Noninvasive near-infrared fluorescent protein-based imaging of tumor progression and metastases in deep organs and intraosseous tissues

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Abstract. Whole-body imaging of experimental tumor growth is more feasible within the near-infrared (NIR) optical window because of the highest transparency of mammalian tissues within this wavelength spectrum, mainly due to improved tissue penetration and lower autofluorescence. We took advantage from the recently cloned infrared fluorescent protein (iRFP) together with a human immunodeficiency virus (HIV)-based lentiviral vector to produce virally transduced tumor cells that permanently express this protein. We then noninvasively explored metastatic spread as well as primary tumor growth in deep organs and behind bone barriers. Intraplume tumor growth was investigated through intracranial and intratibial injections of glioblastoma and osteosarcoma cells, respectively, and metastasis was assessed by tail vein injection of melanoma cells. We found that the emitted fluorescence is captured as sharp images regardless of the organ or tissue considered. Furthermore, by overlaying fluorescence spots with the white light, it was possible to afford whole-body images yet never observed before. This approach allowed us to continuously monitor the growth and dissemination of tumor cells with a small number of animals, minimal animal handling, and without the need for any additive. This iRFP-based system provides high-resolution readouts of tumorigenesis that should greatly facilitate preclinical trials with anticancer therapeutic molecules. © The Authors. Published by SPIE under a Creative Commons Attribution 3.0 Unported License. Distribution or reproduction of this work in whole or in part requires full attribution of the original publication, including its DOI. [DOI: 10.1117/1.JBO.19.1.016019]

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1 Introduction

Animal models allow us to recapitulate many of human pathologies including cancer and, although they do not perfectly mimic all aspects of cancer development, they greatly contributed to our understanding of many tumorigenesis mechanisms. Initial studies consisted of implantation of human cancer cells or tumor biopsies into nude mice and evaluation of tumor progression by palpation and/or measurement of tumor sizes by devices such as a caliper. However, these manipulations could not be applied to study deep tumors or metastatic deposits, unless the animal was dissected to inspect for tumor lesions in several tissues and organs, therefore rendering these approaches laborious and animal consuming. Hence, there was a need to find out alternative methodologies that would ease early tumor detection and evaluation of cancer-linked events such as neoangiogenesis, invasion and metastatic dissemination. Accordingly, interest has increased within the past decade in sensitive noninvasive techniques such as fluorescence-based imaging with proteins of the enhanced green fluorescent protein (EGFP) family, the Discosoma sp. DsRed or their shifted wavelength spectra variants as well as luciferase-based bioluminescence. Yet, for bioluminescence to happen, it is obligatory to inject the substrates luciferin or coelentarazin prior to imaging in order to elicit luciferase reaction. Fluorescence imaging of deep tumors using EGFP, DsRed or their mutants is hampered by the limited light penetration in the body, scattering, and tissue absorption. Another problem encountered with EGFP/DsRed-based imaging is autofluorescence which mostly occurs in the green part of the spectrum. Recently, a series of very bright red-shifted fluorescent proteins, such as mKate, E2-Crimson and mNeptune, with an average excitation/emission in the range of 600/650 nm, have been developed with significantly increased tissue penetration and reduced autofluorescence. With regard to these limitations, probes with the wavelengths in the near-infrared (NIR) to infrared (IR) ranges (i.e., 650 to 900 nm), would be superior to others for deep-tissue imaging. Recently, two NIR fluorescent proteins, IFP1.4 (Ref. 10) and iRFP (Ref. 11), have emerged with fluorescence characteristics laying within an NIR transparency of mammalian tissues. The first described in 2009 was the bacteriophytochrome IFP1.4 followed, 2 years later by iRFP, cloned from the Rhodopsedomonas palustris and engineered through a series of mutations. Although fluorescence spectra of the two molecules are quite similar, IFP1.4 must incorporate its chromophore biliverdin (which needs to be exogenously supplied) to further emit fluorescence, whereas the physiological level of biliverdin is sufficient to fully activate iRFP. This property, added to the high brightness, prompted us to use iRFP as a tool to track primary tumors and metastasis in deep tissues and behind natural barriers such as bones.

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We, herein, used in vivo NIR optical imaging together with a human immunodeficiency virus (HIV)-based lentiviral vector to deliver the iRFP gene into different tumor cell models including glioblastoma, osteosarcoma, and melanoma cells. We then used these cells to explore intracranial and intratibial primary tumor growth as well as metastatic dissemination of melanoma cells.

2 Material and Methods

2.1 Vector Construction

The HIV-derived recombinant lentivirus vector expressing iRFP was made by replacing the deoxyribose nucleic acid (DNA) of EGFP in the lentiviral vector pRLSIN.ePPT.PGK-GFP.WPRE (referred to as pRRL, kindly provided by Dr. Trono, University of Geneva, Switzerland) by the iRFP DNA from the pShuttle-CMV-iRFP plasmid (a gift of Dr. Verkruysse, Addgene plasmid # 31856). To this end, iRFP DNA was excised with BgII and SalI, gel purified and ligated into BamHI/SalI-digested pRRL resulting in pRRL/iRFP construct [Fig. 1(a)].

2.2 Cells and Lentiviral Transduction

The human embryonic kidney cells HEK293T, the glioblastoma U87 MG and the melanoma IGR37 cells were cultured in Dulbecco's modified Eagle medium (Cambrex, Verviers, Belgium). The osteosarcoma U2OS cells were cultured in Roswell Park Memorial Institute (RPMI) medium 1640 (Cambrex). All media were supplemented with 10% fetal calf serum and antibiotics penicillin and streptomycin (Eurobio, Les Ulis, France). Lentiviral particles were produced according to Mathieu et al.14

2.3 Animals and Tumor Implantations

Four- to six-week old male athymic Naval Medical Research Institute nude mice (Harlan, Gannat, France) were maintained in a barrier facility on high efficiency particles arresting (HEPA)-filtered racks and fed with autoclaved alfalfa-free rodent diet to avoid digestive tract autofluorescence.15 Animal experiments were conducted in accordance with the European Directive EEC/2010/63, under the supervision of authorized investigators. Subcutaneous (s.c) and intracranial (i.c) injections of U87 MG glioblastoma cells were performed as described earlier16,17 using 1 x 10^5 cells (s.c) or 5 x 10^5 cells (i.c), respectively. For the experimental metastases, we injected 1.5 x 10^6 melanoma cells IGR37 through the tail veins. For intratibial xenografts, mice were anesthetized and 1 x 10^5 U2OS cells were injected into the tibia with a needle introduced at the top of the tibial metaphysis. Immunohistochemistry of melanoma tumors was performed according to Ref. 18, using S100, HBM-45, and Melan-A as markers.

2.4 In Vivo Imaging and Analyses

Unless otherwise indicated, animals were subjected to NIR imaging every 3 to 4 days with a Pearl60 Impulse Small Animal Imaging System Li-Cor60 Biosciences (LI-COR Biosciences GmbH, Bad Homburg, Germany). Excitation and emission wavelengths were fixed at 690 and 710 nm, respectively. The fluorescence signal was overlaid with the white channel light-emitting diode [LED light] and detected by a cooled CCD camera of the imager. During the in vivo imaging process, mice were kept on the imaging bed by isoflurane inhalation (Abbott Laboratories, Rungis, France) and warmed at 37°C. In some experiments, organs or tumor deposits were retrieved with the minimum surrounding tissue from mice sacrificed by cervical dislocation and subjected to ex vivo imaging. Excised tissues were then placed on the stage and their fluorescence captured as indicated above. Image analysis was performed according to Nakayama et al.19 In brief, regions of interest (ROIs) were targeted on iRFP fluorescent sites and circles with the minimum size were drawn around lesions on each image. The pixel density in the selected ROI was quantitated using the software for image analysis provided with the Pearl60 Imager. The fluorescence signal on the plots was normalized to the background and presented as arbitrary units (a.u.) according to Filonov et al.11

2.5 Statistical Analyses

Data were collected from the three independent experiments. Animal implantation experiments were performed with 3 to 4 mice per each group of glioblastoma, osteosarcoma, or melanoma tumor models and results were expressed as a standard error of the mean (SEM).

3 Results

3.1 Visualization of iRFP-Expressing Tumor Xenografts in Mice

Tumor cells expressing iRFP were produced as described and analyzed for the NIR fluorescence before inoculation to animals [Fig. 1(b)]. As shown in Fig. 1(c), fluorescence of iRFP-expressing U87 MG glioma cells,20 not only clearly distinguishes the tumor from the host but also allows a real-time monitoring of subcutaneous tumor growth, without any additive [Fig. 1(d)]. By subcutaneously inoculating increasing numbers of iRFP-labeled cells, we found that the minimal number of detectable cells by our NIR imaging device is 50,000 cells (data not shown). Filonov et al. advocated the use of iRFP for imaging of deep organs such as the liver.11 We, therefore, wanted to ask whether iRFP would uncover intraosseous tumors using intracranial and intratibial inoculated iRFP-expressing tumor cells. As shown in Fig. 2(A), the growth of an intracranial implant of iRFP-expressing U87 MG cells can be easily detected and monitored in real-time in vivo [Fig. 2(Aa)] and ex vivo, after the brain of this mouse has been retrieved [Fig. 2(Aa)]. The deduced growth curve shows a good correlation with the images [Fig. 2(AB)].

Intratibial growth of U2OS cells is shown in Fig. 2(B). Note that the presence of additional NIR fluorescent spots on the abdomen of mice between days 43 and 51, which could be due to traces of alfalfa in this animal food,15 a problem that we subsequently solved. To further analyze the intrabone tumor progression of iRFP-expressing U2OS cells, two animals from a parallel series and corresponding to those of days 43 and 70 were sacrificed and their tibias retrieved. As shown in Fig. 2(Ba), the tumor that initially localized to the head of the bone at day 43, progressively invaded the rest of the organ a week later (at day 51), eventually leading to fracture of the bone and spreading over surrounding soft tissue at day 70 [Fig. 2(Bb)]. Therefore, the iRFP-emitted fluorescence crosses the bone barrier, thus allowing us to reliably estimate tumor burden without sacrificing the animal [see also Fig. 2(Bc)].
Fig. 1 (a) Schematic diagram of pRRL/iRFP construct obtained by swapping deoxyribose nucleic acid (DNA) of EGFP in pRRLSIN.cPPT.PGK-GFP.WPRE vector by iRFP DNA. The expression of the transgene is under the control of the human phosphoglycerate kinase promoter (PGK). The central polypurine tract (cPPT) and the posttranscriptional Woodchuck hepatitis virus (WPRE) are located upstream and downstream of the transgene, respectively. RSV, rous sarcoma virus; SD, splice donor; SA, splice acceptor; Gag, deleted gag region; RRE, rev-responsive element; LTR, long terminal repeat; EGFP, enhanced green fluorescent protein; iRFP, infrared fluorescent protein. (b) Phase contrast and iRFP-based fluorescence of tumor cells transduced with pRRL/iRFP and observed with the Cy5.5 filter at excitation/emission of 660/710 nm (magnification 400×). (c and d) Sequential images and the deduced growth curve after subcutaneous injection of iRFP-expressing U87 MG cells. The color bar indicates the total fluorescence radiant efficiency.
3.2 Visualization of Metastasis

These data being acquired, we then asked whether iRFP-based imaging could be applied to the study of metastatic spread. For this purpose, we used melanoma cells IGR37, originating from a metastatic spread of the primary human melanoma cells IGR39. As shown in Fig. 3(A), injection of iRFP-expressing IGR37 cells in the tail vein resulted in a tumor deposit in the liver [see m1 series and after dissection of the animal Fig. 3(Aa)], consistent with a previous work showing the liver to be a potential niche for melanoma metastasis. Surprisingly, injection of the same cells to two other mice...
led to tumors that appeared, not in the liver, but in the space between omoplates [Fig. 3(B), see m2 series]. Dissection of the animal revealed the presence of well-defined tumors that strongly adhered to the cervical vertebrae (data not shown) but which were positive for melanoma immunological markers such as S100, HBM-45, and Melan-A [Fig. 3(C)]. These data show that whole-body NIR imaging with iRFP uncovers different metastatic targets for melanomas.

4 Discussion

Considerable progress has been made in recent years to cure cancer. Besides surgery, drug-based therapies are more and more efficient and already allow us to save lives or at least to meaningfully extend patient survival.23 Yet, it is still highly desirable to afford new molecules with an improved bioactivity and decreased toxicity. In order to meet this need, preclinical assays involving small animals and genetic tools for whole-body imaging and tumor-emitted signal quantification would significantly improve anticancer therapeutic strategies. Several previous studies have used fluorescent proteins to track tumor lesions in mice,24 including those related to EGFP, DsRed, or their variants.6 However, the wavelengths of these tracers are in a range of low signal-to-noise background and low coefficient of penetration in the animal body,7 particularly within the green part of the spectrum.25 Autofluorescence, scattering, and absorbance are crucial parameters, particularly when imaging organs that lie millimeters deep under the skin layers.7 With the NIR light, however, these parameters become less important.25 Another problem encountered with many previous studies is the use of cells that have been transfected and selected with drugs such as neomycin.4,24 In addition to the possible clone-to-clone variability, the main challenge with such approaches is whether the transgene expression would be kept stable throughout the study. In fact, once the drug-selected cells are deposited in the animal body, they must grow in a drug-free environment and therefore, might lose the gene of interest due to the lack of selection pressure.

To overcome these difficulties, we herein used a lentivirus-mediated delivery system to permanently express the NIR fluorescent protein iRFP in cells. Another attractive characteristic of
this system is that this human immunodeficiency virus type 1 (HIV-1)-based lentiviral vector can infect virtually all kinds of cells including the nondividing ones. Using this genetically encoded probe and an NIR fluorescence imaging device, we are herein reporting the usefulness of this system for the detection of deeply buried tumor lesions or those hidden behind the bone barrier. Most importantly, metastatic deposits can be easily detected and quantified [see Figs. 3(A) and 3(B)], thereby avoiding cumbersome and tedious dissection steps. Although the detection of tumors by iRFP in deep organs has been demonstrated by Filonov et al. and confirmed in the present study, there is no data available using such a tool for visualization of intraosseous-implanted tumors. Intrabone emitted signals are bright enough to be detected and quantified without any contrast agent or chromophore previously administered to the animal [see Figs. 1(c) and 2(A) and 2(B)]. It is noteworthy that attempts to detect glioblastoma cell growth in the brain using IFP1.4 and biliverdin supplementation have been unsuccessful (data not shown). In conclusion, this iRFP- and lentiviral-based system present many other characteristics that combine the permanent and strong expression of iRFP with the low scattering and absorption by biological liquids and bones. Altogether, these features would allow this HIV-based iRFP-expression system to validly support the emerging therapeutic strategies in cancer treatment.

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References

1. B. Hann and A. Balmain, “Building ‘validated’ mouse models of human cancer,” *Curr. Opin. Cell Biol.* 13(6), 778–784 (2001).
2. R. Kurbacher, “Realistic rodents? Debate grows over new mouse models of cancer,” *J. Natl. Cancer Inst.* 98(17), 1176–1178 (2006).
3. C. P. Kerler et al., “Validity of bioluminescence measurements for non-invasive in vivo imaging of tumor load in small animals,” *Biotechniques* 43(5), 7–13, 30 (2007).
4. M. Bouvet et al., “Real-time optical imaging of primary tumor growth and multiple metastatic events in a pancreatic cancer orthotopic model,” *Cancer Res.* 62(5), 1534–1540 (2002).
5. N. C. Sharer et al., “Improved monomeric red, orange and yellow fluorescent proteins derived from Discosoma sp. red fluorescent protein,” *Nat. Biotechnol.* 22(12), 1567–1572 (2004).
6. R. M. Hoffman, “The multiple uses of fluorescent proteins to visualize cancer in vivo,” *Nat. Rev. Cancer* 5(10), 796–806 (2005).
7. J. Lecocq and M. J. Schnitzer, “An infrared fluorescent protein for deeper imaging,” *Nat. Biotechnol.* 29(8), 715–716 (2011).
8. Y. T. Lim et al., “Selection of quantum dot wavelengths for biomedical assays and imaging,” *Mol. Imaging* 2(1), 50–64 (2003).
9. D. Scherbrob et al., “Bright far-red fluorescent protein for whole-body imaging,” *Nat. Methods* 4(9), 741–746 (2007).
10. X. Shu et al., “Mammalian expression of infrared fluorescent proteins engineered from a bacterial phytochrome,” *Science* 324(5928), 804–807 (2009).
11. G. S. Filonov et al., “Bright and stable near-infrared fluorescent protein for in vivo imaging,” *Nat. Biotechnol.* 29(8), 757–761 (2011).
12. K. Konig, “Multiphoton microscopy in life sciences,” *J. Microsc.* 200(2), 83–104 (2000).
13. C. Mancuso and E. Barone, “The heme oxygenase/biliverdin reductase pathway in drug research and development,” *Curr. Drug Metab.* 10(6), 579–594 (2009).
14. S. Mathieu et al., “Transgene expression of alphafucosyltransferease-I (FUT1) in tumor cells selectively inhibits sialyl-Lewis x expression and binding to E-selectin without affecting synthesis of sialyl-Lewis a or binding to P-selectin,” *Am. J. Pathol.* 164(2), 371–383 (2004).
15. Y. Inoue et al., “Diet and abdominal autofluorescence detected by in vivo fluorescence imaging of living mice,” *Mol. Imaging* 7(1), 21–27 (2008).
16. C. Berenguer et al., “Adrenomedullin, an autocrine/paracrine factor induced by androgen withdrawal, stimulates ‘neuroendocrine phenotype’ in LNCaP prostate tumor cells,” *Oncogene* 27(4), 506–518 (2008).
17. A. Tchoghandjian et al., “A2B5 cells from human glioblastoma have cancer stem cell properties,” *Brain Pathol.* 20(1), 211–221 (2010).
18. B. C. Gleason and A. F. Nascimento, “HMBS and Melan-A are useful in the differential diagnosis between granular cell tumor and malignant melanoma,” *Am. J. Dermatopathol.* 29(1), 22–27 (2007).
19. H. Nakayama et al., “Evaluation by bone scintigraphy of osteogenic activity of commercial bioceramics (porous beta-TCP and HAp particles) subcutaneously implanted in rats,” *J. Biomater. Appl.* 24(8), 751–768 (2010).
20. C. Dai and E. C. Holland, “Glioma models,” *Biochim. Biophys. Acta* 1551(1), M19–M27 (2001).
21. R. Taghizadeh et al., “CXCR6, a newly defined biomarker of tissue-specific stem cell asymmetric self-renewal, identifies more aggressive human melanoma cancer stem cells,” *PLoS One* 5(12), e15183 (2010).
22. V. Gray-Schopfer, C. Wellbrock, and R. Marais, “Melanoma biology and new targeted therapy,” *Nature* 445(7130), 851–857 (2007).
23. C. W. Hong and Q. Zeng, “Awaiting a new era of cancer immunotherapy,” *Cancer Res.* 72(15), 3715–3719 (2012).
24. A. Miyawaki, “Fluorescence imaging in the last two decades,” *J. Electron Microsc.* 62(1), 63–68 (2013).
25. R. Weissleder and V. Ntziachristos, “Shedding light onto live molecular targets,” *Nat. Med.* 9(1), 123–128 (2003).
26. W. Qasim, C. A. Vink, and A. J. Thrasher, “Hybrid lentiviral vectors,” *Mol. Ther.* 18(7), 1263–1267 (2010).

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