

# **Introduction to Fluorescence Correlation Spectroscopy (FCS)**

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***Principles of Fluorescence Techniques***  
***Genova, Italy***  
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# Fluorescence Parameters & Methods

1. Excitation & Emission Spectra
  - Local environment polarity, fluorophore concentration
2. Anisotropy & Polarization
  - Rotational diffusion
3. Quenching
  - Solvent accessibility
  - Character of the local environment
4. Fluorescence Lifetime
  - Dynamic processes (nanosecond timescale)
5. Resonance Energy Transfer
  - Probe-to-probe distance measurements
6. Fluorescence microscopy
  - localization
7. Fluorescence Correlation Spectroscopy
  - Translational & rotational diffusion
  - Concentration
  - Dynamics

# First Application of Correlation Spectroscopy

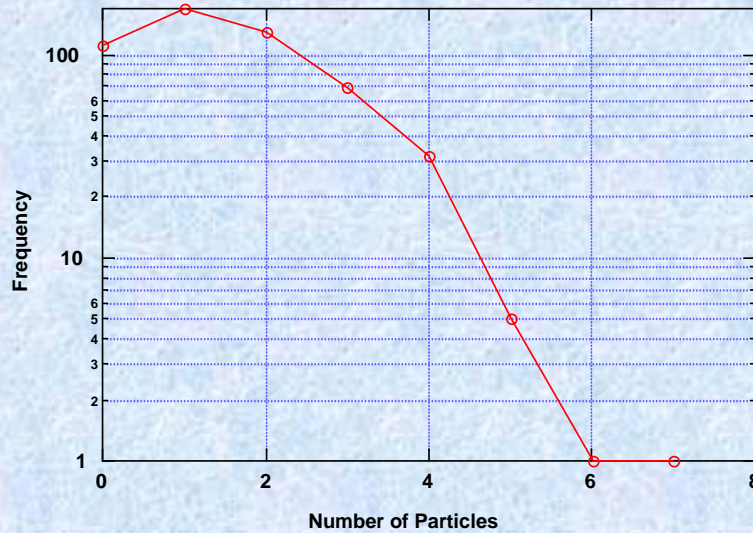
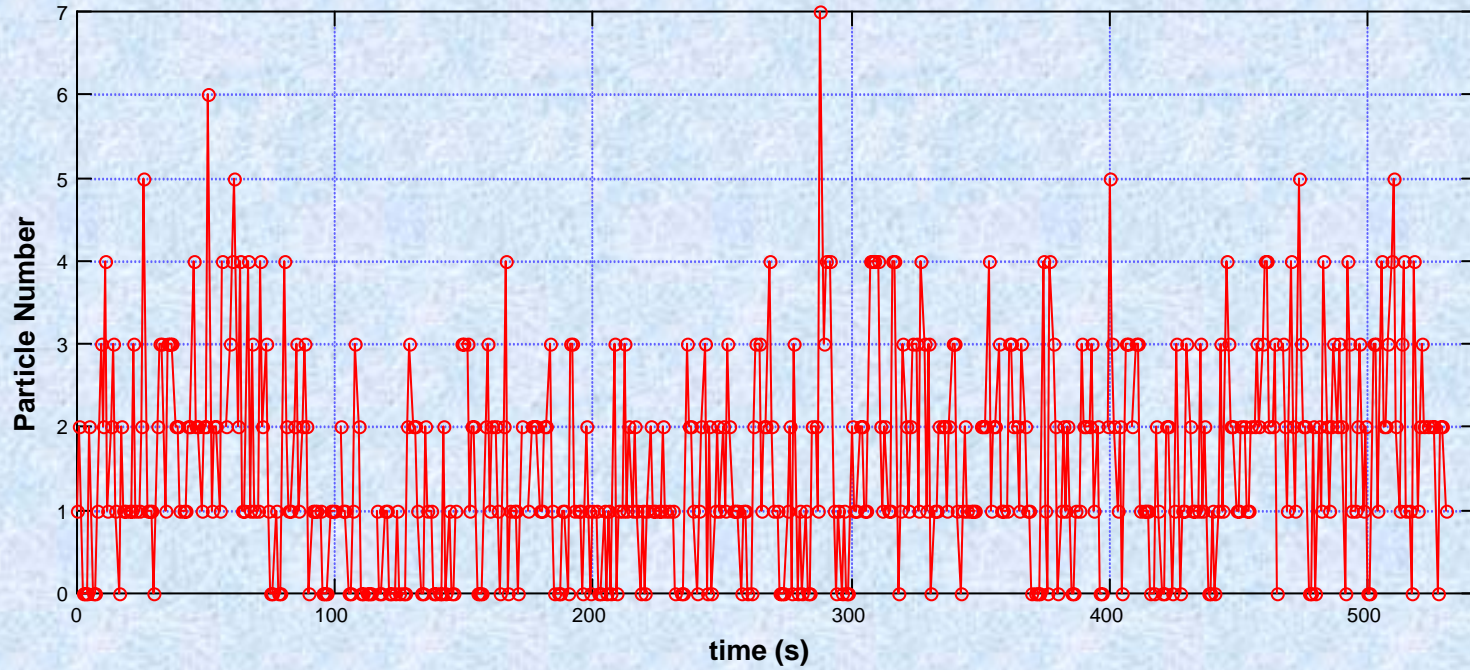
(Svedberg & Inouye, 1911) *Occupancy Fluctuation*

## Experimental data on colloidal gold particles:

```
120002001324123102111131125111023313332211122422122612214
2345241141311423100100421123123201111000111_2110013200000
10011000100023221002110000201001_333122000231221024011102_
1222112231000110331110210110010103011312121010121111211_10
003221012302012121321110110023312242110001203010100221734
410101002112211444421211440132123314313011222123310121111
222412231113322132110000410432012120011322231200_253212033
233111100210022013011321113120010131432211221122323442230
321421532200202142123232043112312003314223452134110412322
220221
```

Collected data by counting (by visual inspection) the number of particles in the observation volume as a function of time

# Particle Correlation



- \*Histogram of particle counts
- \*Poisson behavior
- \*Autocorrelation not available

**In FCS**  
**Fluctuations are in the Fluorescence Signal**

**Diffusion**

**Enzymatic Activity**

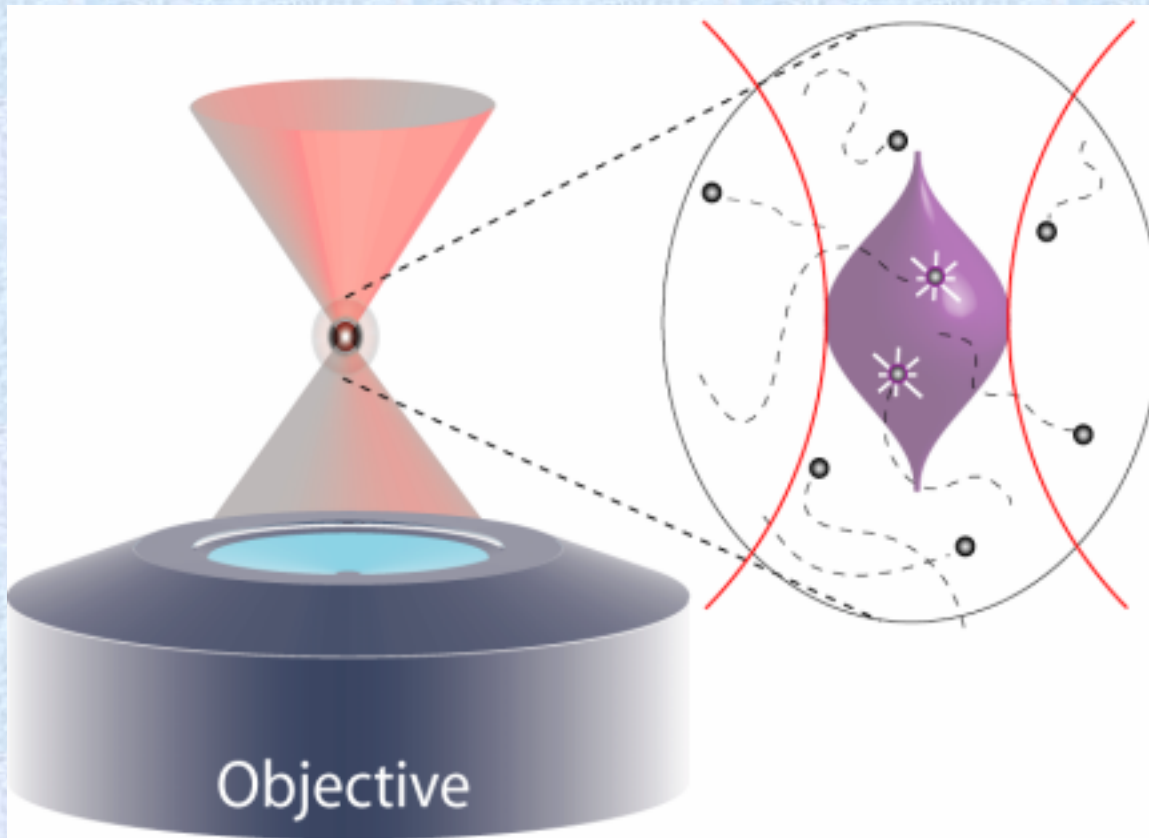
**Phase Fluctuations**

**Conformational Dynamics**

**Rotational Motion**

**Protein Folding**

# Generating Fluctuations By Motion

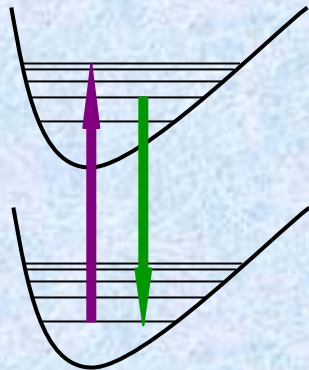


## What is Observed?

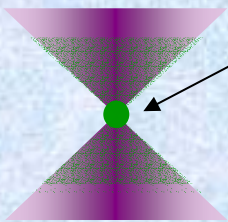
- 1. The rate of motion.**
- 2. The concentration of particles.**
- 3. Changes in the particle fluorescence while under observation.**

# Defining Our Observation Volume: One- & Two-Photon Excitation.

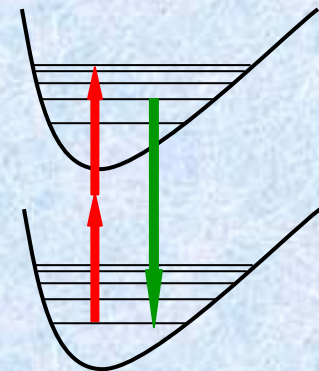
## 1 - Photon



Defined by the pinhole size,  
wavelength, magnification and  
numerical aperture of the  
objective

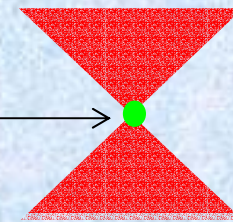


## 2 - Photon



Approximately  $1 \text{ } \mu\text{m}^3$

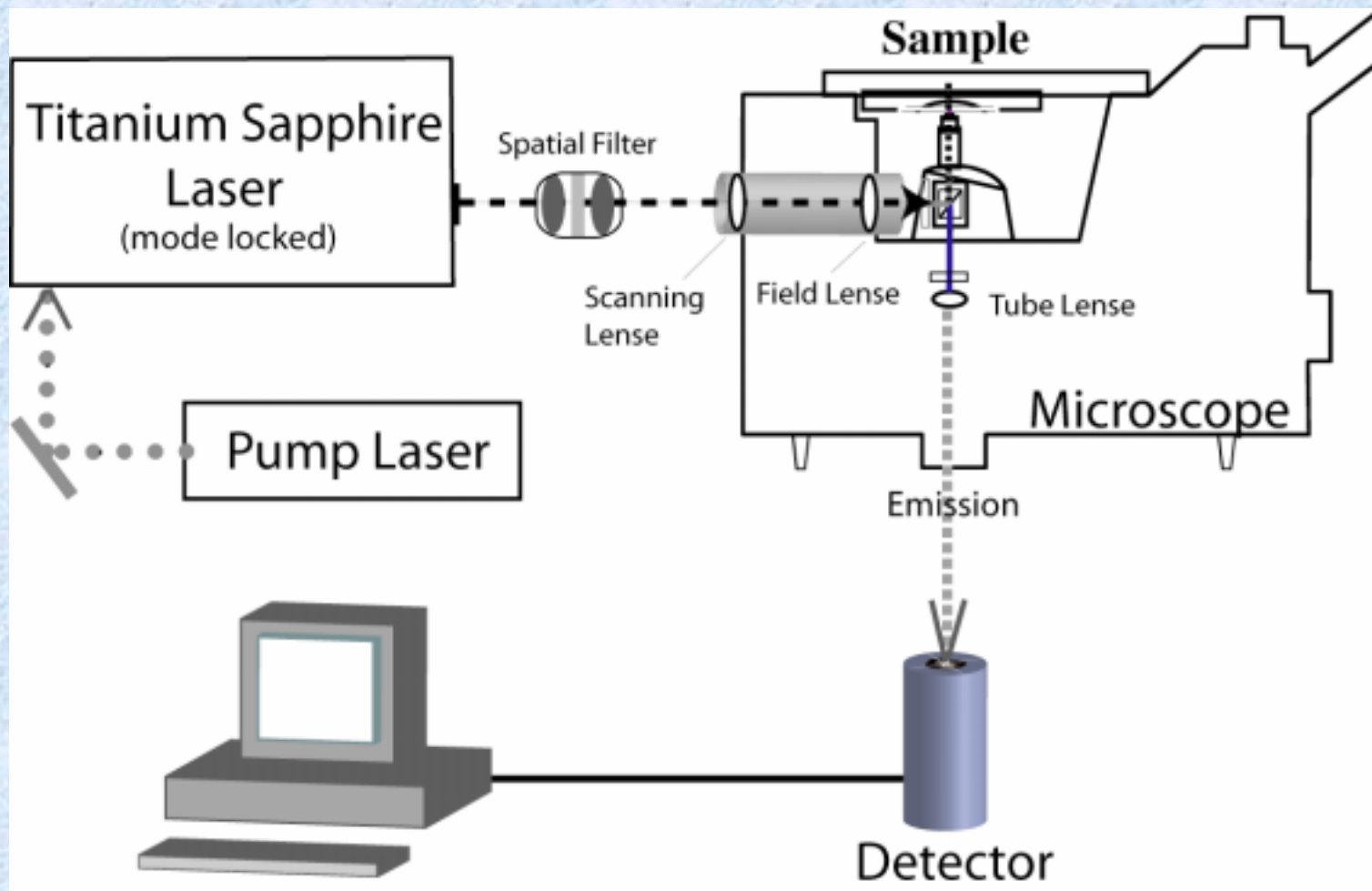
Defined by the wavelength  
and numerical aperture of the  
objective



1-photon

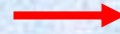
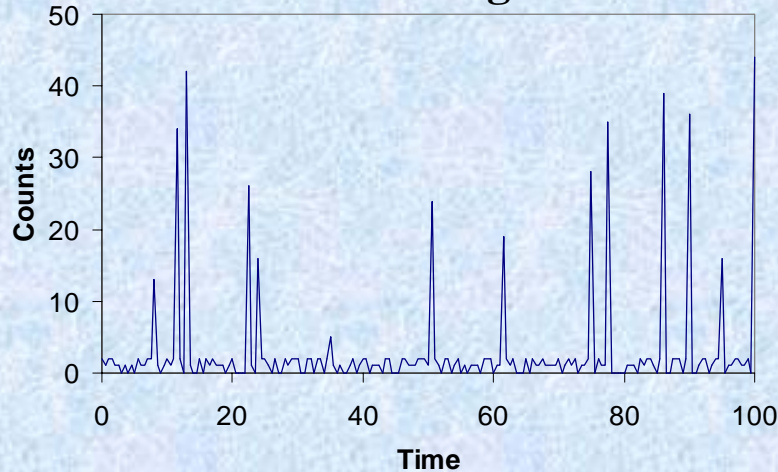


2-photon

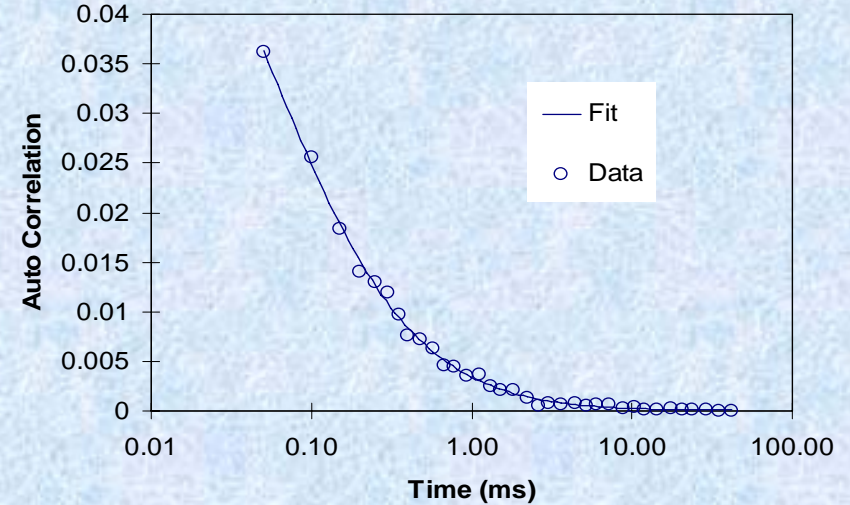


# Data Treatment & Analysis

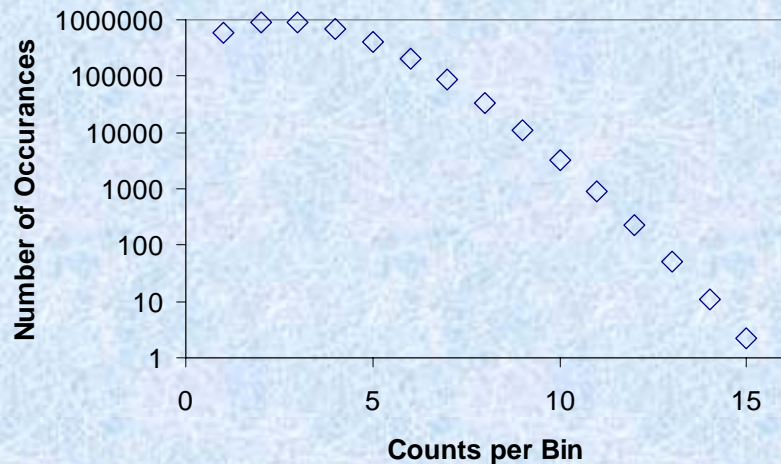
## Time Histogram



## Autocorrelation



## Photon Counting Histogram (PCH)



**Autocorrelation Parameters:  
 $G(0)$  &  $k_{\text{action}}$**

**PCH Parameters:  $\langle N \rangle$  &  $\underline{\varepsilon}$**

# Autocorrelation Function

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

$$\delta F(t) = F(t) - \langle F(t) \rangle$$

Factors influencing the fluorescence signal:

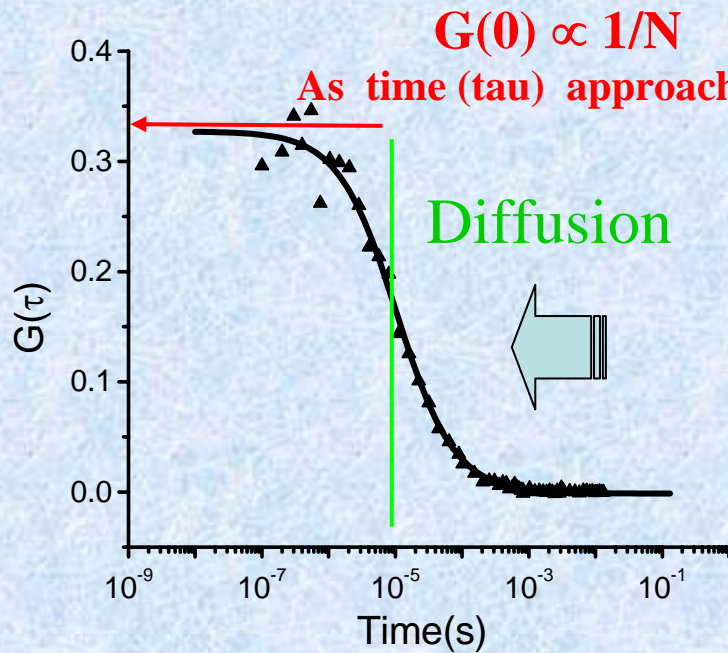
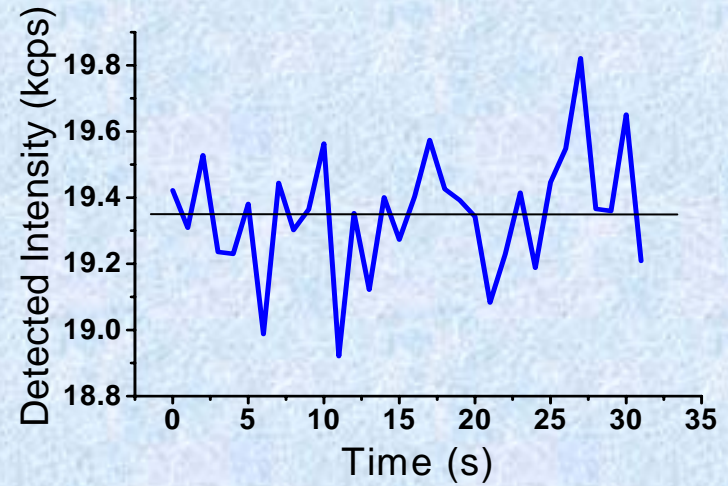
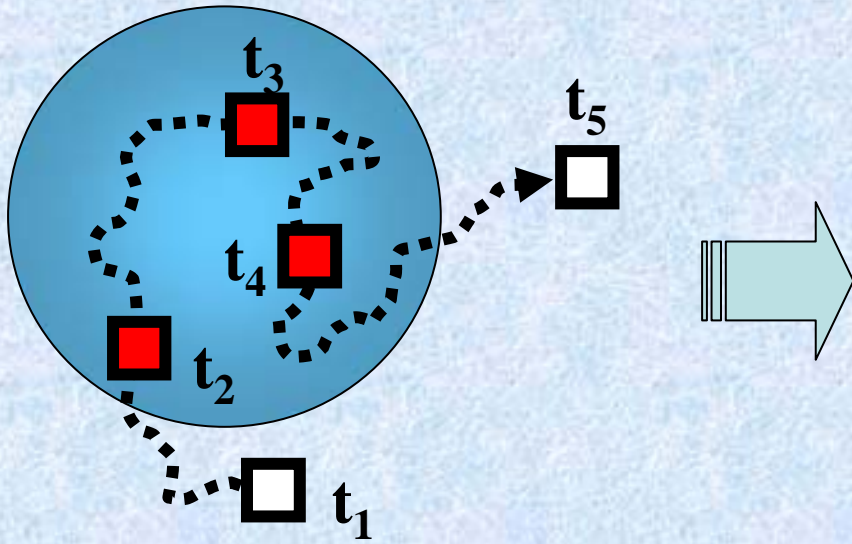
$$F(t) = \kappa Q \int d\mathbf{r} W(\mathbf{r}) C(\mathbf{r}, t)$$

$\kappa Q$  = quantum yield and detector sensitivity (how bright is our probe)

$W(\mathbf{r})$  describes our observation volume

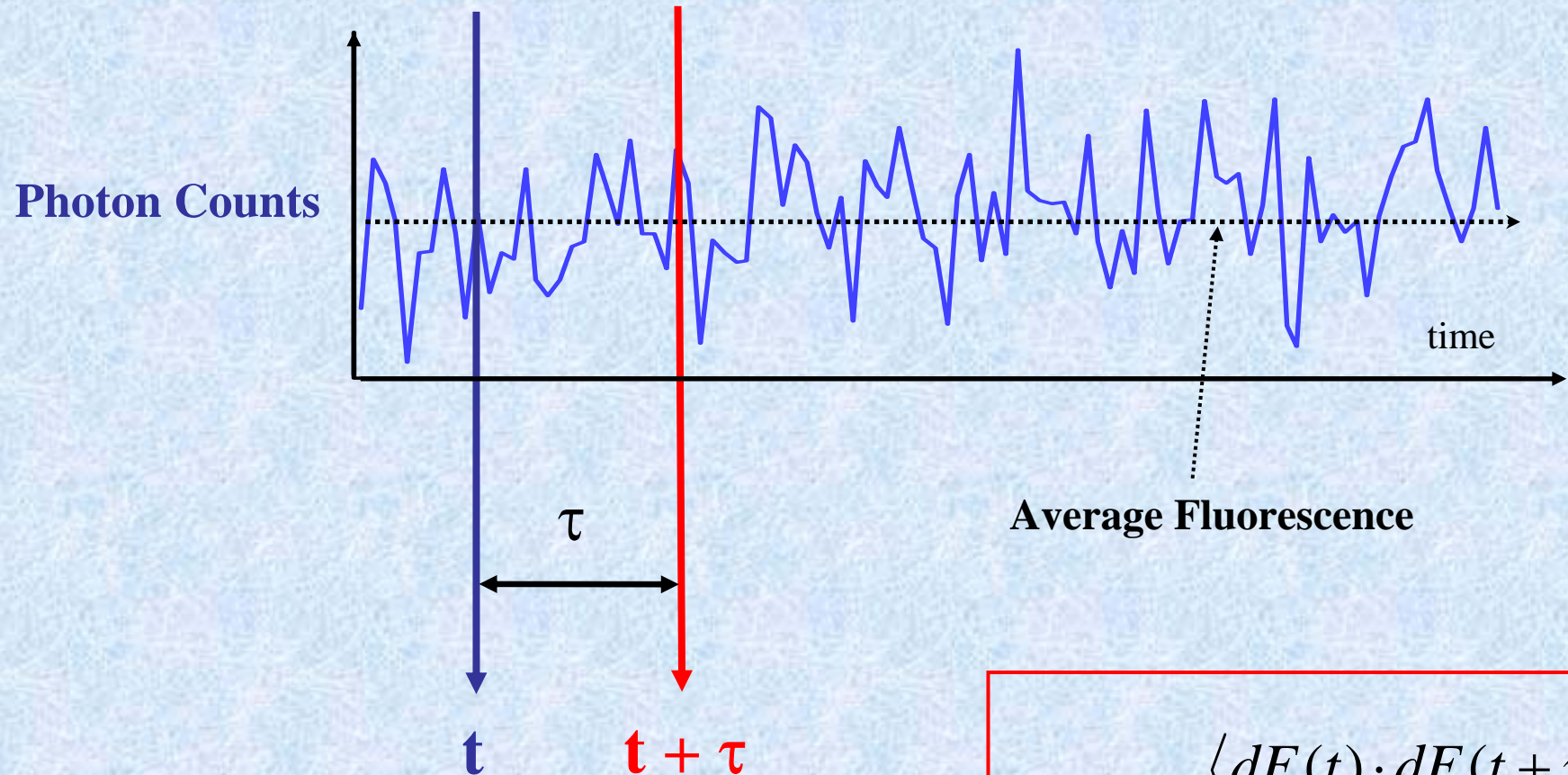
$C(\mathbf{r}, t)$  is a function of the fluorophore concentration over time

# The Autocorrelation Function



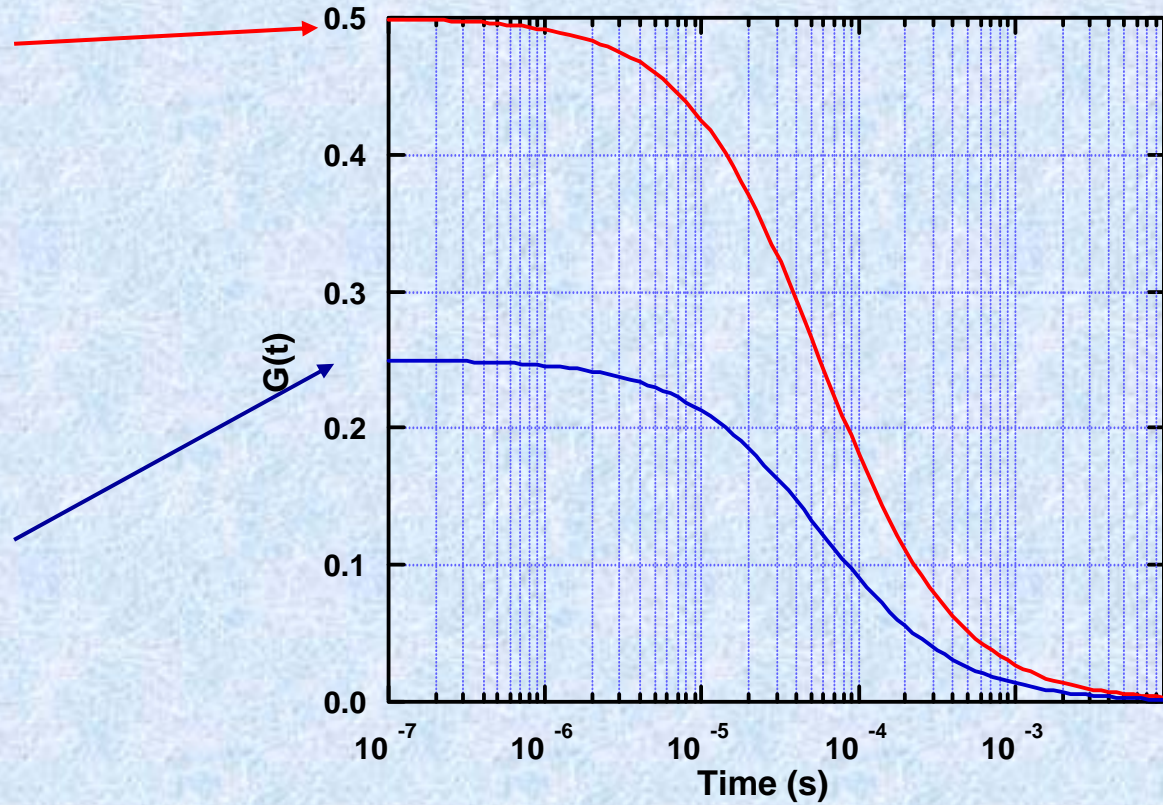
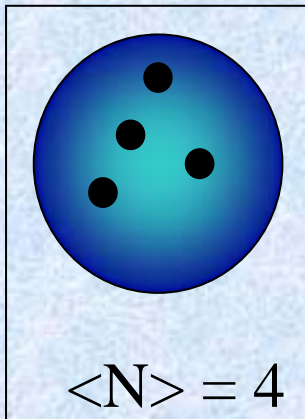
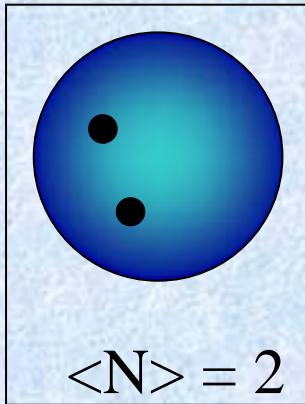
$$G(\tau) = \frac{\langle \delta F(t) \delta F(t + \tau) \rangle}{\langle F(t) \rangle^2}$$

# Calculating the Autocorrelation Function



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

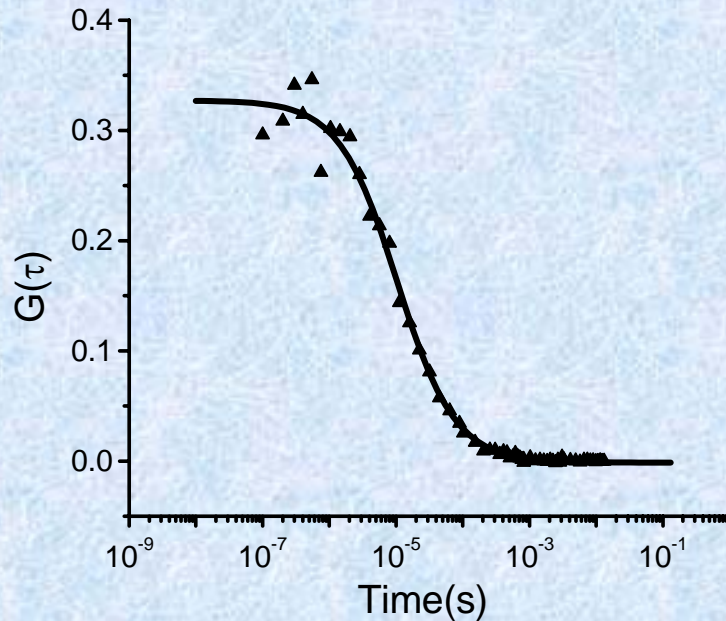
# The Effects of Particle Concentration on the Autocorrelation Curve



# Why Is $G(0)$ Proportional to $1/\text{Particle Number}$ ?

A Poisson distribution describes the statistics of particle occupancy fluctuations. In a Poissonian system the variance is proportional to the average number of fluctuating species:

$$\langle \text{Particle\_Number} \rangle = \text{Variance}$$

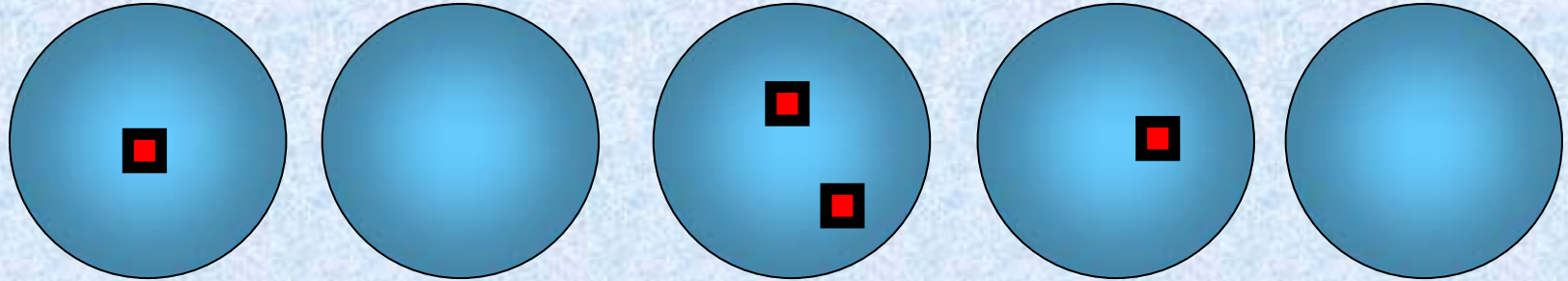


$$G(\tau) = \frac{\langle \delta F(t) \delta F(t + \tau) \rangle}{\langle F(t) \rangle^2}$$

$$G(0) = \frac{\langle \delta F(t)^2 \rangle}{\langle F(t) \rangle^2} = \frac{\langle (F(t) - \langle F(t) \rangle)^2 \rangle}{\langle F(t) \rangle^2}$$

$$G(0) = \frac{\text{Variance}}{\langle N \rangle^2} = \frac{1}{\langle N \rangle}$$

# G(0), Particle Brightness and Poisson Statistics



1 0 0 0 0 0 0 0 2 0 1 1 1 0 0 0 0 0 0 1 0 0 0 0 0 0 0 1 0 1 0 0 0 1 0 0 1 0 0

Time →

Average = 0.275

Variance = 0.256

$$\langle N \rangle \propto \frac{\text{Average}^2}{\text{Variance}} = \frac{0.275^2}{0.256} = 0.296$$

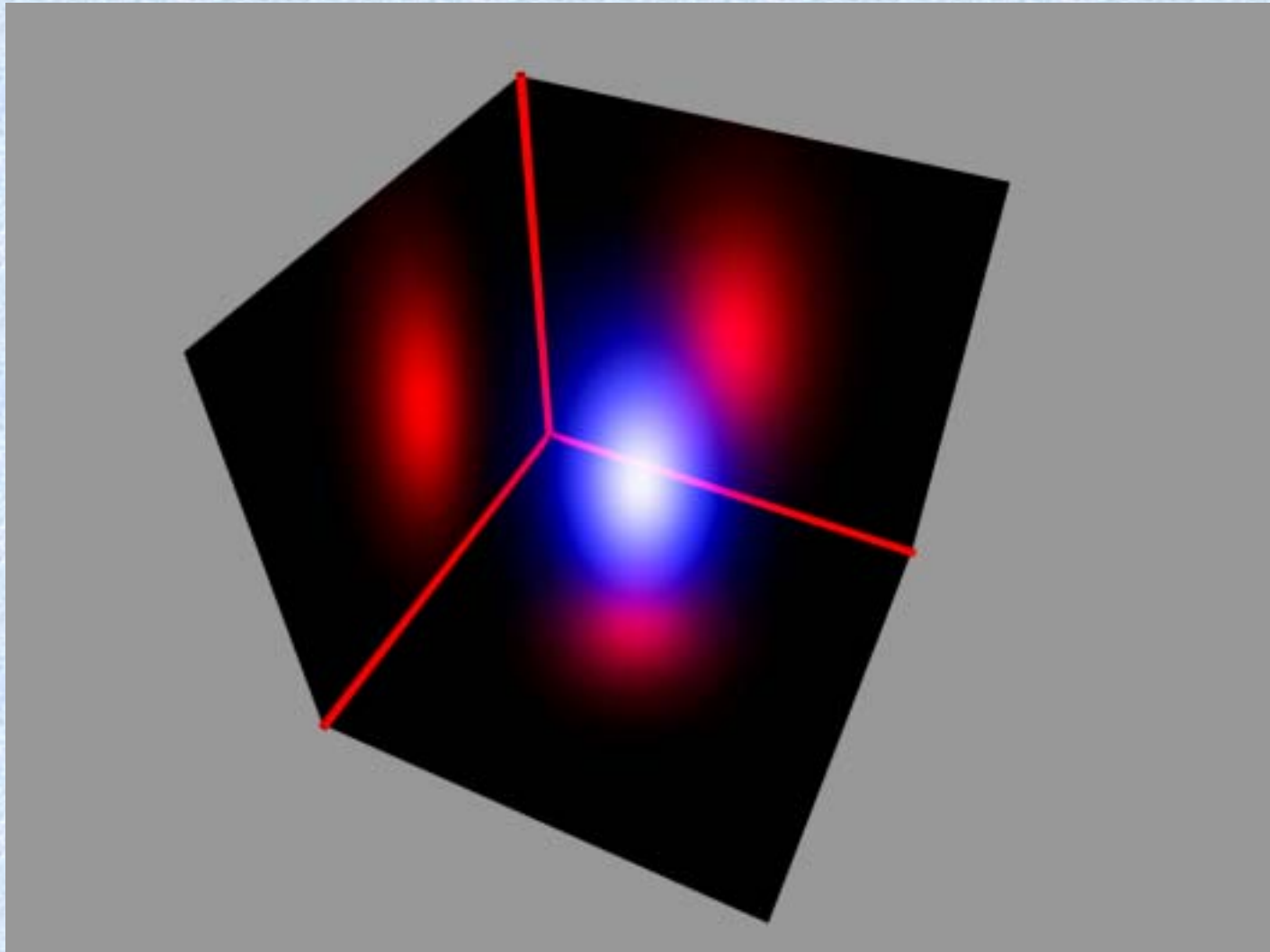
Lets increase the particle brightness by 4x:

4 0 0 0 0 0 0 0 8 0 4 4 4 0 0 0 0 0 0 4 0 0 0 0 0 0 0 4 0 4 0 0 0 4 0 0 4 0 0

Average = 1.1    Variance = 4.09

$$\langle N \rangle \propto 0.296$$

**What about the excitation (or observation) volume shape?**



# Effect of Shape on the (Two-Photon) Autocorrelation Functions:

For a 2-dimensional Gaussian excitation volume:

$$G(\tau) = \frac{\gamma}{N} \left( 1 + \frac{8D\tau}{w_{2DG}^2} \right)^{-1}$$

1-photon equation contains a 4, instead of 8

For a 3-dimensional Gaussian excitation volume:

$$G(\tau) = \frac{\gamma}{N} \left( 1 + \frac{8D\tau}{w_{3DG}^2} \right)^{-1} \left( 1 + \frac{8D\tau}{z_{3DG}^2} \right)^{-1/2}$$

## Additional Equations:

### 3D Gaussian Confocor analysis:

$$G(\tau) = \mathbf{1} + \frac{\mathbf{1}}{N} \left( \mathbf{1} + \frac{\tau}{\tau_D} \right)^{-1} \cdot \left( \mathbf{1} + S^2 \cdot \frac{\tau}{\tau_D} \right)^{-\frac{1}{2}}$$

... where  $N$  is the average particle number,  $\tau_D$  is the diffusion time (related to  $D$ ,  $\tau_D = w^2/8D$ , for two photon and  $\tau_D = w^2/4D$  for 1-photon excitation), and  $S$  is a shape parameter, equivalent to  $w/z$  in the previous equations.

### Triplet state term:

$$\left( 1 + \frac{T}{1-T} e^{-\frac{\tau}{\tau_T}} \right)$$

..where  $T$  is the triplet state amplitude and  $\tau_T$  is the triplet lifetime.

# The Effects of Particle Size on the Autocorrelation Curve

## Diffusion Constants

300  $\mu\text{m}^2/\text{s}$

90  $\mu\text{m}^2/\text{s}$

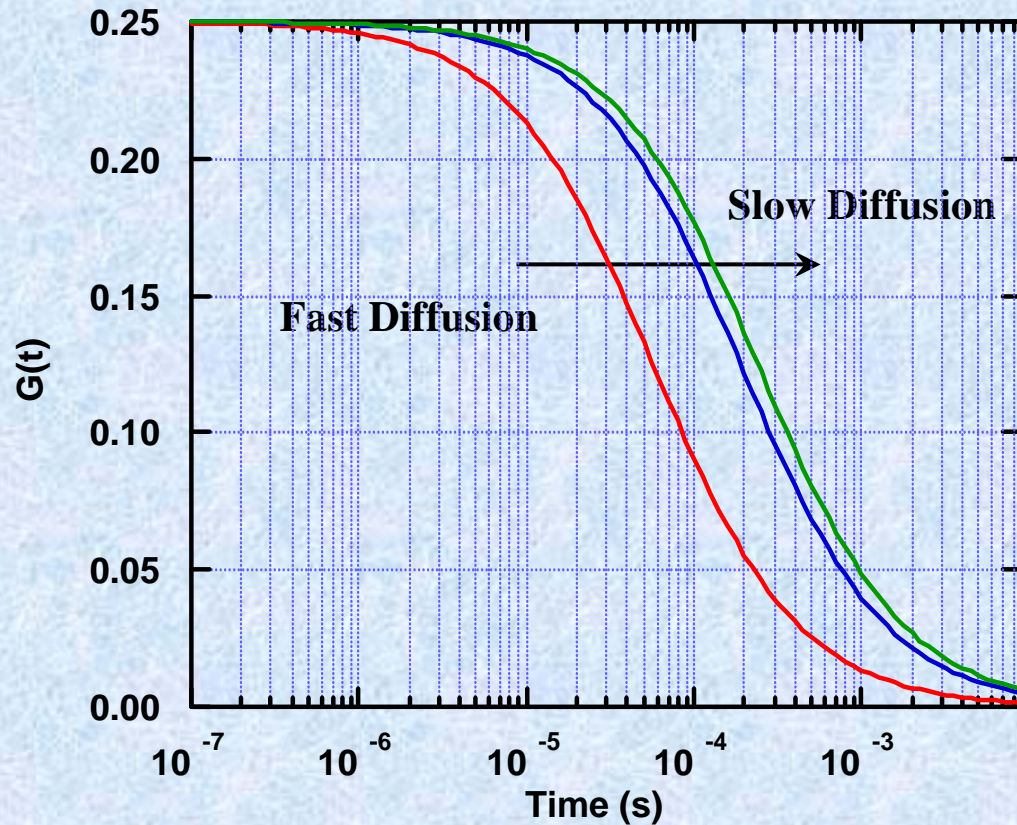
71  $\mu\text{m}^2/\text{s}$

## Stokes-Einstein Equation:

$$D = \frac{k \cdot T}{6 \cdot \pi \cdot \eta \cdot r}$$

and

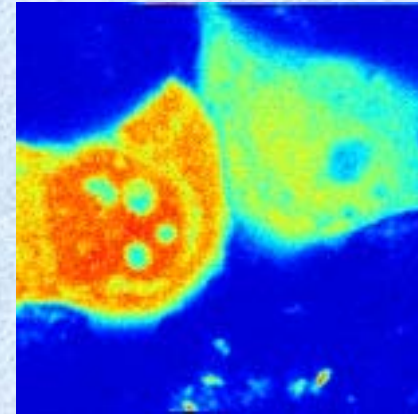
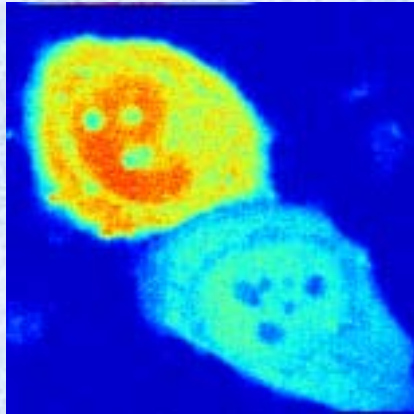
$$MW \propto \text{Volume} \propto r^3$$



Monomer  $\rightarrow$  Dimer

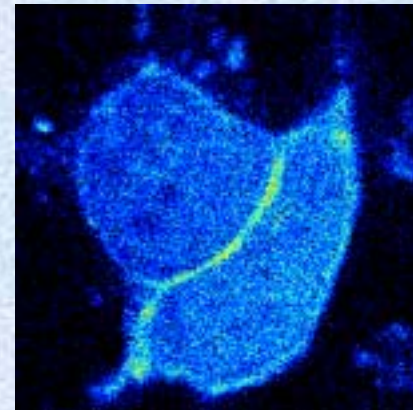
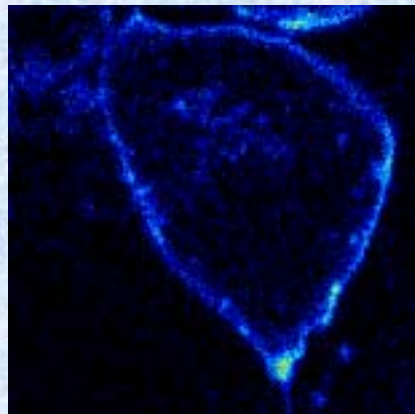
Only a change in  $D$  by a factor of  $2^{1/3}$ , or 1.26

# Autocorrelation Adenylate Kinase -EGFP Chimeric Protein in HeLa Cells



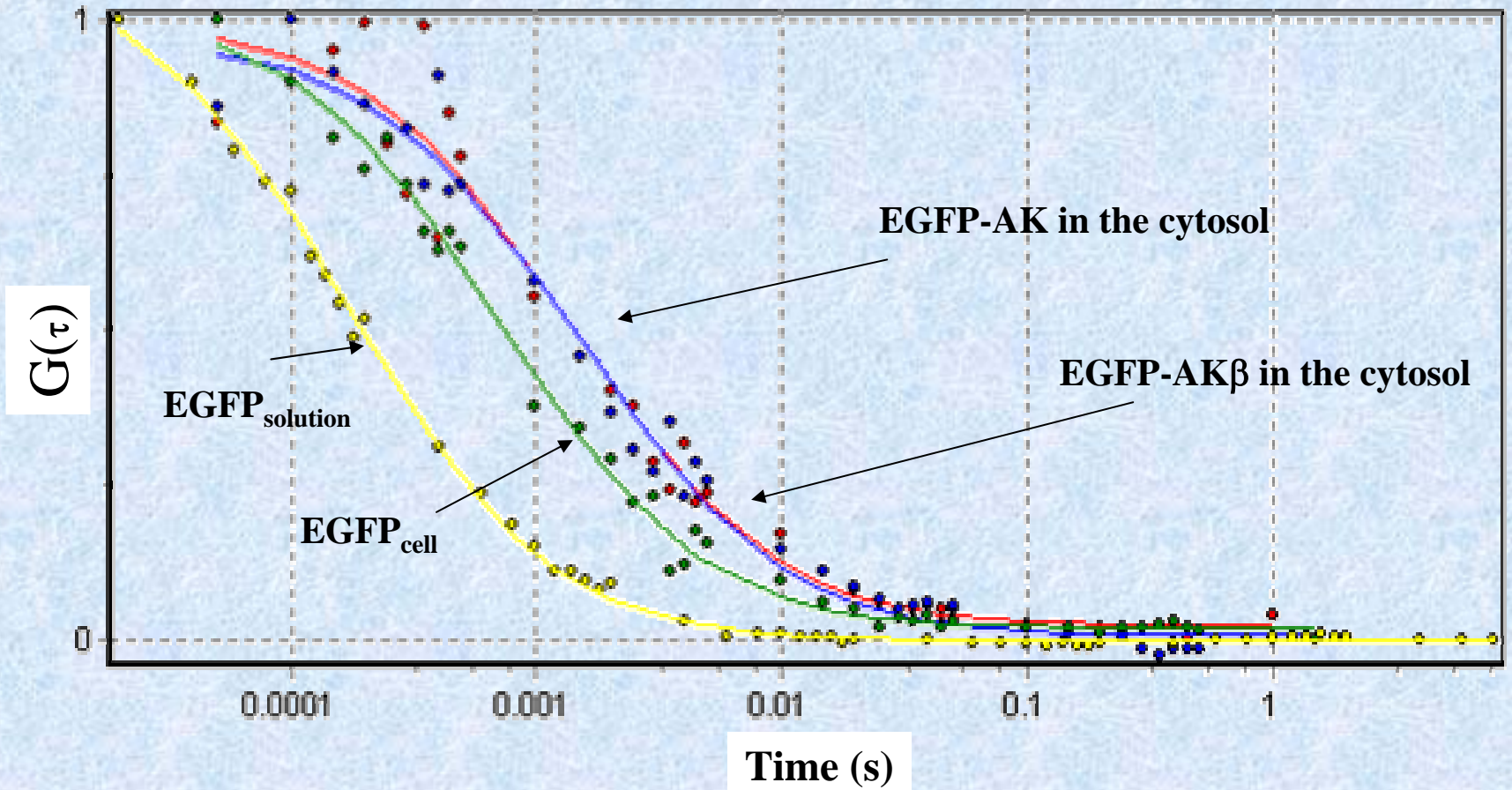
Fluorescence Intensity

Examples of different *HeLa* cells transfected with AK1-EGFP



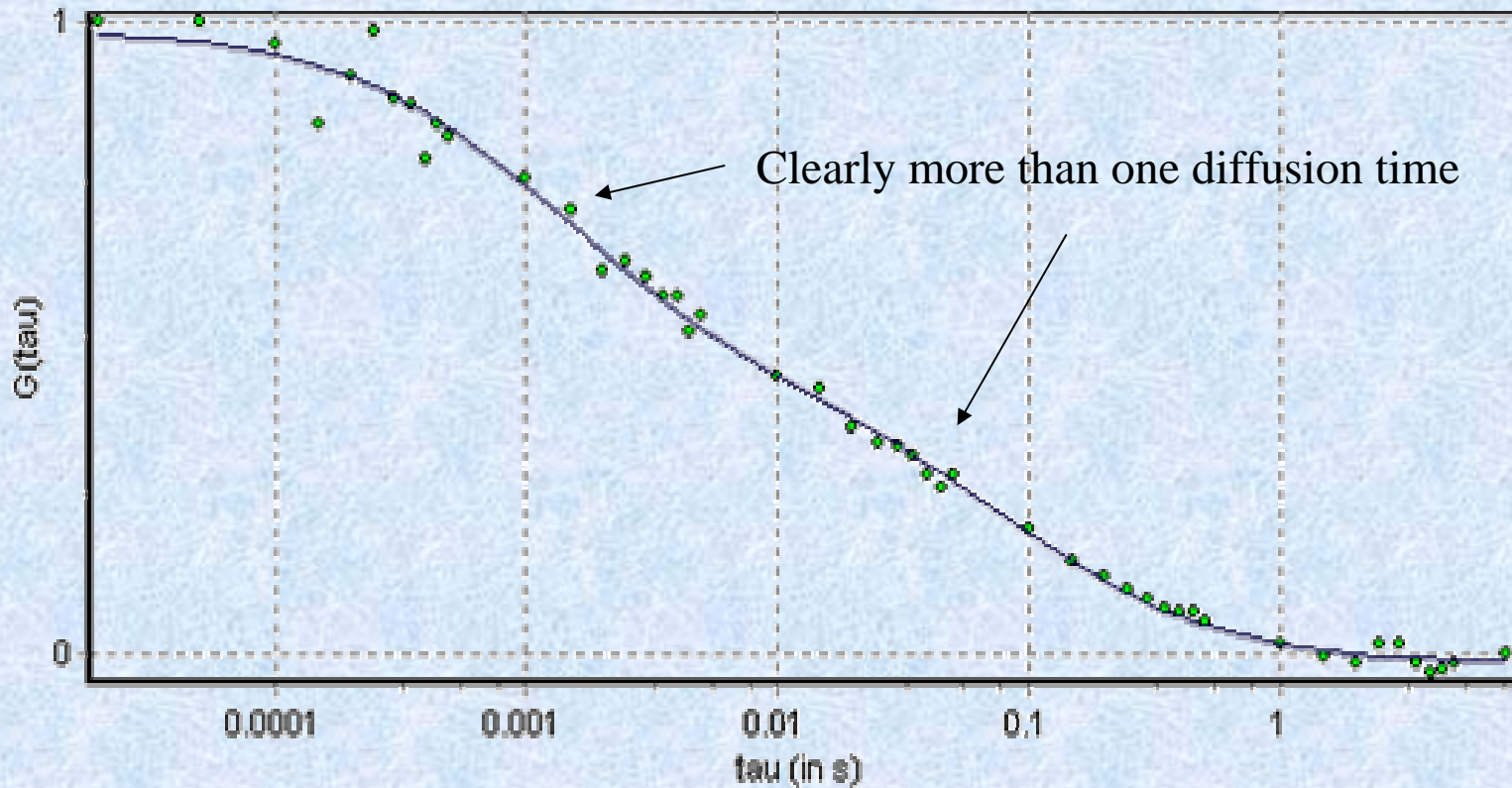
Examples of different *HeLa* cells transfected with AK1 $\beta$  -EGFP

# Autocorrelation of EGFP & Adenylate Kinase -EGFP



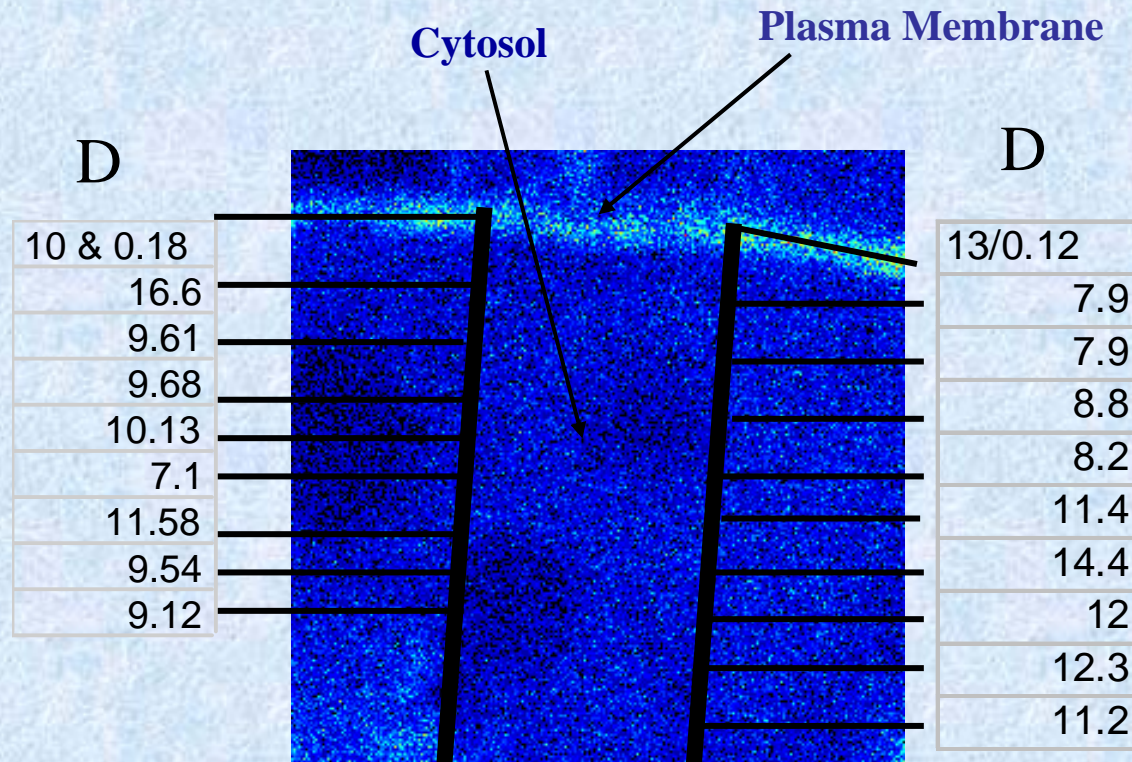
Normalized autocorrelation curve of EGFP in solution ( $\bullet$ ), EGFP in the cell ( $\bullet$ ), AK1-EGFP in the cell( $\bullet$ ), AK1 $\beta$ -EGFP in the cytoplasm of the cell( $\bullet$ ).

# Autocorrelation of Adenylate Kinase –EGFP on the Membrane



**A mixture of AK1b-EGFP in the cytoplasm and membrane of the cell.**

# Autocorrelation Adenylate Kinase $\beta$ -EGFP



Diffusion constants (um<sup>2</sup>/s) of AK EGFP-AK $\beta$  in the cytosol -EGFP in the cell (HeLa). At the membrane, a dual diffusion rate is calculated from FCS data. Away from the plasma membrane, single diffusion constants are found.

# Multiple Species

Case 1: Species vary by a difference in diffusion constant,  $D$ .

*Autocorrelation function can be used:*

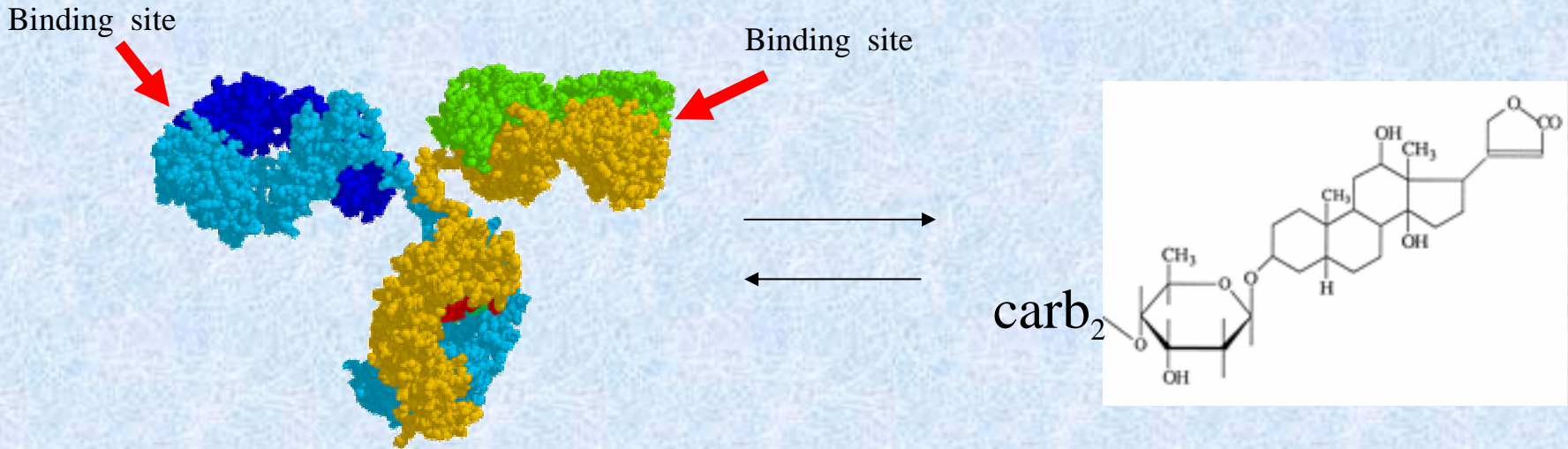
$$G(\tau)_{sample} = \sum_{i=1}^M f_i^2 \cdot G(\mathbf{0})_i \cdot \left(1 + \frac{8D\tau}{w_{2DG}^2}\right)^{-1} \quad (2D\text{-Gaussian Shape})$$

!

$$G(\mathbf{0})_{sample} = \sum f_i^2 \cdot G(\mathbf{0})_i$$

$G(0)_{sample}$  is no longer  $\gamma/N$  !

# Antibody - Hapten Interactions

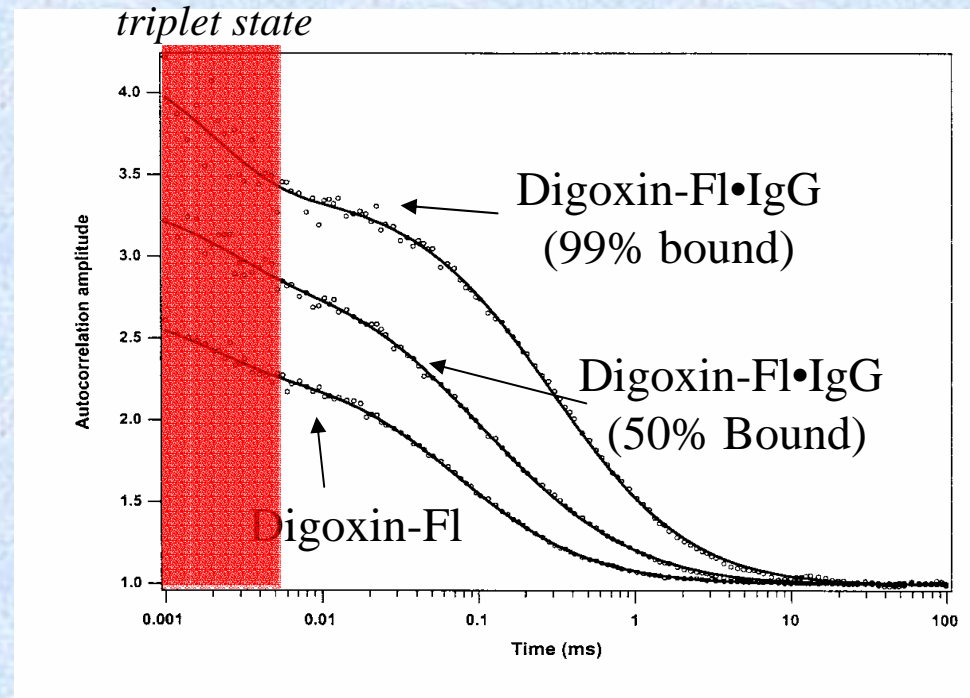


**Mouse IgG:** The two heavy chains are shown in yellow and light blue. The two light chains are shown in green and dark blue..*J.Harris, S.B.Larson, K.W.Hasel, A.McPherson, "Refined structure of an intact IgG2a monoclonal antibody", Biochemistry 36: 1581, (1997).*

**Digoxin:** a cardiac glycoside used to treat congestive heart failure. Digoxin competes with potassium for a binding site on an enzyme, referred to as potassium-ATPase. Digoxin inhibits the Na-K ATPase pump in the myocardial cell membrane.

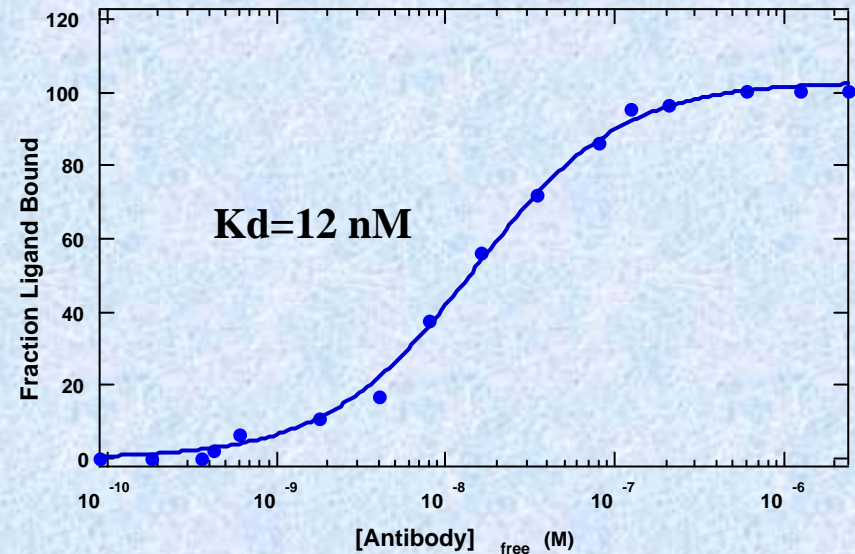
# Anti-Digoxin Antibody (IgG) Binding to Digoxin-Fluorescein

**Autocorrelation curves:**

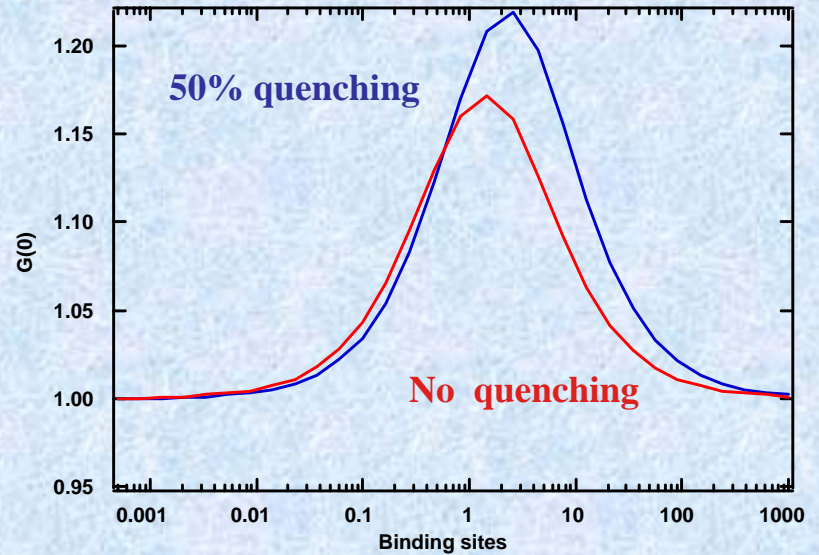
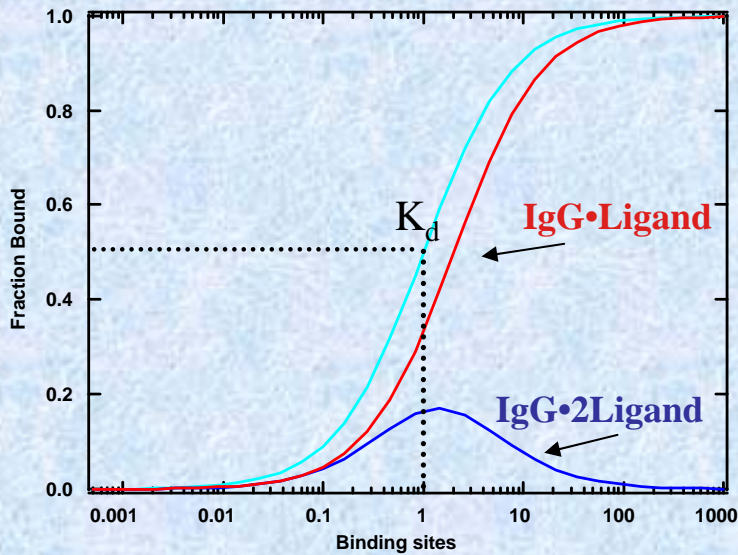


**Binding titration from the  
autocorrelation analyses:**

$$F_b = \frac{m \cdot S_{free}}{K_d + S_{free}} + c$$

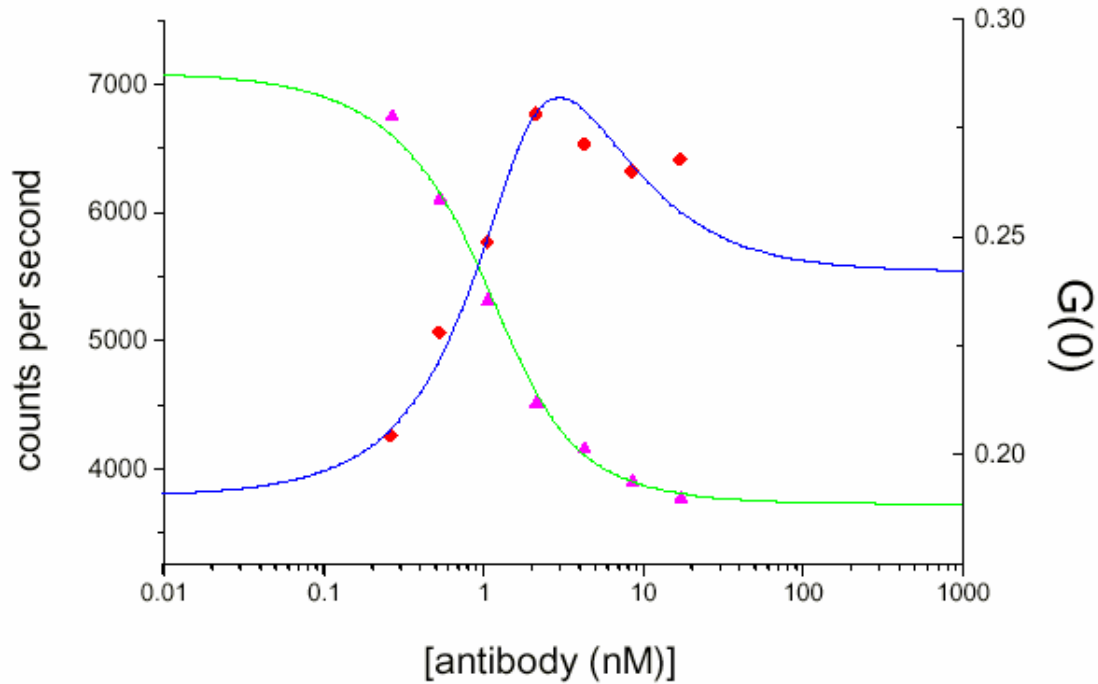


# Two Binding Site Model



$$[\text{Ligand}] = 1, G(0) = 1/N, K_d = 1.0$$

# Digoxin-FL Binding to IgG: G(0) Profile



	Lifetime (nsec)	molecular fraction (lifetime)	<i>cpsm</i>	Molecular fraction ( $G(0)$ )
Digoxin	4.01	100%	29000	100%
Ligated Digoxin( $C_1$ )	4.03	53.6%	23600	52%
Ligated Digoxin( $C_2$ )	1.25	46.4%	7100	48%

## Case 2: Species vary by a difference in brightness

assuming that  $D_1 \approx D_2$

The quantity  $G_0$  becomes the only parameter to distinguish species,  
but we know that:

$$G(\mathbf{0})_{sample} = \sum f_i^2 \cdot G(\mathbf{0})_i$$

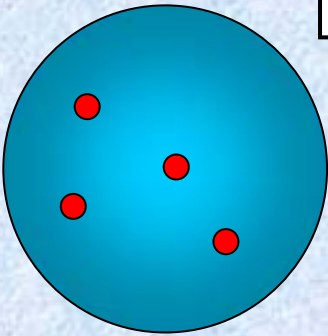
**The autocorrelation function is not suitable  
for analysis of this kind of data without additional information.**

We need a different type of analysis



# Photon Counting Histogram (PCH)

**Aim:** To resolve species from differences in their molecular brightnesses



**Poisson Distribution:** 
$$p(N) = \frac{\langle N \rangle^N \cdot e^{-\langle N \rangle}}{N!}$$

## Sources of Non-Poissonian Noise

Detector Noise

Diffusing Particles in an Inhomogeneous  
Excitation Beam\*

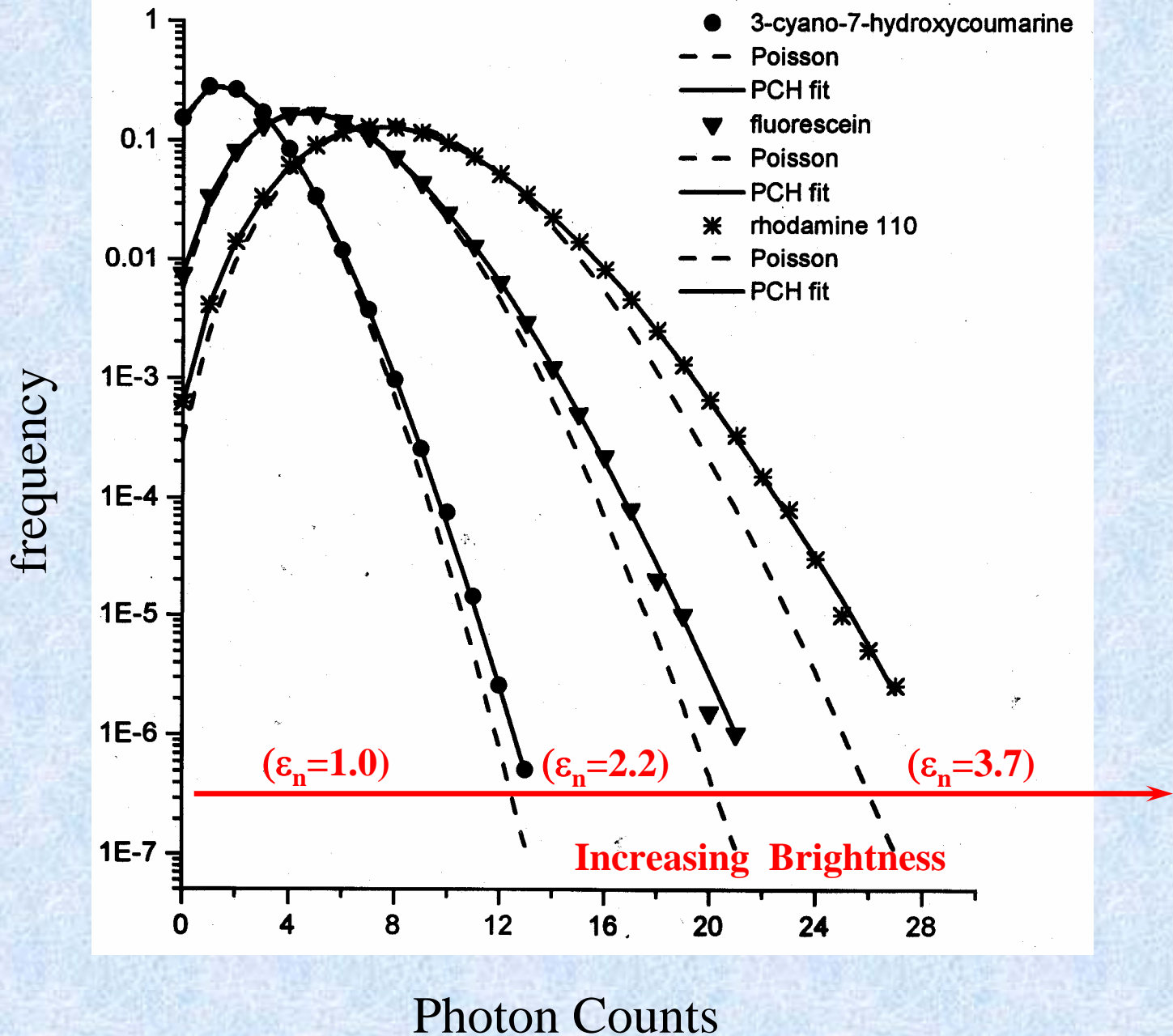
Particle Number Fluctuations\*

Multiple Species\*

**Single Species:** 
$$p(k) = PCH(\varepsilon, \langle N \rangle)$$

Where  $p(k)$  is the probability of observing  $k$  photon counts

# PCH Example: Differences in Brightness



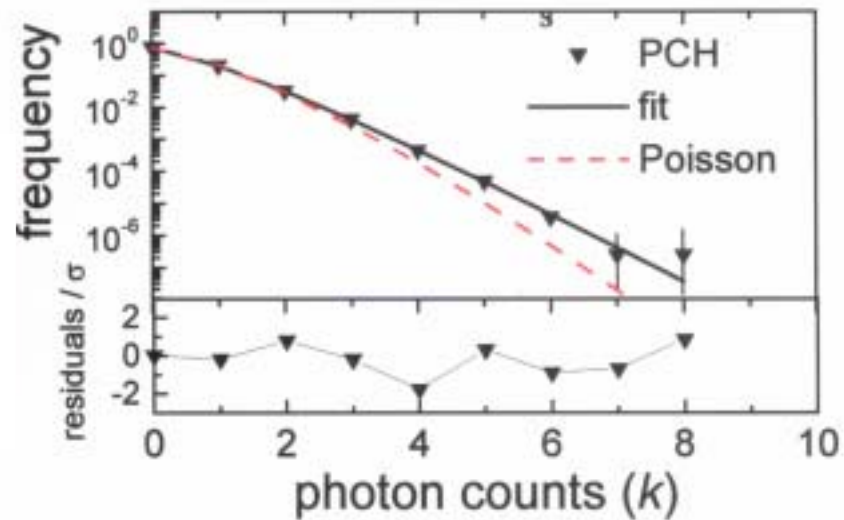
# Single Species PCH: Concentration

5.5 nM Fluorescein

Fit:

$\varepsilon = 16,000$  cpsm

$N = 0.3$

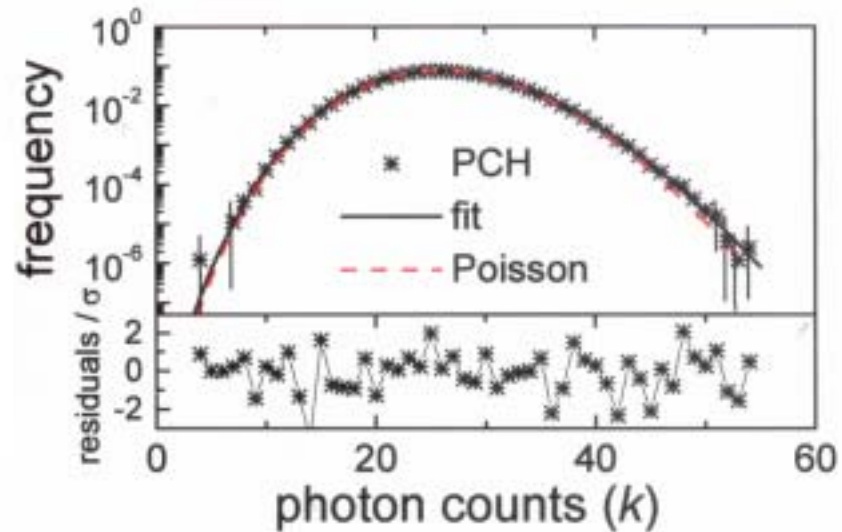


550 nM Fluorescein

Fit:

$\varepsilon = 16,000$  cpsm

$N = 33$

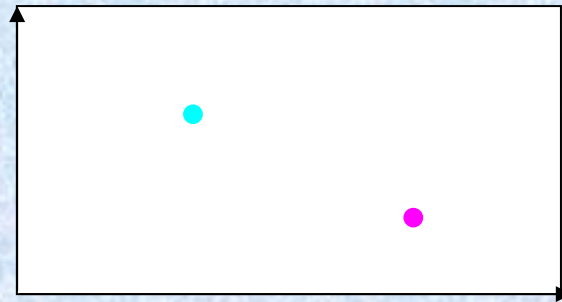


**As particle concentration increases the PCH approaches a Poisson distribution**

# Photon Counting Histogram: Multispecies

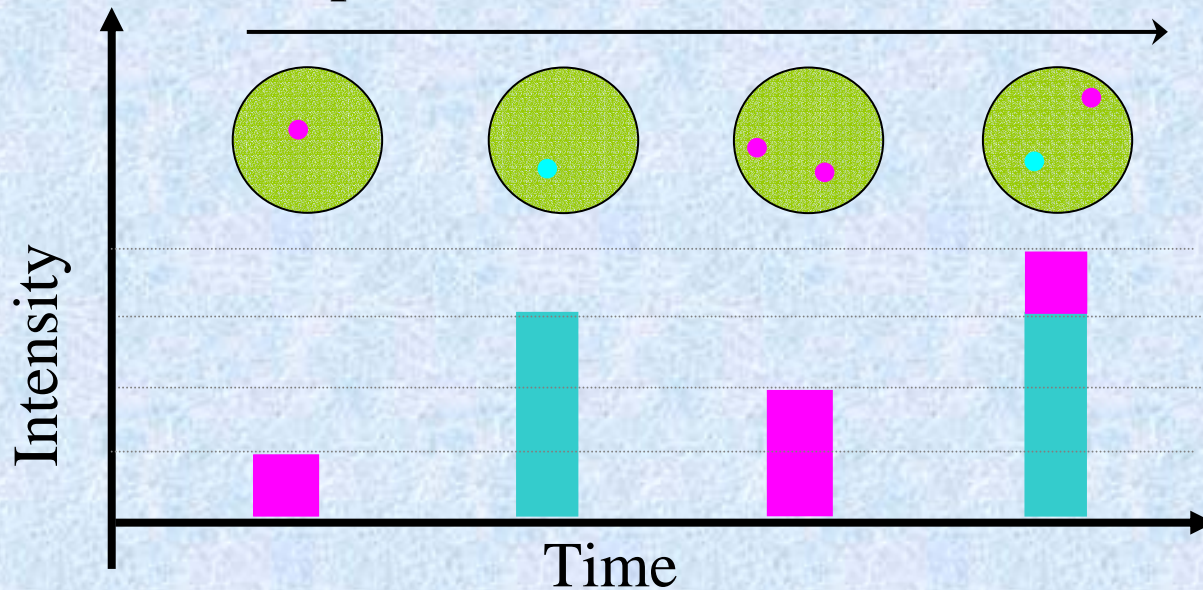
Binary Mixture:  $p(k) = PCH(\varepsilon_1, \langle N_1 \rangle) \otimes PCH(\varepsilon_2, \langle N_2 \rangle)$

Molecular  
Brightness



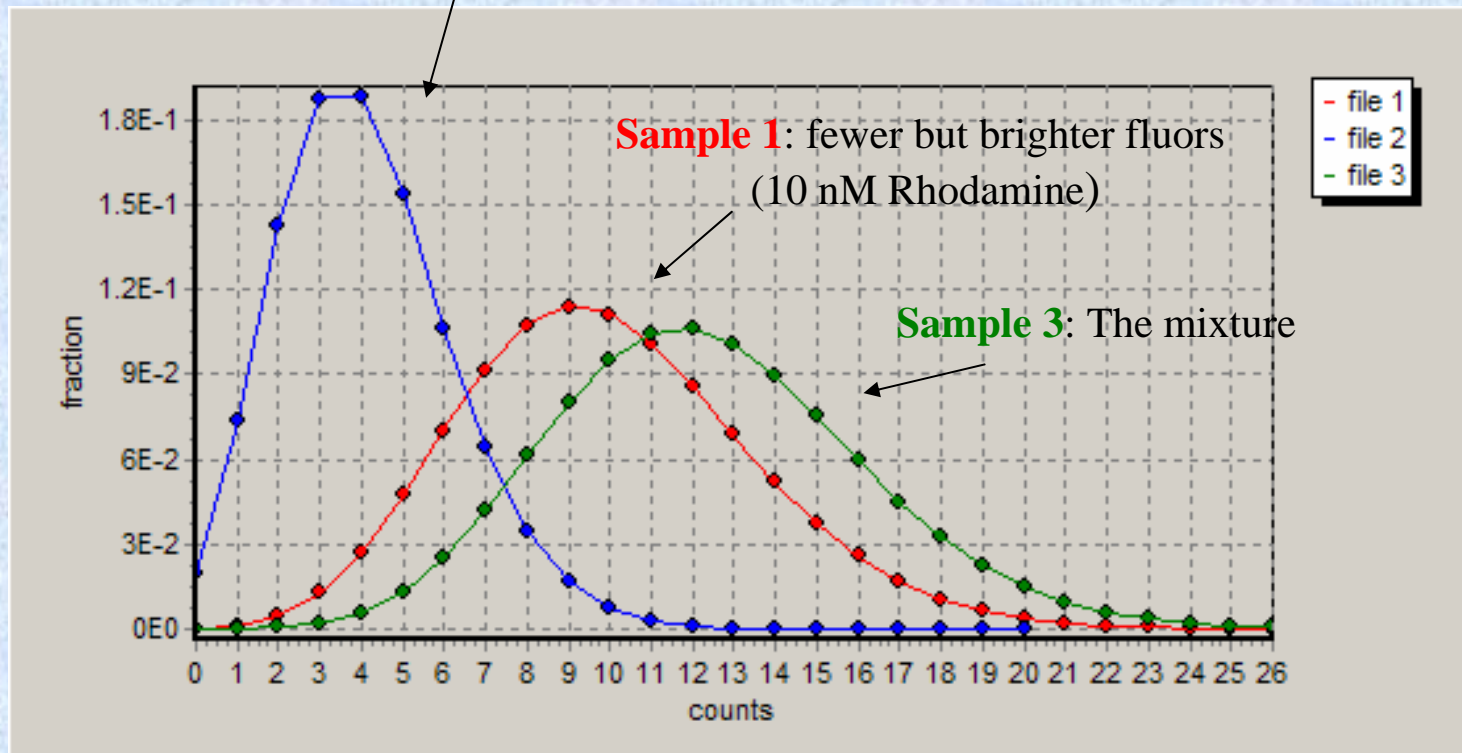
Concentration

Snapshots of the excitation volume



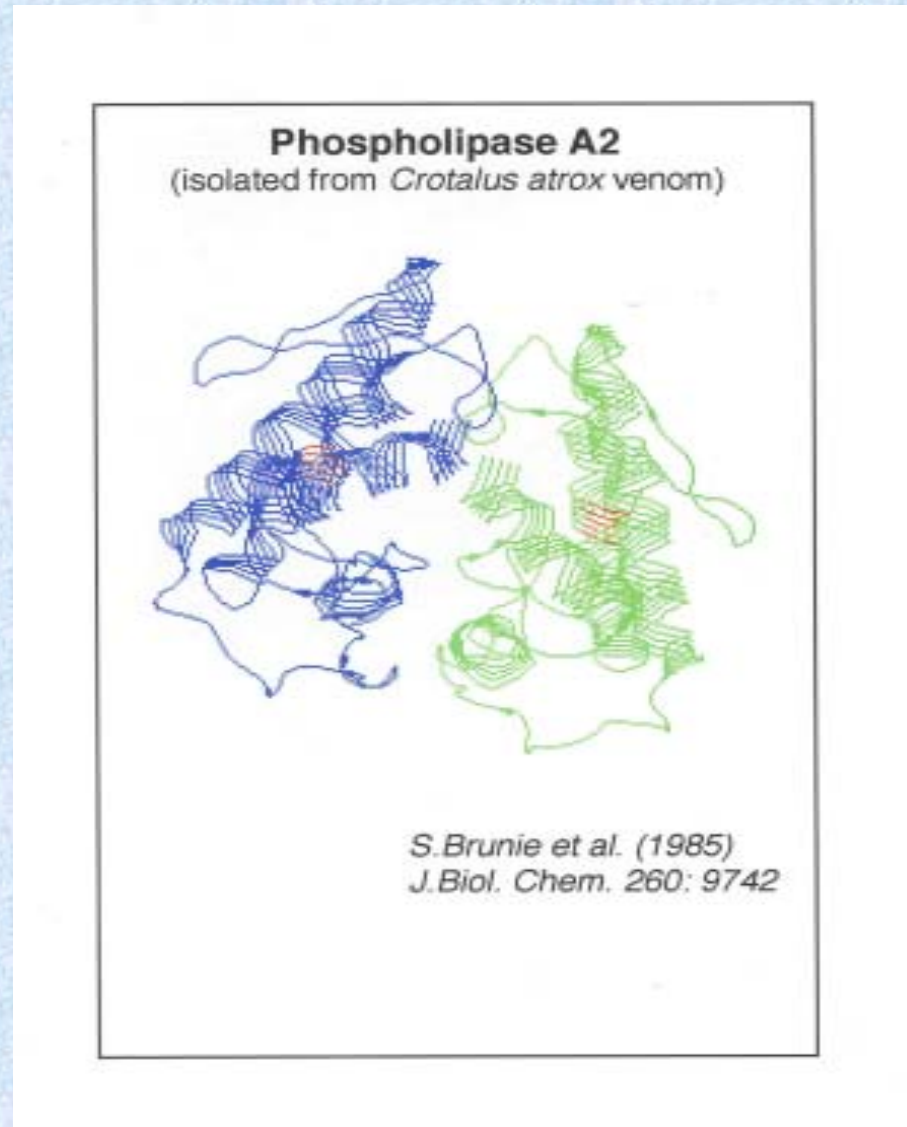
# Photon Counting Histogram: Multispecies

**Sample 2:** many but dim (23 nM fluorescein at pH 6.3)



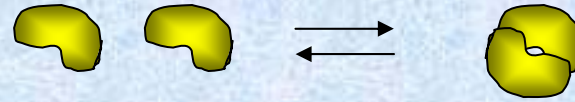
The occupancy fluctuations for each specie in the mixture becomes a convolution of the individual specie histograms. The resulting histogram is then broader than expected for a single species.

# Examination of a Protein Dimer with FCS: Secreted Phospholipase A<sub>2</sub>



# sPLA<sub>2</sub> Interfacial Binding

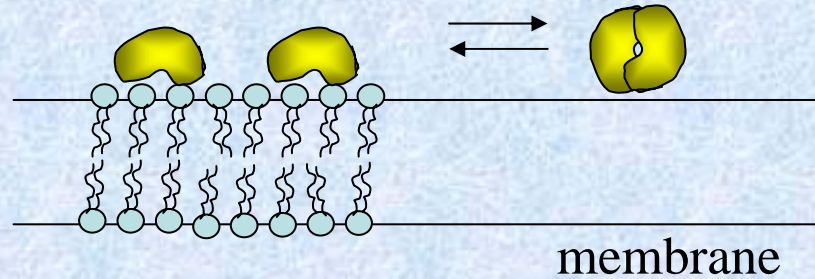
sPLA<sub>2</sub> Self-Association



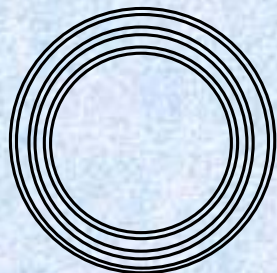
sPLA<sub>2</sub> Membrane Binding



Interfacial sPLA<sub>2</sub> Self-Association

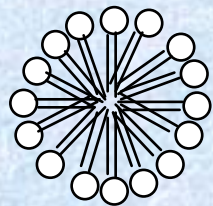
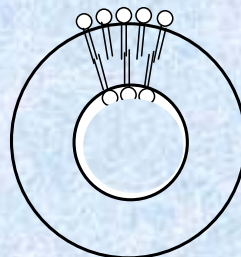


# Lipid Interfaces

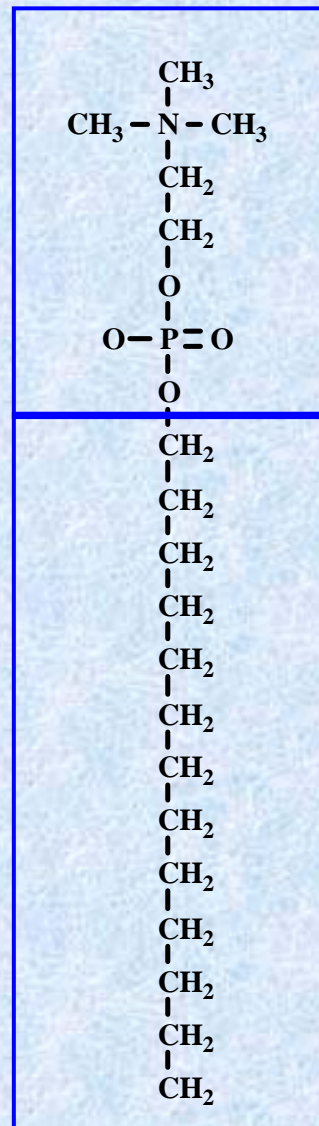


Multibilayers  
(MLVs)

Vesicles  
(SUVs, LUVs  
& GUVs)



Micelles



Choline Group

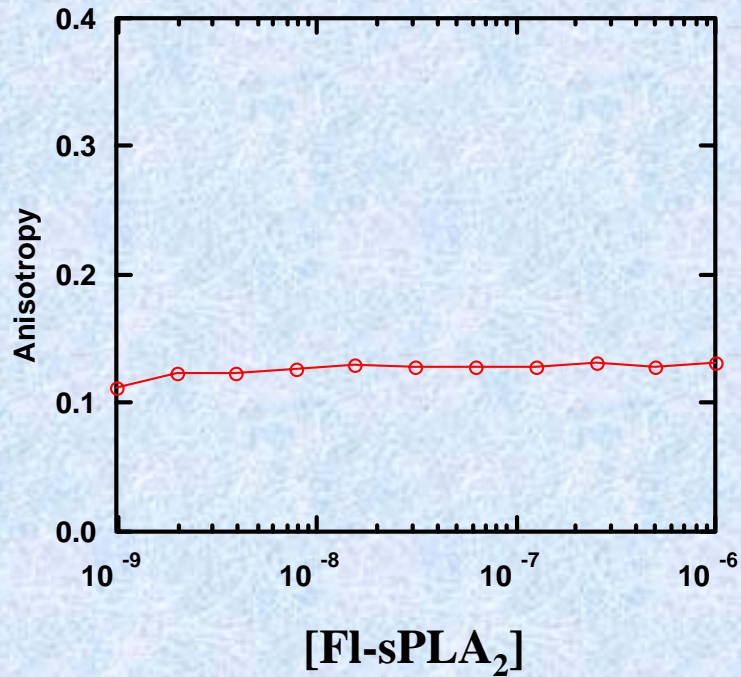
12 Carbon Tail

**Dodecylphosphocholine (DPC)**  
**Micellar Lipid Analog (CMC = 1.1 mM)**

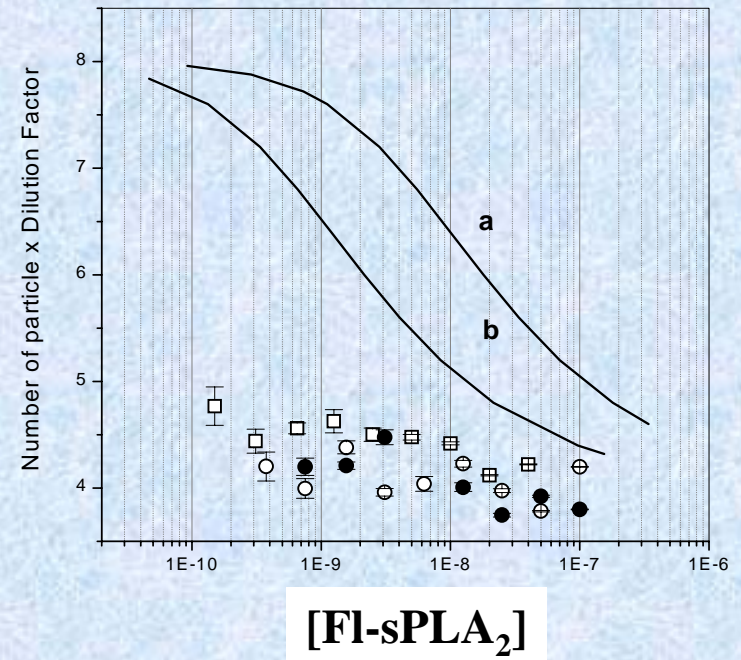
# In Solution: *a Tight Dimer*

## Fluorescein-sPLA<sub>2</sub>

Steady-State Anisotropy



Fluorescence Correlation Spectroscopy



*Time-Resolved Anisotropy:*

**Phi1 = 12.8 ns (0.43)**

**Phi2 = 0.50 ns (0.57)**

# In Solution: Fluorescein-sPLA<sub>2</sub> +/- Urea

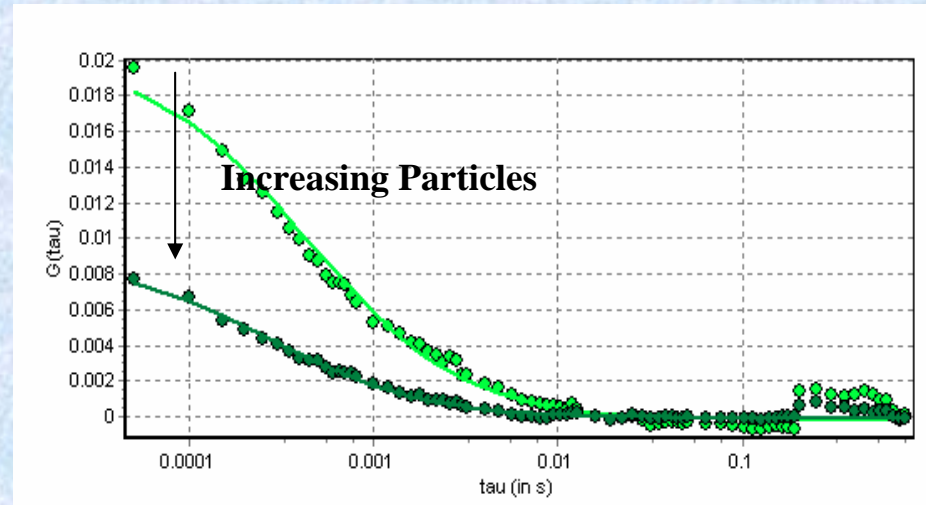
## 1. Autocorrelation

sPLA<sub>2</sub>

$$G(0) = 0.021$$
$$D = 72 \text{ } \mu\text{m}^2/\text{s}$$

sPLA<sub>2</sub> + 3M Urea

$$G(0) = 0.009$$
$$D = 95 \text{ } \mu\text{m}^2/\text{s}$$



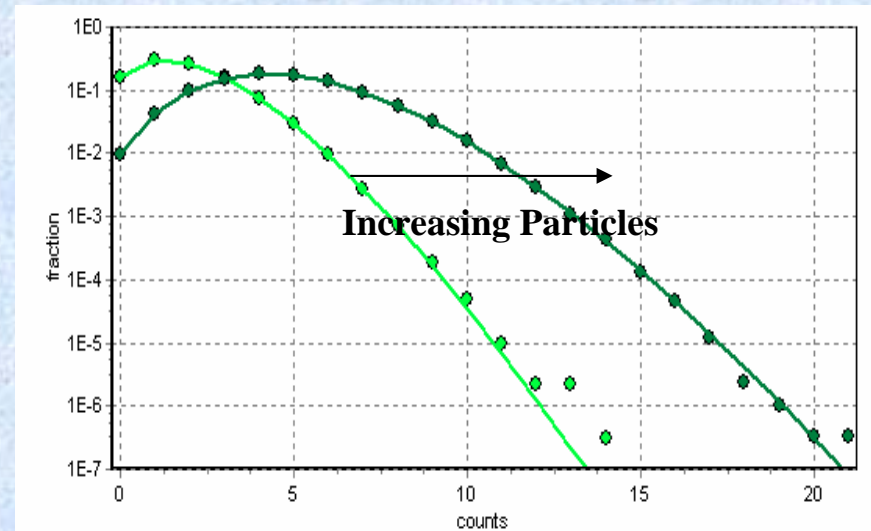
## 2. PCH analysis

sPLA<sub>2</sub>

$$\varepsilon = 0.6$$
$$N = 3.29$$

sPLA<sub>2</sub> + 3M Urea

$$\varepsilon = 0.6$$
$$N = 8.48$$

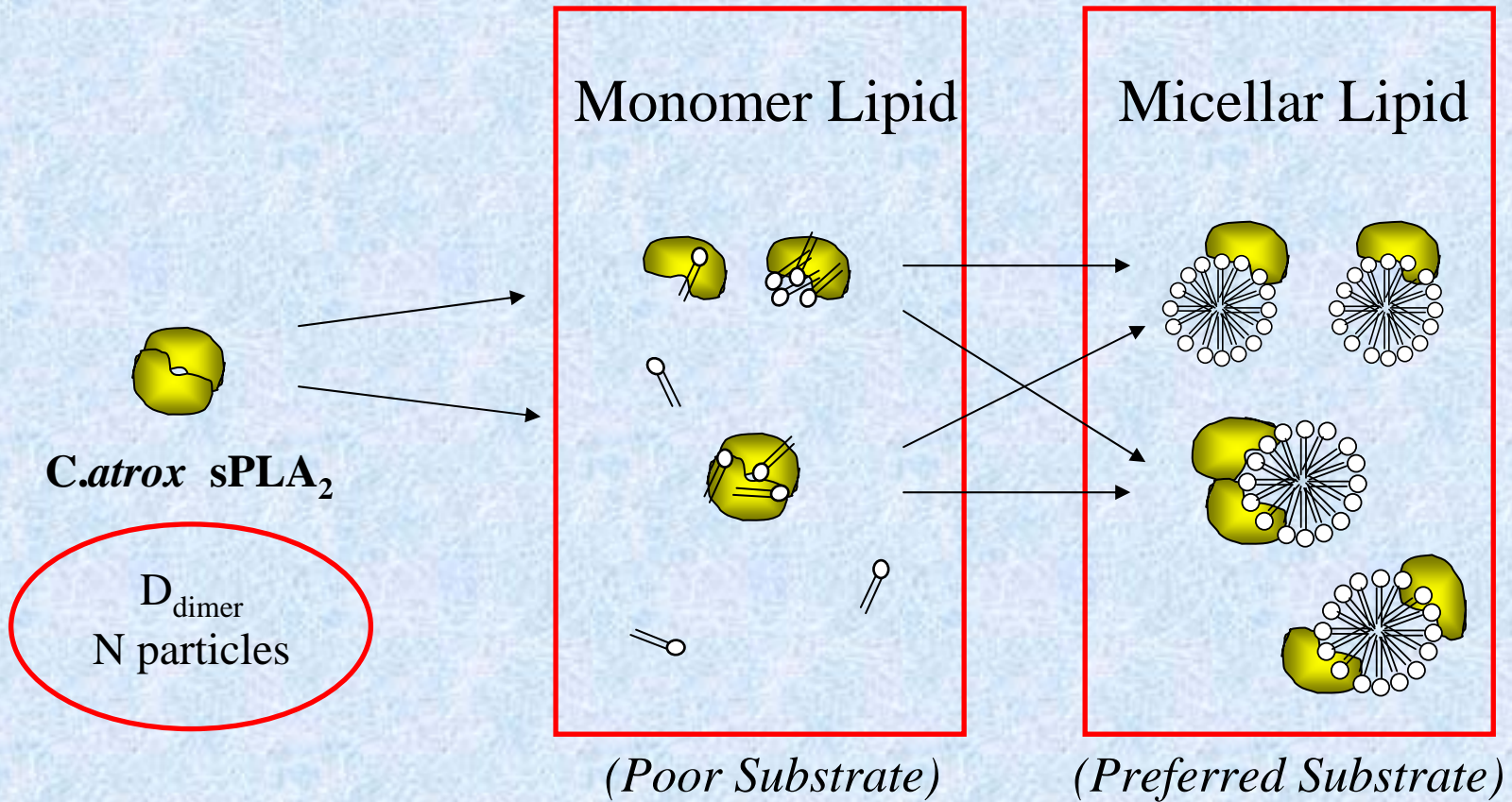


*Adjusted for viscosity differences*

Change in number of particles, little change in brightness

# The Critical Question: Is sPLA<sub>2</sub> a Dimer in the Presence of Interfacial Lipid?

*What Could We Expect to Find in the FCS Data?*



Observing Fluorescein-labeled sPLA<sub>2</sub>

# FCS on Fluorescein - sPLA<sub>2</sub> in Buffer (RED) and with DPC Micelles (BLUE)

## 1. Autocorrelation Analysis

sPLA<sub>2</sub>

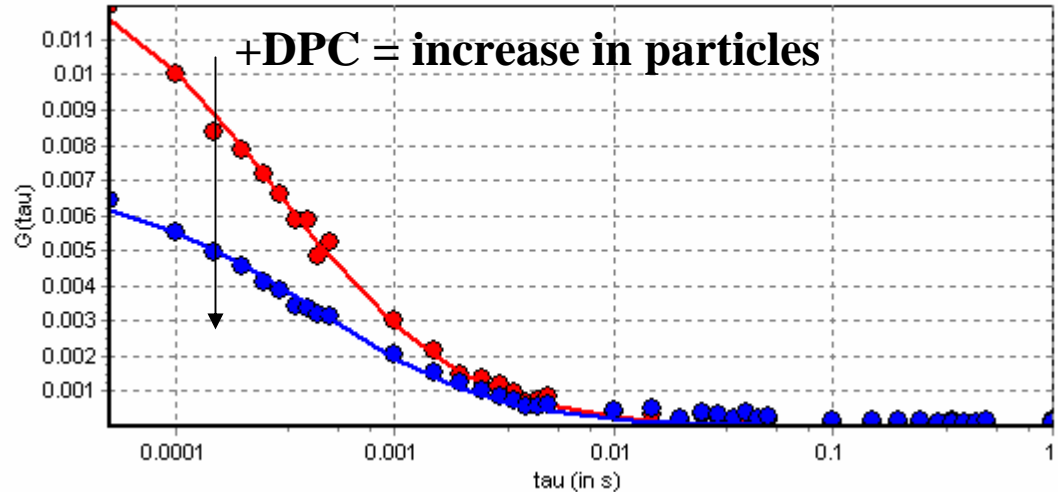
$$G_0 = 0.0137$$

$$D = 75 \text{ } \mu\text{m}^2/\text{s}$$

sPLA<sub>2</sub> + 20 mM DPC

$$G_0 = 0.0069$$

$$D = 55 \text{ } \mu\text{m}^2/\text{s}$$



## 2. PCH Analysis

sPLA<sub>2</sub>

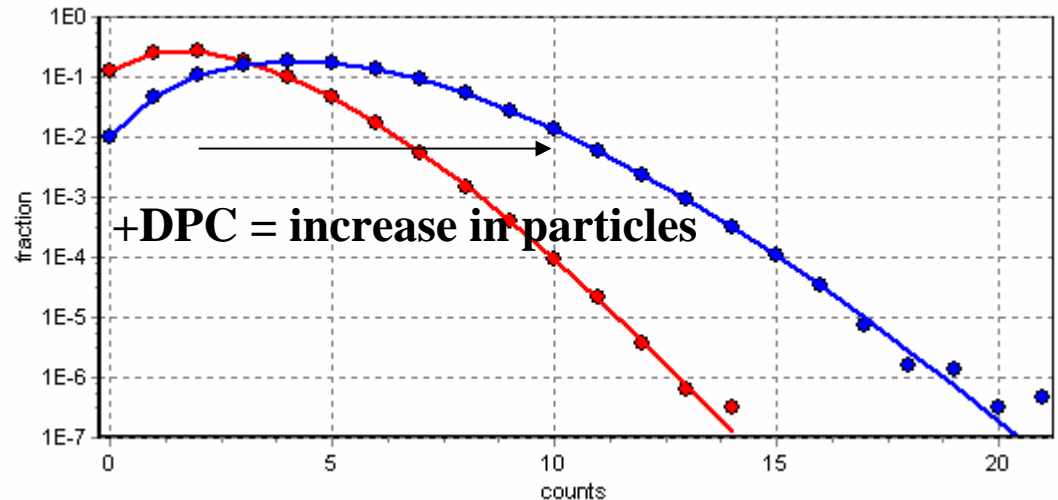
$$\varepsilon = 0.41$$

$$N = 6.5$$

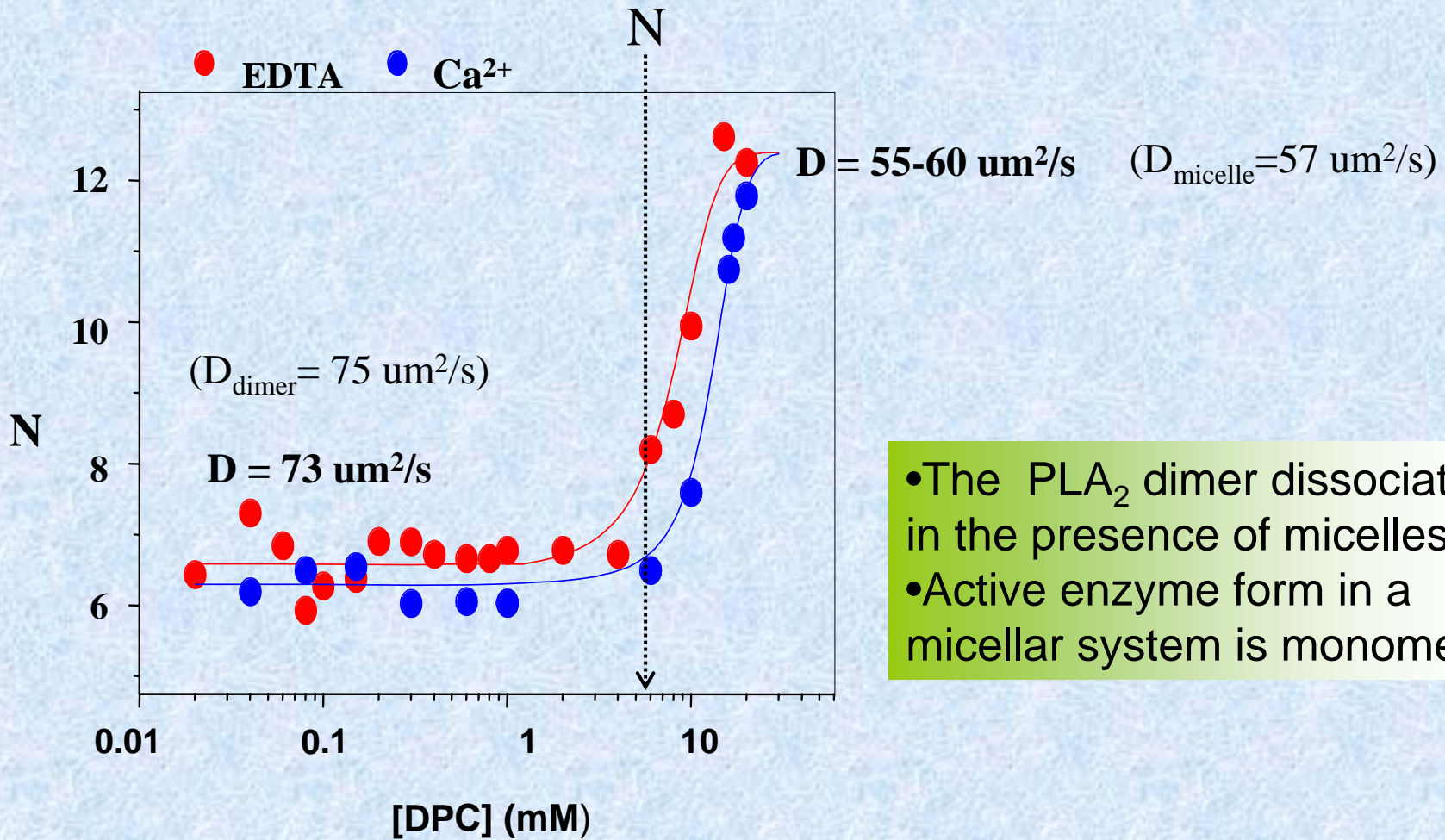
sPLA<sub>2</sub> + 20 mM DPC

$$\varepsilon = 0.45$$

$$N = 12.2$$

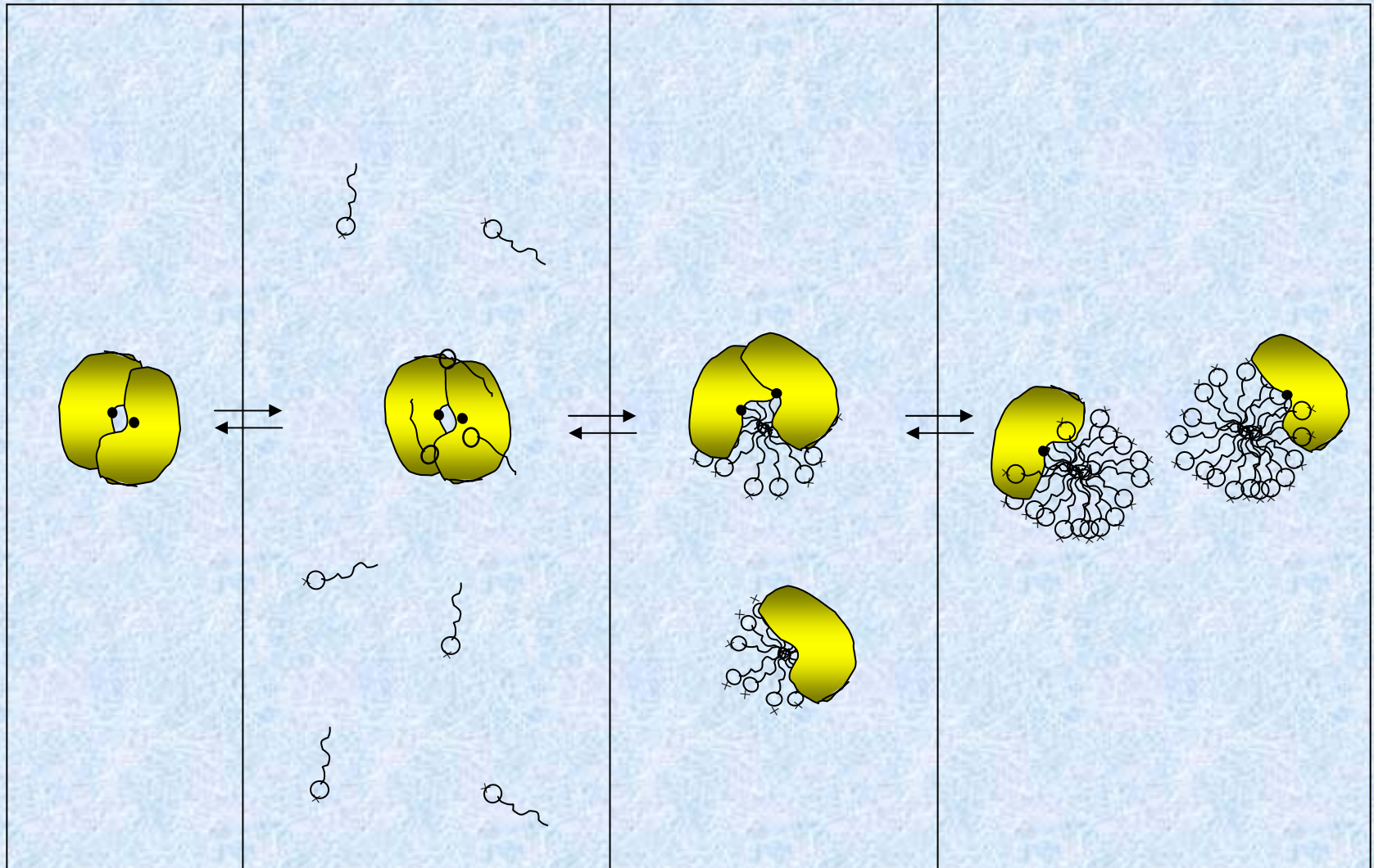


# Fluorescein-sPLA<sub>2</sub> Interaction with DPC



- The PLA<sub>2</sub> dimer dissociates in the presence of micelles.
- Active enzyme form in a micellar system is monomeric.

# Schematic of sPLA<sub>2</sub> - Dodecylphosphocholine Interactions



sPLA<sub>2</sub>

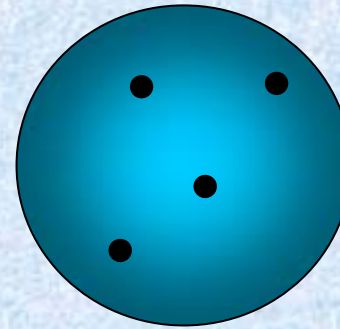
Monomer-Lipid  
Association

Co-Micelle

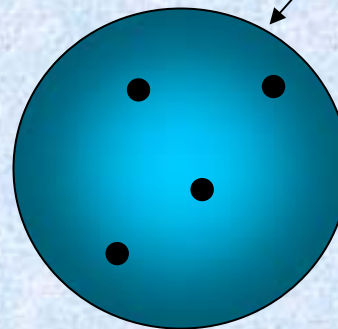
sPLA<sub>2</sub>-Micelle

# Two Channel Detection: Cross-correlation

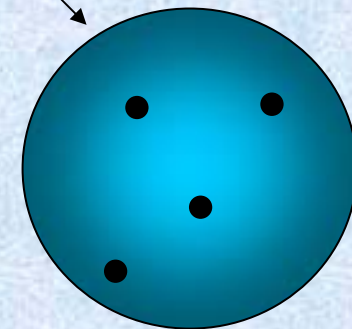
Sample Excitation  
Volume



*Beam Splitter*



**Detector 1**



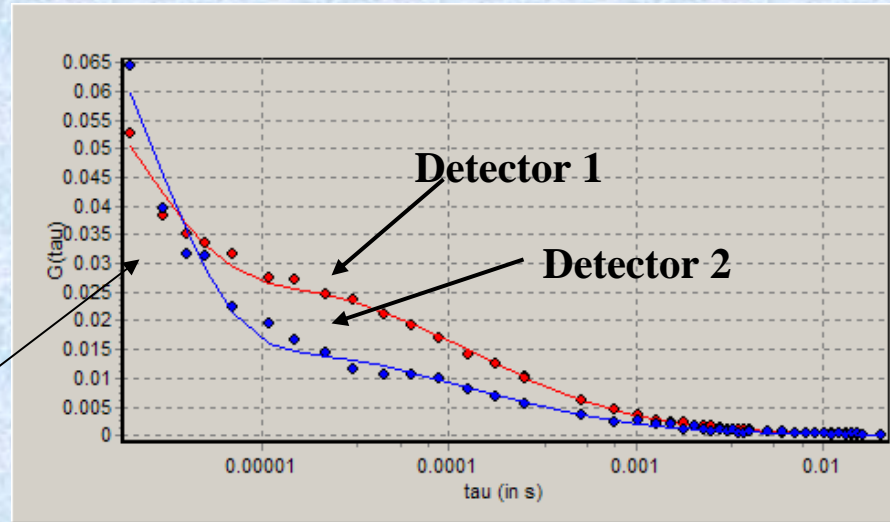
**Detector 2**

1. Increases signal to noise by isolating correlated signals.
2. Corrects for PMT noise

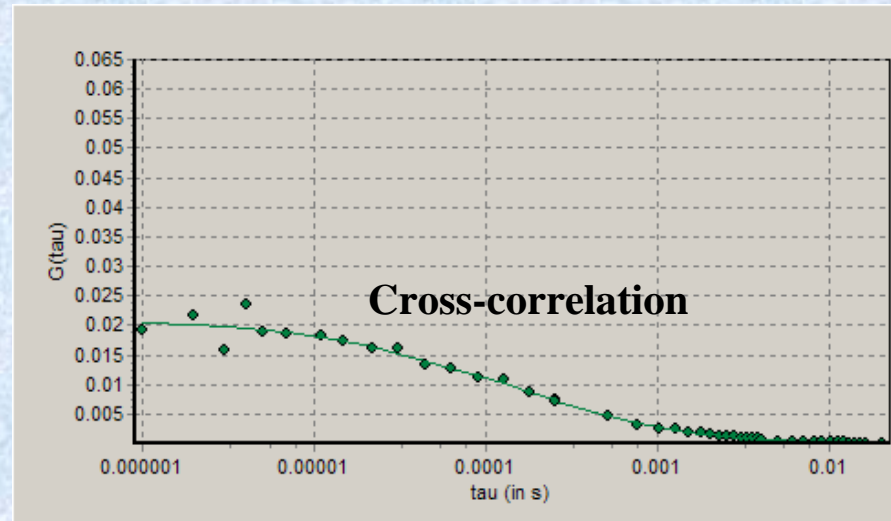
Each detector observes  
the same particles

# Removal of Detector Noise by Cross-correlation

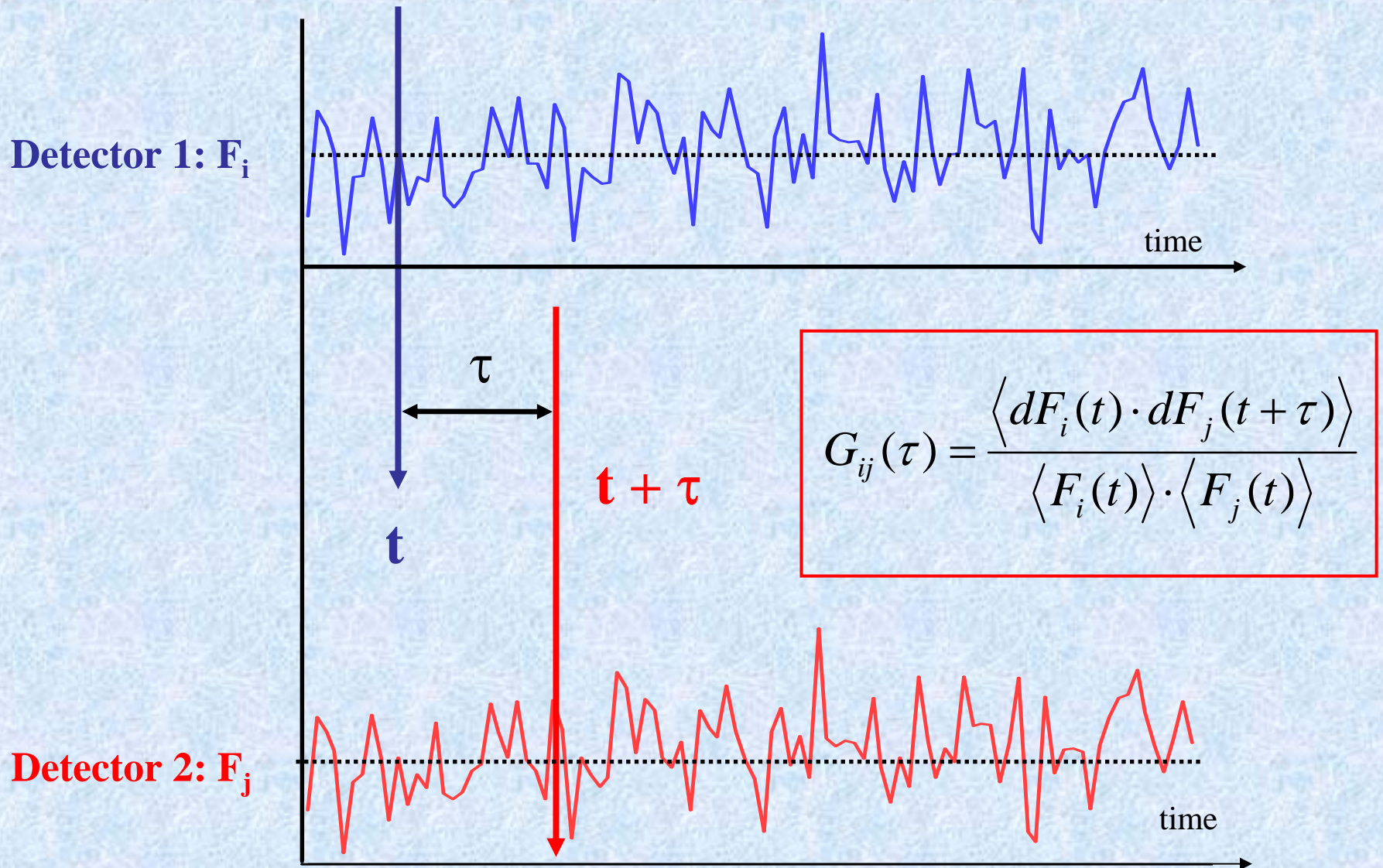
11.5 nM Fluorescein



Detector after-pulsing



# Calculating the Cross-correlation Function



# Cross-Correlation Calculations

One uses the same fitting functions you would use for the standard autocorrelation curves.

**Thus, for a 3-dimensional Gaussian excitation volume one uses:**

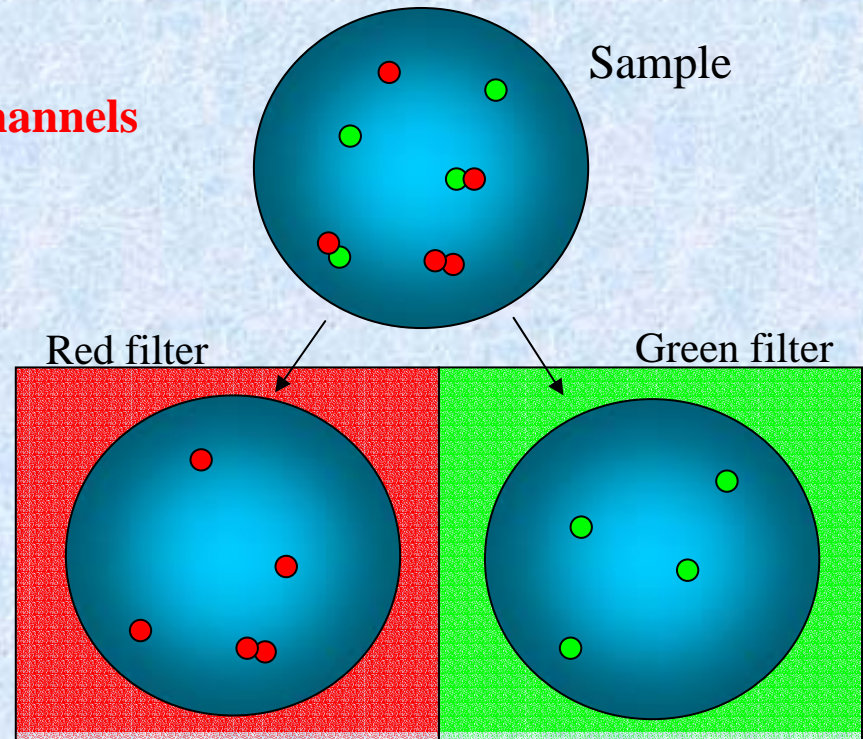
$$G_{12}(\tau) = \frac{\gamma}{N_{12}} \left( 1 + \frac{8D_{12}\tau}{w^2} \right)^{-1} \left( 1 + \frac{8D_{12}\tau}{z^2} \right)^{-1/2}$$

**$G_{12}$  is commonly used to denote the cross-correlation and  $G_1$  and  $G_2$  for the autocorrelation of the individual detectors.** Sometimes you will see  $G_x(0)$  or  $C(0)$  used for the cross-correlation.

# Two-Color Cross-correlation

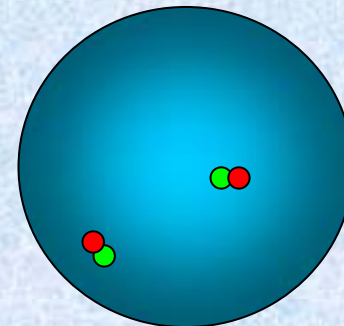
**The cross-correlation  
ONLY if particles are observed in both channels**

Each detector observes  
particles with a particular color



The cross-correlation signal:

**Only the green-red molecules are observed!!**



# Two-color Cross-correlation

Equations are similar to those for the cross correlation using a simple beam splitter:

$$G_{ij}(\tau) = \frac{\langle dF_i(t) \cdot dF_j(t + \tau) \rangle}{\langle F_i(t) \rangle \cdot \langle F_j(t) \rangle}$$

## Information Content

## Signal

Correlated signal from particles having **both colors**.

$$G_{12}(\tau)$$

Autocorrelation from channel 1 on the **green particles**.

$$G_1(\tau)$$

Autocorrelation from channel 2 on the **red particles**.

$$G_2(\tau)$$

# Experimental Concerns: Excitation Focusing & Emission Collection

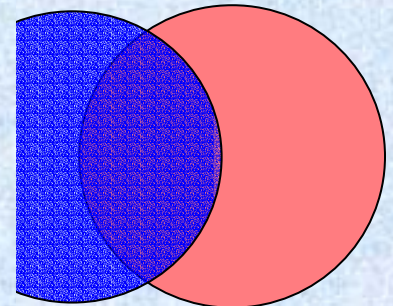
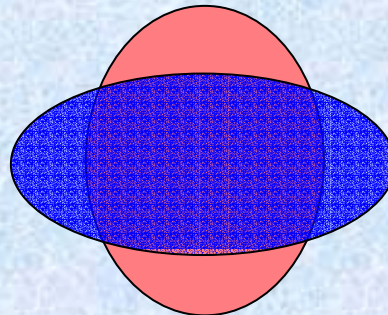
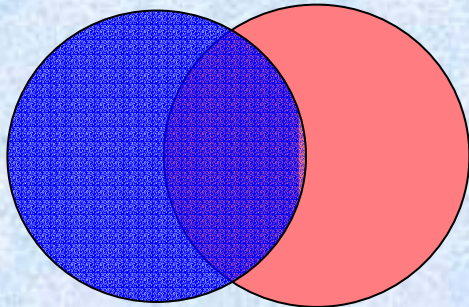
*We assume exact match of the observation volumes in our calculations which is difficult to obtain experimentally.*

## **Excitation side:**

- (1) Laser alignment
- (2) Chromatic aberration
- (3) Spherical aberration

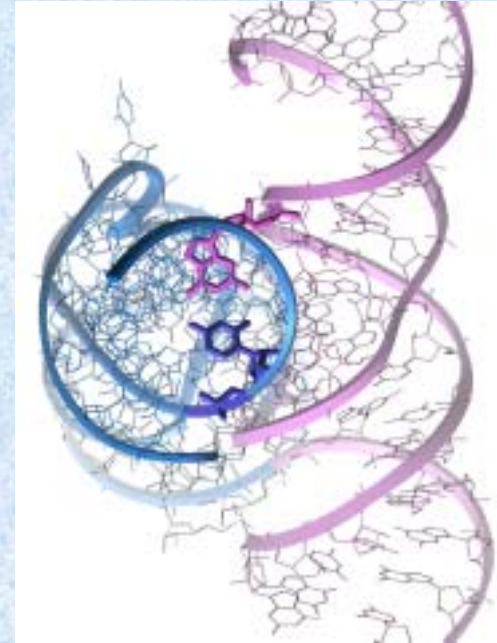
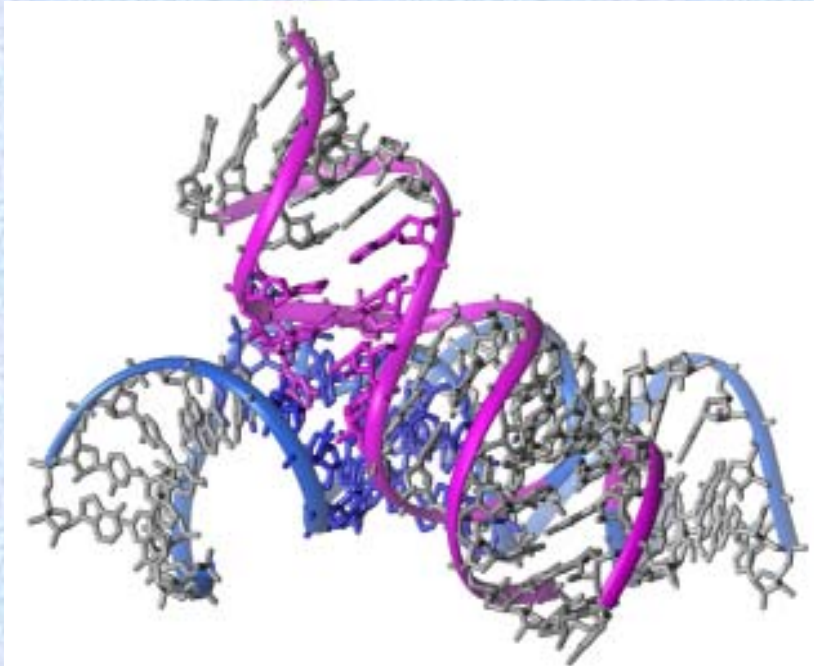
## **Emission side:**

- (1) Chromatic aberrations
- (2) Spherical aberrations
- (3) Improper alignment of detectors or pinhole  
(cropping of the beam and focal point position)



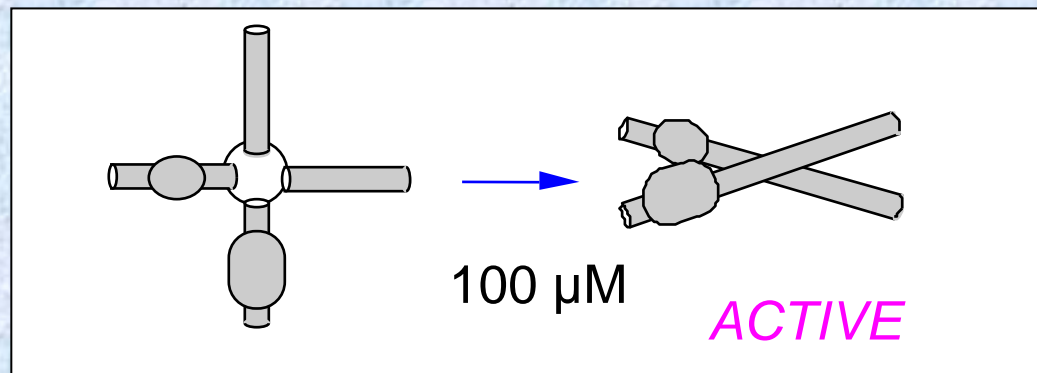
# Single Molecule Application of FCS

*E. Tan, T. Ha & R. Clegg (UIUC, LFD)*

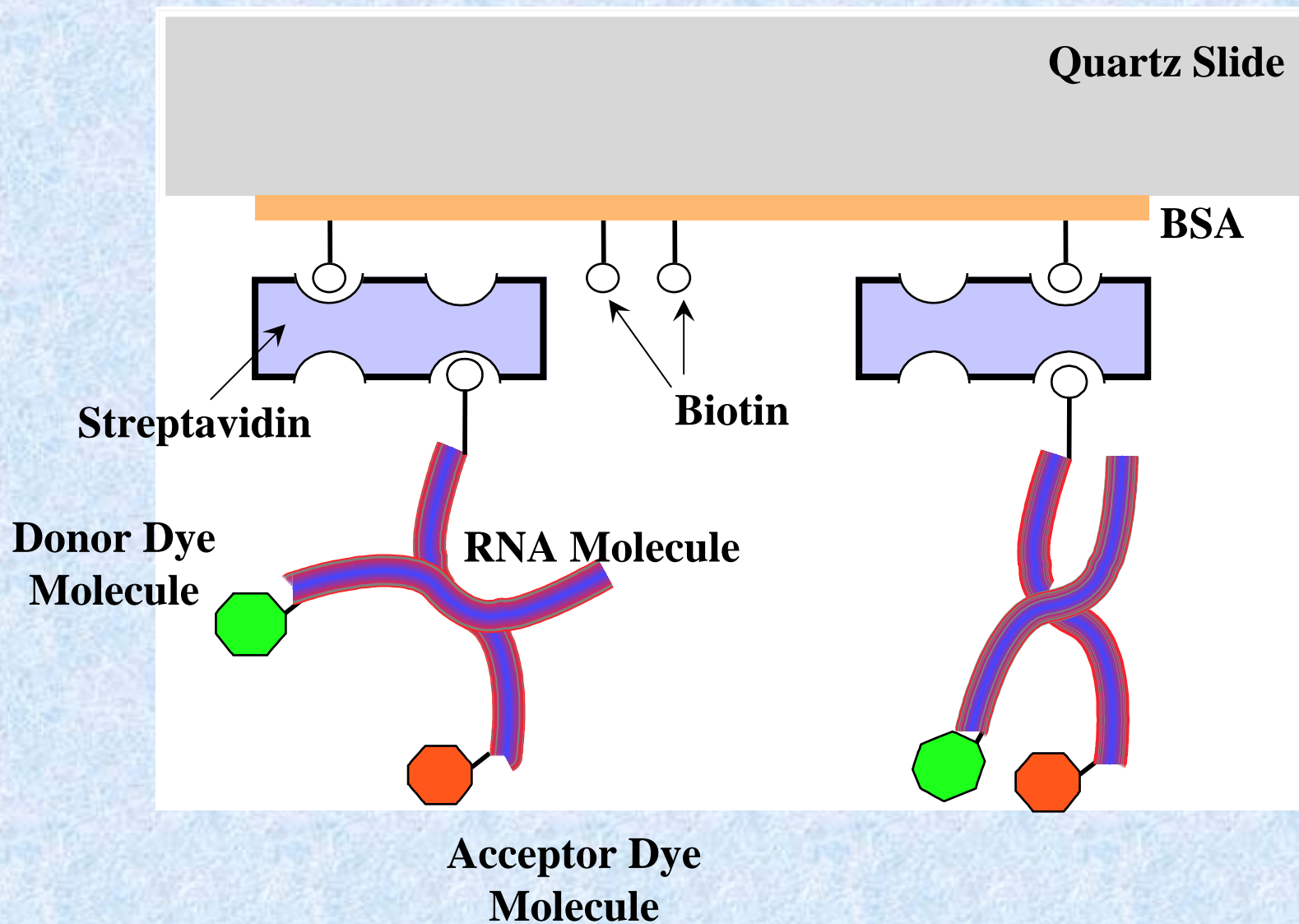


Crystal Structure of Hairpin Ribozyme

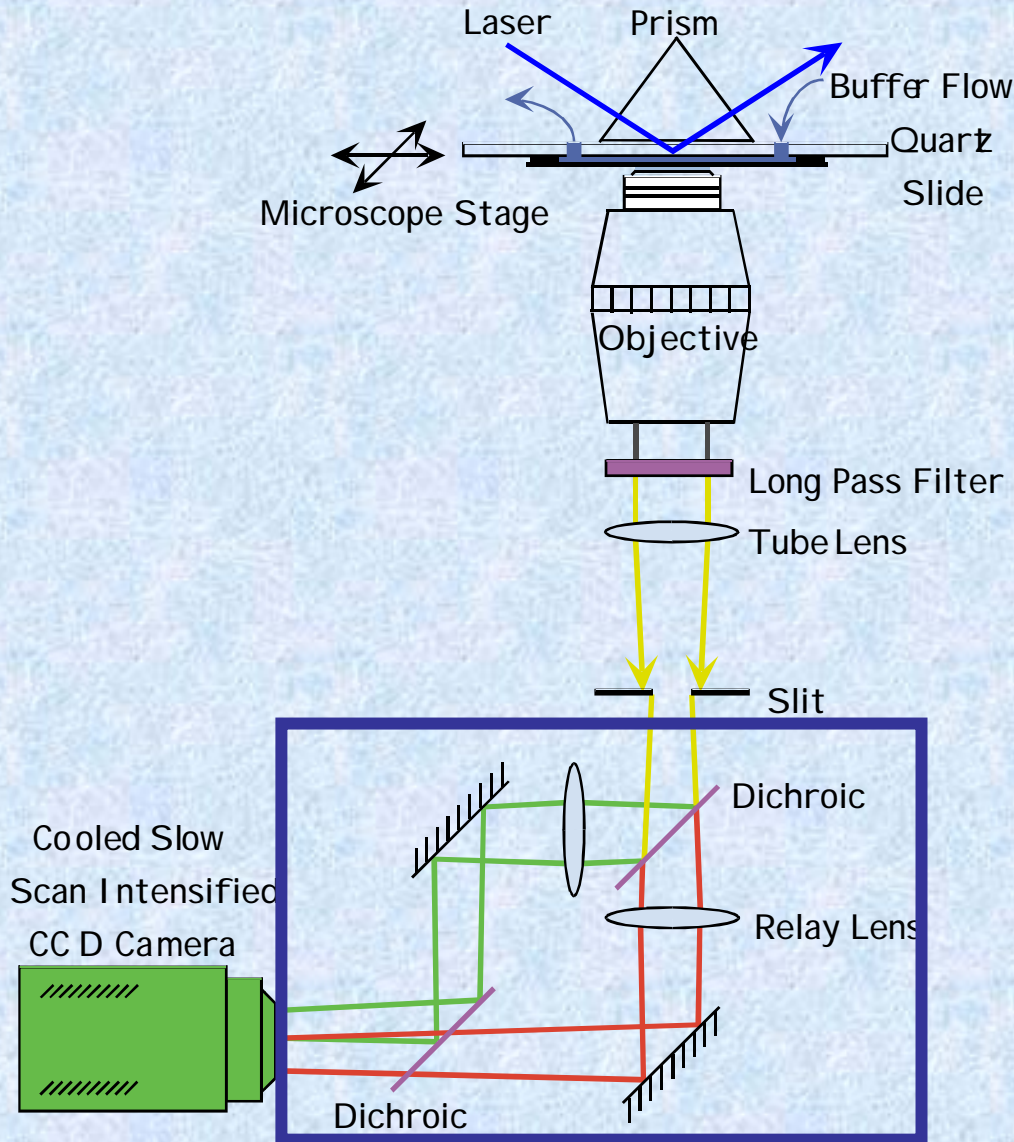
# HAIRPIN RIBOZYME : FOLDING OF THE JUNCTION



# Immobilization Strategy



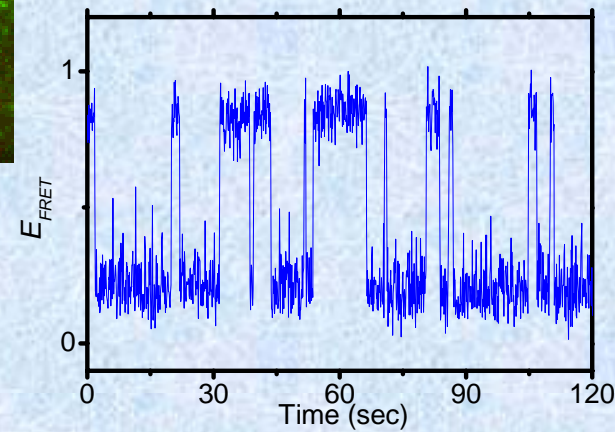
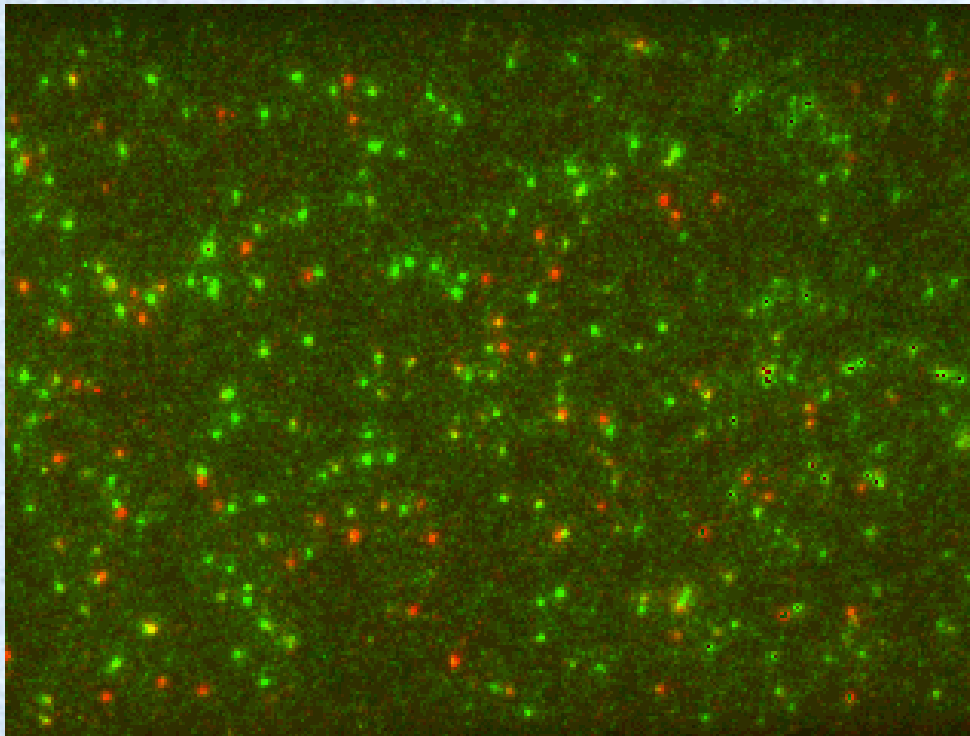
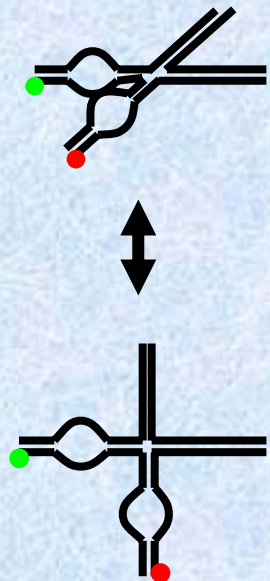
# Dual Channel TIR Imaging



- 100s of molecules
- ~30 ms time resolution
- Flow delivery system
- Dual channel imaging

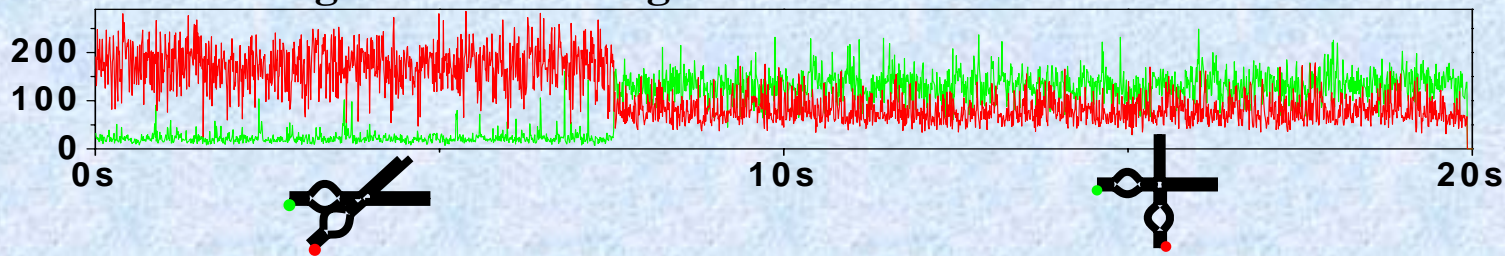
# Hairpin Ribozyme Movie

## Real Time (10 Frames/Sec)

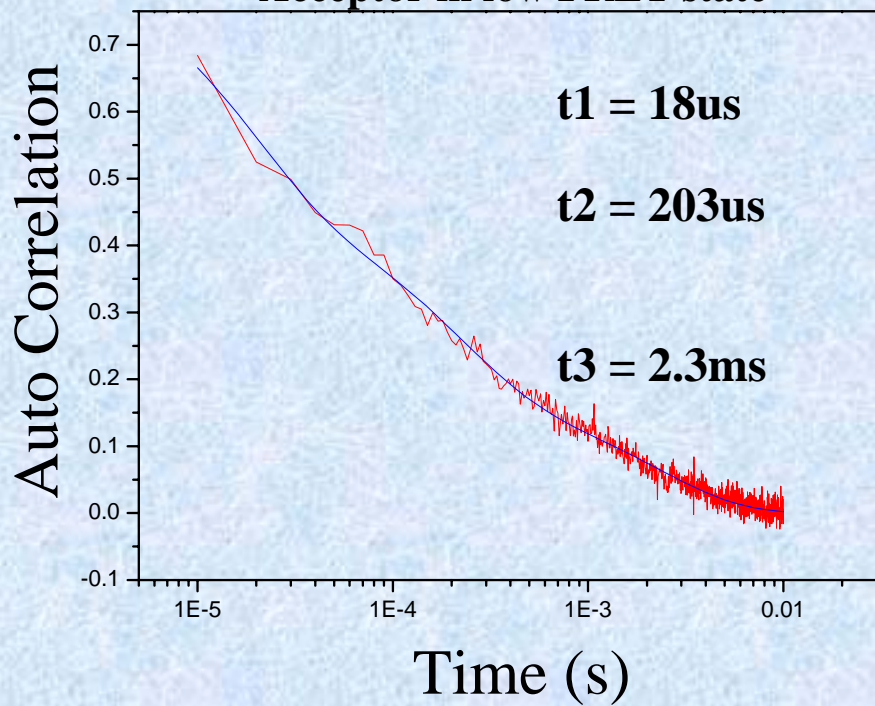


# Dynamics of Hairpin Ribozyme at Low FRET State

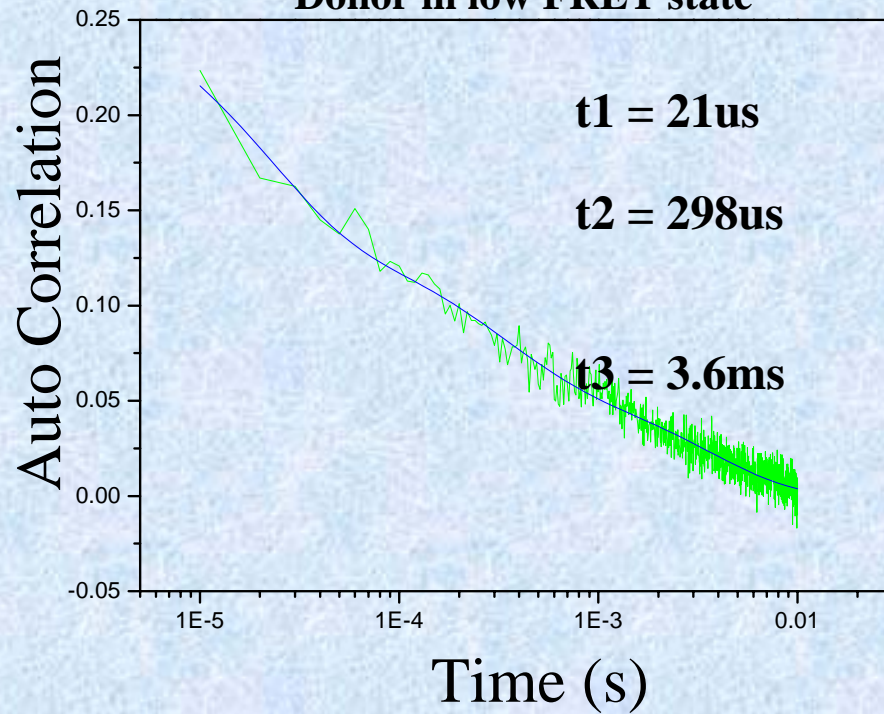
0.2mM Mg<sup>2+</sup> 10 $\mu$ s integration time rebinned to 8ms



Acceptor in low FRET state

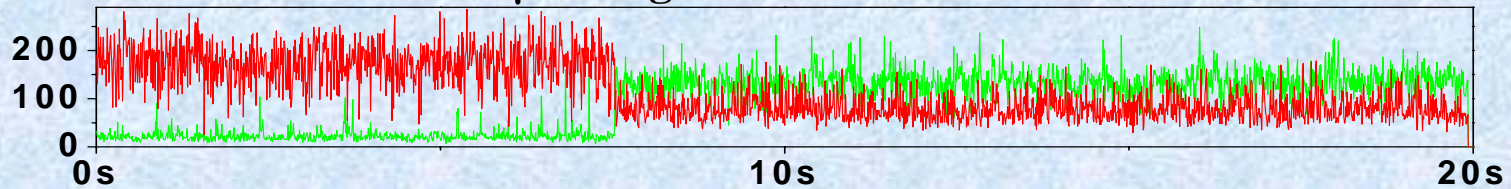


Donor in low FRET state

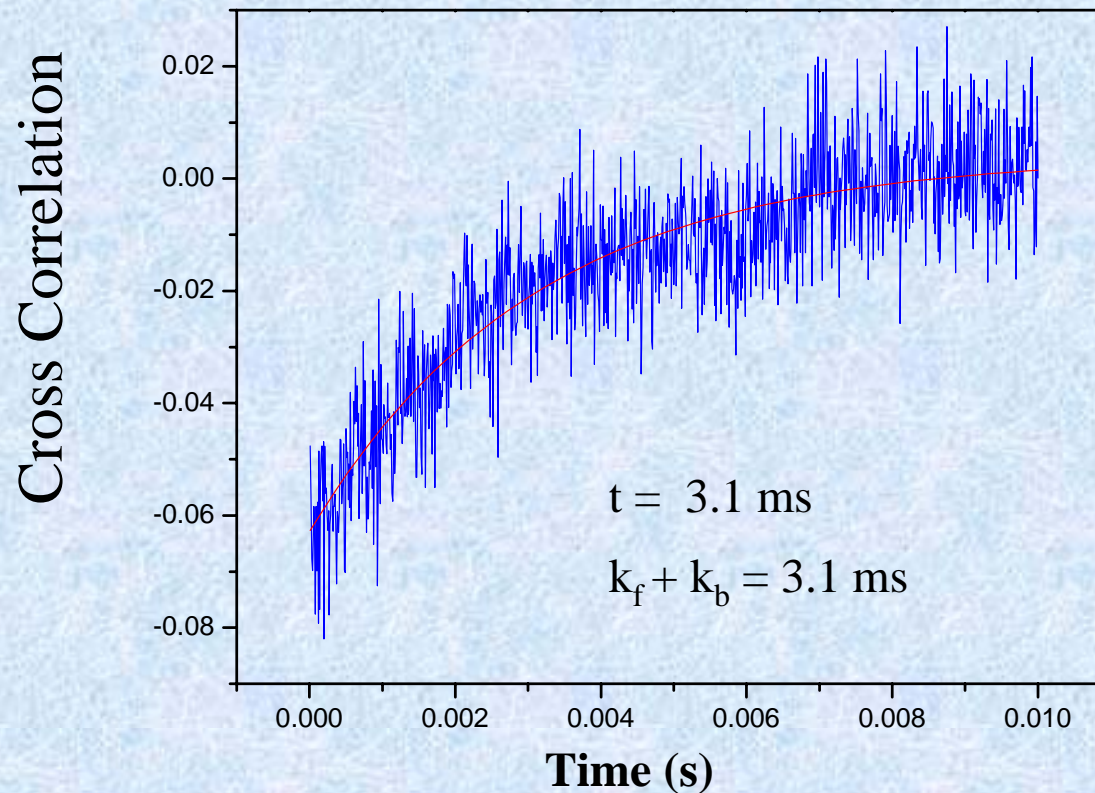


# Dynamics of Hairpin Ribozyme at Low FRET State

10 $\mu$ s integration time rebinned to 8ms



Cross-correlation of acceptor and donor signal from 10 s to 20 s using 10 $\mu$ s data



# FRET Efficiency Distribution in Low $Mg^{++}$

