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Positron Emission Tomography Imaging of Drug-Induced Tumor Apoptosis with a Caspase-Triggered Nano-aggregation Probe**

Bin Shen[†], Jongho Jeon[†], Mikael Palner, Deju Ye, Adam Shuhendler, Frederick T. Chin^{*}, and Jianghong Rao^{*}

Department of Radiology, Molecular Imaging Program at Stanford, Stanford University School of Medicine, Stanford, CA 94305-5484, USA

Abstract

An ¹⁸F-labeled caspase-3 sensitive nano-aggregation PET tracer ([¹⁸F]CSNAT) was prepared and evaluated for imaging caspase-3 activity in doxorubicin-treated tumor xenografts. This enzyme-activatable PET tracer is designed to function in a novel mechanism – enhanced retention of the ¹⁸F activity in apoptotic tumors is achieved through intramolecular macrocyclization and *in situ* aggregation upon caspase-3 activation.

Keywords

PET; apoptosis; cyclization; caspase-3

Positron emission tomography (PET) is a powerful non-invasive molecular imaging technique that uses molecules labeled with positron-emitting isotopes such as carbon-11, fluorine-18, and copper-64, to monitor biochemical processes in living subject in real time with excellent sensitivity.^[1] A contrast-enhanced PET image is produced through selective retention of radioactivity at the location where the molecular target is present, and the common strategy is based on the binding of a radiolabeled ligand to the target receptor molecule. When the molecular target is an enzyme, the methods to generate PET imaging contrast are very few,^[2] such as 2-[¹⁸F]fluoro-2-deoxy-glucose ([¹⁸F]FDG) for imaging hexokinase^[3] and PET tracers for imaging herpes simplex virus-1 thymidine kinase.^[4] This communication reports a new strategy for developing PET tracers for imaging caspase-3 activity in tumors treated with doxorubicin.

One of the most important pathways of chemotherapeutic-induced apoptosis leads to the activation of caspase-3, a central scavenging peptidase that cleaves a specific peptide sequence DEVD-X between X and D (where X is any amino acid), committing the cell to

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jrao@stanford.edu, chinf@stanford.edu.

B. Shen and J. Jeon contributed equally to this work.

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programmed cell death—apoptosis.^[5] While activatable fluorescent probes have been developed for imaging caspase-3 activity in cells and living mice,^[6] there are only reports of radiolabeled caspase-3 inhibitors for PET imaging of caspase-3 in apoptotic cells.^[7] There isn't a general strategy to develop a PET tracer that is mechanistically similar to the activatable fluorescent probes in imaging caspase-3 activity with signal amplification.

We have previously described a biocompatible condensation reaction between 2-cyanobenzothiazole (CBT) and free cysteine for specific labeling of terminal cysteine residues in proteins^[8] and controlled self-assembly of nano-aggregates in cells.^[9] In this communication, we applied it to develop PET tracers for imaging caspase activity, as outlined in Scheme 1. The ¹⁸F-labeled caspase-sensitive nano-aggregation PET tracer ([¹⁸F]C-SNAT, or **1**) has a 2-cyano-6-hydroxyquinoline (CHQ) and a cysteine residue whose amino group is coupled to the peptide substrate of caspase-3, DEVD,^[10] and whose mercapto group of its side chain is converted to a disulfide bond. The ¹⁸F label is introduced to a propargylglycine residue between CHQ and the cysteine residue via the copper-catalyzed azide-alkyne 1,3-dipolar cycloaddition (CuAAC). Upon the activation of caspase-3 that cleaves the peptide DEVD, the amino group of the cysteine residue is released free, and the reductive intracellular environment reduces the disulfide to generate a free thiol. As shown before,^[11] free cysteine and CHQ can undergo fast intramolecular cyclization through the condensation reaction (half-life at 119.8±10.2s). The cyclized products are more hydrophobic and self-assemble *in situ* into nanoparticles with a high density of ¹⁸F activity, resulting in prolonged retention in apoptotic cells and enhanced PET imaging contrast.

The control tracer **1-D** is designed to have a similar structure and pharmacokinetics to **1** but cannot be cleaved by caspase-3 and undergo cyclization and aggregation. Therefore, it contains the same substrate sequence of caspase-3 but made in *D*-amino acids. In addition, **1-D** does not have the cyano substitution on its quinoline and the mercapto group of the cysteine side chain is methylated.

The precursor **3a** for synthesizing **1** was prepared according to Scheme 2. *N*-Boc-propargylglycine was coupled with 6-amino-2-cyanobenzothiazole to afford **4**. Deprotection of *N*-Boc group followed by coupling with protected *D*-cysteine provided the dipeptide **5**. Condensation of **5** and **6a** between the CBT group and cysteine under mild conditions, subsequent deprotection of *N*-Boc and *S*-Trt groups and protection of the thiol group gave intermediate **7a**. The amino group of **7a** was coupled with the protected caspase-3 substrate, Ac-Asp(^tBu)-Glu(^tBu)-Val-Asp(^tBu)-OH **8a**, and final deprotection of the *t*-Butyl group gave the desired precursor **3a**. The precursor **3b** for the control PET tracer **1-D** consisted of *D*-peptide amino acids, and was prepared similarly.

Both precursors were labeled with ¹⁸F through CuAAC, as reported before^[12] (Scheme 3). The prosthetic group **10** was obtained in a one-step substitution of **9** by ¹⁸F in the presence of Kryptofix 222 (4,7,13,16,21,24-Hexaoxa-1,10-diazabicyclo[8.8.8] hexacosane) with a radiochemical yield (RCY) of 37±8% (n=16) (decay corrected to end of bombardment, EOB). The CuAAC labeling of **3a** by **10** occurred in mild conditions (40–60 °C). When CuSO₄ and sodium ascorbate were employed, the RCY (EOB) in this step was 15±6% (n=6) for **1**, and 25±8% (n=4) for **1-D**. When Cu(CH₃)₄PF₆ and BPDS were applied, the RCY

(EOB) increased to $30\pm 5\%$ ($n=4$) for **1** with less side products, probably due to improved stability of **1** in the absence of sodium ascorbate. The final product was purified on a semi-preparative HPLC and formulated in saline containing $<10\%$ EtOH. During the HPLC purification of **1-D**, because both the precursor **3b** and product **1-D** had very close retention times ($t_R < 0.1$ min), 3-azidopropan-1-amine hydrochloride was added to convert remaining **3b** to a product with its retention time shifted to a much earlier time ($t_R > 10$ min), which significantly improved the HPLC purification (Figure S1). In summary, **1** and **1-D** were successfully obtained with an overall RCY (decay corrected to end of synthesis, EOS) and specific radioactivity (EOS) of $3.2\pm 0.1\%$ and 63 ± 7.4 GBq/ μmol (1.7 ± 0.2 Ci/ μmol) for **1**, $1.8\pm 0.8\%$ and 99.9 ± 33.3 GBq/ μmol (2.7 ± 0.9 Ci/ μmol) for **1-D** (radiochemical purity $> 99\%$, chemical purity $> 95\%$, Figure S2) after a total synthesis time of 3–3.5 h from EOB.

The serum stability of **1** and **1-D** was tested in human and mouse serum. Both tracers showed $> 90\%$ intact in human serum after 2 h incubation at 37°C . In mouse serum, **1-D** displayed higher stability than **1**: 90% of intact **1-D** vs. 60% of intact **1** was observed after 1 h incubation—probably due to the replacement by the *D*-amino acids. The cyclization product **2** was also tested and showed $>90\%$ intact in mouse serum after 2 h incubation (Figure S3).

To evaluate the tracer **1** for caspase-3 detection, we first incubated it with recombinant caspase-3 in buffer. Within 1 h, 90% of **1** was shown to cyclize and afford two cyclized isomers based on the Radio-HPLC chromatograph (Figure 1A)^[13]. As expected, under the same condition, **1-D** could not be cleaved by caspase-3 and no cyclization product but **1-D** itself was observed.

Next, we applied **1** to detect caspase-3 in apoptotic cells *in vitro*. HeLa cell was incubated with doxorubicin (Dox) ($2\ \mu\text{M}$) for 24 h to induce apoptosis. Caspase-3 assays confirmed that the lysates of Dox-treated cells showed 9.5 fold higher caspase-3 activity than that of non-treated cells. (Figure S4A) Both Dox-treated and non-treated cells were then incubated with **1** for 4 h. The retained ^{18}F activity in Dox-treated cells was 2.2-fold of that of non-treated cells. As control, a caspase-3 inhibitor (Z-VAD-fmk) was applied to a similar set of Dox-treated cells, which decreased the retained ^{18}F activity to 1.3-fold (Figure 1B). Both Dox-treated and non-treated cells were subject to extraction and analyzed by Radio-HPLC. The major radioactive fractions from Dox-treated cells were cyclized products of **2** (Figure 1A), but it was not observed in the extraction from non-Dox-treated cells. These results demonstrate that caspase-3 activity can trigger the cyclization of **1** in living cells and enhance the retention of the radioactivity.

The biodistribution of **1** was performed in nude mice that were euthanized at 120 minutes post injection. Organs were collected, weighed, and analyzed with a gamma counter for radioactivity uptake (Table S1). Briefly, kidney and bladder showed high levels of absolute uptake, suggesting primary renal clearance of the tracer. Low tracer activity in muscle promises minimal background signal in PET imaging. Uptake in brain was low, suggesting that **1** could not cross the blood-brain barrier.

Evaluation of **1** and **1-D** for PET imaging of caspase-3 activity *in vivo* was performed in HeLa tumor xenograft-bearing nude mice. Tumors were implanted and grown for more than 10 days before intratumoral injection of 0.2 mg Dox (20 μ L). 4 days post treatment, **1** or **1-D** (5–15 MBq/135–405 μ Ci) was injected through the tail vein for PET imaging. Static PET scans (5 min) were performed 65, 125 and 182 min post tracer injection. Figure 2 shows representative PET images of the same mouse injected with **1** before and after Dox treatment and another mouse with **1-D** after Dox treatment.

Quantification of the PET images with the activatable PET tracer **1** revealed that the uptake (%ID/g) of the ^{18}F activity in tumors significantly increased after Dox treatment: from 0.81 ± 0.28 (baseline) to 1.17 ± 0.17 (treated) at 65 min, from 0.67 ± 0.24 (baseline) to 1.29 ± 0.07 (treated) at 182 min (Figure 3A); this result correlates well with the caspase-3 level detected in tumors – a 1.9 fold increase in treated tumors (Figure S4B). The uptake difference between baseline and treated increased from 0.36 ± 0.15 at 65 min to 0.63 ± 0.11 at 182 min (Figure 3B), and the uptake ratio between tumor and muscle (T/M) increased from 3.30 fold at 65 min to 7.00 fold at 182 min in treated tumors (Figure 3C).

In contrast, the uptake of **1-D** in both treated and non-treated tumors was much lower than that of **1** (Figure 3A), and the uptake difference between before and after treatment ($<0.2\%$ ID/g) was also much smaller (Figure 3B). The ratio of T/M did not show significant increases either (Figure 3C). Our PET imaging results demonstrate that **1** can image caspase-3 activity in drug-treated tumors *in vivo* and that both caspase-3 activation and cyclization are required for the enhanced imaging contrast in apoptotic tumors.

$[^{18}\text{F}]\text{C-SNAT}$ (**1**) compares favorably to known apoptosis PET tracers (Table S2) with both high tumor/muscle ratio in apoptotic tumors and high uptake value (%ID/g) in apoptotic tumors. Consistent with the mechanism, $[^{18}\text{F}]\text{C-SNAT}$ showed a trend of increasing uptake over the time (Figure 3A) in apoptotic tumors and thus increased differences between treated apoptotic and non-treated tumors at later time points. This trend has not been observed with other apoptosis PET tracers; for example, with $[^{18}\text{F}]\text{ICMT-11}$, a PET tracer that binds active caspase-3, the uptake at the apoptotic tumors decreased over the time after injection.^[7a] Furthermore, our probe designing principle is not limited to caspase-3 but may serve as a general strategy for developing PET tracers for imaging the activity of other enzymes (i.e. furin, MMPs).

In conclusion, we have successfully designed and synthesized an ^{18}F -labeled caspase-3 triggered nano-aggregation PET tracer ($[^{18}\text{F}]\text{C-SNAT}$), and demonstrated its application for imaging caspase-3 activity in doxorubicin-treated tumor xenografts. This activatable PET tracer undergoes intramolecular cyclization and subsequent aggregation upon caspase-3 activation to achieve enhanced retention in apoptotic tumors. Applications of this strategy for other enzyme targets as well as translation of $[^{18}\text{F}]\text{C-SNAT}$ into clinical studies are currently under investigation.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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13. Two cyclized isomers are likely diastereomers due to two different ring-closing orientations, as observed in our previous study (ref 11).

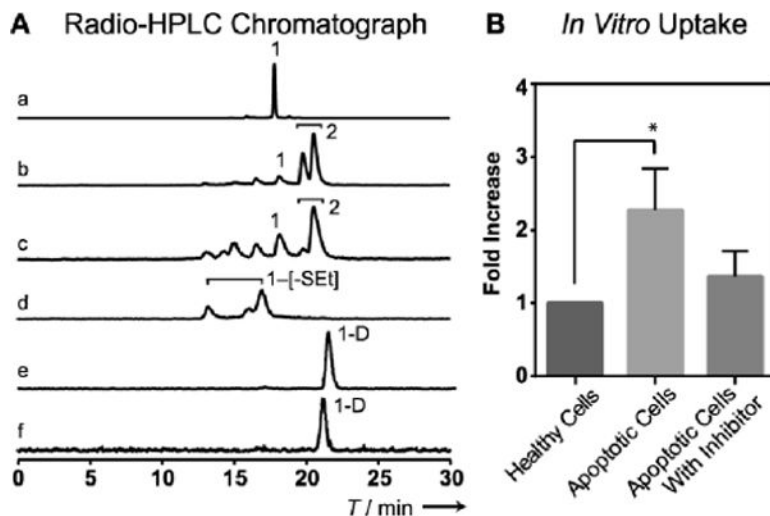


Figure 1.

A) Analytical Radio-HPLC showing the radioactive signal composition: a) **1** in saline; b) **1** incubated with caspase-3 in solution for 1 h; c) extraction from **1** cellular uptake in apoptotic cell (4 h); d) extraction from **1** cellular uptake in healthy cell (4 h); No cyclization product **2** but disulfide bond cleaved intermediates (**1**-[-SEt]) were observed in healthy cells; e) **1-D** in saline; f) **1-D** incubated with caspase-3 for 1 h. **B)** *In vitro* uptake (fold over healthy cell, $n=3$) of **1** in healthy HeLa cells, apoptotic HeLa cells (treated with 2 μ M Doxorubicin) and apoptotic HeLa cells with caspase-3 inhibitor (50 μ M of Z-VAD-FMK) added. * indicates statistically significant $P=0.045$.

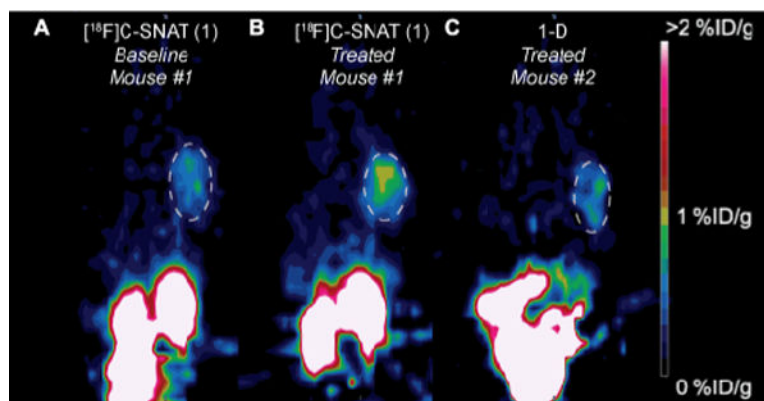


Figure 2. Representative PET images showing HeLa tumor xenografts (white dashed circles) on the right shoulder of mice 125 min after i.v. injection of tracer before (A) and after doxorubicin treatment (B & C). A) Mouse #1 before treatment imaged with **1** (7.8 MBq/211 μ Ci). B) Mouse #1 after treatment imaged with **1** (12 MBq/324 μ Ci). C) Mouse #2 after treatment imaged with **1-D** (5.4 MBq/146 μ Ci). All images are normalized to the same scale.

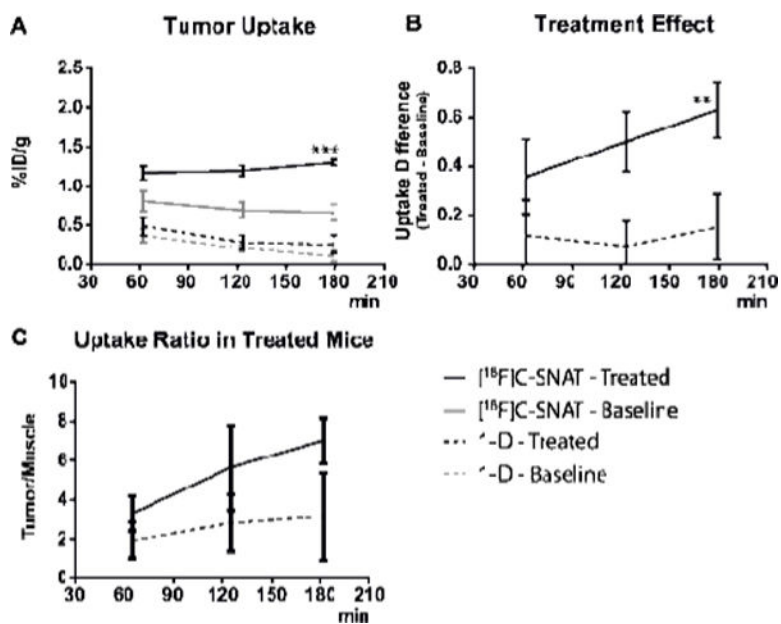
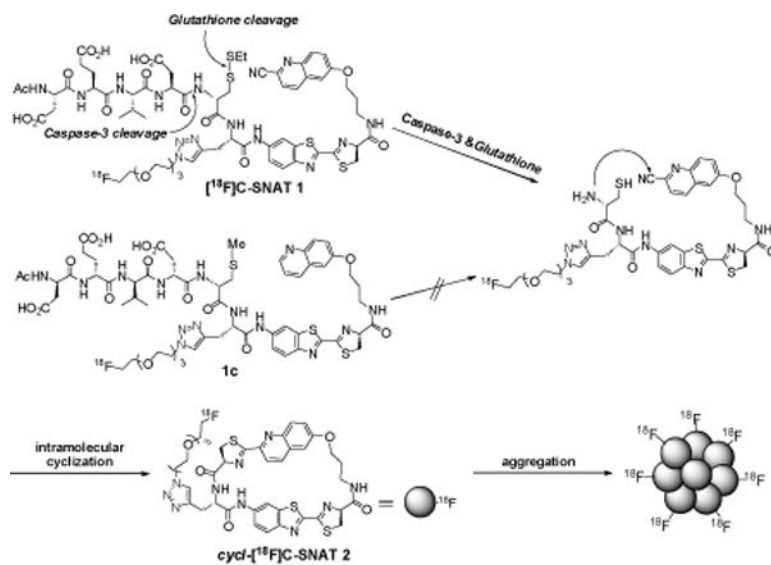
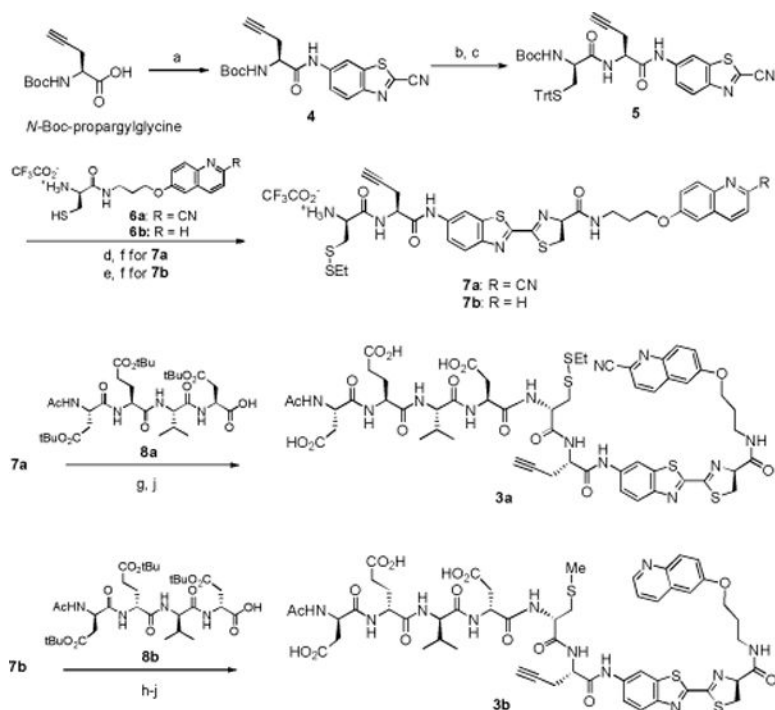


Figure 3.

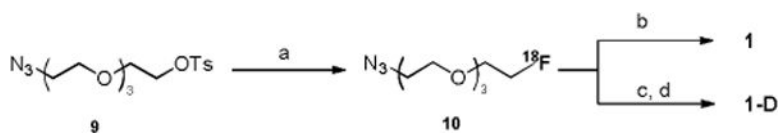
A) Uptake of **1** and **1-D** (%ID/g \pm sem) in xenograft HeLa tumor and muscle, before and after intratumor injection of Dox (0.2 mg) 4 days prior to the imaging. Uptake is calculated based on 5 min static PET scans at 65, 125 and 182 min. *** indicates $p=0.002$ at 182 min. **B)** The uptake of **1** and **1-D** in treated tumors; ** shows a significant $p=0.0037$ at 182 min. **C)** The ratio between tumor and muscle uptake in treated tumors, calculated based on the uptake (average uptake in tumor/average uptake in muscle region).

**Scheme 1.**

Activation, intramolecular cyclization of [¹⁸F] C-SNAT (**1**) and aggregation of cyclized product (**2**).

**Scheme 2.**

Synthesis of precursors **3a** and **3b**. a) *i*-butyl chloroformate, 4-methylmorpholine, THF, 0 °C, 2 h and then 6-amino-2-cyanobenzothiazole, THF, 0 °C to RT, 12 h; 85%. b) 60% TFA in DCM, RT, 1 h. c) *N*-Boc-*S*-Trt-*D*-Cysteine, HBTU, DIPEA, DMF, RT, 2 h; 80% from **4**. d) **6a**, DIPEA, DMF, RT, 30 min. e) **6b**, DIPEA, DMF, 30 min. f) 60% TFA in DCM, RT, 1 h and then 2-(ethyldisulfanyl) pyridine, MeOH, RT, 2 h; 51% from **5** for (**7a**), 40% for **7b**. g) Ac-DEVD-OH (**8a**), HBTU, DIPEA, THF, RT, 1 h. h) Ac-devd-OH (**8b**), HBTU, DIPEA, THF, RT, 1 h. i) CH₃I, TCEP·HCl, DIPEA, DMF, 1 h. j) 50% TFA in DCM, RT, 2 h; 69% (**3a**) from **7a**, 38% (**3b**) from **7b**.

**Scheme 3.**

Radiosynthesis of **1** and **1-D**, a) ^{18}F /Kryptofix 222/ K_2CO_3 , DMSO, 110 °C, 20 min. b) **3a**, $\text{Cu}(\text{CH}_3\text{CN})_4\text{PF}_6$, BPDS, DMSO/water, 60 °C, 30 min. c) **3b**, CuSO_4 , sodium ascorbate, DMSO/water, 40 °C, 30 min. d) 3-azidopropan-1-amine hydrochloride, 40 °C, 30 min.