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Targeted Melanoma Radiotherapy Using Ultrasmall ¹⁷⁷Lu-Labeled α -Melanocyte Stimulating Hormone-Functionalized Core-Shell Silica Nanoparticles

Xiuli Zhang^{a,b,1}, Feng Chen^{c,1}, Melik Z. Turker^d, Kai Ma^e, Pat Zanzonico^f, Fabio Gallazzi^g, Manankumar A. Shah^{a,b}, Austin R. Prater^{a,b}, Ulrich Wiesner^d, Michelle S. Bradbury^{c,h}, Michael R. McDevitt^c, Thomas P. Quinn^{a,b,*}

^aHarry S. Truman Veterans' Hospital, 800 Hospital Dr., Columbia, MO 65201 United States

^bDepartment of Biochemistry, University of Missouri, Columbia, Missouri 65211, United States

^cDepartment of Radiology, Sloan Kettering Institute for Cancer Research, New York, New York 10065, United States

^dDepartment of Materials Science and Engineering, Cornell University, Ithaca, New York 14853, United States

^eElucida Oncology, New York, NY 10016 United States

^fDepartment of Medical Physics, Sloan Kettering Institute for Cancer Research, New York, New York 10065, United States

^gDepartment of Chemistry and Research Core Facilities, University of Missouri, Columbia, Missouri 65211, United States

^hMolecular Pharmacology Program, Sloan Kettering Institute for Cancer Research, New York, NY 10065, United States

*Corresponding author. Thomas P. Quinn, Biochemistry Department, 117 Schweitzer Hall, University of Missouri, Columbia, MO 65211 USA, QuinnT@missouri.edu, *Tel*: +1-573-882-6099, *Fax*: +1-573-882-5635.

Author contributions

T.P.Q., M.R.M., M.S.B., F.C. and U.W. designed the experiments and oversaw data analysis and manuscript writing. X.Z., F.C., M.A.S., A.R.P. performed in vitro nanoparticle cell studies and in vivo nanoparticle imaging and therapy studies. M.Z.T. and K.M. performed nanoparticle synthesis and characterization. P.Z. performed dosimetry calculations. F.G. performed peptide synthesis.

¹Xiuli Zhang and Feng Chen contributed equally to this work

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Data Availability

Data will be made available on request.

Appendix A. Supplementary data

Supplementary data to this article can be found online.

Declaration of interests

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

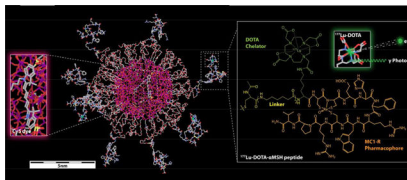
Competing financial interests

M.S.B. and U.W. hold interest in Elucida Technologies, Inc., which has licensed IP from Cornell and MSKCC on C' dots and their application in oncology.

Abstract

Lutetium-177 (^{177}Lu) radiolabeled ultrasmall (~6 nm dia.) fluorescent core-shell silica nanoparticles (Cornell prime dots or C' dots) were developed for improving efficacy of targeted radiotherapy in melanoma models. PEGylated C' dots were surface engineered to display 10–15 alpha melanocyte stimulating hormone (αMSH) cyclic peptide analogs for targeting the melanocortin-1 receptor (MC1-R) over-expressed on melanoma tumor cells. The ^{177}Lu -DOTA- αMSH -PEG-C' dot product was radiochemically stable, biologically active, and exhibited high affinity cellular binding properties and internalization. Selective tumor uptake and favorable biodistribution properties were also demonstrated, in addition to bulk renal clearance, in syngeneic B16F10 and human M21 xenografted models. Prolonged survival was observed in the treated cohorts relative to controls. Dosimetric analysis showed no excessively high absorbed dose among normal organs. Correlative histopathology of *ex vivo* treated tumor specimens revealed expected necrotic changes; no acute pathologic findings were noted in the liver or kidneys. Collectively, these results demonstrated that ^{177}Lu -DOTA- αMSH -PEG-C' dot targeted melanoma therapy overcame the unfavorable biological properties and dose-limiting toxicities associated with existing mono-molecular treatments. The unique and tunable surface chemistries of this targeted ultrasmall radiotherapeutic, coupled with its favorable pharmacokinetic properties, substantially improved treatment efficacy and demonstrated a clear survival benefit in melanoma models, which supports its further clinical translation.

Graphical Abstract



Keywords

ultrasmall silica nanoparticle; α -melanocyte stimulating hormone; melanoma; radionuclide therapy

1. Introduction

Revolutionary advances in metastatic melanoma treatment have been made over the past decade with targeted mutant B-Raf and MEK receptor tyrosine kinase (RTK) inhibitors and by checkpoint blockade immunotherapies [1–3]. Rapid and systemic elimination of tumors is often observed in patients treated with RTK inhibitors, however most patients relapse with lethal drug-resistant disease [4–5]. While systemic checkpoint blockade immunotherapy with cytotoxic T-lymphocyte antigen 4 (CTLA-4) and programmed cell death-1 (PD-1) antibodies has led to dramatic therapeutic responses, progression-free survival for combination immunotherapy has only been about 11.5 months [6]. Initial stereotactic radiotherapy plus combination immunotherapy studies have shown striking results in a subset of patients, fostering new clinical trials. [7]. Melanoma radiotherapy alone has

resulted in the shrinkage of distant tumors through the abscopal effect in a limited number of patients, and its immunological correlate has shown dramatic results in several case reports [8]. Despite exciting clinical results, immunotherapy side effects categorized as immune-related adverse events can lead to reduced efficacy or, in some instances, life-threatening conditions. Although there is considerable excitement around the application of immunotherapy and combination immunotherapy regimens to the treatment of metastatic melanoma, new synergistic therapeutic options are critically needed to simultaneously block multiple pathways controlling cancer cell proliferation.

Targeted radionuclide therapy for the systemic treatment of metastatic melanoma is an important clinical advance in the adjuvant setting. Preclinical studies, as well as several clinical trials, have demonstrated efficacy of small molecule, peptide and antibody targeting radiotherapies for treating melanoma [9]. Melanin targeting benzamides and antibodies have shown therapeutic efficacy in preclinical models and in limited clinical trials [10–11], while novel melanin targeting peptides have shown tumor growth control in melanoma-bearing mice [12]. Melanocortin-1 receptor (MC1-R) targeting alpha-melanocyte stimulating hormone (α MSH) peptides, radiolabeled with alpha- and beta-emitting radionuclides, have shown effective growth control of melanoma or its elimination in murine melanoma models [13–15]. In addition, the very late antigen-4 (VLA-4) integrin-targeting peptides are promising for melanoma-targeted therapy [16–17]. However, dose-limiting toxicity has undermined clinical translation of these radionuclide therapies at doses necessary to achieve tumor eradication. Off-target accumulation in pigmented eyes with radiolabeled melanin-targeting therapeutics can limit administered activities, while extended circulation times for radiolabeled antibodies increases the risk of marrow toxicity [10, 12]. Moreover, non-specific renal uptake of radiolabeled peptides may impose an upper limit on administered therapeutic activities [18]. In addition, normal integrin expression in spleen and bone may be dose-limiting for cognate radioligands [16]. In summary, although studies to date highlight the effectiveness of targeted radiotherapies for treating melanoma, it will be crucial to demonstrate favorable pharmacokinetic (PK) and clearance profiles – without accompanying dose-limiting toxicity – for successful clinical translation.

Ultrasmall fluorescent core-shell silica nanoparticles, referred to as Cornell prime dots (C' dots), are silica core polyethylene glycol (PEG) shell nanoparticles (< 8 nm dia.) that have the potential to effectively deliver therapeutic radionuclides to sites of melanoma while overcoming many of the dose-limiting toxicities associated with current molecular therapeutics [19–22]. The PK and clearance properties of C' dots are situated between those of antibodies and peptides [21, 23]. Radiolabeled C' dots exhibit favorable clearance from the blood, and are primarily excreted through the kidneys, ultimately reducing the potential for marrow, liver, and/or colonic mucosal toxicity by limiting circulation times and largely avoiding hepatobiliary clearance [24]. Uptake of radiolabeled C' dots in liver and other reticuloendothelial system (RES) tissues is low, as is kidney uptake [25]. This property is critically important, given the radiosensitivity of the kidneys. In addition, this overcomes dose-limiting toxicities associated with the high non-specific renal uptake of radiometal-labeled peptides and antibody fragments. Phase 1, first-in-human imaging studies with integrin targeting ^{124}I -cRGDY-PEG-C dots previously demonstrated that the particle tracer was non-toxic and localized at sites of disease [26]. α MSH-bearing C' dots, or α MSH-PEG-

C' dots, which target a key melanoma biomarker, MC1-R, have been synthesized and radiolabeled for imaging [27]. Radio-iodinated and ^{89}Zr radiolabeled $\alpha\text{MSH-PEG-C'}$ dots were found to exhibit receptor-mediated uptake, along with favorable PK and target-to-background ratios [27].

In this study we report, for the first time, on the therapeutic efficacy of complexing agent 1,4,7,10-tetraazacyclododecane-1,4,7,10-tetraacetic acid (DOTA) functionalized $\alpha\text{MSH-PEG-C'}$ dots, or DOTA- $\alpha\text{MSH-PEG-Cy5-C'}$ dots, radiolabeled with the β -emitting radionuclide ^{177}Lu ($E_{\beta\text{max}}$ 0.497 MeV; $t_{1/2}$ =6.65 days) in syngeneic and xenografted melanoma models (Fig. 1). To that end, ultrasmall PEGylated C' dots, covalently encapsulating the near infra-red (NIR) emitting dye Cy5, were engineered to display MC1-R targeting cyclic DOTA- αMSH peptides on their surfaces, according to published procedures [27]. The DOTA- $\alpha\text{MSH-PEG-Cy5-C'}$ dots contained an average of 1.6 dyes and 12.1 DOTA- αMSH peptides per nanoparticle. DOTA- $\alpha\text{MSH-PEG-Cy5-C'}$ dots were directly radiolabeled with ^{177}Lu , and assessed for active cell binding and favorable biodistribution profiles in B16F10 syngeneic and M21 xenograft models. Radiolabeled nanoparticles were radiochemically stable and rapidly internalized in B16F10 and M21 melanoma cells. Biodistribution studies revealed a relatively rapid clearance of radioactivity from the circulation, coupled with bulk renal excretion. Substantial tumor uptake was observed over the first 24 h. Single-photon emission tomography (SPECT) imaging with $^{177}\text{Lu-DOTA-}\alpha\text{MSH-PEG-Cy5-C'}$ dots showed tumor uptake in M21 melanoma-bearing mice that could be blocked with excess non-radiolabeled DOTA- $\alpha\text{MSH-PEG-Cy5-C'}$ dots, suggesting specificity. Therapy studies showed that $^{177}\text{Lu-DOTA-}\alpha\text{MSH-PEG-Cy5-C'}$ dot treatment of B16F10 and M21 melanoma-bearing mice delayed tumor growth and increased median survival. These results support continued development of $^{177}\text{Lu-DOTA-}\alpha\text{MSH-PEG-Cy5-C'}$ dots for melanoma radiotherapy.

2. Materials and methods

2.1. MC1-R targeting DOTA- αMSH peptide synthesis

The MC1-R targeting rhenium (Re) cyclized peptide DOTA- αMSH [Ac-Cys-(Ahx)₂-dLys(DOTA)-Re(Cys-Cys-Glu-His-dPhe-Arg-Trp-Cys)-Arg-Pro-Val-CONH₂] was synthesized using standard fluorenylmethyloxycarbonyl (Fmoc) solid-phase peptide synthesis (SPPS) chemistry on a Tetras peptide synthesizer (Advanced ChemTech). The peptides were purified using semi-preparative mass spectrometry (MS; LCQ-Fleet, Thermo-Fisher Scientific) guided high-performance liquid chromatography (HPLC).

2.2. DOTA- $\alpha\text{MSH-PEG-Cy5-C'}$ dot synthesis

The synthesis of DOTA- $\alpha\text{MSH-PEG-Cy5-C'}$ dot particles was based on procedures previously described [20, 27–28]. The resulting DOTA- $\alpha\text{MSH-PEG-Cy5-C'}$ dots were dialyzed against deionized water, purified using gel permeation chromatography (GPC), and filtered by sterile syringe filters. The final product was characterized and stored at 4°C [29–30].

2.3. Cell culture

B16F10 murine melanoma cells were obtained from American Type Culture Collection (ATCC). The M21 human melanoma cell line was obtained from D. A. Chersesh (University of California San Diego). The cell lines were cultured in RPMI 1640 supplemented with 10% fetal calf serum, 2 mM L-glutamine and Penicillin Streptomycin.

2.4. DOTA- α MSH-PEG-Cy5-C' dot IC₅₀ assays

The concentration of DOTA- α MSH peptide and DOTA- α MSH-PEG-Cy5-C' dots needed to inhibit 50% binding of a MC1-R standard radioligand was determined with B16F10 murine melanoma cells [27]. A competitive binding assay was performed with DOTA- α MSH and DOTA- α MSH-PEG-Cy5-C' dots with ¹²⁵I-NDP over a concentration range of 10⁻¹³ to 10⁻⁵ M using B16F10 melanoma cells. The IC₅₀ values for DOTA- α MSH and DOTA- α MSH-PEG-Cy5-C' dots were calculated using GraphPad Prism V 7.03.

2.5. DOTA- α MSH-PEG-Cy5-C' dot confocal imaging studies

B16-F10 and M21 cells were incubated with DOTA- α MSH-PEG-C' dots (1.5 μ M) for 4 hours, while LysoTracker Green (100 nM; Invitrogen) and Hoechst 33342 (10 μ g/mL; Life Technologies) were added 30 minutes and 5 min prior to imaging. A Leica SP8 WLL confocal microscope (Leica Microsystems, Inc.) equipped with 405 nm and tunable white light lasers, and a 40x/1.1NA water immersion objective was used for imaging. The excitation/emission band-path wavelengths used to detect Hoechst 33342, LysoTracker Green, and DOTA- α MSH-PEG-Cy5-C' dots were set to 405/430–480, 488/500–550, and 650/655–725 nm, respectively. All images were deconvolved with Huygens Professional 19.04 software program (SVI, Netherlands).

2.6. ¹⁷⁷Lu-DOTA- α MSH-PEG-Cy5-C' dot labeling and stability

¹⁷⁷Lu chloride was obtained from the University of Missouri Research Reactor (MURR). For C' dot radiolabeling, ¹⁷⁷Lu in 50 μ L of NH₄Ac buffer (pH 5.5) was added to a vial containing a 10 \times molar excess of 15 μ M DOTA- α MSH-PEG-Cy5-C' dots. The pH of the labeling reaction was adjusted to 5.5 if needed prior to incubation at 70°C for 20 minutes. The labeling reaction was quenched by adding 10 μ L of 0.2 M EDTA and incubated for another 1 h at 37°C prior to use. The radiochemical yield was determined by ITLC. ¹⁷⁷Lu-DOTA- α MSH-PEG-Cy5-C' dot stability was determined by incubating 100 μ L of the ¹⁷⁷Lu-DOTA- α MSH-PEG-Cy5-C' dot solution with 100 μ L of water, phosphate buffered saline (PBS), mouse serum or human serum at 37°C for 1 h, 4 h, and 24 h then assayed by instant thin layer chromatography (ITLC).

2.7. Cellular binding and internalization of ¹⁷⁷Lu-DOTA- α MSH-PEG-Cy5-C' dots

B16F10 murine melanoma cells, M21 human melanoma cells, human embryo kidney-293 (HEK293) cells, retinal epithelia cells (ARPE-19), and normal prostate epithelial cells (RWPE-1) were plated in a 24 well plates (0.2 million/well) and grown overnight at 37°C. Approximately, 100,000 CPM of ¹⁷⁷Lu-DOTA- α MSH-PEG-Cy5-C' dots were added to each well for cell binding studies at 25°C. For specificity studies, 100,000 CPM of ¹⁷⁷Lu-DOTA- α MSH-PEG-Cy5-C' dots or ¹⁷⁷Lu-DOTA- α MSH-PEG-Cy5-C' dots + 5 μ g of NDP

peptide were added to each well. At 6 h, 18 h and 24 h post-incubation, the cells were rinsed and surface-bound radioactivity was obtained with an acidic buffer wash (40 mM sodium acetate pH 4.5, 0.9% NaCl and 0.02% BSA). Internalized activity was obtained by lysing cells with 1 M NaOH after the acid buffer wash. Quantitation of radioactivity in the membrane-bound and the internalized fractions was performed with a γ -counter.

2.8. ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dot biodistribution studies

Animal studies were conducted in compliance with Institutional Animal Care and Use Committee rules for animal experimentation at the University of Missouri, Harry Truman Veterans' Hospital and Memorial Sloan Kettering Cancer Center. Female and male C57 and nude mice were inoculated subcutaneously in the flank or shoulder with 1 million cultured B16/F10 murine melanoma cells or 10 million M21 human melanoma cells (2:1 PBS:matrigel), respectively. For biodistribution studies, a 5 μCi solution of ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots were injected into the tail veins of B16F10 ($n=4$) and M21 ($n=3$) tumor-bearing mice when tumors reached 0.5 g. At prescribed p.i. time points, mice were sacrificed and major organs and tissues removed, weighed, and radioactivity quantitated using a γ -counter. Data were calculated as a percentage of the injected dose (% ID) or percentage of the injected dose per gram (% ID/g) of tissue, plus or minus the standard deviation (\pm s.d.).

2.9. Imaging studies

SPECT imaging studies were performed in M21 tumor-bearing SCID mice injected with 800 μCi ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots or ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots + 200 μg NDP peptide. Imaging was conducted 24 h p.i. on a Siemens Inveon SPECT/CT small animal imaging system (Siemens Medical Solutions).

2.10. Dosimetry

Organ time-activity data were fit to exponential functions and analytically integrated, incorporating the effect of the radioactive decay of the ^{177}Lu , to derive the time-integrated activities. The mouse organ time-integrated activities were scaled by mouse-to-man total-body mass ratios to yield the organ time-integrated activities for the 70-kg Reference Adult. The Reference Adult mean normal absorbed doses were then calculated using the OLINDA computer program [31]. The mean mouse tumor absorbed doses were calculated using the tumor time-activity data and time-integrated activities and assuming complete local absorption of the ^{177}Lu beta particles and ignoring the small contribution of the ^{177}Lu gamma rays.

2.11. Therapy studies

Mice were inoculated with B16F10 or M21 melanoma cells in their flanks, as described in the biodistribution section. Therapy studies began when M21 and B16F10 tumors reached 0.01 cm^3 and 0.005 cm^3 , respectively. Groups of M21 mice ($n = 8$) were i.v.-injected with 1 mCi and 0.5 mCi of ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots, 1 mCi and 0.5 mCi ($n=4$) of ^{177}Lu -DOTA-PEG-Cy5-C' dots, DOTA- α MSH-PEG-Cy5-C' dots, or PBS control. Groups of B16F10 mice ($n = 10$) were also injected with 0.5 mCi of ^{177}Lu -DOTA- α MSH-PEG-

Cy5-C' dots, 0.5 mCi of ^{177}Lu -DOTA-PEG-Cy5-C' dots, DOTA- α -MSH-PEG-Cy5-C' dots, or PBS control. All mice were monitored twice weekly for tumor size, weight, and body score index. Tumor volumes were calculated by caliper measurements of their length, width, and depth. Mice were removed from the study and sacrificed if body weight loss was >15% of the initial weight, tumor size exceeded 1.0 cm³, a body score index of < 2 (0–5 scale), the appearance of skin ulcerations at the tumor site, or abnormal behaviors.

2.12. Statistical analysis

IC₅₀ calculations were performed using Prism (Graphpad, Inc.) using non-linear regression data analysis and 95% confidence intervals. Therapy study survival curve evaluation and graph generation were performed using Prism-7 and Kaplan-Meier analysis [32]. Statistical comparisons between experimental groups were performed by Student's t-test (unpaired, two-tailed) or Gehan-Breslow-Wilcoxon test depending on the analysis [33]. Statistical significance for all tests was set at $p < 0.05$.

3. Results and discussion

3.1. MC1-R targeting peptide and DOTA- α -MSH-PEG-Cy5-C' dot synthesis

The MC1-R targeting DOTA- α -MSH cyclic peptide [Ac-Cys-(Ahx)₂-dLys(DOTA)-Re(Cys³-Cys⁴-Glu⁵-His⁶-dPhe⁷-Arg⁸-Trp⁹-Cys¹⁰)-Arg¹¹-Pro¹²-Val¹³-NH₂] used in C' dot synthesis was produced by solid phase peptide synthesis using standard Fmoc chemistry. Linear DOTA- α -MSH peptide was cyclized on resin via rhenium (Re) coordination between three orthogonally deprotected cysteine residues (Cys^{3,4,10}) flanking the receptor binding region (Fig. S1a). The peptide was deprotected and cleaved from the resin. Crude peptide was purified to homogeneity using liquid chromatography-mass spectrometry (LC-MS) guided high-performance liquid chromatography (HPLC) purification (Fig. S1b). Analytical LC-MS analysis confirmed the final product had the expect mass (Fig. S1c). DOTA was selected as the radiometal chelator due to its ability to stably coordinate a number of therapeutic radionuclides, including ^{177}Lu . The dual 6-amino hexanoic acid (Ahx)₂ linker was employed to separate the receptor binding pharmacophore from the amino-terminal Ac-Cys nanoparticle attachment site.

Cyclic DOTA- α -MSH peptide-bearing C' dots were prepared using a water-based synthesis protocol described previously, yielding ultrasmall (~6 nm dia.) silica nanoparticles (Fig. 1) [20, 27–28]. In short, during the synthetic process silane-modified Cy5 dye molecules were added to the silica sol-gel reaction forming the fluorescent silica core. DOTA- α -MSH peptides were conjugated to maleimide-PEG-silane obtained from reaction of heterobifunctional maleimide-PEG-N-hydroxysuccinimide ester (mal-PEG-NHS) with aminopropyl-silane. The conjugate was then added together with regular PEG-silane to form the PEG shell of the silica nanoparticle. The resulting DOTA- α -MSH-PEG-Cy5-C' dots were purified by gel permeation chromatography (GPC, Fig. 2a). A representative TEM of the GPC purified DOTA- α -MSH-PEG-Cy5-C' dots, demonstrated uniform morphology, is shown in the Fig. 2a-insert. The average hydrodynamic diameter of the DOTA- α -MSH-PEG-Cy5-C' dots was 5.8 nm as determined by fluorescence-correlation spectroscopy (FCS, Fig. 2b). FCS curve fits were also used to determine the concentration of the DOTA- α -MSH-

PEG-Cy5-C' dot particles [29–30]. Optical (UV-Vis) spectroscopy was employed together with FCS determined concentration information to determine the average number of DOTA- α MSH peptides and Cy5 dye molecules per particle (Fig. 2c–e) [34]. To determine the average number of peptides per particle a calibration using absorbance at 256 nm ($\epsilon = 8320 \text{ M}^{-1}, \text{ cm}^{-1}$) was employed (Fig. 2c–d). Absorbance contributions to the DOTA- α MSH-PEG-Cy5-C' dot spectra by DOTA- α MSH and Cy5 components, as revealed by a deconvolution method described earlier, are shown in Fig. 2e [34]. This spectral analysis determined that there were, on average, 12.1 DOTA- α MSH peptides and 1.6 Cy5 dyes per particle.

3.2. DOTA- α MSH-PEG-Cy5-C' in vitro receptor binding, uptake and affinity

A competitive binding assay, using the very high affinity MC1-R agonist ^{125}I -NDP [^{125}I -(Nle⁴, D-Phe⁷)- α MSH], [35] was performed to determine the IC_{50} for binding of DOTA- α MSH-PEG-Cy5-C' dots to MC1-R-expressing B16F10 melanoma cells (Fig. 2f). An IC_{50} value of $4.1 \times 10^{-10} \text{ M}$ was determined for the DOTA- α MSH-PEG-Cy5-C' dots, which was approximately $10 \times$ lower than the DOTA- α MSH peptide ($3.3 \times 10^{-9} \text{ M}$); this result highlights an increased affinity due to a multi-dentate peptide display on the C' dot surface. The IC_{50} value for the DOTA- α MSH-PEG-C' dots was comparable to that of α MSH-PEG-Cy5-C' dots ($6.6 \times 10^{-10} \text{ M}$), demonstrating that addition of the DOTA chelator did not adversely affect MC1-R binding [27].

Internalization of DOTA- α MSH-PEG-Cy5-C' dots in M21 and B16F10 melanoma cells was visualized by confocal microscopy (Fig. 3a). The Cy5-encapsulating silica core allowed direct detection of C' dots within live cells 4 h after addition to the culture media. The DOTA- α MSH-PEG-Cy5-C' dots were visualized within the cytoplasm of the melanoma cells (Fig. 3a(i)). Acidic cytoplasmic vesicles, such as lysosomal vesicles, were visualized using LysoTracker Green (Fig. 3a(ii)). Cellular nuclei were visualized by Hoechst fluorescent staining (Fig. 3a(iii)). Merged images of all three probes demonstrated that the DOTA- α MSH-PEG-Cy5-C' dots colocalized with the LysoTracker Green dye in the cytoplasm (Fig. 3a(iv)). These results showed that the majority of the internalized DOTA- α MSH-PEG-Cy5-C' dots were located in cytoplasmic acidic vesicles of both cell types, such as lysosomes, which is consistent with receptor-mediated particle endocytosis.

3.3. ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dot radiolabeling and cell binding

DOTA- α MSH-PEG-Cy5-C' dots were radiolabeled in $0.2 \text{ M NH}_4\text{OAc}$ at pH 5.5 with $^{177}\text{LuCl}_3$ at 70°C for 20 min. The molar ratio of DOTA- α MSH-PEG-Cy5-C' dots to $^{177}\text{LuCl}_3$ was 10:1. The reaction was quenched by the addition of EDTA (0.01M) and subsequent incubation at 37°C for 60 min. Radiochemical yields were typically $>95\%$. Any remaining free $^{177}\text{LuCl}_3$ was removed by PD-10 (Sephadex G-25) desalting columns yielding a final ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dot specific activity of $5.5 \times 10^6 \text{ Ci/mol}$. The radiochemical stability of ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots remained at $>90\%$ over a 24 h period at 37°C in normal saline and mouse serum (Table S1). *In vitro* cell binding studies were performed with ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots using B16F10 and M21 melanoma cells (Fig. 3b). ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots were bound and internalized by both the B16F10 and M21 melanoma cells. Membrane bound ^{177}Lu -DOTA-

α MSH-PEG-Cy5-C' dots were efficiently internalized over time. Cellular uptake could be blocked to background membrane bound levels using excess (5 μ g) MC1-R-avid NDP peptide demonstrating MC1-R specificity of internalization. Cellular binding and internalization of ^{177}Lu -DOTA- α MSHPEG-Cy5-C' dots were examined in human embryonic kidney (HEK293) cells, human eye epithelial cells (ARPE-19) and normal prostate (RWPE-1) cells that do not express the MC1-R (Fig. S2). Uptake of ^{177}Lu -DOTA- α MSH-PEG-C' dots in normal MC1-R negative cells were below the amount of radioactivity found in the receptor-blocked samples in Fig. 3b. These results showed no significant normal cell line uptake of ^{177}Lu -DOTA- α MSH-PEG-C' dots, which is consistent with the biodistribution data.

Published reports specify the MC1-R densities of M21 and B16F10 cell lines as 1,281 receptors/cell and 2,884 receptors/cell, respectively [36–37]. The difference in MC1-R numbers per cell did not significantly affect cellular uptake of ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots, underscoring the fact that a 2x difference in receptor numbers does not necessarily lead to a meaningful difference in internalized radioactivity. This is likely due to a lower specific activity of the ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dot preparation, which masks the difference in MC1-R numbers at early time points, but is evident at later time points. Both cell lines are in the median range for human melanoma MC1-R expression levels, which range from 900 to 5,500 receptors per cell [38]. In spite of relatively low receptor numbers, first in human melanoma imaging studies with a lactam cyclized α MSH peptide have validated targeting the MC1-R [39]. In summary, *in vitro* studies confirmed MC1-R selective binding and internalization of ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots, highlighting the efficient delivery of a particle-based radiotherapy to tumor cells.

3.4. ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots biodistribution, imaging and dosimetry

The biodistribution and PK profiles of intravenously (i.v.)-injected ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots into B16F10 and M21 tumor-bearing mice were determined over a period of 96 h (Fig. 4). The bolus of activity injected into the bloodstream steadily declined and was nearly undetectable by 96 h post-injection (p.i.). Organs with high blood perfusion initially showed high uptake at 4 h, which subsequently declined over the ensuing 24 h period. For example, cardiac uptake values in B16F10 mice were $10.2 \pm 2.82\%$ ID/g at 4 h and $3.92 \pm 0.76\%$ ID/g at 24 h, while those for lungs were $12.1 \pm 0.83\%$ ID/g at 4 h and $4.86 \pm 1.38\%$ ID/g at 24 h p.i. Activities at 24 h p.i. in the liver and spleen were also low, with values of $5.76 \pm 0.61\%$ ID/g and $5.10 \pm 1.24\%$ ID/g, respectively. The clearance kinetics of activity from the blood, heart, lungs, and RES organs were slightly slower in the M21 xenograft model than in the C57 B16F10 syngeneic model. Blood activity at 24 h p.i. was $17.6 \pm 1.28\%$ ID/g, approximately two times higher than that in B16F10 C57 mice. Uptake values in the lungs, liver, and spleen were nearly 1.5 times higher in the M21 tumor-bearing mouse at 24 h. Renal excretion was the primary route of clearance, with approximately 40% of the dose excreted at 24 h p.i. and nearly 80% excreted at 96 h p.i., in both animal models. Renal uptake values at 4 h, 24 h and 96 h p.i., respectively, were $14.6 \pm 1.14\%$ ID/g, $13.1 \pm 1.57\%$ ID/g and $4.42 \pm 1.06\%$ ID/g in M21 tumor-bearing mice and $14.2 \pm 1.18\%$ ID/g, $15.9 \pm 2.08\%$ ID/g and $13.2 \pm 2.90\%$ ID/g in B16F10 tumor-bearing mice.

We also measured renal uptake at 2 h p.i. ($5.27 \pm 1.18\%$ ID/g) in the B16F10 model for comparison with the α MSH peptide analog ^{177}Lu -DOTA-Re(Arg¹¹)CCMSH (Fig. S3a) [40]. Renal activities measured after injection of the ^{177}Lu -DOTA-Re(Arg¹¹)CCMSH peptide were $17.9 \pm 2.47\%$ ID/g, $19.1 \pm 2.38\%$ ID/g and $13.8 \pm 3.72\%$ ID/g at 2 h, 4 h and 24 h p.i., respectively [40]. Relative to the particle-based probe, radiolabeled peptides showed greater renal retention at the early time points. Retention of radioactivity in the kidneys was non-specific since activity levels did not change in the blocking studies (Fig. S3b–c). These results demonstrate how the unique surface chemistry of the ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots has the potential to mitigate kidney exposure compared to radiolabeled peptide alone. However, the level of renal uptake of the ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots was higher than that observed in prior biodistribution studies using ^{125}I and ^{89}Zr labeled α MSH-PEG-C' dots [27]. On the basis of previously published radiolabeling studies of the DOTARE(Arg¹¹)CCMSH peptide [13, 41–43], it is likely that the unique ^{177}Lu -DOTA chelator solution coordination geometry and net charge could have contributed to the higher renal retention seen with ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots.

Targeted tumor uptake of ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots was evident in both animal models. In M21 xenografts, tumor uptakes were $5.47 \pm 2.50\%$ ID/g and $9.29 \pm 4.74\%$ ID/g at 4 h and 24 h p.i., respectively, while tumor uptake in the B16F10 model was $10.4 \pm 3.56\%$ ID/g at 4 h and $9.56 \pm 1.57\%$ ID/g at 24 h. The progressively higher tumor uptakes from 4 h to 24 h p.i. are consistent with specific localization of the ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots in tumor. Tumor uptake was blocked by about 50% in both mouse models by co-injecting excess NDP peptide (Fig. 4a and Fig. 4b inserts), likewise suggesting that uptake was largely MC1-R mediated.

Tumor uptake and selective tumor blocking of ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots were visualized in the M21 model. The 208 keV gamma emission from ^{177}Lu allowed SPECT imaging of ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots *in vivo*. Mice were administered ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots alone or in the presence of excess i.v.-injected NDP peptide, and imaged 24 h p.i. (Fig. 5). SPECT-CT imaging results showed tumor uptake of ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots (Fig. 5a and Fig. 5b), which could be blocked by excess MC1-R selective NDP peptide (Fig. 5c and Fig. 5d). This imaging study underscored the MC1-R selective uptake *in vivo*. These findings further emphasize the complementary dual-modality imaging capabilities of ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots achievable with this platform, that is, high resolution optical imaging (Fig. 3a) alongside whole-body quantitative SPECT-CT imaging.

Organ absorbed doses were derived for a 70-kg Reference Adult using the MIRD formalism and the mouse ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dot time-activity data (Table 1) [31]. Low doses were found for liver and spleen, which was notable considering that these organs are often sites for non-specific accumulation of nanoparticle probes. Absorbed doses to other radiosensitive tissues, such as the red marrow and intestinal wall, were also low. The kidneys received the highest dose of any non-tumor tissue. Since the ^{177}Lu -DOTA- α MSHPEG-Cy5-C' dots exhibit primary renal clearance, a higher dose to the kidneys was not unexpected, but within tolerable limits. However, dosimetry findings were not significantly different from

those found using standard radiopharmaceuticals, suggesting that ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots are safe and a clinically-promising agent for translation.

3.5. ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots therapy studies

The therapeutic efficacy of ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dot treatment was examined in M21 and B16F10 melanoma-bearing mice (Fig. 6) Initial treatment doses were determined by maximum tolerated dose (MTD) studies in each model system. Mice were injected with 0.4 mCi, 0.7 mCi and 1 mCi of ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots. The mice were observed for changes in weight and behavior over a 7- or 14-day period. Changes in body weight are reported in Table S2. Athymic nude mice bearing M21 tumors were more sensitive to treatment than B16F10 tumor-bearing C57 mice. The MTD was determined to be 1 mCi dose, since it resulted in weight loss that exceeded 15% of the initial weight, and approached the weight criterion for removing an animal from the therapy study. M21 tumor-bearing mice were randomized according to tumor size into five cohorts of mice, each for treatment with: 1 mCi or 0.5 mCi of ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots, 1.0 mCi or 0.5 mCi of non-targeted ^{177}Lu -DOTA-PEG-Cy5-C' dots, non-radioactive DOTA- α MSH-PEG-Cy5-C' dots, or PBS control. All mouse groups receiving radiolabeled C' dots showed reduced tumor growth during the first 18 days of the study compared with groups receiving non-radiolabeled DOTA- α MSH-PEG-Cy5-C' dots or PBS control (Fig. 6a). Kaplan-Meier survival analysis revealed median survival times of 27, 41, 33, 60, 58 and 43 days for PBS control, DOTA- α MSH-PEG-Cy5-C' dots, ^{177}Lu -DOTA-PEG-Cy5-C' dots (0.5 mCi), ^{177}Lu -DOTA-PEG-Cy5-C' dots (1 mCi), ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots (0.5 mCi) and ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots (1.0 mCi), respectively. The Gehan-Breslow-Wilcoxon test demonstrated statistical difference for the survival data ($p=0.029$). A survival comparison of individual treatment groups to PBS control showed statistical significance in M21 tumor-bearing mice that received 0.5 mCi of ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots ($p=0.036$) and 1 mCi ^{177}Lu -DOTA-PEG-Cy5-C' dots ($p=0.01$). There was a significant survival advantage for the mice treated with ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots (0.5 mCi) compared to mice treated with 0.5 mCi of non-targeting ^{177}Lu -DOTA-PEG-Cy5-C' dots ($p=0.021$). Mice that received 1 mCi of ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots, 0.5 mCi ^{177}Lu -DOTA-PEG-Cy5-C' dots or non-radiolabeled DOTA- α MSH-PEG-Cy5-C' dots did not exhibit a statistically significant improvement in survival. Non-radiolabeled DOTA- α MSH-PEG-Cy5-C' dots were not expected to provide an increase in survival compared to PBS control. The lack of a survival benefit for mice receiving 1 mCi of ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots was likely due to its being at the MTD level, resulting in radiotoxicity and removal of mice from the study due to poor body score or weight loss. It is interesting that the mice receiving 1 mCi of non-MC1-R targeted ^{177}Lu -DOTA-PEG-Cy5-C' dots did show a survival benefit without the same degree of radiotoxicity. This is likely due to non-specific enhanced permeability and retention (EPR) effects at the target site, coupled with more rapid renal clearance, mimicking a reduced treatment dose. Cell binding studies with non-targeted ^{177}Lu -DOTA-PEG-Cy5-C' dots showed approximately 38% of the uptake found for targeted ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots at 24 h (Fig. S4 and Fig. 3b). This would equate the 1 mCi administered treatment dose to approximately 400 μCi , which is consistent with the therapy results. Blood chemistries showed no signs of hematological toxicities, and were nearly identical for all treatment groups compared to PBS or non-tumor

bearing controls, except for lymphocytes which were slightly lower in the ^{177}Lu -DOTA labeled C' dot groups (Table S3). Average body weights of mice in treatment groups receiving radioactive C' dots were not significantly different from those groups receiving non-radiolabeled C' dots or PBS controls, even though the numbers decreased due to removal of animals for tumor size and poor body score (Fig. S5a).

Treatment efficacy of ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots was also examined in B16F10 mice (Fig. 6c and Fig. 6d) receiving subcutaneous flank injections (i.e., 1 million B16F10 cells) 7 days prior to treatment. Mice were randomized according to body weights into treatment groups with equal numbers of male and female mice. The treatment dose was set at 0.5 mCi per mouse based on the results of the M21 therapy study. B16F10 melanoma mice were administered 0.5 mCi ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots, 0.5 mCi ^{177}Lu -DOTA-PEG-Cy5-C' dots, DOTA- α MSH-PEG-Cy5-C' dots, or a PBS control via the tail vein. The rate of tumor growth was lower in groups treated with radiotherapeutic C' dots than in groups receiving non-radiolabeled C' dots or PBS over the first 14 days of therapy (Fig. 6c). Kaplan Meier survival analysis yielded median survival times of 16, 14, 21, and 25 days for PBS control, DOTA-PEG-Cy5-C' dots, ^{177}Lu -DOTA-PEG-Cy5-C' dots (0.5 mCi), and ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots (0.5 mCi), respectively. Pairwise Gehan-Breslow-Wilcoxon statistical analyses demonstrated a significant survival improvement in mice treated with 0.5 mCi ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots compared with those receiving PBS ($p=0.0005$) or non-radiolabeled C' dots ($p=0.002$) (Fig. 6d). Likewise, in the B16F10 therapy study, there was a statistically significant difference seen between tumor-bearing mice treated with ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots and those treated with the non-targeted ^{177}Lu -DOTA-PEG-Cy5-C' dots ($p=0.045$). Average body weights of mice remaining in the treatment groups that received radioactive C' dots were not significantly different from those receiving non-radiolabeled C' dots or PBS controls (Fig. S5b).

The in vivo therapy studies mirrored in vitro treatment of B16F10 cells with 1 μCi and 5 μCi of ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots (Fig S6). The higher treatment dose showed no significant difference in tumor cell death between ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots and ^{177}Lu alone, while the lower dose did show a significant treatment advantage at later time points. The higher doses close to MTD resulted in high non-specific toxicities that masked the therapeutic potential of the targeted ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots compared to non-targeted ^{177}Lu -DOTA-PEG-Cy5-C' dots in vivo or ^{177}Lu in vitro.

Tumor growth was initially suppressed over the first two-three weeks post treatment in both the B16F10 and M21 therapy studies, however over time tumor growth resumed and eventually resulted in the mice reaching endpoint criteria. Unless 100% of the malignant cells are killed, the tumors will return. There are several particle-based approaches that could be employed to improve therapeutic efficacy. Multiple low doses of ^{177}Lu -DOTA-PEG-Cy5-C' dots over time would likely allow delivery of a cumulative total dose with less normal tissue toxicities. Combination therapy with chemotherapeutic agents conjugated to the C' dot particles via hydrolyzable linkers have been synthesized and could be used in conjunction with radiolabeled C' dots. The particle itself has been shown to induce ferroptosis in tumor cells, but not in normal cells, and could be used in combination with a chemotherapy agent conjugated C' dots or radiolabeled C' dots [44]. Finally, the C' dots

have been shown to activate immune cell components in the tumor microenvironment, which could leverage more effective combination therapies, such as immunotherapy – C' dot radiotherapy [45].

3.6. Therapy study tissue histology

Harvested tumor, liver, and renal tissue specimens were obtained from ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dot-treated and PBS non-treated control B16F10 tumor bearing mice, H&E stained, and interpreted by a veterinary pathologist (Fig. 7). An analysis of treated and non-treated liver specimens revealed normal hepatic architecture and hepatocellular morphology without evident histopathology. Likewise, an examination of renal tissues revealed normal architecture of the cortex and outer medulla without pathologic findings. No acute renal damage was observed in treated kidneys. A comparison of tumor tissue specimens for treated versus non-treated animals revealed obvious differences. The treated tumors exhibited large areas of tumor necrosis and hemorrhage with intermingled cords of viable neoplastic cells, while non-treated tumors displayed markedly reduced areas of tumor necrosis. These results are consistent with targeted radiotherapy damage to tumor tissues, with relative sparing of major excretory organs (liver and kidney).

A limited histological survey of specific immune cell populations in harvested tumor tissues was also performed at the time of sacrifice using immunohistochemistry. B16F10 tumor tissue samples, obtained from animals treated with ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots, DOTA- α MSH-PEG-Cy5-C' dots, and PBS control, were stained for T-cell markers (CD3, CD8) and the murine F4/80 macrophage marker (Fig. S7). There were no statistically significant differences in the relative percentages of macrophages across specimens, noting that melanin was present in macrophages active in tumor cell turnover. Interestingly, increased percentages of CD3+ T cells were found in tumor tissues treated with ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots or DOTA- α MSH-PEG-Cy5-C' dots relative to those treated with PBS control, while only specimens treated with ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots showed an increased percentage of CD8+ T cells. These findings suggest that both radiolabeled and non-radiolabeled MC1-R targeted C' dots may prime cytotoxic T cells in the tumor microenvironment, offering the potential to synergize with immunotherapy to improve treatment response and outcomes.

4. Conclusion

Fluorescent core-shell silica nanoparticles synthesized in water, *i.e.* C' dots, containing DOTA conjugated MC1-R avid peptides, were radiolabeled with the beta-particle-emitting radionuclide ^{177}Lu and evaluated for their therapeutic properties in melanoma models. DOTA- α MSH-PEG-Cy5-C' dots demonstrated subnanomolar affinity for MC1-R expressing melanoma cells due to the multi-dentate display of α MSH peptides, leading to efficient internalization. The DOTA- α MSH-PEG-Cy5-C' dot particles demonstrated high radiolabeling efficiency and radiochemical stability with ^{177}Lu . *In vivo*, ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots exhibited favorable PK and targeted tumor uptake and retention. Tumor localization of ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots could be visualized by SPECT imaging and confirmed histologically. Melanoma-bearing mice treated with ^{177}Lu -

DOTA- α MSH-PEG-Cy5-C' dots exhibited prolonged survival over control groups, highlighting the potential of C' dot targeted radiotherapy. The ultrasmall melanoma-avid theranostic silica nanoparticles studied here combine many of the targeting properties of peptides and antibodies with more favorable pharmacokinetic, clearance, and safety profiles. These properties, in turn, can enhance targeted tumor penetration, accumulation, and retention while abrogating dose-limiting toxicity.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Highlights

- Unique particle surface chemistry yields favorable targeting and pharmacokinetics
- Nanoparticle therapy results in reduced tumor growth and prolonged survival
- Overcomes dose-limiting toxicities associated with existing therapeutics
- Yields improved treatment efficacy and reduced off-target side effects

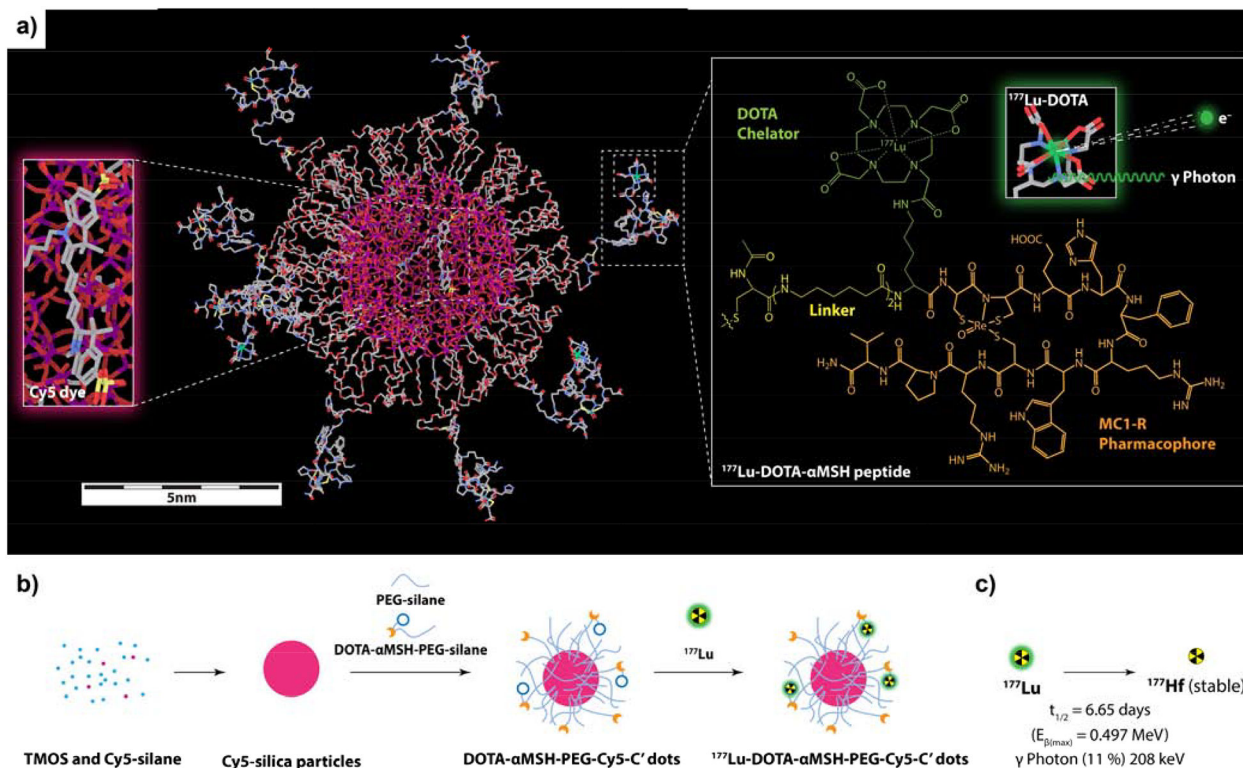


Fig. 1. DOTA-αMSH-PEG-Cy5-C' dots. (a) Molecular rendering of a ^{177}Lu -labeled DOTA-αMSH-PEG-Cy5-C' dot with Cy5 dye-encapsulating silica core (left insert) and the molecular structure of ^{177}Lu -labeled MC1-R targeting cyclic DOTA-αMSH peptides on the right. (b) Schematic of the synthetic approach to DOTA-αMSH-PEG-Cy5-C' dots and its radiolabeling with ^{177}Lu . (c) Radioactive half-life and decay scheme of predominant beta-emitting particles and imageable gamma photons of ^{177}Lu .

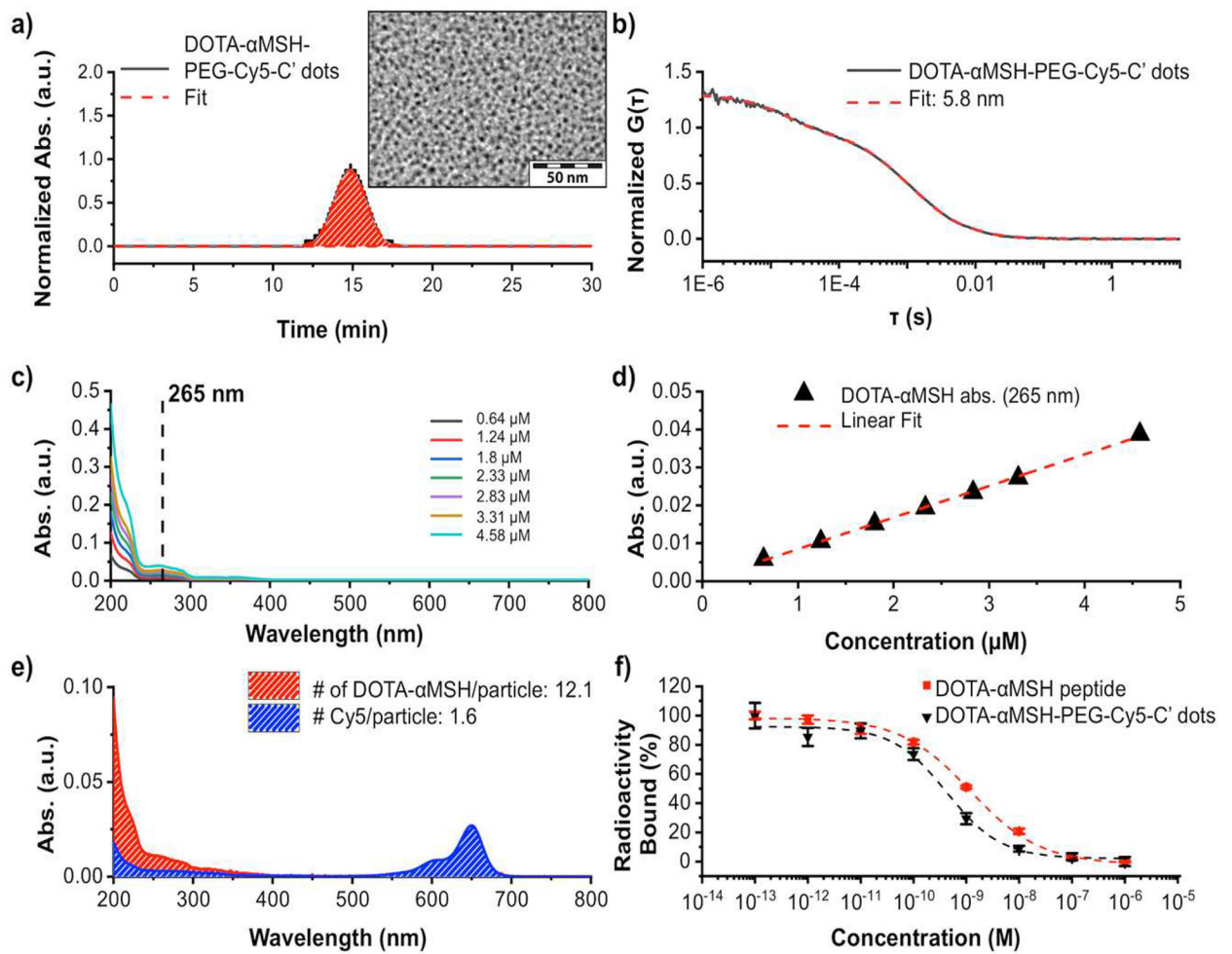


Fig. 2. DOTA- α MSH-PEG-Cy5-C' dot particle characterization. (a) GPC elugram of purified DOTA- α MSH-PEG-Cy5-C' dots, together with fit using a Gaussian distribution function.³⁰ (2a-Insert) is a TEM image of particles associated with the GPC peak. (b) DOTA- α MSH-PEG-Cy5-C' dot FCS correlation curve with fit [29–30]. (c,d) DOTA- α MSH peptide absorbance calibration at 265 nm using UV-VIS spectra at various concentrations. (e) Deconvolution of the UV-vis spectrum of DOTA- α MSH-Cy5-C' dot into DOTA- α MSH peptide and PEG-Cy5-C' dot contributions. (f) Competitive binding studies with DOTA- α MSH-PEG-C' dots and DOTA- α MSH peptide with 125 I-NDP to B16F10 melanoma cells to determine their IC₅₀ values.

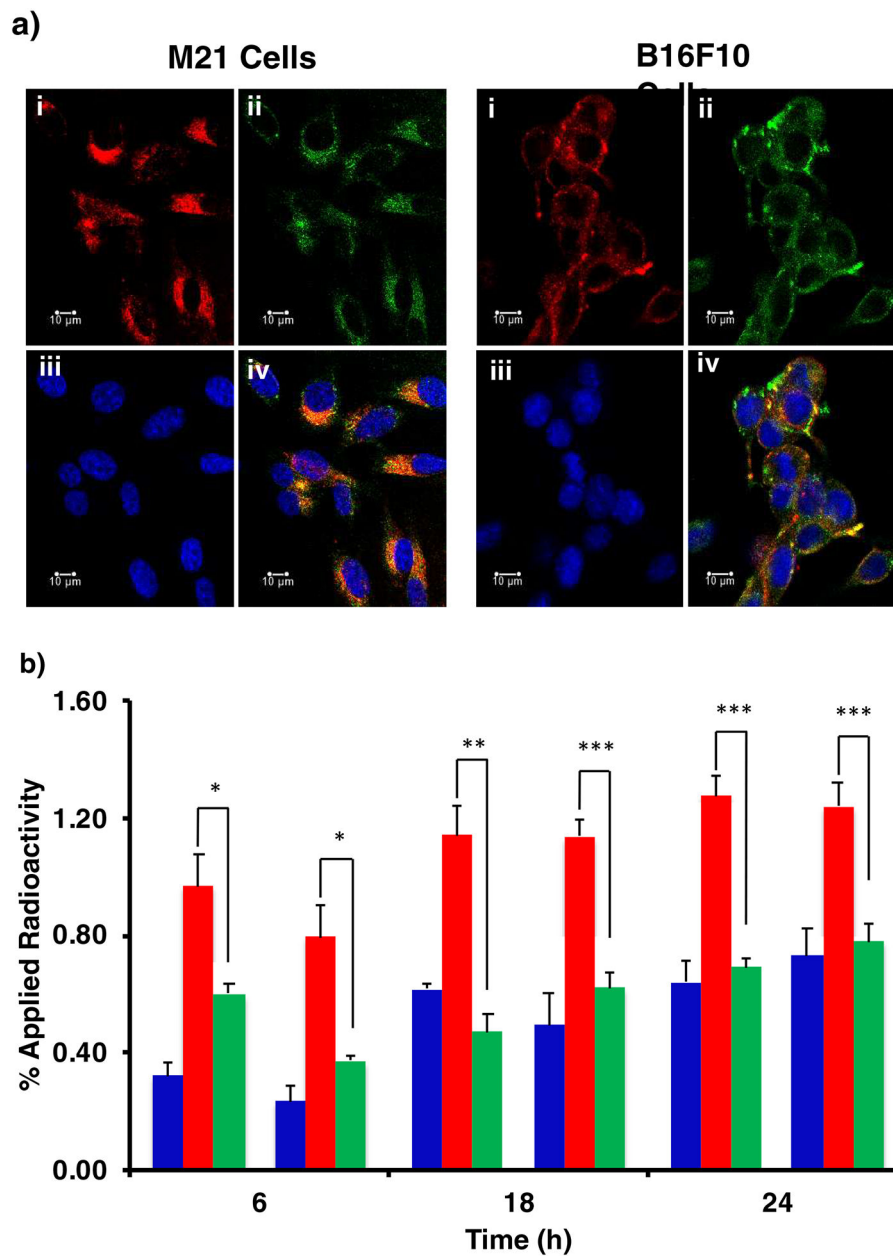


Fig. 3. (a) Confocal fluorescence microscopy visualization of DOTA- α MSH-PEG-Cy5-C' dots internalization in M21 human melanoma and B16F10 murine melanoma cells at 4 h post administration. M21 and B16F10 melanoma cells were imaged in the presence of DOTA- α MSH-PEG-Cy5-C' dots (i), LysoTracker Green (ii) and Hoechst nuclei stain (iii). Merged images displaying all three probes (iv). Scale bar equals 10 μ m. (b) Cell binding and internalization of ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots in B16F10 and M21 melanoma cells. Membrane bound (■) and internalized (■) radioactivity were determined at 6 h, 18 h, and 24 h post incubation with ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots. Co-incubation of ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots with the MC1-R avid NDP blocking peptide (■) was used to assess specificity of internalization. The significance of internalized versus

blocked activity is given as * $p < 0.05$. ** $p < 0.01$ and *** $p < 0.005$. Each data point represents the mean \pm s.d. of 3 replicates.

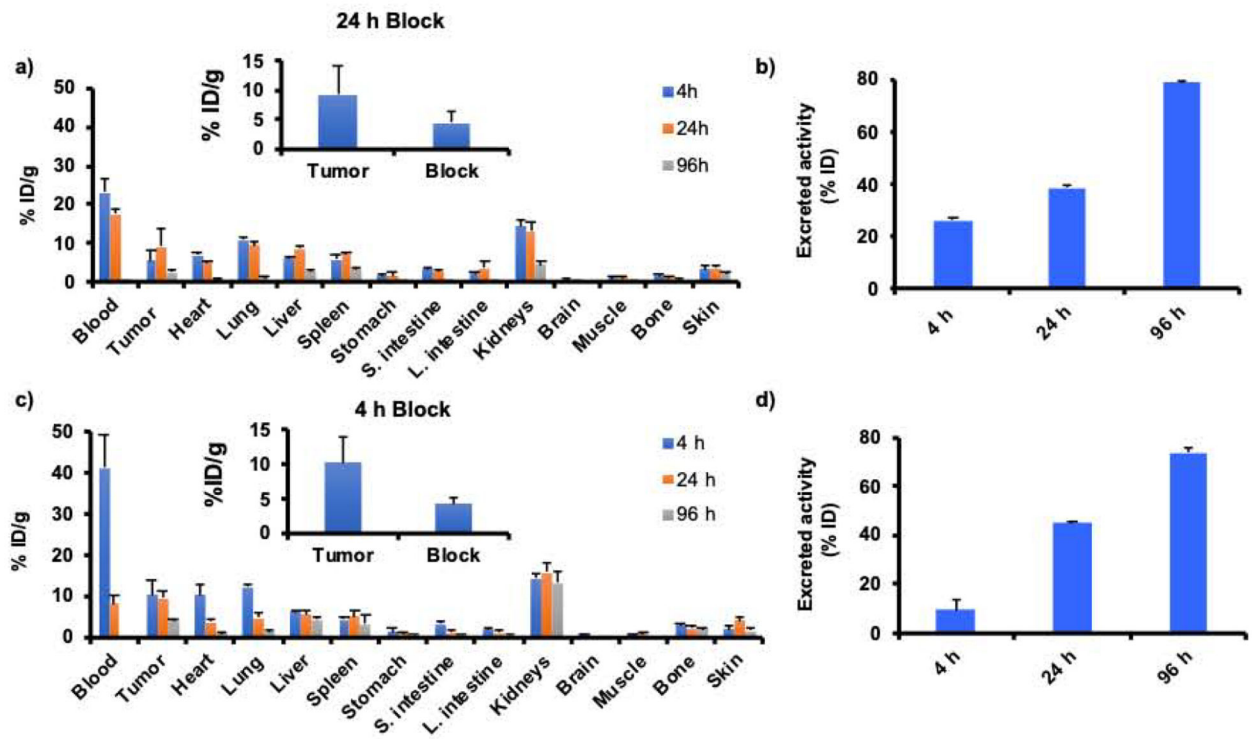


Fig. 4. Biodistribution of ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots in (a-b) M21 and (c-d) B16F10 tumor bearing mice ($n=4$) at 4 h, 24 h and 96 h post injection reported as injected dose per gram of tissue (% ID/g). Clearance of radioactivity in the urine was reported as percent injected dose (% ID). A blocking study was performed by co-injecting ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots + $5\mu\text{g}$ of NDP (inserts). The blocking study was performed at the time point of maximum ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dot uptake. Excreted activity is defined as that measured in urine at 4 h and in urine plus feces at 24 h and 96 h.

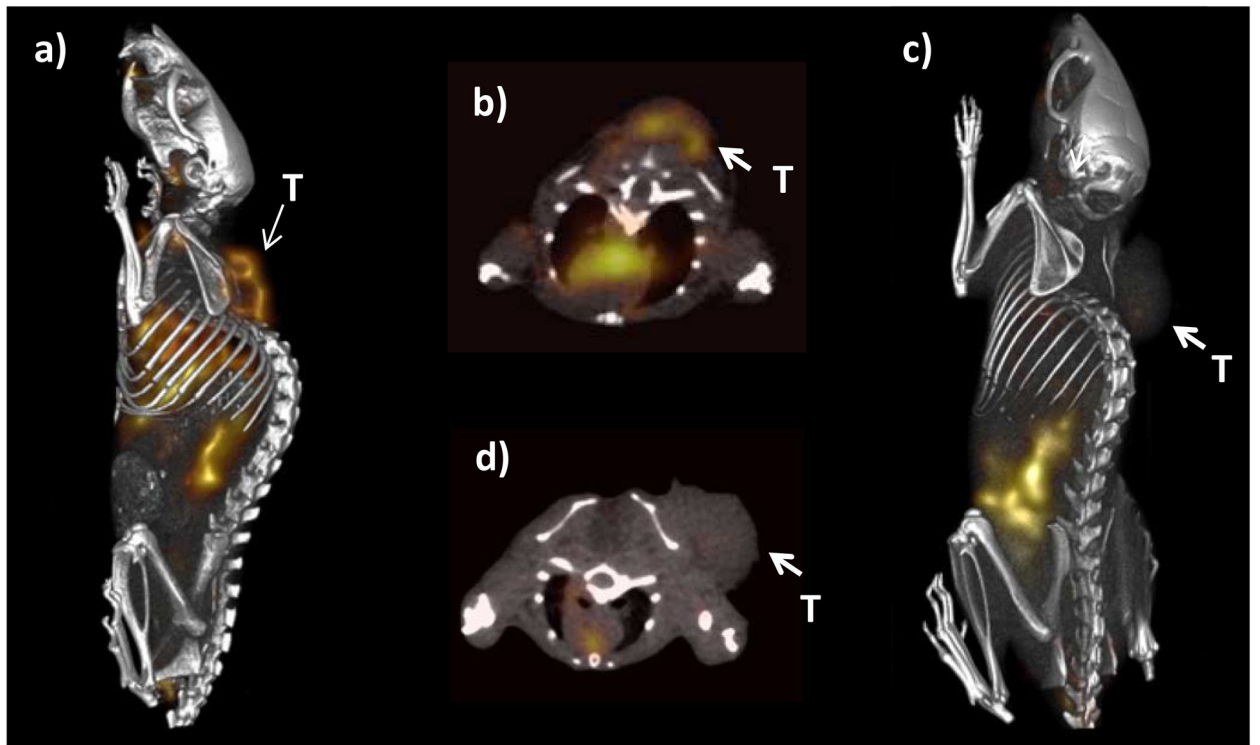


Fig. 5. ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dot SPECT imaging of M21 melanoma tumor bearing mice at 24 h post injection. (a) Whole body and (b) transaxial images with ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots. (c) Whole body and (d) transaxial images showing tumor uptake blocked with ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots + NDP peptide (T = Tumor).

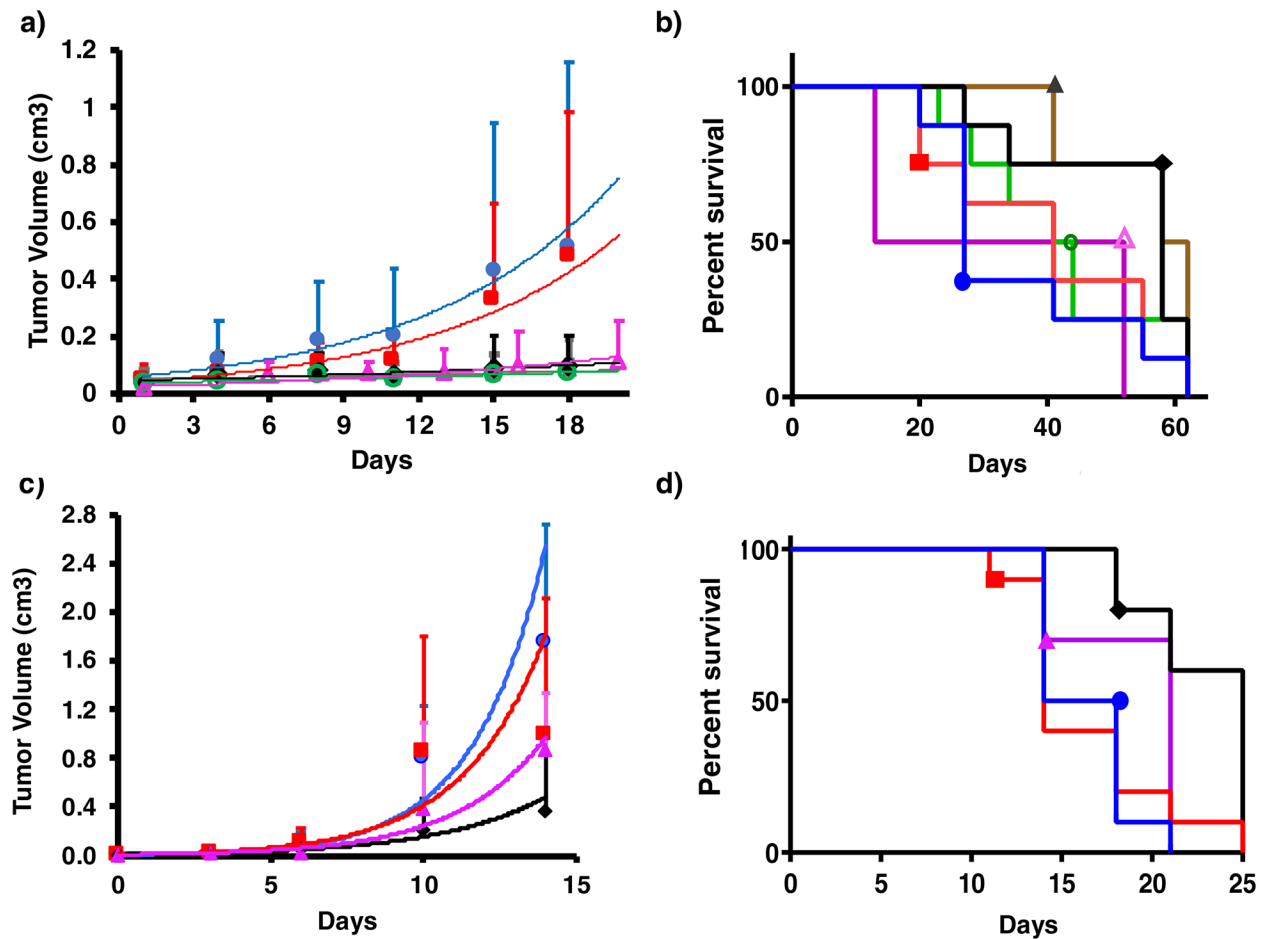


Fig. 6. Tumor volume and survival plots for M21 and B16F10 melanoma-bearing mice treated with ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots. Tumor growth rate plots for (a) M21 xenografted mice and (b) Kaplan-Meier survival plots. M21 tumor mice were administered 1 mCi ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots (○), 0.5 mCi ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots (◆), 1 mCi ^{177}Lu -DOTA-PEG-Cy5-C' dots (▲), 0.5 mCi ^{177}Lu -DOTA-PEG-Cy5-C' dots (◻), DOTA- α MSH-PEG-Cy5-C' dots (■), or PBS (●). (c) B16F10 mice tumor growth rates and (d) Kaplan-Meier survival plots. B16F10 tumor mice were administered 0.5 mCi ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots (◆), 0.5 mCi ^{177}Lu -DOTA-PEG-Cy5-C' dots (▲), DOTA- α MSH-PEG-Cy5-C' dots (■), or PBS (●). M21 tumor bearing mice receiving 0.5 mCi of ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots exhibited significantly improved survival compared to the PBS group $^*(p=0.036)$ and mice receiving 0.5 mCi ^{177}Lu -DOTA-PEG-Cy5-C' dots $^*(p=0.021)$. Mice receiving 1 mCi of the non-targeted ^{177}Lu -DOTA-PEG-Cy5-C' dots also demonstrated a survival advantage $^*(p=0.01)$, while the 1 mCi ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots treated group did not. B16F10 tumor bearing mice receiving 0.5 mCi of ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots demonstrated a significant survival improvement compared to mice receiving PBS $^{***}(p=0.0005)$ or non-radiolabeled DOTA- α MSH-PEG-Cy5-C' dots $^{**}(p=0.002)$. There was a significant enhancement in survival for

tumor bearing mice treated with ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots versus non-targeted ^{177}Lu -DOTA-PEG-Cy5-C' dots *($p=0.045$).

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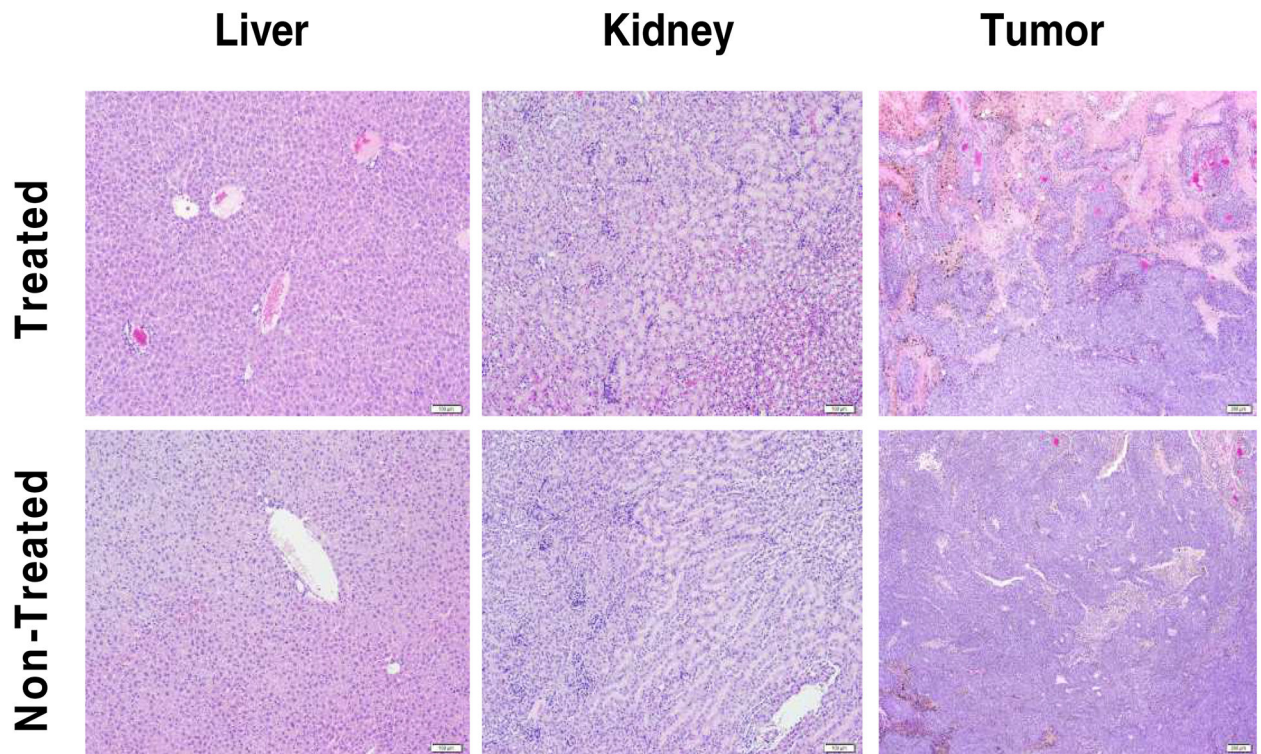



Fig. 7. Therapy study tissue histopathology. H&E stained liver, kidney and tumor tissue from mice treated with 0.5 mCi ^{177}Lu -DOTA- α MSH-PEG-Cy5-C' dots (upper panels) or non-treated PBS controls (lower panels). Scale bar 100 μm .

Table 1.Radiation dosimetry for ^{177}Lu -DOTA- α -MSH-PEG-Cy5-C' dots in a 70-kg standard human.

Tissue	rad/mci
Adrenals	0.029
Blood	
Brain	0.024
Breasts	0.007
Gallbladder Wall	0.032
Lower Large Intestine Wall	0.191
Small Intestine	0.217
Stomach Wall	0.068
Upper Large Intestine Wall	0.128
Heart Wall	0.211
Kidneys	3.78
Liver	0.877
Lungs	0.493
Muscle	0.201
Ovaries	0.015
Pancreas	0.246
Red Marrow	0.122
Bone	0.787
Spleen	0.751
Testes	0.007
Thymus	0.012
Thyroid	0.009
Uterus	0.013
Total Body	0.150