Development of [18F]AmBF3 Tetrazine for Radiolabeling of Peptides:

Preclinical Evaluation and PET Imaging of [¹⁸F]AmBF₃-PEG₇-Tyr³-Octreotide in AR42J Pancreatic Carcinoma Model

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Table of contents

Experimental procedures
Figure S1. ¹ H NMR of AmBF ₃ -Tz (6)
Figure S2. HPLC-DAD of AmBF ₃ -Tz (6) 11
Figure S3. ¹⁹ F NMR of AmBF ₃ -Tz (6) 12
Figure S4. ¹¹ B NMR of AmBF ₃ -Tz (6)
Figure S5. ¹³ C NMR of AmBF ₃ -Tz (6) 14
Figure S6. UHPLC-HRMS ESI(+) extracted ion chromatogram and mass spectrum of AmBF ₃ -Tz (6)
Figure S7. ¹ H NMR of TCO-CHO (9)
Figure S8. ¹³ C NMR of TCO-CHO (9)
Figure S9. HPLC-DAD of TCO-CHO (9)
Figure S10. HPLC-DAD of TCO-functionalized octreotide (TCO-PEG ₄ -TOC, 12)
Figure S11. HPLC-DAD of TCO-functionalized octreotide (TCO-PEG ₇ -TOC, 13)
Figure S12. UHPLC-HRMS ESI(+) mass spectrum of TCO-PEG ₇ -TOC (13) 20
Figure S13. LC-MS spectrum of AmBF ₃ -PEG ₄ -TOC (14)
Figure S14. UHPLC-HRMS ESI(+) extracted ion chromatograms (EICs) and ESI MS spectra of AmBF ₃ -PEG ₇ -TOC (15)
Figure S15 . The influence of temperature and reaction volume on the radiochemical yield (RCY%) of [¹⁸ F]AmBF ₃ -Tz ([¹⁸ F]6)
Figure S16. Radio-TLC of [¹⁸ F]AmBF ₃ -Tz ([¹⁸ F]6)
Figure S17. Radio-HPLC chromatogram of [¹⁸ F]AmBF ₃ -PEG ₄ -TOC ([¹⁸ F]14)
Figure S18. Radio-DAD-HPLC chromatogram of [¹⁸ F]AmBF ₃ -PEG ₇ -TOC ([¹⁸ F]15)25
Figure S19 . Hydrolytic stability of [¹⁸ F]AmBF ₃ -PEG ₄ -TOC ([¹⁸ F]14) at 9 hours in injection formulation
Figure S20 . Hydrolytic stability of [¹⁸ F]AmBF ₃ -PEG ₇ -TOC ([¹⁸ F]15) up to ~6 hours in injection formulation. 27
Injection formulation of [18F]15 comprised of 4% ethanol in 0.01 M phosphate-buffered saline 27
Figure S21. Cell-uptake of [¹⁸ F]AmBF ₃ -PEG ₄ -TOC ([¹⁸ F]14) in AR42J cells
Figure S22. Cell-uptake of [¹⁸ F]AmBF ₃ -PEG ₇ -TOC ([¹⁸ F]15) in AR42J cells
Figure S23 . <i>Ex vivo</i> biodistribution (A , B) and elimination during PET/CT imaging (C) in selected organs and tissue samples of [¹⁸ F]AmBF ₃ -Tz ([¹⁸F]6)
Figure S24 . <i>Ex vivo</i> stability of [¹⁸ F]AmBF ₃ -PEG ₄ -TOC ([¹⁸F]14) in AR42J tumor bearing mouse blood (Radio-HPLC)
Figure S25. Ex vivo stability of [¹⁸ F]AmBF ₃ -PEG ₄ -TOC ([¹⁸ F]14) in urine (Radio-HPLC)
Figure S26 . <i>Ex vivo</i> stability of [¹⁸ F]AmBF ₃ -PEG ₄ -TOC ([¹⁸F]14) in AR42J tumor bearing mouse blood 5 and 30 minutes post-injection (Radio-TLC)

Table S1. Ex vivo blood component distribution analysis from mouse blood after intravenous administration of [18F]14	
Table S2. Biodistribution and statistical analysis of [18F]14 and [18F]15 at 60 minutes after i. v.administration expressed as %ID/g.34Figure S27. Ex vivo biodistribution of [18F]AmBF ₃ -PEG ₄ -TOC ([18F]14) (ex vivo; 60 minutes post-injection and 30, 60 and 120 min post-injection).35Figure S28. Ex vivo biodistribution of [18F]AmBF ₃ -PEG ₇ -TOC ([18F]15) (ex vivo; at 30 min, 60 min and 120 min post-injection, ex vivo after PET/CT; 270 min).36Figure S29. Tumor-to-muscle (T/M) and tumor-to-blood (T/B) ratios of [18F]14 after ex vivo biodistribution ($n \ge 4$).37Figure S30. Tumor-to-muscle (T/M) and tumor-to-blood (T/B) ratios of [18F]AmBF ₃ -PEG ₇ -TOC ([18F]15) after ex vivo biodistribution ($n \ge 2$).38Figure S31. Tumor-to-muscle (T/M) and tumor-to-blood (T/B) ratios of [18F]AmBF ₃ -PEG ₇ -TOC ([18F]15) during PET/CT imaging.39Table S3. Dosimetry calculations.40Figure S34. Standalone single-slice blank-on-white PET images of [18F]6 in a male SCID mouse.43Figure S35. Standalone single-slice blank-on-white PET images of [18F]6 in a healthy female C57BL/6JRj mouse.43Figure S37. Standalone single-slice blank-on-white PET images of [18F]15 in baseline conditions studied in a Rj:NMRI-Foxn1 ^{nu/nu} mouse bearing subcutaneous rat pancreatic AR42J xenografts.45Figure S38. Standalone single-slice blank-on-white PET images of [18F]15 in blocking conditions studied in a Rj:NMRI-Foxn1 ^{nu/nu} mouse bearing subcutaneous rat pancreatic AR42J xenografts.45Figure S38. Standalone single-slice blank-on-white PET images of [18F]15 in blocking conditions studied in Rj:NMRI-Foxn1 ^{nu/nu} mouse bearing subcutaneous rat pancreatic AR42J xenografts.45Figure S38. Standalone single-slice b	Table S1. Ex vivo blood component distribution analysis from mouse blood after intravenous administration of [18F]14. 33
Figure S27. Ex vivo biodistribution of [18F]AmBF3-PEG4-TOC ([18F]14) (ex vivo; 60 minutes post- injection and 30, 60 and 120 min post-injection).35Figure S28. Ex vivo biodistribution of [18F]AmBF3-PEG7-TOC ([18F]15) (ex vivo; at 30 min, 60 min and 120 min post-injection, ex vivo after PET/CT; 270 min).36Figure S29. Tumor-to-muscle (T/M) and tumor-to-blood (T/B) ratios of [18F]14 after ex vivo biodistribution ($n \ge 4$).37Figure S30. Tumor-to-muscle (T/M) and tumor-to-blood (T/B) ratios of [18F]AmBF3-PEG7-TOC ([18F]15) after ex vivo biodistribution ($n \ge 2$).38Figure S31. Tumor-to-muscle (T/M) and tumor-to-blood (T/B) ratios of [18F]AmBF3-PEG7-TOC ([18F]15) during PET/CT imaging.39Table S3. Dosimetry calculations.40Figure S34. Standalone single-slice blank-on-white PET images of [18F]6 in a male SCID mouse.Mouse.43Figure S36. Single-slice coronal PET/CT images of [18F]6 in (A) male SCID and (B) healthy female C57BL/6JRj mouse.44Figure S38. Standalone single-slice blank-on-white PET images of [18F]15 in baseline conditions studied in a Rj:NMRI-Foxn1 ^{nu/nu} mouse bearing subcutaneous rat pancreatic AR42J xenografts.57BL/6JRj mice.45Figure S38. Standalone single-slice blank-on-white PET images of [18F]15 in blocking conditions studied in a Rj:NMRI-Foxn1 ^{nu/nu} mouse bearing subcutaneous rat pancreatic AR42J xenografts.57BL/6JRj mice.45Figure S38. Standalone single-slice blank-on-white PET images of [18F]15 in blocking conditions studied in a Rj:NMRI-Foxn1 ⁿ	Table S2. Biodistribution and statistical analysis of [18F]14 and [18F]15 at 60 minutes after i. v.administration expressed as %ID/g.34
Figure S28. <i>Ex vivo</i> biodistribution of [¹⁸ F]AmBF ₃ -PEG ₇ -TOC ([¹⁸ F]15) (<i>ex vivo</i> ; at 30 min, 60 min and 120 min post-injection, <i>ex vivo</i> after PET/CT; 270 min)	Figure S27 . <i>Ex vivo</i> biodistribution of [¹⁸ F]AmBF ₃ -PEG ₄ -TOC ([¹⁸F]14) (<i>ex vivo</i> ; 60 minutes post-injection and 30, 60 and 120 min post-injection)
Figure S29. Tumor-to-muscle (T/M) and tumor-to-blood (T/B) ratios of [¹⁸ F]14 after <i>ex vivo</i> biodistribution (n ≥ 4)	Figure S28 . <i>Ex vivo</i> biodistribution of [¹⁸ F]AmBF ₃ -PEG ₇ -TOC ([¹⁸F]15) (<i>ex vivo</i> ; at 30 min, 60 min and 120 min post-injection, <i>ex vivo</i> after PET/CT; 270 min)
Figure S30. Tumor-to-muscle (T/M) and tumor-to-blood (T/B) ratios of [¹⁸ F]AmBF ₃ -PEG ₇ -TOC ([¹⁸ F]15) after <i>ex vivo</i> biodistribution (<i>n</i> ≥ 2)	Figure S29 . Tumor-to-muscle (T/M) and tumor-to-blood (T/B) ratios of [¹⁸ F]14 after <i>ex vivo</i> biodistribution ($n \ge 4$)
Figure S31. Tumor-to-muscle (T/M) and tumor-to-blood (T/B) ratios of [¹⁸ F]AmBF ₃ -PEG ₇ -TOC 39 Table S3. Dosimetry calculations. 40 Figure S34. Standalone single-slice blank-on-white PET images of [¹⁸ F]6 in a male SCID 43 Figure S35. Standalone single-slice blank-on-white PET images of [¹⁸ F]6 in a healthy female 43 Figure S36. Single-slice coronal PET/CT images of [¹⁸ F]6 in (A) male SCID and (B) healthy 44 Figure S37. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in baseline conditions 44 Figure S37. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in baseline conditions 45 Figure S37. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in baseline conditions 45 Figure S38. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in blocking conditions 45 Figure S38. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in blocking conditions 45 Figure S38. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in blocking conditions 45 Figure S39. Single-slice coronal PET/CT images of [¹⁸ F]15 in blocking conditions 45 Figure S39. Single-slice coronal PET/CT images of [¹⁸ F]15 in baseline (A) and blocking (B) 60 conditions studied in Rj:NMRI-Foxn1 ^{nu/nu} mice bearing subcutaneous rat pancreatic AR42J xenografts. 46	Figure S30 . Tumor-to-muscle (T/M) and tumor-to-blood (T/B) ratios of [¹⁸ F]AmBF ₃ -PEG ₇ -TOC ([¹⁸ F]15) after <i>ex vivo</i> biodistribution ($n \ge 2$)
Table S3. Dosimetry calculations. 40 Figure S34. Standalone single-slice blank-on-white PET images of [¹⁸ F]6 in a male SCID 43 Figure S35. Standalone single-slice blank-on-white PET images of [¹⁸ F]6 in a healthy female 43 Figure S36. Single-slice coronal PET/CT images of [¹⁸ F]6 in (A) male SCID and (B) healthy 44 Figure S37. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in baseline conditions 44 Figure S37. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in baseline conditions 45 Figure S38. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in blocking conditions 45 Figure S38. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in blocking conditions 45 Figure S38. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in blocking conditions 45 Figure S38. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in blocking conditions 45 Figure S39. Single-slice coronal PET/CT images of [¹⁸ F]15 in baseline (A) and blocking (B) 6 conditions studied in Rj:NMRI-Foxn1 ^{nu/nu} mice bearing subcutaneous rat pancreatic AR42J xenografts. 46	Figure S31 . Tumor-to-muscle (T/M) and tumor-to-blood (T/B) ratios of [¹⁸ F]AmBF ₃ -PEG ₇ -TOC ([¹⁸ F]15) during PET/CT imaging
Figure S34. Standalone single-slice blank-on-white PET images of [¹⁸ F]6 in a male SCID 43 Figure S35. Standalone single-slice blank-on-white PET images of [¹⁸ F]6 in a healthy female 43 Figure S36. Single-slice coronal PET/CT images of [¹⁸ F]6 in (A) male SCID and (B) healthy 43 Figure S37. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in baseline conditions 44 Figure S37. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in baseline conditions 45 Figure S38. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in blocking conditions 45 Figure S38. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in blocking conditions 45 Figure S38. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in blocking conditions 45 Figure S38. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in blocking conditions 45 Figure S38. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in blocking conditions 45 Figure S39. Single-slice coronal PET/CT images of [¹⁸ F]15 in baseline (A) and blocking (B) 6 conditions studied in Rj:NMRI-Foxn1 ^{nu/nu} mice bearing subcutaneous rat pancreatic AR42J 46	Table S3. Dosimetry calculations. 40
Figure S35. Standalone single-slice blank-on-white PET images of [¹⁸ F]6 in a healthy female 43 Figure S36. Single-slice coronal PET/CT images of [¹⁸ F]6 in (A) male SCID and (B) healthy 44 Figure S37. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in baseline conditions 44 Figure S37. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in baseline conditions 45 Figure S38. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in blocking conditions 45 Figure S38. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in blocking conditions 45 Figure S38. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in blocking conditions 45 Figure S38. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in blocking conditions 45 Figure S39. Single-slice coronal PET/CT images of [¹⁸ F]15 in blocking (B) 6 conditions studied in Rj:NMRI-Foxn1 ^{nu/nu} mice bearing subcutaneous rat pancreatic AR42J xenografts. 46	Figure S34. Standalone single-slice blank-on-white PET images of [¹⁸ F]6 in a male SCID mouse
Figure S36. Single-slice coronal PET/CT images of [¹⁸ F]6 in (A) male SCID and (B) healthy female C57BL/6JRj mice. 44 Figure S37. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in baseline conditions 44 Figure S37. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in baseline conditions 44 Figure S38. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in blocking conditions 45 Figure S38. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in blocking conditions 45 Figure S38. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in blocking conditions 45 Figure S38. Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in blocking conditions 45 Figure S39. Single-slice coronal PET/CT images of [¹⁸ F]15 in baseline (A) and blocking (B) 6 conditions studied in Rj:NMRI-Foxn1 ^{nu/nu} mice bearing subcutaneous rat pancreatic AR42J 46	Figure S35 . Standalone single-slice blank-on-white PET images of [¹⁸ F]6 in a healthy female C57BL/6JRj mouse
 Figure S37. Standalone single-slice blank-on-white PET images of [¹⁸F]15 in baseline conditions studied in a Rj:NMRI-Foxn1^{nu/nu} mouse bearing subcutaneous rat pancreatic AR42J xenografts. Figure S38. Standalone single-slice blank-on-white PET images of [¹⁸F]15 in blocking conditions studied in aRj:NMRI-Foxn1^{nu/nu} mouse bearing subcutaneous rat pancreatic AR42J xenografts.45 Figure S39. Single-slice coronal PET/CT images of [¹⁸F]15 in baseline (A) and blocking (B) conditions studied in Rj:NMRI-Foxn1^{nu/nu} mice bearing subcutaneous rat pancreatic AR42J xenografts.46 	Figure S36. Single-slice coronal PET/CT images of [¹⁸ F]6 in (A) male SCID and (B) healthy female C57BL/6JRj mice
Figure S38 . Standalone single-slice blank-on-white PET images of [¹⁸ F] 15 in blocking conditions studied in aRj:NMRI-Foxn1 ^{nu/nu} mouse bearing subcutaneous rat pancreatic AR42J xenografts.45 Figure S39 . Single-slice coronal PET/CT images of [¹⁸ F] 15 in baseline (A) and blocking (B) conditions studied in Rj:NMRI-Foxn1 ^{nu/nu} mice bearing subcutaneous rat pancreatic AR42J xenografts.	Figure S37 . Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in baseline conditions studied in a Rj:NMRI-Foxn1 ^{nu/nu} mouse bearing subcutaneous rat pancreatic AR42J xenografts.
Figure S39 . Single-slice coronal PET/CT images of [¹⁸ F]15 in baseline (A) and blocking (B) conditions studied in Rj:NMRI-Foxn1 ^{nu/nu} mice bearing subcutaneous rat pancreatic AR42J xenografts.	Figure S38 . Standalone single-slice blank-on-white PET images of [¹⁸ F]15 in blocking conditions studied in aRj:NMRI-Foxn1 ^{nu/nu} mouse bearing subcutaneous rat pancreatic AR42J xenografts.45
	Figure S39 . Single-slice coronal PET/CT images of [¹⁸ F]15 in baseline (A) and blocking (B) conditions studied in Rj:NMRI-Foxn1 ^{nu/nu} mice bearing subcutaneous rat pancreatic AR42J xenografts

Experimental procedures

PET/CT imaging

The standardized uptake values (SUVs) were determined from the PET image by drawing regions-of-interest (ROI) around selected organs (heart, liver, kidney, lung, muscle, urinary bladder) with Carimas software (version 2.10, Turku PET Centre, Turku, Finland) and by measuring the ratio of radioactivity per unit volume of that ROI and normalized to the injected dose. Compound [¹⁸F]6 was imaged with Inveon[®] PET/CT (Siemens) and compound [¹⁸F]15 with Molecubes PET (β-CUBE) coupled with a CT (X-CUBE) (MOLECUBES NV, Ghent, Belgium).

Liquid chromatography and mass spectrometry methods

HPLC analysis (Radiodetectotion/Diode-array detection): Shimadzu HPLC system containing a DGU-20A degasser, an LC-20AD UPLC LC unit, a SIL-20A HT autosampler, a CTO-20AC column oven, a CBM-20A communications bus module, a Scionix Holland scintillation detector with a 51 BP 51/2 Nal(TI) crystal and an SPD-M20A diode array detector.

UHPLC-HRMS analysis: The compounds **6**, **13** and **15** were analysed by UHPLC Thermo Scientific Dionex Ultimate 3000 ultrahigh performance liquid chromatography (Germering, Germany) coupled to Thermo Scientific Orbitrap Fusion mass spectrometer (San Jose, CA, USA). Ionization (HESI+), scan range of 120–1200 *m/z*. Data was processed with Xcalibur workstation (Thermo Fisher Scientific, Waltham, MA, USA). Compounds **6** and **13** were analysed for acquiring their exact masses. Compound **15** was analyzed for acquiring its exact mass and isotopic peak patterns.

LC-MS analysis: Compounds **12** and **14** were analyzed with Agilent Technologies 1260 Infinity HPLC-DAD system with Agilent Technologies 6120 Quadruple LC/MS detector.

S6

Ionization HESI⁺, scan range 100-2000 *m/z*. Data was processed with OpenLAB CDS

Workstation.

Liquid chromatography (LC) method gradients:

Method A (HPLC and radio-HPLC): (Shimadzu), Column: Phenomenex Kinetex[®] 5 μ m C18 100 Å, LC Column 250 × 10.0 mm

Eluents: (A) H₂O+0.1% TFA and (B) ACN+0.1% TFA

Gradient 3 mL/min at ambient temperature:

<u>%B</u>
20
50
20
20
20

Method B (HPLC): (Shimadzu), Column: HiChrom (Alltech) Alltima[®] 5 μm C18 100 Å, LC Column 250 × 10.0 mm

Eluents: (A) H₂O+0.1% TFA and (B) ACN+0.1% TFA

Gradient 3 mL/min at ambient temperature: Isocratic

<u>Time (min)</u>	<u>%B</u>
0	80
32	80

Method C (LC-MS): (Agilent 1200 Series), Column: Waters Atlantis® T3 3 μ m C18 100 Å, LC Column 4.6 × 150 mm

Method 1C) for compound **9** ($t_R = 11.5 \text{ min}$)

A: 0.1 % TFA in milli-Q water B: 0.1 % TFA in ACN Flowrate: 0.7 mL/min

<u>Time (min)</u>	<u>%B</u>
0	50
15	90
26	10

Method 2C) for compounds **12** ($t_R = 12.8 \text{ min}$), **14** ($t_R = 11.6 \text{ min}$) and **15** ($t_R = 15.2 \text{ min}$) (Agilent **1200 Series**)

A: 0.1 % FA in milli-Q water B: 0.1 % FA in ACN Flowrate: 0.7 mL/min

<u>Time (min)</u>	<u>%B</u>
0	10
33	90
35	90
40	10

Method D (UHPLC-HRMS): (Orbitrap Fusion)

Column: Waters ACQUITY UPLC® 1.7 μm BEH C18 130Å, UPLC Column 2.1 × 50 mm Eluents: (A) H₂O+0.1% FA and (B) ACN+0.1% FA

Gradient 0.5 mL/min at 40 °C:

Scan range: 120-1200 *m*/*z* for compound **6** and 100-2000 *m*/*z* for compounds **13** and **15**.

For compound 6 and 15:

<u>Time (min)</u>	<u>%B</u>
0	5
3	5
4	25
5	60
6	80
7	100
7.5	5
9	5

Lipophilicity. The lipophilicities were determined with the shake-flask method as a distribution profile between 0.01 M PBS and octanol at pH 7.4. The distribution of the radiolabeled compound between the octanol and the PBS layers was calculated according to equation:

$$Log D_{7.4} = \frac{Ac_{octanol}}{Ac_{PBS}}$$

where Ac_{oct} =radioactivity concentration of octanol and Ac_{PBS} =radioactivity concentration of PBS. The datapoints were presented as mean ± standard deviation.

AR42J cell-uptake assay and mouse tumor model. Rat pancreatic tumor cell line AR42J that expresses SSTR2, was cultured at 37 °C in 95% relative humidity incubator containing 5% CO₂. Cells grown to >90% confluence were used for either *in vitro* or *in vivo* experiments. One million cells/well were seeded overnight on 6-well plates. The growth media was removed and the medium (1 mL/well) containing [18F]14 (11 kBq, 10 pmol per well) or [¹⁸F]15 (97 kBg, 40 pmol per well) was added. For studying the specificity of the cell-uptake, a set of cells were co-incubated in the presence of 2.4 nmol per well of non-modified octreotide representative of blocking conditions (total volume ~1 mL). For determining the amount of the radiotracer in the free fraction, at the designated time-points (15, 30, 60 and 120 min) the incubated media were removed, collected to a 5 mL scintillation tube, cells were washed with 1 mL of cold 0.01 M PBS (1×PBS) and again the supernatant collected into the same scintillation tube. Membrane-bound fraction; ice-cold glycine buffer (1 mL) was added on the cells, incubated for 5 minutes on ice, supernatant removed (procedure repeated once more) and the cells washed with ice-cold 1xPBS, and all washing solutions were collected to the same scintillation tube. Internalized fraction; 1 M NaOH (1 mL) was added on the cells and left incubating at an ambient temperature for 10 min. The supernatant was removed, the wells were washed 2 × 1 mL of cold 0.01 M PBS, and the supernatants collected to the same microtube. The collected samples were measured with a gamma counter for measuring the counts per minute (CPM) of each fraction (n = 3). The internalization is presented in percent (%) normalized to the total radioactivity in all three fractions.



Figure S1. 1 H NMR of AmBF₃-Tz (6).



Figure S2. HPLC-DAD of AmBF₃-Tz (**6**). Purity \geq 99%.



Figure S3. ¹⁹F NMR of AmBF₃-Tz (6).

The signal as quartet demonstrates ¹⁹F-¹¹B coupling with characteristic ratios (1:1:1:1).



Figure S4. ¹¹B NMR of AmBF₃-Tz (**6**).

The signal as quartet demonstrates ¹¹B-¹⁹F coupling with characteristic ratios (1:3:3:1).



Figure S5. ¹³C NMR of AmBF₃-Tz (6).



Figure S6. UHPLC-HRMS ESI(+) extracted ion chromatogram and mass spectrum of AmBF₃-Tz (**6**).

6 detected as a protonated molecule ([M+H]⁺ found m/z 369.18134, mass error Δ =-0.85454 ppm, calculated m/z 369.18165 for C₁₅H₂₁N₆BF₃O⁺



Figure S7. ¹H NMR of TCO-CHO (9).



Figure S8. ¹³C NMR of TCO-CHO (9).



Figure S9. HPLC-DAD of TCO-CHO (9).

Retention time of **9** t_R = 7.8 min. Residual pyridine peak detected at 3.8 min. Purity ≥98%.



Figure S10. HPLC-DAD of TCO-functionalized octreotide (TCO-PEG₄-TOC, **12**) Retention time of **12** t_R = 19.1 min.



Figure S11. HPLC-DAD of TCO-functionalized octreotide (TCO-PEG₇-TOC, **13**) Retention time of **13** t_R = 24.1 min.



Figure S12. UHPLC-HRMS ESI(+) mass spectrum of TCO-PEG₇-TOC (13).

Compound **13** eluted at $t_R = 5.22$ min, detected as two protonated molecule ions: $[M+3H]^{3+}$ at 586,61316 *m/z* with $\Delta = -2,35545$ ppm (calculated 586,61454 *m/z* for C₈₅H₁₂₅O₂₃N₁₃S₂³⁺), and $[M+2H]^{2+}$ at 879,41663 *m/z* with $\Delta = -1,75922$ ppm (calculated 879,41817 *m/z* for C₈₅H₁₂₄O₂₃N₁₃S₂²⁺).

Peak RT : 11.644 min



Figure S13. LC-MS spectrum of AmBF₃-PEG₄-TOC (14).

Retention time of **14** $t_R = 11.644$ min. **14** was detected as molecular peaks corresponding to three charged states (Z=+1 - Z=+3) [M+H]⁺ (found *m/z* 1907.6, calculated *m/z* 1907.9 for C₉₂H₁₂₅BF₃N₁₆O₂₀S₂⁺), [M+2H]²⁺ (found *m/z* 953.9, calculated *m/z* 953.9 for C₉₂H₁₂₆BF₃N₁₆O₂₀S₂²⁺) and [M+3H]³⁺ (found *m/z* 636.6, calculated *m/z* 636.6 for C₉₂H₁₂₇BF₃N₁₆O₂₀S₂³⁺).



Figure S14. UHPLC-HRMS ESI(+) extracted ion chromatograms (EICs) and ESI MS spectra of AmBF₃-PEG₇-TOC (**15**).

Compound **15** was detected as well resolved isotopic molecular ion peaks, with a pattern demonstrating mass differences of 0.33 and 0.50 *m/z* for the detected molecular peaks, indicative of charged state at Z=+3 for 0.33, and a charged state of Z=+2 for 0.50 mass difference. The exact monoisotopic masses for C₁₀₀H₁₄₁BF₃O₂₄N₁₇S₂²⁺ [M+2H]²⁺ was found at *m/z* 1047.99207 (Δ = 1.44731 ppm, calculated *m/z* 1047.99055) and of C₁₀₀H₁₄₂BF₃O₂₄N₁₇S₂³⁺ for [M+3H]³⁺, was found at *m/z* 698.99744 (Δ = 1.87679 ppm, calculated *m/z* 698.99612), correlated with the calculated masses of compound **15**.



Figure S15. The influence of temperature and reaction volume on the radiochemical yield (RCY%) of [¹⁸F]AmBF₃-Tz (**[¹⁸F]6**). Reaction with 0.9% NaCl eluted [¹⁸F]fluoride.



Figure S16. Radio-TLC of [¹⁸F]AmBF₃-Tz (**[**¹⁸**F]6**). RCP ≥98%.



Figure S17. Radio-HPLC chromatogram of [¹⁸F]AmBF₃-PEG₄-TOC ([¹⁸F]14). Retention time of **14** t_R = 18.7 min (mV).



Figure S18. Radio-DAD-HPLC chromatogram of [¹⁸F]AmBF₃-PEG₇-TOC ([¹⁸F]15). Retention time of **15** t_R = 17.3 min (mV).



Figure S19. Hydrolytic stability of [¹⁸F]AmBF₃-PEG₄-TOC ([¹⁸F]14) at 9 hours in injection formulation.

Injection formulation of [¹⁸F]14 comprised of 4% ethanol in 0.01 M phosphate-buffered saline with a retention time of $t_R = 18.7$ min



Figure S20. Hydrolytic stability of $[^{18}F]$ AmBF₃-PEG₇-TOC ($[^{18}F]$ 15) up to ~6 hours in injection formulation.

Injection formulation of [¹⁸F]15 comprised of 4% ethanol in 0.01 M phosphate-buffered saline.





Cell-uptake (% internalized) of [¹⁸F]14 demonstrating specific uptake of the radiotracer in AR42J tumor cells. (*p < 0.05, ** $p \le 0.01$, ns= not statistically significant, n = 3). The datapoints are presented as mean ± standard deviation.



Figure S22. Cell-uptake of [¹⁸F]AmBF₃-PEG₇-TOC ([¹⁸F]15) in AR42J cells.

Cell-uptake (% internalized) of [¹⁸F]15 demonstrating specific uptake of the radiotracer in AR42J tumor cells (*** p < 0.001, **** $p \le 0.0001$, n = 3). The datapoints are presented as mean ± standard deviation.



Figure S23. *Ex vivo* biodistribution (**A**, **B**) and elimination during PET/CT imaging (**C**) in selected organs and tissue samples of [¹⁸F]AmBF₃-Tz (**J**¹⁸F]6).

Ex vivo in healthy C57BL/6JRj (control) and SCID mice (control) at t = 270 minutes after intravenous administration of [¹⁸F]6 in healthy SCID mice (control) after intravenous administration (**A**: biodistribution, **B**: elimination to urine and feces from *ex vivo* biodistribution samples **C**: SUV in selected organs during PET/CT acquisition,). G.B. = gallbladder; S.I. = small intestine; LI. = large intestine. TACs from PET/CT imaging presented as SUVs. The datapoints are presented as mean ± standard deviation (n = 4).



Figure S24. Ex vivo stability of [18F]AmBF₃-PEG₄-TOC ([18F]14) in AR42J tumor

bearing mouse blood (Radio-HPLC).

HPLC chromatogram of a sample extracted from mouse blood (*ex vivo*). The radiopeak corresponding to intact [¹⁸F]14 was detected at a retention time of t_R = 18.6 min.



Figure S25. Ex vivo stability of [18F]AmBF₃-PEG₄-TOC ([18F]14) in urine (Radio-

HPLC).

HPLC chromatogram of the tracer extracted from mouse urine (*ex vivo*). The radiopeak corresponding to intact [¹⁸F]14 was detected at a retention time of t_R = 18.6 min.



Figure S26. Ex vivo stability of [¹⁸F]AmBF₃-PEG₄-TOC ([¹⁸F]14) in AR42J tumor

bearing mouse blood 5 and 30 minutes post-injection (Radio-TLC).

Radio-TLC revealed a slightly polar metabolite (4%) in blood 5 minutes post-injection with a retardation factor (R_f) of R_f = 40 mm. At 30 minutes post-injection the radiometabolite at 40 mm had disappeared and instead a highly polar metabolite (2.8%) at 4.0 mm was visible, and was accompanied by another metabolite (2.0%) eluting in the immediate vicinity of [¹⁸F]14 at R_f = 58 mm. At 60 minutes post-injection the tracer accounted for ≥99% of the radioactivity both in blood and urine (*ex vivo*).

 Table S1. Ex vivo blood component distribution analysis from mouse blood after intravenous administration of [18F]14.

	Red blood	d cells	Protein bound		Free fraction	
	Average (%)	Stdv.(%)	Average (%)	Stdv.(%)	Average (%)	Stdv.(%)
5 min	21.6	4.4	5.5	0.9	72.9	5.1
60 min	24.5	5.4	7.00	1.5	68.5	5.3
60 min	29.7	2.9	14.6	1.4	55.7	11.4
(B)						

B=blocked with octreotide, (n = 2).

	[¹⁸ F]14	[¹⁸ F]14	Statistical	[¹⁸ F]15	[¹⁸ F]15	Statistical
	Baseline	Blocked	significance	Baseline	Blocked	significance
	(/@D/g)	(/@D/9)	(p-value)	(/@D/g)	(/@D/9)	(p-value)
			+ (0, 0, 1, 1, 0)			
Urine	84.9 ±	41.9 ±	* (0.0449)	174.2 ± 73.4	303.4 ±	ns (0.2654)
	58.6	15.7			145.8	
Blood	10.2 ± 2.3	7.4 ± 1.5	** (0.0079)	9.6 ± 3.3	7.2 ± 2.5	ns (0.2918)
Spleen	1.2 ± 0.5	0.9 ± 0.1	* (0.0473)	1.4 ± 0.3	0.9 ± 0.2	* (0.0375)
Pancreas	1.1 ± 0.2	0.8 ± 0.2	** (0.0087)	1.6 ± 0.5	0.9 ± 0.5	ns (0.1655)
Kidney	14.3 ± 4.5	11.4 ± 3.7	ns (0.1806)	21.0 ± 4.5	15.9 ± 2.7	ns (0.0919)
Gallbladder	17.8 ± 5.2	10.7 ± 6.3	ns (0.0530)	9.1 ± 7.0	18.8 ± 10.3	ns (0.3923)
Liver	30.0 ± 9.4	26.2 ±	ns (0.6075)	19.0 ± 5.4	11.8 ± 4.7	ns (0.1032)
		15.8				
Lung	6.2 ± 2.4	5.5 ± 2.7	ns (0.5991)	5.4 ± 2.3	3.6 ± 1.4	ns (0.1948)
Heart	2.4 ± 0.7	1.9 ± 0.4	ns (0.0565)	2.6 ± 0.6	2.1 ± 0.5	ns (0.2824)
Muscle	0.6 ± 0.2	0.5 ± 0.1	* (0.0404)	1.1 ± 0.9	0.5 ± 0.1	ns (0.2078)
Bone (t.)	1.3 ± 0.6	1.0 ± 0.6	ns (0.2563)	1.1 ± 0.4	0.8 ± 0.2	ns (0. 1095)
Bone (occ.)	1.3 ± 0.6	1.1 ± 0.5	ns (0.5751)	0.9 ± 0.3	1.0 ± 0.3	ns (0.7031)
Brain	0.3 ± 0.1	0.3 ± 0.1	ns (0.3096)	0.3 ± 0.1	0.2 ± 0.1	ns (0.0655)
Stomach	0.6 ± 0.2	0.6 ± 0.2	ns (0.7522)	1.4 ± 0.8	0.6 ± 0.1	ns (0.1197)
(+cont.)						
S. I.	2.6 ± 0.8	1.9 ± 0.4	* (0.0266)	3.6 ± 2.1	5.0 ± 1.7	ns (0.3371)
(+cont.)						
L. I.	1.1 ± 1.1	0.7 ± 0.3	ns (0.2071)	1.0 ± 0.7	0.7 ± 0.3	ns (0.3750)
(+cont.)						
Skin	2.1 ± 0.9	1.6 ± 0.4	ns (0.1348)	2.4 ± 0.9	1.6 ± 0.2	ns (0.1041)
Tumor	3.1 ± 0.7	2.2 ± 0.6	* (0.0120)	4.5 ± 1.0	3.1 ± 0.5	* (0.0143)
				1		

Table S2. Biodistribution and statistical analysis of [¹⁸F]14 and [¹⁸F]15 at 60 minutes after i. v. administration expressed as %ID/g.

ns = not significant, *p<0.05, **p<0.01, and ***p<0.001. Differenced between groups were considered significant when $p \le 0.05$.

t.; tibia, occ.; occipital, cont.; content, S.I.; small intestines, L.I.; large intestine. Unpaired *t*-test with Welch's correction was used to calculate statistical significances (significant p<0.05). The datapoints are presented as mean ± standard deviation.





Biodistribution of radioactivity after intravenous administration of [¹⁸F]AmBF₃-PEG₄-TOC ([¹⁸F]14) in AR42J tumor bearing Rj:NMRI-*Foxn1*^{nu/nu} mice. (*p < 0.05). S.I. = small intestine; L.I. = large intestine, occip. = occipital, cont. = contents, B = blocked. The datapoints are presented as mean ± standard deviation.





Biodistribution of radioactivity in AR42J rat pancreatic tumor bearing Rj:NMRI-Foxn1^{nu/nu} mice at t = 60 min after intravenous administration. The uptake was blocked using a dose (44 nmol) of octreotide that was co-administered with the radiotracer, and the mice were euthanized at selected time points. (*p < 0.05). S.I. = small intestine; L.I. = large intestine, cont. = contents, B = blocked. The datapoints are presented as mean ± standard deviation.



Figure S29. Tumor-to-muscle (T/M) and tumor-to-blood (T/B) ratios of [¹⁸F]14 after *ex vivo* biodistribution ($n \ge 4$).

The datapoints are presented as mean ± standard deviation.





The datapoints are presented as mean ± standard deviation.



Figure S31. Tumor-to-muscle (T/M) and tumor-to-blood (T/B) ratios of [¹⁸F]AmBF₃-PEG₇-TOC ([¹⁸F]15) during PET/CT imaging.

Blood radioactivity was determined by drawing a region-of-interest (ROI) for left ventricle of the heart (n = 2). The datapoints are presented as mean ± standard deviation.

Table S3. Dosimetry calculations.

Results after intravenous administration of [¹⁸F]15 from OLINDA/EXM (Version 2.1, Vanderbilt University, 2012), where ICRP 89 reference adult male (73 kg) and ICRP 103 radiation weighting factors were used. Absorbed doses to each target organ is given in units mGy/MBq and the effective dose in units mSv/MBq. The difference in radiation burden based on baseline (non-block) and blocked mice is negligible.

Target organ	Baseline Avg. (mGy/MBq)	Baseline Std. (mGy/MBq)	Blocked Avg. (mGy/MBq)	Blocked Std. (mGy/MBq)
Adrenals	0.0190	0.0008	0.0185	0.0001
Brain	0.0102	0.0003	0.0103	0.0000
Esophagus	0.0129	0.0004	0.0129	0.0002
Eyes	0.0102	0.0003	0.0103	0.0000
Gallbladder Wall	0.0194	0.0011	0.0190	0.0011
Left colon	0.0143	0.0001	0.0144	0.0001
Small Intestine	0.0144	0.0001	0.0145	0.0000
Stomach Wall	0.0141	0.0002	0.0141	0.0001
Right colon	0.0151	0.0001	0.0150	0.0001
Rectum	0.0136	0.0003	0.0137	0.0000
Heart Wall	0.0148	0.0008	0.0150	0.0010
Kidneys	0.0366	0.0016	0.0337	0.0043
Liver	0.0334	0.0050	0.0313	0.0040
Lungs	0.0109	0.0030	0.0106	0.0030
Pancreas	0.0156	0.0002	0.0154	0.0001
Prostate	0.0135	0.0003	0.0136	0.0000
Salivary Glands	0.0118	0.0002	0.0119	0.0000
Red Marrow	0.0113	0.0000	0.0114	0.0001
Osteogenic Cells	0.0120	0.0001	0.0121	0.0001
Spleen	0.0137	0.0001	0.0136	0.0001
Testes	0.0112	0.0003	0.0113	0.0000
Thymus	0.0122	0.0002	0.0122	0.0003
Thyroid	0.0118	0.0000	0.0119	0.0003
Urinary Bladder Wall	0.0134	0.0003	0.0135	0.0000
Total Body	0.0121	0.0001	0.0121	0.0000
Effective Dose (mSv/MBq)	0.0112	6.36E-04	0.011	2.83E-04



Figure S32. Absorbed doses (mGy/MBq) in selected tissues in AR42J tumor-bearing mice at baseline and blocking conditions (co-injection with 45 μg, 44 nmol octreotide) after intravenous administration of [¹⁸F]15 demonstrating highest absorbed doses to kidneys and liver. Individual values plotted as a single datapoint.



Figure S33. Maximum intensity projections of PET/CT at 20–80 min post-injection of $[^{18}F]AmBF_3$ -PEG₇-TOC (**[**¹⁸**F]15**) in AR42J tumor bearing mice. Blocked (co-injection with 45 µg, 44 nmol octreotide) on the left.



Figure S34. Standalone single-slice blank-on-white PET images of [¹⁸F]6 in a male SCID mouse.



Figure S35. Standalone single-slice blank-on-white PET images of [¹⁸F]6 in a healthy female C57BL/6JRj mouse.



Figure S36. Single-slice coronal PET/CT images of [¹⁸F]6 in (A) male SCID and

⁽B) healthy female C57BL/6JRj mice.



Figure S37. Standalone single-slice blank-on-white PET images of [¹⁸F]15 in baseline conditions studied in a Rj:NMRI-Foxn1^{nu/nu} mouse bearing subcutaneous rat pancreatic AR42J xenografts.



Figure S38. Standalone single-slice blank-on-white PET images of [¹⁸F]15 in blocking conditions studied in aRj:NMRI-Foxn1^{nu/nu} mouse bearing subcutaneous rat pancreatic AR42J xenografts.



Figure S39. Single-slice coronal PET/CT images of [¹⁸F]15 in baseline (**A**) and blocking (**B**) conditions studied in Rj:NMRI-Foxn1^{nu/nu} mice bearing subcutaneous rat pancreatic AR42J xenografts.