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Preclinical evaluation of [111In]MICA-401, an activity-based probe for SPECT imaging of in vivo uPA activity

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Abstract
Urokinase-type plasminogen activator (uPA) and its inhibitor PAI-I are key players in cancer invasion and metastasis. Both uPA and PAI-1 have been described as prognostic biomarkers, however non-invasive methods measuring uPA activity are lacking. We developed an indium-111 ($^{111}\text{In}$) - labelled activity-based probe to image uPA activity in vivo by single photon emission computed tomography (SPECT). A DOTA-conjugated uPA inhibitor was synthesized and radiolabelled with $^{111}\text{In}$ ([$^{111}\text{In}$]MICA-401), together with his inactive, hydrolysed form ([$^{111}\text{In}$]MICA-402). A biodistribution study was performed in mice (healthy and tumour-bearing) and tumour-targeting properties were evaluated in two different cancer xenografts (MDA-MB-231 and HT29) with respectively high and low levels of uPA expression in vitro, with either the active or hydrolysed radiotracer. MicroSPECT was performed 95 h post injection (pi) followed by ex vivo biodistribution. Tumor uptake was correlated with human and murine uPA expression determined by ELISA and immunohistochemistry (IHC). Biodistribution data with the hydrolysed probe [$^{111}\text{In}$]MICA-402 showed almost complete clearance 95 h pi. The ex vivo biodistribution and SPECT data with [$^{111}\text{In}$]MICA-401 demonstrated similar tumour uptake in the two models: ex vivo 5.68 ±1.41%ID/g versus 5.43 ± 1.29%ID/g and in vivo 4.33 ± 0.80 versus 4.86 ± 1.18 for MDA-MB-231 and HT-29 respectively. Human uPA ELISA and IHC showed significant higher uPA expression in the MDA-MB-231 tumours, while mouse uPA staining revealed similar staining intensity of both tumours. Our data demonstrate the non-invasive imaging of uPA activity in vivo although the moderate tumour uptake and hence potential clinical translation of the radiotracer warrants further investigation.

1. Introduction
Enzymatic degradation of the extracellular matrix (ECM) plays an important role in cancer invasion and metastasis. The urokinase-type plasminogen activator (uPA) and its inhibitor PAI-I are key players in a proteolytic cascade, involved in physiological and pathophysiological degradation and remodelling of the ECM. These processes promote tumour progression, invasion and metastasis [1, 2]. uPA is a serine protease, which after binding with the urokinase-type plasminogen activator receptor (uPAR) initiates the conversion of the inactive zymogen plasminogen to plasmin. The protease plasmin triggers fibrinolysis, cleaves a range of ECM components and by
activation of matrix metalloproteinase (MMPs) facilitates the degradation of the ECM. Plasmin will also indirectly facilitate release of growth factors from the interstitial matrix and basement membrane [3]. uPA is expressed by a variety of cells (including tumour cells) as a pro-enzyme (pro-uPA) which is converted to its active form by proteolysis. Interestingly this conversion is catalysed by plasmin, thus creating a positive feedback loop [4, 5]. The activation of pro-uPA is significantly accelerated upon binding of the pro-enzyme to its specific cell surface receptor [6]. Numerous studies examining the uPA system indicate that high levels of uPA and/or PAI-1 in tumours are correlated with poor prognosis. Also, these have been validated at the highest level of evidence as clinical biomarkers of prognosis in breast cancer [7–10]. Moreover, uPA is also suggested as a prognostic biomarker in other tumour types including lung, prostate, ovarian, cervical, pancreatic, colorectal and gastric cancer [11-17]. Determination of uPA and PAI-1 levels is currently done using standardized enzyme-linked immunosorbent assays (ELISA), thus fresh tissue samples are needed. More importantly, this method lacks the ability to discriminate between active and inactive uPA (pro-uPA) [18]. Therefore, the development of non-invasive imaging methods to visualize and quantify uPA activity in vivo could be a valuable tool in the clinical practice towards patient stratification and therapy monitoring [19-22]. In particular, molecular imaging techniques in nuclear medicine serve as highly sensitive tools that are ideally suited for non-invasive imaging of cancer and small micrometastatic lesions. Targeting uPA with positron emission tomography (PET) or single photon emission computed tomography (SPECT) activity-based imaging probes would hence offer the ability to monitor uPA activity in vivo. So far, non-invasive imaging of the uPA-system has been limited to imaging probes targeting the uPAR, which may not be an accurate indication of the actual uPA activity in the tumour [23-25] as binding of uPA/pro-uPA to its receptor and activation of pro-uPA are critical steps for cancer cell dissemination [26]. A first-in-human clinical trial with PET imaging of uPAR in patients with breast, prostate and bladder cancer, is recently published and high uptake in both primary tumor lesions and lymph node metastases was seen and paralleled high uPAR expression in excised tumor tissue [27].

Previously, our research group reported the synthesis and in vivo evaluation of an activity-based [18F]-labeled uPA probe [28]. This probe [18F]-1 was designed based
on a phosphonate scaffold (which binds covalently to uPA) bearing a guanidine group to enhance the selectivity for uPA [29] and a PEG linker to connect to the \(^{18}\text{F}\)-4-fluorobenzoyl group (Fig. 1). However, the aforementioned radiotracer displayed low absolute tumour uptake values due to unfavourable stability and pharmacokinetic properties. The same uPA activity-based probe has now been linked to a DOTA-chelator which is used as a complexing agent for radioisotope indium-111 (\(^{111}\text{In}\)) (MICA-401, Fig. 1). The affinity and covalent bond formation of MICA-401 for uPA and the selectivity over a set of related trypsin-like serine proteases was analysed.

Finally, the functional activity-based probe \(\text{[^{111}\text{In}]MICA-401}\) was evaluated for its capacity to image uPA activity in vivo by SPECT in two different cancer models, with a confirmed distinct uPA expression in vitro. As a control, a corresponding hydrolysed inactive probe (\(\text{[^{111}\text{In}]MICA-402, Fig. 1}\)) was used in vivo in both cancer models. This probe serves as a negative control since we confirmed that MICA-402 has no affinity for uPA and is not able to form a covalent bond with uPA and related proteases.

2. Experimental section

2.1. Organic synthesis
A detailed description of the organic synthesis of compounds MICA-401 and MICA-402 and their corresponding precursor (compound 3 and 5, respectively) is provided in Supplemental data.

2.2. In vitro evaluation of binding kinetics
Previously described methods were used to determine the IC\(_{50}\) and k\(_{\text{app}}\) values of compounds uPA-PET, MICA-401 and MICA-402 for uPA, tPA, Thrombin, plasmin, FXa Plasma KLK, KLK1, KLK4, KLK8, HNE and AChE. For HNE we used the same method as described for uPA but with Neutrophil elastase, (100 \(\mu\)g (Enzo) dissolved in 200 \(\mu\)L 50 mM NaAc, 150 mM NaCl pH 5.5) as the enzyme solution MeOSuc-Ala-Ala-Pro-Val-pNA, (Enzo, Km: 125 \(\mu\)M) as substrate and a Tris buffer at pH 7.8 [29-31].

2.3. Radiochemistry
Radiosynthesis of \(\text{[^{111}\text{In}]MICA-401}\)
0.5 M ammonium acetate buffer (pH 5.5, 450 µL) was added to 342.62±2.22 MBq [\(^{111}\text{In}\)] - InCl\(_3\) (Mallinckrodt, Petten, The Netherlands). Thirty µL of a solution of compound 3 (Supplemental data; 1 mg/ml in MeOH) were added and the solution was heated for 45 min at 60°C. The reaction mixture was diluted with 200 µL of water and injected into the HPLC loop. The crude reaction mixture was purified on a Chromolith® Performance RP-18e, 5 µm, 100 x 4.6 mm eluted with a solution of NaOAc 0.05M, pH 5.5 / EtOH (75/25, V/V) at 1 ml/min. [\(^{111}\text{In}\)]MICA-401 has a retention time of 21-22 min and was collected and the solution was evaporated at 40°C under a mild flow of compressed air. The volume was adjusted to 1 mL with EtOH and the solution was filtered through a 0.22 µm sterile filter (Pall Acrodisc syringe filter, 13 mm, 0.2 µm) and diluted with 0.9% sodium chloride solution to reach a concentration of EtOH below 10%. [\(^{111}\text{In}\)]MICA-401 was synthesized in a 59% (n=4) yield. Chemical and radiochemical purity (>98%) of the tracer were measured by analytical HPLC (Chromolith® Performance RP-18e, 5 µm, 100 x 4.6 mm) eluted with 0.1% TFA in H\(_2\)O / 0.1% TFA in CH\(_3\)CN (78/22, V/V) at flow rate of 1 ml/min. The specific activity was determined using a UV calibration curve (\(\lambda = 204\) nm) and was in the range of 6–12 GBq/µmol.

Radiosynthesis of [\(^{111}\text{In}\)]MICA-402

0.5 M ammonium acetate buffer (pH 5.5, 450 µL) was added to 354.46±15.54 Mbq [\(^{111}\text{In}\)] - InCl\(_3\) (Mallinckrodt, Petten, The Netherlands). Thirty µL of a solution of compound 5 (Supplemental data; 1 mg/ml in MeOH) were added and the solution was heated for 45 min at 60°C. The reaction mixture was diluted with 200 µL of water and injected into the HPLC loop. The crude reaction mixture was purified on a Chromolith® Performance RP-18e, 5 µm, 100 x 4.6 mm eluted with a solution of NaOAc 0.05M, pH 5.5 / EtOH (90.5/9.5, V/V) at 1 ml/min. [\(^{111}\text{In}\)]MICA-402 has a retention time of 15-16 min and was collected and filtered through a 0.22 µm sterile filter (Pall Acrodisc syringe filter, 13 mm, 0.2 µm). [\(^{111}\text{In}\)]MICA-402 was obtained in a radiochemical yield of 71% (n=2). Chemical and radiochemical purity (>98%) of the tracer were measured by analytical HPLC (Chromolith® Performance RP-18e, 5 µm, 100 x 4.6 mm) eluting with 0.1% TFA in H\(_2\)O / 0.1% TFA in CH\(_3\)CN (90/10, V/V) at flow rate of 1 ml/min. The specific activity was determined using a UV calibration curve (\(\lambda = 207\) nm) and was in the range of 8–10 GBq/µmol.
For the measurement of the partition coefficient, n-octanol (1 ml) and PBS 0.01 M (pH = 7.4) were added to a test tube and allowed to equilibrate for 10 min. Five microliters of $[^{111}\text{In}]$MICA-401 or $[^{111}\text{In}]$MICA-402 was added and vortexed for 2 min at RT. The tube was centrifuged at 3000 rpm for 10 min. Fractions of 500 µL of both phases were put in pre weighted tubes and measured in the gamma counter. The partition coefficient D was calculated as: [radioactivity (cpm/mL) in n-octanol/radioactivity (cpm/mL) in PBS]. The log D(n-octanol/PBS pH = 7.4) value is given as an average of three measurements. The partition coefficient of $[^{111}\text{In}]$MICA-401 and $[^{111}\text{In}]$MICA-402 were measured and log D values of -2.73 ± 0.01 and -4.15 ± 0.02 were respectively obtained.

2.4. Cell cultures and animal model

Female 6-week-old CD1 −/− Foxn1nu mice (n=37) were obtained (25.6 ± 2.1 g) from Charles River Laboratories (L’arbresle, France). The animals were kept under environmentally controlled conditions (12 h light/dark cycle, 20–24°C and 40–70% relative humidity) in individually ventilated cages with food and water ad libitum. The study protocol was approved by the local Animal Experimental Ethical Committee of the University of Antwerp, Belgium (2011-20). All animal studies were ethically reviewed and carried out in accordance with European Directive 86/609/EEC Welfare and Treatment of Animals.

MDA-MB-231-luc2-GFP Bioware® (human breast cancer cell line, Perkin Elmer, Waltham, USA) and HT-29 (human colon adenocarcinoma, ATCC, Rockville, USA) were selected for their different uPA expression. Secreted uPA has been demonstrated in the culture medium of MDA-MB-231, but not in HT-29 [32, 33]. Cells were routinely cultured in Dulbecco’s Modified Eagle Medium supplemented with 10% heath inactivated fetal bovine serum (FBS), 2 mM glutamine, 1% sodium pyruvate and 1% penicillin/streptomycin (Invitrogen, Merelbeke, Belgium). To establish an orthotopic MDA-MB-231 model, approximately 5.10^6 viable MDA-MB-231 cells were resuspended in 100 µl phosphate buffered saline (PBS) containing Matrigel (1:2 in PBS; BD Bioscience, San Jose, USA) and were subsequently injected in the second left abdominal mammary fat pad. For the HT-29 subcutaneous model, approximately 1.10^6 viable cells, suspended in 100 µl PBS, were inoculated in the right hind leg of the mice. Tumour growth was monitored twice a week with digital calliper measurements starting upon detection of palpable tumours. Tumour volume was
calculated using the modified ellipsoid formula \( \frac{1}{2} \) (length \( \times \) width\(^2\)). Four to six weeks after MDA-MB-231 tumour inoculation, when tumours reached a volume of 150 mm\(^3\), and 2-3 weeks after HT-29 tumour inoculation, when tumours reached a volume of 400-500 mm\(^3\), mice were subjected to \( \mu \)SPECT/CT imaging.

2.5. Selection of imaging window

To obtain the optimal time point for \( \mu \)SPECT imaging, a biodistribution study was performed in female nude mice (n=12) including both healthy animals (n=9) as MDA-MB-231 tumour-bearing mice (n=3). These animals were injected with 30-37 MBq of the radiolabelled uPA activity-based probe \([^{111}\text{In}]\text{MICA-401}\) via the lateral tail vein. At, 30 min, 2 h and 18 h post injection, each time three healthy animals were sacrificed by cervical dislocation and an \textit{ex vivo} biodistribution was performed. The tumour-bearing animals underwent longitudinal static SPECT scans (cfr. infra) at 2 h, 4 h, 18 h and 95 h post injection and were sacrificed at the end of the last scan. Blood was collected via cardiac puncture and the main organs were thereafter rapidly removed and rinsed in PBS. Radioactivity in the samples was measured using an automatic gamma-counter (WIZARD2 Automatic Gamma counter 2480, Perkin Elmer, Waltham, USA) and expressed as percentage of the injected dose per gram of tissue plus or minus the standard deviation (%ID/g \( \pm \) SD).

2.6. Imaging study with active and hydrolysed \(^{111}\text{In}\)-labeled uPA activity-based probe

\( \mu \)SPECT/CT scans of the uPA activity-based probe \([^{111}\text{In}]\text{MICA-401}\) were performed in a HT-29 (n=5), MDA-MB-231 (n=5) and a healthy control group (n=5). Similarly, \( \mu \)SPECT/CT scans with the hydrolysed inactive uPA probe \([^{111}\text{In}]\text{MICA-402}\) were performed also in a HT-29 (n=5) and MDA-MB-231 (n=5) group. These static SPECT scans were acquired 95 h post injection and \textit{ex vivo} biodistribution, as described above, was additionally performed. The radiotracer was injected in the lateral tail vein, with an average of 20.35\( \pm \) 8.14 MBq (9.41 \( \pm \) 3.36 GBq/\( \mu \)mol) for \([^{111}\text{In}]\text{MICA-401}\), and 35.15\( \pm \) 4.81 MBq (specific activity not determined for inactive probe) for \([^{111}\text{In}]\text{MICA-402}\), depending on the yield of each radiosynthesis. The \( \mu \)SPECT scans had a duration of 30, 45 or 90 min, depending on the absolute injected dose, to obtain similar signal-to-noise ratios in the reconstructed images for every performed \( \mu \)SPECT scan. To undergo the scan procedure, animals were anaesthetized with isoflurane (3-5% for induction, 1-2% for maintenance). Further, all
mice were scanned on a VECTor system (Milabs, Utrecht, The Netherlands) with the animal bed translated in three dimensions using an XYZ stage. A 20% main photopeak was centered at 171 and 245 keV to reconstruct the $^{111}$In images on 1.2-mm$^3$ voxels by 20 iterations of 16 POSEM (pixel-based ordered-subset expectation maximization) subsets. For whole body imaging, 8 bed positions were used with a head first prone orientation and anatomical CT images were acquired (45kV, 615 µA). Region of interest (ROI) were drawn in PMOD Software 3.3 (PMOD technologies, Zürich, Switzerland) on the CT images manually. Subsequently, tracer uptake was expressed as %ID/g tissue and by standardized uptake values (SUV) mean, which was generated by summation of voxels within the tomographic planes. SUV was defined as $(CT*W)/D_{\text{inj}}$, where CT is radioactivity counts in the tissue (kBq/cc), W is weight of the animal (g) and $D_{\text{inj}}$ is injected dose (kBq). A technical consideration to make is that despite the differences in tumour volumes between the two different xenograft models (150 versus 450 mm$^3$), the smallest tumor diameter (i.e. around 4 mm) is 4 times the resolution of the scanner (i.e. around 1mm) hereby excluding partial volume effects. Therefore no signal recovery differences are to be expected between the different tumor diameters.

2.7. Validation by anti-human uPA ELISA and ex vivo histology

The human uPA protein level of both models was determined in the tumour tissue ($n=5$ for MDA-MB-231 and $n=5$ for HT-29) using ELISA. At the end of each scan of the aforementioned imaging study, tumour tissue was collected and part of the tumour was snap-frozen in liquid nitrogen and stored at −80 °C until analysis. The remaining part of the tumours was used for histological examination. All tumour tissues were homogenized for ELISA analysis using a GentleMACS tissue dissociator (Miltenyi Biotec, North Ryde, New South Wales, Australia). Total protein content of the tumour tissue samples was measured using the BCA Protein Assay Kit (Thermo Fisher Scientific, Waltham, USA) according to the protocol provided. The total human uPA content was measured using the Imubind® uPA ELISA kit (American Diagnostica Inc., Stamford, USA) according to the manufacturer’s instructions and normalized to the total protein content of the tumour lysates (ng uPA/mg protein).

The remaining of all tumours ($n=20$) and lymph nodes (LNs; inguinal: $n=23$; axillary: $n=24$) were excised and fixed in formalin. Immunohistochemistry (IHC) was
performed using two different primary rabbit polyclonal antibodies, one targeting specific human (ab24121, Abcam, Cambridge, UK) and a second one targeting specific mouse uPA (LS-C150831, Lifespan BioSciences, Inc, Seattle, USA). Negative control sections were processed identically, except for the lack of the primary antibody. A human prostate carcinoma slide and a mouse pancreatic carcinoma (with confirmed uPA expression) was used as positive control for human and mouse uPA immunostaining respectively and to evaluate species cross reactivity of both uPA-targeting antibodies. In addition, to investigate proliferation status of the tumours and for the detection of human tumour cells in mouse lymph nodes (micrometastases) a rabbit polyclonal antibody was used targeting human Ki67 (#12202, Cell Signaling Technology, Danvers, USA). A tumour slide (already stained positively for Ki67) was included as positive control tissue. A description of the protocol and scoring system is provided in Supplemental data.

2.8. Statistical analysis

GraphPad Prism 6.0c (GraphPad Inc, La Jolla, USA) was applied for all statistical analysis. To verify if the data were normally distributed, a Shapiro-Wilk normality test was applied. Differences between the biodistribution groups was statistically evaluated by one-way ANOVA and Tukey’s multiple comparison test. Differences between two unpaired independent groups were analysed with non-parametric Mann Whitney U-test. For a correlation study, a non-parametric Spearman test was performed. Data are reported as mean ± SEM and p < 0.05 was considered statistically significant here.

3. Results

3.1. Chemistry and in vitro evaluation

The cold standard MICA-401 was synthesized in two steps starting from the amine 2. [28]. In the first step, the addition of the DOTA chelator was achieved in a moderate yield (42%, Figure S1) and gave access to precursor 3. Complexation of the DOTA ring with InCl3 led to the standard MICA-401 also in moderate yield (42%,). The same synthetic strategy was applied to prepare compounds 5 and MICA-402 after hydrolysis of amine 2 (Figure S2). Both compounds MICA-401 and MICA-402 were evaluated in vitro to determine their affinity for uPA (human and mouse) and for
a panel of 9 related serine proteases and acetylcholinesterase. The uPA SPECT probe MICA-401 appeared to be a potent, selective and irreversible uPA inhibitor \( (k_{\text{app}} = 4.80 \times 10^3 \text{ M}^{-1} \text{s}^{-1}) \) for human uPA) while the hydrolyzed probe MICA-402 lost all affinity for uPA (Table 1) as anticipated. In conclusion, the SPECT-probe MICA-401 is able to make a covalent bond selectively with uPA compared to a broad panel of related enzymes, and therefore should allow the selective imaging of uPA activity in vivo. Since the hydrolysed compound MICA-402 shows no binding to any of the enzymes, this compound is well suited to serve as negative control to demonstrate that the observed imaging is not due to any aspecific interaction.

3.2. Radiochemistry

The radiosynthesis of \[^{111}\text{In}]\text{MICA-401}\) and \[^{111}\text{In}]\text{MICA-402}\) was performed by complexation of \[^{111}\text{In}]\text{InCl}_3\) at 60°C for 45 min (Fig. 2). The crude reaction mixture was purified by analytical HPLC. The column was eluted with a biocompatible mobile phase (NaOAc 0.05M, pH 5.5 / EtOH) and the collected radiotracer was filtered through a sterile filter and diluted with sterile solution 0.9% NaCl to reduce the ethanol concentration to below 10% for in vivo evaluation. \[^{111}\text{In}]\text{MICA-401}\) and \[^{111}\text{In}]\text{MICA-402}\) were respectively produced with an average radiochemical conversion of 90% (n=4) and 92% (n=2). After purification and formulation the yields dropped to 49% (n=4) for \[^{111}\text{In}]\text{MICA-401}\) and 71% (n=2) for \[^{111}\text{In}]\text{MICA-402}\). The radiochemical purity was >95% and the tracers were stable in solution for at least 5 h post radiosynthesis as determined by HPLC analysis.

3.3. Time window selection

The radiotracer \[^{111}\text{In}]\text{MICA-401}\) slowly cleared from the blood of healthy animals, with still 7.54 ± 0.78% ID/g present at 18 h post injection. Clearance was mainly through the renal system with a kidney uptake of 7.14 ± 0.77 %ID/g at 30 min post injection and decreasing to 2.59 ± 0.31 %ID/g at 18 h post injection (Fig. 3). Other organs showing moderate tracer uptake were lungs, liver, spleen and skin. Pancreas, stomach, small and large intestine, muscle, bone and brain showed negligible uptake (< 4%ID/g) at any of the time points. Further, ex vivo biodistribution of tumour bearing animals 95 h post injection indicated an almost complete clearance from the blood (1.40 ±0.35 %ID/g). Uptake was seen in the liver (7.10 ±1.15%ID/g), spleen (5.87 ±0.19%ID/g), skin (5.83 ±1.67%ID/g) and adjacent (12.08 ± 7.13%ID/g) and
axillary (17.17%ID/g, n=1) LNs. Tumour uptake was 5.16 ± 1.04 %ID/g (Fig. 3).

SPECT images in Fig. 4 show good tumour visualization from 18 h post injection onwards, with a more pronounced radiotracer uptake in the rim of the tumour. The optimal tumour contrast is reached at the 95 h p.i. time point. High uptake by skin and active drainage to and uptake by lymph nodes are also clearly visible (Fig. 4).

3.4. In vivo imaging study

Post-scan ex vivo biodistribution in healthy control animals, as well as the MDA-MB-231 and HT29 group showed a comparable tracer distribution of the uPA inhibitor [111In]MICA-401, with high uptake in the liver (5.55 ± 1.01 %ID/g), spleen (4.50 ± 1.22 %ID/g) and skin (6.22 ±1.20 %ID/g) as averaged over the three groups (Fig. 5). High tumour uptake (5.68 ±1.41%ID/g and 5.43 ± 1.29%ID/g) in the MDA-MB-231 and HT-29 model, respectively, was demonstrated. Also, a remarkably high uptake was observed in the LNs for all animals, up to 12.21 ± 6.70 %ID/g in the inguinal LNs of the MDA-MB-231 group.

In vivo SPECT images for [111In]MICA-401 are shown in Fig. 6. Both tumour types are easily detectable and moderate tumour uptake was observed for the MDA-MB-231 and HT-29 model: 1.18 ± 0.18 and 1.10 ± 0.29 respectively for SUVmean, and 4.33 ± 0.80 and 4.86 ± 1.18 respectively for %ID/g. Uptake of [111In]MICA-401 in liver, LNs and skin were also confirmed on the SPECT images. Further, a high and significant correlation was obtained between the in vivo measured tumour SUVs and the ex vivo %ID/g tumour tissue (r = 0.6070 and p = 0.0214).

For the hydrolysed inactive uPA probe, [111In]MICA-402, an almost complete removal from the mouse body was observed ex vivo and in vivo (Fig. 7A and B resp) 95 h p.i. as uptake in all organs was less than 1%ID/g. Further, these SPECT images equally indicated a renal clearance of [111In]MICA-402. As anticipated no tumours could be detected with this hydrolysed inactive uPA probe (Fig. 7B).

3.5. Ex vivo validation

uPA levels determined by ELISA in both tumour types showed a significantly higher human uPA expression in MDA-MB-231 tumour samples (21.42 ± 3.40 ng uPA/mg
protein) compared to HT-29 tumours (1.96 ± 0.36ng uPA/mg protein, p<0.0001).

Target specificity and species specificity for both anti-human and anti-mouse uPA antibodies were investigated. The positive control tissue (human prostate carcinoma) for anti-human uPA showed a clear, specific uPA staining pattern of the glands, while the negative control without the primary antibody proved no background staining. The positive control tissue (mouse pancreatic carcinoma) for the anti-mouse uPA antibody showed clear and specific uPA staining while the negative control showed only a slight background staining. To investigate possible cross-reactivity, staining of the mouse uPA positive control tissue with the anti-human uPA antibody however did show weak staining of tumour cells and staining of mouse lymphoid cells and macrophages, so species-cross reactivity occurs with the anti-human uPA antibody. Also staining of the human prostate carcinoma tissue with the anti-mouse uPA antibody demonstrated a clear staining pattern comparable to that of the human uPA staining, confirming species-cross reactivity of this antibody as well, however staining differences was still present when human antibodies were compared versus mouse uPA and vice versa.

The two tumour types showed a clear staining with the anti-human uPA antibody in both MDA-MB-231 and HT-29 tumours. An overall, high staining was observed in the MDA-MB-231 tumours, with both pericellular and intracellular staining of tumour cells, staining positively over the entire section of the tumours. By morphological criteria also tumoural stroma and some macrophages showed uPA immunoreactivity. (Fig. 8A and B). In the HT-29 tumours a weaker staining pattern was observed (which is in line with the ELISA data), with primarily granulated pericellular uPA staining with a more pronounced staining at the rim of the tumours (Fig. 8D and E). No (or little) stromal uPA staining and no intracellular staining of the tumour cells were observed. IHC scores for the MDA-MB-231 were significantly higher compared to HT-29 scores (Fig. 9, p = 0.008).

Staining with the anti-mouse antibody revealed a more comparable staining intensity and pattern between the two tumour types (Fig. 8C and F). In the MDA-MB-231 tumours a more heterogeneous staining pattern could be observed with a mix of intense and negative stained regions in the tumours. uPA immunoreactivity was found intracellularly, in tumoural stroma and macrophages. In the HT29 tumours, a typical
granulated, pericellular staining was again observed, with more stromal staining compared to the anti-human uPA antibody. Also macrophages and some endothelial cells showed positive uPA staining. No significant differences were demonstrated between the IHC scores (Fig. 9).

Due to the high LN uptake measured *ex vivo* and in vivo by the SPECT images, LN sections were also stained with H&E and antibody directed to human Ki67 was used to identify micro metastases. No Ki67 staining was visible and also no micro metastasis could be revealed on the H&E stained LN sections. In addition, LNs showed clear anti-mouse uPA staining in the medulla of the LN, where - among other cell types - histiocytes, macrophages and dendritic cells are present (Figure S3 A-C). More remarkable is the intense uPA staining of the surrounding brown adipose tissue, sometimes present on the dissected LN sections. A strong, specific staining of the adipocytes, small blood vessels and macrophages present in the brown fat is illustrated in Figure S3 A and D.

4. Discussion

Our objective is to develop a molecular imaging probe for discrimination between high and low uPA activity in tumours with a view on prediction of tumour invasiveness, therapeutic intervention and personalized treatment [20–23]. In a previous study with [18F]-1 (Fig. 1) [28], we observed an unfavourable stability profile combined with slow blood clearance. Scanning at later time points increases the tumour/blood ratio of the uPA activity based probe and therefore, 111- indium with a half-life of 2.80 days was selected to design a SPECT probe. The aforementioned PET probe showed a similar inactivation rate constant (k<sub>app</sub>) for human uPA (k<sub>app</sub> = 6.80 x 10<sup>3</sup> ± 0.25 x 10<sup>3</sup> M<sup>-1</sup> s<sup>-1</sup>) as the current SPECT probe MICA-401 (k<sub>app</sub> = 4.80 x 10<sup>3</sup> ± 0.32 x 10<sup>3</sup> M<sup>-1</sup> s<sup>-1</sup>), whereas both probes had 40-fold reduced inactivation rate constants for mouse uPA (Table 1). Both the SPECT probe MICA-401 and the PET probe demonstrated similar affinities for two closely related members of trysin-like protease, kallikrein (KLK)4 and KLK8 compared to uPA. However, rigorous analysis of the binding kinetics showed that the binding to KLK4 and KLK8 is reversible. This is a further demonstration that potent binding of a molecule with a warhead designed to be irreversible does not always result in irreversible binding [30].
Our *ex vivo* and *in vivo* biodistribution data with the uPA activity-based probe \[^{111}\text{In}]\text{MICA-401}\) demonstrated a high blood activity 18 h post injection (about 7 %ID/g), which declined to approximately 1 %ID/g at 95 h p.i. At this time point, tumours were clearly detectable and *ex vivo* biodistribution indicated MDA-MB-231 tumour uptake of approximately 5 %ID/g. Therefore, 95h p.i. was chosen as the ideal imaging time point for further experiments. Compared to our previously developed \[^{18}\text{F}]\text{-1}\) (Fig. 1) radiotracer, the uPA activity-based probe \[^{111}\text{In}]\text{MICA-401}\) showed substantially lower uptake in the large intestines and blood while a higher liver and spleen activity was observed [28]. In the MDA-MB-231 model we found a tumour uptake measured *ex vivo* of 5.7 % ID/g and *in vivo* of 4.3 %ID/g, while we obtained a relatively low tumour uptake with our previous \[^{18}\text{F}]\text{-1}\) uPA probe (peak uptake of 2.4 %ID/g *ex vivo* and 2.5 %ID/g *in vivo* at 4 h p.i) [28]. A direct comparison of the quantitative tumour uptake between previous published studies targeting uPAR [23-25, 34] is not feasible, as in our study enzymatically active uPA is targeted and quantified. Recently, an \[^{111}\text{In}\text{-labeled U33}, an antibody directed to uPA, showed a remarkable tumour uptake of 43.2% ID/g 72h p.i. in a prostate cancer xenograft which was attributed to the internalization through an uPAR mediated mechanism [35].

A similar uptake of \[^{111}\text{In}]\text{MICA-401}\) was observed for both tumour models *ex vivo* as well as *in vivo*, despite the known differential *in vitro* uPA expression between the two cell lines. Additionally, uPA ELISA and IHC using the anti-human uPA antibody performed on tumour tissue revealed a significant higher human uPA expression of the MDA-MB-231 model than the HT-29 model. However, IHC staining with the anti-mouse uPA antibody on the tumour sections showed no significant difference in mouse uPA expression between the tumour types suggesting that mouse uPA levels were more comparable. Nevertheless, caution should be taken in the interpretation of these IHC data as we demonstrated species cross-reactivity of both uPA-targeting antibodies. This discrepancy of *in vitro* and *in vivo* uPA expression has already been demonstrated by Hsiao et al., who evaluated an uPA sensitive fluorescence imaging probes in two different models, including the HT-29 and a high uPA expressing fibrosarcoma model. Similar fluorescence intensity in both models was observed and it was hypothesised that HT-29 tumour cells need to recruit host uPA expressed by neighbouring stroma cells through its surface uPAR to grow *in vivo* [36]. In this case,
the amount of active uPA may be similar in both models, but therefore not the antigen levels (pro-uPA plus uPA). Another point to consider is the possible binding of the uPA SPECT probe to other serine proteases present in tumours. However, we demonstrated in an in vitro panel of highly related serine proteases, that the SPECT probe MICA-401 can only bind covalently to human and mouse uPA. Since imaging is performed at 95 h p.i. it is highly unlikely that any non-covalently bound probe is still present. This is confirmed by the absence of the inactivated probe MICA-402 at 95 h p.i., of which we confirmed that it is unable to make covalent bonds to the tested serine proteases.

Remarkably, a very high uptake was also observed in the LNs in both cancer models and the healthy control group. Both H&E and staining with a specific anti-human Ki67 antibody were thoroughly analysed but no proliferating human tumour cells could be detected in the mouse LNs, excluding the presence of micro metastases. Immunohistochemical staining with anti-mouse uPA of all the LNs revealed specific uPA presence in the cortex and medulla of the LNs, and in particular histiocytes, macrophages and dendritic cells stained positively. Noteworthy, an intense uPA staining of the brown fat, occasionally present on the section of the dissected LNs, was present. A specific staining of the adipocytes and small blood vessels in the adipose tissue was visible. Indeed, high expression levels of PAI-1 and uPA have been described in adipocyte tissue and furthermore uPAR is known to be preferentially expressed in infiltrating adipose tissue macrophages [37, 38].

5. Conclusion

We have synthesized and characterized an $^{111}$In labelled activity-based SPECT probe that selectively targets active uPA and we have proven its usefulness in two xenograft mouse models. Moderate tumour uptake was found in both tumour models with clear, detectable tumours on µSPECT images at 95 h p.i. The high uptake in non invaded lymph nodes reported here may represent a limitation to the clinical translation of this tracer for oncological applications. Therefore, translational capabilities of this $[^{111}\text{In}]{\text{MICA-401}}$ radiotracer for future clinical cancer patient management remains to be further explored.

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References


33. Reiter LS, Kruijthof EKO, Cajot J-F, Sordat B. The role of the urokinase receptor in extracellular matrix degradation by the HT29 human colon...
### Table 1: In vitro evaluation of compounds uPA-PET [28] and MICA-401 as well as negative control compound MICA-402. (*: Reversible binding kinetics; **: Irreversible binding kinetics)

<table>
<thead>
<tr>
<th>IC$_{50}$ (µM)</th>
<th>PET probe (28)</th>
<th>SPECT probe (MICA-401)</th>
<th>Hydrolyzed probe (MICA-402)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Human uPA</td>
<td>0.019 ± 0.002**</td>
<td>0.022 ± 0.002**</td>
<td>&gt;10</td>
</tr>
<tr>
<td>Mouse uPA</td>
<td>0.058 ± 0.056**</td>
<td>1.29 ± 0.08**</td>
<td>&gt;25</td>
</tr>
<tr>
<td>tPA</td>
<td>46.1 ± 4.6</td>
<td></td>
<td>&gt;10</td>
</tr>
<tr>
<td>Thrombin</td>
<td>13.2 ± 0.6</td>
<td>&gt;10</td>
<td>&gt;250</td>
</tr>
<tr>
<td>Plasmin</td>
<td>5.9 ± 0.8</td>
<td>&gt;10</td>
<td>&gt;250</td>
</tr>
<tr>
<td>Fxa</td>
<td>62</td>
<td>&gt;10</td>
<td>&gt;10</td>
</tr>
<tr>
<td>Plasma KLK</td>
<td>&gt;2.5</td>
<td>&gt;10</td>
<td>&gt;10</td>
</tr>
<tr>
<td>KLK1</td>
<td>&gt;10</td>
<td>&gt;10</td>
<td>&gt;10</td>
</tr>
<tr>
<td>KLK4</td>
<td>0.013 ± 0.001 *</td>
<td>0.036 ± 0.003*</td>
<td>&gt;25</td>
</tr>
<tr>
<td>KLK8</td>
<td>0.050 ± 0.004*</td>
<td>0.033 ± 0.004*</td>
<td>&gt;2.5</td>
</tr>
<tr>
<td>HNE</td>
<td>&gt;2.5</td>
<td>&gt;2.5</td>
<td>&gt;10</td>
</tr>
<tr>
<td>AChE</td>
<td>&gt;20</td>
<td>&gt;20</td>
<td>&gt;20</td>
</tr>
</tbody>
</table>
Figure 1: Chemical structures of fluor-18 and indium-111 labeled uPA inhibitors $[^{18}F]1$, $[^{111}In]MICA-401$ and the uPA hydrolysed probe $[^{111}In]MICA-402$. 

$[^{18}F]1$

uPA (kapp) = $6.80 \times 10^3 \pm 0.25 \times 10^3$ M$^{-1}$ s$^{-1}$  
uPA IC$_{50}$ = 19 nM

$[^{111}In]MICA-401$

uPA (kapp) = $4.80 \times 10^3 \pm 0.32 \times 10^3$ M$^{-1}$ s$^{-1}$  
uPA IC$_{50}$ = 22 nM

$[^{111}In]MICA-402$

uPA IC$_{50}$ > 10 µM
Figure 2: Radiosynthesis of $[^{111}\text{In}]$MICA-401 and $[^{111}\text{In}]$MICA-402; Reagents and conditions: (a) $[^{111}\text{In}]$InCl$_3$, NH$_4$OAc 0.5M, pH 5.5, 45 min at 60°C, 49-71%.
Figure 3. *Ex vivo* biodistribution of $[^{111}\text{In}]\text{MICA-401}$ at different time points post injection in healthy and tumour-bearing mice
*data from adjacent LN, axillary LN and tumours were only collected at 95 h p.i. Results are shown as mean ±SD (n=3/group).
Figure 4. Representative SPECT images of an MDA-MB-231-bearing mice longitudinally imaged with the activity-based uPA probe [$^{111}$In]MICA-401. Sagittal images at the different time points (2 h, 4 h, 18 h and 95 h p.i. resp.). White arrows indicate tumours, yellow arrow indicates liver and red arrows indicate active drainage to lymph nodes (axillary and inguinal). (SUV scale from 0 – 2).
Figure 5. *Ex vivo* biodistribution of the uPA activity-based probe $[^{111}\text{In}]$MICA-401 at 95h p.i. in MDA-MB-231, HT-29 and healthy mice. Results are shown as mean ±SD (n=5/group).
Figure 6. Representative SPECT images at 95h post injection of the uPA activity-based probe $[^{111}\text{In}]{\text{MICA-401}}$ in the three models. A) A healthy control mice B) HT-29 tumour bearing mice, C) MDA-MB-231 tumour bearing mice. D) White arrows indicate tumours and yellow arrows indicate liver. All images scaled to the same color scale (SUV scale from 0 – 2).
Figure 7. Ex vivo biodistribution and a representative SPECT image at 95 h post injection of the hydrolysed, inactive uPA probe [\(^{111}\text{In}\)]MICA-402. A) Ex vivo biodistribution data of MDA-MB-231 and HT-29 tumour bearing mice. Results are shown as mean ±SD (n=5/group). B) SPECT image of a HT-29 (left) and MDA-MB-231 (right) tumour bearing mice imaged 95 h p.i. of the inactive radiotracer. Two images scaled to the same color scale (SUV scale from 0 – 0.1).
Figure 8. IHC staining of human and mouse uPA in tumour sections. A and B) MDA-MB-231 tumour stained for human uPA, C) MDA-MB-231 tumour stained for mouse uPA, D and E) HT-29 tumour stained for human uPA, F) HT-29 tumours stained for mouse uPA.

Figure 9. IHC scores of human and mouse uPA of both tumour models. Grading was performed by two independent observers and as described in material and methods. The difference in score between the two tumour models is significant for the anti-human uPA staining (p = 0.008), but not for the anti-mouse uPA staining.