is required to establish therapeutic efficacy. Still, the uptake of tracer by xenograft tumors at different time points and faster renal clearance confirms the suitability of the DOTA-integrin $\alpha\beta 6$ knottin as a tracer and of integrin $\alpha\beta 6$ as a valid target for diagnosis and potential targeted radionuclide therapy in PDAC.

CONCLUSION

In this study, binding of the peptide tracer ¹⁷⁷Lu-DOTA-integrin $\alpha\beta 6$ knottin to its target both *in vitro* and *in vivo* was investigated. ¹⁷⁷Lu-DOTA-integrin $\alpha\beta 6$ knottin exhibited high affinity and specific binding to target-positive cells and tumors. The study demonstrated the translational potential of this tracer for imaging and therapy of integrin $\alpha\beta 6$ -overexpressing tumors like PDAC.

DATA AVAILABILITY STATEMENT

The datasets presented in this study can be found in online repositories. The names of the repository/repositories and accession number(s) can be found in the article/**Supplementary Material**.

ETHICS STATEMENT

The animal study was reviewed and approved by Landesamt für Gesundheit und Soziales, Land Berlin.

AUTHOR CONTRIBUTIONS

Concept and experimental design: BW, SG, RK, SR, and CG. Development of methodology: RK and NB. Acquisition of data: SS, TH, SE, SP, and NB. Analysis and interpretation of data:

SS and CG. Writing, review, and/or revision of the manuscript: SS, CG and all other authors. Study supervision: BW, SG, and CG. All authors contributed to the article and approved the submitted version.

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SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fonc.2021.684713/full#supplementary-material>

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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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