

HDYDI

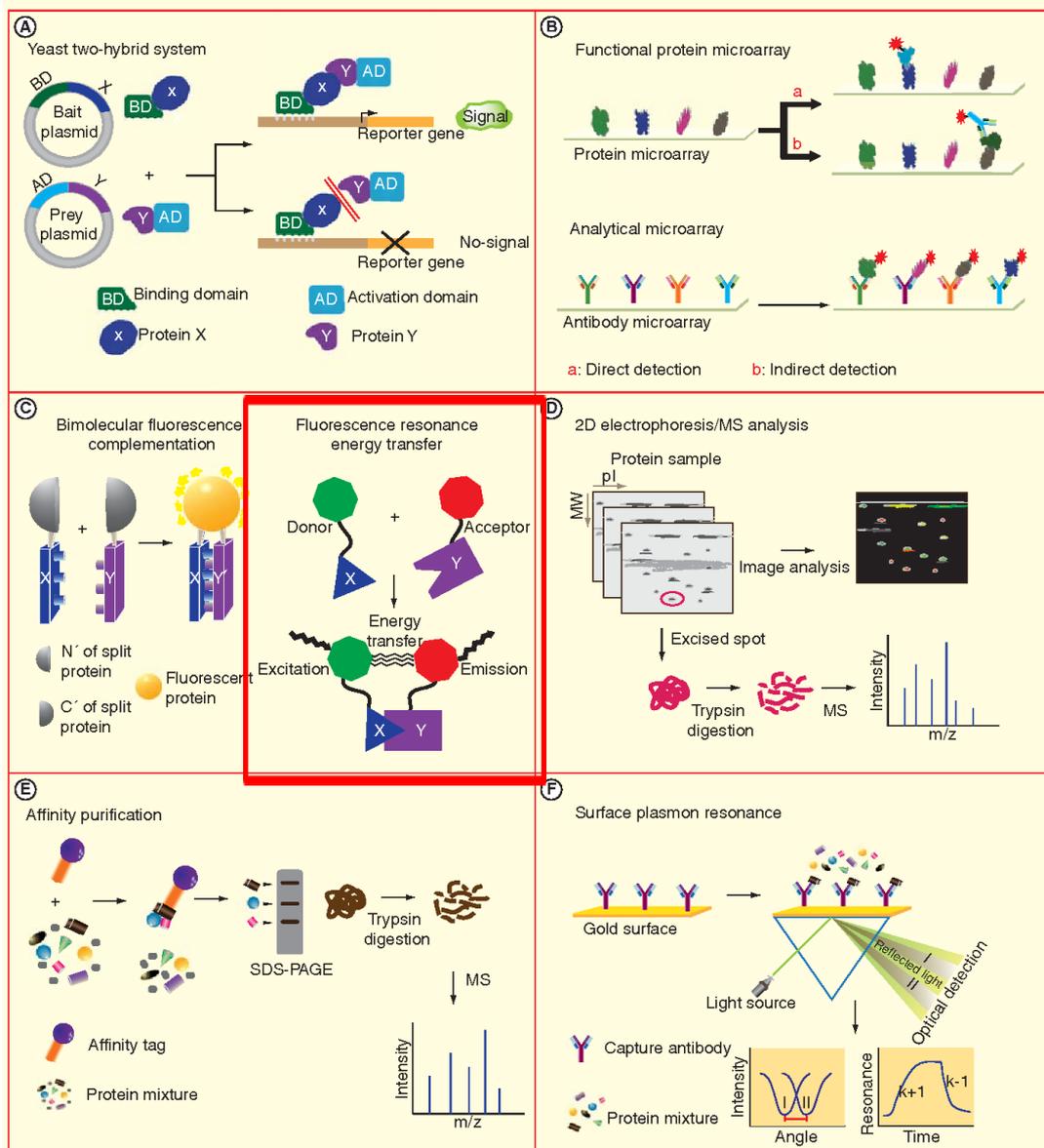
Visualization of Protein Interactions in Living Cells by Homo- and Hetero-FRET

**Paula Dlugosz
Johannes Nimpf Lab**

A joint venture of

Part of

Methods to investigate protein-protein interactions



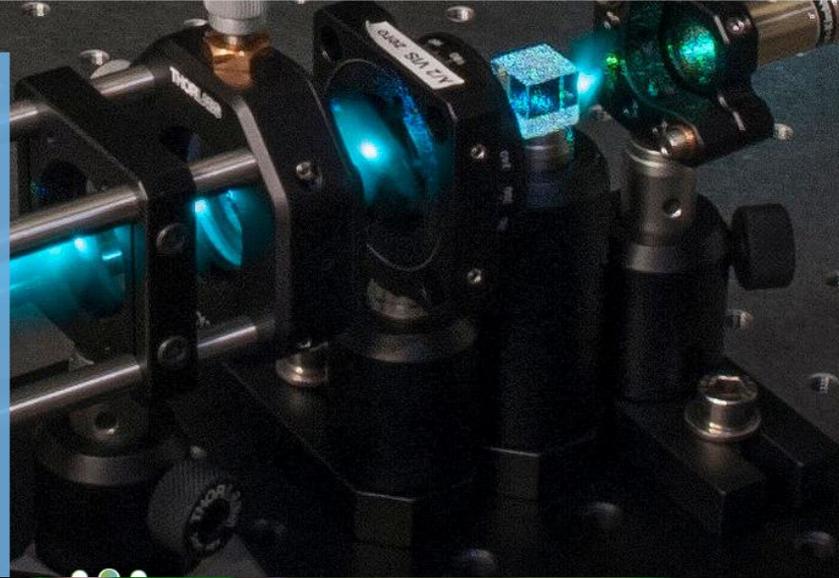
- Bimolecular fluorescence complementation (BiFC)
- Fluorescence cross-correlation spectroscopy (FCCS)
- Number&Brightness (N&B)
- MicroScale Thermophoresis (MST)
- Isothermal Titration Calorimetry (ITC)
- Proximity Ligation Assay (PLA)

Direct vs. Indirect
 Live vs. fixed/lysed

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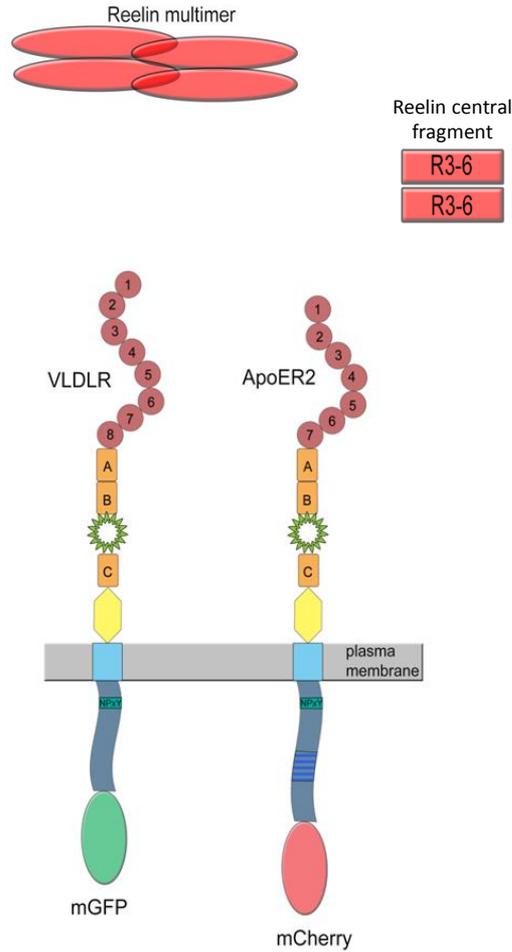
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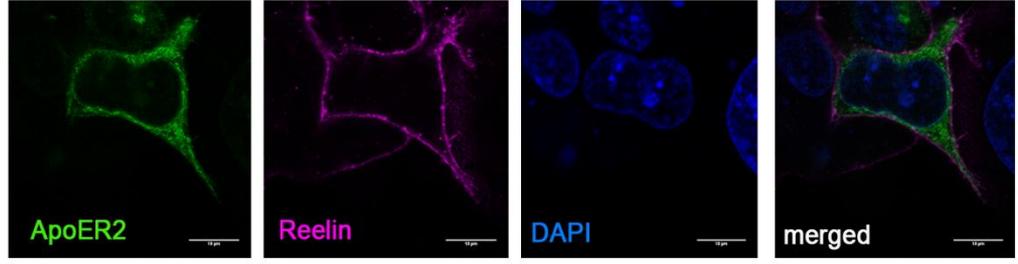
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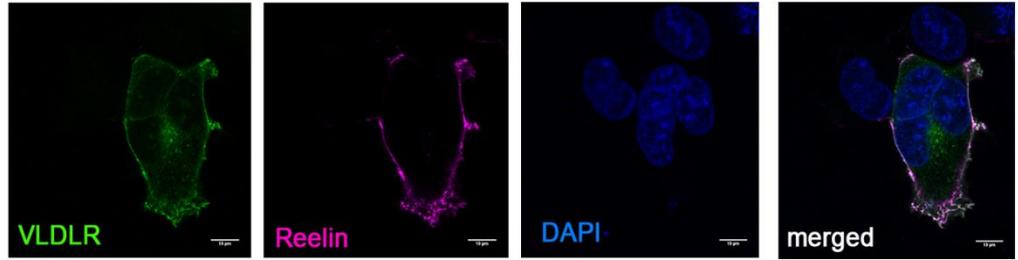
Localization and functionality of fluorescently tagged proteins of interest



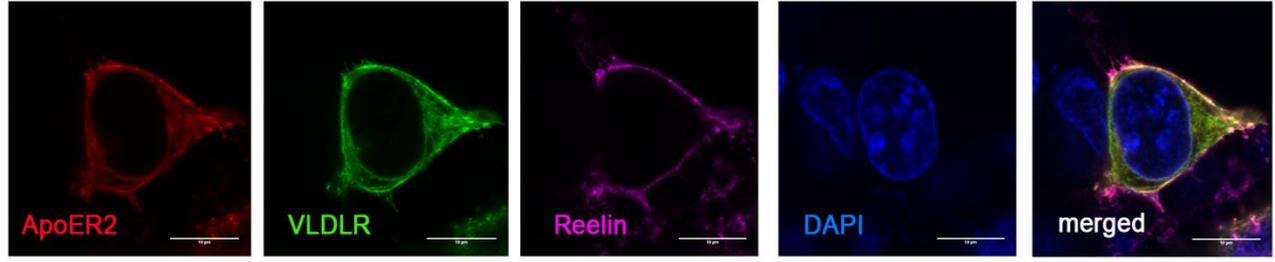
D ApoER2_mGFP+Reelin



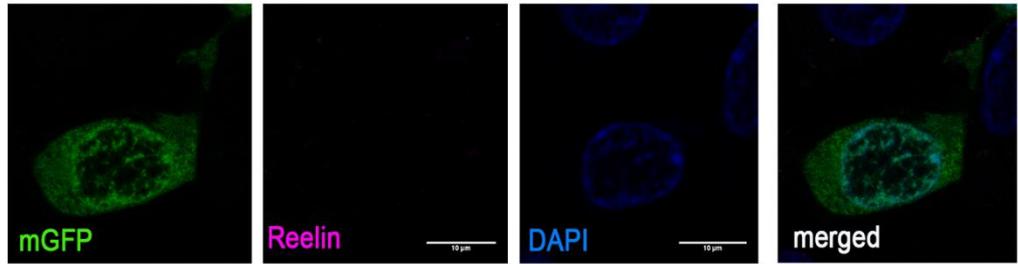
E VLDLR_mGFP+Reelin



F ApoER2_mCherry+VLDLR_mGFP+Reelin



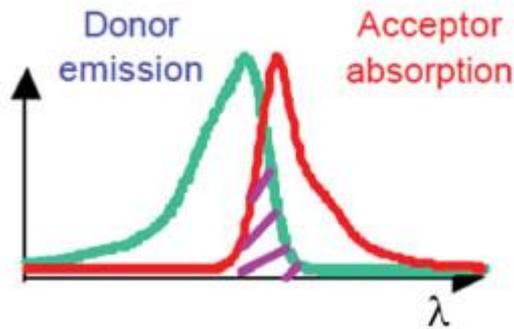
G mGFP+Reelin



HEK293 cells

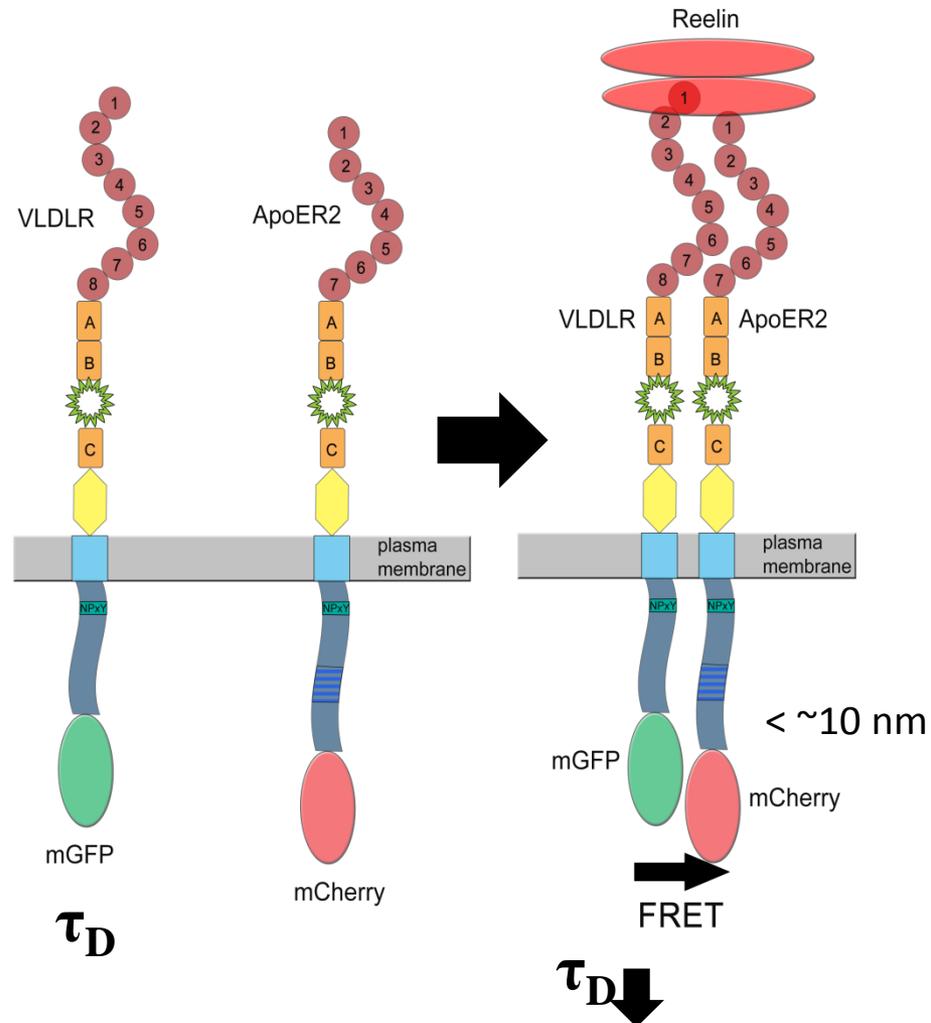
Hetero-FRET (Förster resonance energy transfer between a pair of different species of fluorophores) by FLIM (Fluorescence Lifetime Imaging Microscopy)

1) Spectral overlap

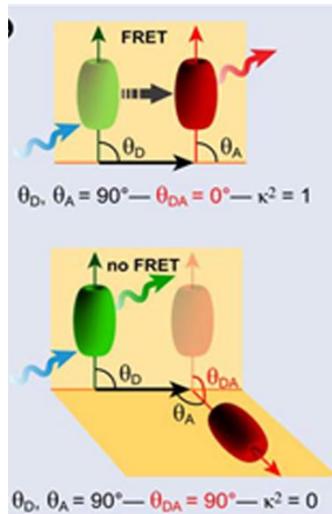


(Padilla-Parra & Tramier, 2012)

2) Proximity

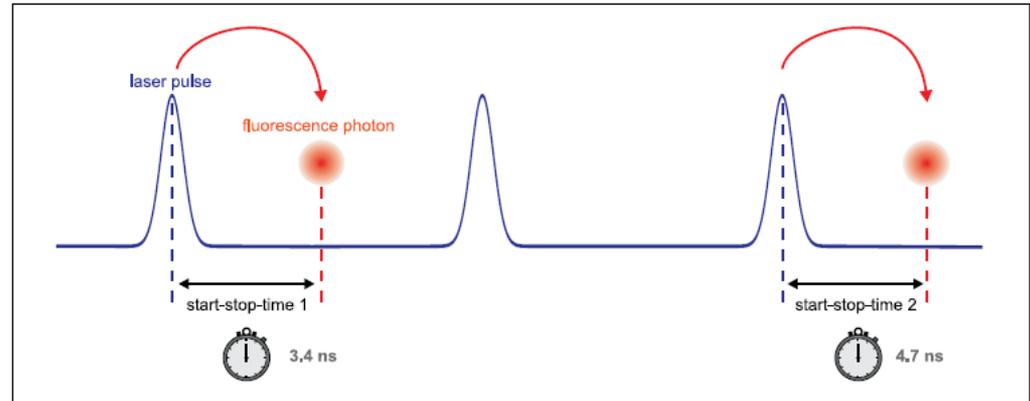
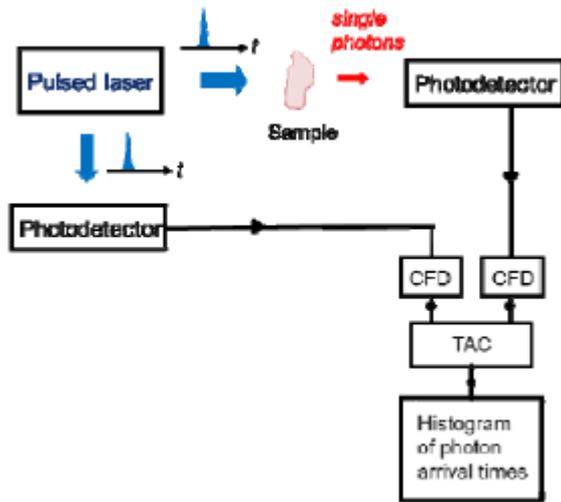


3) Orientation of fluorophores

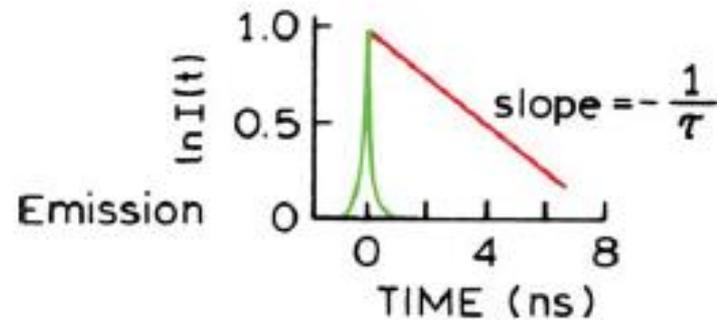
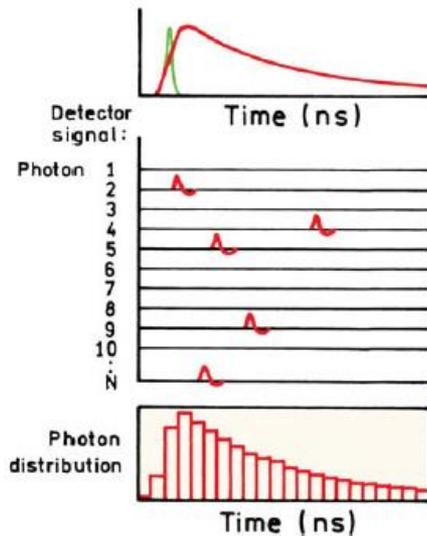


(Preus & Wilhelmsson, 2012)

Time-Correlated Single Photon Counting

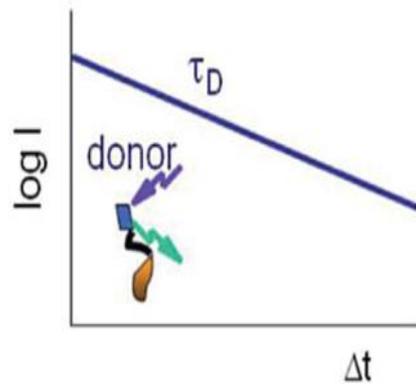


(Picoquant, Germany)

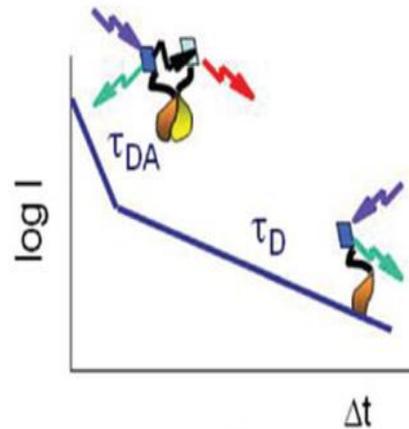


$$I(t) = I_0 \exp(-t/\tau)$$

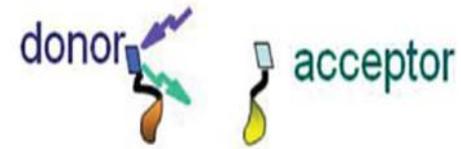
(Principles of Fluorescence Spectroscopy, Lakowicz)



$$I(t) = \alpha e^{-t/\tau_D}$$



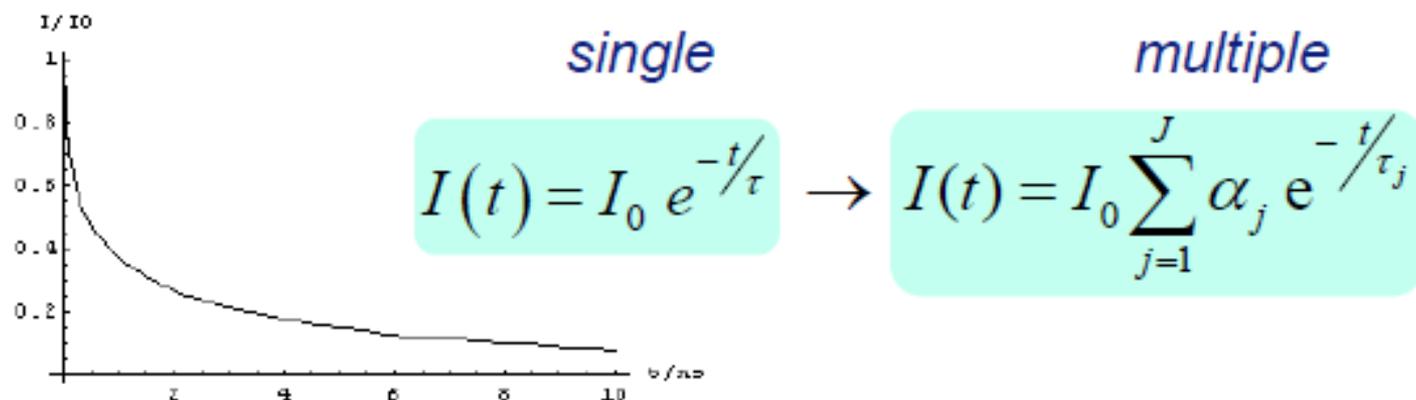
$$I(t) = \alpha_D e^{-t/\tau_D} + \alpha_{DA} e^{-t/\tau_{DA}}$$



$$\Rightarrow \text{FRET efficiency} = 1 - \frac{\tau_{DA}}{\tau_D}$$

$$\Rightarrow \text{Proportion bound donor} = \frac{\alpha_{DA}}{\alpha_{DA} + \alpha_D}$$

Analysis of fluorescence decay profiles



Fitting complex decay profiles requires more detected photons

- achieving ~10% accuracy for single exponential fit requires 100's photons
- to fit double exponential decay model requires 1000's photons

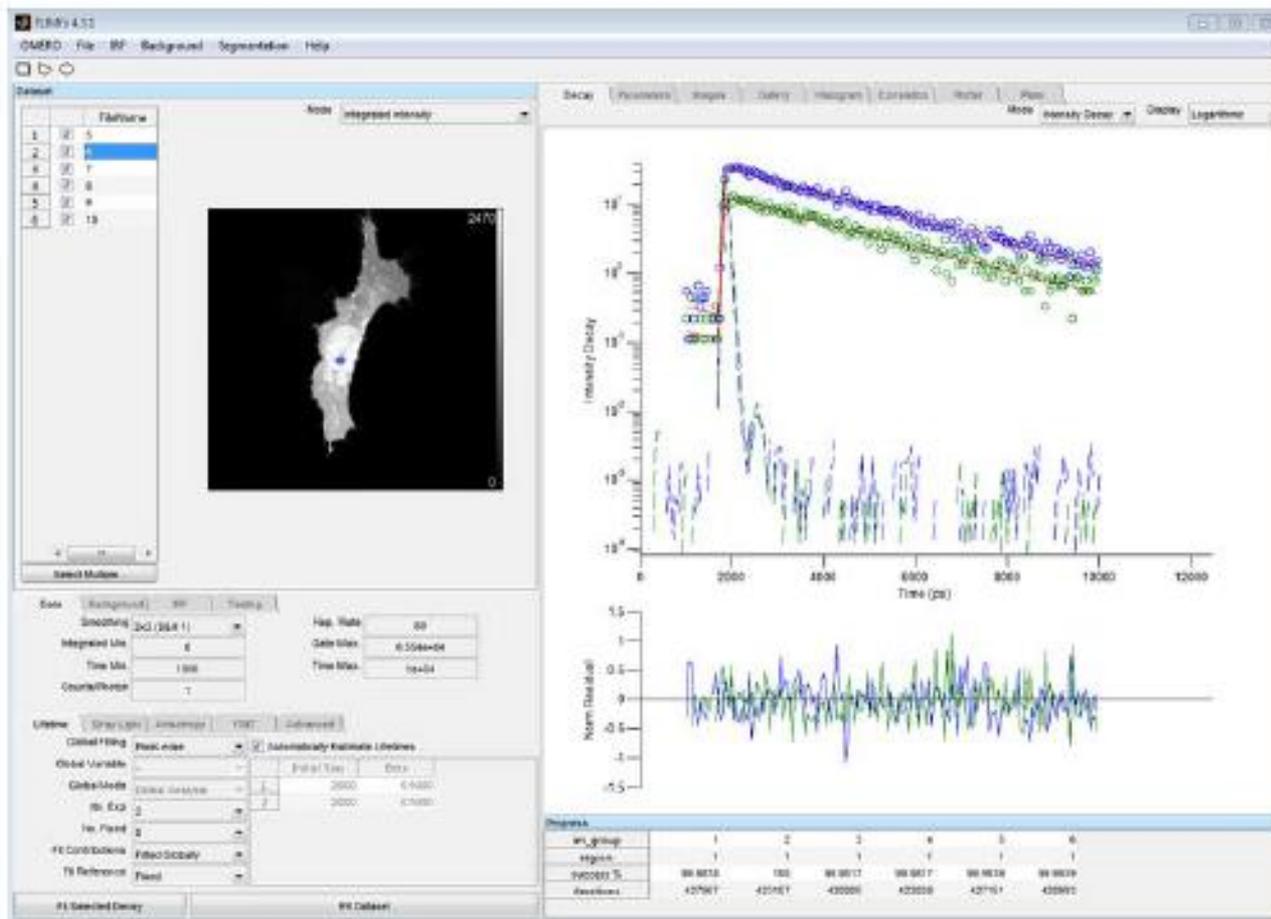
BUT # photons/pixel is constrained in many biological experiments by photobleaching/toxicity, time resolution ...

⇒ fit to single exponential decay model anyway

⇒ segment ROI, bin pixels and fit to (complex) model

⇒ global fitting to complex model

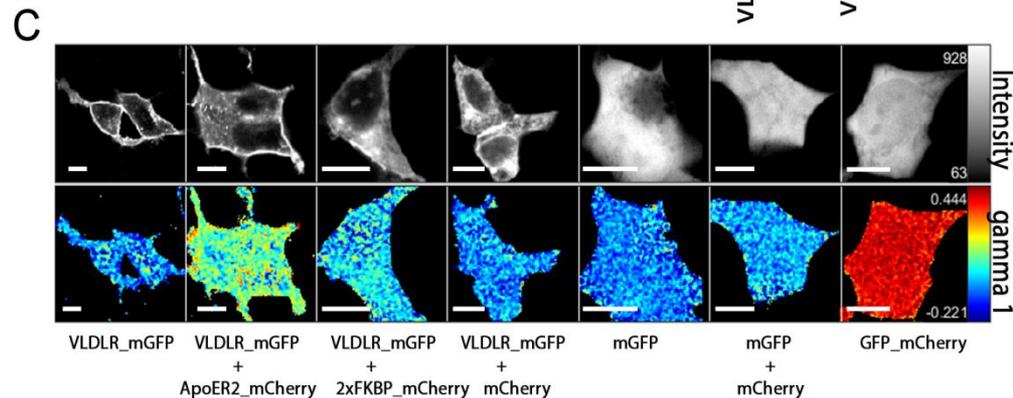
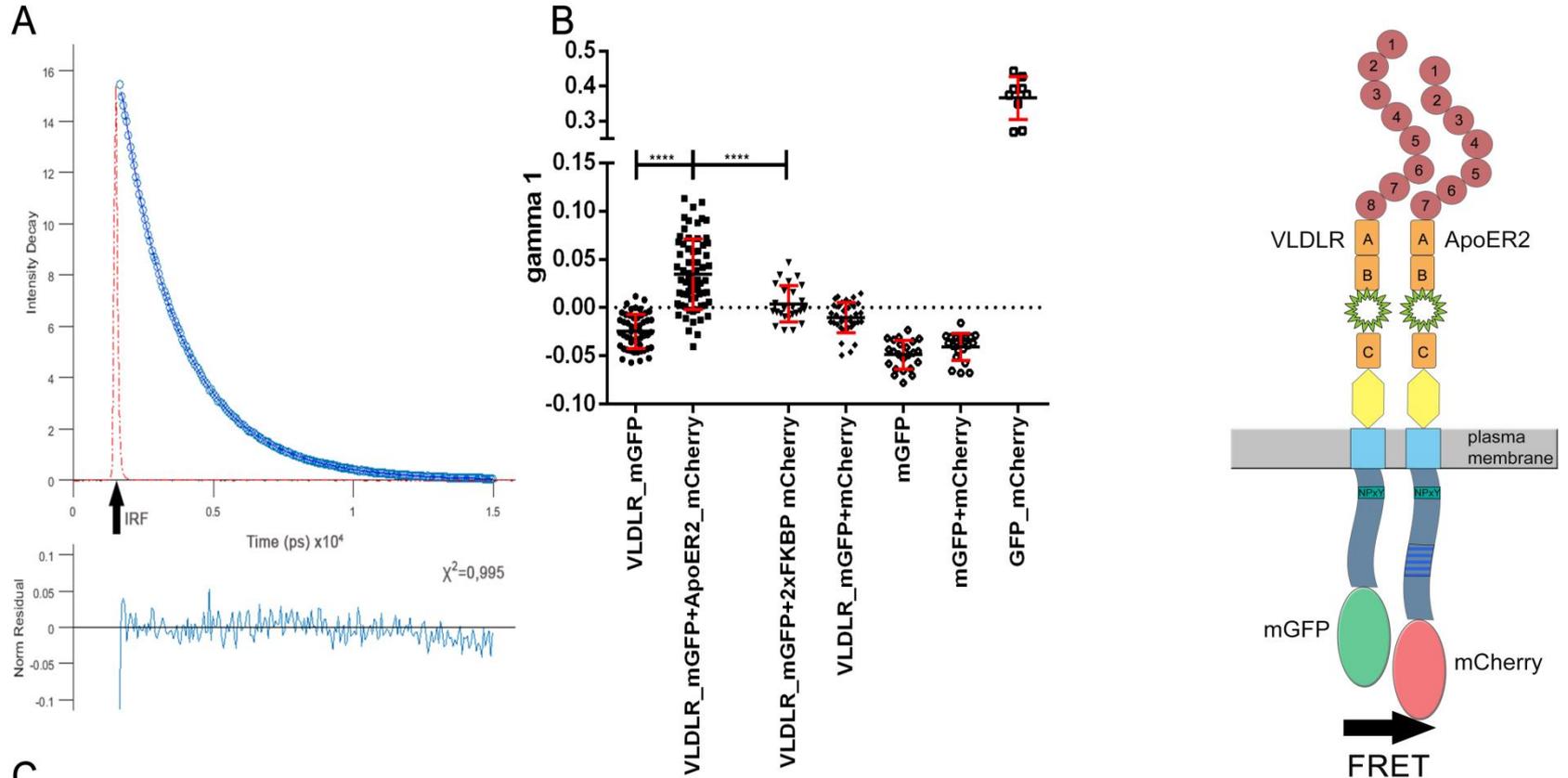
⇒ no fitting (phasor analysis)



Open-source software available via Open Microscopy Environment

<http://downloads.openmicroscopy.org/latest/flimfit>

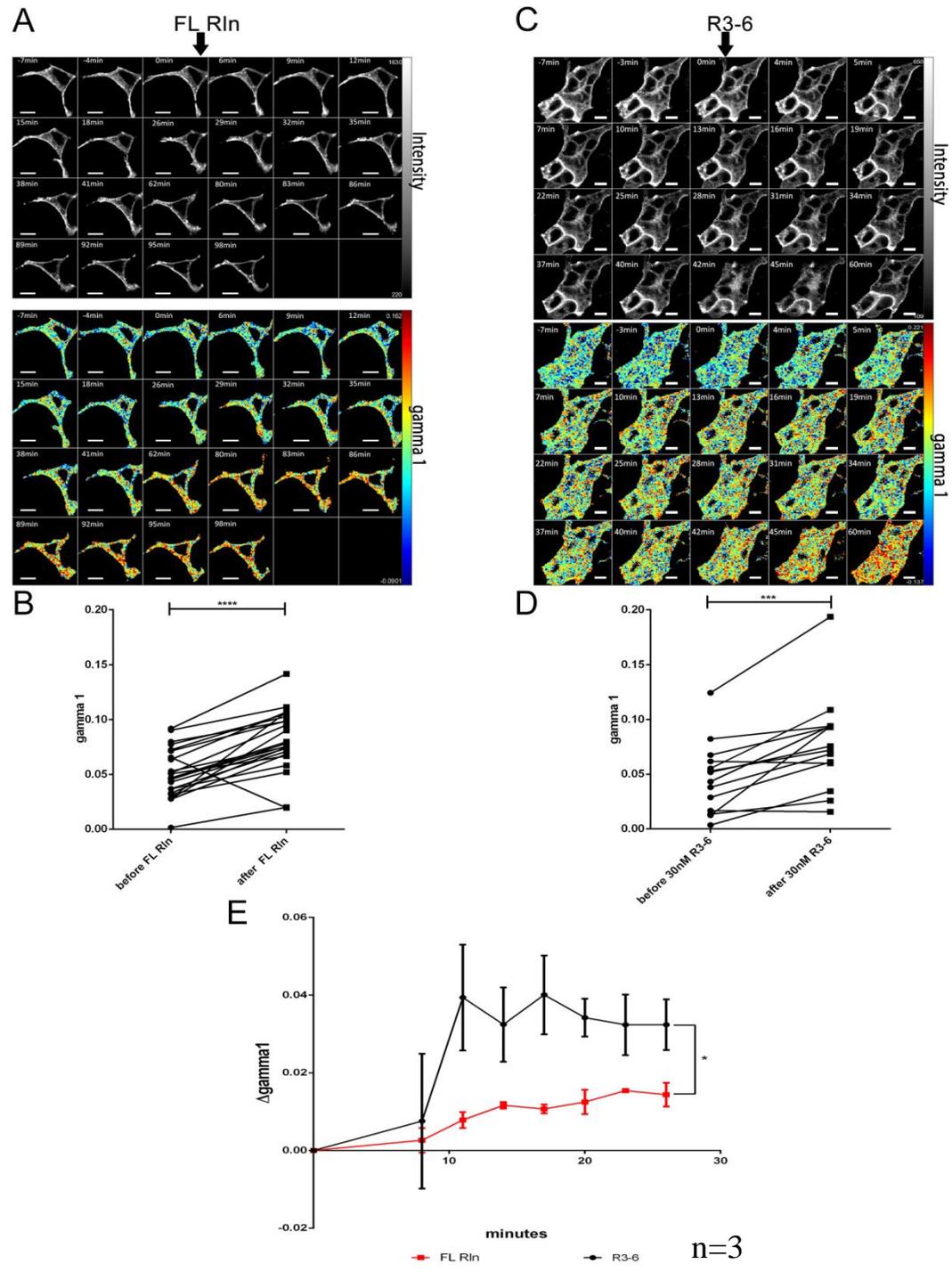
ApoER2 and VLDLR form hetero-oligomers as assessed by fluorescence-lifetime imaging microscopy



Gamma 1 =
contribution of FRET
with globally fitted
efficiency (E) = 0.5

IRF= instrument response function

Differential effect of full length Reelin and R3-6 on the increase of cluster size of VLDLR/ApoER2 hetero-oligomers

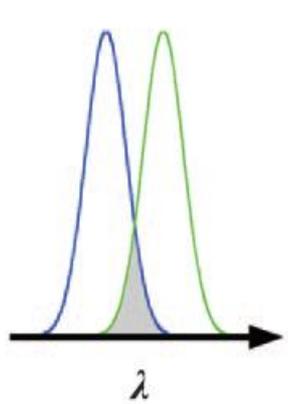


Homo-FRET Fluorescence Anisotropy Imaging as a Tool to study Molecular Self-Assembly in Live Cells

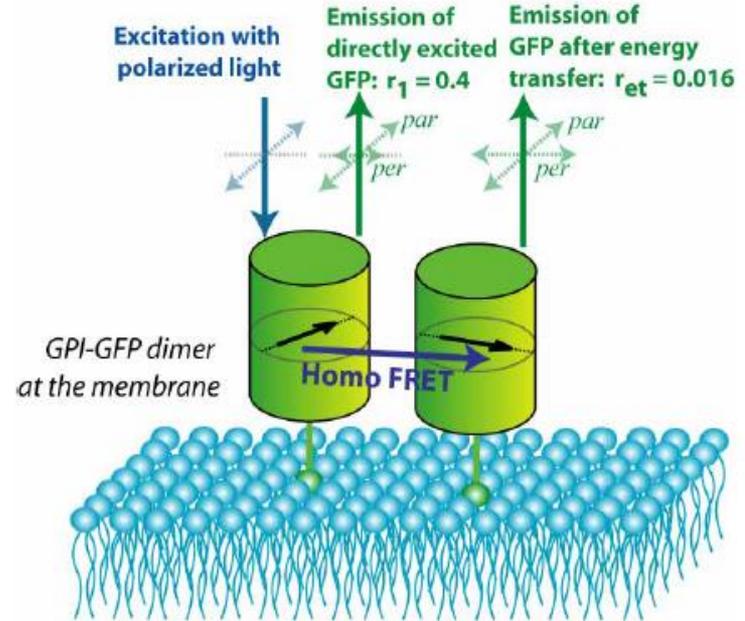
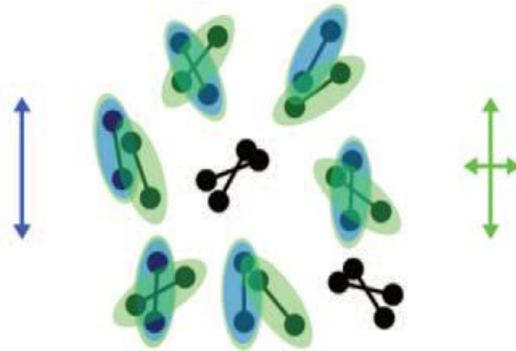
Using only one fluorophore brings advantages over hetero-FRET

- 1) all the fluorescent molecules are analyzed by this method comparing to hetero-FRET where dimers composed of donor-donor or acceptor-acceptor cannot be identified;
- 2) easier fluorescent labelling as only one fluorescent probe is required;
- 3) problem with differential expression levels of donor and acceptor-tagged proteins does not have to be taken into consideration;
- 4) possibility to perform multiplexed imaging which allows to follow spatio-temporal correlation of signaling pathways.

Homo-FRET Fluorescence Anisotropy Imaging as a Tool to study Molecular Self-Assembly in Live Cells



GFP excitation and emission spectra

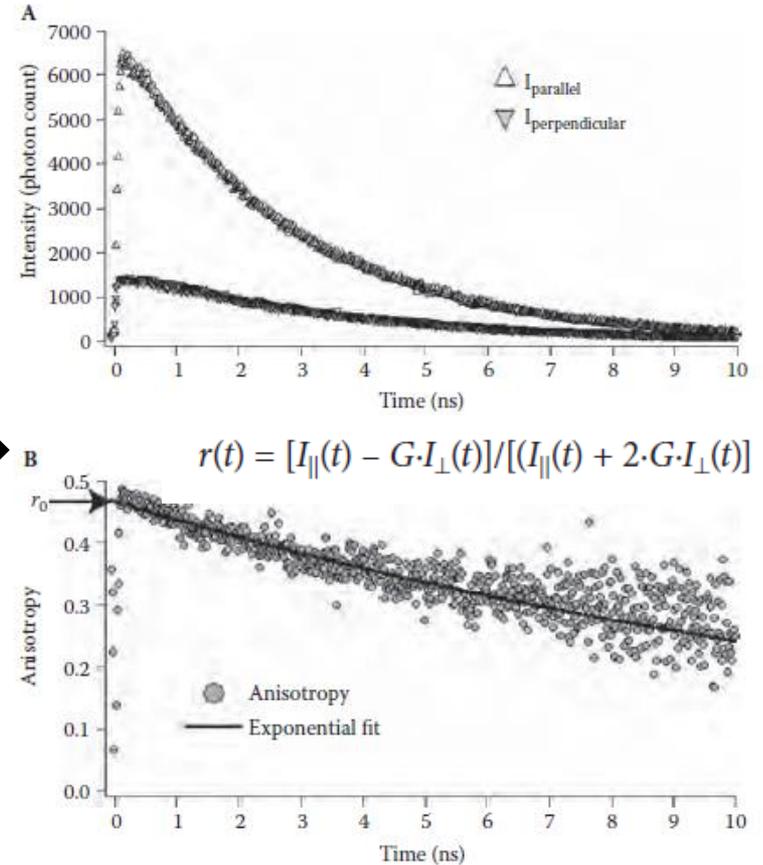
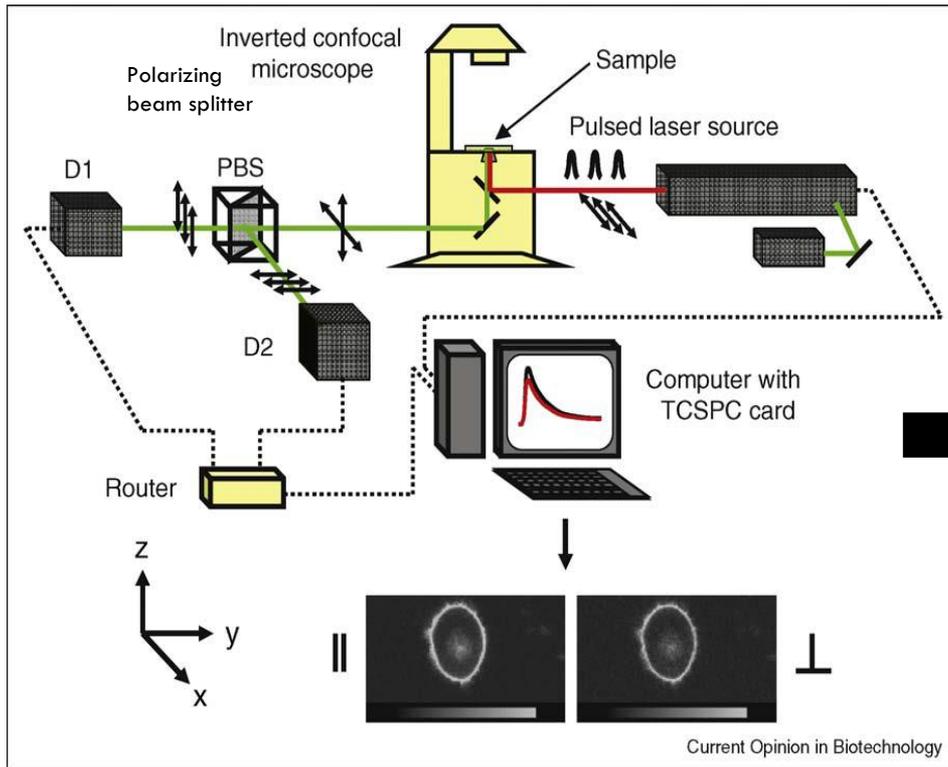


(Bader, Hofman et al., 2007)
(Chan, Kaminski et al., 2011)

HomoFRET depolarisation increases with the number of participating fluorophores.

This effect presents an opportunity for cluster size quantification by measuring the degree of depolarisation.

Time Resolved Fluorescence Anisotropy

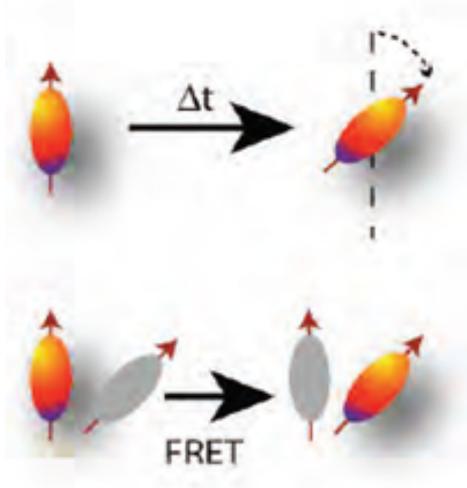


(Chan, Kaminski, and Kaminski Schierle, 2011)
(Vogel, Thaler et al. 2009)

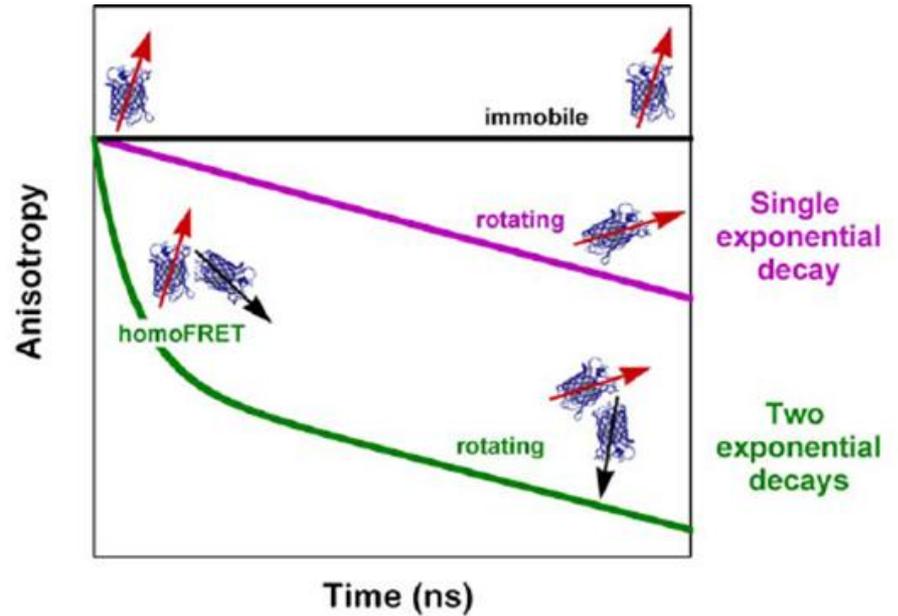
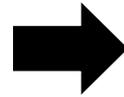
Depolarization of emitted light can arise from

rotation

energy transfer



(Vogel, Thaler et al., 2009)

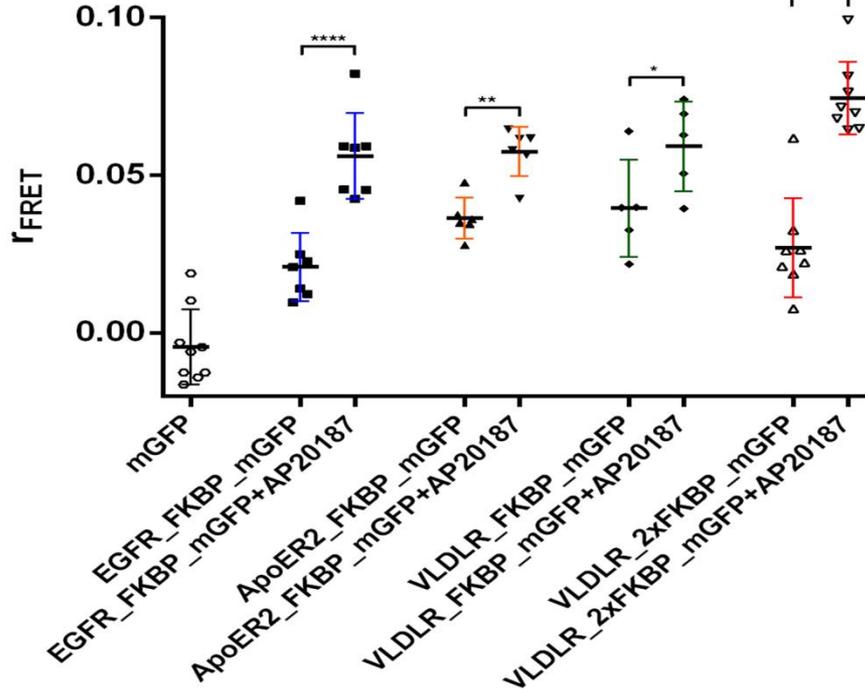


(Altman, Goswami et al., 2007).

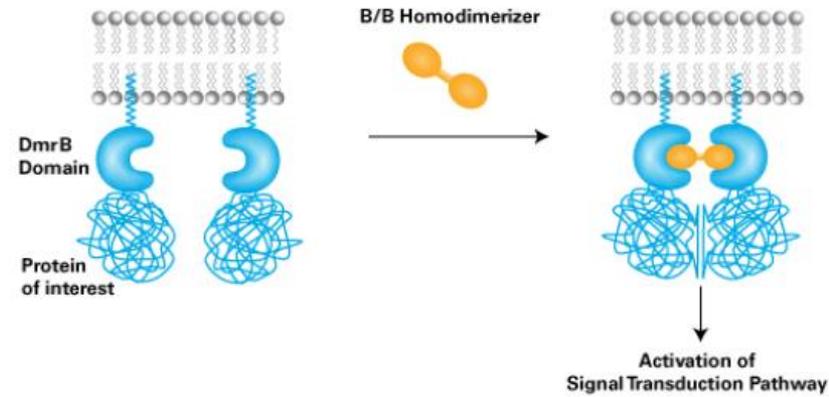
Depolarization due to FRET by forced oligomerization of receptor molecules

Figure S2

A



Homodimerization



B

HEK293	ApoER2_FKBP_mGFP		EGFR_FKBP_mGFP		VLDLR_FKBP_mGFP	
Parameter	before Homodim.	after Homodim.	before Homodim.	after Homodim.	before Homodim.	after Homodim.
tau_1 [ns]			2,453			
tau_2 [ns]			1,309			
beta_1			0,685			
beta_2			0,315			
theta_1 [ns]			102,600			
theta_2 [ns]			1,944			
r_2	0,036	0,0574	0,0209	0,0561	0,0395	0,0591
Number of cells	n=6	n=6	n=7	n=7	n=5	n=5

C

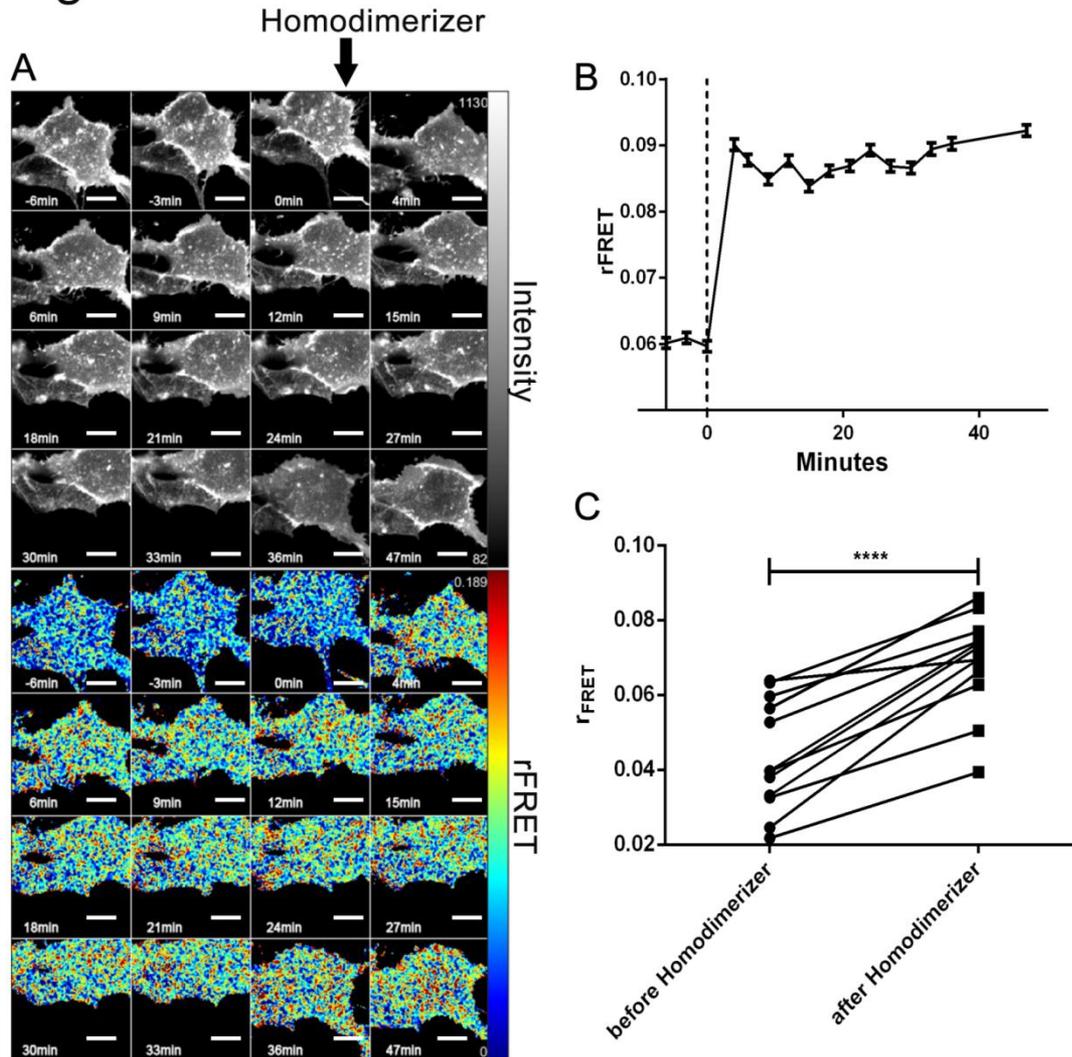
Parameter	mGFP
tau_1 [ns]	2,6708
tau_2 [ns]	1,552
beta_1	0,527
beta_2	0,473
theta_1 [ns]	18,825
theta_2 [ns]	2,503
r_2	-0,0044
Number of cells	n=9

From Clontech company

DmrB domain=FKBP domain
B/B Homodimerizer= AP20187

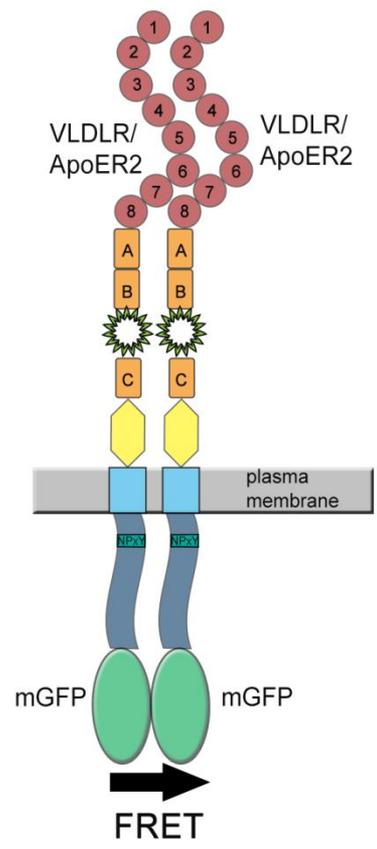
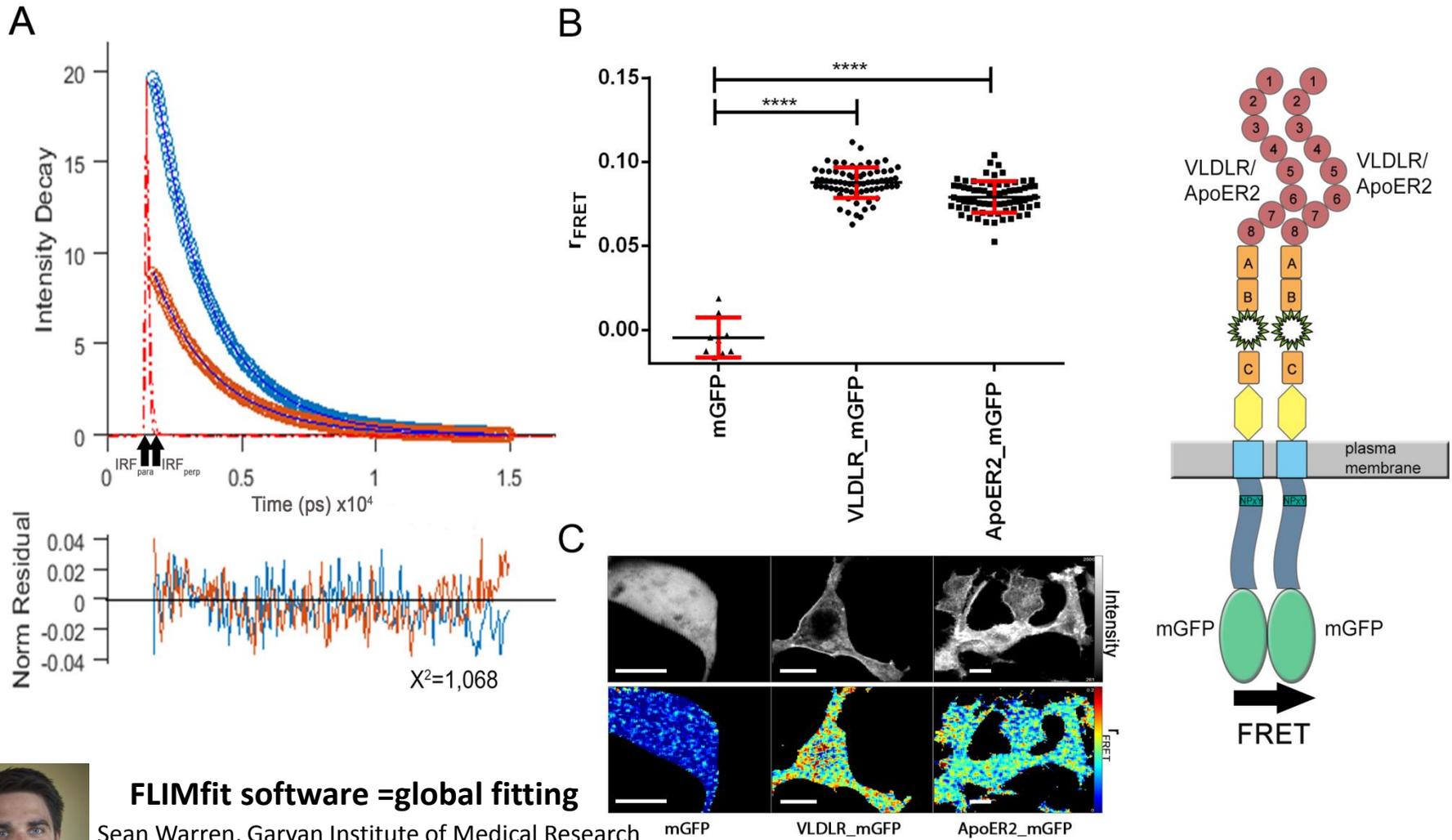
Depolarization due to FRET by forced oligomerization of receptor molecules

Figure S3



VLDLR_FKBP_mGFP was expressed in HEK293 cells and anisotropy associated with homo-FRET was measured

ApoER2 and VLDLR form homo-oligomers as assessed by time-resolved fluorescence anisotropy imaging



FLIMfit software =global fitting

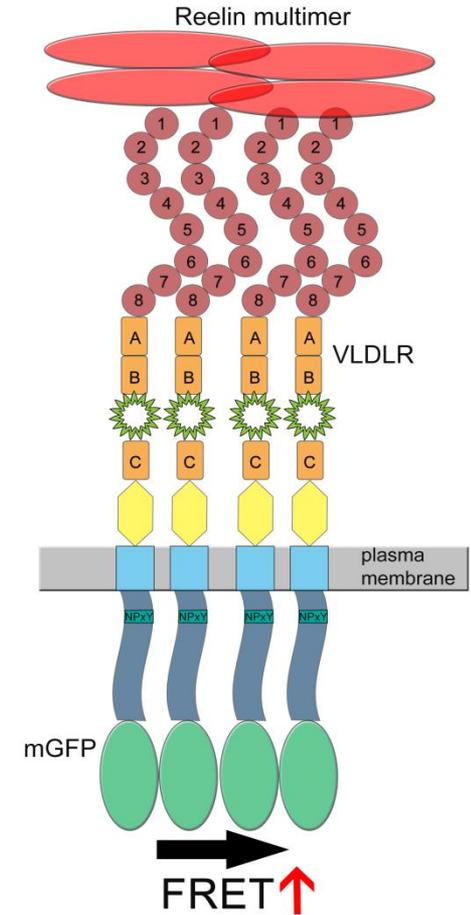
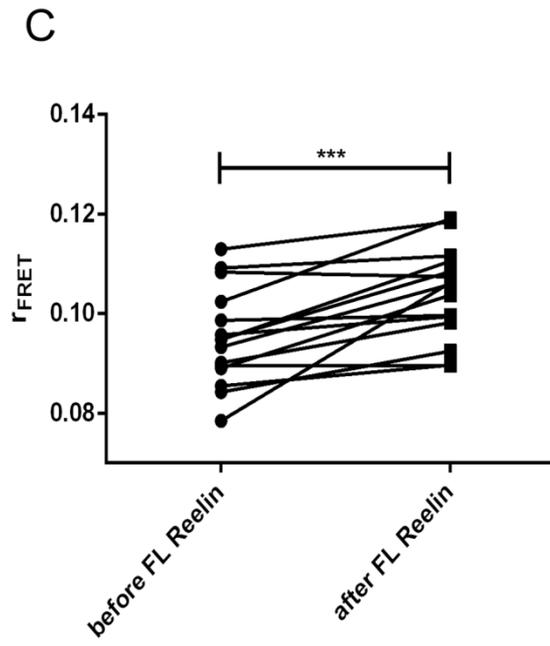
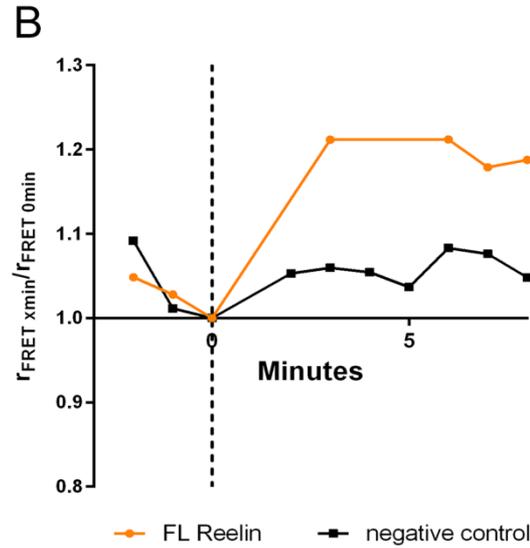
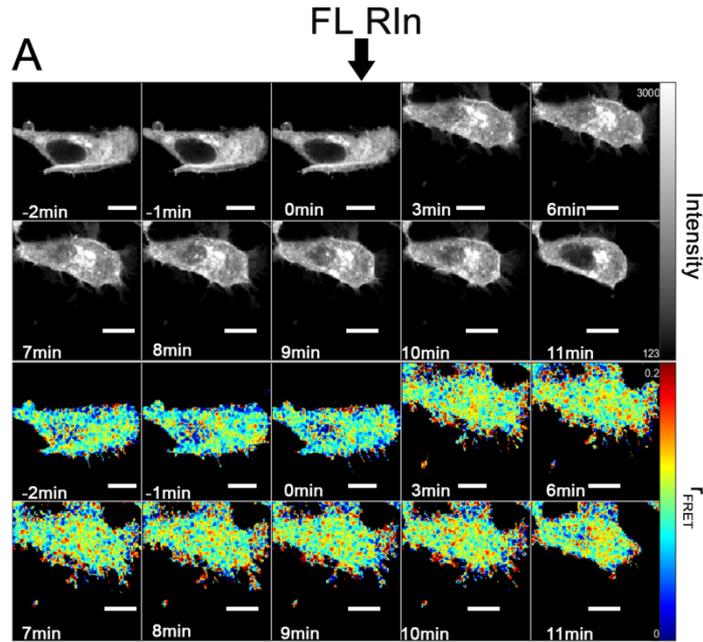
Sean Warren, Garvan Institute of Medical Research

“Homo-FRET Based Biosensors and Their Application to Multiplexed Imaging of Signalling Events in Live Cells” , *Int. J. Mol. Sci.*

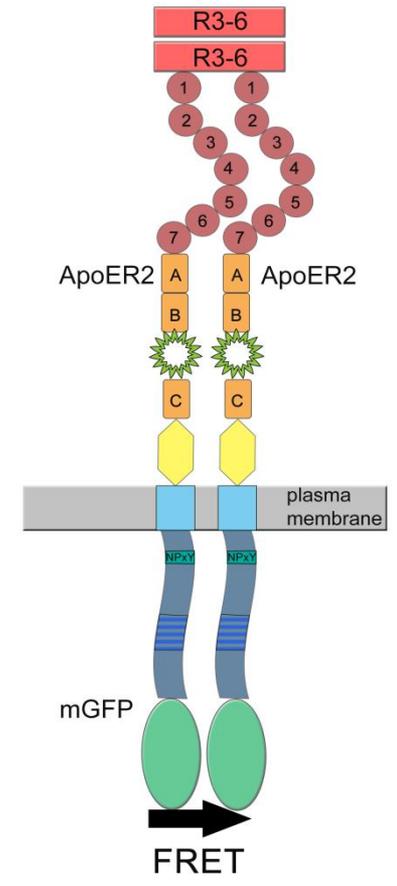
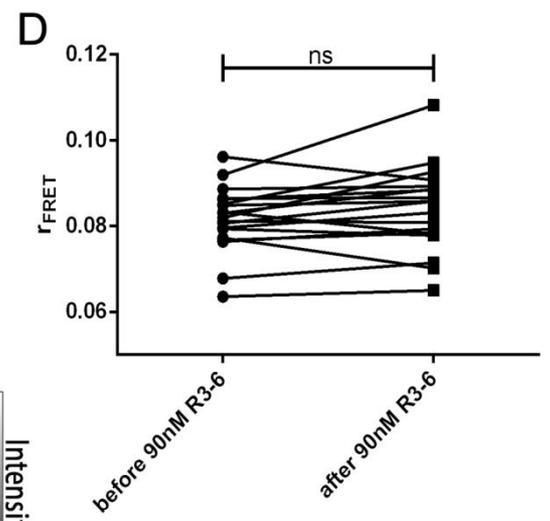
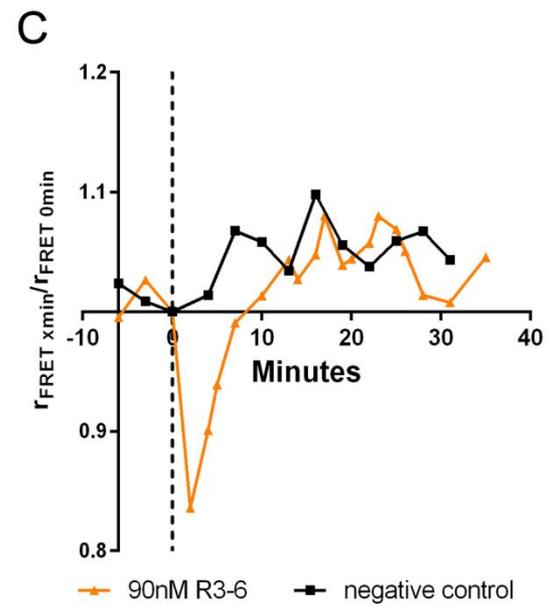
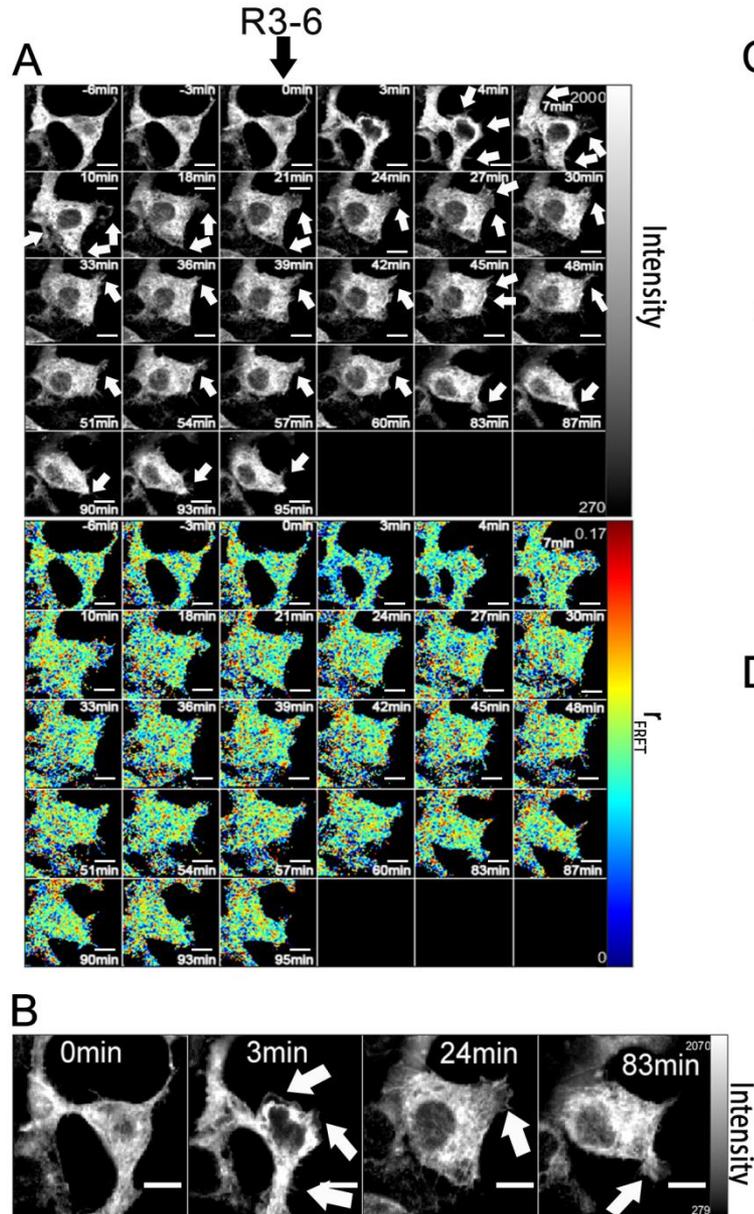
2015



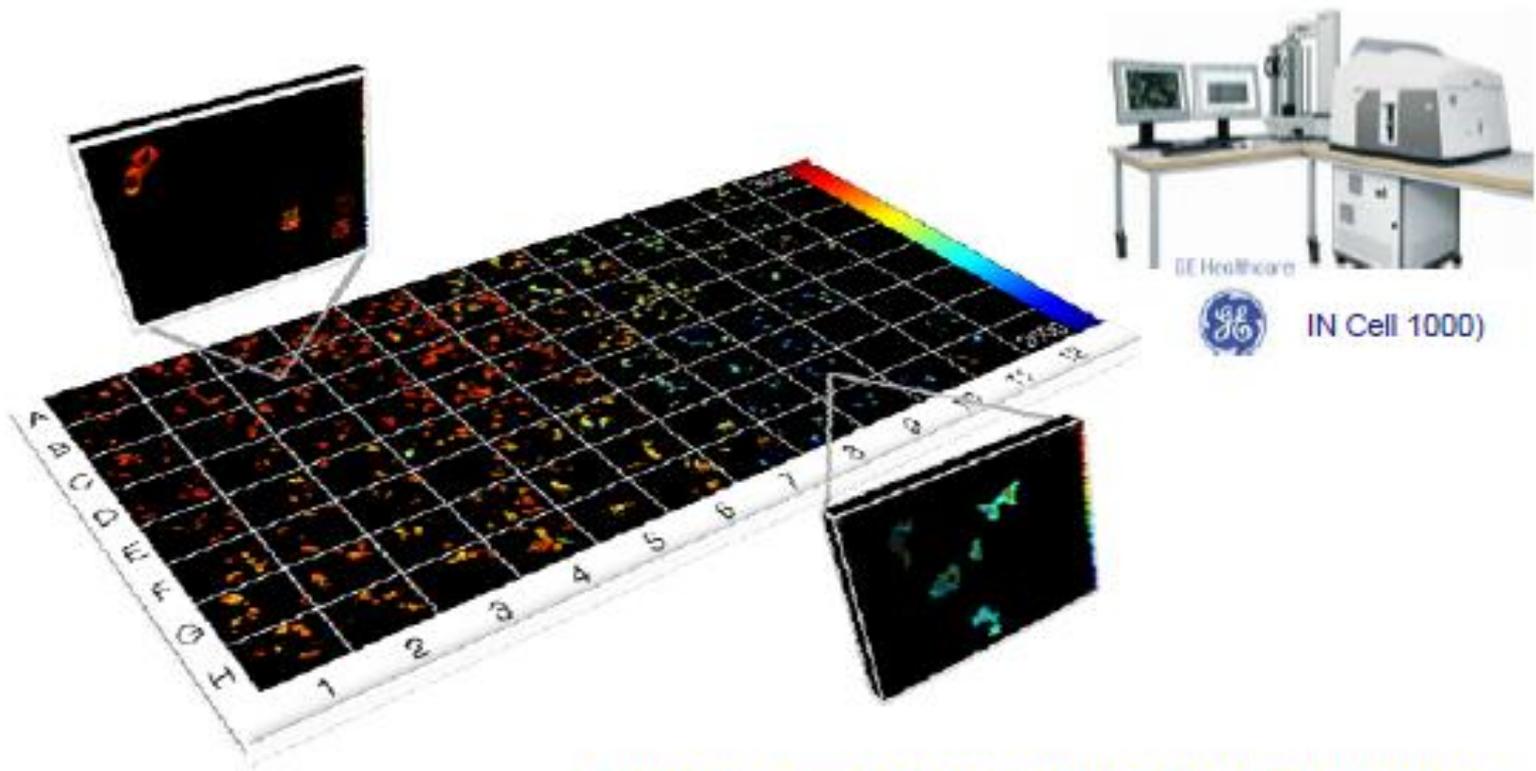
Full length Reelin increases cluster size of VLDLR homo-oligomers



Central fragment of Reelin (R3–6) does not increase cluster size of ApoER2 homo-oligomers



HCA: automated FLIM multiwell plate reader



Screening protein interactions for drug discovery

Technology Strategy Board
Driving Innovation

Healthcare

AstraZeneca

GE Healthcare

gsk
GlaxoSmithKline

Imperial College
London

Kentech Instruments Ltd

From Paul French, Imperial Collage London

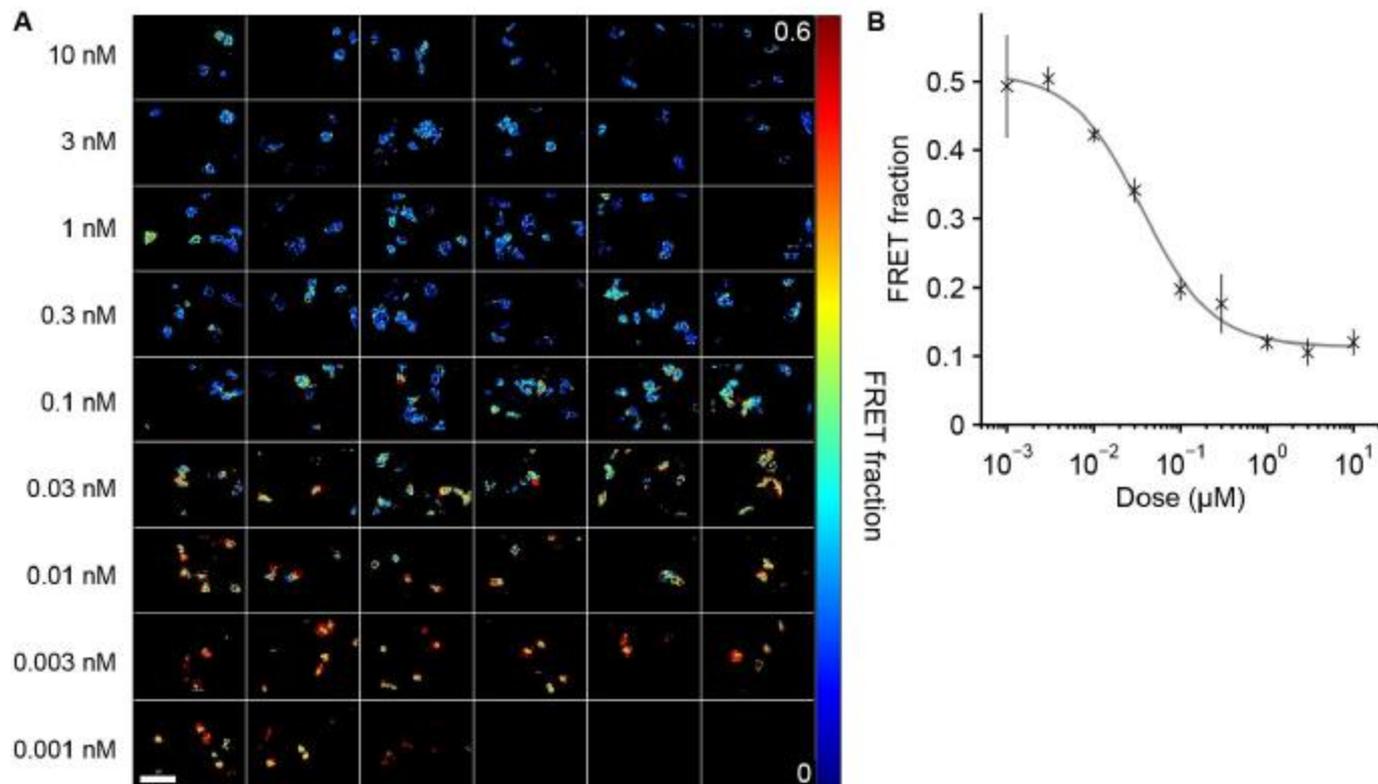


Figure 4. Global analysis of an NMT inhibitor dose-response dataset modulating Gag aggregation. Global analysis was performed across a multiwell plate dataset with HeLa cells expressing ECFP-Gag and EYFP-Gag with increasing levels of a NMT inhibitor using a bi-exponential donor FRET model. A) representative images from each inhibitor dose showing distribution of fraction Gag-CFP undergoing FRET. B) plot of fraction of Gag-CFP undergoing FRET against inhibitor concentration, averaged across wells with fitted dose-response curve. Error bars indicate 95% confidence intervals across wells. This dataset was collected as part of a previous study [23]. White scale bar represents 100 μm. doi:10.1371/journal.pone.0070687.g004

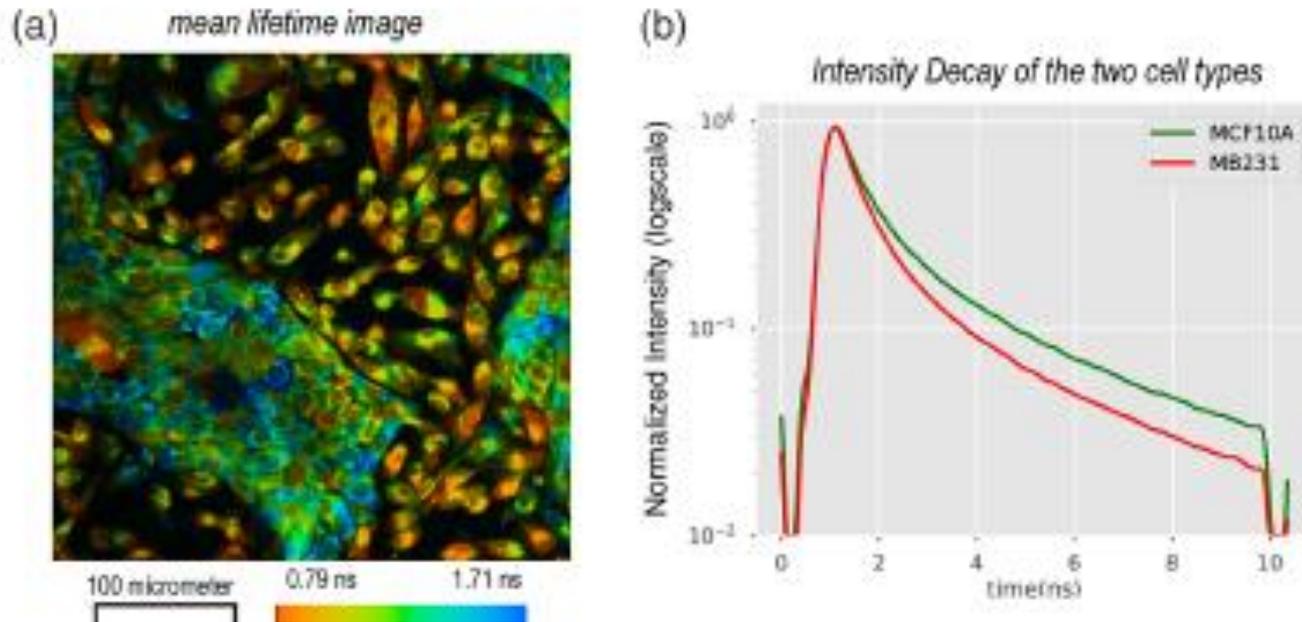
Rapid global fitting of large fluorescence lifetime imaging microscopy datasets. Warren SC et al., 2013

„We fitted a dose-response curve to the average FRET population across wells using nonlinear fitting to the Hill equation, giving an

EC50 value of 0.037 μM.

Label-free FLIM

Fluorescence lifetime imaging microscopy (FLIM) has emerged as one of the more powerful autofluorescence imaging tools for NADH and FAD characterization due to its unique ability to noninvasively detect metabolite bound and free states and quantitate cellular redox ratio.



Co-culture of MCF10A and MD-MB231

MCF 10A cell line is a non-tumorigenic epithelial cell line.

MB231 cancerous cell line

From Chacko JV and Eliceiri KW, 2019

Thank you for attention!

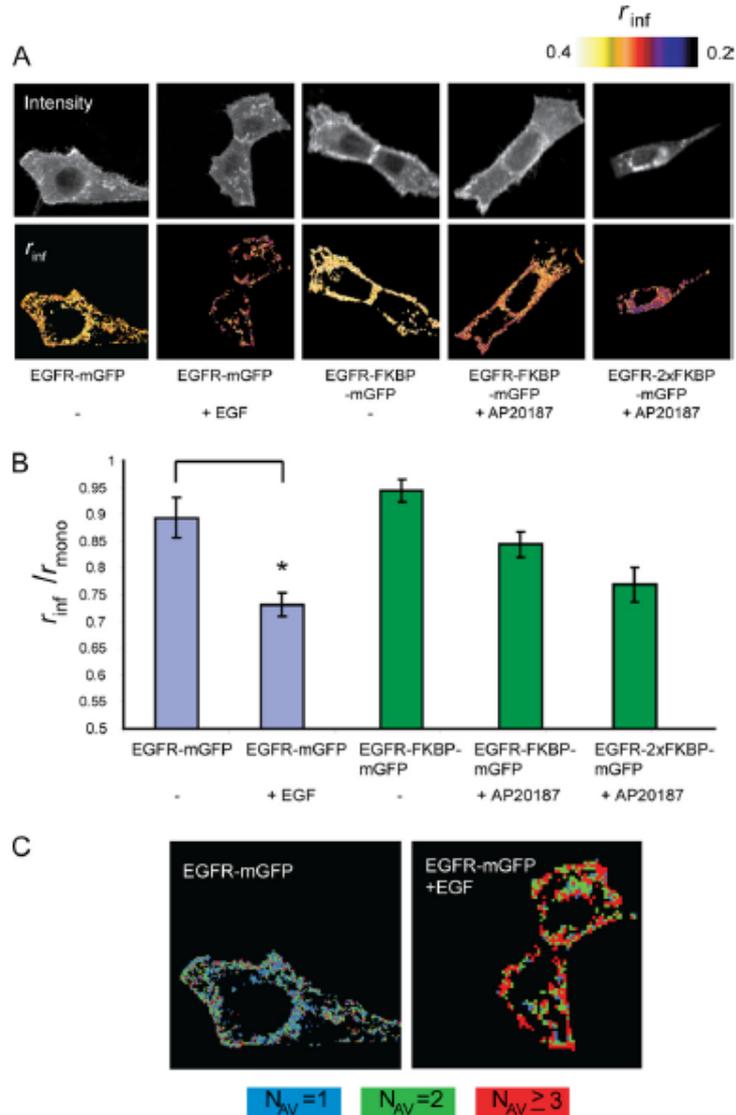


FIGURE 2. EGFR-mGFP is oligomerized after EGF stimulation. *A*, cellular distribution of GFP intensities and anisotropy values. NIH 3T3 2.2 cells expressing the indicated EGFR-mGFP constructs were left untreated or were stimulated with 8 nM EGF for 10 min or with 1 μ M AP20187 for 2 h. Limiting anisotropy values (r_{inf}) were measured as described under “Experimental Procedures” and are expressed in false colors. *B*, average anisotropy values expressed as fraction of r_{mono} of the indicated constructs (S.E.). *, $p < 0.005$. *C*, representation of the cluster size values of EGFR-mGFP before and after EGF stimulation in false colors. The anisotropy values of the cells shown in *A* were classified as monomers ($N_{AV} = 1$ (blue)), dimers ($N_{AV} = 2$ (green)), and oligomers ($N_{AV} \geq 3$ (red)) as described under “Experimental Procedures.”

(Hofman, Bader et al., 2010)

Using only one fluorophore brings advantages over hetero-FRET

- 1) all the fluorescent molecules are analyzed by this method comparing to hetero-FRET where dimers composed of donor-donor or acceptor-acceptor cannot be identified;
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