



# Design of a dual Ir-Eu tag for fluorescent visualization and ICP-MS quantification of SIRP $\alpha$ and its host cells

Chunlan Liu<sup>1,2</sup> · Pengtao Li<sup>3</sup> · Xiaowen Yan<sup>1</sup> · Limin Yang<sup>1</sup> · Pingguo Liu<sup>3</sup> · Qiuquan Wang<sup>1</sup>

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## Abstract

With the expansion of ICP-MS application into the field of bioanalysis, there is an urgent need for novel element tags today. Here, we report the design of a dual-element Ir-Eu tag, opening the door to simultaneous fluorescent imaging and ICP-MS quantification. The ratio of  $^{153}\text{Eu}/^{193}\text{Ir}$  may serve as a precision control of the labeling process, allowing internal validation of the quantitative results obtained. As for SIRP $\alpha$  and its host cell analysis exemplified here, the Ir-Eu tag demonstrated superior figures of ICP-MS quantification with the LOD ( $3\sigma$ ) down to 0.5 ( $^{153}\text{Eu}$ ) and 1.1 ( $^{193}\text{Ir}$ ) pM SIRP $\alpha$  and 220 ( $^{153}\text{Eu}$ ) and 830 ( $^{193}\text{Ir}$ ) RAW264.7 cells more than 130 times more sensitive compared with the LOD ( $3\sigma$ ) of 65.2 pM SIRP $\alpha$  at 612 nm using fluorometry. Not limited to these demonstrations, we believe that the design ideas of the dual Ir-Eu tags should be applicable to various cases of bioanalysis when dual optical profiling and ICP-MS quantification are indispensable.

**Keywords** ICP-MS · Fluorescent imaging · Dual-element tag · SIRP $\alpha$  · Cell

## Introduction

So far fluorometry and mass spectrometry have been recognized as two of the main tools for bioanalysis. Moreover, the synergistic use of them should be a more powerful strategy

for easily optic-profiling and accurately quantifying biomolecules and/or their host cells in complex biological samples, providing more comprehensive information of targeted bioanalytes. Although some targeted analytes could be directly detected, a consensus has been reached within chemical measurement science community that many of them should be labeled first with suitable optical and/or mass tags for improving not only sensitivity but also selectivity [1]. This is almost a prerequisite especially when inductively coupled plasma mass spectrometry (ICP-MS) is applied for quantification, in which the  $m/z$  of labeled element/isotope ions is determined to reflect the content of the targeted bioanalytes [1–4]. In such a circumstance, both stable and sensitive optical and  $m/z$  signals simultaneously from the same tag itself are much more desired [5, 6] rather than, respectively, from different tags [7–9], launching the development of an expanded element-tag toolbox that can be more efficiently used for not only optical spectrometric but also ICP-MS bioanalysis. Fortunately, lanthanides (Ln) have very charming fluorescent properties of stable and sharp signals with millisecond lifetime ranging from the visible (for example, Eu and Tb) to near-infrared (Yb, Er, Nd) region because the inner-shell 4f-4f transitions of the Ln are formally parity forbidden [10, 11]. More importantly, on the other hand, all the Ln are very sensitive on ICP-MS [12, 13] due to their relatively low first ionization potentials of around 5.5 eV

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Chunlan Liu and Pengtao Li have equal contribution.

✉ Pingguo Liu  
pqliu@xmu.edu.cn

✉ Qiuquan Wang  
qqwang@xmu.edu.cn

<sup>1</sup> Department of Chemistry & the MOE Key Laboratory of Spectrochemical Analysis and Instrumentation, College of Chemistry and Chemical Engineering, Xiamen University, Xiamen 361005, China

<sup>2</sup> College of Chemistry and Bioengineering, Yichun University, Yichun 336000, China

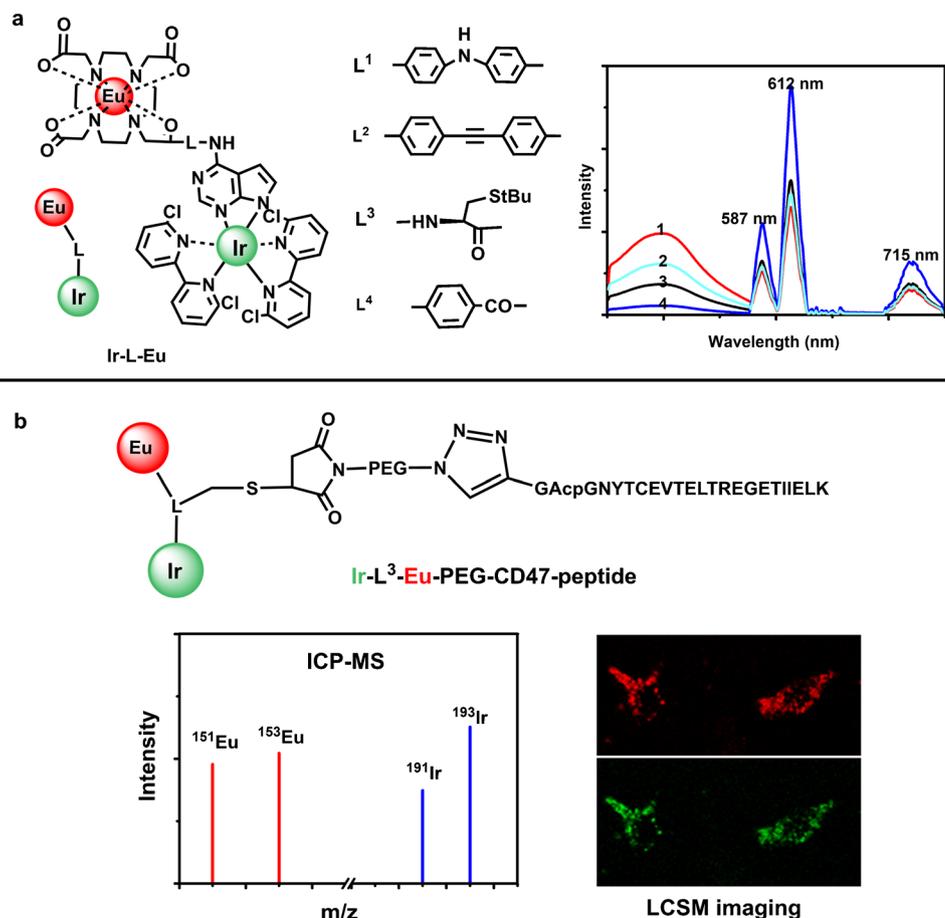
<sup>3</sup> Department of Hepatobiliary Surgery & Fujian Provincial Key Laboratory of Chronic Liver Disease and Hepatocellular Carcinoma, Xiamen Key Laboratory of Translational Medical of Digestive System Tumor, Zhongshan Hospital of Xiamen University, School of Medicine, Xiamen University, Xiamen 361000, China

[14], thus a high ionization efficiency in Ar (15.8 eV)-based ICP. Therefore, dual fluorescent and ICP-MS signals can be simultaneously obtained from the same Ln element through sophisticated molecular design. However, it should be noted that Ln do not have high enough molar absorption coefficient mainly because of their small light-absorbed diameters. An energy antenna moiety as the donor sensitizer for Ln-cored complex chromophores must be employed, which can absorb energy and then transfer to the Ln-cored complex via a rapid internal conversion process for pumping the Ln to excited states [6, 11]. Rational selection of the light-antenna sensitizers is very important to match the energy-requested by the Ln used that emit light signals at their characteristic wavelength. Previous studies showed that a luminescent organometallic complex compared with the conventional organic dyes has many desirable properties for the use as a light-antenna that transfers energy to the Ln [15]. The advantages include the intense light absorption and long-lived excited states with a triplet metal-to-ligand charge-transfer character, thus maximizing the likelihood of energy transfer to the Ln-complex, and photochemical stability and kinetic inertness as well as luminescence of their own, not only providing a basis to monitor energy transfer to the Ln acceptor [16], but also, more meaningfully, offering another ICP-MS signal

source along with the Ln when they conjugate with each other to form a dual-element tag, thus resulting in a precision control for a more accurate analysis. Among numerous fluorescent organometallic complexes, the Ir-cored complex formed between Ir(III) and phenylpyridine-type ligands should be a judicious choice as a light-antenna for the Ln-cored complexes due to the fact that its characteristic excited-state energies are well more than  $19\,000\text{ cm}^{-1}$  over that of, for example, Eu ( $^5D_0$ ,  $17,200\text{ cm}^{-1}$ ) [17, 18].

Here, we design and synthesize a series of dual Ir-Eu tags with Ir and/or Eu fluorescent and ICP-MS signals by linking Eu-DOTA (DOTA stands for 1,4,7,10-tetraazacyclododecane-1,4,7,10-tetraacetic acid) and DCBP-Ir-APP (DCBP stands for 6, 6'-dichloro-3, 3'-bipyridine and APP 4-amino-7H-pyrrole pyrimidine) with different energy-transferring linkers ( $L=L^1, L^2, L^3$ , and  $L^4$ ; Fig. 1a). To demonstrate their feasibility and benefits, Ir- $L^3$ -Eu was conjugated with CD47 peptide GNYTCEVTELTREGETIIEK that can target signal regulatory protein alpha (SIRP $\alpha$ ) overexpressed on immunocytes [19] for the visualization and quantification of SIRP $\alpha$  and its host cells (Fig. 1b). The SIRP $\alpha$  amount overexpressed on the immunocytes in blood samples quantified by ICP-MS provides an indication of the degree of immune response from different diseased people.

**Fig. 1** Structures of the dual Ir-Eu tags linked by different linkers,  $L^1, L^2, L^3$ , and  $L^4$ , and their emission spectra under excitation at 400 nm (a). Ir- $L^3$ -Eu-PEG-CD47-peptide for visualization of the SIRP $\alpha$  overexpressed on RAW264.7 macrophages at 475 nm (Ir) and 612 nm (Eu) under the excitation of 400 nm on LCSM and quantification by  $^{191}\text{Ir}$  and/or  $^{193}\text{Ir}$  and  $^{151}\text{Eu}$  and/or  $^{153}\text{Eu}$  on ICP-MS (b)



## Materials and methods

### Design and synthesis of the dual Ir-Eu tags and the CD47 peptide-conjugated Ir-L<sup>3</sup>-Eu for targeting SIRP $\alpha$ and its host cells

In order to obtain an Ir-Eu tag with effective fluorescent and ICP-MS signals, we started from the synthesis of DCBP-Ir-APP complex and then linked to Eu-DOTA with the linker molecules of bis(4-bromophenyl)amine (L<sup>1</sup>), bis(4-bromophenyl) acetylene (L<sup>2</sup>), N-(9-fluorenylmethoxycarbonyl)-*s*-tert-butylcystine (N-Fmoc-S-tBu-L-cystine, L<sup>3</sup>), and 4-bromobenzoic acid (L<sup>4</sup>) (Scheme S1-S2), respectively.

We first mixed IrCl<sub>3</sub> and DCBP in a solution of 2-ethoxyethanol:H<sub>2</sub>O (3:1) and stirred the mixture for 8 h in room temperature and then heated to 125 °C and reflux for 48 h to obtain DCBP-Ir. Afterwards, APP was mixed with DCBP-Ir to obtain amino-modified Ir complex, DCBP-Ir-APP. L<sup>1</sup> was used to link DCBP-Ir-APP and DO3AtBu through the reaction between amino group and bromine to obtain Ir-L<sup>1</sup>-Eu tag. Ir-L<sup>2</sup>-Eu and Ir-L<sup>4</sup>-Eu tags were synthesized via similar routes (Scheme S1 and see the detailed synthetic procedures, purification, and characterization in the Supplementary Material, Fig. S1-6).

To synthesize Ir-L<sup>3</sup>-Eu tag, we used L<sup>3</sup> first to react with APP via the reaction between carboxyl and amino group to obtain a pyridine ligand containing the post reaction groups and then mixed with DCBP-Ir to obtain the DCBP-Ir-APP tag containing the amino group with Fmoc protection and the *t*Bu-protected sulfhydryl group. After deprotecting the Fmoc from the amino group, and reacting with NHS-DOTA-Eu to obtain Ir-L<sup>3</sup>-Eu-S-tBu. Subsequently, Mal-PEG (2000)-N<sub>3</sub> was used to react with the deprotected Ir-L<sup>3</sup>-Eu-SH to obtain Ir-L<sup>3</sup>-Eu-PEG-N<sub>3</sub>. It was used to label alkynyl-CD47-peptide (GAcP<sub>6</sub>NYTCEVTELTREGETIIEELK) that targeted SIRP $\alpha$  via click reaction (Scheme S2 and see the detailed synthetic procedures, purification, and characterization in the Supplementary Material, Fig. S7-10).

### LCSM imaging and ICP-MS quantification of SIRP $\alpha$ and its host cells

Macrophage cell line RAW264.7 (SIRP $\alpha$ -positive) and cervical carcinoma cell line HeLa (SIRP $\alpha$ -negative), which were cultured in DMEM medium containing 10% FBS, 1% penicillin, and 1% streptomycin at 37 °C in a 5% CO<sub>2</sub> and 95% air incubator, were used to investigate the specificity of the Ir-Eu tags and alkynylated CD47-peptide towards the cells. Each kind of cells was seeded in a 12-well glass plate (1.0 × 10<sup>4</sup> cells per well) and incubated overnight. After washing with PBS buffer, added 20  $\mu$ L 1  $\mu$ M alkynylated CD47-peptide to each well and incubated for 3 h to label SIRP $\alpha$  on the surface of cells. After washing, 50  $\mu$ L Ir-L<sup>3</sup>-Eu-PEG-N<sub>3</sub>, Ir-PEG-N<sub>3</sub>, and Ir-L<sup>3</sup>-Eu (1  $\mu$ M each) as well as PBS buffer were added, respectively, and then 25

$\mu$ L 10  $\mu$ M CuSO<sub>4</sub>, 27.5  $\mu$ L 10  $\mu$ M BIM, and 25  $\mu$ L 100  $\mu$ M sodium ascorbate were added and incubated for another 0.5 h for the click reaction between alkynylated CD47-peptide labeled and the azide group if available in the tags added. After washed five times using PBS buffer, the cells were first imaged on LCSM and then quantified using ICP-MS after dissolving the cells using an appropriate amount of ultrapure concentrated HNO<sub>3</sub> overnight and an appropriate dilution with ultrapure water. To further confirm the specificity towards SIRP $\alpha$ -positive cells, RAW264.7 cells were pre-incubated with 20  $\mu$ L 1  $\mu$ M CD47 protein but not the alkynylated CD47-peptide for 3 h prior to the addition of 50  $\mu$ L 1  $\mu$ M Ir-L<sup>3</sup>-Eu-PEG-N<sub>3</sub> for a blocking experiment, while SIRP $\alpha$ -negative HeLa cells were used as a control. Moreover, the SIRP $\alpha$  per cell in a series of known number RAW264.7 cells were quantified using ICP-MS. Based on this information obtained, in turn, number of the SIRP $\alpha$ -positive RAW264.7 cells can be counted.

We collected blood samples from the Affiliated Zhongshan Hospital of Xiamen University, including 27 blood samples from liver cancer patients, 26 cholecystitis, gastritis and pneumonia patients, and 12 lithiasis (detailed information is listed in Table S1). All the blood samples were stored at -80 °C until use. These obtained samples were approved by the hospital ethics committee, and the patients signed the informed consents (see the Supplementary Material). Prior to quantification of SIRP $\alpha$  and SIRP $\alpha$ -positive cells in blood using ICP-MS, the samples of 1 mL each were appropriately diluted with PBS buffer and incubated with the alkynylated CD47-peptide for 3 h followed by a click labeling with Ir-L<sup>3</sup>-Eu-PEG-N<sub>3</sub> for 5 min and washing and centrifugation 5 times to get rid of the excess unreacted Ir-L<sup>3</sup>-Eu-PEG-N<sub>3</sub> tag and alkynylated CD47-peptide.

Sources of the chemical and biochemical reagents used and instrumentation information were described in the Supplementary Material. Detailed synthesizing routes (Scheme S1-2) and conditions as well as their semi-preparative HPLC purification and fluorescence, NMR, and ESI-MS characterizations (Fig. S1-10) can be found in the Supplementary Material.

## Results and discussion

**Fluorescence and ICP-MS signals from the Ir-Eu tags** After ESI-TOF-MS and NMR characterizations of the designed and synthesized Ir-Eu tags, we evaluated their fluorescent and mass spectroscopic properties. DCBP-Ir-APP has an absorption spectrum from 310 to 420 nm and emits a corresponding fluorescent peak from 450 to 500 nm of 0.1  $\mu$ s lifetime ( $\tau_{Ir_0}$ ) (Fig. S11a), which can match the energy level of Eu-DOTA. When DCBP-Ir-APP was linked to Eu-DOTA

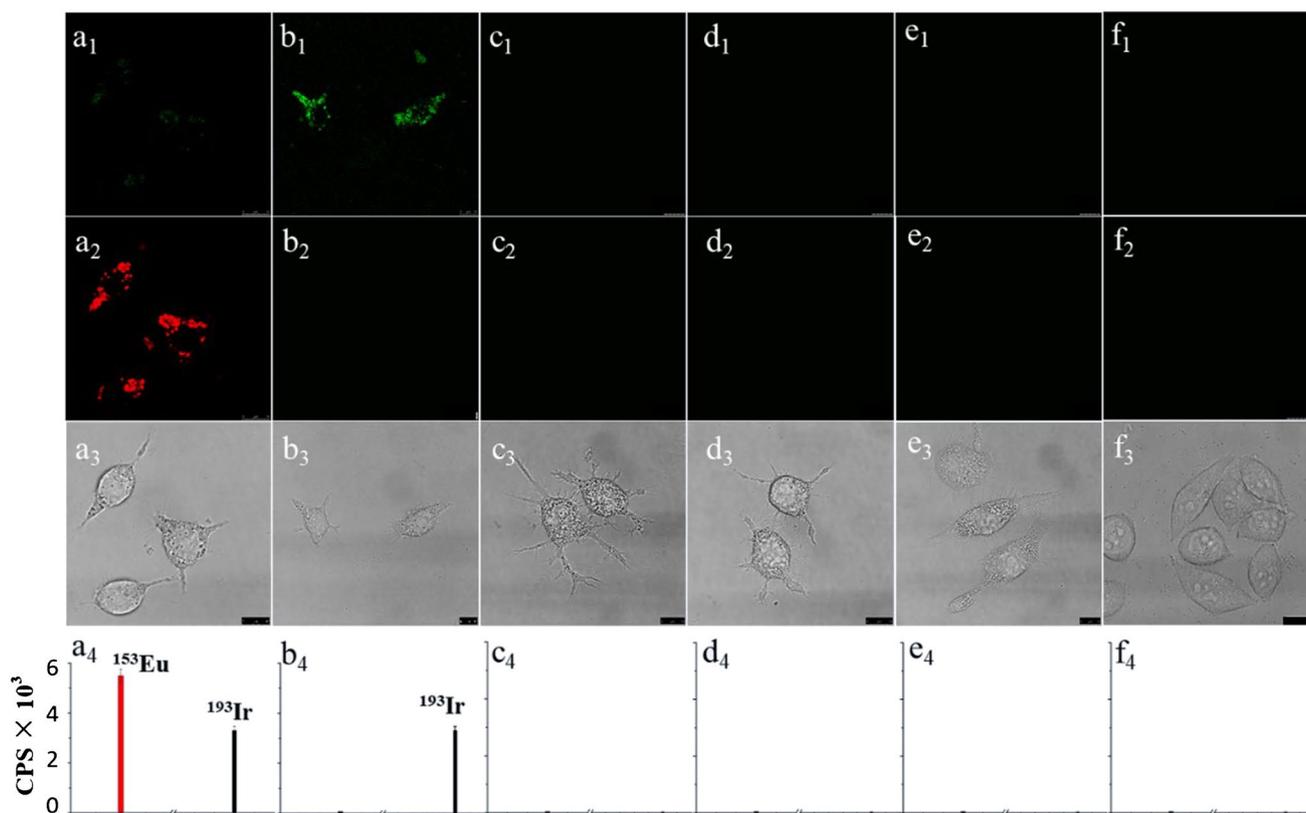
via  $L^1$ ,  $L^2$ ,  $L^3$ , and  $L^4$  of different molecular structure and excited at 400 nm, the emission intensity from DCBP-Ir-APP at 475 nm decreased along with the increase in emission intensity of Eu-DOTA at 591 nm ( ${}^5D_0 \rightarrow {}^7F_1$ ), 612 nm ( ${}^5D_0 \rightarrow {}^7F_2$ ), and 710 nm ( ${}^5D_0 \rightarrow {}^7F_5$ ) (Fig. 1a and Fig. S11b). These phenomena observed confirmed the emission light from DCBP-Ir-APP can be absorbed by Eu-DOTA and profitably transferred the energy to Eu-DOTA. Theoretical calculation ( $\eta = 1 - \tau_{Ir}/\tau_{Ir_0}$ ) according to the fluorescent lifetime ( $\tau_{Ir}$ ) determined at 475 nm [59 ns ( $L^1$ ), 37 ns ( $L^2$ ), 25 ns ( $L^3$ ), and 18 ns ( $L^4$ )] of DCBP-Ir-APP linked to Eu-DOTA by the different linkers suggests that the energy transfer efficiency between DCBP-Ir-APP and Eu-DOTA, which is corresponding to the fluorescent efficiency, was 41% ( $L^1$ ), 63% ( $L^2$ ), 75% ( $L^3$ ), and 82% ( $L^4$ ). These results indicated that both length and structure of the linkers have significant impact on the fluorescent signals from Eu-DOTA in the Ir-Eu tags. In contrast to the fluorescent properties of the Ir-Eu tags, the ICP-MS signal intensities of  ${}^{153}\text{Eu}$  and  ${}^{193}\text{Ir}$  depend simply on their first ionization potential of Eu (5.67 eV) and Ir (8.97 eV) rather than the linkers (Fig. 1b), as we know that ICP is a very hard ionization source breaking the bonds between the atoms and making the atomic ions. When we use the Ir-Eu tags for ICP-MS application, the signals were not affected by the length and structure of the linkers. On the other hand, we must consider the possibility to conjugate a warhead into the tags for labeling the targeted bioanalytes. Regarding SIRP $\alpha$  and its host cells exemplified in this study, we selected Ir- $L^3$ -Eu because it has a better fluorescent efficiency, more importantly, a tBu-protected -SH in Ir- $L^3$ -Eu benefiting the chemical assembly of a post-conjugation moiety (MMA-PEG- $N_3$ ) as we desired for the labeling purpose (Fig. 1b). Subsequently, we further measured a series of Ir- $L^3$ -Eu-PEG- $N_3$  solutions of different concentration from 0 to 500 nM to make a calibration curve at 612 nm (Fig. S12). The linear dynamic range was up to 200 nM, and the limit of detection (LOD,  $3\sigma$ ) was 6.3 pM with an RSD of 3.1% at 50 nM ( $n=3$ ). While the obtained ICP-MS results (Fig. S12) indicated that the LODs ( $3\sigma$ ) when monitoring  ${}^{153}\text{Eu}$  and  ${}^{193}\text{Ir}$  were down to 0.4 and 0.8 pM, the linear dynamic range was up to 500 nM (higher concentrations were not tested) ( $R^2=0.995$  and  $0.997$ ) with an RSD of 2.6% ( ${}^{153}\text{Eu}$ ) and 3.1% ( ${}^{193}\text{Ir}$ ) at 50 nM ( $n=5$ ). All these results suggested that although both fluorescence and ICP-MS signals could be obtained from the same Ir- $L^3$ -Eu-PEG- $N_3$  tag, ICP-MS is superior to fluorescence from the viewpoint of analytical merits.

**Specificity of Ir- $L^3$ -Eu-CD47 peptide towards SIRP $\alpha$  and SIRP $\alpha$ -host cells** In our previous work, we found that SIRP $\alpha$  could be selectively labeled by CD47-peptide with a stoichiometry of 1:1 [20]. Here, we set SIRP $\alpha$  and its host cell as examples and used CD47-peptide to label SIRP $\alpha$  and then

azido Ir- $L^3$ -Eu-PEG- $N_3$  to conjugate the SIRP $\alpha$ -targeted alkynyl CD47-peptide through a click chemistry reaction to demonstrate their LCSM imaging and ICP-MS quantification. Prior to this, the fluorometric and ICP-MS analysis figures of SIRP $\alpha$  were tested. The obtained fluorometric results at 612 nm (Fig. S13) showed that the LOD ( $3\sigma$ ) for SIRP $\alpha$  was 65.2 pM, and the linear dynamic range was from 0.07 to 100 nM ( $R^2=0.920$ ) with an RSD of 3.2% at 50 nM ( $n=5$ ). While monitoring  ${}^{153}\text{Eu}$  and  ${}^{193}\text{Ir}$  on ICP-MS, the LOD was down to 0.5 and 1.1 pM SIRP $\alpha$  with a linear dynamic range up to 200 nM ( $R^2=0.996$  and  $0.997$ ; higher concentrations were not tested) and an RSD of 2.6 and 3.2% at 50 nM ( $n=5$ ) (Fig. S13). The LOD of SIRP $\alpha$  using ICP-MS comparing to that of fluorometry is about 130 times lower, and the sensitivity remains almost unchanged as that of Ir- $L^3$ -Eu-PEG- $N_3$  itself, indicating again the matrix-independent advantage of ICP-MS.

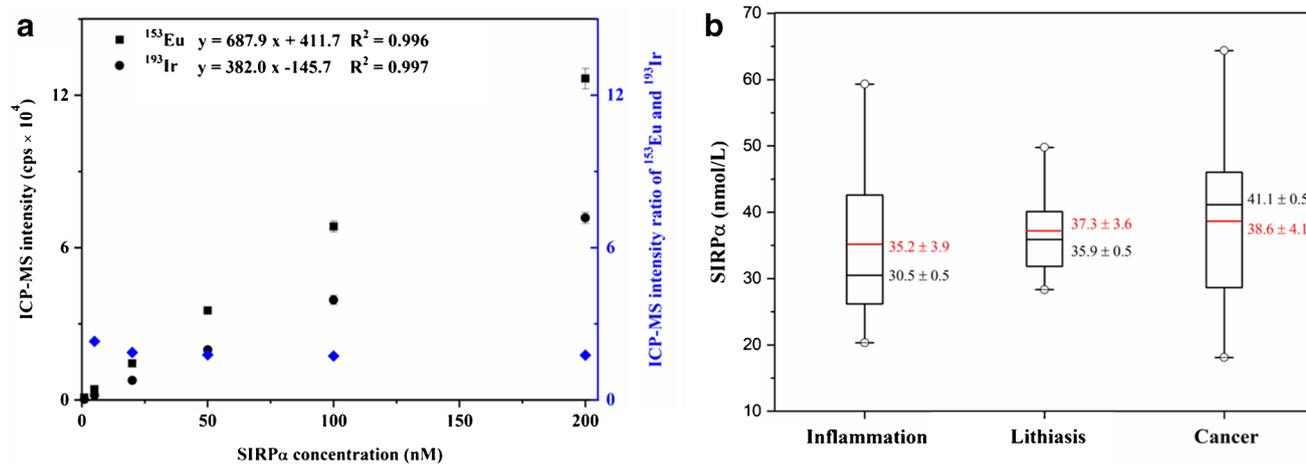
To further verify the targeting specificity of the alkynylated CD47-peptide towards the cells and the click labeling of Ir- $L^3$ -Eu-PEG- $N_3$ , we designed cell experiments using SIRP $\alpha$ -positive RAW264.7 macrophage as a model cell line and SIRP $\alpha$ -negative HeLa cell line as a control. When we added alkynylated CD47-peptide into the cell culture and incubated for 3 h to target the SIRP $\alpha$  overexpressed on the cell surface, and then added Ir- $L^3$ -Eu-PEG- $N_3$ , Ir-PEG- $N_3$ , and Ir- $L^3$ -Eu tags, respectively, to label the SIRP $\alpha$ -targeted alkynylated CD47-peptide. The results obtained indicated that only RAW264.7 could be detected using ICP-MS monitoring  ${}^{153}\text{Eu}$  and/or  ${}^{193}\text{Ir}$  and visualized at 612 nm on LCSM in the cases of Ir- $L^3$ -Eu-PEG- $N_3$  and 475 nm Ir-PEG- $N_3$  as expected, but not the cases of Ir- $L^3$ -Eu and/or HeLa cells (Fig. 2). Moreover, when the alkynylated CD47-peptide was not added or when RAW264.7 cells were pre-incubated with CD47 protein to block the SIRP $\alpha$  on RAW264.7, no fluorescent and ICP-MS signals from RAW264.7 could be detected. All these results obtained suggested that the specificity of alkynylated CD47-peptide towards the SIRP $\alpha$ -positive RAW264.7 cells and that of the click labeling reaction.

Subsequently, different known numbers of RAW264.7 cells were targeted first with alkynylated CD47-peptide and then labeled by Ir- $L^3$ -Eu-PEG- $N_3$  for quantifying the amount of SIRP $\alpha$  overexpressed on per RAW264.7 cell using ICP-MS. The obtained results indicated that the average SIRP $\alpha$  overexpressed on each RAW264.7 cell was  $4.71 \pm 0.04$  amol when 50,000 cells were employed ( $n=5$ ). Based on the LOD of SIRP $\alpha$ , down to 220 ( ${}^{153}\text{Eu}$ ) and 830 ( ${}^{193}\text{Ir}$ ) RAW264.7 cells could be directly counted using ICP-MS. It is worthy of mentioning that the ratio of  ${}^{153}\text{Eu}/{}^{193}\text{Ir}$  remained constant even though the SIRP $\alpha$  concentration and/or the cell number increased (Fig. 3a and Fig. S14), which implying that it can serve as a precision control of the labeling process, allowing internal validation of the quantitative results obtained.



**Fig. 2** LCSM images of SIRPα on the surface of RAW264.7 incubated first with alkynylated CD47-peptide and then labeled with Ir-L<sup>3</sup>-Eu-PEG-N<sub>3</sub> (a<sub>1</sub>-a<sub>4</sub>), Ir-PEG-N<sub>3</sub> (b<sub>1</sub>-b<sub>4</sub>), and Ir-L<sup>3</sup>-Eu (c<sub>1</sub>-c<sub>4</sub>); RAW264.7 without alkynylated CD47-peptide incubation and labeled directly with Ir-L<sup>3</sup>-Eu-PEG-N<sub>3</sub> (d<sub>1</sub>-d<sub>4</sub>); RAW264.7 pre-incubated

with CD47 protein to block SIRPα on the surface of the cells and then incubated with alkynylated CD47-peptide and labeled by Ir-L<sup>3</sup>-Eu-PEG-N<sub>3</sub> (e<sub>1</sub>-e<sub>4</sub>); SIRPα-negative Hela cells were used as a control (f<sub>1</sub>-f<sub>4</sub>)



**Fig. 3** Calibration curves of SIRPα concentration against ICP-MS intensities of <sup>153</sup>Eu and <sup>193</sup>Ir labeled as well as their ratio (a); boxplots of SIRPα concentrations in the blood samples from the patients of liver cancer (*n* = 27), lithiasis (*n* = 12), and inflammation (*n* = 26).

The boxes represent the 25th and 75th percentiles, the whiskers represent the 5th and 95th percentiles, and outliers are displayed by circles. The black lines represent the median value; the red lines represent the mean value (b)

**Quantification of SIRP $\alpha$  and estimation of SIRP $\alpha$ -host immunocytes in real blood samples using ICP-MS** After verifying the specificity of alkynylated CD47-peptide towards SIRP $\alpha$  and its host cells as well as the click-labeling of Ir-L<sup>3</sup>-Eu-PEG-N<sub>3</sub>, SIRP $\alpha$  in real blood samples were quantified using ICP-MS. The obtained results (Fig. 3b and Table S1) showed that the SIRP $\alpha$  quantified in liver cancer blood samples ranged from  $18.1 \pm 0.2$  to  $64.4 \pm 1.6$   $\mu\text{mol/mL}$  with the mean and median values of  $38.6 \pm 4.1$   $\mu\text{mol/mL}$  and  $41.1 \pm 0.5$   $\mu\text{mol/mL}$  ( $n = 27$ ). When further analyzing these data in details, we could see that the mean and median concentrations of the SIRP $\alpha$  in liver cancer blood samples are the biggest, following by  $37.3 \pm 3.6$   $\mu\text{mol/mL}$  (mean) and  $35.9 \pm 0.5$   $\mu\text{mol/mL}$  (median) from  $28.3 \pm 0.1$  to  $49.8 \pm 0.8$   $\mu\text{mol/mL}$  in the lithiasis blood samples ( $n = 12$ ) and  $35.2 \pm 3.9$   $\mu\text{mol/mL}$  (mean) and  $30.5 \pm 0.5$   $\mu\text{mol/mL}$  (median) from  $20.3 \pm 0.2$  to  $59.3 \pm 1.3$   $\mu\text{mol/mL}$  in the inflammation blood samples ( $n = 26$ ). It should be noted that the SIRP $\alpha$ -positive cells in blood sample are not limited to macrophage (for example, RAW264.7 demonstrated here), but also include monocyte, dendritic cell, neutrophil, eosinophil, and basophilic granulocyte, which are all immunocytes. If we might roughly estimate the SIRP $\alpha$ -host immunocyte number that may reflect the degree of immune response of the patients based on the average content of SIRP $\alpha$  on each RAW264.7, the SIRP $\alpha$ -host cell number ranged from  $3.84 \pm 0.04$  to  $13.67 \pm 0.33 \times 10^6$  with  $8.20 \pm 0.86 \times 10^6$  (mean) and  $8.73 \pm 0.16 \times 10^6$  (median) in the liver cancer blood samples,  $6.02 \pm 0.02$  to  $10.57 \pm 0.16 \times 10^6$  with  $7.91 \pm 0.76 \times 10^6$  (mean) and  $7.62 \pm 0.10 \times 10^6$  (median) in the lithiasis blood samples and  $4.32 \pm 0.02$  to  $12.59 \pm 0.28 \times 10^6$  with  $7.46 \pm 0.82 \times 10^6$  (mean) and  $6.47 \pm 0.11 \times 10^6$  (median) in the inflammation blood samples (Table S1). Clearly, although the diagnostic meaning behind these data presented here needs further investigation together with professionally medical doctors in the future when considering individual differences even among patients with the same disease, the design of a dual-element Ir-Eu tag did provide an alternatively selectable way for biomarkers quantification and cells counting using ICP-MS, which allowing internal validation of the quantitative results obtained.

## Conclusions

We designed and synthesized a dual Ir-Eu tag that offers both fluorescence and ICP-MS signals for bioanalysis, enabling the imaging and quantification of a biomarker and its host cells. Although the LOD using the dual Ir-Eu tag is similar to that of our previous work [6], the two different ICP-MS detectable Ir and Eu are available for the internal validation of the quantitative results obtained. Such an idea is not limited to the analysis of SIRP $\alpha$  and

its host cells exemplified here; we envision that more applications to bioanalysis could be expected when more dual-element and even multi-element tags are designed and prepared towards targeting more biomarkers and their host cells in the future. On the other hand, we believe that the individual SIRP $\alpha$ -host immunocytes, rather than a little bit rough estimation of their total numbers in the blood samples demonstrated here, will be recognized and counted on a single-cell multi-parameter ICP-MS platform as the development of the Ln-encoded signal-amplification strategies [21–25], in which not only dual- but also multi-element tags will play a unique role.

**Supplementary Information** The online version contains supplementary material available at <https://doi.org/10.1007/s00216-023-05108-5>.

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## Declarations

**Ethics approval** Sixty-five blood samples from 27 liver cancer patients, 12 lithiasis patients and 26 inflammation patients were provided by Xiamen University affiliated Zhongshan Hospital and permitted by Medical Ethics Council.

**Consent to participate** Informed consent was given by the patients donating the blood sample for analysis.

**Conflict of interest** The authors declare that they have no competing interests. Qiuquan Wang is co-editor of *Analytical and Bioanalytical Chemistry* but was not involved in the peer review of this paper.

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