

# 1 *In vitro* and *in vivo* NIR Fluorescence Lifetime 2 Imaging with a time-gated SPAD camera

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12 **Abstract**—Near-infrared (NIR) fluorescence lifetime imaging (FLI) provides a unique contrast mechanism to monitor  
13 biological parameters and molecular events *in vivo*. Single-photon avalanche photodiode (SPAD) cameras have been  
14 recently demonstrated in FLI microscopy (FLIM) applications, but their suitability for *in vivo* macroscopic FLI (MFLI)  
15 in deep tissues remains to be demonstrated. Herein, we report *in vivo* NIR MFLI measurement with SwissSPAD2, a large  
16 time-gated SPAD camera. We first benchmark its performance in well-controlled *in vitro* experiments, ranging from  
17 monitoring environmental effects on fluorescence lifetime, to quantifying Förster Resonant Energy Transfer (FRET)  
18 between dyes. Next, we use it for *in vivo* studies of target-drug engagement in live and intact tumor xenografts using  
19 FRET. Information obtained with SwissSPAD2 was successfully compared to that obtained with a gated-ICCD camera,  
20 using two different approaches. Our results demonstrate that SPAD cameras offer a powerful technology for *in vivo*  
21 preclinical applications in the NIR window.

22 Index terms—Fluorescence Lifetime Imaging, single photon avalanche diode, preclinical imaging, molecular imaging

## 23 I. Introduction

24 Preclinical molecular imaging is used in early drug development<sup>1,2</sup> and as a research tool to better understand the biology of  
25 drug resistance. Two imaging techniques provide the high sensitivity needed to detect biomarkers during and after drug delivery:  
26 nuclear (PET) and optical imaging. They both allow non-invasive assessment of delivery efficacy, pharmacokinetics, and  
27 response in longitudinal studies<sup>1–3</sup>. PET is good at quantitative molecular imaging of targeted drug delivery in live subjects<sup>4</sup>,  
28 providing spatial and temporal distribution of labeled probes in living animals<sup>4</sup>, but is limited to a single targeted radiotracer. Co-  
29 localization of a radiotracer-labeled antibody-drug conjugates with the pathological site unfortunately does not provide  
30 unequivocal evidence of actual binding to the target protein i.e. receptor engagement, which is essential to elicit the cellular  
31 response necessary to kill cancer cells<sup>5–8</sup>. By contrast, optical imaging methods<sup>9,10</sup> and in particular fluorescence imaging<sup>11,12</sup>,  
32 which offer the possibility to monitor several probes simultaneously, can lift this ambiguity. Fluorescence lifetime imaging (FLI)  
33 can additionally report on numerous intracellular parameters such as metabolic status<sup>13</sup>, reactive oxygen species<sup>14</sup> and  
34 intracellular pH<sup>15</sup> or to quantify Förster Resonant Energy Transfer (FRET), a powerful technique used to study protein-protein  
35 interactions and biosensor activity<sup>16</sup>.

36 We use macroscopic FLI-FRET (MFLI-FRET) to quantify *in vivo* drug-target engagement in live, intact animals over large  
37 fields of view<sup>17–19</sup>: the reduction of donor fluorophore probe lifetime upon drug-target engagement results from the proximity of  
38 an another acceptor fluorophore-labeled probe bound to the same target. Donor lifetime can be measured with high sensitivity  
39 and dispense with corrections required in other FRET techniques, such as sensitized-emission FRET<sup>20</sup>. On the other hand, MFLI-  
40 FRET data acquisition requires more complex instrumentation compared to intensity-based fluorescence imaging. Specifically,  
41 time-gated ICCD cameras, which are the detectors of choice for MFLI applications, are expensive, prone to photocathode  
42 degradation, susceptible to damage from overexposure, and as a dated technology, have limited room for technical  
43 improvements. By contrast, time-resolved CMOS SPAD arrays (SPAD cameras) have undergone tremendous developments over  
44 the past decade, and are poised to become a competitive solution for fluorescence lifetime imaging as discussed here<sup>21</sup>.

45 SwissSPAD2 (SS2) is a very large time-resolved SPAD camera with single-photon sensitivity, developed specifically for  
46 FLI applications<sup>22–24</sup>. This time-gated imaging sensor is comprised of 512×512 SPAD pixels, each associated with a 1-bit

47 memory, readout as a whole at up to 97,700 frames per second<sup>22</sup>. SS2's capabilities for FLIM applications in the visible spectrum  
48 have been recently described<sup>22-24</sup>. Although these studies demonstrated SS2's potential for microscopic biological applications,  
49 there are outstanding challenges involved with using it for pre-clinical macroscopic imaging applications. First, NIR dyes used  
50 for *in vivo* small animal imaging are challenging to detect, due to the reduced photon detection probability of silicon SPADs in  
51 the NIR. Next, most NIR dyes exhibit lifetimes far shorter than visible dyes (few hundreds of picoseconds – ps – compared to  
52 few nanoseconds). Lastly, MFLI-FRET involves quantifying fluorescence decays from two or more fluorophore species or states  
53 simultaneously, which requires larger signal-to-noise compared to mono-exponential cases, leading to additional challenges<sup>25</sup>.

54 Herein, we report the first application of SS2 in a variety of MFLI measurements of NIR fluorescent samples *in vitro* and *in*  
55 *vivo*, systematically comparing it to a state-of-the-art gated-ICCD (Supplementary Fig. S1). First, we study the sub-nanosecond  
56 lifetime NIR dye Alexa Fluor 750 with both detectors, using two distinct methods: nonlinear least square fit (NLSF) and phasor  
57 analysis. We then quantify the lifetime of the clinically relevant NIR dye IRDye 800CW<sup>26</sup> as a function of molecular  
58 microenvironment. MFLI measurements of NIR-FRET pairs in different ratios characterized by multi-exponential decays  
59 conclude these *in vitro* benchmarks. Next, we use SS2 for noninvasive preclinical MFLI-FRET imaging of live mice carrying  
60 HER2 positive tumor models. Two clinical cancer drugs, the anti-HER2 monoclonal antibody (mAb) Trastuzumab (TZM) and  
61 the anti-EGFR mAb Cetuximab (CTM), both labeled with NIR donor and acceptor dyes are used as FRET probes to visualize  
62 HER2-and EGFR-positive tumors in live mice. We successfully characterize organs of interest across the mouse body based on  
63 lifetime information, and support these characteristics by a systematic comparison between detectors (SS2 and ICCD) and  
64 methods (NLSF and phasor analysis), demonstrating the suitability of SS2 for these challenging *in vitro* and *in vivo* applications.

## 65 II. RESULTS

### 66 *NIR-MFLI lifetime measurement as a function of molecular environment (IRDye 800CW-2DG)*

67 To assess SS2's ability to resolve minute differences in NIR lifetimes, such as those expected for fluorescent dyes exposed  
68 to different microenvironments, solutions of IRDye 800CW conjugated to 2-deoxyglucose (2DG) were prepared in aqueous  
69 buffers with different pH, as well as in DMSO. IRDye 800CW-2DG is typically used as a reporter of metabolic activity in small  
70 animal model. Although no systematic study of environmental effects on IRDye 800CW has been published, there is evidence of  
71 pH effects on fluorescence quantum yield, and hence radiative lifetime, in other cyanine dyes; for instance, some ICG derivatives  
72 exhibit higher quantum yields and lifetimes at lower pH<sup>27</sup>. Similarly, DMSO has been observed to increase the quantum  
73 yield/radiative lifetime of ICG, among other indocyanine derivatives<sup>28</sup>. We thus performed MFLI studies of solutions of IRDye  
74 800CW-2DG in various environments with both SS2 and ICCD, and compared the resulting lifetimes obtained with either NLSF  
75 or phasor analysis.

76 Nonlinear Least Square Fit (NLSF, see *Online Methods*) is a standard technique which only requires an instrument response  
77 function (IRF) measurement, as well as good signal to noise ratio (SNR) to obtain accurate results, but can be computationally  
78 demanding<sup>25,29</sup>. A user-friendly and fit-free alternative to NLSF analysis is the phasor approach<sup>18,30-34</sup>, which is based on the  
79 computation of a pair of Fourier coefficients of the decay (Fourier harmonic  $n$  or phasor frequency  $f = n/T$ ), which are then  
80 represented as a point in the *phasor plot*. In the case of species characterized by mono-exponential decays, phasors are located on  
81 the *universal circle* (UC) and their lifetime can be retrieved geometrically (referred to as the *phase lifetime*, see *Online Methods*).  
82 Phasor analysis of *time-gated* decays proceeds similarly, with the universal circle replaced by a modified curve<sup>36</sup> (dubbed *single-*  
83 *exponential phasor locus* or SEPL following ref. <sup>34</sup>), which is barely distinguishable from the UC as the resolution (number of  
84 gates) increases. Other differences can appear in the case of (i) incomplete or truncated decays (as is the case for the ICCD data  
85 obtained in this study), where a different phasor frequency  $f = 1/D$  (decay support window length  $D < T$ ) is preferable, or (ii)  
86 gate with non-ideal shapes (which characterize both detectors used here)<sup>34</sup>. Details on the best way to handle these differences  
87 can be found in ref. <sup>34</sup> and are summarized in the *Online Methods*.

88 We first verified that both methods retrieve lifetimes accurately independently from signal intensity by analyzing solutions  
89 of a single NIR dye prepared at different concentration (see Supplementary Note 1 and Fig. S2).

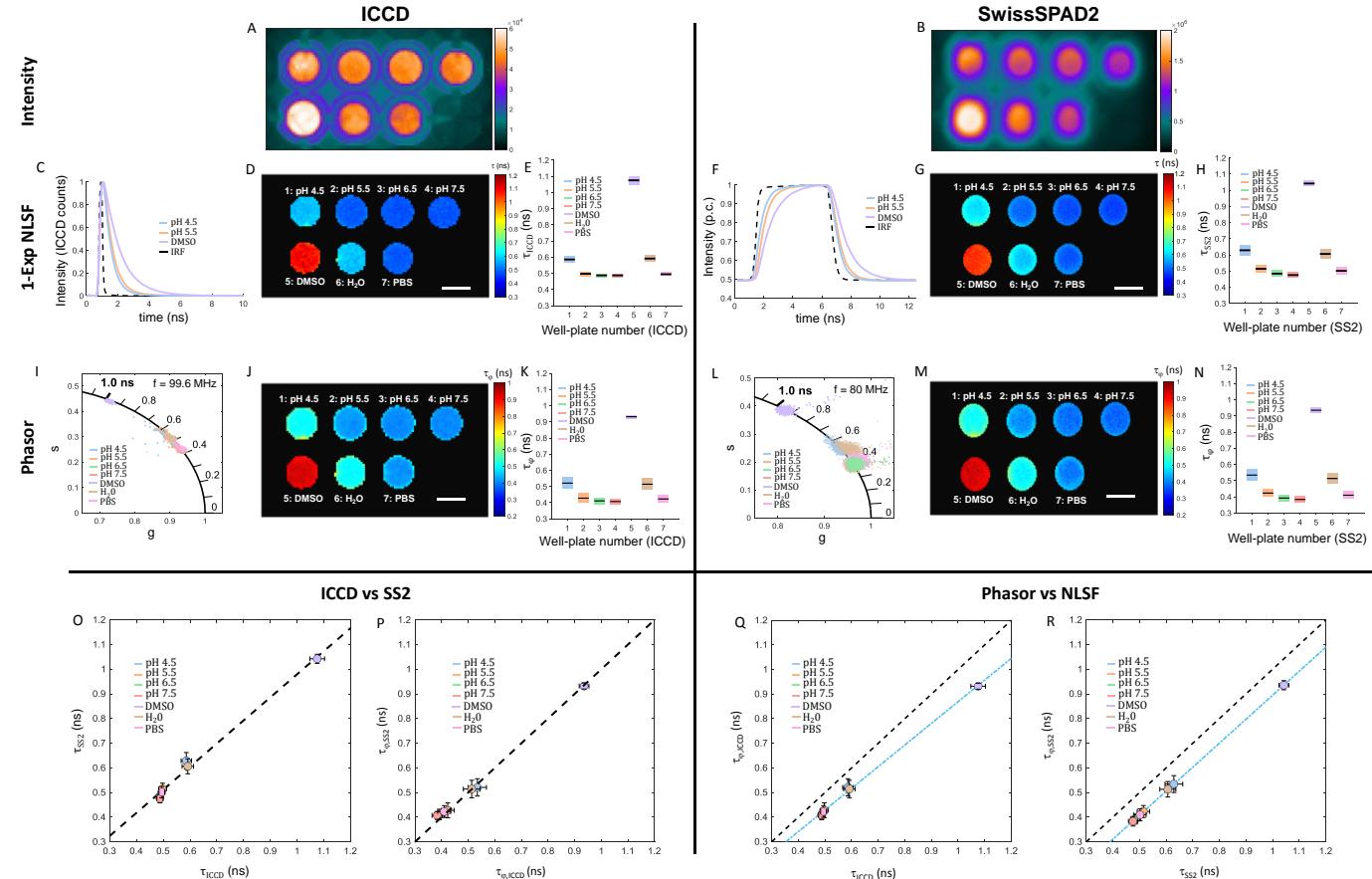
### 90 NLSF Analysis

91 As shown for the AF750 measurements (Fig. S2), intensity maps obtained with both ICCD and SS2 for IRDye 800CW (Fig.  
92 1A & B, respectively) are comparable despite setup differences. However, very distinct normalized decays are observable in  
93 some wells, indicative of different lifetimes (Fig. 1C and 1F for the ICCD and SS2, respectively). Lifetime results obtained for  
94 both imagers (Fig. 1E,H) indicate no variation of IRDye 800CW's lifetime ( $\tau = 0.5$  ns) as function of pH in the range 5.5 to 7.5,  
95 but a measurable increase at lower pH ( $\tau = 0.6$  ns at pH 4.5), which is also the pH of our distilled water ( $\text{H}_2\text{O}$  sample). Interestingly,  
96 the lifetime in DMSO more than doubles ( $\tau = 1.1$  ns) compared to the values in aqueous buffers, likely reflecting the increased  
97 orientation polarizability of DMSO, as observed for similar dyes<sup>37,38</sup>. As observed before, both detectors provide  
98 essentially identical results (Fig. 1O), with a maximum relative difference between the two measurements of 2.7%.

99

## Phasor analysis

100 Phasor plots for all IRDye 800CW-2DG mixtures (Fig. 1I,L), as well as corresponding pixel-wise phase lifetime maps (Fig. 101 1J,M), are shown in Fig. 1 for the ICCD and SS2. The phase lifetime results obtained with both cameras are in excellent 102 qualitative and quantitative agreement with one another (Fig. 1P). Comparison between phase lifetime ( $\tau_p$ ) and NLSF lifetime ( $\tau$ ) 103 for each well shows a small positive bias (<100 ps) for NLSF lifetimes (Fig. 1Q,R). Because it is common to both cameras, this 104 suggests that the IRF data used for convolution (NLSF) and calibration (phasor analysis) might not be perfectly adequate (see 105 *Online Methods*). In any case, once characterized, this systematic bias can easily be corrected for post-analysis and will be 106 neglected in the remainder of this study.



107 Fig. 1: Environment effects on IRDye 800CW-2DG. A, B: Fluorescence intensity images. A, ICCD: MCP voltage = 400 V, integration time = 108 308 ms, illumination power = 0.76 mW/cm<sup>2</sup>. B, SS2: integration time = 1.02 s, illumination power = 1.53 mW/cm<sup>2</sup>; All wells were prepared at 109 constant fluorophore concentration (15  $\mu$ M) and are labeled with their respective buffer/pH. C, F: Representative ICCD and SS2 normalized 110 single-pixel decays for 3 of the wells, plotted with the corresponding IRF. D, G: lifetime maps obtained by NLSF. E, H: Boxplot summarizing 111 lifetime results for all wells. J, M: Phasor plot for each well and K, N: pixel-wise phase lifetime maps obtained with the ICCD and SS2 112 detectors. L, O: Boxplot quantifying phase lifetime results for all wells. O: Scatter plot of averaged lifetime results for ICCD and SS2. P: 113 Scatter plot of SS2 phase lifetime versus ICCD phase lifetime. Q, R: Scatter plot of averaged NLSF lifetime results versus averaged phase 114 lifetime results for both ICCD and SS2, respectively. Error bars represent 1-standard deviation in O-R. Scale bar in A, B, D, G, J, M: 3 mm. 115 116

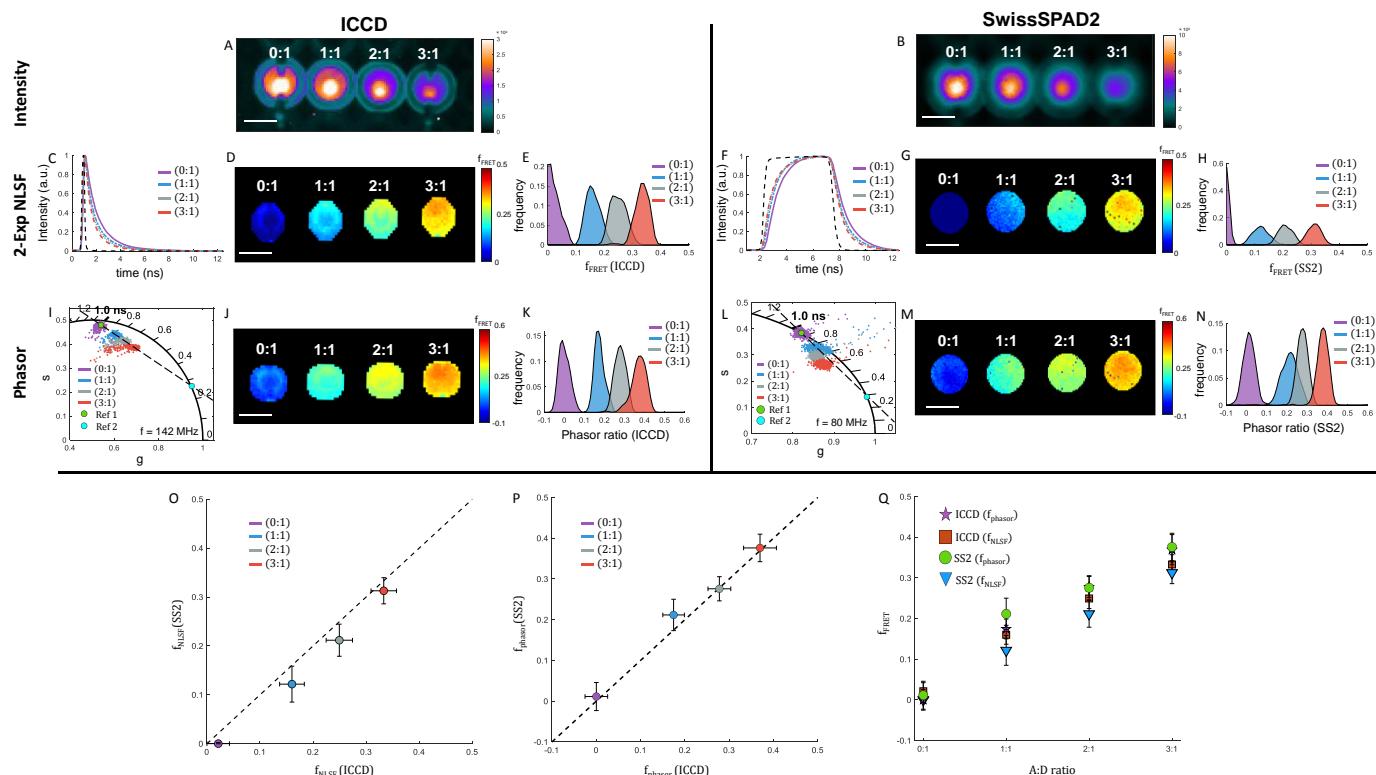
## 117 *In vitro* NIR MFLI-FRET measurements: IgG-AF700/anti-IgG-AF750 mixtures

118 FRET between neighboring fluorescent dyes having overlapping emission (donor molecule) and absorption spectra 119 (acceptor molecule) is widely used as a molecular ruler<sup>39</sup> but also as a qualitative reporter of protein-protein interactions. FRET 120 acts as a proximity assay thanks to its nanoscale range, set by the Förster radius  $R_0$ , of the order of a few nm, which depends on 121 the donor and acceptor photophysical properties. One of the characteristic signatures of FRET is a decrease of the donor 122 fluorescence lifetime,  $\tau_D$ , in the presence of a nearby acceptor molecule ( $\tau_{DA} < \tau_D$ ). In an ideal mixture of interacting donor- 123 and acceptor-labeled molecules, donor molecules are either engaged in FRET with an acceptor molecule at a certain distance (FRET 124 fraction:  $f_{FRET}$ , lifetime:  $\tau_{DA}$ ) or isolated (donor-only fraction: 1-  $f_{FRET}$ , lifetime:  $\tau_D$ ), resulting in an observed bi-exponential 125 fluorescence decay. The FRET fraction  $f_{FRET}$  can then be quantified by a fit of the measured decay to a bi-exponential model, or

126 indirectly, by measuring the average lifetime. Increasing FRET fraction decreases the effective donor lifetime, measured as either  
 127 the average donor lifetime or approximated as the best single-exponential decay fit<sup>40,41</sup>.  
 128 AF700/AF750 is a NIR FRET pair suitable for *in vivo* optical imaging applications, which we have used to assess  
 129 transferrin<sup>17-19</sup> and TZM<sup>42,43</sup> tumor delivery and efficacy in small animal models. To verify SS2's ability to quantify FRET, a  
 130 controlled *in vitro* NIR MFLI-FRET experiment was first carried out. A multiwell-plate was prepared using the AF700/AF750  
 131 FRET pair (Förster radius  $R_0 \sim 8 \text{ nm}$ <sup>44</sup>), with each dye conjugated to a complementary IgG and anti-IgG pair (IgG-AF700 and  
 132 anti-IgG-AF750 respectively). Each well is characterized by a different acceptor-to-donor ratio (A:D, of the form  $n:1$  with  $n = 0 -$   
 133 3). As before, we acquired data with both detectors using similar conditions, and processed data using both NLSF and phasor  
 134 analysis.

### 135 NLSF analysis

136 *Single-exponential* NLSF analysis reveals the expected trend of a decreasing donor lifetime with increasing A:D ratio (Fig.  
 137 S4), as well as close agreement between both detectors. While this agreement is important, single-exponential fits are not  
 138 sufficient for the quantitative study of FRET mixtures, as there is no simple relationship between fitted single-exponential  
 139 lifetime  $\tau$  and FRET fraction  $f_{\text{FRET}}$ . Hence, *bi-exponential* NLSF analysis of the data provides a more accurate interpretation of  
 140 the data despite relying on some simplifying assumptions. Here, the lifetimes of donor-only ( $\tau_1 = \tau_D = 1 \text{ ns}$ ) and quenched-donor  
 141 ( $\tau_2 = 0.265 \text{ ns}$ ) were retrieved through ROI-level decay fitting (Fig. S5).



142  
 143 Fig. 2: AF700/AF750 FRET pair series. A, B: Fluorescence intensity images. A: ICCD; MCP voltage = 450 V, integration time = 359 ms,  
 144 illumination power = 1.52 mW/cm<sup>2</sup>. B: SS2; integration time = 4.08 s, illumination power = 2.29 mW/cm<sup>2</sup>; Wells contained solutions of  
 145 labeled antibodies in PBS buffer with acceptor to donor A:D ratio from 0:1 to 3:1 indicated above each well, with a constant donor fluorophore  
 146 concentration of 32  $\mu\text{M}$ . C, F: Representative ICCD and SS2 normalized single-pixel decays for the different wells, plotted with the  
 147 corresponding IRF. D, G: MFLI FRET-fraction maps obtained by bi-exponential NLSF for both cameras. E, H: Corresponding  $f_{\text{FRET}}$  KDE  
 148 distributions for each well obtained for both cameras. I, L: phasor scatter plots overlaid with linear fit (dashed black line) and reference  
 149 lifetimes set as the centroid of the donor-only well's cluster (green dot, reference 1) and as the linear fit intersection with the SEPL (blue dot,  
 150 reference 2) – resulting in  $\tau_{1,ICCD} = 0.98 \text{ ns}$ ,  $\tau_{2,ICCD} = 0.27 \text{ ns}$  and  $\tau_{1,SS2} = 1.01 \text{ ns}$ ,  $\tau_{2,SS2} = 0.17 \text{ ns}$ . J, M: pixel-wise phasor ratio maps and K, N:  
 151 phasor-ratio KDE distributions for the different wells (acceptor-donor ratios indicated A:D in parenthesis, color code matching that of panels J,  
 152 L). O: Scatter plot (mean  $\pm$  standard deviation) showing the FRET-fraction measured via bi-exponential NLSF for each well with the ICCD  
 153 versus that measured with SS2. P: Scatter plot of phasor ratio results (mean  $\pm$  standard deviation) for ICCD and SS2. R: Scatter plot of FRET  
 154 fractions obtained by NLSF (panels C-H) and phasor ratio results for ICCD and SS2 (panels I-N) as a function of A:D ratio. Scale bar in A, B,  
 155 D, G, J, M: 3 mm.

156 Following conventional practices, these values of  $\tau_1$  and  $\tau_2$  were fixed for all subsequent pixel-level analyses, leaving only 3  
 157 free parameters for the NLSF analysis: baseline  $B$  and amplitudes  $A_1$  and  $A_2$  (*Online Methods*, Eq. (1)). The FRET intensity

158 fraction  $f_{FRET}$  (*Online Methods*, Eq. (4)) values yield similar results for both imagers (Fig. 2D,G,E,H). Importantly, both results  
159 reveal the expected linear increase of the calculated FRET fraction as a function of acceptor-to-donor ratio (Fig. 2O).

## 160 Phasor analysis

161 Although NLSF analysis as performed above provides important information, it relies on assumptions that may not always  
162 be fulfilled. By contrast, phasor analysis of mixtures does not rely on any specific functional form for the reference decays  
163 (donor only species or fully quenched donor decay). However, because the decays of mixtures are linear combinations of donor  
164 and quenched donor decays, their phasors are also linear combinations of the donor and quenched donor phasors, and are  
165 therefore aligned along a segment connecting donor and quenched donor phasors<sup>18,45</sup>.

166 As hypothesized in the NLSF analysis above, in the ideal case where the donor-only species (reference 1) and the FRET pair  
167 species (reference 2) can be assumed to be single-exponential decays, the phasor of each species is located on the SEPL. The  
168 relative distance of a particular sample's phasor with respect to reference 2 (called phasor ratio<sup>18,46</sup>), is equal to the intensity  
169 fraction of reference 1 (Eq. (8)). Hence, in that ideal case, the calculated intensity fraction should be equal to that obtained by  
170 bi-exponential NLSF analysis of the recorded decay (*Online Methods*, Eq. (4)).

171 In panels I-N of Fig. 2, we show the result of phasor analysis of the datasets discussed above. The references used for phasor  
172 ratio quantification were retrieved by computing the intersection of the SEPL with a line obtained as best fit through phasors  
173 from all four wells (Fig. 2I,L). The resulting phasor ratio analysis (Fig. 2J,K,M,N) shows a remarkable agreement for both  
174 cameras (Fig. 2P).

175 FRET fraction results retrieved through both cameras using NLSF and phasor are in good agreement (Fig. 2Q). Noticeably,  
176 in the case of phasor ratio analysis, although both datasets are computed with different phasor frequencies and with automatically  
177 defined references, both analyses result in reference phasors which are remarkably similar to those chosen for NLSF analysis  
178 (donor-only:  $[\tau_{1,ICCD} = 0.98$  ns,  $\tau_{1,SS2} = 1.01$  ns] and quenched donor:  $[\tau_{2,ICCD} = 0.27$  ns and  $\tau_{2,SS2} = 0.17$  ns], respectively).

## 179 *In vivo* NIR MFLI-FRET measurements: Trastuzumab-HER2 engagement in mouse xenografts

180 Having established the *in vitro* equivalence of time-gated ICCD data and SS2 data, we extended this comparison to *in vivo*  
181 small animal experiments. We used xenograft models of human breast cancer (AU565) and ovarian cancer (SK-OV-3) selected  
182 due to their HER2 overexpression and distinct challenges regarding effective drug delivery<sup>47</sup>, to compare drug-receptor  
183 engagement efficiency across two tumor models. AU565 and SK-OV-3 cells overexpress HER2 receptors, which are targeted by  
184 Trastuzumab (TZM), a monoclonal antibody used as anti-HER2 breast cancer therapy. In our approach, AF700 or AF750 are  
185 conjugated to TZM and intravenously injected into mice carrying HER2 positive tumor xenografts. Thus, the amount of recorded  
186 FRET between the two probes provides a measure of TZM-receptor binding. Noninvasive MFLI-FRET imaging of nude mice  
187 bearing AU565 and SK-OV-3 tumor xenografts was performed upon intravenous injection of NIR-labeled TZM pair with both  
188 time-resolved detectors in two different mice (see *Online Methods*). Mice are imaged at 24 hr post-injection (p.i.) using the  
189 ICCD (mouse 1) or SS2 (mouse 2), followed by two other imaging sessions of mouse 2 at 48 hr (SS2) and 51 hr (ICCD) p.i.,  
190 respectively.

## 191 NLSF Analysis

192 NLSF analysis of the 24 hr p.i. data was limited to the two tumors and urinary bladder regions of interests (ROIs) in which  
193 significant AF700 signal was observed (Fig. 3A,B). As for the *in vitro* FRET analysis, pixel-wise single-exponential NLSF  
194 analysis of the observed decays (Fig. 3C,F) was first performed to obtain a qualitative understanding of the data (Supplementary  
195 Fig. S6C-F). The lifetime of AF700-TZM in the urinary bladders is close to 1 ns and distributed uniformly. These results are  
196 consistent with the expected detection of donor-only labeled TZM as measured in different pH conditions in Fig. S7. By contrast,  
197 AF700-TZM's lifetime is noticeably shorter in the AU565 tumors, as previously reported<sup>43</sup>, but not in SK-OV-3, suggesting that  
198 FRET and thus TZM-HER2 binding is occurring at a higher level in AU565 tumors than SK-OV-3.

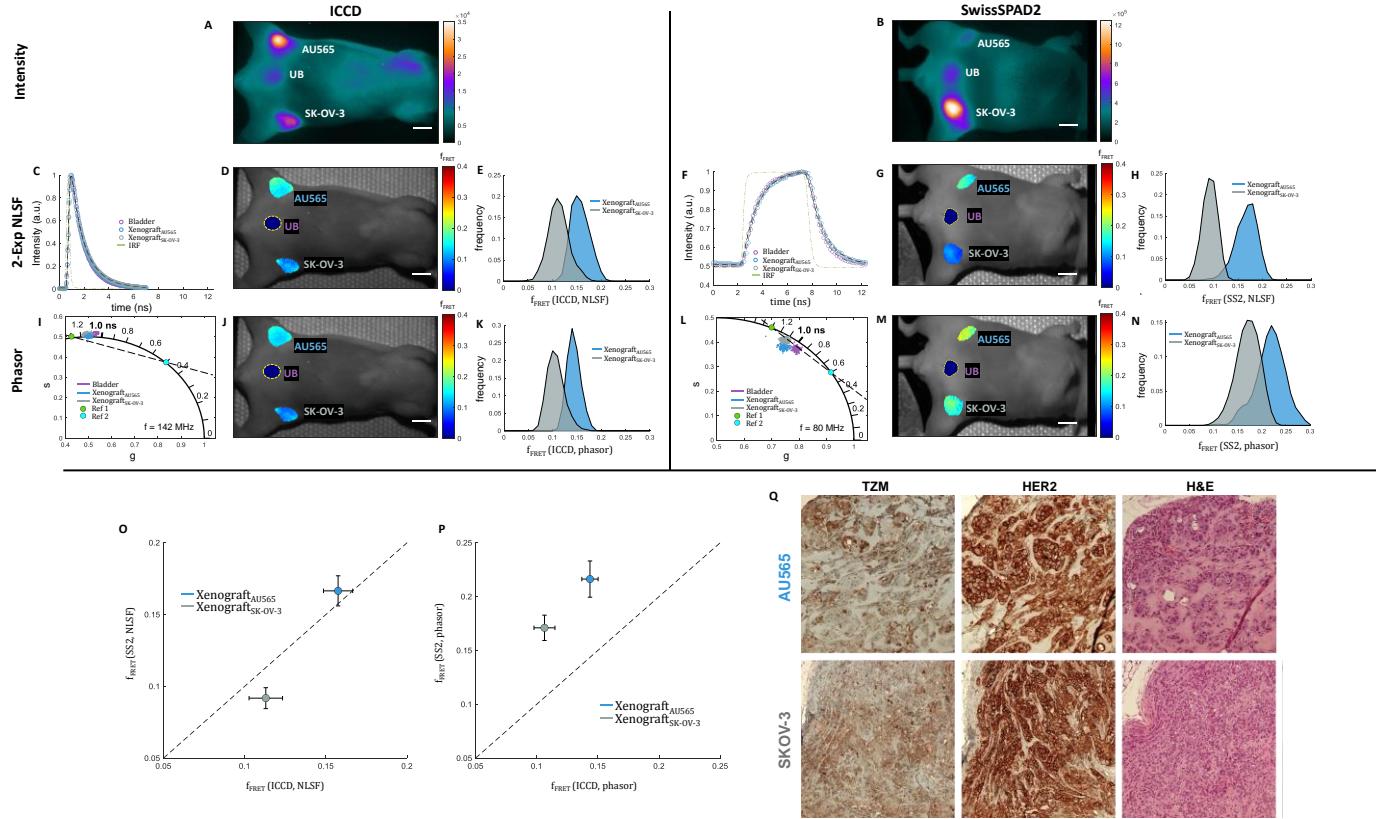
199 For FRET quantification, bi-exponential NLSF analysis of the decays was performed in similar fashion to that of *in vitro*  
200 before (Fig. 2). Donor-only and quenched-donor lifetimes were retrieved through unconstrained full ROI decay fitting ( $\tau_1 = 1.3$   
201 ns  $\tau_2 = 0.5$  ns). FRET fraction results obtained for mouse 1 (ICCD: Fig. 3D,E) and mouse 2 (SS2: Fig. 3G,H) are in high  
202 agreement. A direct comparison (Fig. 3O) shows that the two tumors behave remarkably similarly in both mice, with the AU565  
203 xenograft exhibiting close to 50% more FRET ( $15.9 \pm 1.9\%$  vs  $10.4 \pm 2.1\%$ ) than the SK-OV-3 (ovarian cancer) xenograft (Fig.  
204 S6 and Supplementary Note 2). Importantly, this pattern is preserved over time as shown in Figs. S8-S9, which summarize the  
205 analysis of mouse 2, observed at 48 hr p.i. (with SS2), and at 51 hr p.i. (with the ICCD).

## 206 Phasor Analysis

207 As *in vitro*, phasor ratio analysis (Fig. 3I-N) was performed using the reference lifetimes obtained in the NLSF analysis ( $\tau_1 =$   
208  $1.3$  ns  $\tau_2 = 0.5$  ns). Noticeably, phasor ratio maps (Fig. 3J,M) and corresponding distributions for each tumor (Fig. 3K,N)  
209 demonstrate a remarkable consistency between methods and across mice.

210 Immunohistochemistry

211 IHC analysis of excised tumors (Fig. 3Q) is in general agreement with MFLI data, showing more intracellular TZM  
 212 accumulation in the AU565 tumors than in SK-OV-3 (Fig. 3Q left) despite high HER2 expression in both tumors (Fig. 3Q  
 213 center). Reduced TZM accumulation in SK-OV-3 vs. AU565 tumors is consistent with decreased HER2-TZM binding as  
 214 indicated by the lower FRET signal observed in SK-OV-3 vs. AU565 tumors via MFLI imaging with ICCD or SS2 detectors.



215

216 Fig. 3: *in vivo* Trastuzumab-Her2 receptor engagement. Mice were injected with 20  $\mu$ g of AF700-TZM and 40  $\mu$ g of AF750-TZM and imaged  
 217 by MFLI at 24 h post-injection (p. i.). Mouse 1 was imaged with the ICCD and mouse 2 with SS2. A, B: Fluorescence intensity images. A:  
 218 ICCD; MCP voltage = 500V, integration time = 500 ms, illumination power = 2.13 mW/cm<sup>2</sup>. B: SS2; integration time = 2.65 s, illumination  
 219 power = 3.2 mW/cm<sup>2</sup>; C, F: ICCD and SS2 normalized whole ROI decays for the different organs, plotted with the corresponding IRF. D, G:  
 220 MFLI FRET-fraction maps obtained by bi-exponential NLSF for both cameras. The urinary bladder (yellow dashed outline) was analyzed by 1-  
 221 Exp NLSF and is therefore not included. E, H: Corresponding  $f_{FRET}$  KDE distributions for each well obtained for both cameras. I, L: phasor  
 222 scatter plots color-coded by ROI, with overlaid reference lifetimes (green dot, reference 1; blue dot, reference 2) and dashed black line  
 223 connecting them. J, M: pixel-wise phasor ratio maps and K, N: phasor-ratio KDE distributions for the two xenografts. O: Scatter plot (mean  $\pm$   
 224 standard deviation) showing the FRET-fraction measured for each tumor with SS2 (mouse 2) versus that measured with the ICCD (mouse 1)  
 225 retrieved through bi-exponential NLSF. P: Scatter plot of phasor ratio results (mean  $\pm$  standard deviation) for ICCD and SS2. Scale bar in A,  
 226 B, D, G, J, M: 6 mm. Q: *ex vivo* IHC of intracellular accumulation of TZM in AU565 and SK-OV-3 tumors in mouse 2. Consecutive sections  
 227 were processed for H&E (showing cell localization and context), anti-HER2, and anti-TZM immunohistochemical staining. NovaRED was  
 228 used as peroxidase substrate (brown stain), tissue was counterstained with methyl green. Scale bar = 100  $\mu$ m.

229 In summary, our experiments confirm our previous observation of a significant level of TZM-HER2 engagement in AU565  
 230 (breast cancer) tumor xenografts<sup>43</sup>. Moreover, we show that MFLI imaging can discriminate between different types of HER2  
 231 tumor models (breast AU565 and ovarian SK-OV-3) exhibiting distinct levels of TZM tumor delivery and HER2 binding,  
 232 despite both tumor models showing HER2 overexpression. Our results suggest that these tumor models may display distinct  
 233 tumor microenvironment challenges regarding effective drug delivery<sup>47</sup>. A manuscript focusing on the biological mechanisms  
 234 underlying these differences in drug delivery between HER2 positive tumor models is currently in preparation.

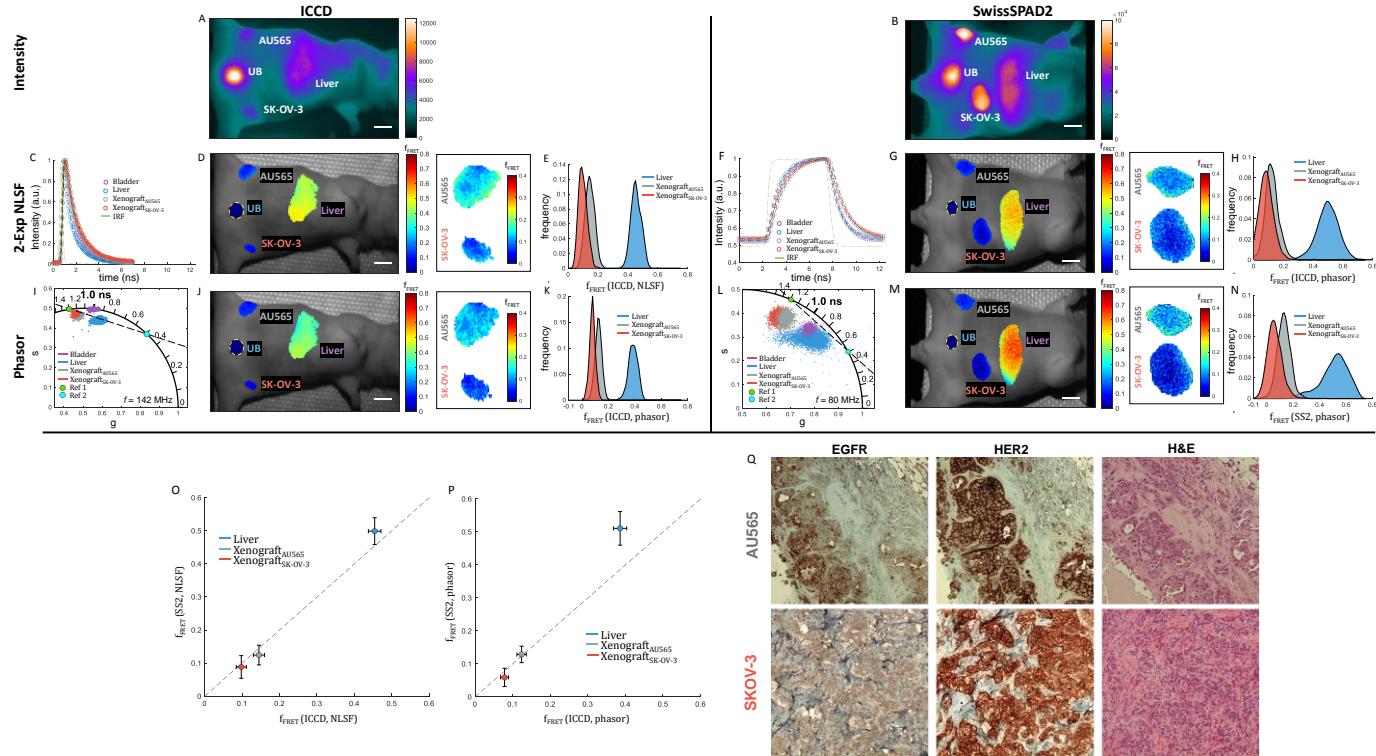
235 *In vivo* NIR MFLI-FRET measurements: Cetuximab-EGFR engagement in mouse xenografts

236 Cetuximab (CTM) is another monoclonal antibody used as EGFR inhibitor in clinical treatment of breast, colon, and head-  
 237 and-neck cancers. As for TZM, AF700- and AF750-labeled CTM can be used in a NIR FLI-FRET assay to monitor CTM-EGFR  
 238 drug-receptor engagement, as we recently demonstrated *in vitro*<sup>48</sup>. Here, similarly to the TZM experiment reported above,  
 239 noninvasive MFLI-FRET imaging of nude mice bearing AU565 and SK-OV-3 tumor xenografts was performed. As described in

240 *Online Methods*, two different mice were imaged at 48 hr p.i., either with the ICCD (mouse 1) or with SS2 (mouse 2). Fig. 4A,B  
 241 shows the observed fluorescence intensity maps. In contrast to the TZM fluorescence experiment (Fig. 3), four regions are  
 242 clearly visible: the liver as well as both tumor xenografts and the urinary bladder.  
 243

### NLSF Analysis

244 Following the workflow outlined in the *in vivo* TZM-HER2 experiment (Fig. 3), bi-exponential NLSF analysis of pixel-wise  
 245 decays was performed using fixed component lifetimes obtained from whole ROI decay analysis (Fig. 4C,F:  $\tau_1 = 1.3$  ns  $\tau_2 = 0.5$   
 246 ns). As observed in the case of TZM FRET pair, the resulting FRET fraction quantification (Fig. 4D,G,E,H) in the SK-OV-3  
 247 tumor ( $f_{SK-OV-3} \sim 8.9 \pm 4.7\%$ ) are smaller than those measured in the AU565 tumors ( $f_{AU565} \sim 13.5 \pm 4.4\%$ ) and significantly  
 248 lower than what is observed in the liver of both mice ( $f_{Liver} \sim 48.5 \pm 7.3\%$ ), irrespective of the camera used. Comparison of  
 249 results obtained in both mice using two different cameras (Fig. 4O) show once again exceptional agreement.



250

251 Fig. 4: *in vivo* Cetuximab-EGFR engagement. Mice were injected with 20  $\mu$ g AF700-CTM and 40  $\mu$ g AF750-CTM and subjected to MFLI  
 252 imaging at 48 h p.i. (mouse 1: ICCD, mouse 2: SS2). A, B: Whole body fluorescence intensity images for ICCD and SS2. A: ICCD; MCP  
 253 voltage = 520 V, integration time = 500 ms, illumination power = 2.13 mW/cm<sup>2</sup>. B: SS2; integration time = 2.45 s, illumination power = 3.2  
 254 mW/cm<sup>2</sup>. C, F: ICCD and SS2 normalized whole ROI decays for the different organs, plotted with the corresponding IRF. D, G: MFLI FRET-  
 255 fraction maps obtained by bi-exponential NLSF for both cameras. The urinary bladder (yellow dashed outline) was analyzed by 1-Exp NLSF  
 256 and is therefore included as a constant 0 fraction. A zoomed in view of  $f_{FRET}$  quantification retrieved for both xenografts with color scale  
 257 adjusted to match that used in Fig. 3 is shown on the right. E, H: Corresponding  $f_{FRET}$  KDE distributions for each well obtained for both  
 258 cameras. I, L: phasor scatter plots color-coded by ROI, with overlaid reference lifetimes (green dot, reference 1; blue dot, reference 2) and  
 259 dashed black line connecting them. J, M: Pixel-wise phasor ratio maps. A zoomed in view of phasor ratio quantification retrieved for both  
 260 xenografts with color scale adjusted to match that used in Fig. 3 is shown on the right. K, N: Phasor-ratio KDE distributions for the two  
 261 xenografts and the liver. O: Scatter plot (mean  $\pm$  standard deviation) showing the FRET-fraction measured for each tumor and the liver with  
 262 SS2 (mouse 2) versus that measured with the ICCD (mouse 1). P: Scatter plot of phasor ratio results (mean  $\pm$  standard deviation) for ICCD and  
 263 SS2. Scale bar in A, B, D, G, J, M: 6 mm. Q: *ex vivo* IHC validation of intracellular localization of EGFR and HER2 in AU565 and SK-OV-3  
 264 tumors in mouse 2. Consecutive sections were processed for H&E (showing cell localization and context), anti-HER2, and anti-EGFR  
 265 immunohistochemical staining. NovaRED was used as peroxidase substrate (brown stain), tissue was counterstained with methyl green. Scale  
 266 bar = 100  $\mu$ m.

### Phasor Analysis

267 Pixel-wise phasor ratio analysis for each mouse (Fig. 4I-N) shows a similar pattern to that observed previously for the  
 268 TZM-HER2 experiment (Fig. 3-NL), with the notable difference of the presence of a liver phasor cluster. It is clear from both the  
 269 phasor clusters themselves, as well as from the superimposed references (green dot, reference 1; blue dot, reference 2) and

271 dashed black line connecting them, that tumors and liver phasors can be interpreted as a linear combination of a short (blue dot)  
272 and a long (green dot) single-exponential component.

273 Comparison of the FRET fractions obtained by this method with those obtained by NLSF analysis shows a truly remarkable  
274 correspondence, both between methods (compare Fig. 4D,E to 4J,K and 4G,H to 4M,N) and across mice (compare Fig. 4D,E to  
275 4G,H and 4J,K to 4M,N). While there still exists a small offset between results obtained by the two methods, as indicated in Fig.  
276 4O-P, the difference between tumors, and between tumor and liver is crucially preserved, further validating our approaches.

277 **Immunohistochemistry**

278 The differences between tumors observed by MFLI-FRET are supported by IHC analysis (Fig. 4Q), which looked at anti-  
279 EGFR (left) and anti-HER2 (center) immunostaining, in addition to standard H&E staining (right). In contrast to the HER-TZM  
280 experiment, in which we monitored both the HER2 receptor and its respective antibody (TZM), here we examine only the  
281 receptors (EGFR and HER2) and not the actual probe (CTM). Therefore, we cannot directly confirm the uptake of CTM by  
282 AU565 or SK-OV-3 tumors. However, the comparatively low anti-EGFR staining in the SK-OV-3 tumor is consistent with the  
283 relatively lower FRET signal observed in SK-OV-3 xenografts.

284 In summary, our experiments confirm our previous observation of a significant level of CTM-HER2 engagement in AU565  
285 (breast cancer) tumor xenografts<sup>48</sup>. The detection of a strong CTM FRET signal in the liver is also consistent with the known  
286 EGFR expression in mice livers. MFLI imaging with both ICCD and SS2 can therefore clearly discriminate different levels of  
287 CTM-EGFR drug-receptor tumor binding in two distinct EGFR-positive tumor models.

288 **III. Discussion**

289 MFLI provides a unique contrast mechanism that enables monitoring key biological parameters dynamically and  
290 noninvasively. To perform such measurements in live specimen requires using NIR probes characterized by very short lifetimes,  
291 a challenge for current time-resolved detectors, which have lower detection efficiency in the NIR spectral range. In the past,  
292 time-gated ICCDs have been the workhorse for this type of MFLI studies. However, their high cost and limitations justify the  
293 exploration of alternative technologies. New development in SPAD technology has led to the integration of large time-gated  
294 CMOS SPAD cameras such as SwissSPAD2. These new cameras represent the next generation of imager for FLI studies across  
295 numerous biomedical applications. Using such cameras in the NIR spectral range is a priori challenging, and compounded by  
296 additional specificities of SS2 such as gate duration or rise- and fall-times, which are longer or of the order of magnitude of the  
297 lifetimes to be measured. We show here that these are not obstacles to accurate lifetime measurements. Indeed, the gate step size  
298 and the photon count, more than the gate duration or its rise- and fall-times, are the primary parameters affecting the measured  
299 lifetime precision<sup>18,24,25</sup>.

300 SS2 recovered FLI parameters of interest with high accuracy and in remarkable agreement with a state-of-the-art gated-  
301 ICCD. The outstanding performance of SS2 for MFLI applications ranges from resolving lifetimes of only a few hundreds of  
302 picosecond and lifetime differences of a few tens of ps. More importantly, this performance extends to multi-exponential decays  
303 analysis, where SS2 provides accurate estimates of FRET donor fraction.

304 These observations are further supported by *in vivo* studies which provided quantitative measurements of FRET donor  
305 fraction, a direct readout of drug-target engagement, over the whole animal body. SS2-measured FRET fractions were consistent  
306 with the molecular biology expected in the different organs in which signal was detected, including tumor xenografts with  
307 various target expression levels. Importantly, all *in vivo* experiments were performed at low illumination power (~2.5 mW/cm<sup>2</sup>),  
308 well beyond the Maximum Permissible Exposure (MPE) limit, and with excellent SNR (~400-1200, see Supplementary Tables  
309 2-5)—even in deep seated tissues. This suggests that faster imaging rates can be achieved for applications with less demanding  
310 SNR constraints, potentially achieving live imaging capabilities.

311 By contrast with ICCDs, SS2's technology offers room for significant future improvements: micro lenses can be added for  
312 up to 4-fold increased collection efficiency<sup>24</sup>, and upcoming next-generation SwissSPAD detectors will feature (i) shorter gate  
313 duration for better resolution, (ii) dual-gate architecture for more efficient photon collection and (iii) rolling-shutter recording, as  
314 well as (iv) higher recording rate to match that of the excitation laser, resulting in higher duty cycle. Moreover, the built-in field-  
315 programmable gate-array used to control the detector can be used for additional data pre-processing, opening the door for  
316 additional gains in speed and performance.

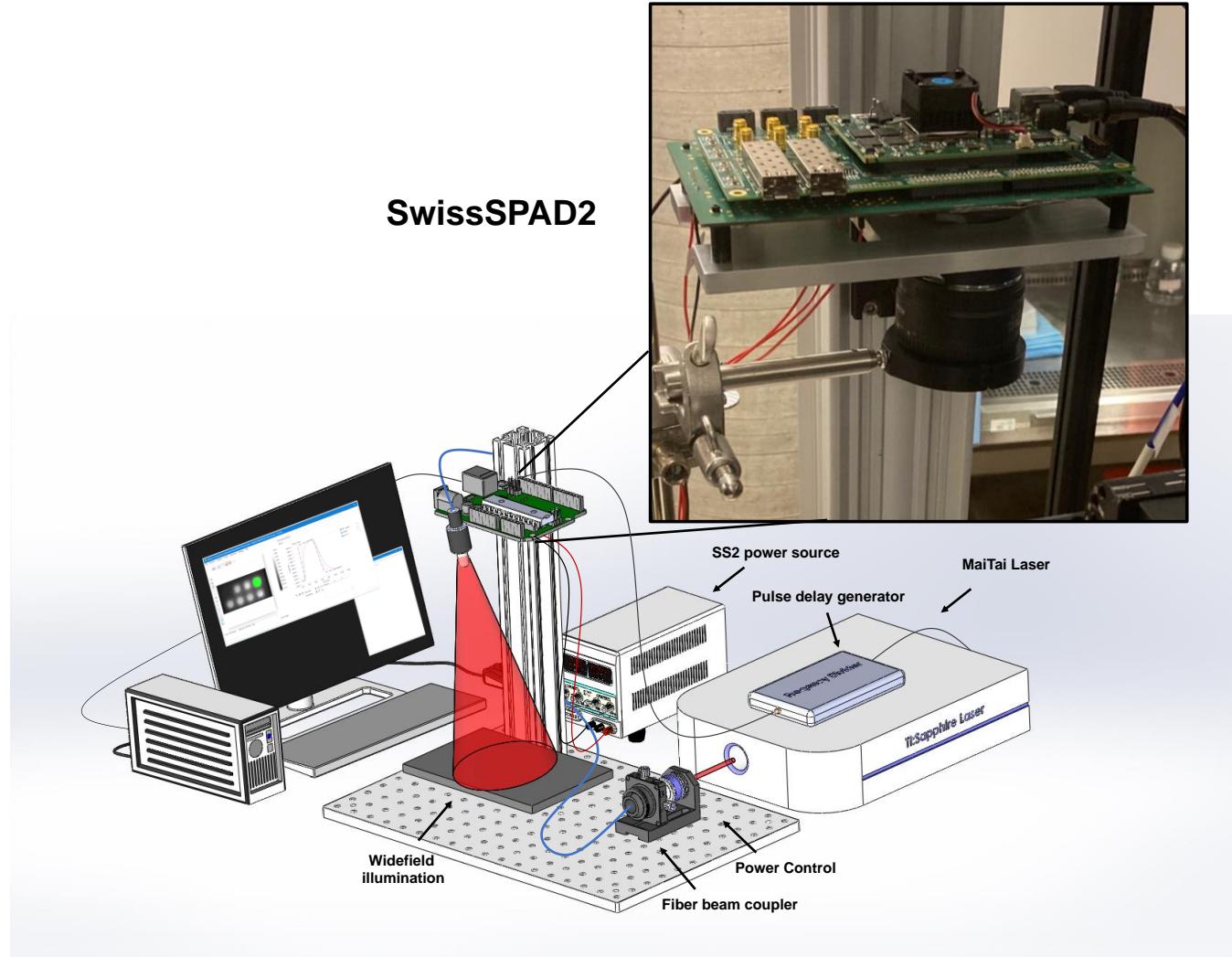
317 Our work presents a systematic comparison of two improved MFLI analysis approaches: (i) single- and bi-exponential  
318 NLSF analysis using full-period periodic convolution using a measured IRF and global lifetime constraints and (ii) phasor  
319 analysis based on calibration using a measured IRF. We demonstrate that these improvements capture the effects of sample  
320 topography and enable robust phasor analysis of target-receptor engagement *in vivo*.

321 In summary, we have validated a new large-area time-resolved SPAD camera for *in vitro* and *in vivo* macroscopic  
322 fluorescence lifetime imaging, supported by two complementary analysis approaches. This comprehensive investigation  
323 establishes the quantitative performances of SwissSPAD2 for NIR FLI imaging in complex scenarios. Due to its excellent

324 performance, room for significant improvements, small form factor footprint and reduced cost, SwissSPAD2 and the next  
325 generation of time-gated detectors are expected to become the technology of choice for macroscopic FLI imaging with impact on  
326 a wide-range of applications – ranging from preclinical studies in drug development to optical guided surgery.

## IV. Online Methods

### 328 Macroscopic FLI with SwissSPAD2



329  
330 Fig. 5: Schematic illustration of the widefield time-resolved imaging system equipped with the SwissSPAD2 camera. All imaging was  
331 performed in reflectance geometry. A fs pulsed, tunable Ti:Sapphire laser beam was directed through a power control module before  
332 coupling into a multimode fiber (blue line). The divergent output of the fiber was directed toward the sample to achieve a Gaussian  
333 illumination profile. Light emitted by the sample was collected using an overhead macroscopic objective lens directly attached to the C-  
334 mount port in front of SwissSPAD2 (SS2, green PCB boards and photograph inset). SS2 is powered by two regulated external power  
335 supplies (red lines) and a small DC adapter. A TTL pulse derived from the laser trigger signal by a frequency divider module is used to  
336 synchronize data acquisition (thick black line). Data is transferred to a PC via a USB 3 cable (thin black cable).

337 An illustration of the SwissSPAD2 MFLI configuration (reflectance geometry) used herein is provided in Fig. 5.  
338 Technical details about the camera can be found in earlier publications<sup>22-24</sup> and are briefly summarized below. The system's  
339 excitation source was a tunable Ti-Sapphire laser (Mai Tai HP, Spectra-Physics, CA, USA). Laser excitation was directed to  
340 the sample plane directly from a multimode optical fiber (QP200-2-VIS-NIR, Ocean Optics, FL, USA) output. Emitted  
341 fluorescence was collected through an application-specific bandpass emission filter (Supplementary Fig. S10) by a  
342 macroscopic photographic lens (AF Nikkor 50mm f/1.8D, Nikon, Tokyo, Japan). IRFs (instrument response functions) were  
343 acquired similarly, using a sheet of white paper as sample (for *in vitro* experiments) or the mouse itself (in the case of *in vivo*  
344 measurements), after removing the emission filter. Slightly lower laser power and integration time were also used. For use

345 with SS2, which requires synchronization with a source signal at around 20 MHz, a frequency divider (TOMBAK, Laser Lab  
346 Source Corporation, MT, USA) was used to divide the Mai Tai's laser repetition rate ( $\sim$ 80 MHz) by a factor of four ( $f_{SYNC} =$   
347 19.77 MHz reported). SS2 was set to acquire 10-bit gate images consisting of 1,020 accumulated 1-bit gate images, each 1-  
348 bit image resulting from exposure of each SPAD pixel to the incoming photon flux for a user-specified duration of  $n_E \times 400$   
349 ns, where  $n_E$  is some number typically between 1 and 100<sup>23</sup>. During that 1-bit accumulation period, each SPAD is "on" (*i.e.*  
350 capable of detecting a photon) for a duration  $W_{SS2}$  (gate width) and "off" the remainder of each  $T_{sync} = 1/f_{SYNC} = 50.6$  ns  
351 period, and able to detect at most one photon. Each full gate image differs from the previous one by its distance to the laser  
352 pulse, or gate offset, by a user-specified amount  $\delta t$  ("gate step"), multiple of 1/56 ns = 17.857 ps.  $\delta t = 178.57$  ps was used in  
353 this work. Although gate images covering the whole 50 ns window of SwissSPAD2 max sync rate (*i.e.* 280 gates in total)  
354 were acquired, resulting in the acquisition of four laser periods-worth of data, a single laser period-worth of data (70 first  
355 gates) was used in most analyses, unless mentioned otherwise. Because the laser period  $T = 12.65$  ns is comparable to the  
356 shortest gate implemented in SS2 ( $W_{SS2} = 10.7$  ns), which results in severely deformed periodic decays, a larger gate duration  
357 ( $W_{SS2} = 17.9$  ns) was chosen to minimize this artifact (Supplementary Note 3 & Supplementary Fig. S11). To increase the  
358 total effective integration time for each gate, each acquisition was repeated multiple times (50 - 100 repetitions). Integration  
359 time for each gate image is calculated following ref.<sup>23</sup> as:  $T_{int} = (8n_G - 1)bNT_{sync}$ , where  $n_G$  is a gate sequence parameter  
360 (typically  $n_G = 100$ ),  $b$  is the number of 1-bit gate images per final image ( $b = 1,020$ ),  $N$  is the number of accumulations (e. g.  
361  $N = 70$ ) and  $T_{sync} = 1/f_{SYNC}$  is the period of the synchronization signal.  $T_{int}$  was set between 1.02-4.08 s depending on the  
362 sample brightness. Actual values of these parameters are reported when needed and are available in the raw data provided on  
363 Figshare<sup>49</sup>. Data is transferred asynchronously via a USB 3.0 connection from the SS2's field-programmable gate-array  
364 (FPGA) to the computer under control of a dedicated open-source LabVIEW program, SwissSPAD Live<sup>50</sup>.

### 365 Macroscopic FLI via Gated-ICCD

366 MFLI was also performed using a widefield time-resolved FLI apparatus equipped with a gated-ICCD (described  
367 previously<sup>51</sup>). In brief, the system used the same excitation source as described previously, but a digital micro-mirror device  
368 (Dli 4110, Texas Instruments, TX, USA) was used for widefield illumination over the sample plane. The time-gated ICCD  
369 (Picostar HR, LaVision, GmbH, Bielefeld, Germany) was set to acquire gate images with a gate width of  $W_{ICCD} = 300$  ps,  
370 separated gate steps  $\delta t = 40$  ps (details provided elsewhere<sup>52</sup>). Data was acquired over a window of duration shorter than the  
371 full laser period (generally  $G = 176$  total gate images per acquisition, *i.e.*  $D = 7$  ns, but occasionally longer, see details in each  
372 figure caption). As with SS2, IRFs were acquired with equivalent conditions to those used for fluorescence imaging. During  
373 fluorescence imaging, fluorophore-dependent filters were inserted, the ICCD's microchannel plate (MCP) voltage was  
374 increased for sufficient signal amplification (between 350-550 V depending on sample brightness) and integration time  
375 adjusted within the range 300-500 ms per gate image. Details are provided in each figure caption. Because no calibration of  
376 the camera gain (photon per camera unit signal) as a function of MCP voltage was performed for this device, no signal-to-  
377 noise ratio is reported.

### 378 Well-plate sample preparation and imaging

- 379 - *AF750 serial dilution.* To test SS2's capability to quantitatively resolve short-lifetime NIR dye at low concentrations, a  
380 serial dilution of AF750 in two distinct aqueous buffers (H<sub>2</sub>O and PBS) was prepared in a well-plate (Fig. S2).  
381 Concentrations ranged from 25  $\mu$ g/mL to 3  $\mu$ g/mL. Laser excitation was set to 750 nm and the emission filter used was  
382  $780 \pm 10$  nm (Semrock, FF01-780/12-25).
- 383 - *IRDye 800CW-2DG.* A well-plate sample of IRDye 800CW-2DG (LI-COR Biotechnology, NA, USA) was prepared at a  
384 constant concentration of 15  $\mu$ M using seven different buffers: Intracellular pH Calibration Buffer Kit (pH 4.5, pH 5.5, pH  
385 6.5, pH 7.5 – ThermoFisher, USA) distilled H<sub>2</sub>O, PBS (pH 7) and DMSO. Laser excitation was set to 760 nm and the  
386 emission filter used was 800  $\pm$  10 nm (Semrock, FF01-800/12-25).
- 387 - *AF700/AF750 IgG FRET.* A well-plate was prepared using the FRET pair AF700/AF750 IgG (ThermoFisher Scientific,  
388 MA, USA) at four acceptor-donor ratios (0:1-3:1) in PBS buffer. Each well contained AF700 prepared at an equivalent  
389 concentration (30  $\mu$ g/mL) across all wells. The well-plate was imaged  $\sim$ 24 hours following preparation. Laser excitation  
390 was set to 700 nm and the emission filter used was 740  $\pm$  10 nm (Semrock, FF01-740/13-25). Laser power was set to 1.75  
391 mW for the SS2 and 0.8 mW for the ICCD.
- 392 - *TZM-AF700 and CTM-AF700 (donor only).* A well-plate was prepared using AF700 conjugated to either TZM or CTM  
393 (Fig. S7). The concentration was kept constant for both TZM-AF700 and CTM-AF700 (33  $\mu$ g/mL) across separate rows.  
394 Buffer pH was varied from 4.5 to 7.5 as explained previously for the IRDye 800CW-2DG well-plate experiment. Laser  
395 excitation was set to 700 nm and the emission filter used was a 715 nm longpass filter (Semrock, FF01-715/LP). Laser  
396 power was set to 1.2 mW.

## 397 Animal experiments

398 All animal procedures were conducted with the approval of the Institutional Animal Care and Use Committee at both  
399 Albany Medical College and Rensselaer Polytechnic Institute. Animal facilities of both institutions have been accredited by  
400 the American Association for Accreditation for Laboratory Animals Care International. HER2-overexpressing cell lines  
401 AU565 and SK-OV-3 used in this study were obtained from ATCC (Manassas, VA, USA) and cultured in RPMI and  
402 McCoy's media respectively supplemented with 10% fetal bovine serum (ATCC) and 50 Units/mL/50  $\mu$ g/mL  
403 penicillin/streptomycin (Thermo Fisher Scientific, Waltham, MA, USA). Tumor xenografts were generated by injecting  
404  $10 \times 10^6$  AU565 cells and  $4 \times 10^6$  SK-OV-3 in phosphate-buffered saline (PBS) mixed 1:1 with Cultrex BME (R&D Systems  
405 Inc, Minneapolis, MN, USA) on the opposite sides of inguinal mammary fat pads of female 5-week-old athymic nude mice  
406 (CrTac:NCr-Foxn1nu, Taconic Biosciences, Rensselaer, NY, USA). Tumors were monitored daily over a period of 3-4  
407 weeks.

408 Two xenograft-bearing mice were injected with 20  $\mu$ g AF700-TZM and 40  $\mu$ g AF750-TZM. Each mouse was imaged  
409 separately using either the SS2 or the ICCD at 24 h post-injection (Fig. 3, ~1hr between). Additionally, one of these mice  
410 was imaged at 48 h (SS2) and 51 h (gated-ICCD) post-injection (i.e., three hours after SS2 imaging, Supplementary Fig.  
411 S8&S9). Another two xenograft-bearing mice were injected with 20  $\mu$ g AF700-CTM and 40  $\mu$ g AF750-CTM. Each mouse  
412 was imaged separately using either the SS2 or the ICCD at 48 h post-injection (Fig. 4, ~1 hr between session). All injections  
413 were performed retro-orbitally on anesthetized mice. During mouse imaging, isoflurane anesthesia was performed, and the  
414 body temperature of each animal was maintained using a warming pad (Rodent Warmer X2, Stoelting, IL, USA) on the  
415 imaging plane.

## 416 Immunohistochemical analysis

417 Excised tumors were fixed in formalin, paraffin embedded, and processed for IHC. Epitope retrieval was performed by  
418 boiling deparaffinized and rehydrated 5  $\mu$ m sections in 1 mM EDTA pH 8.0 for 30 min. IHC staining was carried out using a  
419 standard protocol from Vectastain ABC Elite kit (Vector Labs, Burlingame, CA, P/N: PK-6101). Vector NovaRED (Vector  
420 Labs) was used as a peroxidase substrate. Tissue sections were counterstained with Methyl Green (Sigma, P/N: M8884).  
421 Hematoxylin Eosin stain was used for basic histology. Primary antibodies were as followed: rabbit monoclonal HER2 1:800  
422 (Cell Signaling, P/N: 2165), rabbit monoclonal EGFR 1:50 (Cell Signaling, P/N: D38B1), rabbit monoclonal TZM 1:100  
423 (R&D Systems, P/N: MAB95471-100). Brightfield images were acquired using Olympus BX40 microscope equipped with  
424 Infinity 3 camera (Lumenera Inc., Ottawa, ON, Canada).

## 425 Preprocessing of SS2 data

426 All SS2 data was processed using Alligator<sup>18</sup> software as described in Supplementary Note 4. Briefly, raw gate images  
427 (consisting in the sum of four 8-bit accumulation of individual 1-bit frames) were corrected for pile-up effects as described  
428 previously<sup>24</sup>. Next, subtraction of a detector background files acquired similarly as the data were used to correct for detector  
429 background (dark count noise) inhomogeneities and minimize the influence of “hot” SPADs/pixels characterized by  
430 intrinsically high dark count noise (Supplementary Fig. S12). Only one laser period (12.5 ns) worth of data was retained from  
431 each dataset (first 12.5 ns i.e. first 70 gates) except for one experiment (CTM *in vivo*). For the CTM *in vivo* experiment,  
432 significant photobleaching was observed during the last half of imaging. Hence, for this experiment's data processing, three  
433 12.5 ns sections (first 70 gates, 71-140 gates and 141-210 gates) of MFLI data were extracted from the sum of the first twenty  
434 acquisitions. These three data were summed together to provide 60 total accumulations. 2 $\times$ 2 binning was used for all SS2  
435 data herein. IRF data was processed in the same manner as fluorescence data (pile-up correction and background correction).  
436 SNR numbers reported in the text correspond to the square-root of the resulting pile-up and background-corrected pixel  
437 values.

## 438 Preprocessing of gated-ICCD data

439 All ICCD data was binned spatially (using either 2 $\times$ 2 or 4 $\times$ 4 binning). No further preprocessing was undertaken for  
440 phasor analysis. For NLSF analysis, because the decays were truncated to cover only 7 ns out of the 12.5 ns of a full laser  
441 period, the IRFs of individual pixels were extrapolated to cover the full-period (see Supplementary Note 4 for details). No  
442 further preprocessing of fluorescence decays was performed.

## 443 Nonlinear Least Square Fit of Fluorescence Decays

444 All full period (SS2) or partial (ICCD) decays  $\{G_i\}_{i=1,\dots,N}$  were fitted using the Levenberg-Marquardt non-linear least  
445 square fit (NLSF) algorithm implemented in Alligator<sup>18</sup>. Briefly, experimentally acquired pixel-wise IRF data (SS2) or  
446 extrapolated  $T$ -periodic IRF data (ICCD, see Supplementary Fig. S5),  $I_T(t)$ , was utilized for  $T$ -periodic convolution with a

447 single- or bi-exponential  $T$ -periodic decay model  $F_T(t)$ , including an optional IRF offset parameter  $t_0$  (set to 0 in all our  
448 analyses) and a baseline parameter  $B$  accounting for residual uncorrelated background (Eq. (1)) (Supplementary Note 4):

$$449 \quad \begin{cases} S_T(t) = B + I_{T|t_0} * F_T(t) \\ F_T(t) = \sum_{i=1}^n A_i \exp\left(-\frac{t[T]}{\tau_i}\right) \end{cases} \quad (1)$$

450 where the notations of ref.<sup>46</sup> are used. In particular,  $*$  denotes the cyclic convolution product,  $I_{T|t_0}(t)$  designates the  
451 experimental  $T$ -periodic IRF (with unknown offset  $t_0$ ) and  $x[T]$  denotes  $x$  modulo  $T$ . Index  $T$  in the function notation indicates  
452 a  $T$ -periodic function.  $A_i$  and  $\tau_i$  are the amplitude and lifetime of component  $i$ , where the number of components  $n$  is either 1  
453 or 2 depending on the experiment. Weighted fits were performed using the minimization function (Eq. (2)):

$$454 \quad \chi^2 = \frac{1}{dof} \sum_{p=1}^G \frac{(F_T(t_p) - G_p)^2}{|G_p|} \quad (2)$$

455 where  $dof$  is the number of free parameters of the fit,  $t_p$  is the  $p^{\text{th}}$  gate location in the period and  $G_p$  the gate value. If  $G_p = 0$ , a  
456 weight of 1 replaces the factor  $|G_p|$  in Eq. (2).

457 Note that in the *in vivo* case, in which the mouse itself was used for IRF acquisition, the corresponding recorded decay is not  
458 expected to be the true IRF, as the collected signal corresponds mostly to photons scattered off the mouse surface (or the  
459 most superficial layers of the skin). It does not account for propagation and scattering of the excitation light deep into tissues,  
460 nor does it account for propagation and scattering of the emitted fluorescence through the same layers of tissue<sup>53</sup>.  
461 Nevertheless, as discussed in Supplementary Note 5 and illustrated in Supplementary Fig. S13, which represents the local  
462 delay of the recorded scattered laser signal (relative to that measured with a white sheet of paper) as an equivalent  
463 topographic map (see below), the corresponding signal does differ significantly from a mere “paper IRF” and is therefore  
464 expected to be a better approximation of the true IRF than a simple paper IRF.

#### 465 FRET fraction analysis using decay fits

466 As discussed in ref.<sup>46</sup>, a periodic decay  $F_T(t)$  expressed according to Eq. (1), is the infinite summation of the non-periodic  
467 decay:

$$468 \quad F(t) = \sum_{i=1}^n A_i \left(1 - e^{-T/\tau_i}\right) \exp\left(-\frac{t}{\tau_i}\right), \quad t \geq 0 \quad (3)$$

469 whose component  $i$  contributes  $A_i \tau_i \left(1 - e^{-T/\tau_i}\right)$  to the total integral over  $[0, +\infty]$  and whose intensity fraction  $f_i$  is therefore:

$$470 \quad f_i = \frac{A_i \tau_i \left(1 - e^{-T/\tau_i}\right)}{\sum_{j=1}^n A_j \tau_j \left(1 - e^{-T/\tau_j}\right)} \quad (4)$$

471 This intensity fraction can be directly compared to that obtained by phasor ratio analysis (Eq. (8) below).

#### 472 Phasor Analysis

473 All phasor analysis was performed using Alligator<sup>18</sup> as detailed elsewhere<sup>23,34</sup>. In brief, for every pixel with coordinate  
474  $(x, y)$  within the MFLI region of interest, the uncalibrated *discrete* phasor  $z(x, y) = g(x, y) + i s(x, y)$ , where  $i$  the complex root  
475 of -1, was retrieved from the time-gated decay  $\{G_p(x, y)\}_{p=1, \dots, G}$  according to Eq. (5).

$$476 \quad z(x, y) = \sum_{p=1}^G G_p(x, y) e^{i 2\pi f t_p} \Big/ \sum_{p=1}^G G_p(x, y) \quad (5)$$

477 where  $f$  is the phasor frequency,  $G$  the number of gates,  $G_p$  the  $p^{\text{th}}$  gate value at pixel  $(x, y)$ . In particular, phasors computed  
478 for ICCD data, where truncated decays were recorded, correspond to the truncated (and offset) decay case discussed in ref.<sup>46</sup>.  
479 The harmonic frequency  $f$  was chosen equal to  $1/D \sim 142$  MHz – where  $D$  is the time span of the recorded decay. The  
480 resulting appearance of the phasor plot differs from the standard case, where the phasors of single-exponential decays are  
481 located on the so-called universal semicircle (or universal circle, UC)<sup>45</sup>. Instead, in the case of a truncated decay, the single-  
482 exponential decay’s phasor locus (or SEPL<sup>34</sup>) is in general a complex curve, which, after calibration, may or may not overlap  
483 partially with the UC (see next).

484 For SS2, one laser period-worth of data was used, therefore  $D = T$ , and  $f = 80$  MHz. However, since the number of gates ( $G =$   
 485 70) is finite, the SEPL is also expected to depart from the UC and be closer to an arc of circle with larger radius than the  
 486 UC<sup>45</sup>. In practice,  $G$  is sufficiently large for this difference to be negligible, and the UC was used in all representation of  
 487 calibrated phasors.

488 **Phasor calibration**

489 Visual depiction of how phasor calibration was undertaken is illustrated in the *Supplementary Material* (Supplementary  
 490 Fig. S14 and S15 for SS2 and ICCD, respectively) and follows ref.<sup>46</sup>. Briefly, the uncalibrated phasor of the IRF data  
 491 corresponding to the sample of interest (a sheet of white paper for the *in vitro* samples, and the mouse itself for *in vivo*  
 492 measurements [Supplementary Fig. S16]) was calculated using Eq. (5) on a pixel-by-pixel basis. These *calibration* phasors,  
 493  $z_{I_r}(x, y)$ , associated with lifetime  $\tau_{IRF} = 0$  were used to compute the calibrated phasor of sample data,  $\tilde{z}(x, y)$ , according to  
 494 Eq. (6):

$$495 \quad \tilde{z}(x, y) = \frac{z(x, y)}{z_{I_r}(x, y)} \quad (6)$$

496 As discussed above, due to their discrete nature and the nontrivial shape of the IRFs involved in their calculation, calibrated  
 497 phasors of single-exponential decays, normally expected to be located on the so-called universal circle (UC), are instead  
 498 located on a slightly different curve<sup>18</sup> (dubbed SEPL – or Single-Exponential Phasors Locus - in ref.<sup>34</sup>), which is expected to  
 499 depart from the UC for large enough lifetimes. In practice, for the number of gates used and the short lifetimes studied in this  
 500 work, the SEPL is indistinguishable from the UC, which was used instead.

501 **Phase lifetime calculation**

502 The phase lifetime,  $\tau_\phi$ , was calculated using the components ( $g, s$ ) of the calibrated phasor (Eq. (6)) according to Eq. (7),  
 503 which is only correct for an infinite number of gates  $G$ , but is a good approximation in the experimental situations described  
 504 in this work<sup>34</sup>:

$$505 \quad \tau_\phi = \frac{1}{2\pi f} \frac{s}{g} \quad (7)$$

506 **Phasor ratio calculation**

507 In the case of decays comprised of the contribution of two different species, such as encountered for the mixtures of  
 508 quenched donor and unquenched donor lifetimes in the FRET assays presented here, the phasor of the mixture is a linear  
 509 combination of the phasor of each species<sup>18,45,46</sup>. The intensity fraction of each species in the mixture can be graphically  
 510 recovered from the location of the phasor with respect to the pure species (reference) phasors. In the case studied here, one of  
 511 the reference phasor, the donor-only – or unquenched donor – phasor can be measured experimentally with the pure donor  
 512 sample, while the second reference can be inferred from the observed *linear arrangement* of the phasors of the different  
 513 acceptor to donor ratio samples (see Fig. 2): while there is no guarantee – and in fact it is unlikely – that the decay of a  
 514 maximally quenched donor is characterized by a strictly single-exponential decay, a natural reference to use for quantification  
 515 of the mixture is the intersection of the best-fit line on which all samples are aligned, with the SEPL (see Supplementary Fig.  
 516 S17 for a graphical illustration). Since there are two intersections, that with the shortest lifetime is chosen as the second  
 517 reference (the other intersection should correspond to the donor-only sample if it is characterized by a single-exponential  
 518 decay – this is approximately verified here, but not exactly, which is why the actual donor phasor is used as the first  
 519 reference). Because of shot noise and other sources of variance, each phasor is projected orthogonally onto the line  
 520 connecting both references to extract the so-called phasor ratio  $r$  expressing the relative distance of the phasor to the two  
 521 reference phasors (Eq. (8)):

$$522 \quad \begin{cases} r = \frac{g - g_2}{d} \cos \theta + \frac{s - s_2}{d} \sin \theta \\ d = \left( (g_1 - g_2)^2 + (s_1 - s_2)^2 \right)^{\frac{1}{2}} \\ \sin \theta = \frac{s_1 - s_2}{d}, \quad \cos \theta = \frac{g_1 - g_2}{d} \end{cases} \quad (8)$$

523 where  $(s_i, g_i)$ ,  $i = 1$  or  $2$ , correspond to the phasor of the two references and  $(g, s)$  is the mixture phasor. The phasor ratio  $r$   
 524 defined in Eq. (8) corresponds to the intensity fraction of species 1 in the mixture<sup>18</sup> and can thus be compared to the intensity  
 525 fraction extracted by NLSF of decays (Eq. (4)), assuming that each component species is characterized by a single-  
 526 exponential decay.

527 Statistical Analysis

528 Raw analysis data exported from AlliGator as text files were subsequently processed in MATLAB to obtain publication-  
529 quality plots and statistical representations. Boxplots used in Fig. 1 were created using MATLAB with all solid black lines  
530 marking the average value and each box width marking 1-standard deviation. Kernel density estimation distributions were  
531 calculated in MATLAB using the lower and upper bounds of the listed x-axis and a  $10^{-2}$  interval (e.g., [0:0.01:0.8] for Fig.  
532 4E) for all cases herein. All average and standard deviation results used for comparative scatter plots were calculated using  
533 MATLAB (dashed black line marks the diagonal in all cases). Linear regression used in Fig. 1Q&R (light blue dashed line)  
534 was performed in MATLAB. All average and standard deviation results were computed using MATLAB. Example script is  
535 provided for reproduction purposes<sup>49</sup>.

536 Fluorescent probes

537 IRDye 800CW 2-DG was purchased at Li-Cor (Lincoln, Nebraska, USA). Cetuximab (CTM) was purchased at  
538 MedChemExpress (Monmouth Junction, NJ, USA), trastuzumab (TZM) was obtained through Albany Medical Center  
539 pharmacy. Fluorescent labeling of TZM is described in depth elsewhere<sup>43</sup>. Fluorescent labeling of CTM was performed  
540 following protocol described elsewhere<sup>54</sup>. Mouse IgG AF700 and rabbit IgG AF750 were purchased at Thermo Fisher  
541 Scientific (Waltham, MA, USA).

542 V. Associated content

543 The supplementary information is available free of charge at <https://XXXXXXX>

544 Software and Data Availability: All analyses are performed using freely available software<sup>18</sup>, and data as well as analysis  
545 details and results are available on a public cloud repository in order to ensure reproducibility<sup>49</sup>.

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552 Competing interests: Edoardo Charbon holds the position of Chief Scientific Officer of Fastree3D, a company making  
553 LiDARs for the automotive market, and Claudio Bruschini and Edoardo Charbon are co-founders of Pi Imaging Technology.  
554 Neither company has been involved with the work or manuscript. The authors declare no other competing financial interests.

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561 VIII. References

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