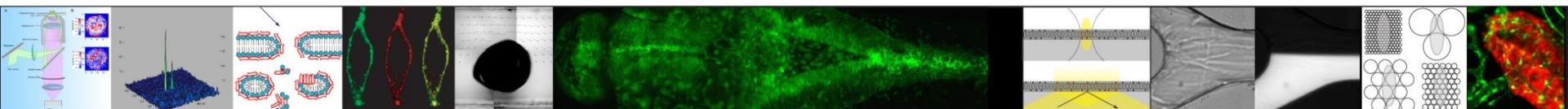


Biophysical Fluorescence Laboratory



Department of Chemistry



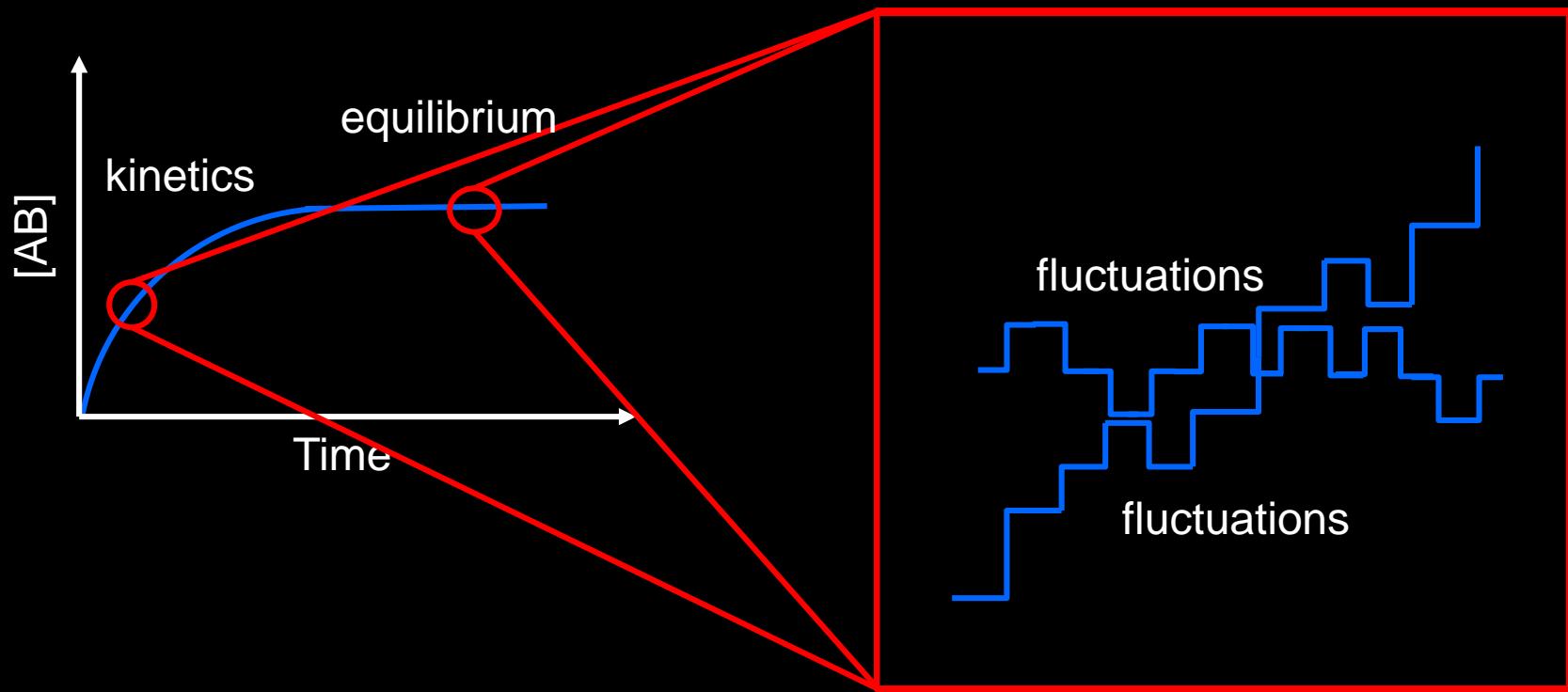
**Fluorescence Correlation and Cross-Correlation
Spectroscopy for the measurement of molecular
dynamics and interactions**

Thorsten Wohland

Outline

- Fluorescence Correlation Spectroscopy (FCS)
 - Introduction to basics of FCS
 - How to use amplitude, width and shape to obtain quantitative information
- Fluorescence Cross-Correlation Spectroscopy (FCCS)
 - Measurement of interactions and affinity constants (K_d s)
- FCS limitations and workarounds
- Imaging FCS
 - Motivation and principles
 - Example: Organization of Wnt3 in zebrafish membranes

Fluctuations



Correlations

$$\langle a \cdot b \rangle \neq \langle a \rangle \langle b \rangle$$

$$g = \frac{\langle a \cdot b \rangle}{\langle a \rangle \langle b \rangle}$$

Anti-correlation

$$g < 1$$

No correlation

$$g = 1$$

Correlation

$$g > 1$$

Autocorrelations

$$\langle a(t) \cdot a(t) \rangle \geq \langle a(t) \rangle \langle a(t) \rangle$$

$$\langle a(t) \cdot a(t + \tau) \rangle \geq \langle a(t) \rangle \langle a(t + \tau) \rangle$$

$$G(\tau) = \frac{\langle a(t) \cdot a(t + \tau) \rangle}{\langle a(t) \rangle \langle a(t + \tau) \rangle}$$

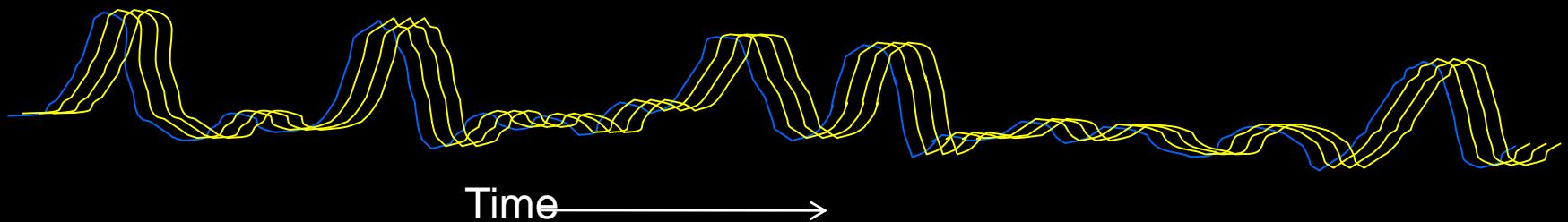
$$G(\tau) = \frac{\langle F(t + \tau) F(t) \rangle}{\langle F(t + \tau) \rangle \langle F(t) \rangle} = \frac{\langle F(t + \tau) F(t) \rangle}{\langle F(t) \rangle^2}$$

Stationary
Processes

Short time shifts τ

$$\langle F(t) \cdot F(t + \tau) \rangle ? \langle F(t) \rangle \langle F(t + \tau) \rangle$$

Blue: $F(t)$
Yellow: $F(t + \tau)$



$$\langle F(t) \cdot F(t + \tau_3) \rangle$$

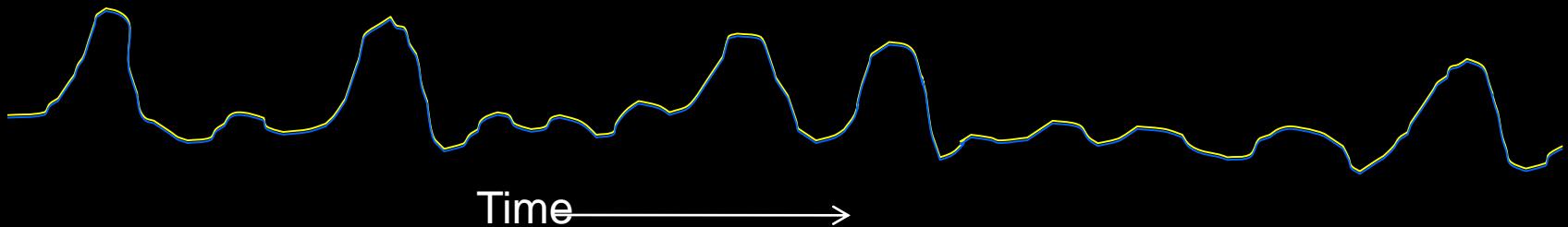
The intensity peaks always overlap to some extent and thus

$$\langle F(t) \cdot F(t + \tau) \rangle \geq \langle F(t) \rangle \langle F(t + \tau) \rangle$$

Long time shifts τ

$$\langle F(t) \cdot F(t + \tau) \rangle ? \langle F(t) \rangle \langle F(t + \tau) \rangle$$

Blue: $F(t)$
Yellow: $F(t + \tau)$



$$\langle F(t) \cdot F(t + \tau_3) \rangle$$

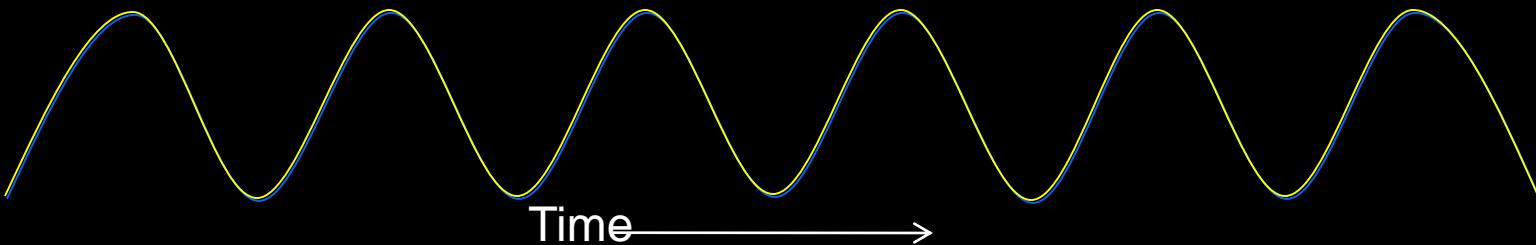
The intensity trace contains a random pattern of intensity peaks. Therefore an overlap of all/many peaks is only achievable at short times.

$$\langle F(t) \cdot F(t + \tau) \rangle = \langle F(t) \rangle \langle F(t + \tau) \rangle$$

Periodic signals

$$\langle F(t) \cdot F(t + \tau) \rangle ? \langle F(t) \rangle \langle F(t + \tau) \rangle$$

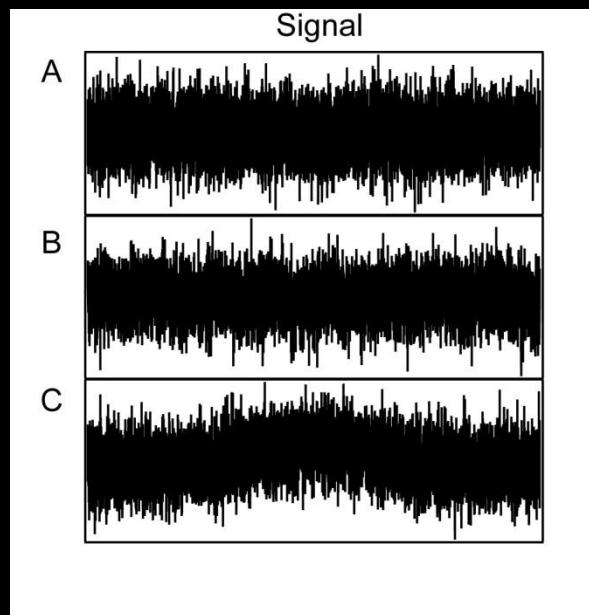
Blue: $F(t)$
Yellow: $F(t + \tau)$



$$\langle F(t) \cdot F(t + \tau_3) \rangle$$

The intensity trace contains a regular pattern of intensity peaks (i.e. it is repeated). Therefore an overlap of all/many peaks is achievable periodically and the correlation function will show that periodicity.

ACF: Autocorrelation
Function (the
correlation of a
variable with itself)



How is an ACF calculated practically?

Intensity values recorded every nanosecond



...



...

To calculate the correlation for the range of seconds you would need 1 billion values ...



...

If we make the time bins larger then we lose the information at short times.

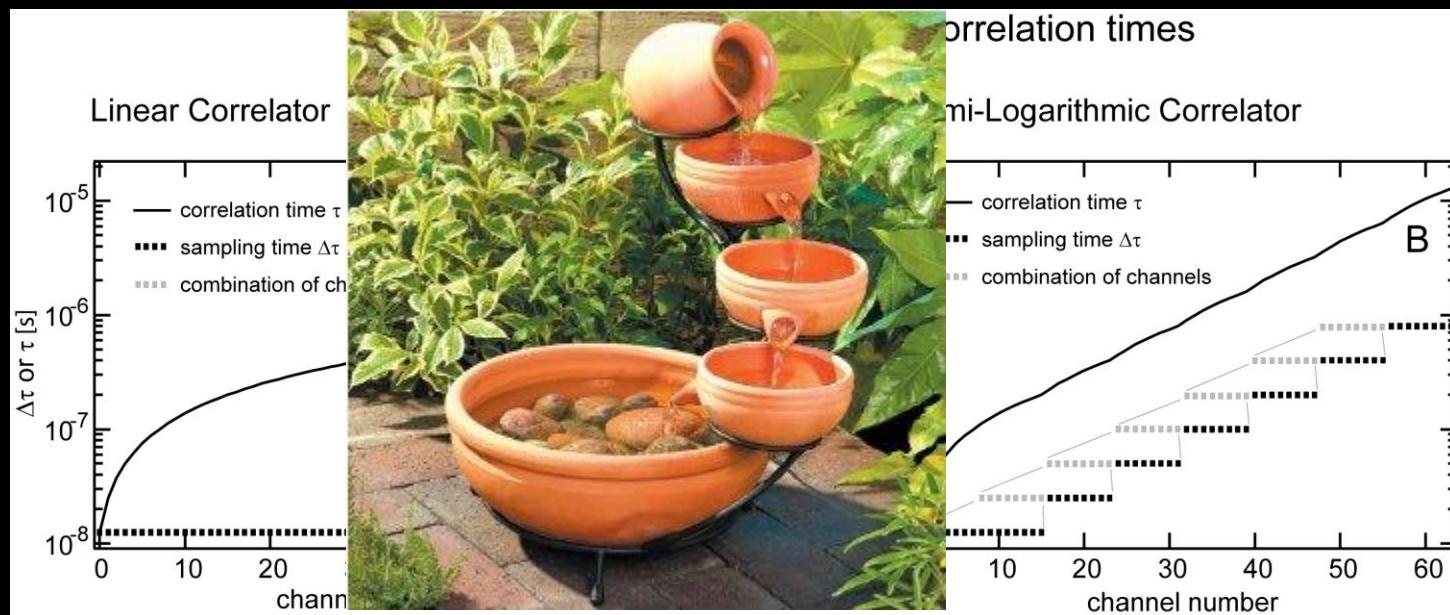


...

So best would be to use a varying time scheme.

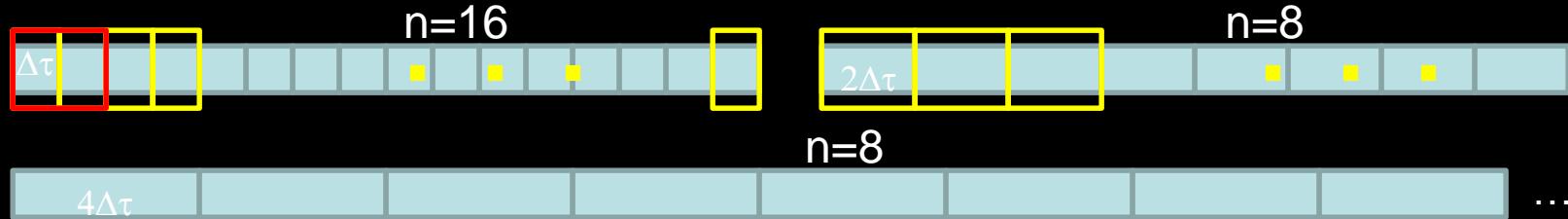
Correlation Time Schemes

The typical scheme used is called the semi-logarithmic time scale. The first n channels have a time $\Delta\tau$. The second group contains $n/2$ channels with $2 \Delta\tau$. The next group $n/2$ channels with $4 \Delta\tau$.



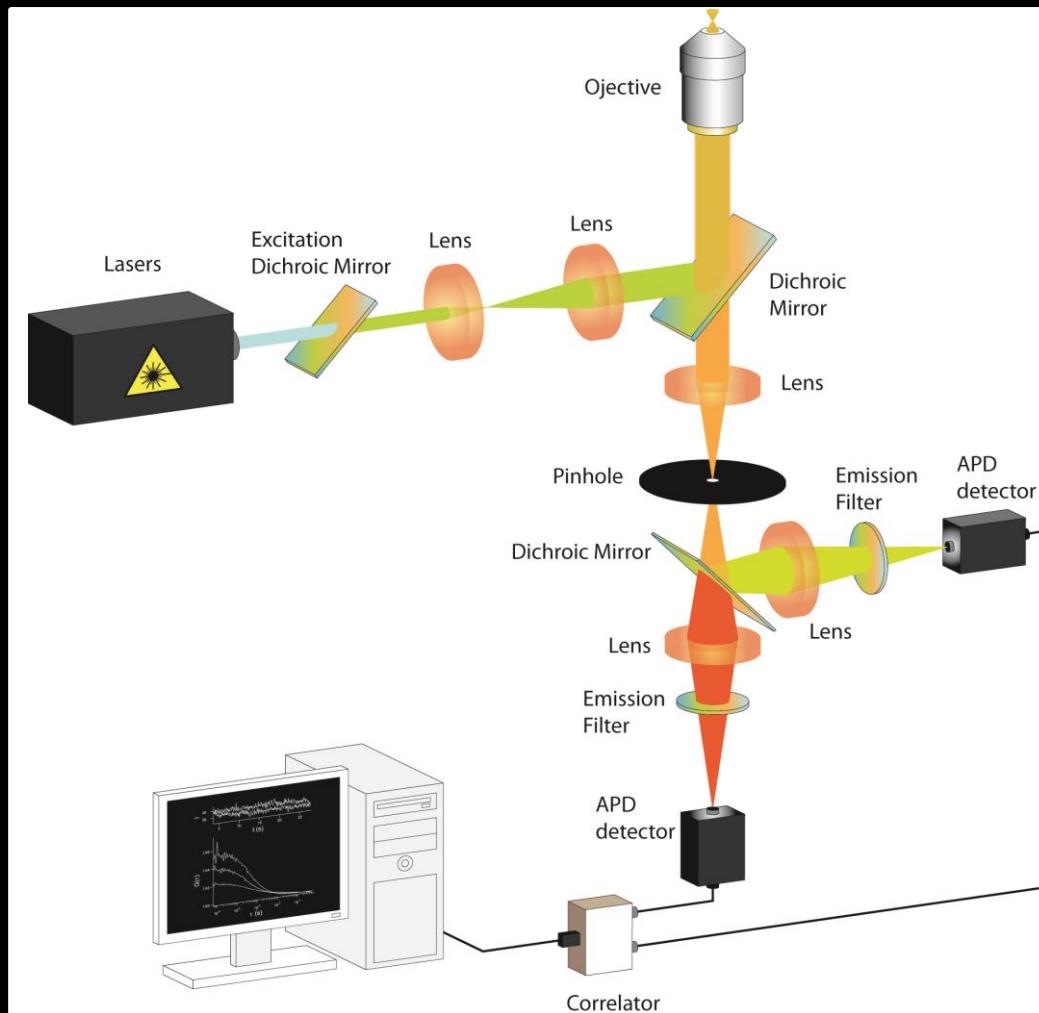
Correlation Time Schemes

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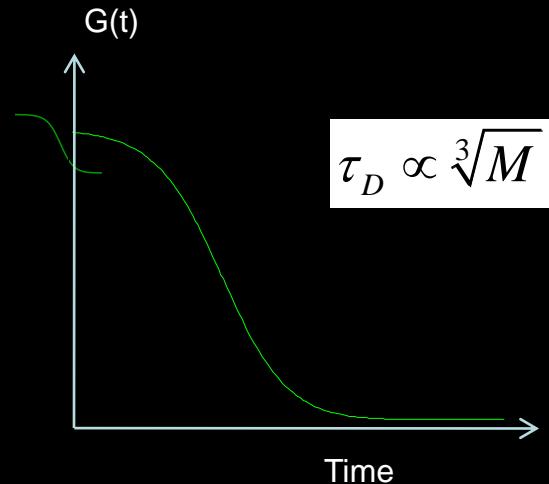
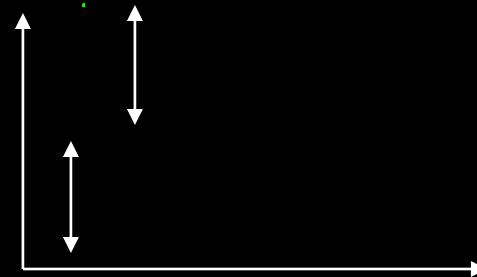
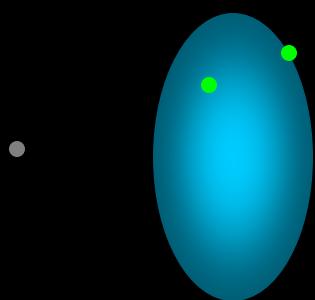
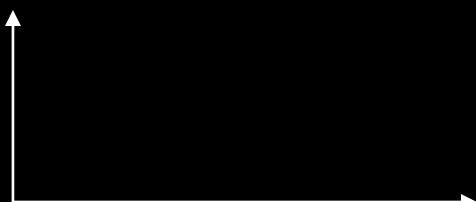
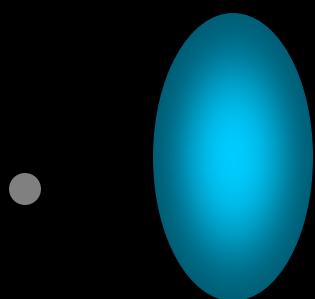
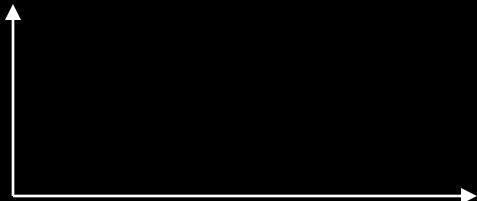
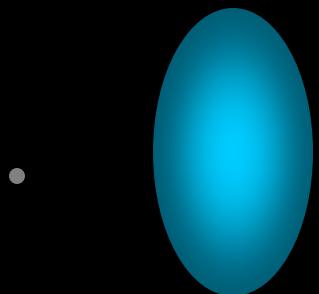


- 1) Each time a new measurement of length $\Delta\tau$ comes in, calculate all ACF values for lag times 0 to $16\Delta\tau$.
- 2) After 2 measurements of $\Delta\tau$, correlate the last two newest measurements with all channels in group 2. Then take the last two channels of group 1 and combine them into one channel with width $2\Delta\tau$ of group 2 and shift.

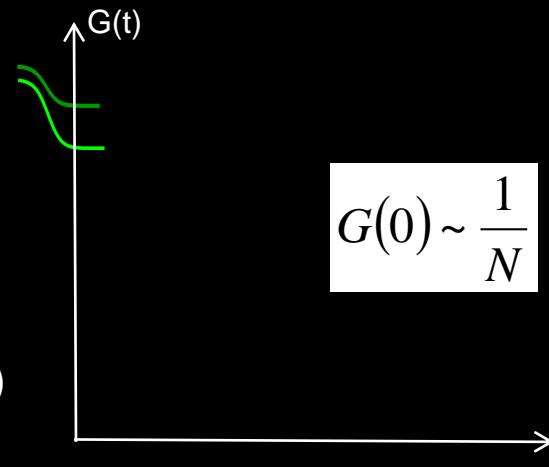
Confocal FCS setup



FCS: Characteristic Parameters



$$\tau_D \propto \sqrt[3]{M}$$

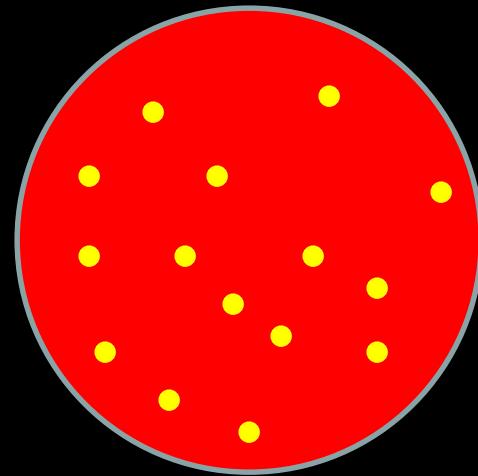
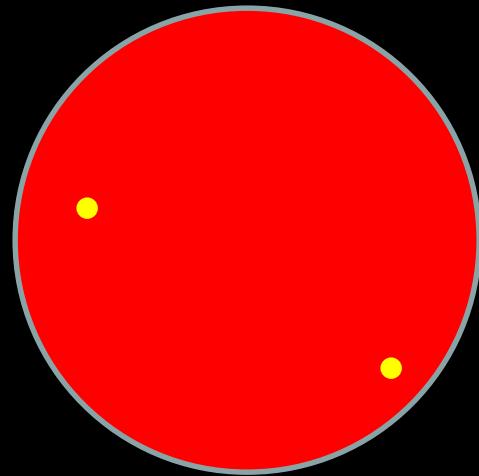
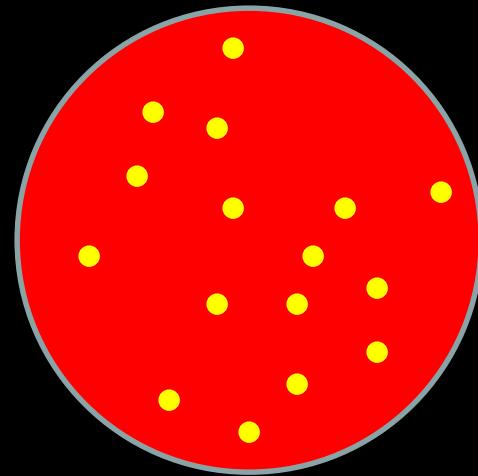
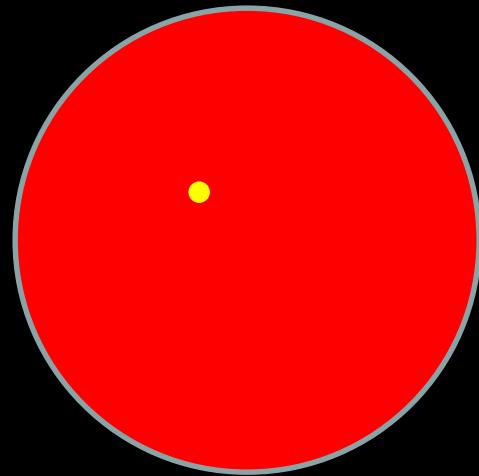


$$G(0) \sim \frac{1}{N}$$

$\sim F(1)$

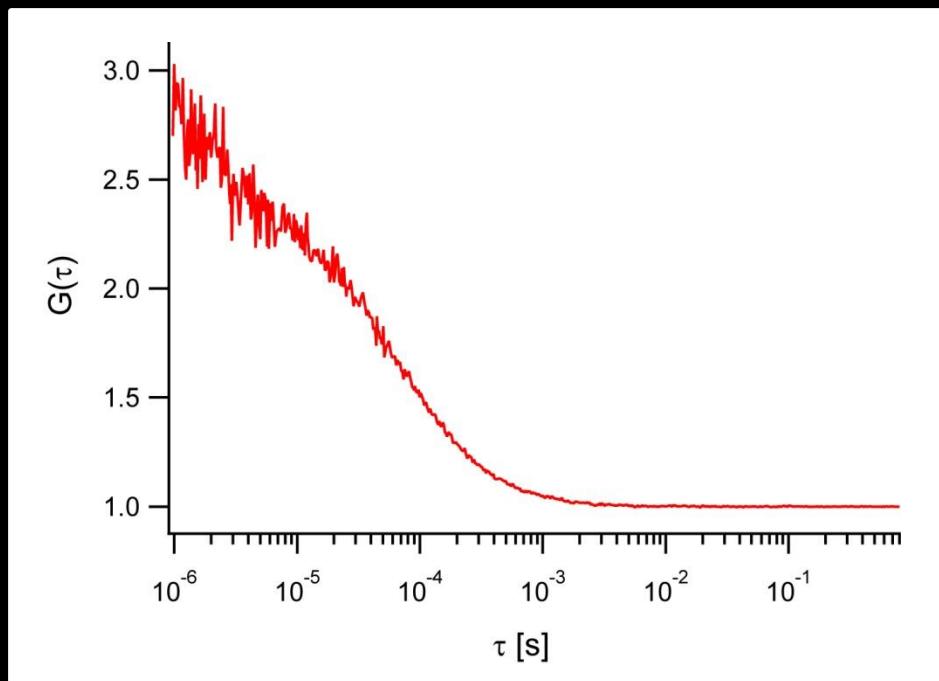
$\sim F(N)$

Time



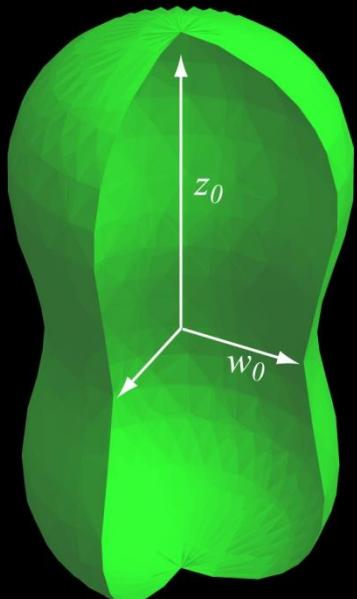
Correlation Functions

$$G(\tau) = \frac{\langle F(t + \tau)F(t) \rangle}{\langle F(t) \rangle^2} = \frac{\langle \delta F(t)\delta F(t + \tau) \rangle}{\langle F(t) \rangle^2} + 1$$



Correlation Functions

$$G(\tau) = \frac{1}{\langle C \rangle \pi^{3/2} w_0^2 z_0} \left(1 + \frac{4D\tau}{w_0^2} \right)^{-1/2} \left(1 + \frac{4D\tau}{w_0^2} \right)^{-1/2} \left(1 + \frac{4D\tau}{z_0^2} \right)^{-1/2} + 1$$



$$z_0 = K w_0$$

Number of particles

$$N = \langle C \rangle V_{eff} = \langle C \rangle \pi^{3/2} w_0^2 z_0$$

Correlation time

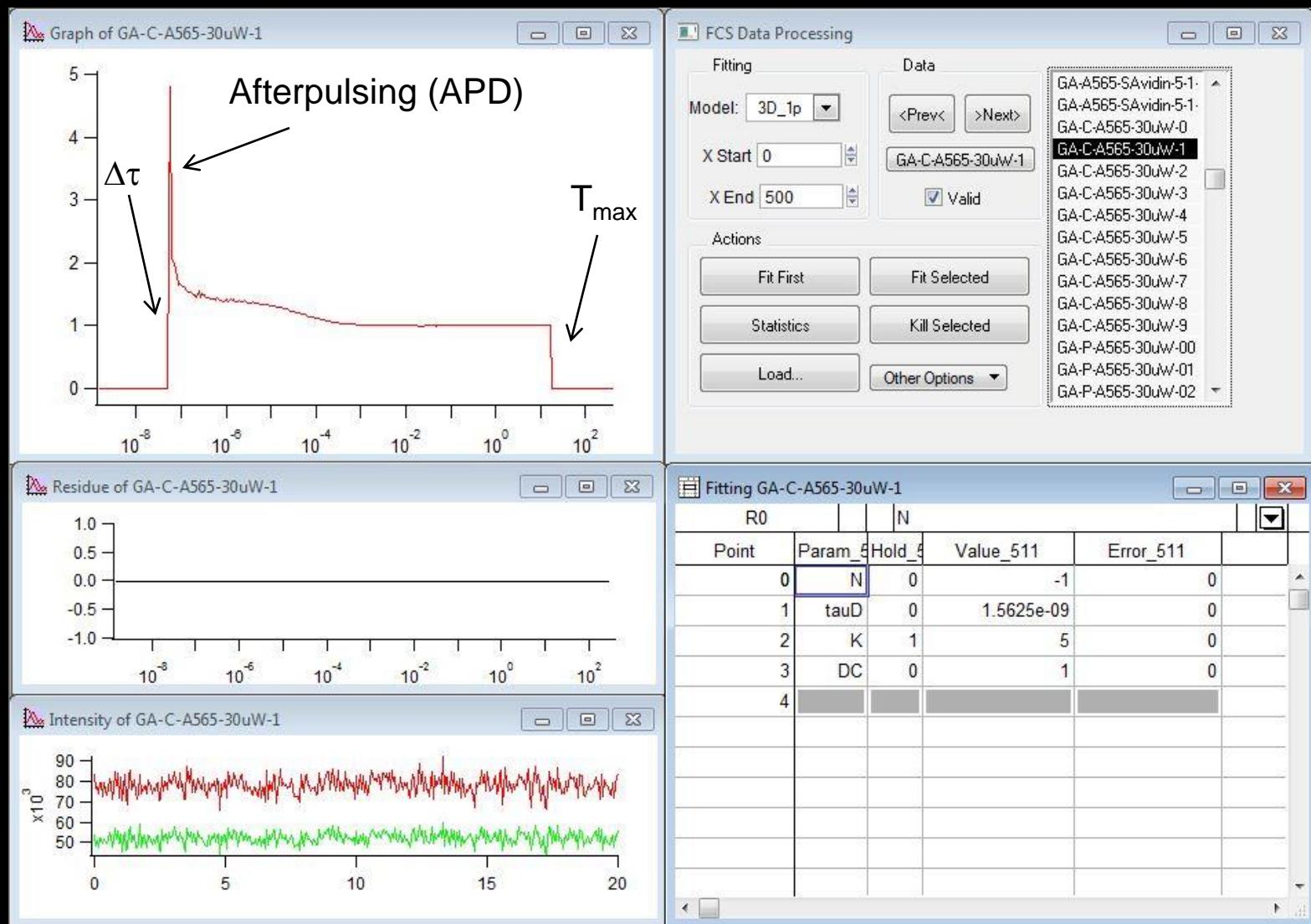
$$\tau_D = \frac{w_0^2}{4D}$$

Structure factor

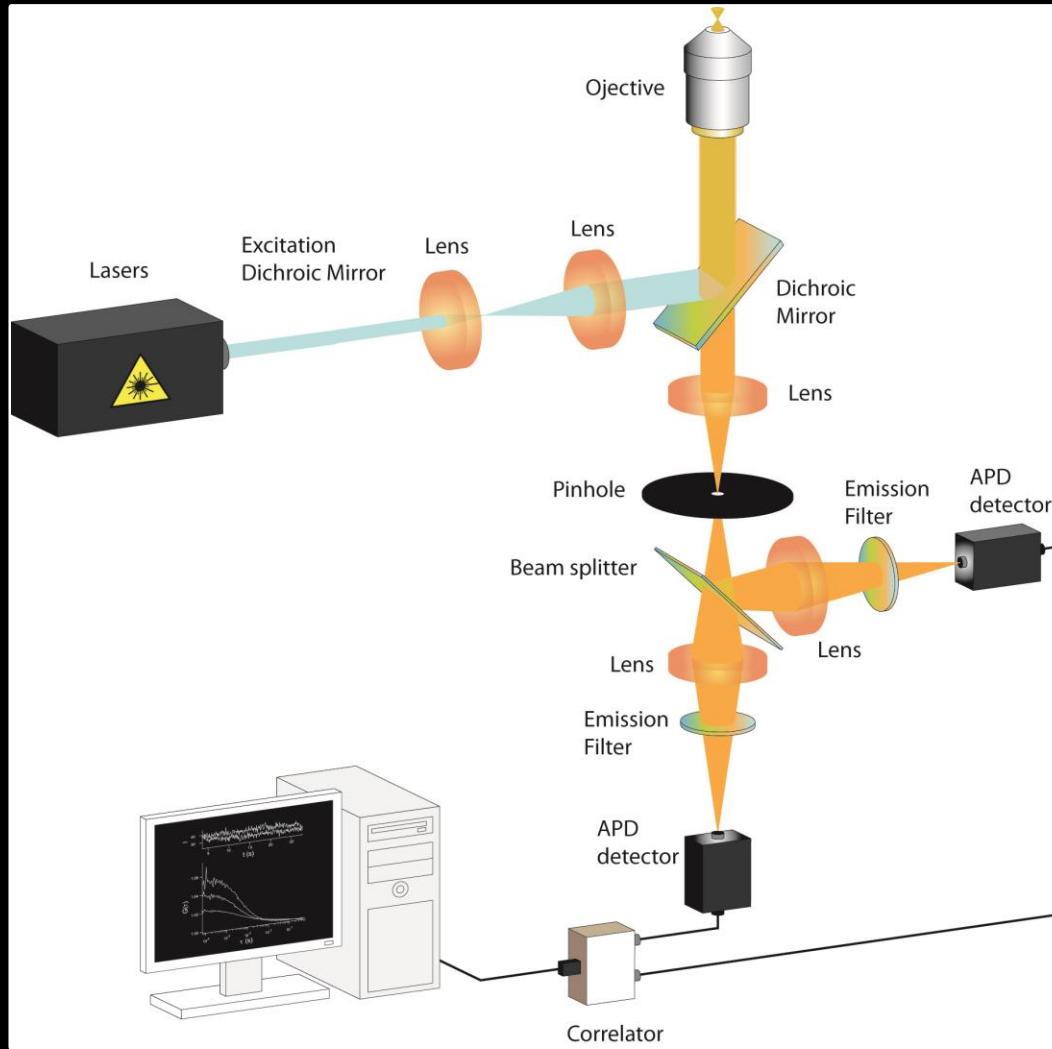
$$K = \frac{z_0}{w_0}$$

$$G(\tau) = \frac{1}{N} \left(1 + \frac{\tau}{\tau_D} \right)^{-1} \left(1 + \frac{\tau}{K^2 \tau_D} \right)^{-1/2} + G_\infty$$

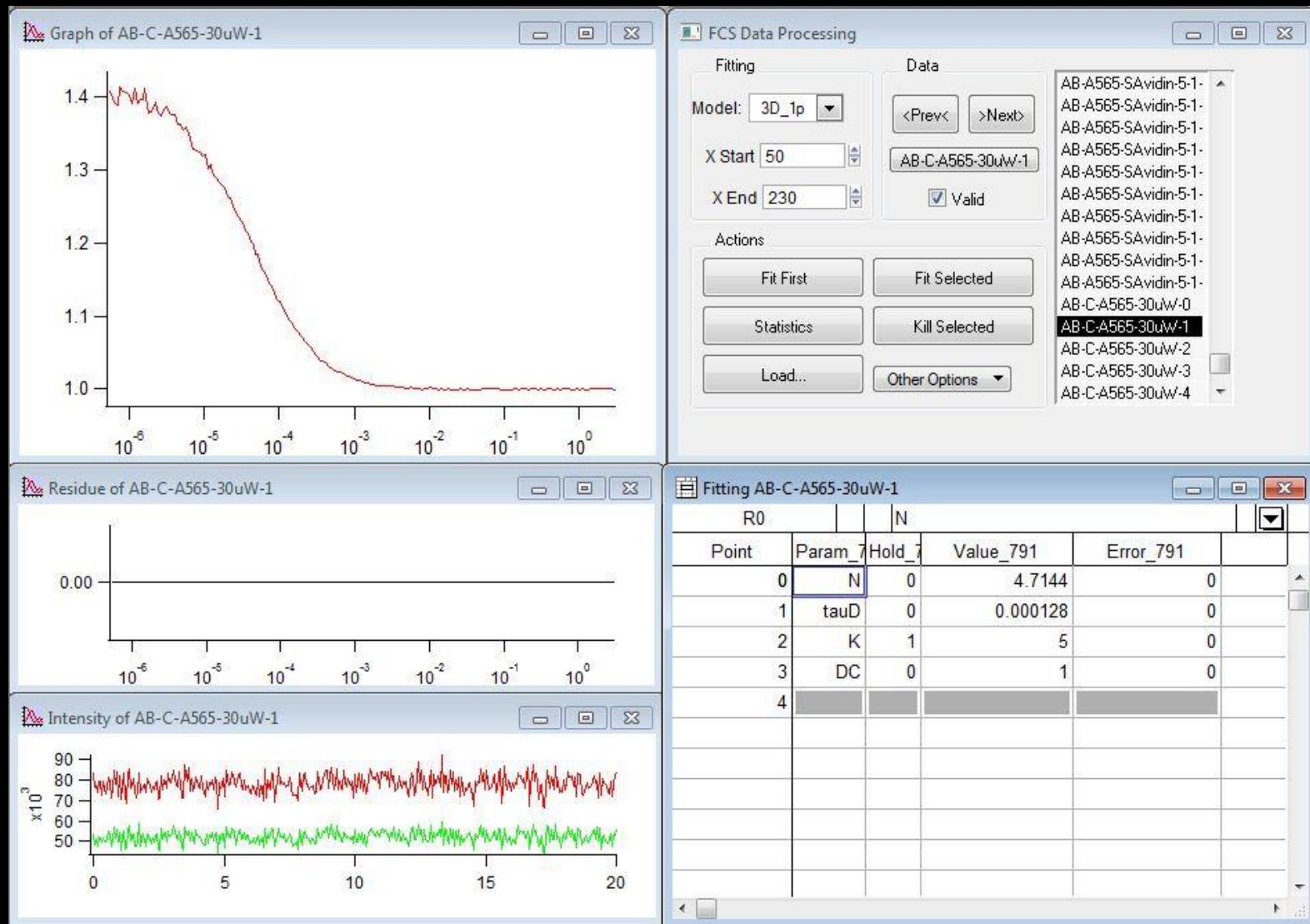
Data Fitting: Raw data



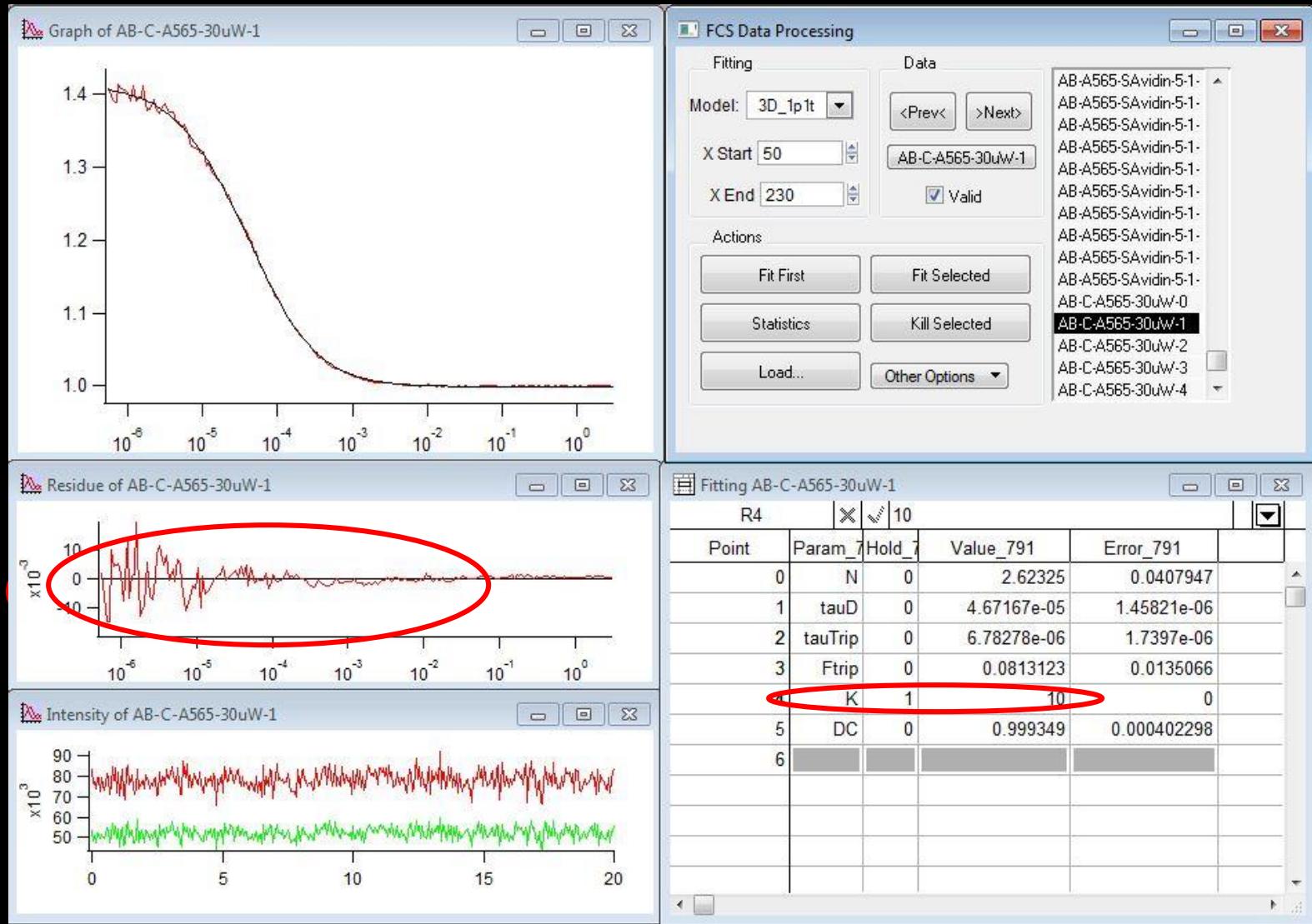
Removal of afterpulsing by cross-correlation



Data Fitting: Raw data



Data Fitting



Weighted data fits

	non-weighted fit	weighted fit
N	2.601	2.591
τ_D	4.781e-05	4.739e-05
τ_{Trip}	4.992e-06	4.406e-06
F_{Trip}	1.010e-01	8.187e-02
K	6.089	5.843
G_∞	1.000	1.000
Residue of A		
	Standard deviation	
N	9.377e-02 (3.6 %)	6.294e-02 (2.4 %)
τ_D	4.254e-06 (8.9 %)	2.699e-06 (5.7 %)
τ_{Trip}	4.128e-06 (83 %)	2.439e-06 (55 %)
Intensity of A		
F_{Trip}	3.780e-02 (37 %)	1.853e-02 (23 %)
K	1.940 (32 %)	1.219 (21 %)
G_∞	8.698e-05	5.586e-05

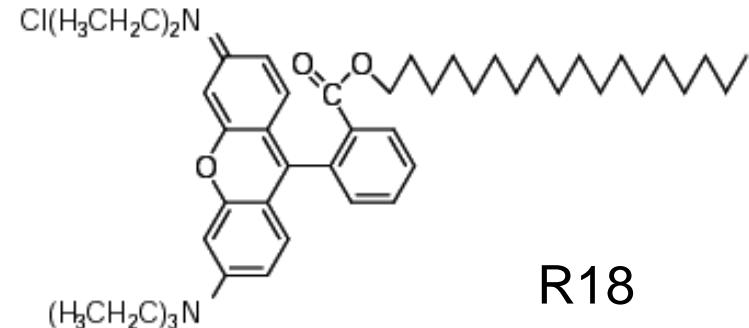
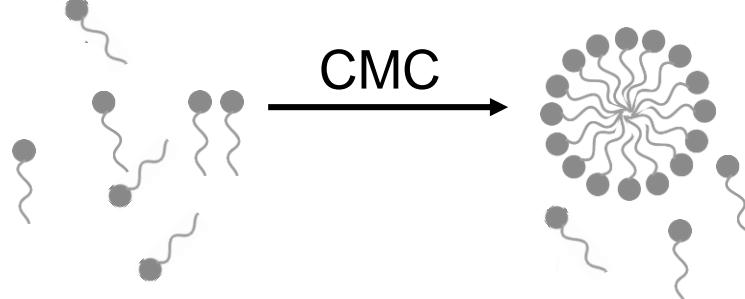
Measured over 10 experiments

How to use the FCS amplitude

Aggregation numbers of
detergent/lipid micelles

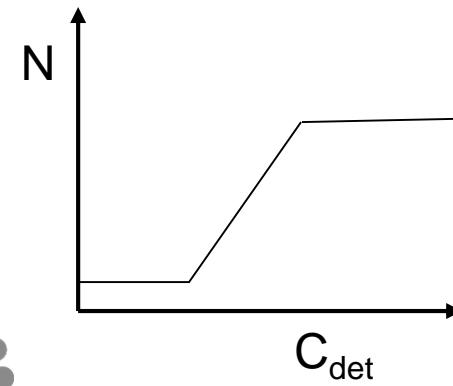
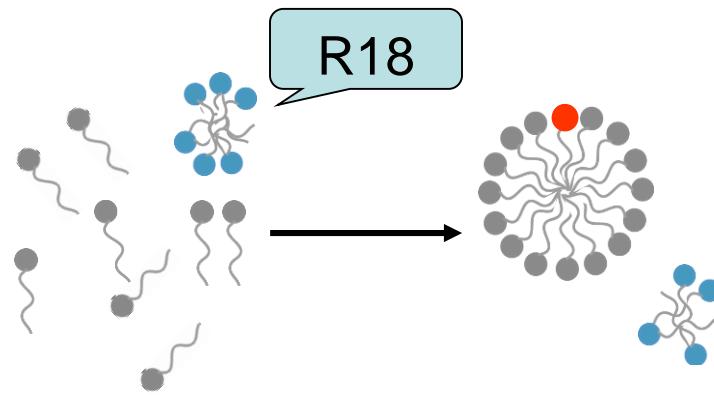
Determination of the aggregation of detergent and LPS

Micelle Formation

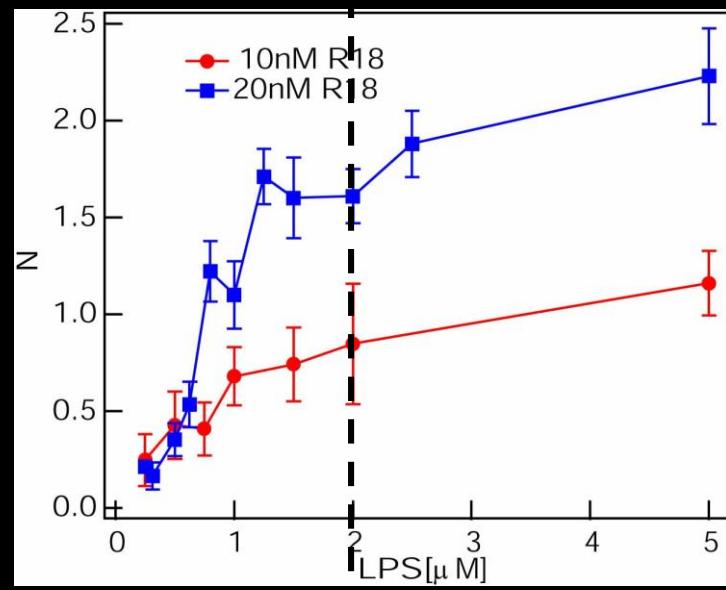
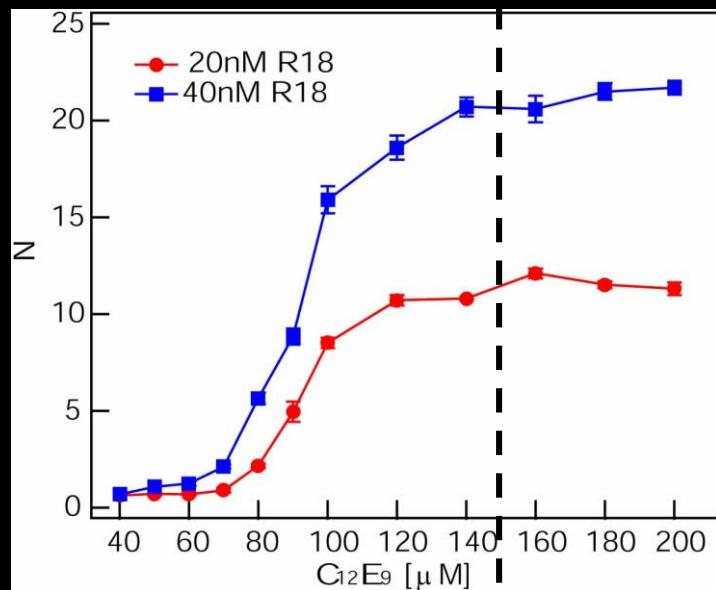


R18

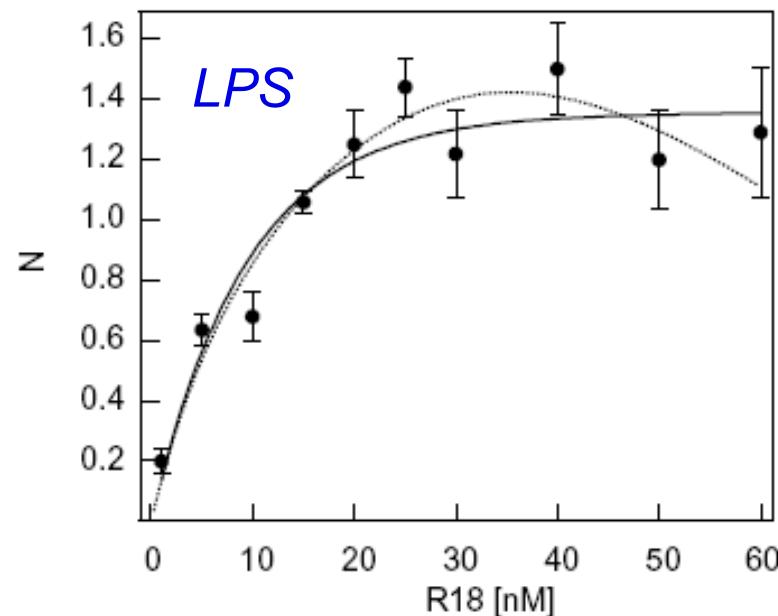
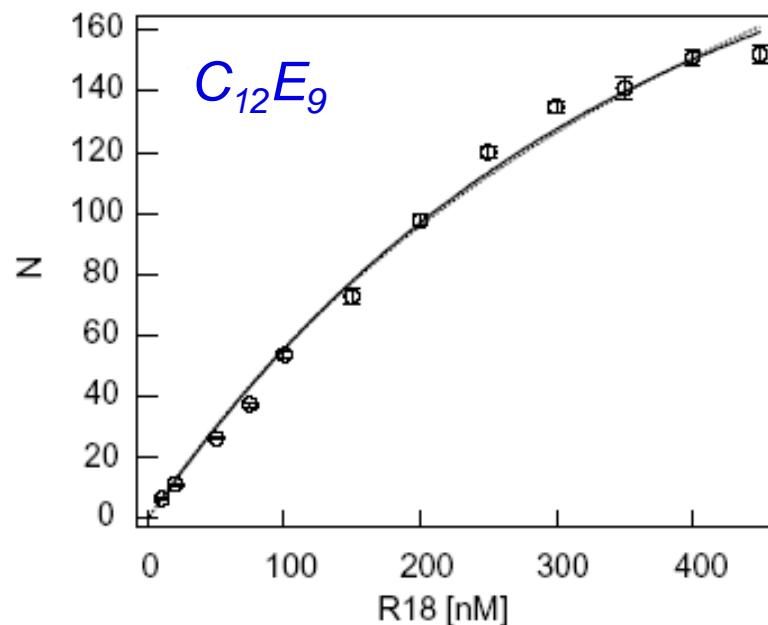
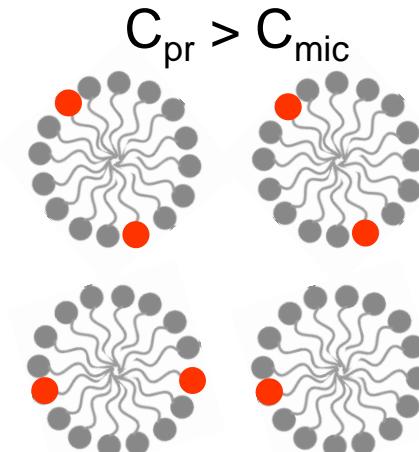
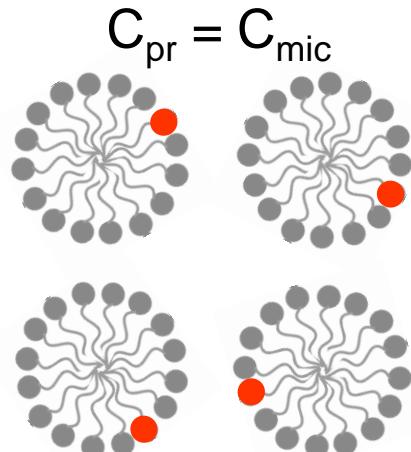
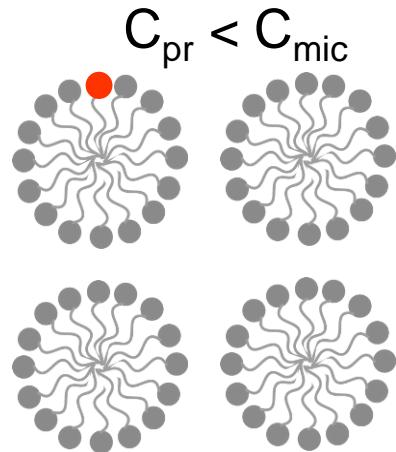
R18 non-flu oligomers \rightarrow Micelle formation \rightarrow dissolution of R18 oligomers and incorporation into micelles with fluorescence increase



Determination of the aggregation of detergent and LPS



Determination of the aggregation of detergent and LPS



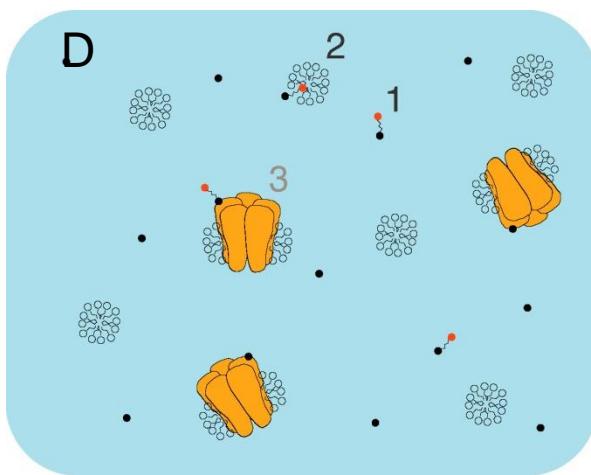
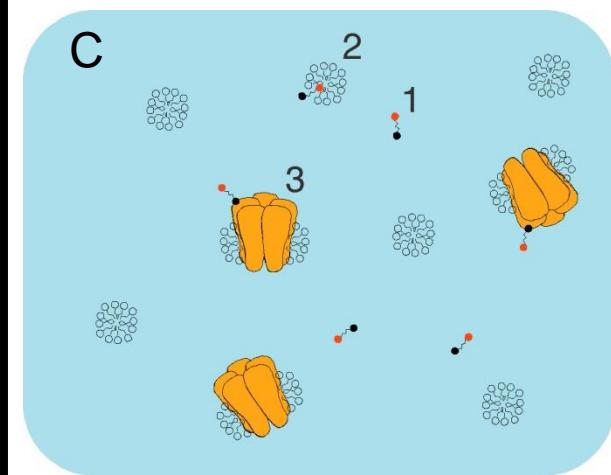
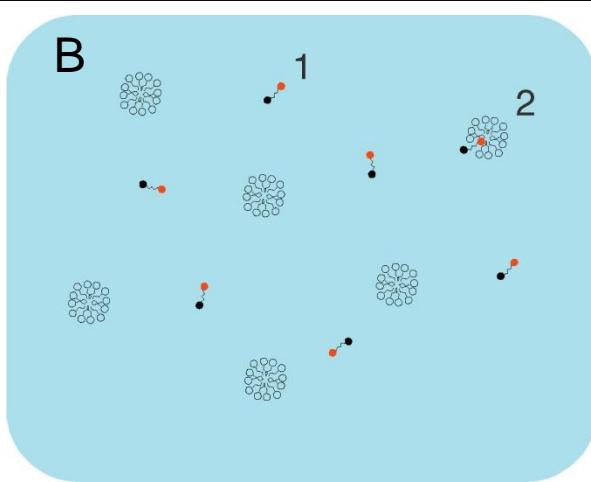
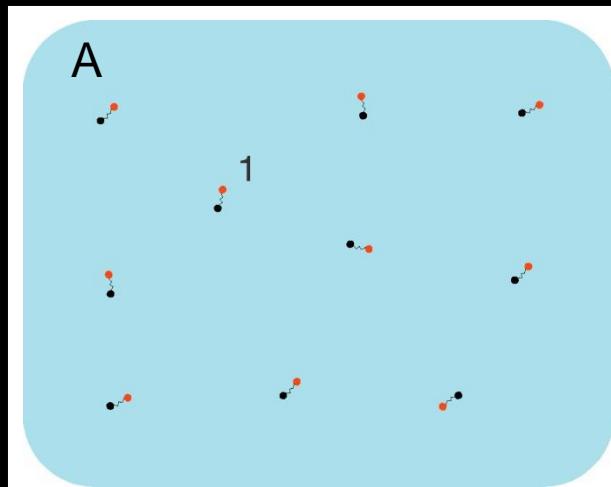
$c_{cmc}=93-105 \mu M$; $N_{agg}=112-132$
($c_{cmc}=80 \mu M$; $N_{agg}=120$)

$c_{cmc}=1.3-1.6 \mu M$; $N_{agg}=43-49$

How to use the FCS width

Ligand affinities for the 5HT₃
receptor

Measurements in Solution



Parameters: correlation times (τ_1, τ_2, τ_3),

fraction of particles (Y_1, Y_2, Y_3)

A: Ligand in Buffer solution

B: Ligand + Detergent

C:
Ligand+Detergent
+Receptor

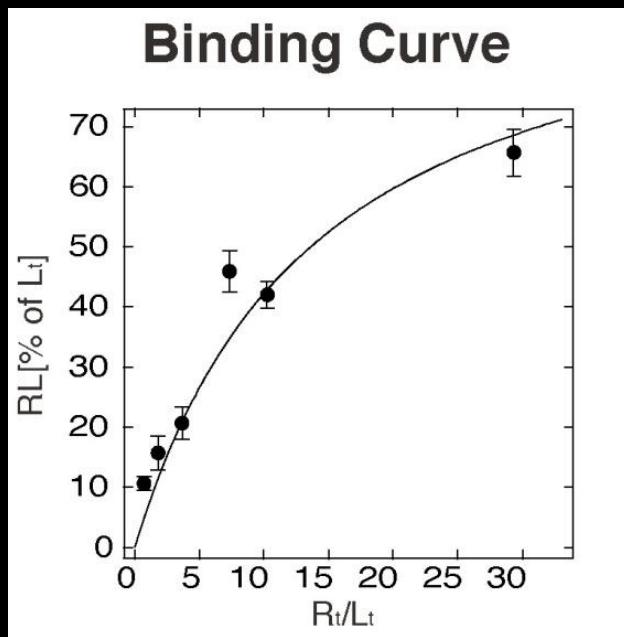
D:
Ligand+Detergent+
Receptor
+Competitor

Ligand-Receptor Interactions

Ligands: 0.5 – 1.1 kDa $425 \text{ } \mu\text{m}^2/\text{s}$

C_{12}E_9 micelle: 60 - 70 kDa $73 \text{ } \mu\text{m}^2/\text{s}$

$5\text{HT}_{3\text{As}}\text{-R} + \text{micelle}$: ~320 kDa $35 \text{ } \mu\text{m}^2/\text{s}$



$$K_d^{FCS} = 15.7 \pm 8.0 \text{ nM}$$

$$K_d^{RBA} = 18.0 \pm 2.0 \text{ nM}$$

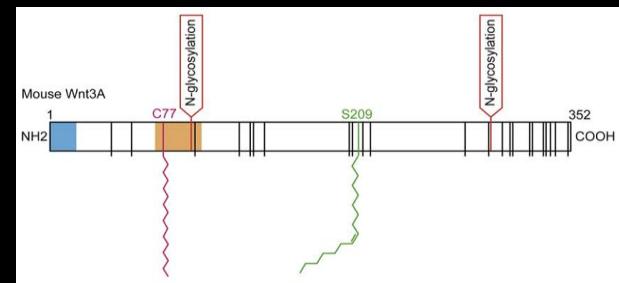
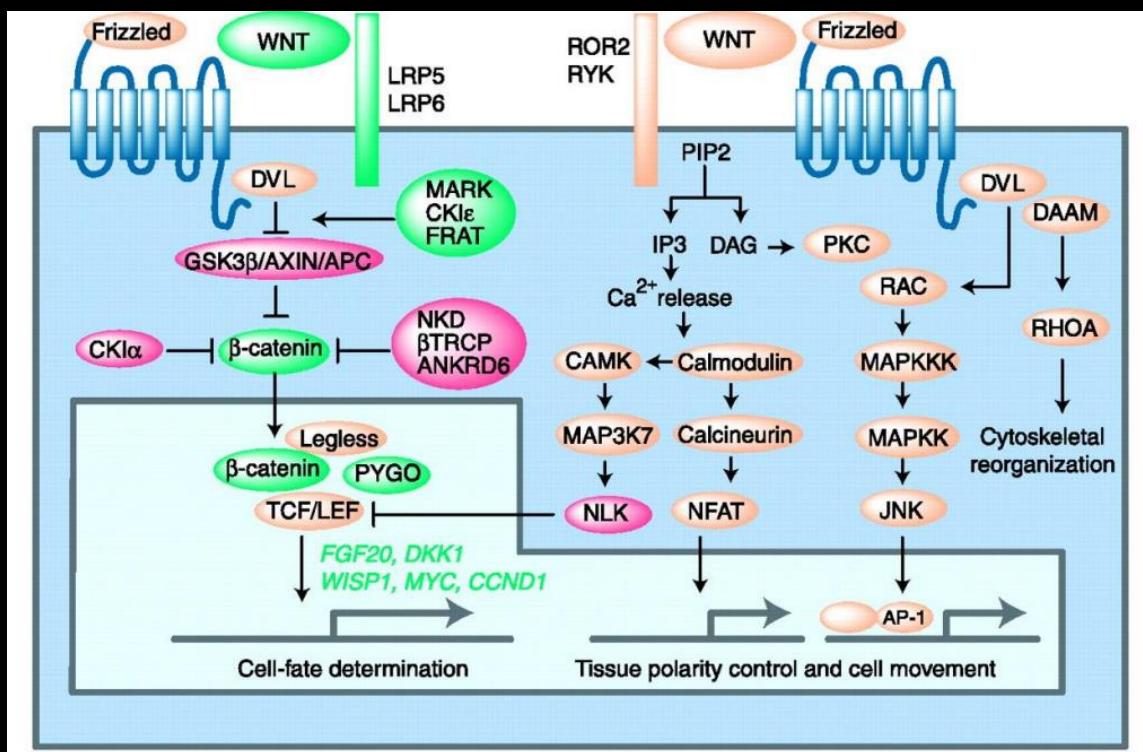
How to use the FCS shape

Determination of morphogen
secretion in live zebrafish

Wnt Signaling

Canonical Wnt Pathway
Wnt/β-catenin signaling

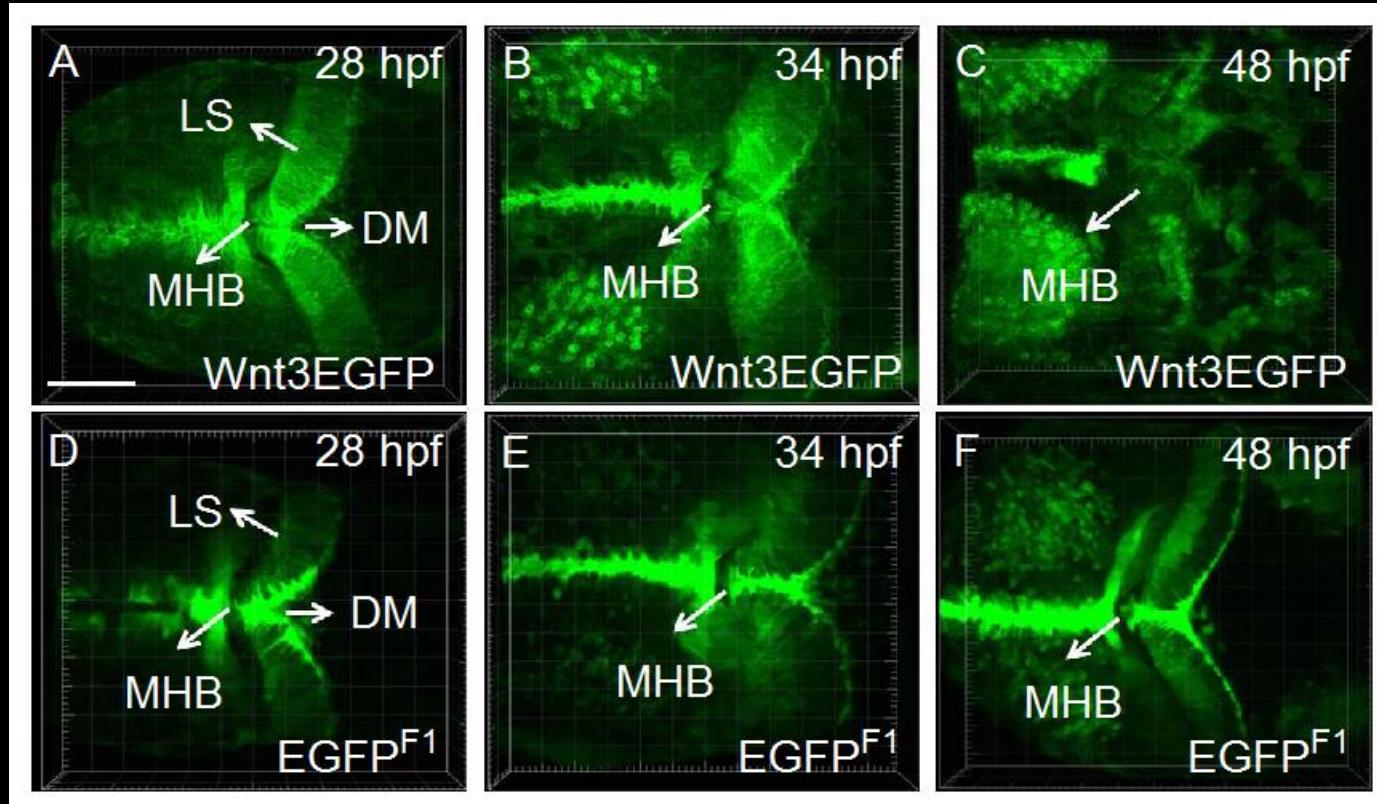
Non-Canonical Wnt Pathway
β-catenin independent Wnt signaling



1. Is Wnt3 secreted?
2. Where in the membrane does Wnt3 reside?

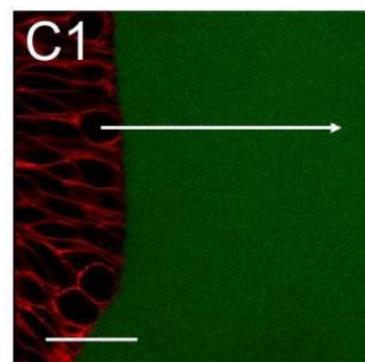
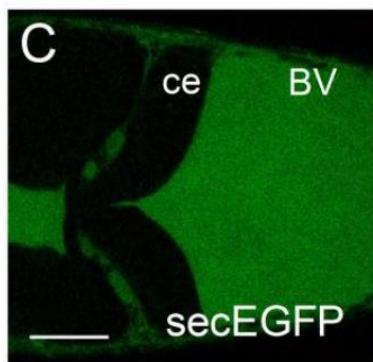
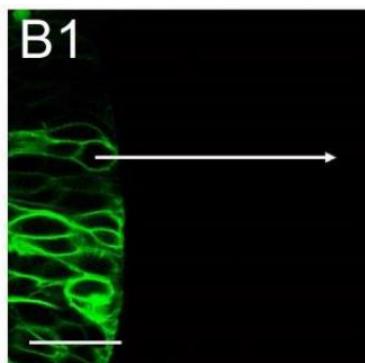
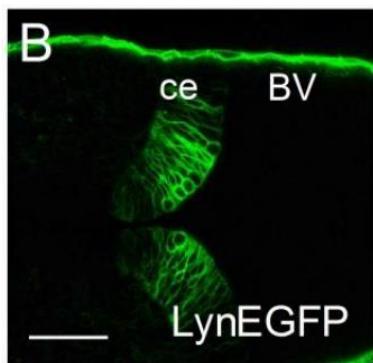
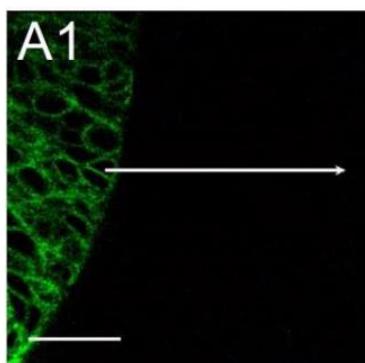
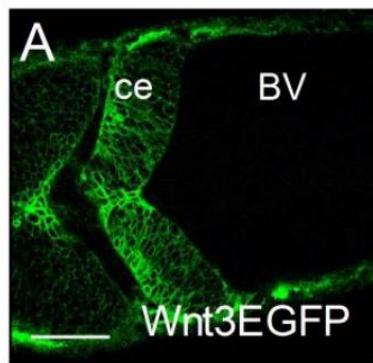
Katoh, M. *Clinical Cancer Research* 2007, 13, 4042-4045.

Wnt3EGFP Expression in the Cerebellum

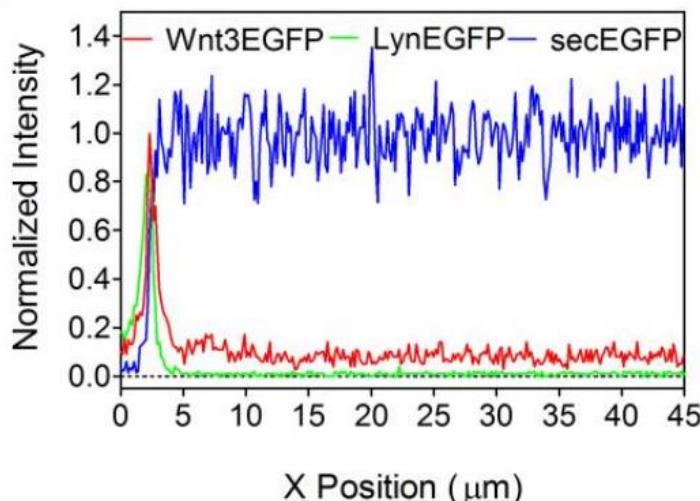


ce: cerebellum; ot: optic tectum; MHB: midbrain/hindbrain boundary;
BV: brain ventricle; DM: dorsal midline; LS: lateral side.

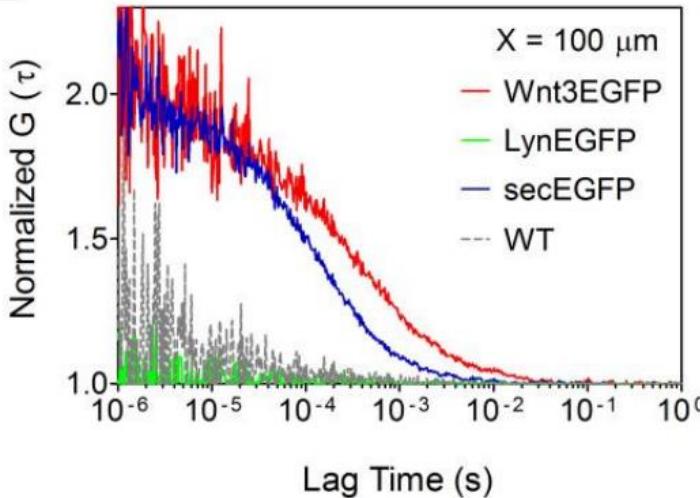
Wnt3EGFP Secretion to the Brain Ventricle



D



E



ce: cerebellum; BV: brain ventricle

Bayesian Model Selection

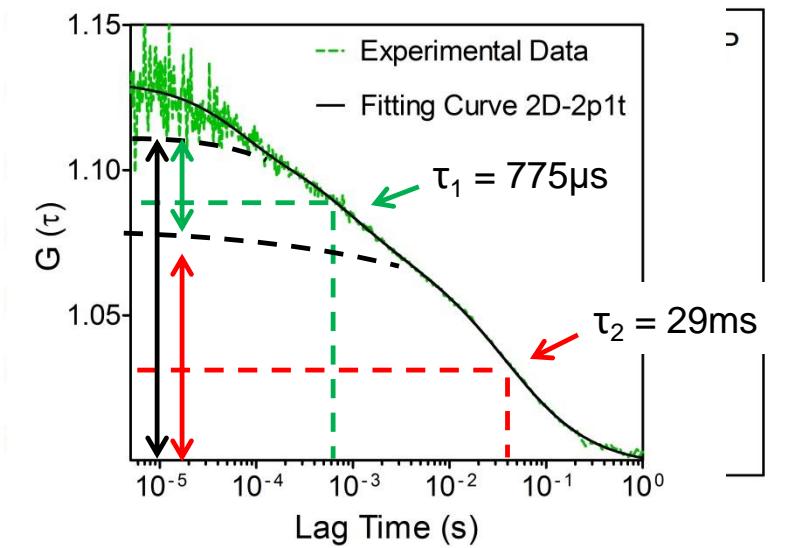
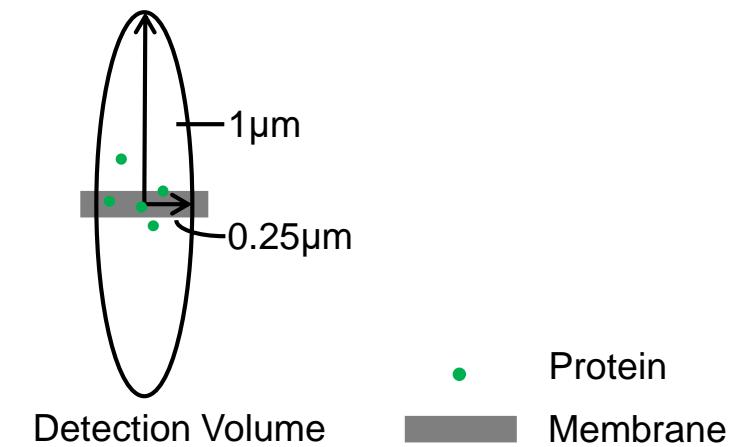
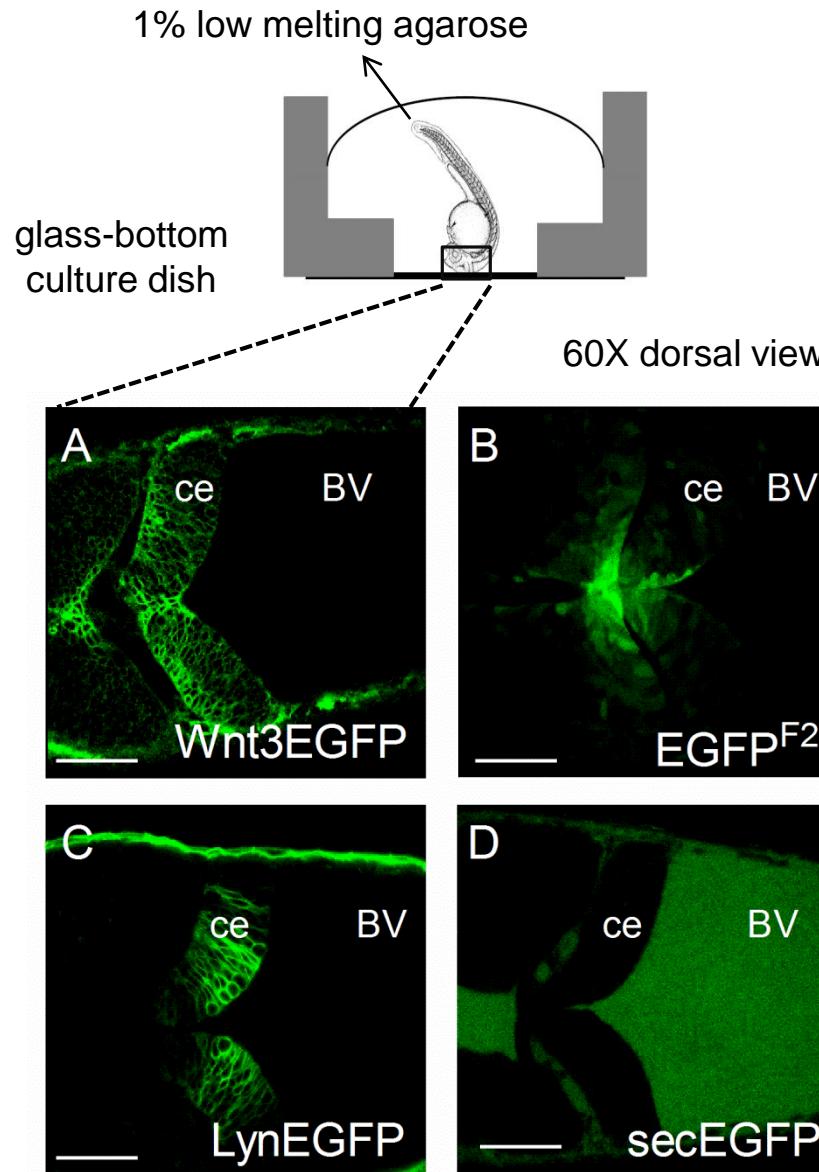
- Bayes' Theorem

$$P(M_k|\mathbf{y}) = \frac{P(\mathbf{y}|M_k)P(M_k)}{P(\mathbf{y})}$$

$$P(\mathbf{y}|M_k) = \int_{\beta} P(\mathbf{y}|\boldsymbol{\beta}, M_k)P(\boldsymbol{\beta}|M_k) d\boldsymbol{\beta}$$

$$\begin{aligned} P(\mathbf{y}|\boldsymbol{\beta}) &= \frac{1}{(2\pi)^{n/2} \sqrt{\det(\mathbf{C})}} \exp \left\{ -\frac{1}{2} [\mathbf{y} - \mathbf{f}(\mathbf{x}, \boldsymbol{\beta})]^T \mathbf{C}^{-1} \right. \\ &\quad \left. \times [\mathbf{y} - \mathbf{f}(\mathbf{x}, \boldsymbol{\beta})] \right\} \end{aligned}$$

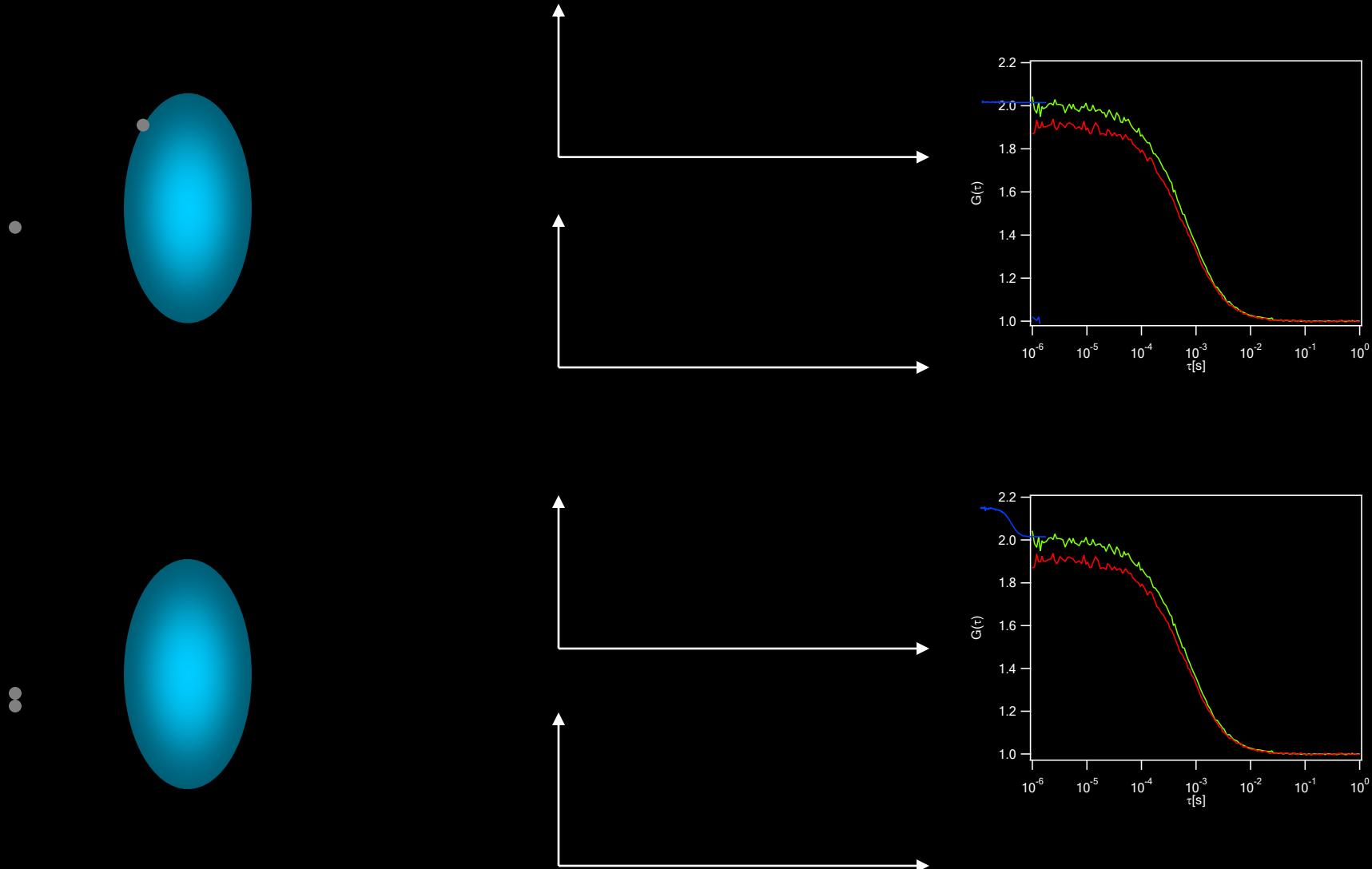
Zebrafish FCS measurements



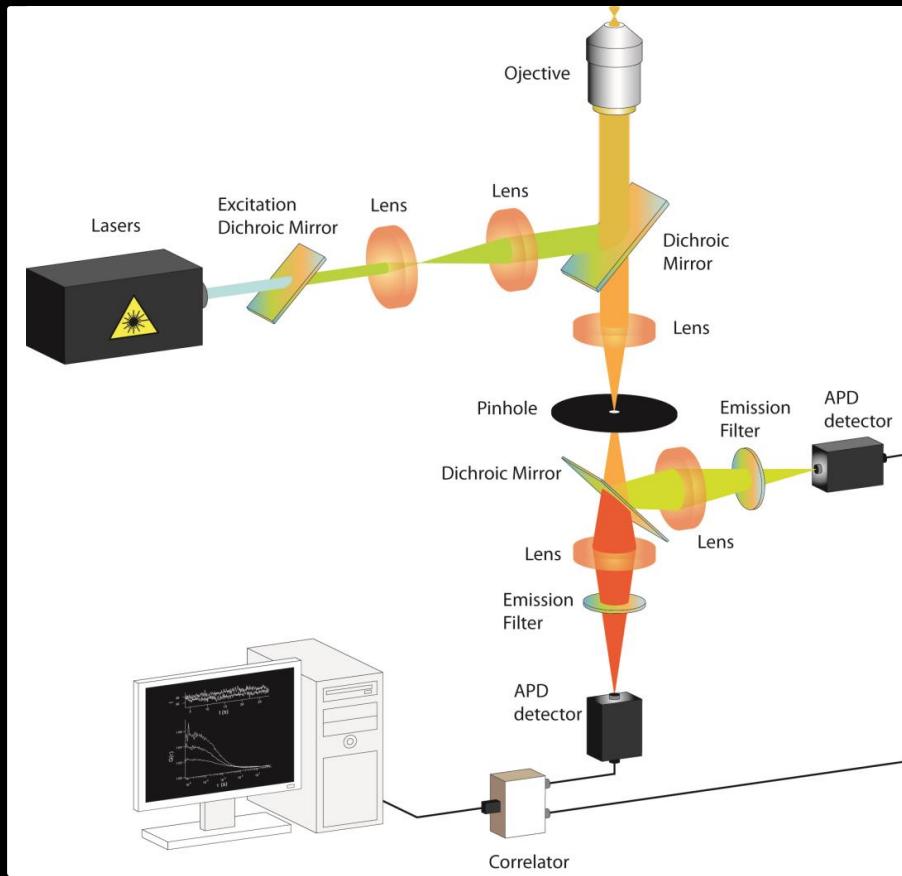
ce: cerebellum; BV: brain ventricle

Fluorescence Cross-Correlation Spectroscopy

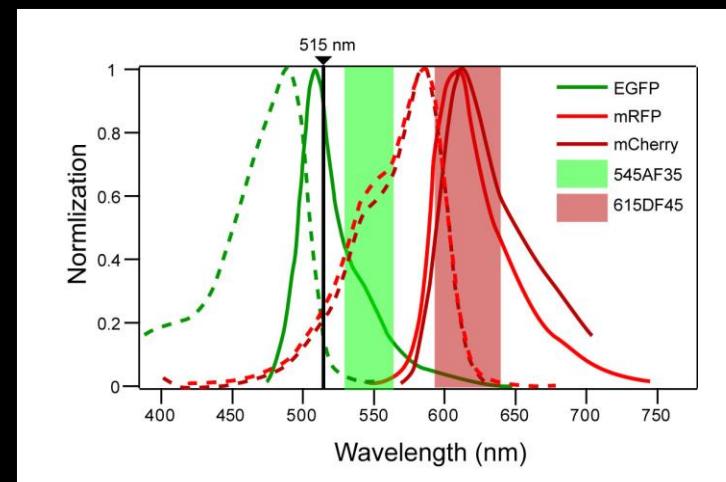
Fluorescence Cross-correlation Spectroscopy (FCCS)



SW-FCCS



Fluorophores:
Quantum dots
Tandem dyes (energy transfer dyes)
Organic dyes
Fluorescent proteins



~2000 counts per second and particle

Ricka and Binkert, *Phys Rev A*, 39(5) :2646-52 (1989)

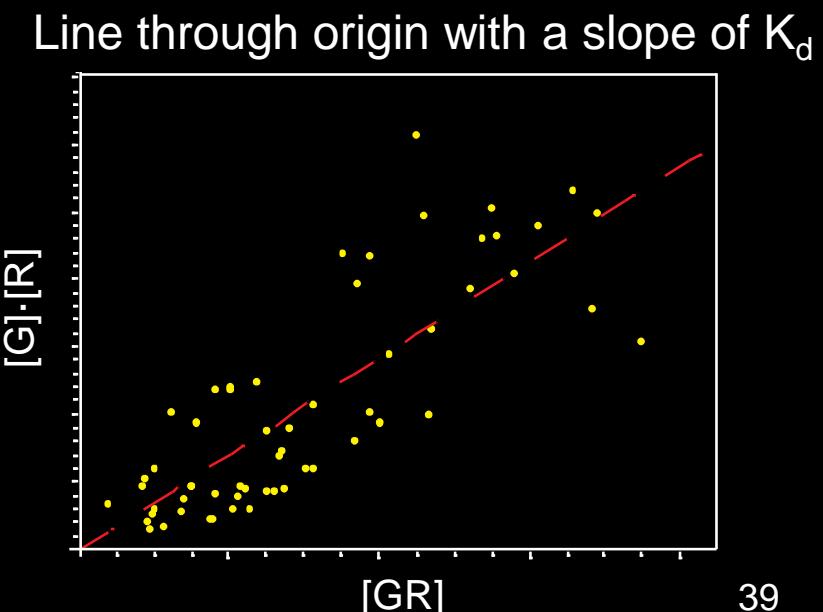
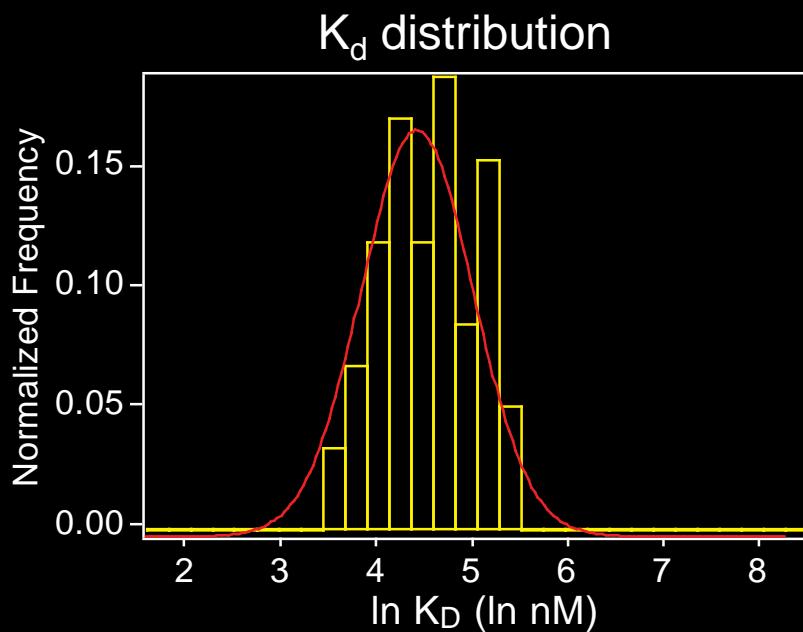
Hwang and Wohland, *ChemPhysChem* 5, 549-551 (2004)

Hwang and Wohland, *JCP*, 122, 114708 (2005)

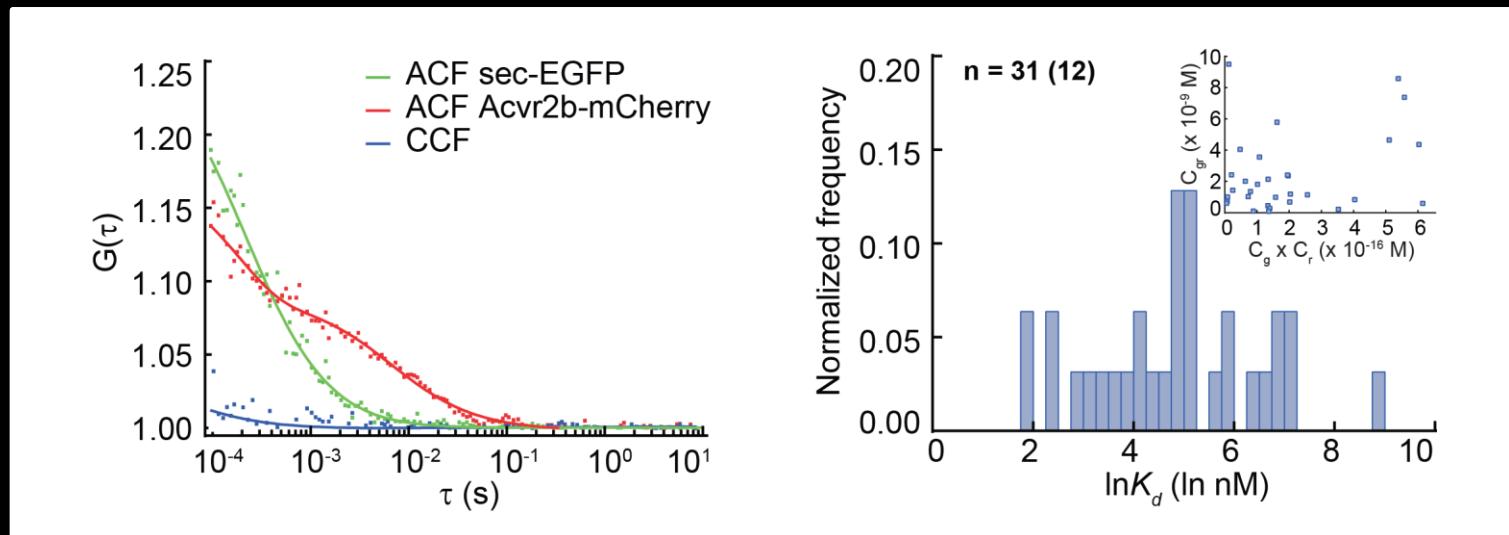
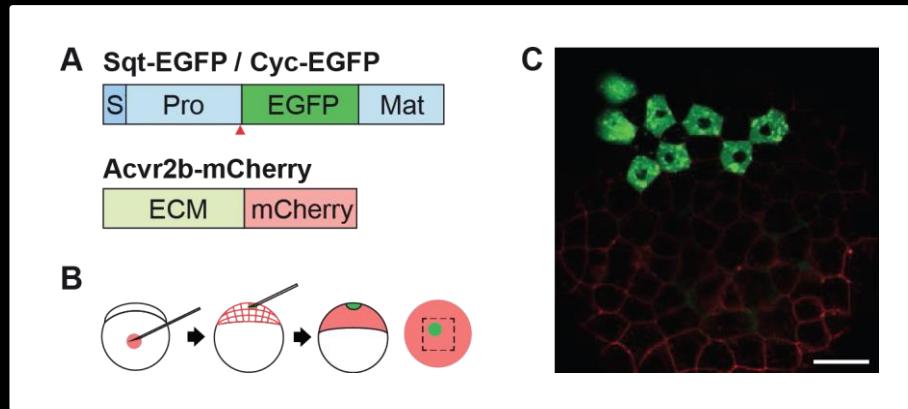
How to determine the K_d



$$K_d = \frac{[G][R]}{[GR]} \longrightarrow [G][R] = K_d [GR]$$



Nodal/Acvr2b interactions in live zebrafish



Examples of Applications

Membrane proteins: EGFR dimerization

Liu et al., *Biophys. J.* (93): 684-698 (2007).

Yavas et al., *Biophys. J.* 111(10) - 2241-2254 (2016)

Membrane and cytosolic proteins: EGFR activation

Ma et al *Front. Biosci.* Jan 1;3:22-32 (2011).

Cytosolic protein (cdc42 and effectors: IQGAP1, N-WASP etc.)

Shi et al., *Biophys. J.* (97)2:678-686 (2009).

Sudhaharan et al., *JBC* 284: 13602-13609 (2009).

Protein - DNA: Sox2/Oct4 DNA motif binding and cooperativity

EMBO Reports 2015 16(9)p1177

bioarxiv 052530

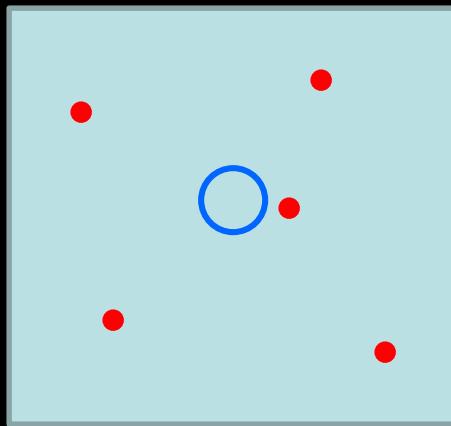
Nuclear proteins: K_ds of kinetochore protein interactions (CENPs)

Hoischen et al. 2018 PLoS ONE. 13: e0192572–26.

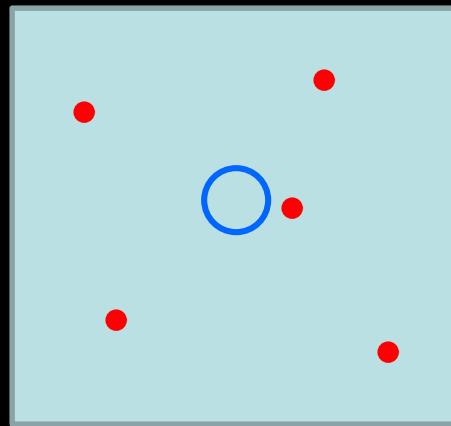
Some FCS limitations and solutions

Scanning FCS: The problem of immobile particles

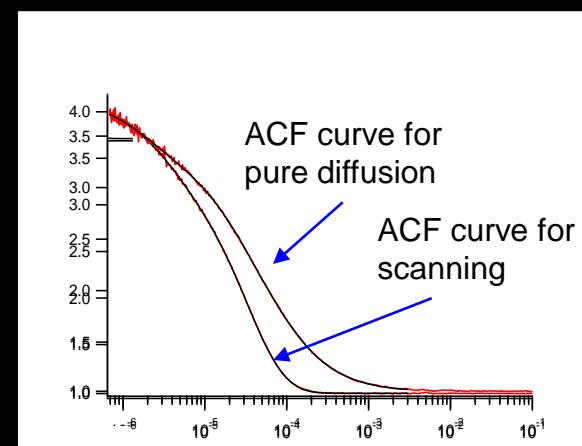
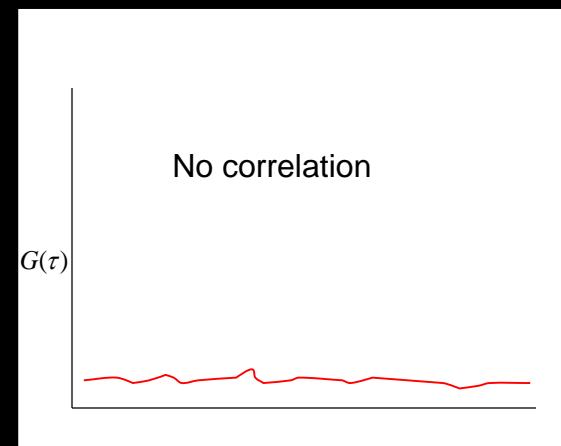
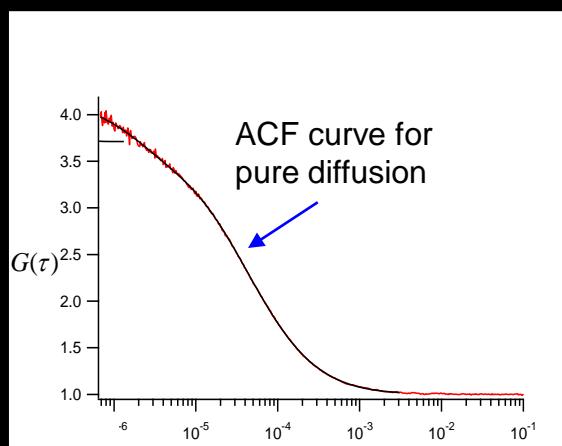
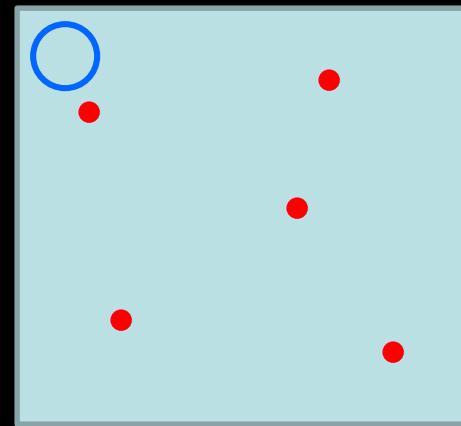
Moving particles



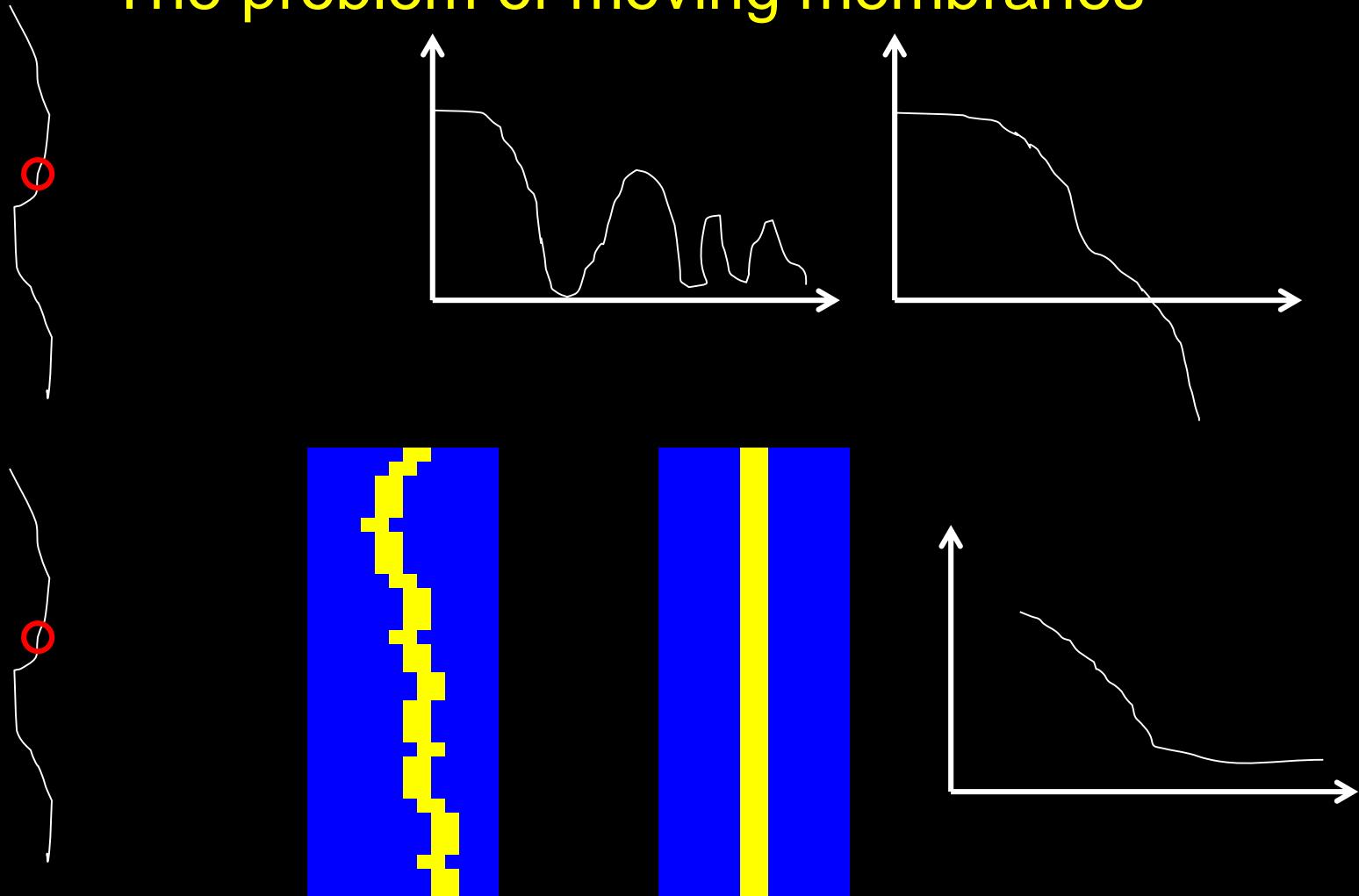
Immobile particles



Scanning Beam

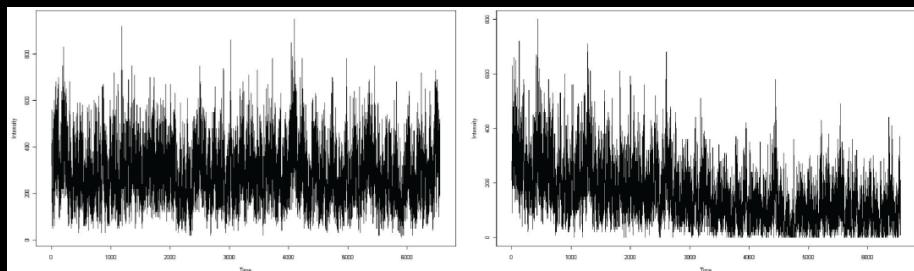


Scanning FCS: The problem of moving membranes

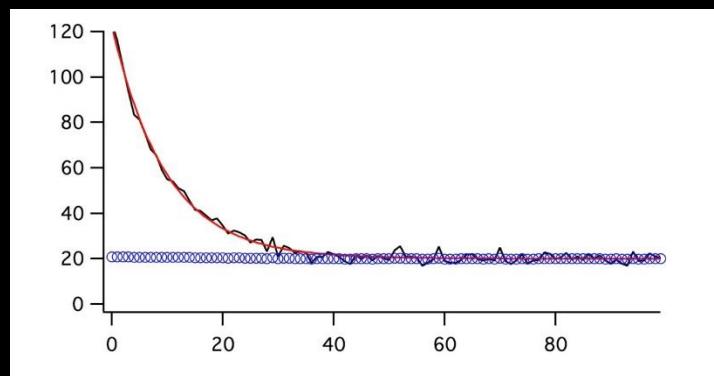
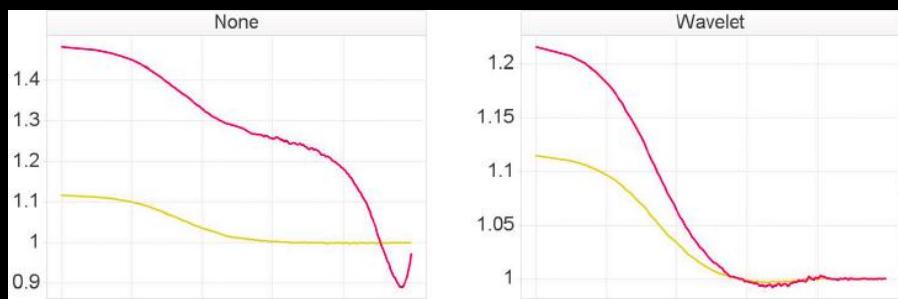
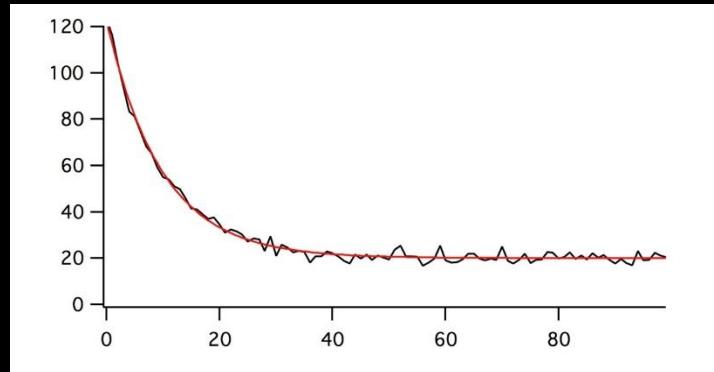


Bleach Correction

Wavelet Shrinking



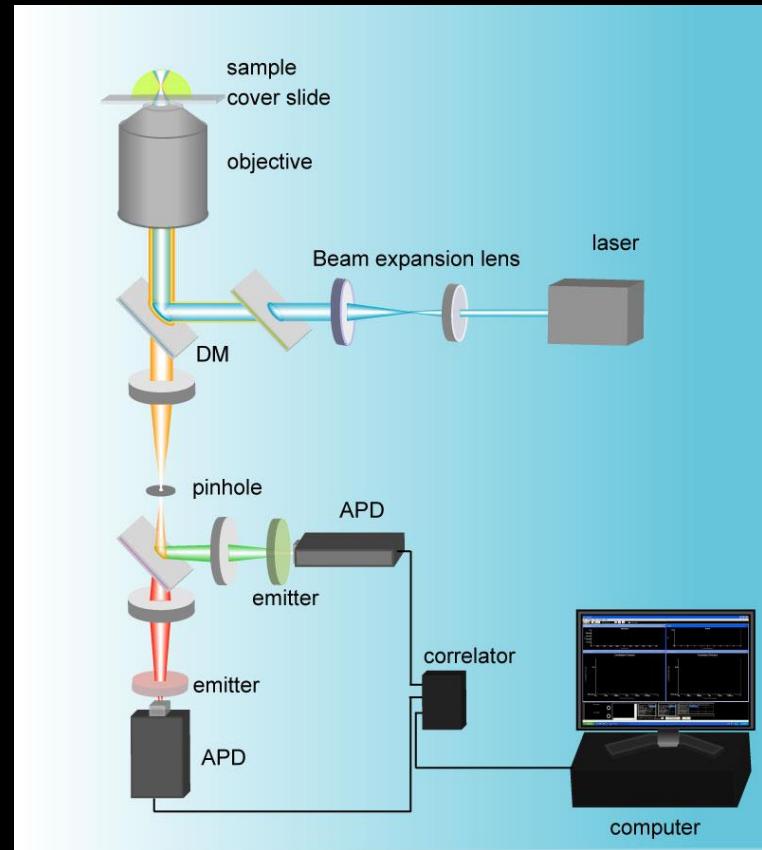
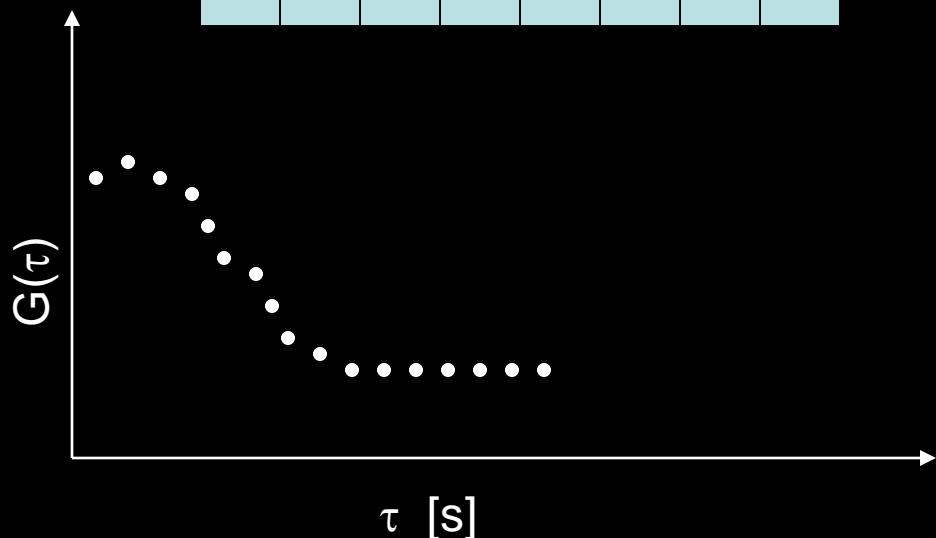
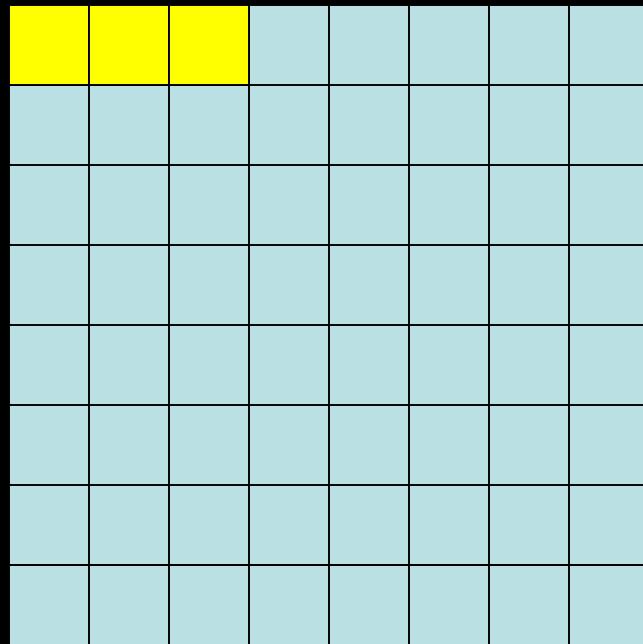
Fitting of a multi-exponential decay



Lange, J.J. et al. 2013. In: Enderlein J, et al. editors. SPIE. pp. 859006–859006–15.

Imaging FCS

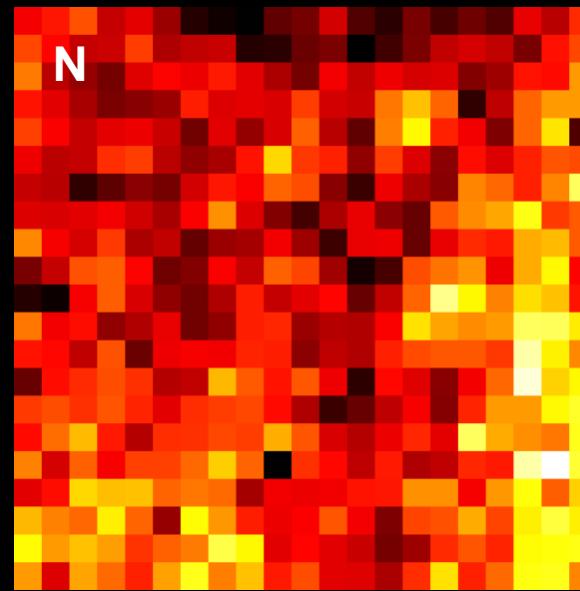
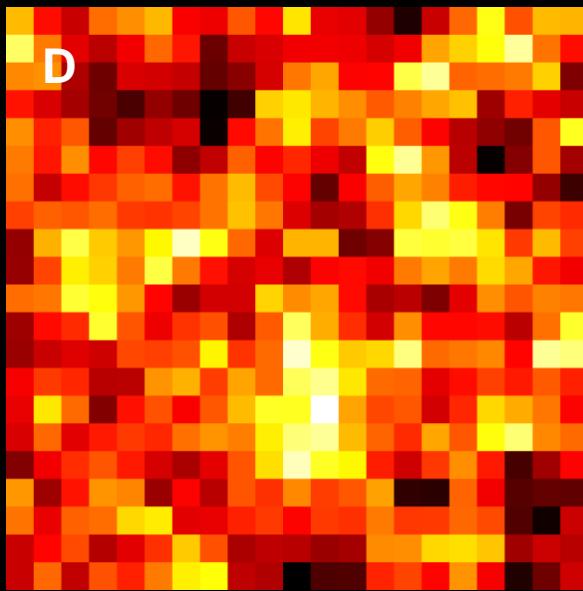
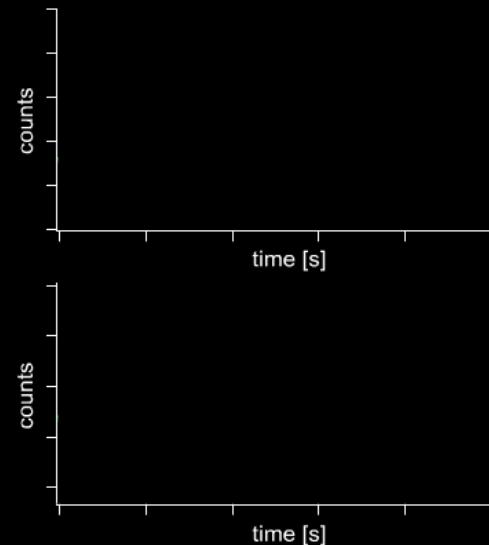
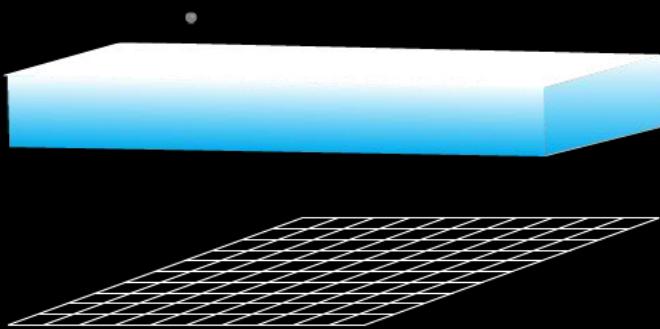
FCS in a confocal system



1) Measurements are not simultaneous

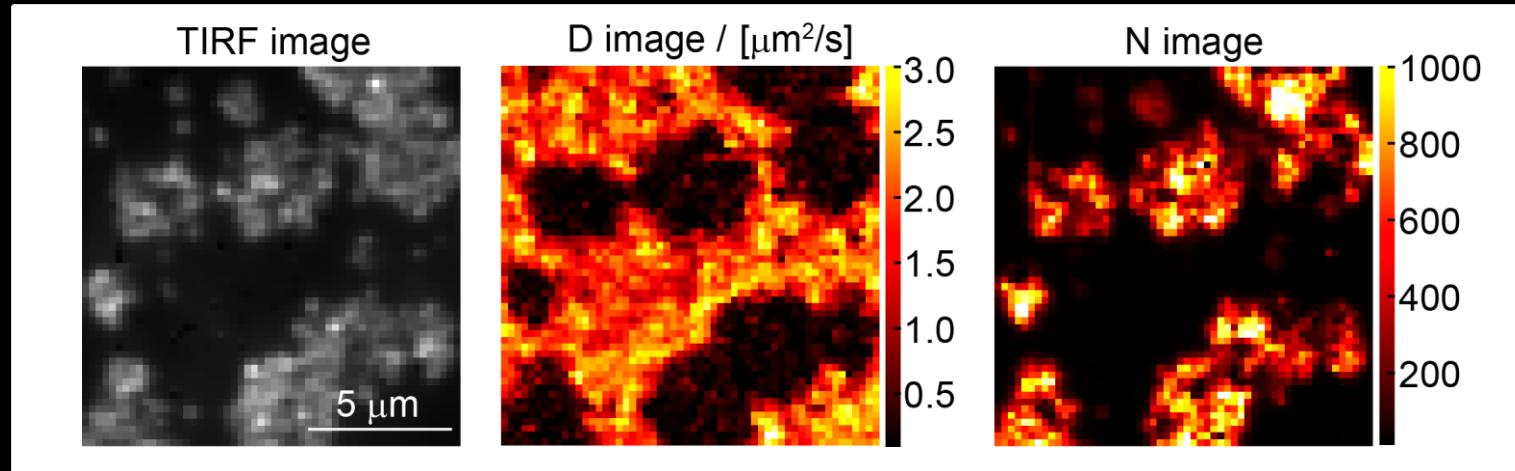
2) Cell damage by long illumination times

Imaging FCS

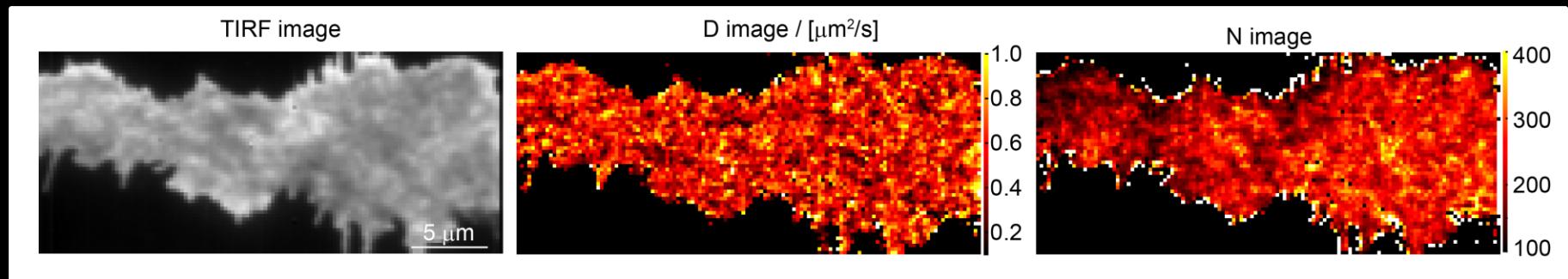


Examples

DLPC/DSPC bilayer on glass

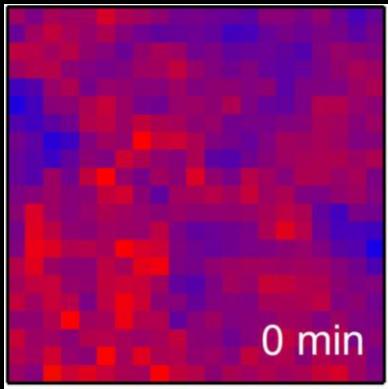


GFP-GPI on SH-SY5Y cells

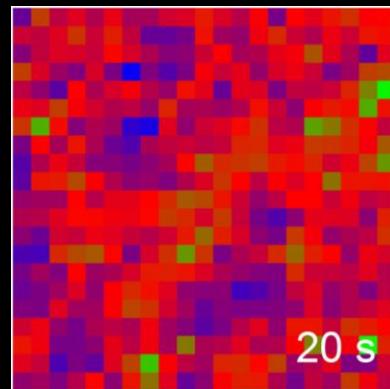


FCS videos

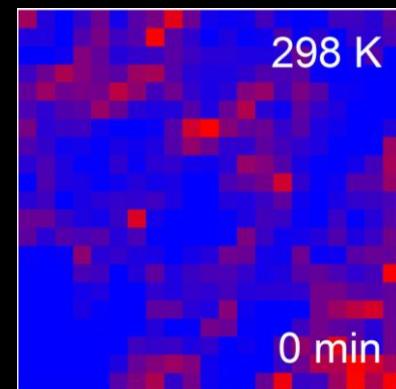
hIAPP treated Dil-C18
labelled SH-SY5Y cell



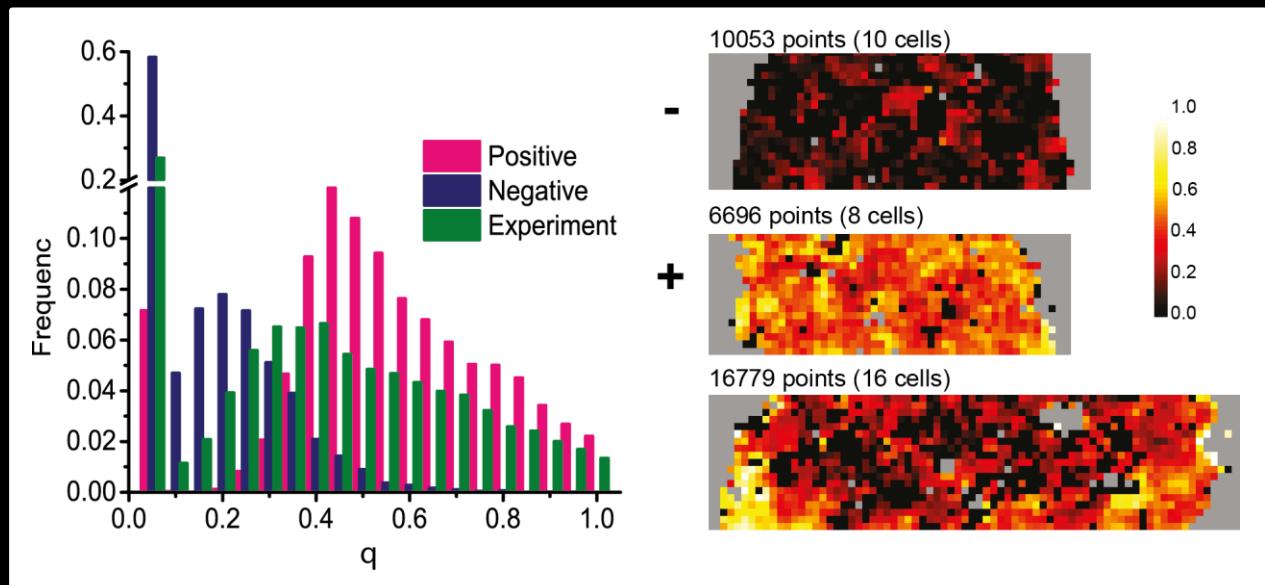
RhoPE labelled DOPC
bilayer



GFP-GPI transfected SH-SY5Y
cell at different temperature

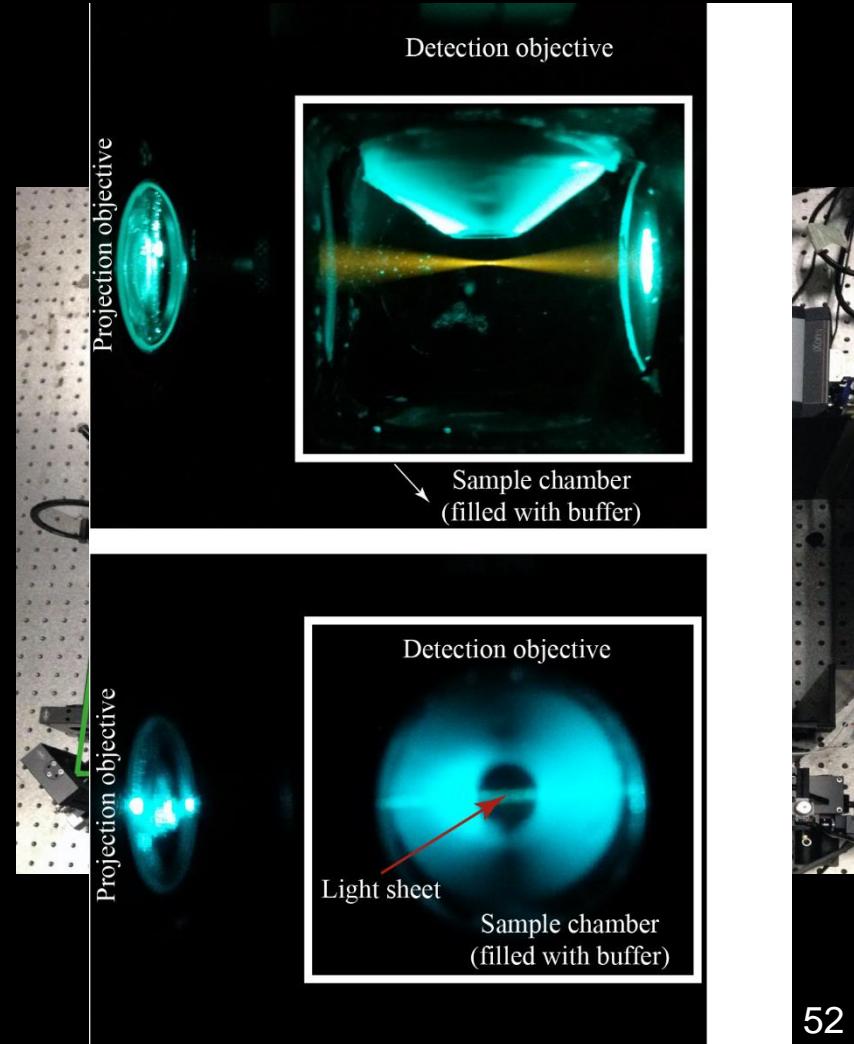
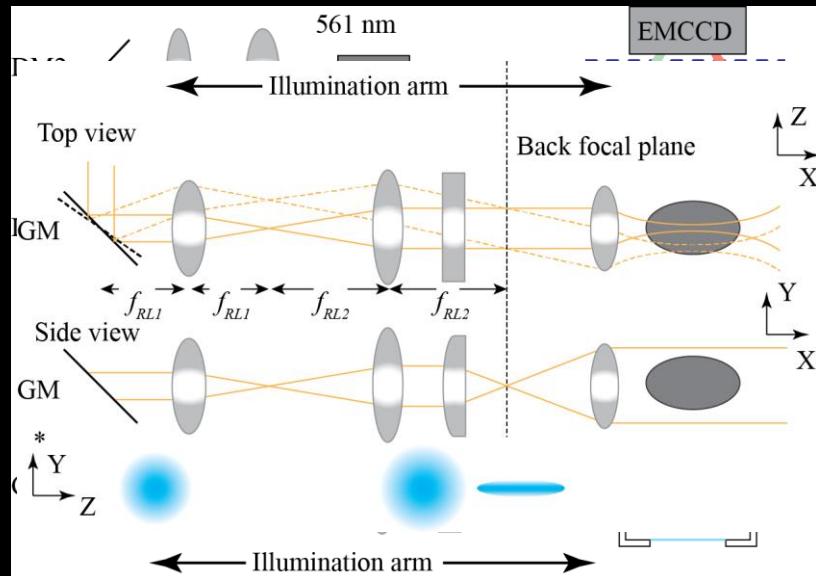


Imaging FCCS on EGFR



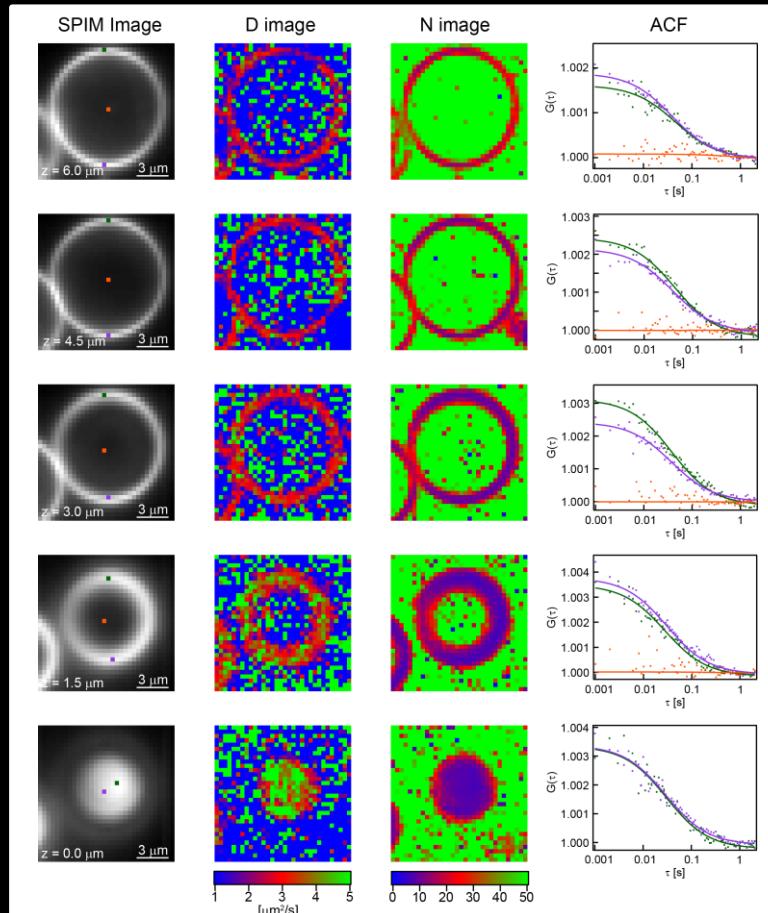
Degree of dimerization
 $q = G_{GR}(0)/\text{Min}\{G_G(0), G_R(0)\}$

Single Plane Illumination Microscopy (SPIM)



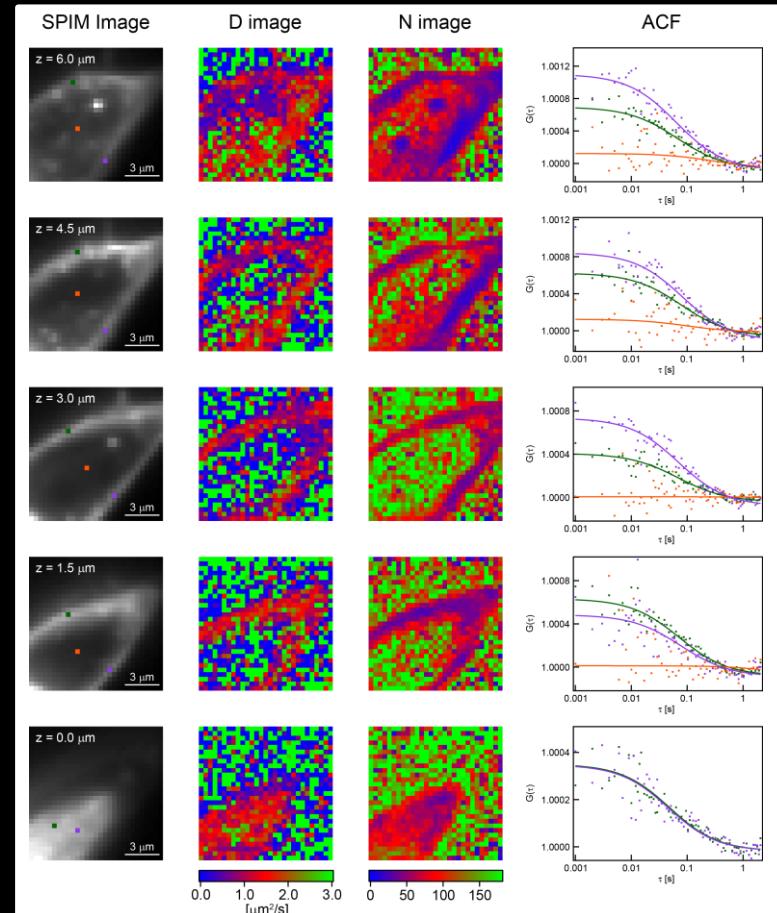
SPIM-FCS in 3D

Giant Unilamellar Vesicles (GUV)



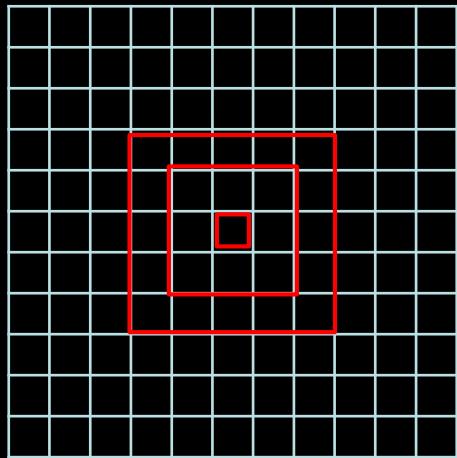
RhoPE-labelled DOPC:DOPG (10:1)

Dil-C₁₈ labelled live SH-SY5Y cell



Imaging FCS diffusion law

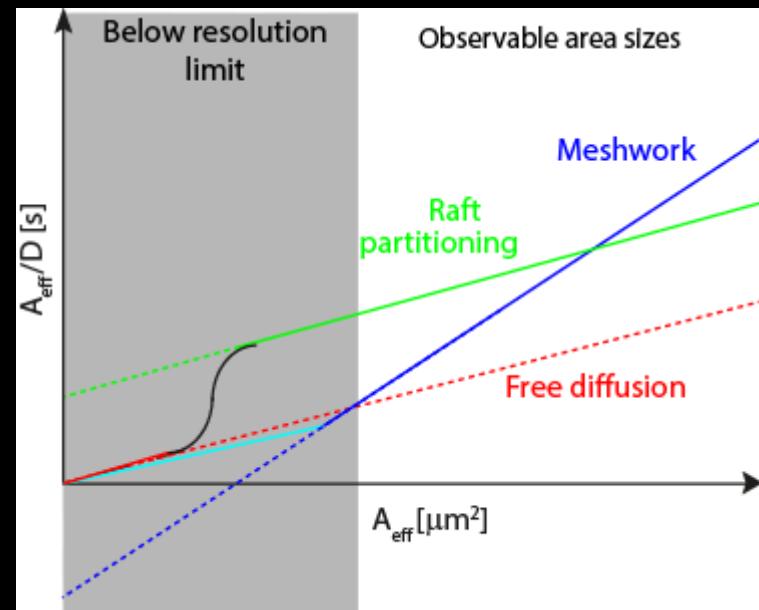
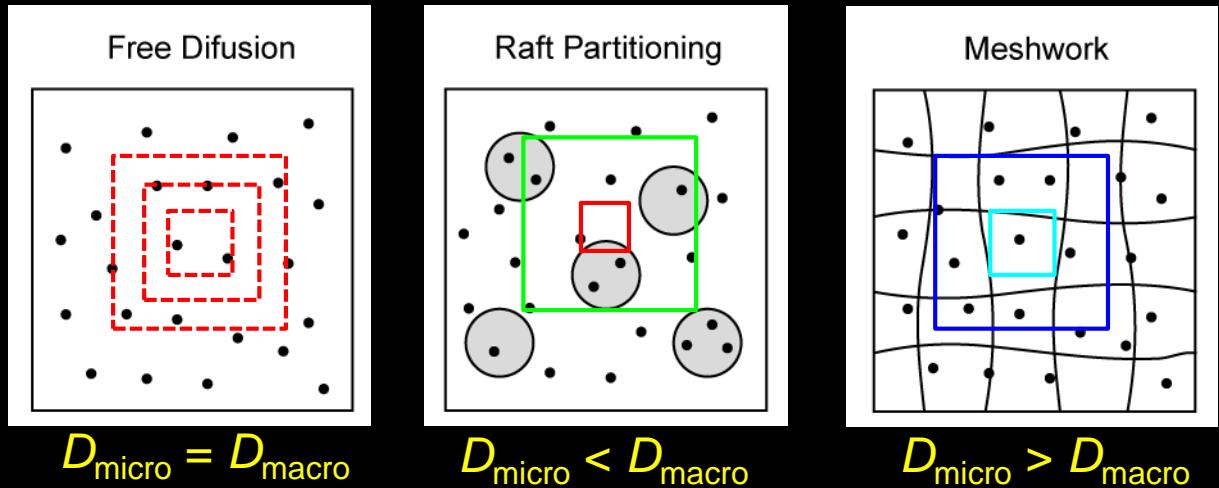
Pixel binning



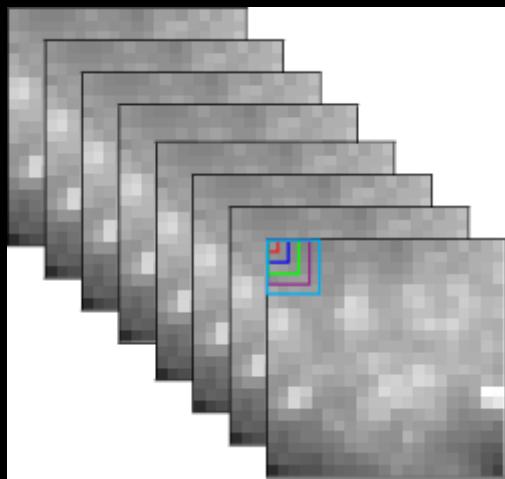
EMCCD chip

$$\tau_D(A_{eff}) = \tau_0 + \frac{A_{eff}}{D}$$

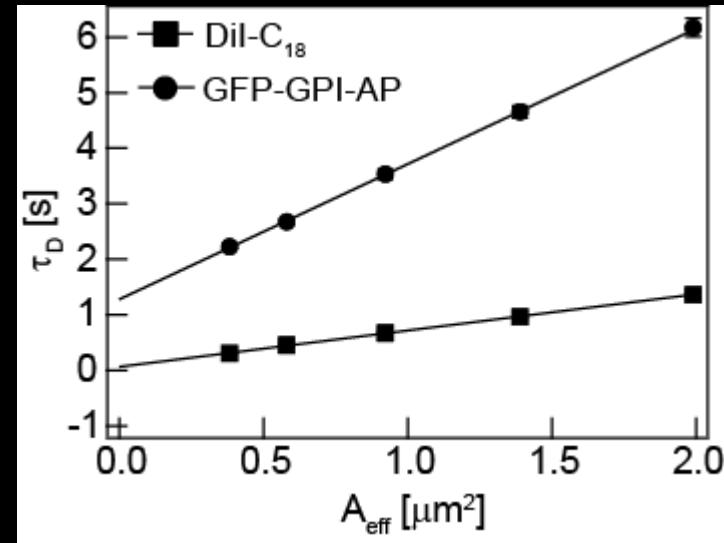
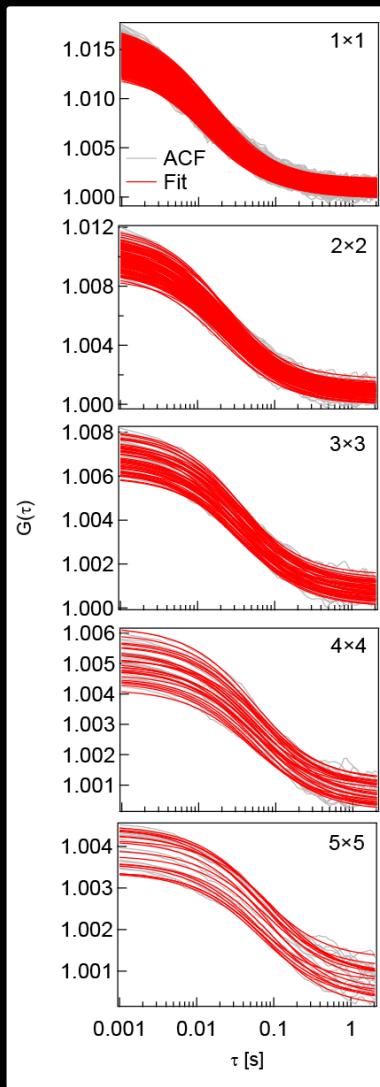
Wawrzinieck *et al.* *Biophysical Journal*, 2005
Huang and Pralle, *arXiv:1101.5087*



Imaging FCS diffusion law

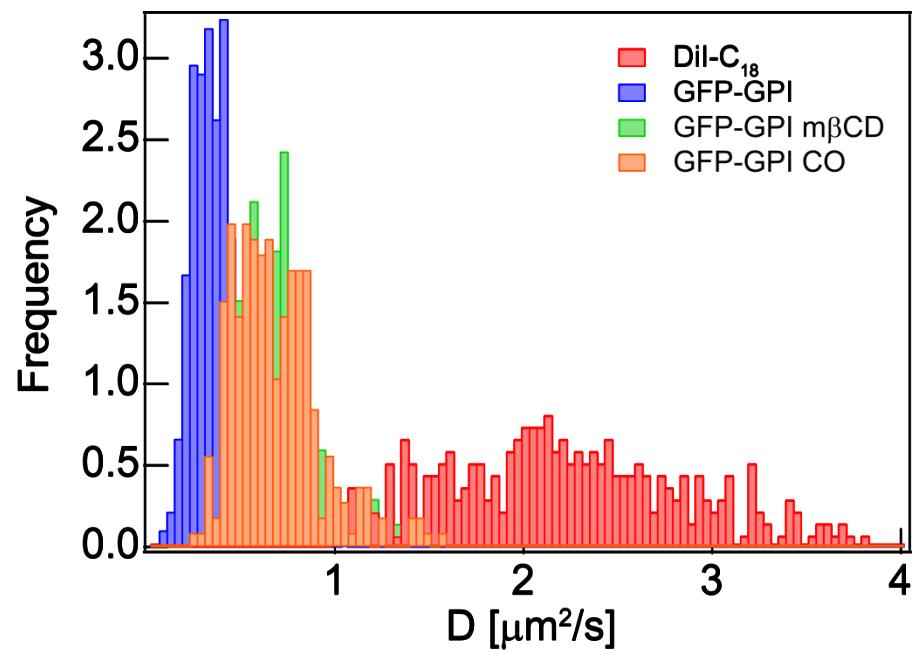
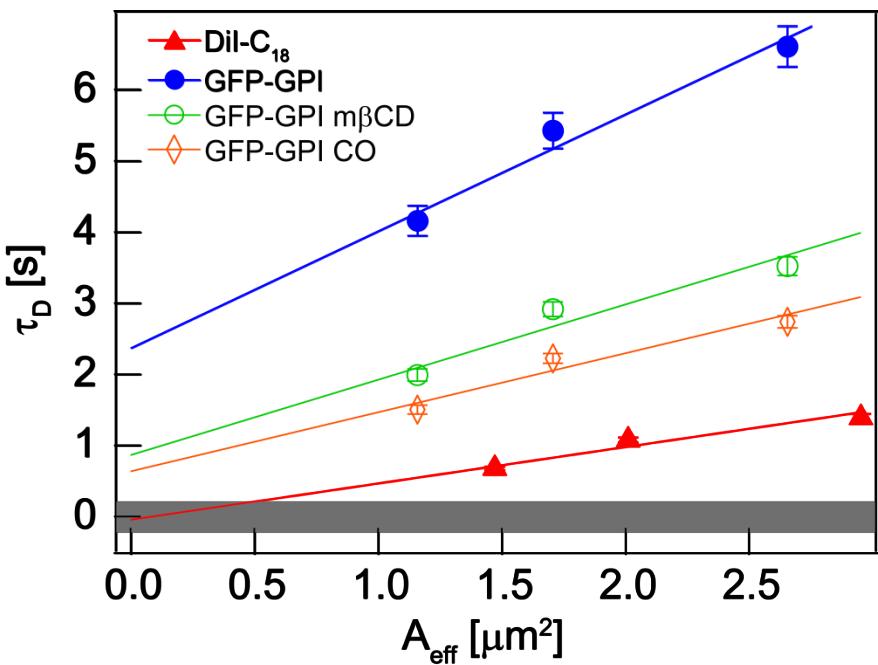


$$\tau_D(A_{eff}) = \tau_0 + \frac{A_{eff}}{D_{eff}}$$

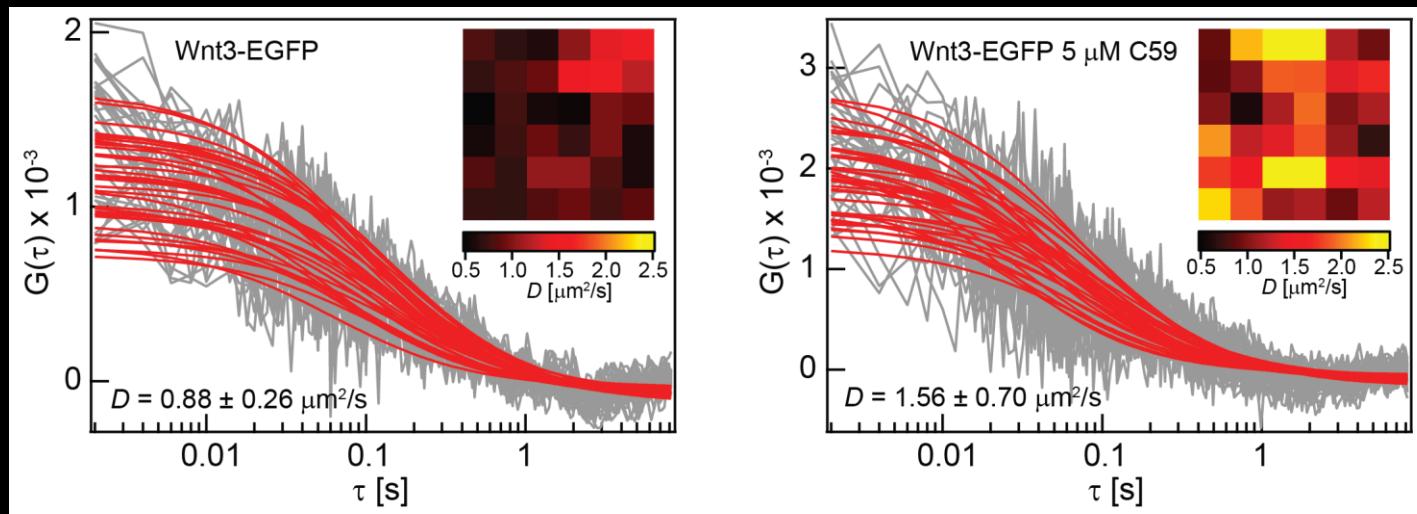
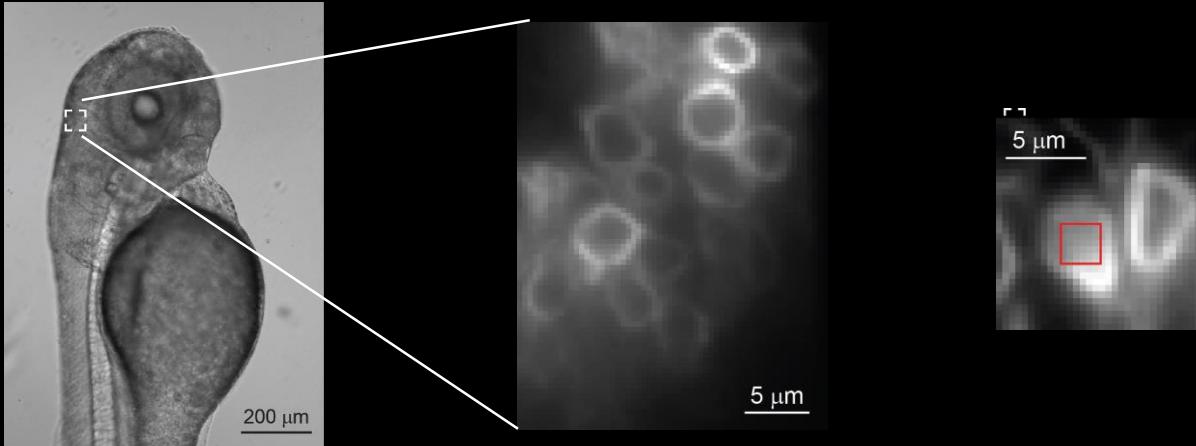


Diffusion mode	τ_0 [s]	D_{eff}/D_{ACF}
Free	0	1
Domain confined	> 0	> 1

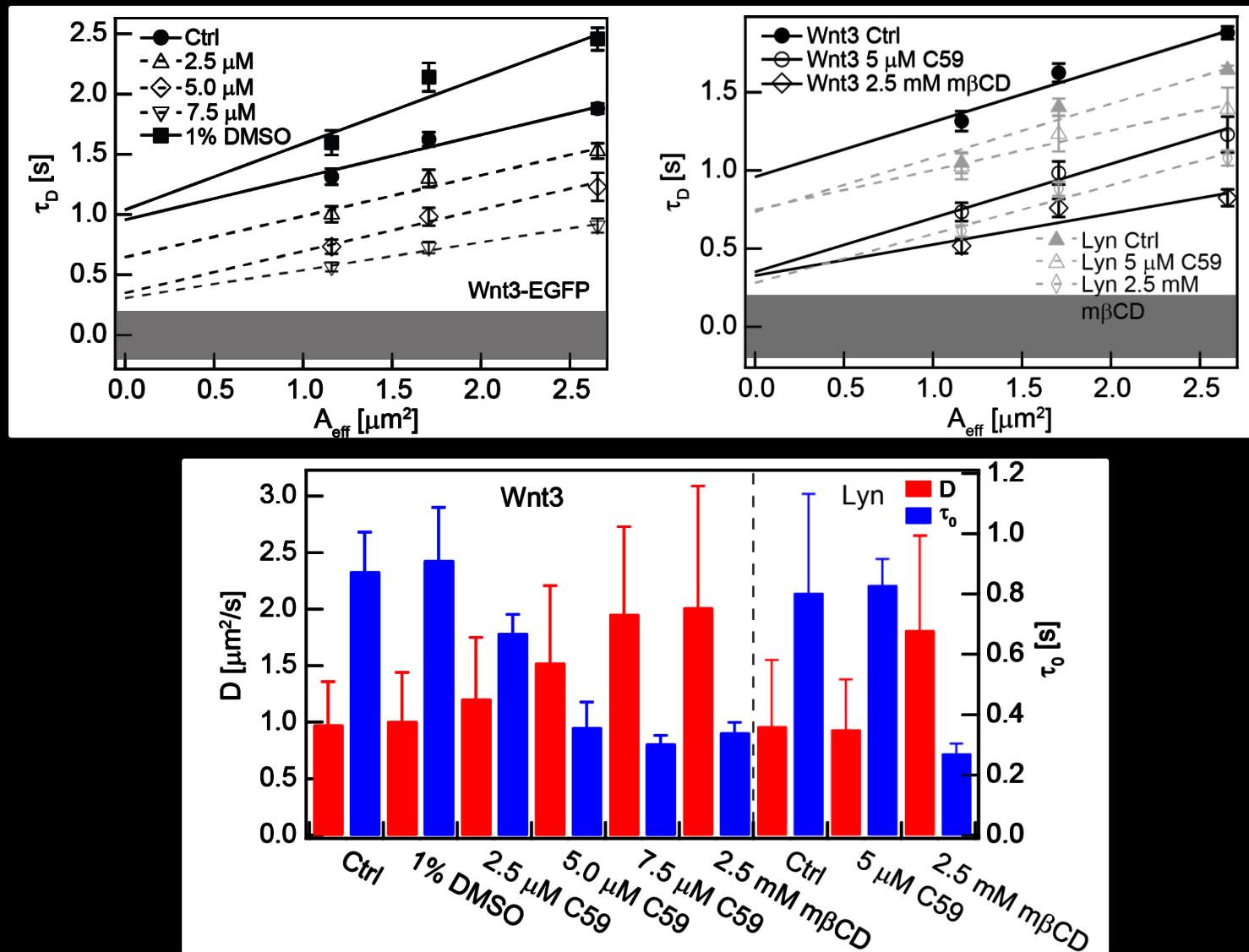
SH-SY5Y membrane organization



Wnt3 membrane localization in zebrafish

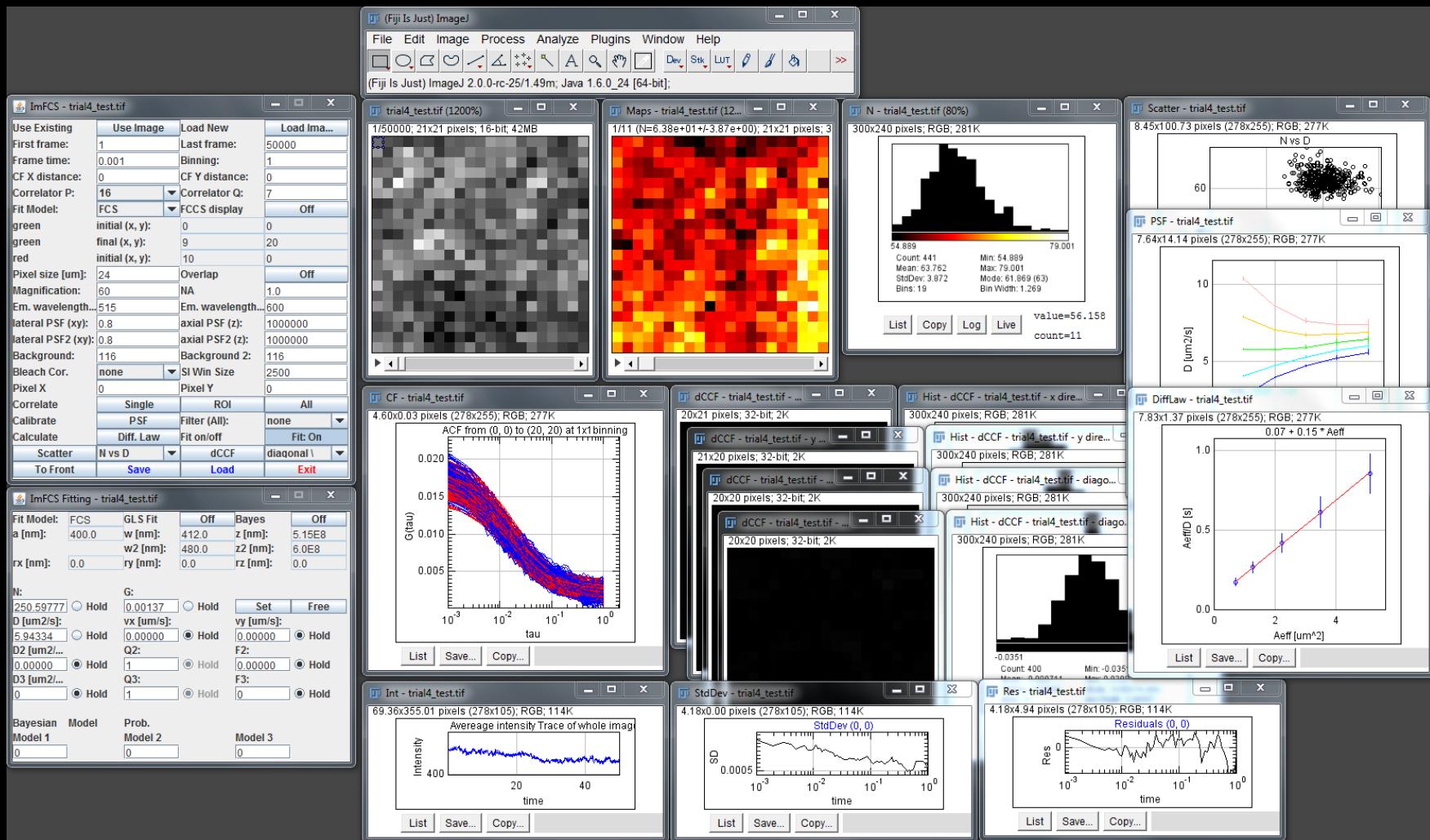


Inhibition of palmitoylation



ImageJ Plugin for Imaging FCS

<http://dbs.nus.edu.sg/lab/BFL>



Sankaran et al. *Opt. Exp.* 2010, 18 (24): 25468-25481

Bag and Wohlhand, *Ann. Rev. Phys. Chem.* 2014, 65: 225-48

Krieger et al. *Nat. Prot.* 2015, 10 (12) 1948-1972

Summary

- FCS provides measures for concentrations and diffusion coefficients
- These parameters can be quantified and can be used to derive secondary parameters (affinity, stoichiometry etc.)
- (SW-) FCCS provides an easy readout for interactions via ACF and CCF amplitudes
- Imaging FCS multiplexes FCS and FCCS measurements and can be used to make time lapse FCS videos
- TIRF and SPIM modes provide high S/N 2D and 3D measurements, respectively
- The spatiotemporal information in imaging FCS provides information beyond the diffraction limit via the diffusion laws

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Singapore Stem Cell Consortium (SSCC)



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