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A fluorine-18 labeled radiotracer for PET imaging of γ -glutamyltranspeptidase in living subjects†

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The expression level of γ -glutamyltranspeptidase (GGT) in some malignant tumors is often abnormally high, while its expression is low in normal tissues. Therefore, GGT is considered as a key biomarker for cancer diagnosis. Several GGT-targeting fluorescence probes have been designed and prepared, but their clinical applications are limited due to their shallow tissue penetration. Considering the advantages of positron emission tomography (PET) such as high sensitivity and deep tissue penetration, we designed a novel PET imaging probe for targeted monitoring of the expression of GGT in living subjects, $[(^{18}\text{F})\gamma\text{-Glu-Cys-PPG}(\text{CBT})\text{-AmBF}_3)_2$, hereinafter referred to as $[(^{18}\text{F})\text{GCPA}]_2$. The non-radioactive probe (GCPA)₂ was synthesized successfully and ^{18}F fluorinated rapidly *via* the isotope exchange method. The radiotracer $[(^{18}\text{F})\text{GCPA}]_2$ could be obtained within 0.5 h with the radiochemical purity over 98% and the molar activity of $10.64 \pm 0.89 \text{ GBq } \mu\text{mol}^{-1}$. It showed significant difference in cellular uptake between GGT-positive HCT116 cells and GGT-negative L929 cells ($2.90 \pm 0.12\%$ vs. $1.44 \pm 0.15\%$ at 4 h, respectively). *In vivo* PET imaging showed that $[(^{18}\text{F})\text{GCPA}]_2$ could quickly reach the maximum uptake in tumor ($4.66 \pm 0.79\% \text{ ID g}^{-1}$) within 5 min and the tumor-to-muscle uptake ratio was higher than 2.25 \pm 0.08 within 30 min. Moreover, the maximum tumor uptake of the control group co-injected with the non-radioactive probe (GCPA)₂ or pre-treated with the inhibitor GGsTop decreased to $3.29 \pm 0.24\% \text{ ID g}^{-1}$ and $2.78 \pm 0.32\% \text{ ID g}^{-1}$ at 10 min, respectively. *In vitro* and *in vivo* results demonstrate that $[(^{18}\text{F})\text{GCPA}]_2$ is a potential PET probe for sensitively and specifically detecting the expression level of GGT.

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Introduction

γ -Glutamyltranspeptidase (GGT) is a member of the N-terminal nucleophile hydrolases anchored to the cell membrane.¹ It is vital in the γ -glutamine cycle of glutathione synthesis and degradation, and can functionally catalyze GSH transformation into other amino acids or peptides *via* cleaving the γ -glutamate (γ -Glu) bond.^{2–5} GGT binds glutathione as a donor substrate and initially forms a γ -glutamyl enzyme which reacts with a water molecule or an acceptor substrate (dipeptides, L-methionine, L-alanine, L-glutamine and L-cystine) to form glutamate or a new isopeptide bond, respectively, thus regenerating the free enzyme.⁶ Several studies have demonstrated that GGT is a pro-oxidant in intracellular transduction and its catalysis is notably

enhanced in the liver.⁷ A high GGT level in plasma is often associated with some adverse symptoms such as viral hepatitis,⁸ bone disease⁹ and inflammatory bowel disease.¹⁰ Therefore, the expression level of GGT in human serum is considered as a key indicator in the clinical diagnosis of several diseases.¹¹ Moreover, abnormally high expression level of GGT could lead to the breakdown of redox homeostasis in intracellular environment and eventually result in cancerization.^{5,12} GGT can promote the growth and proliferation of tumor cells by mediating the circulation of γ -Glu in tumor cells.¹³ It also can promote cancer metastasis and drug resistance of tumors.¹⁴ Preclinical studies have indicated that GGT is overexpressed in certain malignant tumor cells, such as liver, breast, neck, cervical, colon and ovarian cancers, while it is not or low expressed in normal tissues except liver and kidney.^{13–17} Accordingly, GGT could be used as an important biomarker for cancer diagnosis.¹⁸ Therefore, sensitively and specifically monitoring the expression level of GGT in living subjects should be of great significance for the clinical cancer diagnosis and treatment.

Up to now, a series of molecular probes have been successfully developed to detect the expression level of GGT. According to their different imaging modality, they can be divided into near-infrared fluorescent probes,¹⁴ bioluminescent probes,¹⁹ chemiluminescent probes,²⁰ magnetic resonance imaging

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(MRI) probes,¹⁵ positron emission tomography (PET) probes,²¹ etc. Optical imaging, owing the characteristics of excellent spatial resolution and sensitivity, has been applied to direct the resection of lesions in clinical surgery.^{22,23} However, its application also be limited due to the poor tissue penetration.²⁴ Due to higher sensitivity, deeper tissue penetration and can provide more quantifiable information, PET imaging may be superior to other imaging technologies in monitoring the expression level of GGT in living subject.^{22,24–26} At present, GGT-targeting probes under preclinical study are mainly optical probes, while PET probes are rarely reported for detecting the GGT expression.²⁷ Recently, our group reported a GGT-targeting PET probe ¹⁸F-1G to monitor the expression level of GGT in living subjects.²¹ However, due to its poor water solubility and high-dose requirement, its further clinical applications were limited. Therefore, it is necessary to develop new GGT-targeting PET probes to sensitively and specifically monitor the expression level of GGT in living subjects.

In this study, a novel GGT-targeting radiotracer (¹⁸F)GCPA₂ was designed and synthesized. As shown in Scheme 1, (¹⁸F)GCPA₂ contains two γ -glutamate (γ -Glu) groups in response to GGT and two moieties of [¹⁸F]ammoniomethyl-trifluoroborate ([¹⁸F]AMBF₃) conjugated to the CBT-PPG-Cys(γ -Glu)-Cys(γ -Glu)-PPG-CBT scaffold. The incorporation of two hydrophilic groups (γ -Glu and AMBF₃) would be conducive to enhance the water solubility, and the introduction of two γ -Glu might be beneficial to enhance the specificity of the probe to GGT. *In vitro* stability, water solubility and GGT-targeting ability as well as sensitivity and specificity of the probe for detecting GGT *in vivo* were evaluated systematically.

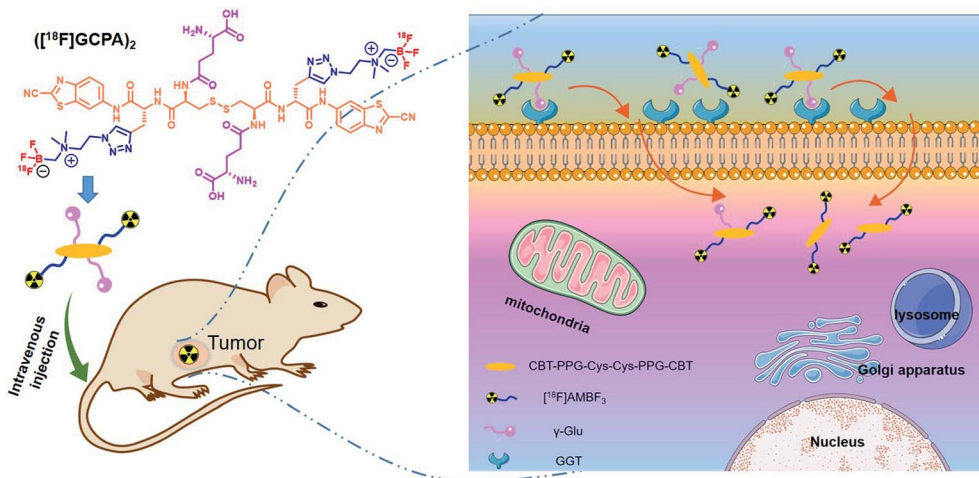
Results and discussion

Chemistry and radiochemistry

The non-radioactive probe (GCPA)₂ was synthesized successfully according to the synthesis route as shown in Scheme 2. At first, compound 1 was successfully synthesized based on the method reported previously.^{28,29} Then it was coupled with *N,N'*-

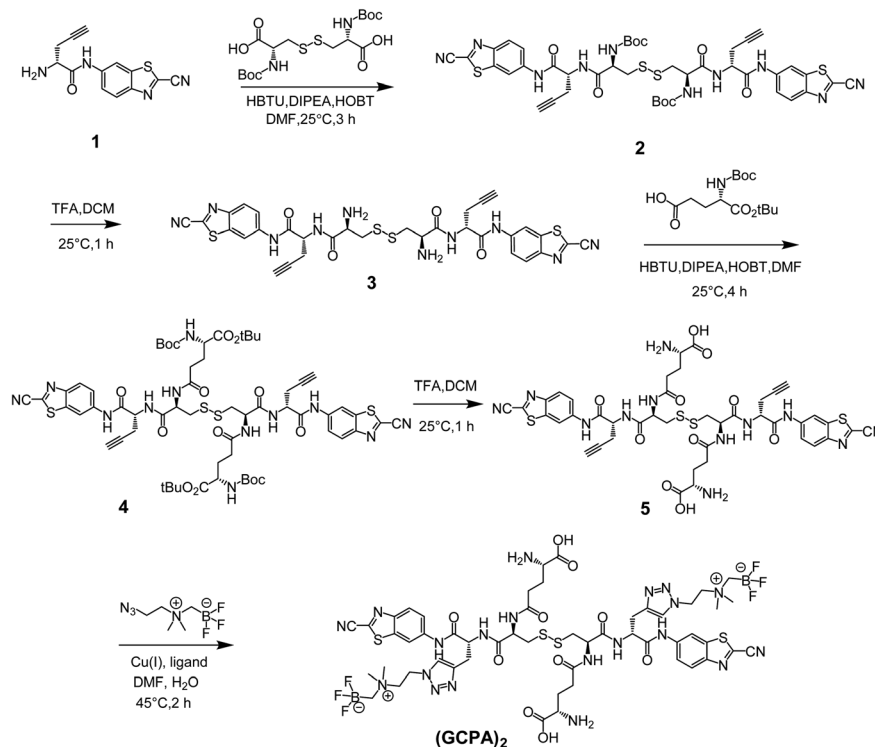
bis(*tert*-butoxycarbonyl)-L-cystine to yield compound 2. After deprotection, compound 3 was obtained and then coupled with Boc-Glu-OtBu through a condensation reaction to yield compound 4. Subsequently, two protecting groups were removed to produce compound 5. At last, AMBF₃ was attached to the alkynyl group of compound 5 *via* the click reaction to yield the non-radioactive probe (GCPA)₂. The compound 5 and non-radioactive probe (GCPA)₂ were purified with semi-preparative high performance liquid chromatography (HPLC) whose conditions were listed in ESI (Tables S1–S2†). All compounds were characterized by HPLC (see Fig. S1–S5 in ESI section†) and electrospray ionization-mass spectrometry (ESI-MS) (see Fig. S6–S10 in ESI section†), and the HPLC profiles of all compounds were recorded at wavelength of 254 nm. Analytical HPLC conditions were display in Table S3.† The HPLC purity of compound 2 to compound (GCPA)₂ are 97%, 97%, 98%, 95% and 97%, respectively. The chemical structures of all the new compounds were further verified by ¹H NMR, ¹³C NMR and ¹⁹F NMR (see Fig. S11–S21 in ESI†).

The radiochemical synthesis of GGT-targeting PET probe (¹⁸F)GCPA₂ was successfully completed *via* a single-step isotope exchange reaction (see Fig. 1). The non-radioactive probe (GCPA)₂ (0.75 μ mol) was efficiently labelled with fluorine-18 (~9.25 GBq) within 30 min. As shown in Fig. 1b, the decay-corrected radiochemical yield (RCY) of (¹⁸F)GCPA₂ was 76 \pm 3% and the radiochemical purity (RCP) was over 98% after a simple and fast purification. The molar activity was measured to be 10.64 \pm 0.89 GBq μ mol⁻¹ (*n* = 5). The retention time of (¹⁸F)GCPA₂ was 14.7 min (Fig. 1a), which was consistent with the retention time of non-radioactive probe (GCPA)₂ (Fig. S5 in ESI section†). All results suggested that (GCPA)₂ could be quickly and conveniently radiolabelled with fluorine-18 *via* the isotope exchange (IEX) method. Then, the stability of the target probe (¹⁸F)GCPA₂ was measured. As shown in Fig. 1c and d, when the probe was incubated with phosphate-buffered saline (PBS) or fetal bovine serum (FBS) for 4 h, no other impurity was observed. It is suggested that the probe was stable enough for further biological evaluation.



Scheme 1 Proposed action mechanism of the GGT-targeting radiotracer (¹⁸F)GCPA₂.





Scheme 2 Synthesis route of non-radioactive probe (GCPA)₂. Synthesis of compound 2 to compound (GCPA)₂: (1) *N,N'*-bis(*tert*-butoxycarbonyl)-*L*-cystine, HBTU, HOBT, DIPEA, DMF, 25 °C, 3 h, 97%; (2) 50% TFA/DCM, 25 °C, 1 h, 85%; (3) Boc-Glu-OtBu, HBTU, HOBT, DIPEA, DMF, 25 °C, 4 h, 71%; (4) 50% TFA/DCM, 25 °C, 1 h, 46%; (5) 66% DMF/H₂O, Cu(I), ligand, AMBF₃, 45 °C, 2 h, 83%.

Octanol-water partition coefficient (log *P*)

The log *P* of ([¹⁸F]GCPA)₂ was determined to be -1.66 ± 0.02 (*n* = 3), indicating that the radiotracer is hydrophilic. Compared with ¹⁸F-1G, log *P* value of ([¹⁸F]GCPA)₂ is significantly less than that of ¹⁸F-1G (-1.66 ± 0.02 vs. 0.042 ± 0.0056), indicating that the hydrophilicity of ([¹⁸F]GCPA)₂ is significantly higher than that of ¹⁸F-1G.

Biocompatibility assay

Biocompatibility of the probe ([¹⁸F]GCPA)₂ was studied by measuring the cytotoxicity of the non-radioactive probe (GCPA)₂ against normal cells L929 and tumor cells HCT116 using the traditional MTT assay. As displayed in Fig. S22,† when L929 and HCT116 cells were incubated with (GCPA)₂ at a low concentration (<50 μM), the cell viability of both cells was more than 90%.

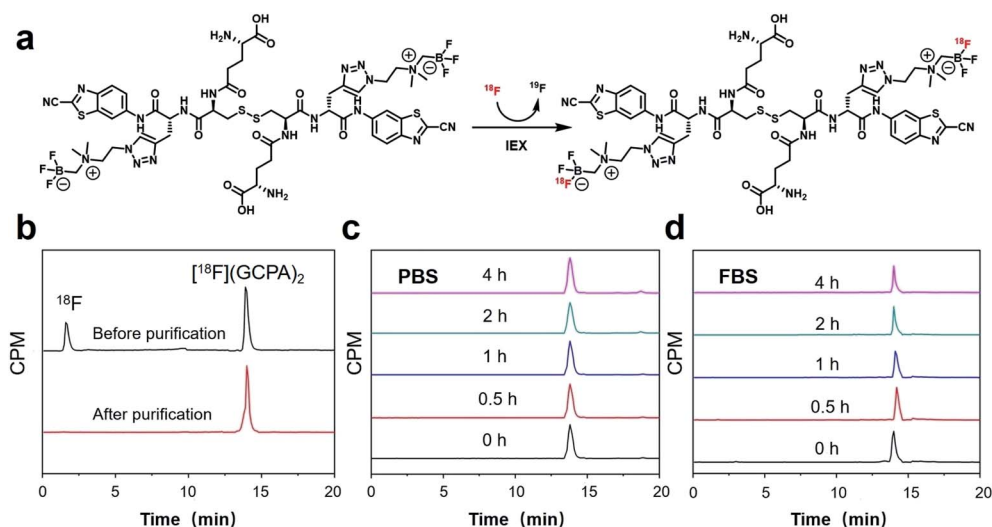


Fig. 1 (a) Radiosynthesis of ([¹⁸F]GCPA)₂ at 80 °C via the isotope exchange (IEX) method. (b) HPLC analysis of ([¹⁸F]GCPA)₂ before and after purification. (c and d) Stability assay of ([¹⁸F]GCPA)₂ in PBS and FBS.



After both cells were incubated with (GCPA)₂ (100 μM) for 24 h, the viability of HCT116 cells was more than 80% and that of L929 cells was 70.29 ± 0.03%. Considering that the actual concentration of the radiotracer used in the living subjects was far less than 50 μM, the tracer possesses good biocompatibility and could be safely used in the following biological evaluation.

GGT-targeting responsive ability

In order to confirm that the probe ([¹⁸F]GCPA)₂ could be recognized and cleaved by GGT, the specific response of the non-radioactive probe (GCPA)₂ towards GGT was studied (see Fig. 2). After the non-radioactive probe (GCPA)₂ (250 μM) was incubated with GGT (100 U L⁻¹) for 30 min, the peak of (GCPA)₂ disappeared and two new peaks were clearly observed (see Fig. 2b). In order to verify the two new products after incubating the non-radioactive probe with GGT, we synthesized two enzymatic products (C1 and C2). The ESI-MS and HPLC analysis were displayed in Fig. S23–S26.† The half retention time of C1 (10.7 min) and C2 (8.7 min) were consistent with that of these two new products, confirming that non-radioactive probe could target and respond to GGT. Moreover, the reaction rate of (GCPA)₂ respond to GGT was studied. As shown in Fig. S27,† (GCPA)₂ has been almost converted into cleavage products C1 (73%) and C2 (27%) after incubating with GGT for 20 min. This results indicated that the probe ([¹⁸F]GCPA)₂ could be specifically recognized and cleaved quickly by GGT.

Cellular uptake

The GGT-targeting ability of ([¹⁸F]GCPA)₂ was further investigated by monitoring the uptake of the radiotracer in GGT-positive HCT116 cells and GGT-negative L929 cells. As shown in Fig. 3, when the probe ([¹⁸F]GCPA)₂ (37 KBq) was incubated with HCT116 cells, the cellular uptake of ([¹⁸F]GCPA)₂ gradually increased with the prolonged incubation time and arrived at a plateau at 2 h with the uptake value of 2.81 ± 0.04%. At 4 h, the cellular uptake still maintained at a high level (2.90 ± 0.12%).

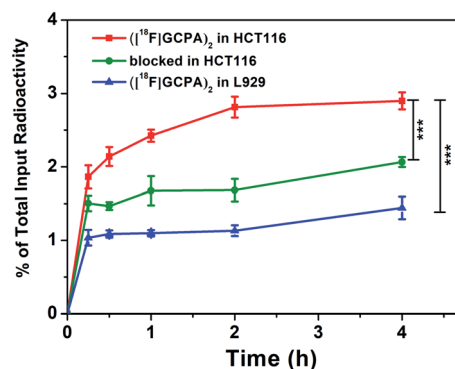


Fig. 3 Cellular uptake of ([¹⁸F]GCPA)₂ in HCT116 cells (red), HCT116 cells pre-treated with 10 nM (GCPA)₂ (green), and L929 cells (blue) incubated for different time (*n* = 3). ****P* < 0.001.

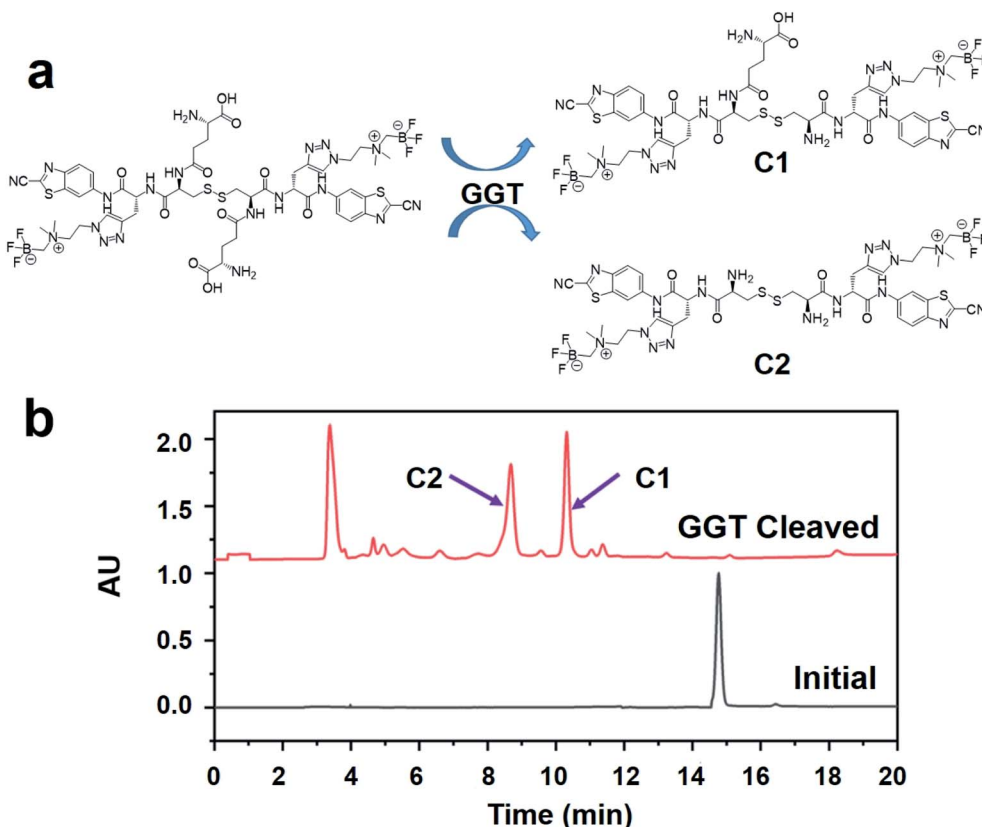


Fig. 2 (a) Schematic illustration of GGT-controlled cleavage of (GCPA)₂ to form two enzymatic products. (b) HPLC analysis of 250 μM (GCPA)₂ incubated with 0 (black) or 100 U L⁻¹ of GGT (red) at 37 °C for 30 min.



On the contrary, the maximum cellular uptake of $([^{18}\text{F}]\text{GCPA})_2$ in L929 cells was only $1.44 \pm 0.15\%$ at 4 h, which was significantly lower than that in the GGT-positive HCT116 cells. High uptake of radioactivity in HCT116 cells was observed and the uptake value of HCT116 cells was twice as high as that of L929 cells, suggesting that the tracer $([^{18}\text{F}]\text{GCPA})_2$ could target tumor cells with high expression level of GGT. The small amount of $([^{18}\text{F}]\text{GCPA})_2$ in L929 cells was likely due to the non-specific uptake by normal cells with low expression level of GGT. The significant difference of cellular uptake between HCT116 cells and L929 cells was related to the different GGT expression in the two kind cells. These results also suggested that the tracer could be selectively recognized by GGT.

In addition, to further study the specificity of $([^{18}\text{F}]\text{GCPA})_2$ for GGT, the cellular uptake in the blocked group was also investigated, in which GGT-positive HCT116 cells were pre-treated with the non-radioactive probe $(\text{GCPA})_2$ (10 nmol) for 30 min before incubated with the tracer $([^{18}\text{F}]\text{GCPA})_2$. As expected, cellular uptake was significantly inhibited and the maximum decreased remarkably from $2.90 \pm 0.12\%$ to $1.89 \pm 0.07\%$. This further demonstrated that the tracer $([^{18}\text{F}]\text{GCPA})_2$ could specifically target to GGT and can be used for sensitively and specifically monitoring the expression level of GGT.

MicroPET imaging

Sensitivity and specificity of the tracer $([^{18}\text{F}]\text{GCPA})_2$ in detecting the GGT expression was investigated by microPET imaging of HCT116 xenografts. In Fig. 4a, HCT116 tumor could be clearly visualized from 5 min to 30 min after injection of $([^{18}\text{F}]\text{GCPA})_2$ and the tumor uptake rapidly reached the maximum value $4.66 \pm 0.79\%$ ID g^{-1} at 5 min post injection (Fig. 4b). The rapid

increase of tumor uptake within such a short time might be due to that two γ -Glu groups greatly enhanced the GGT-targeting ability of the radiotracer so that the tracer could be efficiently recognized and cleaved by GGT and undergoes caveolae-mediated endocytosis and transcytosis rapidly into tumor cells.³⁰ This also indicated that the tracer $([^{18}\text{F}]\text{GCPA})_2$ could sensitively detect the GGT expression in living subjects. As time went on, the tumor uptake decreased to $2.97 \pm 0.29\%$ ID g^{-1} and $1.35 \pm 0.20\%$ ID g^{-1} at 20 min and 60 min post injection, respectively. Since the disulfide bond in the tracer was not easily to be reduced, a small amount of dimer formed during the whole imaging could not significantly prolong the retention time of the probe in the tumor.³¹ In addition, the elimination half-life of the tracer $([^{18}\text{F}]\text{GCPA})_2$ was determined to be 30.5 min, suggesting that the tracer could be quickly cleared from the body (Fig. S28 in ESI†), which was consistent with the PET imaging results. Although both the tracer $([^{18}\text{F}]\text{GCPA})_2$ and ^{18}F -1G were cleared rapidly from the tumor site, the tracer $([^{18}\text{F}]\text{GCPA})_2$ could be absorbed by tumor more quickly and the tumor uptake value was higher than the probe ^{18}F -1G. The tracer $([^{18}\text{F}]\text{GCPA})_2$ reached the maximum tumor uptake ($4.66 \pm 0.79\%$ ID g^{-1}) at 5 min post injection, while the probe ^{18}F -1G only reached the maximum tumor uptake ($3.20 \pm 0.29\%$ ID g^{-1}) at 10 min post injection.

Compared with tumor uptake, the muscle uptake of tracer was relatively low, which ranged from $1.66 \pm 0.27\%$ ID g^{-1} to $0.61 \pm 0.21\%$ ID g^{-1} within 60 min (see Fig. 4b). Accordingly, the tumor-to-muscle uptake ratio (T/M) was always high and the maximum T/M ratio of 2.82 ± 0.10 was achieved at 5 min post injection (see Fig. 4c). This indicated that $([^{18}\text{F}]\text{GCPA})_2$ could serve as a potential tracer for sensitively visualizing GGT expression in living subjects. In addition, we also compared the

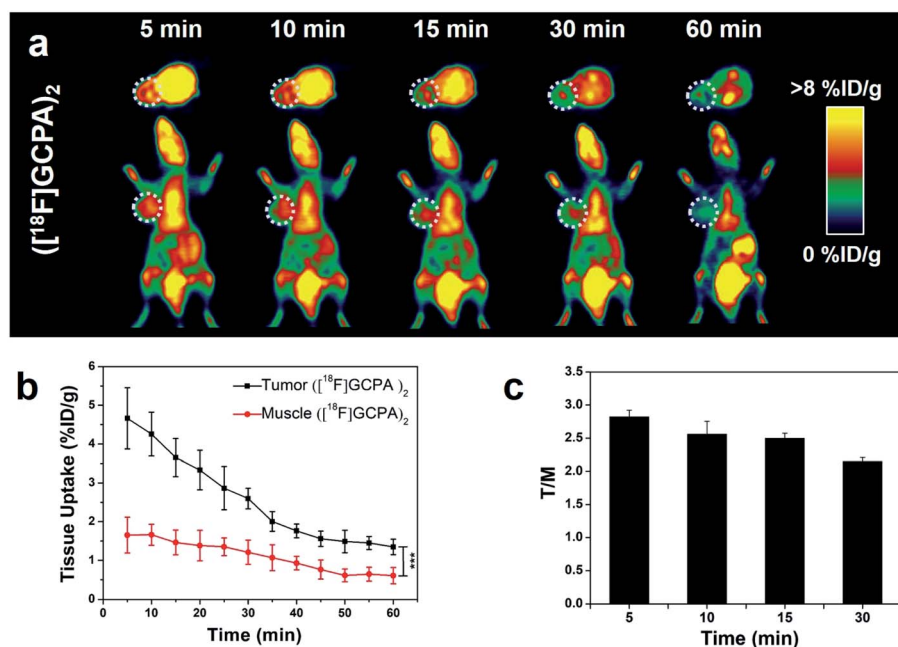


Fig. 4 (a) MicroPET imaging of $([^{18}\text{F}]\text{GCPA})_2$ in HCT116 tumor-bearing mice ($n = 4$). Tumor was circled by the white dotted line. (b) Quantitative analysis of tumor uptake and muscle uptake from microPET imaging. (c) Time course of tumor-to-muscle uptake ratio (T/M).



T/M ratio between (^{18}F] GCPA)₂ and ^{18}F -1G. The T/M value of ^{18}F -1G was about 2 at 10 min and 30 min post injection, while that of this probe was 2.56 ± 0.19 and 2.15 ± 0.06 at 10 min and 30 min post injection, respectively. These results also proved that the GGT-targeting ability of the tracer (^{18}F] GCPA)₂ was higher than the probe ^{18}F -1G.

From the PET imaging, one could observe that the PET signal of the liver gradually weakens over the imaging time. At the same time, it could be also observed that there was a strong PET signal in the bladder at 5 min post injection, and the PET signal further enhanced as time went on. However, there was no significant reduction in the liver uptake of probe ^{18}F -1G during the whole imaging process.²¹ In addition, no strong PET signal was observed in the bladder at 10 min post injection of ^{18}F -1G. These difference could be attributed to the fact that the tracer (^{18}F] GCPA)₂ was more hydrophilic than the probe ^{18}F -1G, resulting in faster clearance of the tracer (^{18}F] GCPA)₂ through the liver and kidneys.^{32–34}

To further demonstrate the specificity of (^{18}F] GCPA)₂ to the expression of GGT *in vivo*, microPET imaging of two control groups was also studied. One group of HCT116 tumor-bearing mice were co-injected with (^{18}F] GCPA)₂ and the non-radioactive (GCPA)₂ during the microPET scanning, and the other group were pre-treated with GGsTop before injection of the tracer (^{18}F] GCPA)₂. PET images and time course of tumor and muscle uptake quantified from PET images have been shown in Fig. 5a–c. As expected, low radioactive signal in the tumor region could be detected for both groups within 1 h. For the control group with co-injection of (^{18}F] GCPA)₂ and (GCPA)₂, the tumor uptake derived from PET images was $2.71 \pm 0.19\%$ ID g⁻¹ at 5 min post injection. For the other control group with pre-treatment of GGsTop, tumor uptake also decreased significantly

and it was only $1.83 \pm 0.22\%$ ID g⁻¹ at 5 min post injection. Compared with the mice injected with only (^{18}F] GCPA)₂, the T/M ratio of two control groups also decreased significantly (see Fig. 5d). For example, T/M ratio of the control group with co-injection of (^{18}F] GCPA)₂ and (GCPA)₂ was 2.12 ± 0.20 and 1.60 ± 0.09 at 5 min and 30 min post injection, respectively, while T/M ratio of the other control group with pre-treatment of GGsTop was only 1.58 ± 0.09 and 1.04 ± 0.02 at 5 min and 30 min post injection, respectively, which was obviously smaller than that of the group with injection of only (^{18}F] GCPA)₂ (2.82 ± 0.10 at 5 min and 2.15 ± 0.06 at 30 min). This phenomenon was consistent with the cellular uptake in HCT116 cells co-cultured with (^{18}F] GCPA)₂ and non-radioactive probe (GCPA)₂. As shown in Fig. S29a,† the tumor uptake of the group injected with only (^{18}F] GCPA)₂ was significantly more than the group co-injected with (^{18}F] GCPA)₂ and (GCPA)₂ ($P < 0.05$) and that of the group pretreated with GGsTop ($P < 0.001$). As for the T/M ratio (Fig. S29b in ESI†), the group treated with the tracer (^{18}F] GCPA)₂ was significantly different from that co-injected with (^{18}F] GCPA)₂ and (GCPA)₂ ($P < 0.01$) and that pre-treated with GGsTop ($P < 0.001$). These results fully verified that (^{18}F] GCPA)₂ is a promising GGT-specific and responsive PET tracer, which could be used to sensitively and specifically detect GGT expression in living subjects.

Conclusions

In summary, a novel GGT-targeting radiotracer (^{18}F] GCPA)₂ was successfully developed for sensitive and specific detection of GGT expression in living subjects. The radiotracer exhibited good molar activity, high radiochemical yield and purity as well as outstanding stability and excellent biocompatibility.

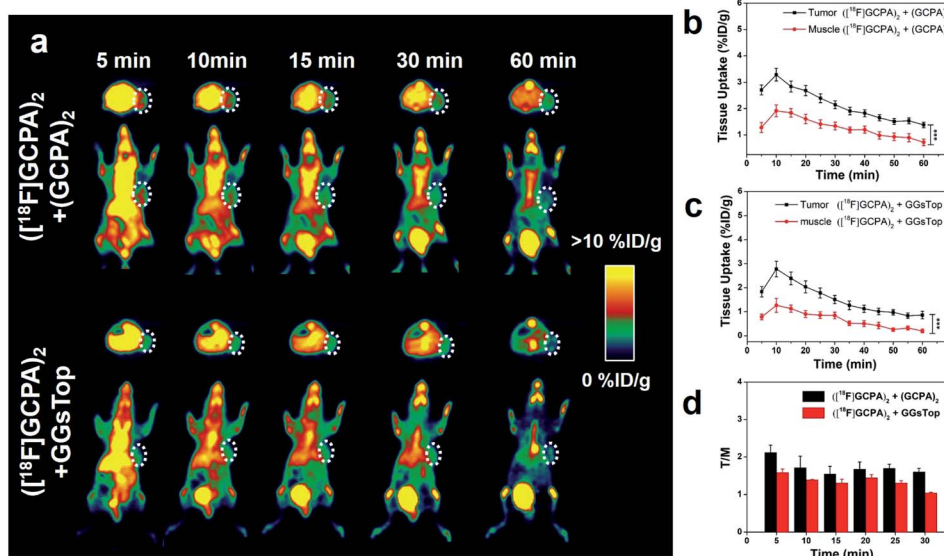


Fig. 5 (a) Representative coronal and transverse PET images of two control groups ($n = 4$). One control group was co-injected with (^{18}F] GCPA)₂ and (GCPA)₂ during microPET scanning, and another group was pre-treated with GGsTop for 30 min before injection with (^{18}F] GCPA)₂. (b and c) Quantitative analysis of tumor uptake and muscle uptake in two control groups. (d) T/M ratio of two control groups ((^{18}F] GCPA)₂ + (GCPA)₂ and (^{18}F] GCPA)₂ + GGsTop).



Biological evaluations both confirmed that the tracer possessed high sensitivity and specificity to the expression level of GGT. MicroPET imaging of HCT116 xenografts indicated that [^{18}F]GCPA₂ has great potential for sensitively and specifically monitoring the expression of GGT in living subjects, which may provide important information for the clinical diagnosis of GGT-related diseases.

Experimental section

General information

All reagents were purchased from suppliers, such as Aladdin and Sangon Biotech. All chemicals could be used for experiment without additional purification. γ -Glutamyltranspeptidase (GGT, G9270-100UN) originating from equine kidney was afforded by Sigma-Aldrich (Shanghai, China). The enzyme inhibitor GGTsTop was purchased from TRC (Toronto Research Chemicals, Canada). [^{18}F]fluoride ion was produced by irradiation of ^{18}O -enriched water using a medical cyclotron (Sumitomo HM-7, Japan). Human colon cancer cell line HCT116 and mouse fibroblast cell line L929 were purchased from the Cell Bank of the Chinese Academy of Sciences (Shanghai, China). Female BALB/c nude mice (5 weeks old) were purchased from Cavens Laboratory Animal Co. Ltd (Changzhou, China).

ESI-MS analysis was performed with a quadrupole tandem mass spectrometer (SQ-detect 2) (Waters, USA). NMR spectra (^1H , ^{19}F , and ^{13}C) were recorded on Bruker DRX-400 spectrometer (Bruker, Germany). Analytical HPLC is constituted with a Waters 1525 pump, a Waters 2487 UV/visible detector and a C18 column (10 μm , 250 \times 4.6 mm, Elite). Semi-preparative HPLC (Waters, USA) equipped with a Waters 2489 UV/visible detector, a Waters 1525 pump and a RP-C18 column (5 μm , 250 \times 10 mm, Elite) was used for the purification of intermediates and the final product. Radio-HPLC equipped with a radioactivity detector (Radiomatic 610TR; Perkin Elmer; MA; USA). The radioactivity was measured using a γ counter (1470, Perkin-Elmer corporation, USA). MicroPET imaging was carried out on a Small Animal micro-PET scanner (Siemens, Germany).

Chemical synthesis of non-radioactive probe (GCPA)₂

Synthesis of compound 2. Compound 1 was synthesized with previously reported method.^{28,29} Compound 1 (65 mg, 0.24 mmol), *N,N'*-bis(*tert*-butoxycarbonyl)-L-cystine (43 mg, 0.098 mmol), *O*-benzotriazole-*N,N,N',N'*-tetramethyl-uronium-hexafluorophosphate (HBTU, 110 mg, 0.29 mmol), and 1-hydroxybenzotriazole (HOBT, 39 mg, 0.29 mmol) were added to 5 mL of anhydrous *N,N*-dimethylformamide (DMF), and then *N,N*-diisopropylethylamine (DIPEA, 96 μL , 0.58 mmol) was added to adjust the pH (8–9). The solution was stirred at 25 $^\circ\text{C}$ for 3 h under N_2 . The solvent was evaporated under reduced pressure and crude product was purified by silica gel chromatography with the eluent (hexane/ethyl acetate, 2 : 1) to afford the compound 2 as white powder (90 mg, yield ratio 97%). The melting points (mp) of compound 1 and compound 2 are 119–121 $^\circ\text{C}$ and 144–145 $^\circ\text{C}$. ^1H NMR of compound 2 (400 MHz, $\text{DMSO}-d_6$) δ 10.59 (s, 2H), 8.74 (s, 2H), 8.45 (d, J = 7.8 Hz, 2H),

8.21 (d, J = 9.0 Hz, 2H), 7.80 (d, J = 9.1 Hz, 2H), 7.20 (d, J = 8.1 Hz, 2H), 4.67 (q, J = 7.0 Hz, 2H), 4.30 (q, J = 8.8 Hz, 2H), 3.14 (d, J = 9.3 Hz, 2H), 2.91 (t, J = 2.6 Hz, 2H), 2.87 (d, J = 12.5 Hz, 2H), 2.67 (dd, J = 16.9, 6.3 Hz, 4H), 1.37 (s, 18H). ^{13}C NMR of compound 2 (100 MHz, $\text{DMSO}-d_6$) δ 171.0, 169.5, 155.9, 148.3, 139.4, 137.1, 135.8, 125.3, 121.4, 114.0, 112.2, 80.3, 79.0, 74.0, 54.0, 52.8, 28.6, 22.3. ESI-MS (m/z) of compound 2 calculated for $\text{C}_{42}\text{H}_{45}\text{N}_{10}\text{O}_8\text{S}_4$ ($[\text{M} + \text{H}]^+$) 945.22, observed as 945.17.

Synthesis of compound 3. To remove the protecting group Boc, trifluoroacetic acid (TFA, 2 mL) was added to the solution of compound 2 in dichloromethane (DCM, 2 mL), and the solution was stirred at 25 $^\circ\text{C}$ for 1 h. Cold diethyl ether (30 mL) was then added into the reaction instrument when solvent had been removed, and the resulting precipitates were dried to obtain compound 3 as white powder (60 mg, yield ratio 85%), mp 135–137 $^\circ\text{C}$. ^1H NMR of compound 3 (400 MHz, $\text{DMSO}-d_6$) δ 10.89 (s, 2H), 9.30 (d, J = 7.8 Hz, 2H), 8.73 (s, 2H), 8.45 (s, 4H), 8.24 (d, J = 9.0 Hz, 2H), 7.83 (d, J = 11.0 Hz, 2H), 4.80 (q, J = 7.4 Hz, 2H), 4.24 (s, 2H), 3.32 (d, J = 10.1 Hz, 2H), 3.08 (d, J = 5.9 Hz, 2H), 2.97 (s, 2H), 2.79–2.60 (m, 4H). ^{13}C NMR of compound 3 (100 MHz, $\text{DMSO}-d_6$) δ 169.2, 167.6, 148.4, 139.3, 137.2, 136.0, 125.4, 121.4, 114.0, 112.2, 80.0, 74.3, 53.1, 51.5, 38.6, 22.6. ESI-MS (m/z) of compound 3 calculated for $\text{C}_{32}\text{H}_{29}\text{N}_{10}\text{O}_4\text{S}_4$ ($[\text{M} + \text{H}]^+$) 745.12, observed as 745.31.

Synthesis of compound 4. Compound 3 (60 mg, 0.081 mmol) was solved in anhydrous DMF (5 mL). Then Boc-Glu-OtBu (73 mg, 0.24 mmol), HBTU (91 mg, 0.24 mmol), HOBT (32 mg, 0.24 mmol) and DIPEA (79 μL , 0.48 mmol) were added, and then the solution was stirred at 25 $^\circ\text{C}$ for 4 h under N_2 . The crude product was purified by silica gel flash chromatography using the eluent (hexane/ethyl acetate, 1 : 2) to yield compound 4 as white solid (75 mg, yield ratio 71%), mp 137–139 $^\circ\text{C}$. ^1H NMR of compound 4 (400 MHz, $\text{DMSO}-d_6$) δ 10.49 (s, 2H), 8.78 (s, 2H), 8.64 (d, J = 7.8 Hz, 2H), 8.34 (d, J = 7.3 Hz, 2H), 8.20 (d, J = 9.0 Hz, 2H), 7.84 (d, J = 10.7 Hz, 2H), 7.12 (d, J = 7.7 Hz, 2H), 4.68–4.63 (m, 2H), 4.63–4.57 (m, 2H), 3.84–3.76 (m, 2H), 3.13 (d, J = 8.7 Hz, 2H), 2.93 (d, J = 9.8 Hz, 2H), 2.90 (s, 2H), 2.68 (dd, J = 25.0, 5.7 Hz, 4H), 2.25 (t, J = 7.4 Hz, 4H), 1.83 (dt, J = 64.1, 6.5 Hz, 4H), 1.37 (s, 18H), 1.36 (s, 18H). ^{13}C NMR of compound 4 (100 MHz, $\text{DMSO}-d_6$) δ 172.0, 170.6, 169.6, 156.0, 148.3, 139.4, 135.8, 125.3, 121.4, 112.1, 80.8, 78.5, 73.9, 54.4, 53.0, 52.7, 32.1, 28.6, 28.1, 27.0, 22.1. ESI-MS (m/z) of compound 4 calculated for $\text{C}_{60}\text{H}_{75}\text{N}_{12}\text{O}_{14}\text{S}_4$ ($[\text{M} + \text{H}]^+$) 1316.44, observed as 1316.23.

Synthesis of compound 5. Compound 4 was dissolved in 4 mL TFA and DCM (1 : 1, v/v). Then the solution was stirred at 25 $^\circ\text{C}$ for 1 h. After evaporation of the solvent and precipitation with cold diethyl ether (30 mL), the crude product 5 was obtained as white powder. Subsequently, the crude product was further purified by semi-preparative HPLC to obtain a white powder (26 mg, yield ratio 46%), mp 137–140 $^\circ\text{C}$. ^1H NMR of compound 5 (400 MHz, $\text{DMSO}-d_6$) δ 10.67 (s, 2H), 8.78 (d, J = 1.9 Hz, 2H), 8.72 (d, J = 6.6 Hz, 2H), 8.46 (d, J = 6.0 Hz, 2H), 8.21 (d, J = 9.0 Hz, 2H), 7.85 (d, J = 9.1 Hz, 2H), 4.68–4.63 (m, 2H), 4.63–4.58 (m, 2H), 3.86 (s, 2H), 3.17 (d, J = 8.9 Hz, 2H), 2.90 (s, 2H), 2.89 (s, 2H), 2.68 (dd, J = 23.1, 6.9 Hz, 4H), 2.37 (dd, J = 16.9, 8.2 Hz, 4H), 2.05–1.96 (m, 4H). ^{13}C NMR of compound 5 (100 MHz, $\text{DMSO}-d_6$) δ 172.0, 171.4, 170.6, 169.6, 148.3, 139.5,



137.1, 135.7, 125.3, 121.4, 114.0, 112.1, 80.4, 73.9, 53.1, 52.7, 52.4, 36.3, 31.2, 26.6, 22.2. ESI-MS (m/z) of compound 5 calculated for $C_{42}H_{43}N_{12}O_{10}S_4$ ($[M + H]^+$) 1003.20, observed as 1003.50.

Synthesis of non-radioactive probe (GCPA)₂. The labeling group AMBF₃ was prepared with previous method.³⁵ Compound 5 (26 mg, 0.026 mmol) was dissolved in a solution of DMF (4 mL) and H₂O (2 mL), followed by AMBF₃ (46 mg, 0.23 mmol), tri(2-benzimidazole methyl)amines (ligand, 2.1 mg, 0.0052 mmol) and tetrakis(acetonitrile)copper(i)hexafluorophosphate (Cu(i), 12 mg, 0.031 mmol). The solution was heated to 45 °C for 2 h under N₂. The crude product was separated by semi-preparative HPLC to give non-radioactive probe (GCPA)₂ (30 mg, yield ratio 83%) as white solid, mp 130–132 °C. ¹H NMR of (GCPA)₂ (400 MHz, DMSO-*d*₆) δ 10.49 (s, 2H), 8.72 (s, 2H), 8.58 (d, *J* = 7.7 Hz, 2H), 8.45 (d, *J* = 7.6 Hz, 2H), 8.31 (s, 2H), 8.19 (d, *J* = 8.9 Hz, 4H), 7.94 (s, 2H), 7.78 (d, *J* = 7.8 Hz, 2H), 4.83 (t, *J* = 6.6 Hz, 6H), 4.75 (d, *J* = 7.1 Hz, 2H), 4.57–4.51 (m, 2H), 3.71 (t, *J* = 6.7 Hz, 4H), 3.25–3.18 (m, 2H), 3.08 (d, *J* = 12.7 Hz, 4H), 2.95 (d, *J* = 3.4 Hz, 16H), 2.85 (d, *J* = 13.1 Hz, 4H), 2.40–2.36 (m, 6H), 2.07–1.97 (m, 4H). ¹³C NMR of (GCPA)₂ (100 MHz, DMSO-*d*₆) δ 172.0, 171.3, 170.5, 170.4, 148.3, 143.4, 139.5, 137.1, 135.7, 125.3, 124.3, 121.4, 114.0, 112.1, 63.9, 54.2, 53.5, 52.6, 52.1, 44.0, 31.1, 28.4, 26.4. ¹⁹F NMR of (GCPA)₂ (375 MHz, DMSO-*d*₆) δ –135.8. ESI-MS (m/z) of (GCPA)₂ calculated for $C_{52}H_{67}B_2F_6N_{19}O_{10}S_4$ ($[M + H]^+$) 1395.42, observed as 1395.96.

Radiolabeling of (¹⁸F)GCPA₂

Radiolabeling of the non-radioactive probe (GCPA)₂ was carried out *via* ¹⁸F–¹⁹F isotope exchange reaction.^{35–37} In brief, [¹⁸F] fluoride in ¹⁸F/H₂¹⁸O (7.4–9.25 GBq) was captured by an activated Sep-Pak light QMA cartridge. Then it was eluted into a reaction tube using pyridazine-HCl buffer (pH 2.5, 300 μL). Subsequently, the non-radioactive probe (GCPA)₂ (25 mM, 30 μL) was added. The reaction was at 80 °C for 25 min to 30 min. The target product was capture using an activated Sep-Pak light C18 cartridge and then free [¹⁸F]fluoride was removed by deionized water (20 mL). Finally, the trapped ([¹⁸F]GCPA)₂ was eluted into a penicillin bottle using absolute ethanol (0.5 mL). Radiochemical yield, radiochemical purity of the target tracer ([¹⁸F]GCPA)₂ were measured by radio-HPLC. The RCY value was obtained with decay corrected. The molar activity was measured at the end of radiochemical synthesis.

Determination of log *P*

The log *P* value of ([¹⁸F]GCPA)₂ was determined according to the reported method.³⁸ In brief, ([¹⁸F]GCPA)₂ (0.74 MBq) was added to a saturated solution of octanol/distilled water (v/v, 1 : 1) and mixed sufficiently. And then each phase was separated *via* centrifugation at a speed of 3000 rpm min^{−1}. The distribution of ([¹⁸F]GCPA)₂ in each phase was quantified using a γ counter. $\log P = \log(C_o/C_w)$, where *C*_o represented the concentration of ([¹⁸F]GCPA)₂ in the octanol phase and *C*_w represented the concentration of ([¹⁸F]GCPA)₂ in the water phase.

In vitro stability assay

To evaluate the stability of the tracer ([¹⁸F]GCPA)₂, the tracer (37 KBq μL^{−1}, 20 μL) was incubated with PBS (pH = 7.4, 180 μL) and FBS (180 μL) at 37 °C for 1, 2 and 4 h, respectively. The tracer incubated in PBS was directly analyzed by the radio-HPLC without additional processing to assess the stability of ([¹⁸F]GCPA)₂ in PBS. For serum sample, the serum protein should be removed before radio-HPLC analysis. In detail, approximately an equal volume of acetonitrile was added to the serum sample (20 μL) to precipitate the protein, and then the mixture was centrifuged at 12 000 rpm for 5 min. Finally, the supernatant was analyzed by radio-HPLC to evaluate the stability of ([¹⁸F]GCPA)₂ in FBS.

GGT-responsive ability assay

The response of the probe ([¹⁸F]GCPA)₂ to GGT was investigated and monitored by analyzing whether the non-radioactive probe (GCPA)₂ could be specifically recognized and cleaved by GGT. At first, (GCPA)₂ (2.5 μmol) was dissolved in 100 μL of DMF to obtain a stock solution (25 mM), and GGT was dissolved in water to achieve a stock solution (10 000 U L^{−1}). The enzyme-reaction buffer consisted of Na₂HPO₄ (100 mM), NaH₂PO₄ (3.2 mM), KH₂PO₄ (1.8 mM), NaCl (27.5 mM) and 0.025% TritonX-100 (v/v). Then, (GCPA)₂ (25 mM, 1 μL) was incubated with 100 U L^{−1} of GGT in enzyme-reaction buffer (pH 7.4, 98 μL) at 37 °C for 30 min.

Cell culture and animal model

According to previous studies, human colon cancer cells HCT116 with high expression level of GGT and mouse fibroblast cells L929 with low expression level of GGT were selected for evaluating the biological performance of the probe.^{3,39,40} HCT116 and L929 cells were cultured in DMEM, and incubated at 37 °C and 5% CO₂. HCT116 cells (5 × 10⁶ cells) were inoculated subcutaneously. The mice were used for biological experiments when the tumor xenograft reached 8–10 mm in diameter. All animal procedures were followed in accordance with the Guidelines of the National Technical Committee for Standardization of Laboratory Animals of China (GB/T 358922018) and approved by the Animal Ethics Committee of the Jiangsu Institute of Nuclear Medicine.

In vitro biocompatibility assay

The biocompatibility of ([¹⁸F]GCPA)₂ was evaluated by measuring the cytotoxicity of non-radioactive probe (GCPA)₂ against HCT116 and L929 cells using MTT method. In brief, HCT116 and L929 cells were seeded in 96-well plate (8 × 10³ cells per well). Then cells were cultured in an incubator overnight to allow attachment. Then cells were incubated with DMEM medium containing different concentrations of non-radioactive probe (GCPA)₂ (0, 12.5, 25, 50 and 100 μM). After incubation for 6, 12 and 24 h, MTT (5 mg mL^{−1}, 20 μL) was added and then cells were incubated under the same conditions for another 4 h. Then the medium was removed and the formazan crystal was dissolved using 150 μL of dimethyl sulfoxide.



The optical density (OD) was measured with an ELISA microplate reader at the wavelength of 490 nm, and cells viability was evaluated. The well without drug treatment served as the control.

Cellular uptake study

The probe ($[^{18}\text{F}]\text{GCPA}$)₂ (37 KBq) was dispersed in 100 μL of FBS-free medium. Subsequently, it was added to a tube containing 1×10^6 HCT116 or L929 cells and 200 μL of medium ($n = 3$). Then cells were incubated at 37 °C for different time (15 min, 30 min, 1 h, 2 h and 4 h). Meanwhile, HCT116 cells pre-treated with non-radioactive probe (GCPA)₂ (10 nmol) for 30 min before incubation with ($[^{18}\text{F}]\text{GCPA}$)₂ served as the blocked group, and only the tracer ($[^{18}\text{F}]\text{GCPA}$)₂ (37 KBq) in tube ($n = 3$) served as the blank control group. At different time points (15 min, 30 min, 1 h, 2 h and 4 h), 500 μL of cold PBS was added to each tube to terminate cellular uptake and each sample was centrifuged at 4000 rpm min^{-1} . After supernatant was carefully pipetted out, each sample was carefully rinsed by 500 μL of cold PBS for three times. The activity of each tube was measured using a γ counter. Cellular uptake% = (mean counts in experimental group/mean counts in blank control group) \times 100%.

MicroPET imaging

PET imaging was performed using a small-animal PET scanner. The mice were anesthetized with isoflurane (2% isoflurane mixed with oxygen) at the flow rate of 2 L min^{-1} . HCT116 tumor-bearing mice were injected with 5.6 MBq of ($[^{18}\text{F}]\text{GCPA}$)₂ *via* tail vein after mice were anesthetized. Two control groups were used to verify the specificity of the radiotracer for imaging GGT. One group was co-injected with ($[^{18}\text{F}]\text{GCPA}$)₂ (5.6 MBq) and non-radioactive probe (GCPA)₂ (250 nmol), and another group was pretreated with GGsTop (500 nmol) for 30 min before injection of ($[^{18}\text{F}]\text{GCPA}$)₂ (5.6 MBq). Subsequently, dynamic microPET scanning was performed for 60 min. The imaging profile was reconstructed using three-dimensional ordered subsets expectation maximization (OSEM3D) model. Quantitative analysis of PET images was processed using ASIProVM software (Siemens Preclinical Solutions, USA). The tissue uptake was analyzed using the region of interest (ROI) method and expressed as % ID g^{-1} . The muscle ROI was drawn in the armpit muscle of the upper limb at the opposite side of the tumor.

In vivo pharmacokinetics

The tracer ($[^{18}\text{F}]\text{GCPA}$)₂ (~ 7.4 MBq) was dispersed in 200 μL of saline solution. Subsequently, it was intravenously injected into normal mice ($n = 3$). Meanwhile, ($[^{18}\text{F}]\text{GCPA}$)₂ (~ 1.5 MBq) was placed in a tube as control ($n = 3$). Blood samples were collected at 1, 2, 5, 7, 10, 15, 20, 30, 45, 60 and 120 min post injection, respectively. After measuring the weight and the radioactivity of every samples, the pharmacokinetics was calculated by DAS2.1 software.

Statistical analysis

Results were presented as mean \pm standard error (SD). All statistical analysis was performed using the SPSS 17.0 software.

Comparison between groups was analyzed by one-way ANOVA and *t*-test. * $P < 0.05$, ** $P < 0.01$ and *** $P < 0.001$ were considered to be statistically significant.

Author contributions

Dingyao Gao: methodology, software, investigation, formal analysis, visualization, writing-original draft. Yinxing Miao: software, investigation, formal analysis, visualization. Siqin Ye: methodology, investigation, formal analysis, visualization. Chunmei Lu: software, formal analysis, visualization. Gaochao Lv: methodology, writing-review & editing. Ke Li: formal analysis, methodology. Chunjing Yu: conceptualization, resources, supervision. Jianguo Lin: conceptualization, methodology, writing-review & editing, supervision. Ling Qiu: conceptualization, methodology, resources, writing-review & editing, supervision.

Conflicts of interest

There are no conflicts of interest to declare.

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References

- 1 Y. J. Reo, M. Dai, Y. J. Yang and K. H. Ahn, *Anal. Chem.*, 2020, **92**, 12678–12685.
- 2 T. Nishihara, H. A. Yoshihara, H. Nonaka, Y. Takakusagi, F. Hyodo, K. Ichikawa, E. Can, J. A. Bastiaansen, Y. Takado, A. Comment and S. Sando, *Angew. Chem.*, 2016, **55**, 10626–10629.
- 3 J. Ou-Yang, Y. F. Li, P. Wu, W. L. Jiang, H. W. Liu and C. Y. Li, *ACS Sens.*, 2018, **3**, 1354–1361.
- 4 Y. Zhang, G. Zhang, P. Yang, B. Moosa and N. M. Khashab, *ACS Appl. Mater. Interfaces*, 2019, **11**, 27529–27535.
- 5 F. Liu, Z. Wang, W. Wang, J. G. Luo and L. Kong, *Anal. Chem.*, 2018, **90**, 7467–7473.
- 6 J. W. Keillor, R. Castonguay and C. J. M. E. Lherbet, *Methods Enzymol.*, 2005, **401**, 449–467.
- 7 B. Flahou, F. Haesebrouck, K. Chiers, K. Van Deun, L. De Smet, B. Devreese, I. Vandenberghe, H. Favoreel, A. Smet, F. Pasmans, K. D'Herde and R. Ducatelle, *Cell. Microbiol.*, 2011, **13**, 1933–1955.
- 8 S. Ahmed, A. Zahoor, M. Ibrahim, M. Younus, S. Nawaz, R. Naseer, Q. Akram, C. L. Deng and S. C. Ojha, *Oxid. Med. Cell. Longevity*, 2020, **2020**, 7087921.
- 9 H. S. Choi, K. J. Kim, Y. Rhee and S. K. Lim, *Endocrinol. Metab.*, 2016, **31**, 64–71.



- 10 D. N. Abarbanel, S. M. Seki, Y. Davies, N. Marlen, J. A. Benavides, K. Cox, K. C. Nadeau and K. L. Cox, *J. Clin. Immunol.*, 2013, **33**, 397–406.
- 11 L. W. Chen, P. R. Huang, C. H. Chien, C. L. Lin and R. N. Chien, *J. Formosan Med. Assoc.*, 2020, **119**, 173–181.
- 12 Y. Chen, Q. Zheng, X. Jia, K. Chen, Y. Wang, T. Wu, X. Xu, Z. Han, Z. Zhang and X. Zhang, *Front. Plant Sci.*, 2019, **10**, 994.
- 13 Y. J. Kim, S. J. Park, C. S. Lim, D. J. Lee, C. K. Noh, K. Lee, S. J. Shin and H. M. Kim, *Anal. Chem.*, 2019, **91**, 9246–9250.
- 14 Z. Luo, L. Feng, R. An, G. Duan, R. Yan, H. Shi, J. He, Z. Zhou, C. Ji, H. Y. Chen and D. Ye, *Chem.–Eur. J.*, 2017, **23**, 14778–14785.
- 15 Z. Hai, Y. Ni, D. Saimi, H. Yang, H. Tong, K. Zhong and G. Liang, *Nano Lett.*, 2019, **19**, 2428–2433.
- 16 I. S. Fentiman and D. S. Allen, *Br. J. Cancer*, 2010, **103**, 90–93.
- 17 H. Li, Q. Yao, F. Xu, N. Xu, R. Duan, S. Long, J. Fan, J. Du, J. Wang and X. Peng, *Biomaterials*, 2018, **179**, 1–14.
- 18 J. Liu, S. Zhang, B. Zhao, C. Shen, X. Zhang and G. Yang, *Biosens. Bioelectron.*, 2019, **142**, 111497.
- 19 S. Li, R. Hu, C. Yang, X. Zhang, Y. Zeng, S. Wang, X. Guo, Y. Li, X. Cai, S. Li, C. Han and G. Yang, *Biosens. Bioelectron.*, 2017, **98**, 325–329.
- 20 R. An, S. Wei, Z. Huang, F. Liu and D. Ye, *Anal. Chem.*, 2019, **91**, 13639–13646.
- 21 S. Ye, S. Wang, D. Gao, K. Li, Q. Liu, B. Feng, L. Qiu and J. Lin, *Bioconjugate Chem.*, 2020, **31**, 174–181.
- 22 M. A. Klenner, G. Pascali, M. Massi and B. H. Fraser, *Chem.–Eur. J.*, 2021, **27**, 861–876.
- 23 J. Zhao, J. Chen, S. Ma, Q. Liu, L. Huang, X. Chen, K. Lou and W. Wang, *Acta Pharm. Sin. B*, 2018, **8**, 320–338.
- 24 Y. Zhan, F. Ai, F. Chen, H. F. Valdovinos, H. Orbay, H. Sun, J. Liang, T. E. Barnhart, J. Tian and W. Cai, *Small*, 2016, **12**, 2872–2876.
- 25 J. A. Hendricks, E. J. Keliher, D. Wan, S. A. Hilderbrand, R. Weissleder and R. Mazitschek, *Angew. Chem.*, 2012, **51**, 4603–4606.
- 26 D. Ni, E. B. Ehlerding and W. Cai, *Angew. Chem.*, 2019, **58**, 2570–2579.
- 27 Z. Luo, R. An and D. Ye, *ChemBioChem*, 2018, **20**, 474–487.
- 28 J. Lin, W. Wang, K. Li, H. Huang, G. Lv, Y. Peng, S. Luo and L. Qiu, *Chem. Commun.*, 2017, **53**, 6476–6479.
- 29 B. Shen, J. Jeon, M. Palner, D. Ye, A. Shuhendler, F. T. Chin and J. Rao, *Angew. Chem.*, 2013, **52**, 10511–10514.
- 30 Q. Zhou, S. Shao, J. Wang, C. Xu, J. Xiang, Y. Piao, Z. Zhou, Q. Yu, J. Tang, X. Liu, Z. Gan, R. Mo, Z. Gu and Y. Shen, *Nat. Nanotechnol.*, 2019, **14**, 799–809.
- 31 J. Huang and S. Whitman, *J. Liq. Chromatogr. Relat. Technol.*, 2006, **29**, 2513–2520.
- 32 Z. Li, Z. Wu, K. Chen, F. T. Chin and X. Chen, *Bioconjugate Chem.*, 2007, **18**, 1987–1994.
- 33 Y. Chen, A. Lisok, S. Chatterjee, B. Wharram, M. Pullambhatla, Y. Wang, G. Sgouros, R. C. Mease and M. G. Pomper, *Bioconjugate Chem.*, 2016, **27**, 1655–1662.
- 34 M. Wang, H. Wang, C. Q. Niu, T. Zhang, Z. Wu and Z. Li, *Bioconjugate Chem.*, 2020, **31**, 1795–1803.
- 35 Z. Liu, M. Pourghiasian, F. Benard, J. Pan, K. S. Lin and D. M. Perrin, *J. Nucl. Med.*, 2014, **55**, 1499–1505.
- 36 A. Roxin, C. Zhang, S. Huh, M. Lepage, Z. Zhang, K. S. Lin, F. Benard and D. M. Perrin, *Bioconjugate Chem.*, 2019, **30**, 1210–1219.
- 37 C. Zhang, Z. Zhang, K. S. Lin, J. Lau, J. Zeisler, N. Colpo, D. M. Perrin and F. Benard, *Mol. Pharm.*, 2018, **15**, 2116–2122.
- 38 Y. Zhang, L. Wang, S. Yu, K. Hu, S. Huang, Y. Li, H. Wu, H. Li and Q. Wang, *Contrast Media Mol. Imaging*, 2019, **2019**, 6315954.
- 39 J. Ou-Yang, Y. Li, W. L. Jiang, S. Y. He, H. W. Liu and C. Y. Li, *Anal. Chem.*, 2019, **91**, 1056–1063.
- 40 X. Hou, F. Zeng and S. Wu, *Biosens. Bioelectron.*, 2016, **85**, 317–323.

